



Full wwPDB EM Validation Report ⓘ

Oct 13, 2024 – 10:49 pm BST

PDB ID : 4V8M
EMDB ID : EMD-2239
Title : High-resolution cryo-electron microscopy structure of the Trypanosoma brucei ribosome
Authors : Hashem, Y.; des Georges, A.; Fu, J.; Buss, S.N.; Jossinet, F.; Jobe, A.; Zhang, Q.; Liao, H.Y.; Grassucci, R.A.; Bajaj, C.; Westhof, E.; Madison-Antenucci, S.; Frank, J.
Deposited on : 2012-12-09
Resolution : 5.57 Å(reported)
Based on initial models : 3U5G, 2XZM, 4A1B, 4A1A, 4A19, 3U5B, 3U5I, 3U5F, 3IZ6, 4A1E, 3IZR, 2XZN, 3U5C, 3U5E, 4A1C, 3IZ7, 3IZ9, 4A17, 4A18, 4A1D, 3U5D, 3U5H

This is a Full wwPDB EM Validation Report for a publicly released PDB entry.

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

The following versions of software and data (see [references ⓘ](#)) were used in the production of this report:

EMDB validation analysis : 0.0.1.dev113
MolProbity : 4.02b-467
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.39

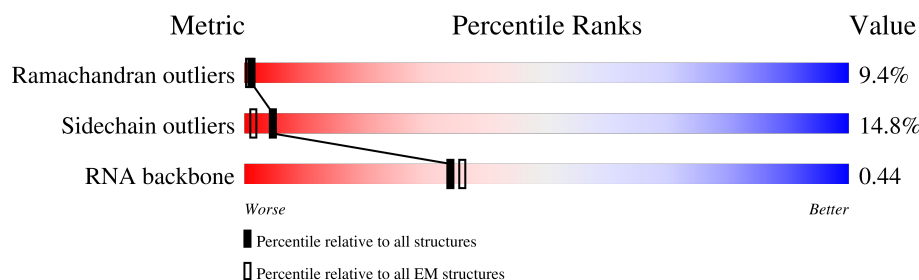
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 5.57 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.



Metric	Whole archive (#Entries)	EM structures (#Entries)
Ramachandran outliers	207382	16835
Sidechain outliers	206894	16415
RNA backbone	6643	2191

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A0	256	<div> <div>30%</div> <div>55%</div> <div>22%</div> <div>7%</div> <div>14%</div> </div>
2	A1	273	<div> <div>24%</div> <div>58%</div> <div>26%</div> <div>5%</div> <div>9%</div> </div>
3	A2	190	<div> <div>62%</div> <div>69%</div> <div>18%</div> <div>10%</div> <div>..</div> </div>
4	A3	250	<div> <div>25%</div> <div>64%</div> <div>26%</div> <div>8%</div> <div>.</div> </div>
5	A4	202	<div> <div>17%</div> <div>52%</div> <div>29%</div> <div>11%</div> <div>5%</div> </div>
6	A5	220	<div> <div>35%</div> <div>60%</div> <div>20%</div> <div>6%</div> <div>11%</div> </div>
7	A6	190	<div> <div>17%</div> <div>61%</div> <div>25%</div> <div>9%</div> <div>..</div> </div>
8	A7	318	<div> <div>68%</div> <div>74%</div> <div>21%</div> <div>..</div> </div>

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Mol	Chain	Length	Quality of chain
9	A8	57	
10	A9	153	
11	AC	277	
12	AD	172	
13	AE	174	
14	AF	144	
15	AG	151	
16	AH	144	
17	AI	152	
18	AJ	130	
19	AK	149	
20	AL	142	
21	AM	153	
22	AO	167	
23	AP	266	
24	AQ	117	
25	AR	194	
26	AS	143	
27	AT	137	
28	AU	113	
29	AV	111	
30	AW	86	
31	AX	214	
32	AY	66	
33	AZ	103	

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Mol	Chain	Length	Quality of chain
34	BA	1847	
35	BB	1465	
36	BC	169	
37	BD	119	
38	BE	210	
39	BF	73	
40	BG	182	
41	BH	135	
42	BI	193	
43	BJ	214	
44	BK	213	
45	BL	194	
46	BM	164	
47	BN	218	
48	BO	222	
49	BP	189	
50	BQ	221	
51	BR	166	
52	BS	179	
53	BT	260	
54	BU	159	
55	BV	130	
56	BW	139	
57	BX	164	
58	BY	125	

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Mol	Chain	Length	Quality of chain
59	BZ	143	
60	Ba	133	
61	Bb	145	
62	Bc	146	
63	Bd	71	
64	Be	260	
65	Bf	429	
66	Bg	105	
67	Bh	188	
68	Bi	132	
69	Bj	170	
70	Bk	127	
71	Bl	149	
72	Bm	109	
73	Bn	84	
74	Bo	93	
75	Bp	82	
76	Bq	51	
77	Br	374	
78	Bs	128	
79	Bt	106	
80	Bu	308	
81	Bv	192	
82	Bw	257	
83	Bx	276	

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Mol	Chain	Length	Quality of chain
84	By	189	<div><div></div><div>9%</div><div>56%</div><div>28%</div><div>13%</div><div></div></div>
85	AA	2251	<div><div></div><div>17%</div><div>19%</div><div>51%</div><div>28%</div><div></div></div>
86	AB	73	<div><div></div><div>45%</div><div>21%</div><div>51%</div><div>25%</div><div></div></div>

2 Entry composition

There are 86 unique types of molecules in this entry. The entry contains 232955 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, the AltConf column contains the number of residues with at least one atom in alternate conformation and the Trace column contains the number of residues modelled with at most 2 atoms.

- Molecule 1 is a protein called 40S RIBOSOMAL PROTEIN S3A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A0	219	Total	C	N	O	S	0	1
			1782	1124	337	313	8		

- Molecule 2 is a protein called 40S RIBOSOMAL PROTEIN S4, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	A1	248	Total	C	N	O	S	0	1
			1940	1232	360	339	9		

- Molecule 3 is a protein called 40S RIBOSOMAL PROTEIN S5, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	A2	187	Total	C	N	O	S	0	0
			1484	928	286	265	5		

- Molecule 4 is a protein called 40S RIBOSOMAL PROTEIN S6.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	A3	250	Total	C	N	O	S	0	0
			2003	1243	415	341	4		

- Molecule 5 is a protein called RIBOSOMAL PROTEIN S7, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	A4	192	Total	C	N	O	S	0	1
			1592	1014	310	263	5		

- Molecule 6 is a protein called 40S RIBOSOMAL PROTEIN S8.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	A5	195	Total	C	N	O	S	0	1
			1551	975	315	259	2		

- Molecule 7 is a protein called 40S RIBOSOMAL PROTEIN S9, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	A6	187	Total	C	N	O	S	0	1
			1518	951	307	253	7		

- Molecule 8 is a protein called GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT-LIKE PROTEIN.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	A7	315	Total	C	N	O	S	0	1
			2412	1508	429	462	13		

- Molecule 9 is a protein called RIBOSOMAL PROTEIN S29, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	A8	42	Total	C	N	O	S	0	0
			334	204	69	57	4		

- Molecule 10 is a protein called UBIQUITIN/RIBOSOMAL PROTEIN S27A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	A9	66	Total	C	N	O	S	0	1
			530	330	102	91	7		

- Molecule 11 is a protein called 40S RIBOSOMAL PROTEIN SA, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	AC	204	Total	C	N	O	S	0	1
			1620	1034	293	282	11		

- Molecule 12 is a protein called 40S RIBOSOMAL PROTEIN S10, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	AD	104	Total	C	N	O	S	0	1
			853	553	148	147	5		

- Molecule 13 is a protein called 40S RIBOSOMAL PROTEINS S11, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	AE	160	Total	C	N	O	S	0	0
			1300	812	262	220	6		

- Molecule 14 is a protein called 40S RIBOSOMAL PROTEIN S12.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	AF	121	Total	C	N	O	S	0	0
			940	578	169	184	9		

- Molecule 15 is a protein called 40S RIBOSOMAL PROTEIN S13, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	AG	141	Total	C	N	O	S	0	0
			1148	724	227	190	7		

- Molecule 16 is a protein called 40S RIBOSOMAL PROTEIN S14.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	AH	126	Total	C	N	O	S	0	1
			922	572	167	174	9		

- Molecule 17 is a protein called 40S RIBOSOMAL PROTEIN S15, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	AI	134	Total	C	N	O	S	0	1
			1074	679	211	181	3		

- Molecule 18 is a protein called 40S RIBOSOMAL PROTEIN S15A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	AJ	129	Total	C	N	O	S	0	0
			1018	645	191	174	8		

- Molecule 19 is a protein called 40S RIBOSOMAL PROTEIN S16, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	AK	148	Total	C	N	O	S	0	0
			1190	757	225	205	3		

- Molecule 20 is a protein called 40S RIBOSOMAL PROTEIN S17, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	AL	127	Total	C	N	O	S	0	1
			1021	641	198	177	5		

- Molecule 21 is a protein called 40S RIBOSOMAL PROTEIN S18, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	AM	153	Total	C	N	O	S	0	0
			1229	764	244	215	6		

- Molecule 22 is a protein called RIBOSOMAL PROTEIN S19, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	AO	149	Total	C	N	O	S	0	0
			1181	746	230	196	9		

- Molecule 23 is a protein called 40S RIBOSOMAL PROTEIN S2, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	AP	224	Total	C	N	O	S	0	1
			1731	1103	309	310	9		

- Molecule 24 is a protein called RIBOSOMAL PROTEIN S20, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	AQ	105	Total	C	N	O	S	0	1
			827	522	153	149	3		

- Molecule 25 is a protein called 40S RIBOSOMAL PROTEIN S21, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	AR	81	Total	C	N	O	S	0	1
			603	374	108	118	3		

- Molecule 26 is a protein called 40S RIBOSOMAL PROTEIN S23, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
26	AS	142	Total	C	N	O	S	0	0
			1116	706	219	189	2		

- Molecule 27 is a protein called 40S RIBOSOMAL PROTEIN S24.

Mol	Chain	Residues	Atoms					AltConf	Trace
27	AT	131	Total	C	N	O	S	0	0
			1050	666	206	174	4		

- Molecule 28 is a protein called 40S RIBOSOMAL PROTEIN S25, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
28	AU	86	Total	C	N	O	S	0	1
			673	427	127	114	5		

- Molecule 29 is a protein called RIBOSOMAL PROTEIN S26, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
29	AV	101	Total	C	N	O	S	0	1
			809	498	172	131	8		

- Molecule 30 is a protein called 40S RIBOSOMAL PROTEIN S27, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
30	AW	83	Total	C	N	O	S	0	1
			636	396	120	111	9		

- Molecule 31 is a protein called 40S RIBOSOMAL PROTEIN S3, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
31	AX	206	Total	C	N	O	S	0	1
			1628	1020	307	289	12		

- Molecule 32 is a protein called 40S RIBOSOMAL PROTEIN S30, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
32	AY	65	Total	C	N	O	S	0	0
			514	322	107	84	1		

- Molecule 33 is a protein called 40S RIBOSOMAL PROTEIN S33, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
33	AZ	68	Total	C	N	O	S	0	0
			526	315	107	100	4		

- Molecule 34 is a RNA chain called ALPHA CHAIN OF THE LARGE RIBOSOMAL SUB-UNIT 28S RRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
34	BA	1847	Total	C	N	O	P	0	0
			39395	17589	7008	12952	1846		

There are 16 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
BA	?	-	C	deletion	GB X14553
BA	?	-	U	deletion	GB X14553
BA	?	-	U	deletion	GB X14553
BA	?	-	C	deletion	GB X14553
BA	?	-	C	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	C	deletion	GB X14553
BA	?	-	C	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	U	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	?	-	G	deletion	GB X14553
BA	799	A	-	insertion	GB X14553

- Molecule 35 is a RNA chain called BETA CHAIN OF THE LARGE RIBOSOMAL SUBUNIT 28S RRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
35	BB	1465	Total	C	N	O	P	0	0
			31164	13918	5476	10306	1464		

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
BB	484	G	-	insertion	GB X14553
BB	485	U	-	insertion	GB X14553
BB	486	G	-	insertion	GB X14553
BB	487	A	-	insertion	GB X14553

- Molecule 36 is a RNA chain called 5.8S RRNA CHAIN OF THE LARGE RIBOSOMAL SUBUNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
36	BC	169	Total	C	N	O	P	0	0
			3584	1604	629	1183	168		

- Molecule 37 is a RNA chain called 5S RRNA CHAIN OF THE LARGE RIBOSOMAL SUBUNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
37	BD	119	Total	C	N	O	P	0	0
			2533	1131	449	835	118		

- Molecule 38 is a RNA chain called SHORT RRNA-I OF THE LARGE RIBOSOMAL SUB-UNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
38	BE	210	Total	C	N	O	P	0	0
			4441	1986	768	1478	209		

- Molecule 39 is a RNA chain called SHORT RRNA-II OF THE LARGE RIBOSOMAL SUB-UNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	BF	73	Total	C	N	O	P	0	0
			1521	682	247	520	72		

- Molecule 40 is a RNA chain called SHORT RRNA-III OF THE LARGE RIBOSOMAL SUBUNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	BG	182	Total	C	N	O	P	0	0
			3896	1737	706	1272	181		

- Molecule 41 is a RNA chain called SHORT RRNA-IV OF THE LARGE RIBOSOMAL SUBUNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
41	BH	135	Total	C	N	O	P	0	0
			2867	1280	502	951	134		

- Molecule 42 is a protein called 60S RIBOSOMAL PROTEIN L18.

Mol	Chain	Residues	Atoms					AltConf	Trace
42	BI	192	Total	C	N	O	S	0	0
			1527	956	315	248	8		

- Molecule 43 is a protein called RIBOSOMAL PROTEIN.

Mol	Chain	Residues	Atoms					AltConf	Trace
43	BJ	214	Total	C	N	O	S	0	0
			1717	1086	308	307	16		

- Molecule 44 is a protein called 60S RIBOSOMAL PROTEIN L10, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
44	BK	212	Total	C	N	O	S	0	0
			1725	1086	338	287	14		

- Molecule 45 is a protein called 60S RIBOSOMAL PROTEIN L11, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
45	BL	170	Total	C	N	O	S	0	1
			1363	859	258	239	7		

- Molecule 46 is a protein called 60S RIBOSOMAL PROTEIN L12, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
46	BM	139	Total	C	N	O	S	0	1
			1022	642	187	188	5		

- Molecule 47 is a protein called 60S RIBOSOMAL PROTEIN L13.

Mol	Chain	Residues	Atoms					AltConf	Trace
47	BN	216	Total	C	N	O	S	0	1
			1762	1097	366	292	7		

- Molecule 48 is a protein called 60S RIBOSOMAL PROTEIN L13A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
48	BO	201	Total	C	N	O	S	0	1
			1627	1035	323	262	7		

- Molecule 49 is a protein called PROBABLE 60S RIBOSOMAL PROTEIN L14.

Mol	Chain	Residues	Atoms					AltConf	Trace
49	BP	184	Total	C	N	O	S	0	1
			1484	934	299	247	4		

- Molecule 50 is a protein called RIBOSOMAL PROTEIN L15.

Mol	Chain	Residues	Atoms					AltConf	Trace
50	BQ	203	Total	C	N	O	S	0	0
			1716	1077	370	264	5		

- Molecule 51 is a protein called 60S RIBOSOMAL PROTEIN L17, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
51	BR	155	Total	C	N	O	S	0	1
			1245	782	247	208	8		

- Molecule 52 is a protein called 60S RIBOSOMAL PROTEIN L18A.

Mol	Chain	Residues	Atoms					AltConf	Trace
52	BS	179	Total	C	N	O	S	0	0
			1473	931	290	244	8		

- Molecule 53 is a protein called 60S RIBOSOMAL PROTEIN L19, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
53	BT	200	Total	C	N	O	S	0	1
			1672	1025	366	273	8		

- Molecule 54 is a protein called 60S RIBOSOMAL PROTEIN L21E, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
54	BU	158	Total	C	N	O	S	0	0
			1260	802	246	206	6		

- Molecule 55 is a protein called 60S RIBOSOMAL PROTEIN L22, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
55	BV	104	Total	C	N	O	S	0	1
			863	558	152	150	3		

- Molecule 56 is a protein called 60S RIBOSOMAL PROTEIN L23, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
56	BW	138	Total	C	N	O	S	0	0
			1042	659	198	180	5		

- Molecule 57 is a protein called 60S RIBOSOMAL PROTEIN L23A.

Mol	Chain	Residues	Atoms					AltConf	Trace
57	BX	121	Total	C	N	O	S	0	0
			990	629	186	173	2		

- Molecule 58 is a protein called 60S RIBOSOMAL PROTEIN L24, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
58	BY	100	Total	C	N	O	S	0	0
			836	530	171	130	5		

- Molecule 59 is a protein called 60S RIBOSOMAL PROTEIN L26, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
59	BZ	125	Total	C	N	O	S	0	1
			1008	623	213	167	5		

- Molecule 60 is a protein called 60S RIBOSOMAL PROTEIN L27, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
60	Ba	132	Total	C	N	O	S	0	0
			1091	691	222	175	3		

- Molecule 61 is a protein called 60S RIBOSOMAL PROTEIN L27A.

Mol	Chain	Residues	Atoms					AltConf	Trace
61	Bb	144	Total	C	N	O	S	0	0
			1137	717	228	186	6		

- Molecule 62 is a protein called 60S RIBOSOMAL PROTEIN L28, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
62	Bc	141	Total	C	N	O	S	0	1
			1129	704	226	191	8		

- Molecule 63 is a protein called 60S RIBOSOMAL PROTEIN L29, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
63	Bd	70	Total	C	N	O	S	0	0
			571	349	128	93	1		

- Molecule 64 is a protein called 60S RIBOSOMAL PROTEIN L2, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
64	Be	186	Total	C	N	O	S	0	1
			1390	859	284	237	10		

- Molecule 65 is a protein called RIBOSOMAL PROTEIN L3, MITOCHONDRIAL, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
65	Bf	414	Total	C	N	O	S	0	1
			3317	2084	661	559	13		

- Molecule 66 is a protein called 60S RIBOSOMAL PROTEIN L30.

Mol	Chain	Residues	Atoms					AltConf	Trace
66	Bg	96	Total	C	N	O	S	0	0
			735	457	132	141	5		

- Molecule 67 is a protein called 60S RIBOSOMAL SUBUNIT PROTEIN L31, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
67	Bh	188	Total	C	N	O	S	0	0
			1526	961	309	250	6		

- Molecule 68 is a protein called 60S RIBOSOMAL PROTEIN L32, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
68	Bi	129	Total	C	N	O	S	0	1
			1054	664	215	171	4		

- Molecule 69 is a protein called 60S RIBOSOMAL PROTEIN L34, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
69	Bj	162	Total	C	N	O	S	0	1
			1293	801	286	202	4		

- Molecule 70 is a protein called 60S RIBOSOMAL PROTEIN L35, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
70	Bk	84	Total	C	N	O	S	0	0
			719	448	161	108	2		

- Molecule 71 is a protein called 60S RIBOSOMAL PROTEIN L35A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
71	Bl	116	Total	C	N	O	S	0	0
			936	589	189	155	3		

- Molecule 72 is a protein called RIBOSOMAL PROTEIN L36, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
72	Bm	107	Total	C	N	O	S	0	1
			849	530	178	139	2		

- Molecule 73 is a protein called RIBOSOMAL PROTEIN L37.

Mol	Chain	Residues	Atoms					AltConf	Trace
73	Bn	83	Total	C	N	O	S	0	0
			699	425	161	107	6		

- Molecule 74 is a protein called 60S RIBOSOMAL PROTEIN L37A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
74	Bo	92	Total	C	N	O	S	0	1
			715	442	148	119	6		

- Molecule 75 is a protein called 60S RIBOSOMAL PROTEIN L38, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
75	Bp	81	Total	C	N	O	S	0	0
			656	411	130	111	4		

- Molecule 76 is a protein called 60S RIBOSOMAL PROTEIN L39, PUTATIVE.

Mol	Chain	Residues	Atoms				AltConf	Trace
76	Bq	50	Total	C	N	O	0	0
			457	297	98	62		

- Molecule 77 is a protein called 60S RIBOSOMAL PROTEIN L4.

Mol	Chain	Residues	Atoms					AltConf	Trace
77	Br	368	Total	C	N	O	S	0	1
			2883	1802	576	488	17		

- Molecule 78 is a protein called UBIQUITIN-60S RIBOSOMAL PROTEIN L40.

Mol	Chain	Residues	Atoms					AltConf	Trace
78	Bs	52	Total	C	N	O	S	0	0
			427	265	88	67	7		

- Molecule 79 is a protein called 60S RIBOSOMAL PROTEIN L44.

Mol	Chain	Residues	Atoms					AltConf	Trace
79	Bt	105	Total	C	N	O	S	0	0
			866	547	170	144	5		

- Molecule 80 is a protein called 60S RIBOSOMAL PROTEIN L5, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
80	Bu	299	Total	C	N	O	S	0	1
			2354	1485	447	416	6		

- Molecule 81 is a protein called 60S RIBOSOMAL PROTEIN L6, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
81	Bv	158	Total	C	N	O	S	0	1
			1222	776	228	215	3		

- Molecule 82 is a protein called 60S RIBOSOMAL PROTEIN L7, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
82	Bw	257	Total	C	N	O	S	0	0
			2066	1316	394	345	11		

- Molecule 83 is a protein called 60S RIBOSOMAL PROTEIN L7A, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
83	Bx	240	Total	C	N	O	S	0	0
			1908	1198	375	329	6		

- Molecule 84 is a protein called 60S RIBOSOMAL PROTEIN L9, PUTATIVE.

Mol	Chain	Residues	Atoms					AltConf	Trace
84	By	189	Total	C	N	O	S	0	0
			1540	975	284	277	4		

- Molecule 85 is a RNA chain called 18S RRNA OF THE SMALL RIBOSOMAL SUBUNIT.

Mol	Chain	Residues	Atoms					AltConf	Trace
85	AA	2227	Total	C	N	O	P	0	0
			47370	21162	8354	15629	2225		

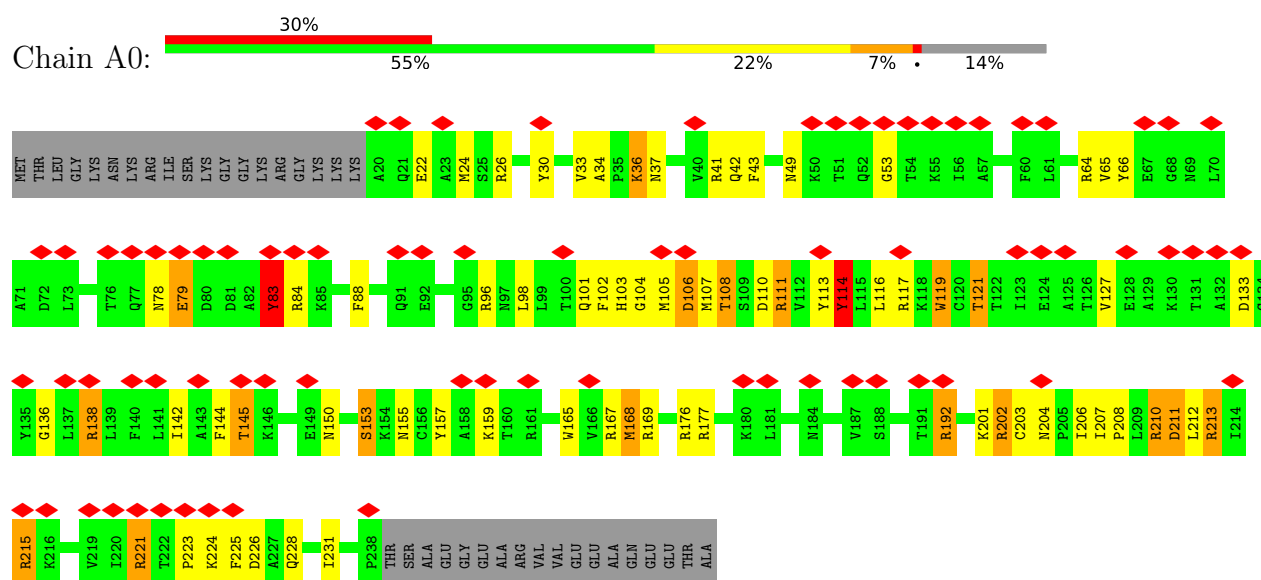
- Molecule 86 is a RNA chain called E-SITE TRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	P		
86	AB	73	1557	695	279	511	72	0	0

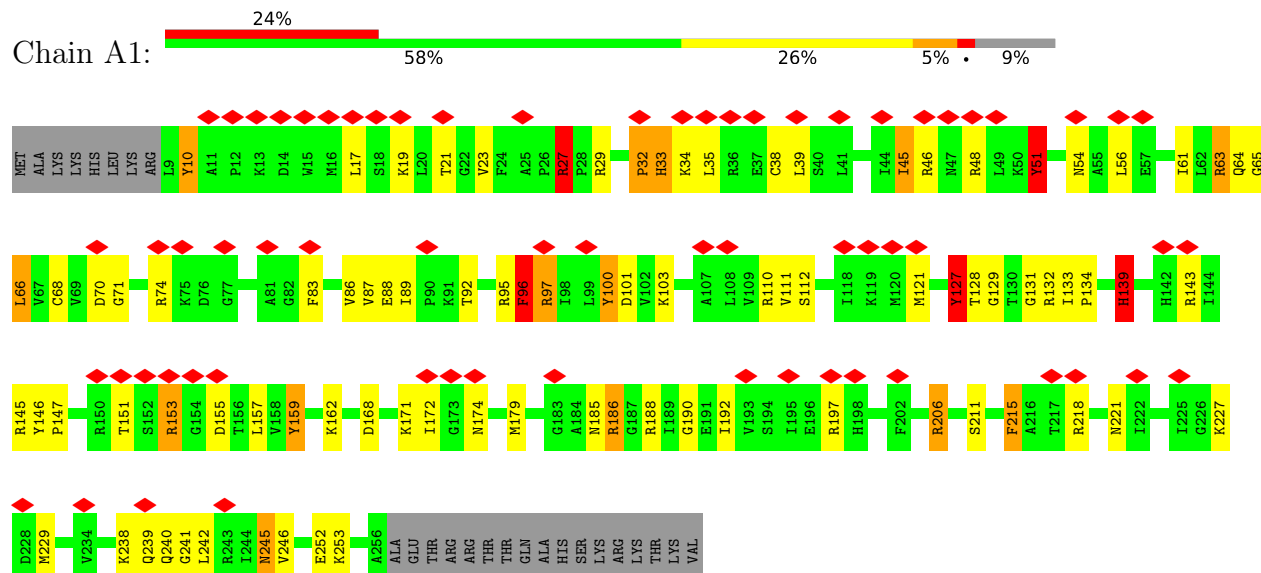
3 Residue-property plots

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion in map density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red diamond above a residue indicates a poor fit to the EM map for this residue (all-atom inclusion < 40%). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

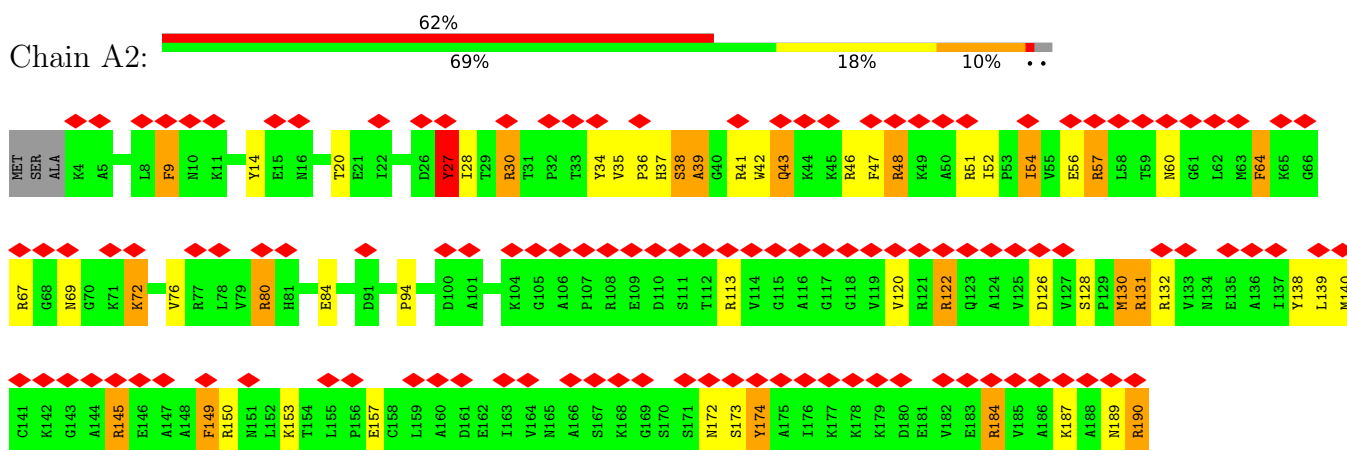
- Molecule 1: 40S RIBOSOMAL PROTEIN S3A, PUTATIVE



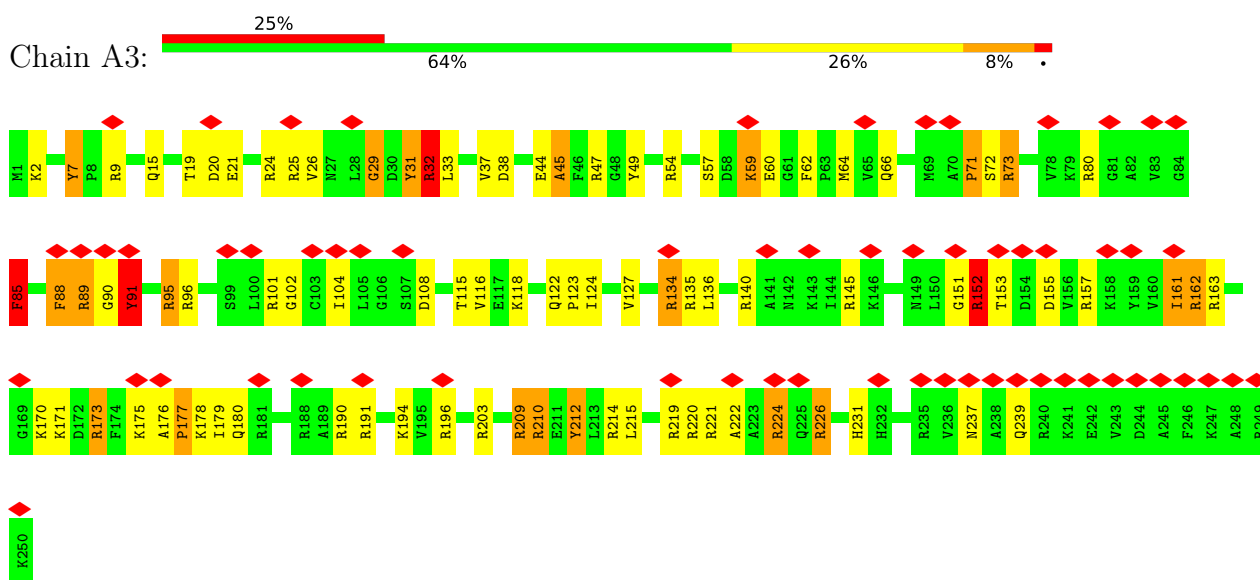
- Molecule 2: 40S RIBOSOMAL PROTEIN S4, PUTATIVE



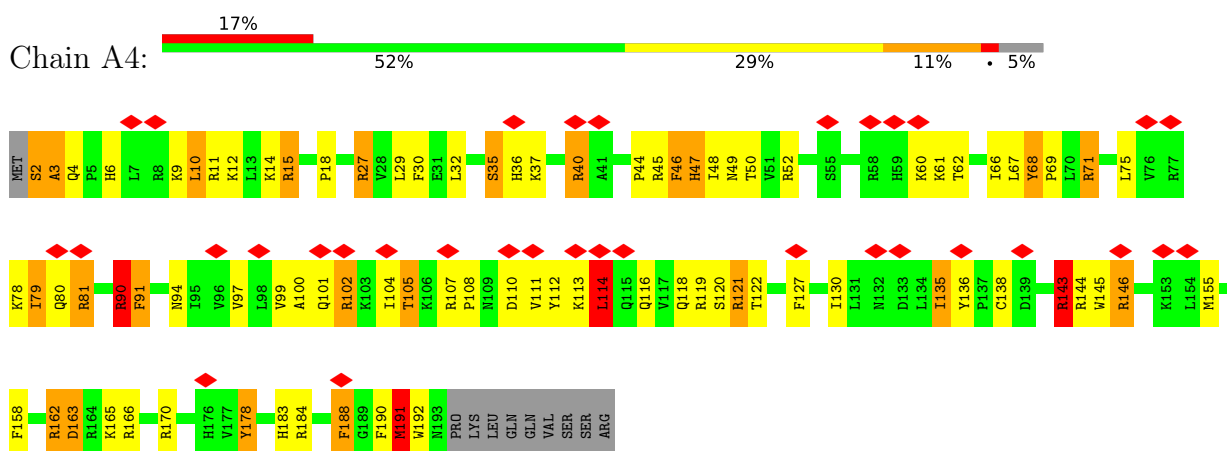
- Molecule 3: 40S RIBOSOMAL PROTEIN S5, PUTATIVE



• Molecule 4: 40S RIBOSOMAL PROTEIN S6

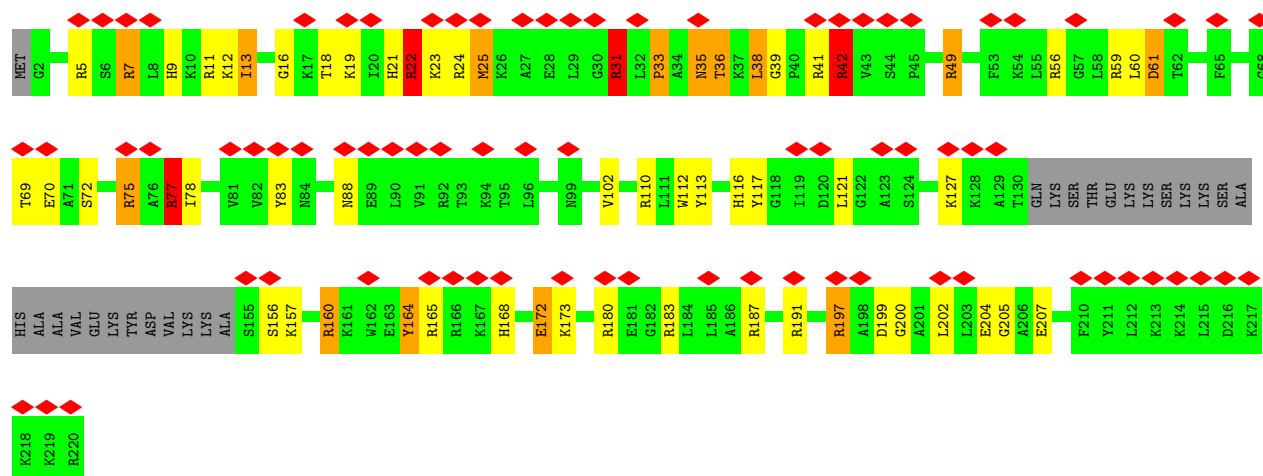


• Molecule 5: RIBOSOMAL PROTEIN S7, PUTATIVE

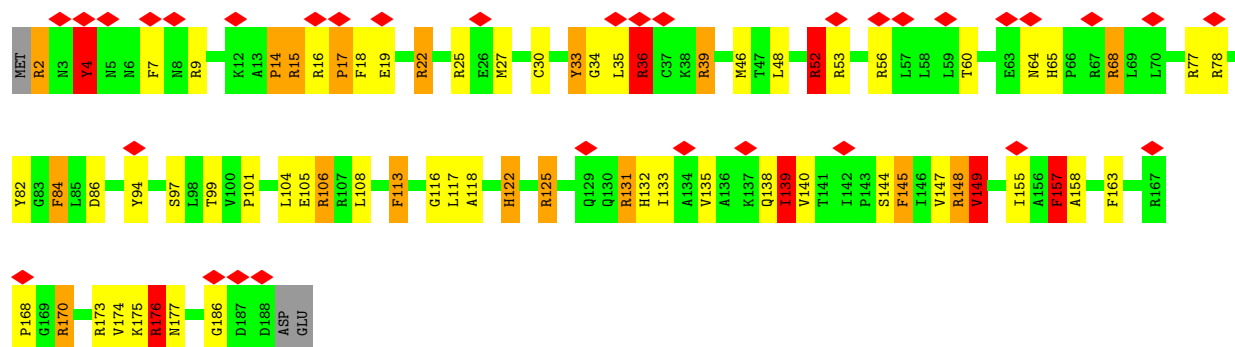


• Molecule 6: 40S RIBOSOMAL PROTEIN S8

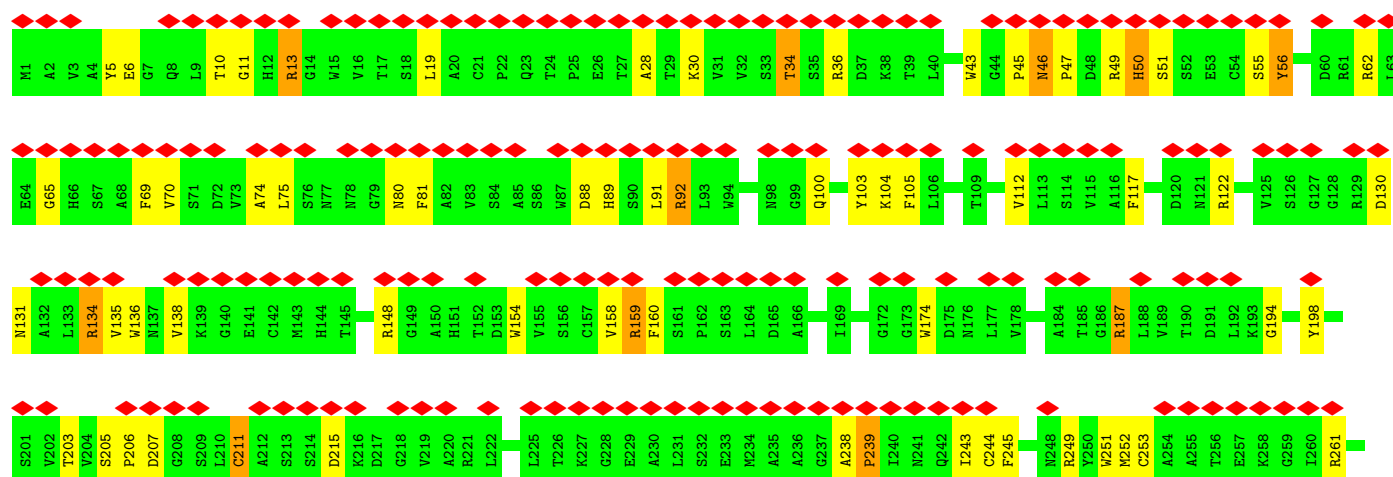
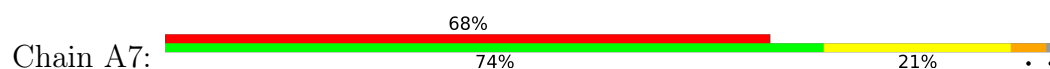


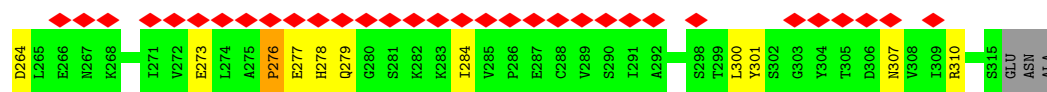


• Molecule 7: 40S RIBOSOMAL PROTEIN S9, PUTATIVE

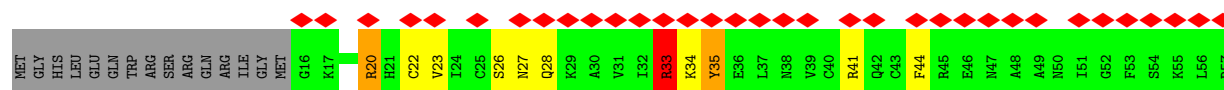


• Molecule 8: GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT-LIKE PROTEIN

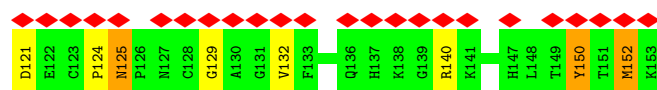
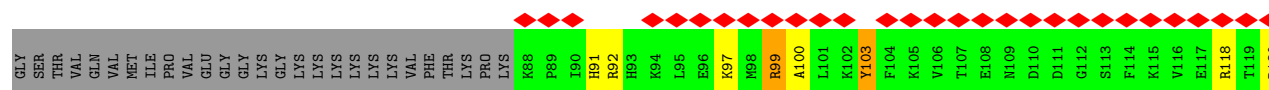
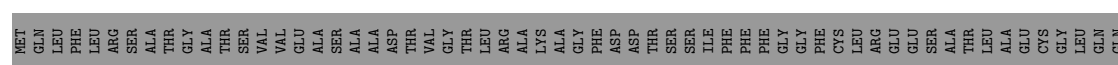
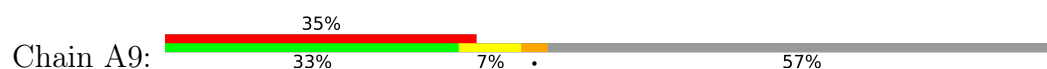




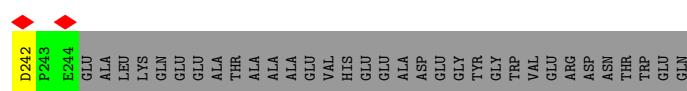
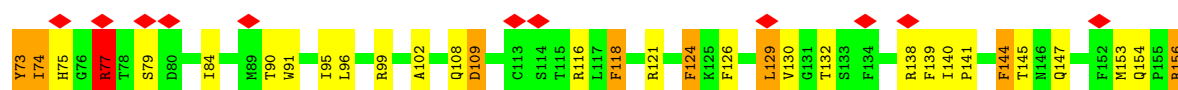
• Molecule 9: RIBOSOMAL PROTEIN S29, PUTATIVE



• Molecule 10: UBIQUITIN/RIBOSOMAL PROTEIN S27A, PUTATIVE

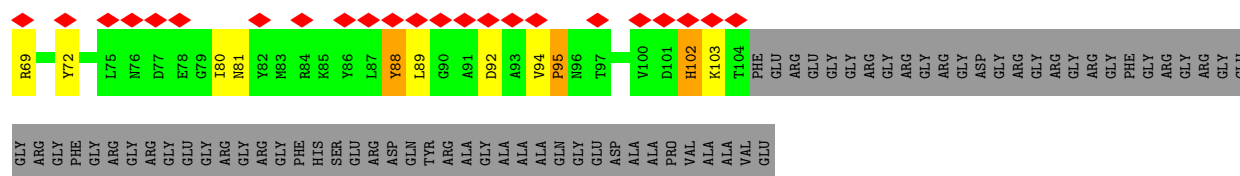


• Molecule 11: 40S RIBOSOMAL PROTEIN SA, PUTATIVE

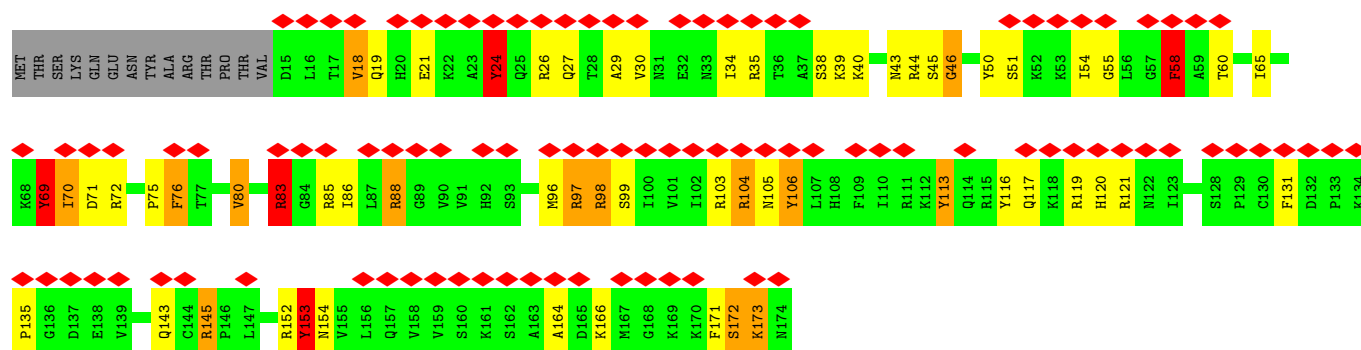


• Molecule 12: 40S RIBOSOMAL PROTEIN S10, PUTATIVE

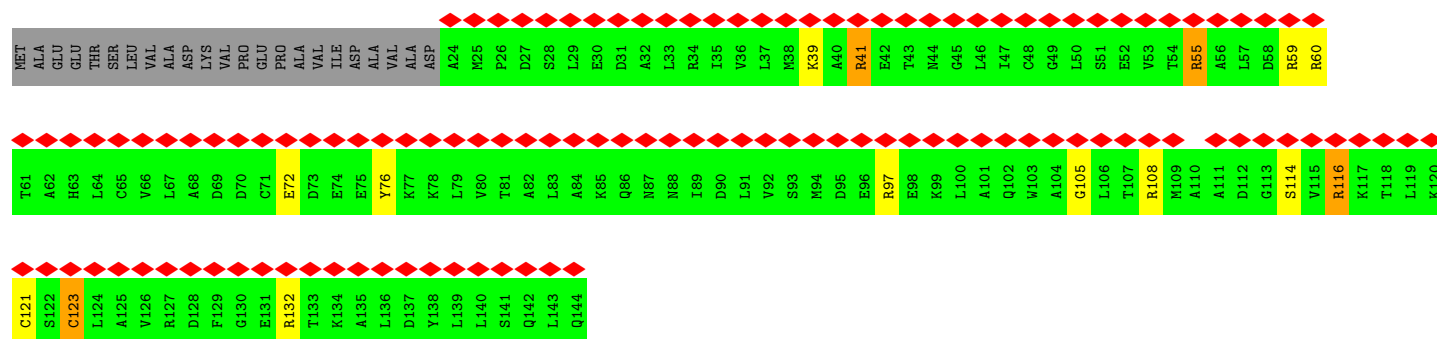
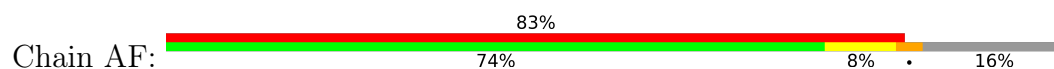




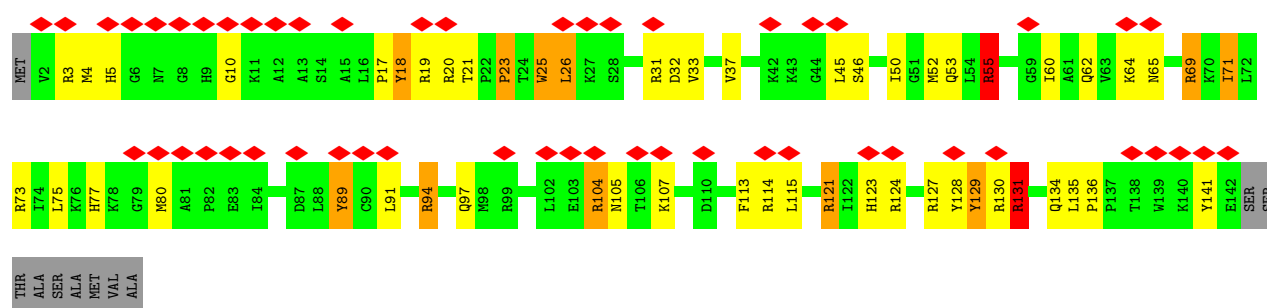
• Molecule 13: 40S RIBOSOMAL PROTEINS S11, PUTATIVE



• Molecule 14: 40S RIBOSOMAL PROTEIN S12

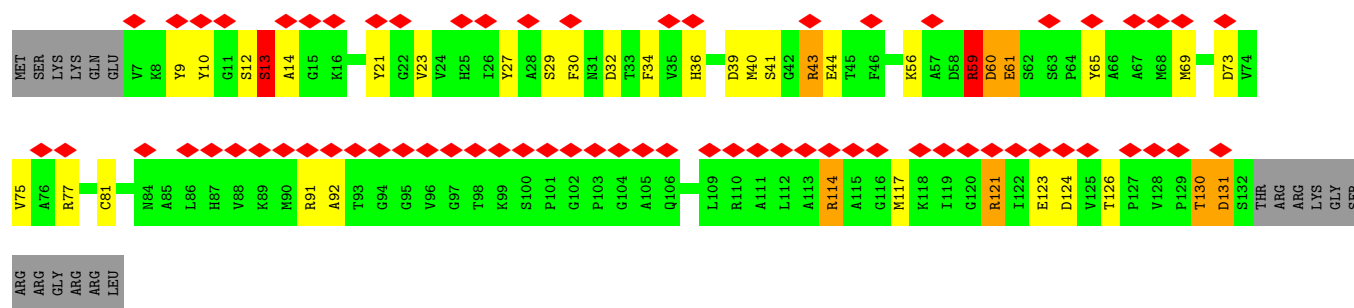


• Molecule 15: 40S RIBOSOMAL PROTEIN S13, PUTATIVE



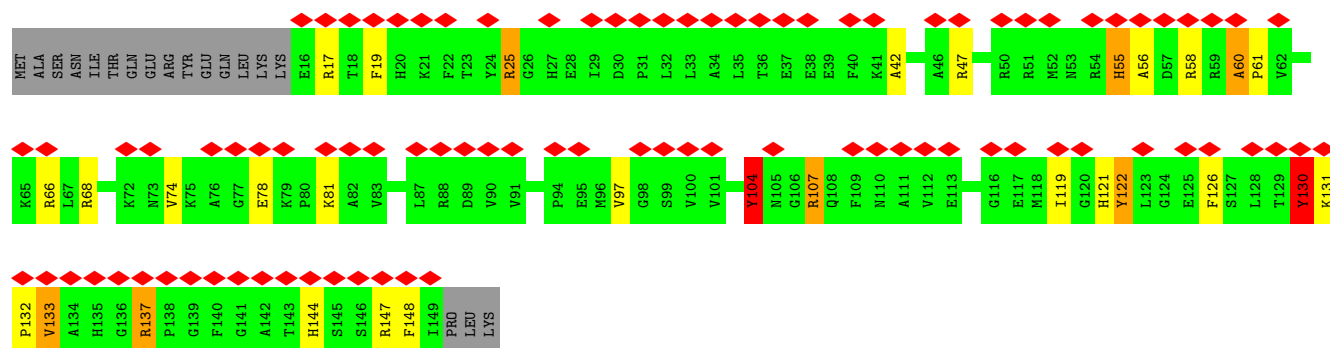
• Molecule 16: 40S RIBOSOMAL PROTEIN S14

Chain AH: 




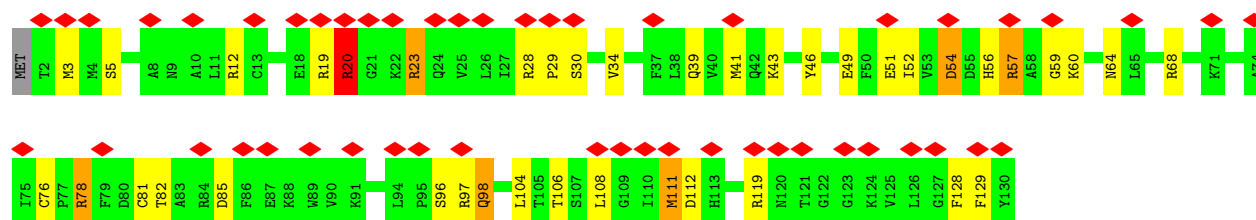
- Molecule 17: 40S RIBOSOMAL PROTEIN S15, PUTATIVE

Chain AI: 



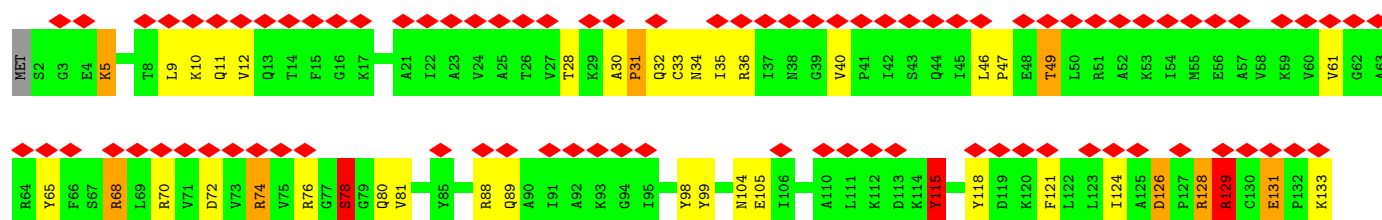
- Molecule 18: 40S RIBOSOMAL PROTEIN S15A, PUTATIVE

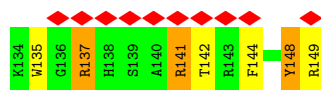
Chain AJ: 



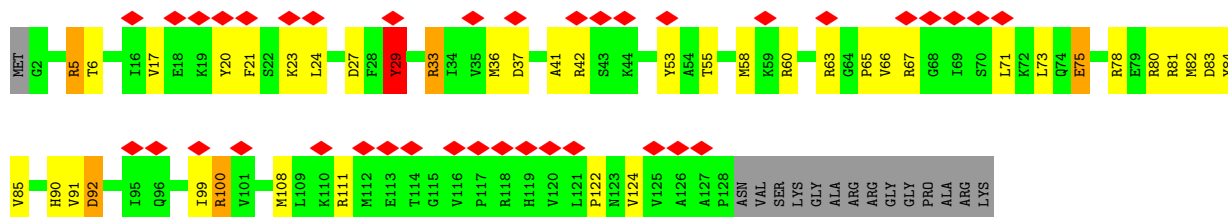
- Molecule 19: 40S RIBOSOMAL PROTEIN S16, PUTATIVE

Chain AK: 

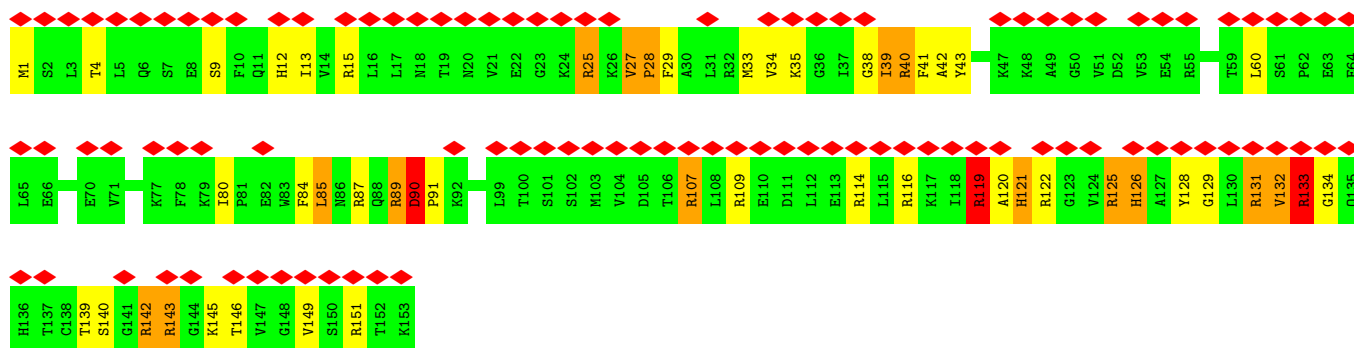




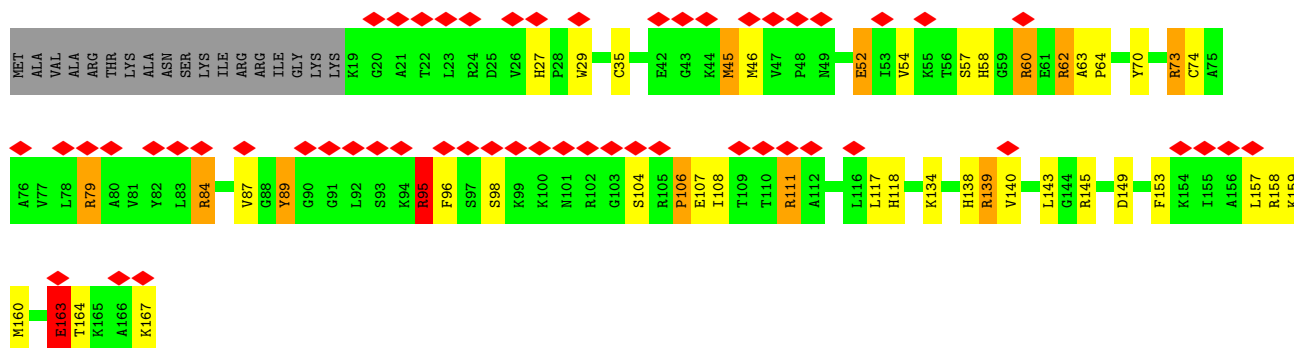
• Molecule 20: 40S RIBOSOMAL PROTEIN S17, PUTATIVE



• Molecule 21: 40S RIBOSOMAL PROTEIN S18, PUTATIVE

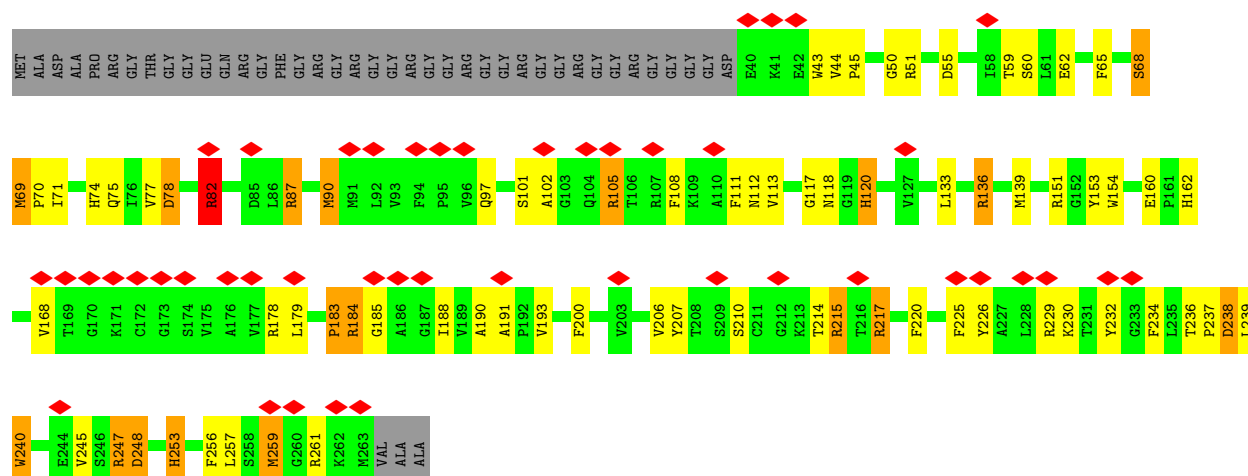


• Molecule 22: RIBOSOMAL PROTEIN S19, PUTATIVE

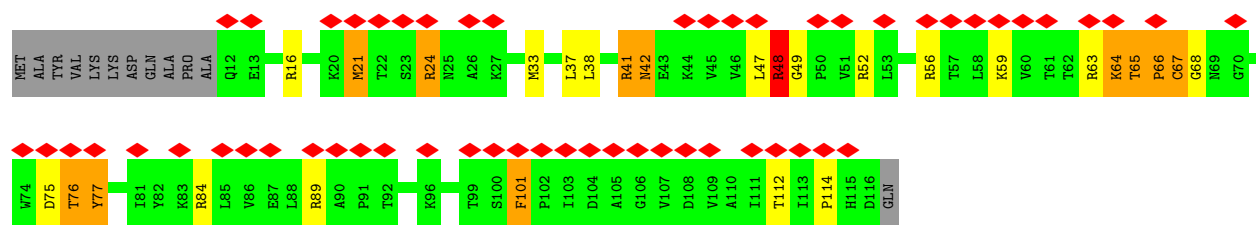


• Molecule 23: 40S RIBOSOMAL PROTEIN S2, PUTATIVE

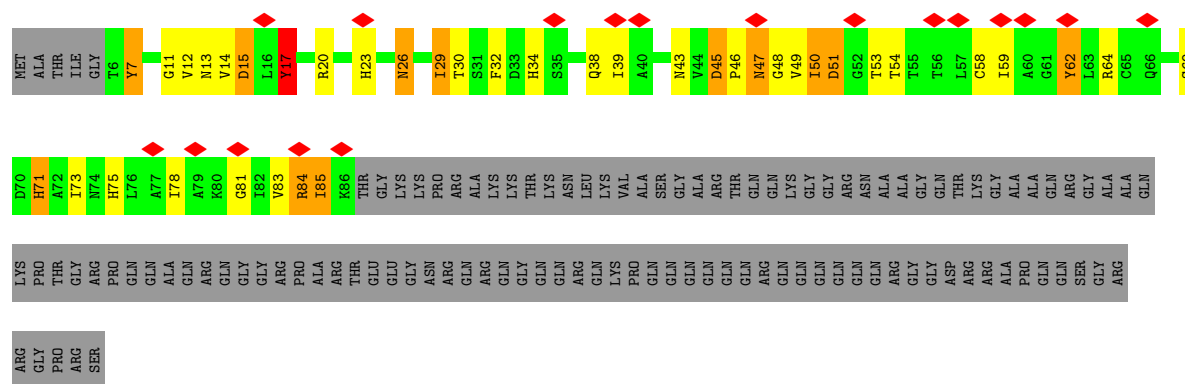
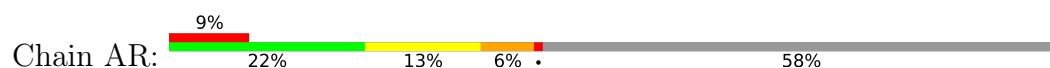




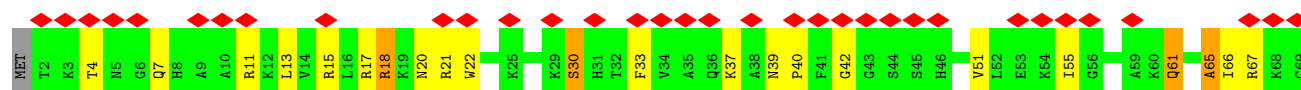
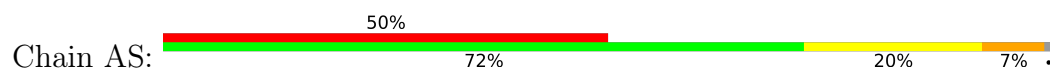
• Molecule 24: RIBOSOMAL PROTEIN S20, PUTATIVE

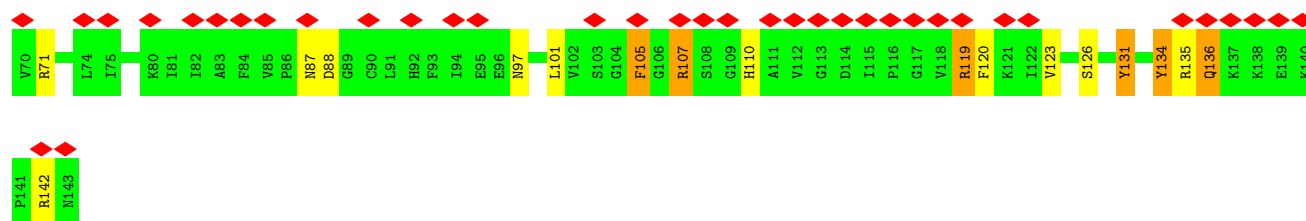


• Molecule 25: 40S RIBOSOMAL PROTEIN S21, PUTATIVE

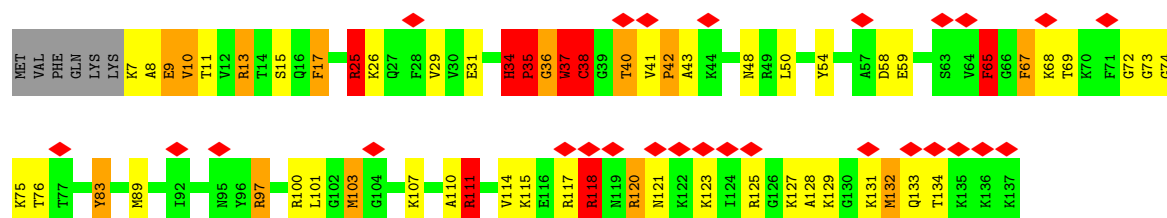


• Molecule 26: 40S RIBOSOMAL PROTEIN S23, PUTATIVE

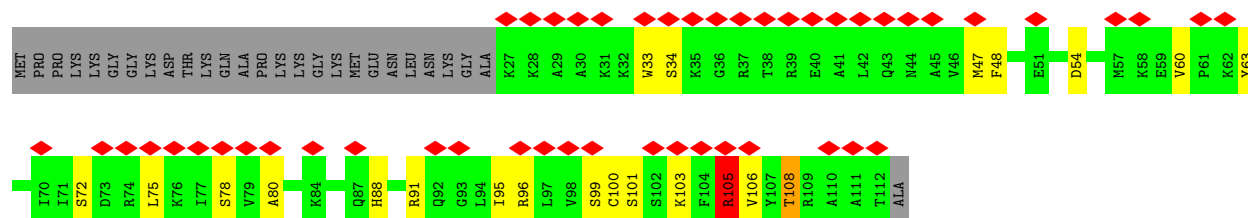
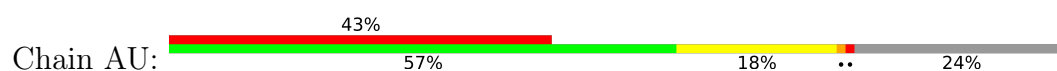




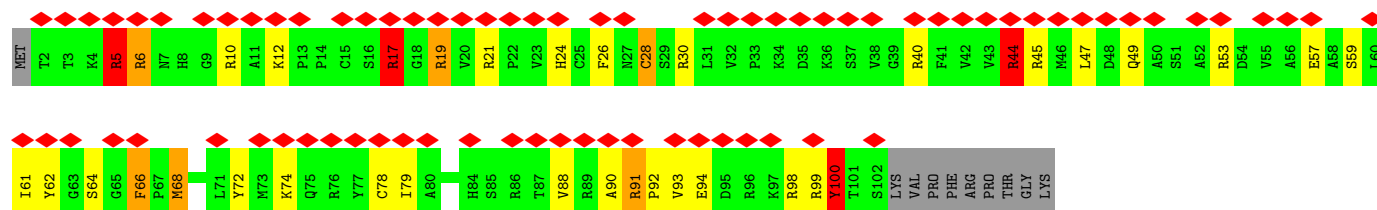
• Molecule 27: 40S RIBOSOMAL PROTEIN S24



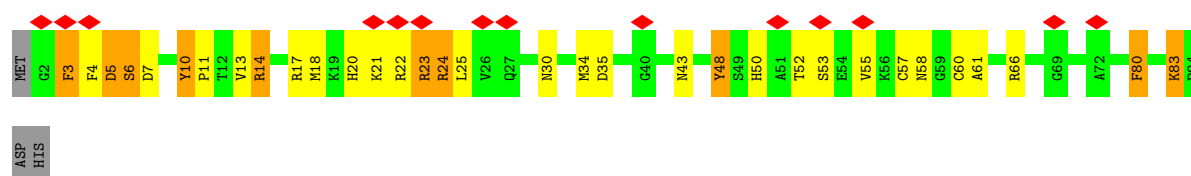
• Molecule 28: 40S RIBOSOMAL PROTEIN S25, PUTATIVE



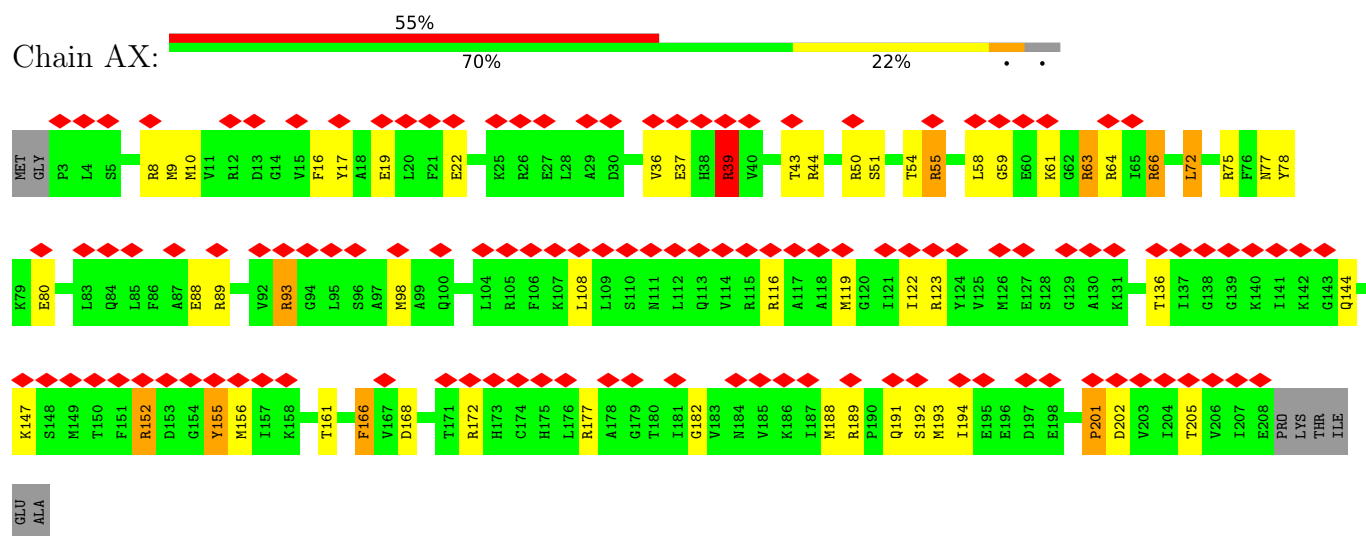
• Molecule 29: RIBOSOMAL PROTEIN S26, PUTATIVE



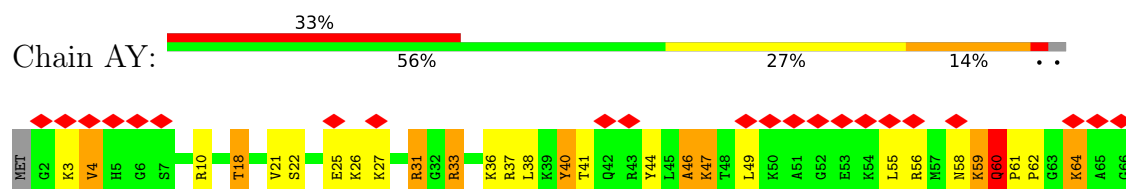
• Molecule 30: 40S RIBOSOMAL PROTEIN S27, PUTATIVE



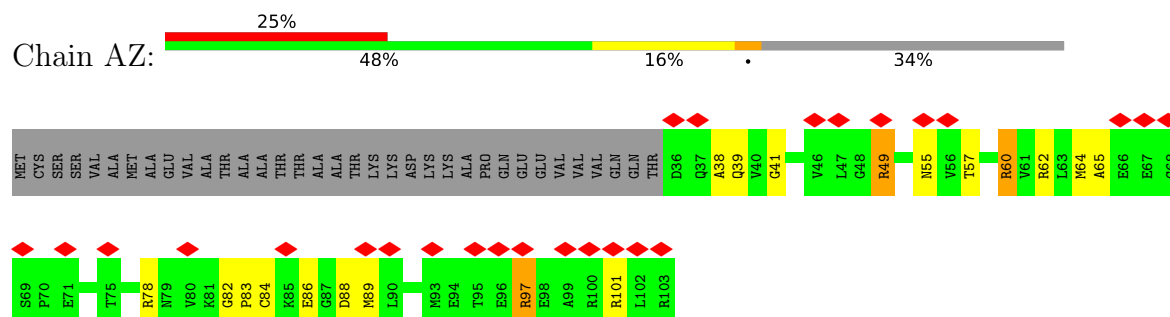
• Molecule 31: 40S RIBOSOMAL PROTEIN S3, PUTATIVE



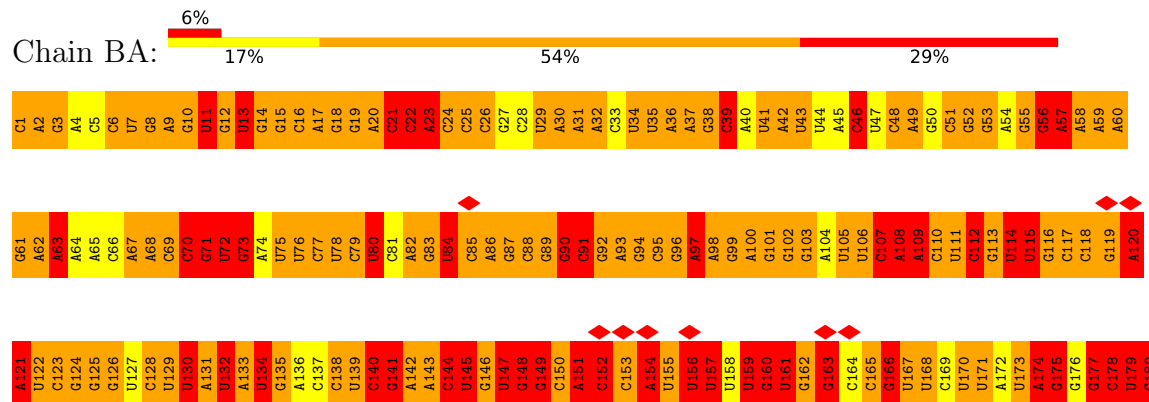
• Molecule 32: 40S RIBOSOMAL PROTEIN S30, PUTATIVE



• Molecule 33: 40S RIBOSOMAL PROTEIN S33, PUTATIVE

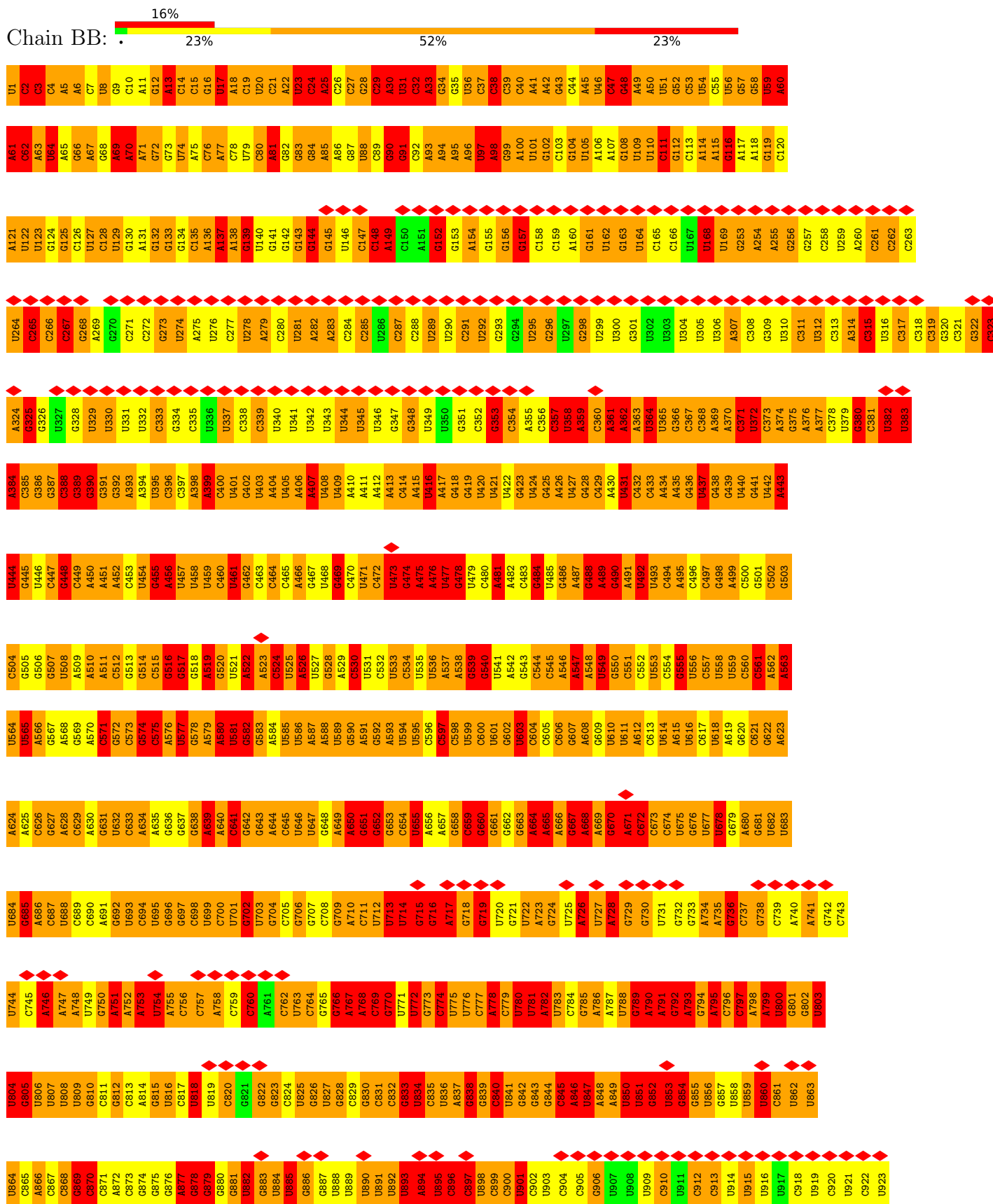


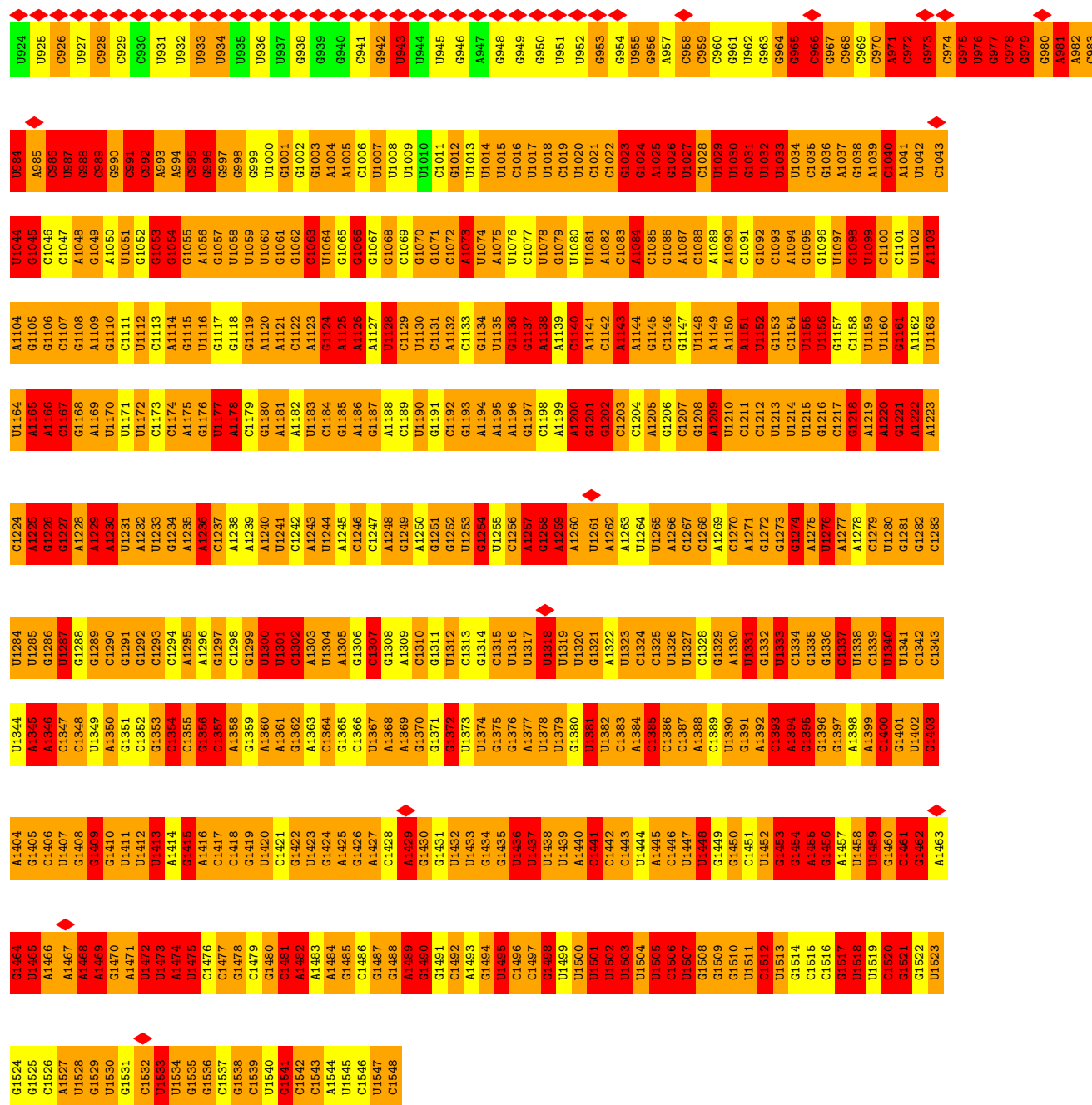
• Molecule 34: ALPHA CHAIN OF THE LARGE RIBOSOMAL SUBUNIT 28S RRNA



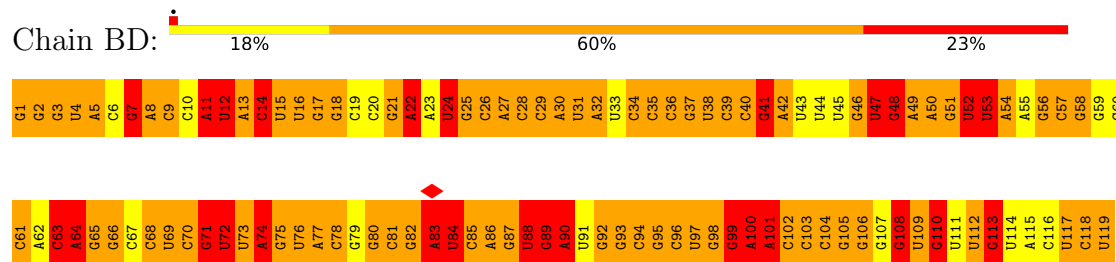
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C961	U962	G963	U964	A965	G966	C967	G968	A969	U970	C971	C972	U973	G974	A975	C976	G977	U978	G979	C980	A981	A982	A983	U984	C985	G986	C987	U988	C989	A990	C991	U992	A993	U994	A995	U996	U997	U998	G999	C1000	G1001	U1002	A1003	U1004	A1005	G1006	C1007	A1008	U1009	C1010	G1011	A1012	A1013	U1014	G1015	A1016	C1017	U1018	C1019	A1020				
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U781	C782	G783	C784	G785	U786	U787	C788	U789	C790	A791	A792	G793	C794	G795	C796	A797	C798	U799	C800	U801	C802	U803	C804	A805	U806	U807	U808	U809	A810	C811	A812	C813	C814	C815	C816	C817	C818	C819	C820	C821	U822	C823	C824	C825	C826	A827	A828	U829	U830	G838	U831	U832	C833	C834	U835	U836	U837	U838	U839	U840			
A721	A722	C723	A724	C725	C726	C727	C728	A729	C730	A731	A732	G733	C734	A735	C736	C737	C738	A739	A740	A741	C742	A743	C744	A745	C746	C747	C748	C749	C750	A751	A752	C753	C754	C755	A756	C757	C758	A759	C760	U761	A762	U763	C764	C765	A766	U767	C768	U769	C770	A771	C772	C773	A774	C775	U776	C777	U778	U779	U780				
C661	U662	C663	U664	C665	C666	U667	C668	U669	C670	C671	C672	U673	C674	C675	C676	U677	C678	U679	C680	C681	A682	C683	C684	C685	U686	C687	C688	C689	C690	A691	U692	C693	C694	A695	A696	A697	U698	C699	C700	U701	C702	U703	C704	C705	C706	C707	C708	C709	A710	C711	C712	C713	C714	U715	C716	U717	U718	C719	A720				
A601	C602	U603	C604	C605	C606	C607	C608	C609	A610	A611	U612	A613	A614	A615	C616	C617	C618	C619	C620	C621	C622	U623	C624	U625	C626	U627	U628	C629	U630	C631	C632	C633	U634	C635	C636	C637	U638	C639	U640	U641	U642	U643	C644	U645	C646	C647	C648	A649	C650	U651	C652	U653	C654	U655	U656	C657	C658	U659	C660				
C541	A542	C543	U544	U545	U546	C547	C548	C549	U550	U551	C552	A553	A554	C555	C556	C557	C558	C559	U560	C561	C562	A563	U564	U565	C566	U567	C568	U569	U570	C571	C572	C573	U574	U575	C576	C577	C578	U579	C580	U581	C582	C583	A584	C585	C586	U587	C588	A589	U590	C591	C592	C593	G594	U595	C596	C597	C598	U599	C600				
A481	C482	A483	A484	C485	G486	A487	C488	A489	A490	U491	C492	G493	A494	A495	G496	U497	A498	C499	C500	U501	C502	C503	A504	U505	U506	U507	C508	U509	U510	U511	U512	U513	U514	U515	U516	A517	C518	G519	C520	C521	C522	A523	C524	A525	C526	C527	C528	A529	A530	C531	U532	U533	C534	C535	U536	C536	C537	G538	C539	C540			
G421	C422	G423	U424	G425	A426	G427	C428	A429	U430	A431	A432	G433	U434	U435	U436	G437	A438	C439	A440	A441	C442	G443	A444	A445	U446	U447	U448	U449	A450	A451	A452	A453	G454	A455	G456	A457	G458	U459	G460	A461	C462	A463	U464	A465	G466	A467	A468	C469	C470	U471	G472	A473	A474	A475	U476	C477	G478	U479	G480				
C361	G362	G363	C364	A365	G366	C367	G368	A369	U370	U371	C372	G373	U374	C375	U376	G377	C378	C379	A380	A381	C382	G383	C384	U385	A386	A387	A388	U389	A390	U391	A392	G393	A394	G395	U396	A397	G398	G399	U399	A400	A401	U342	G402	G343	A403	C404	C405	G406	A407	U348	C409	A409	G410	C411	C412	A413	A414	C415	A416	A417	G418	U419	A420
U241	U242	C243	A244	U245	G246	U247	G248	A249	C250	U251	U252	U253	U254	C255	A256	G257	C258	C259	A260	A261	C262	G263	C264	A265	C266	U267	U268	C269	A270	C271	A272	U273	C274	C275	C276	A277	U278	U279	A280	C281	A282	C282	U283	U284	C285	A286	C287	U287	U288	A289	C290	C291	U292	U293	A294	C295	A296	A297	C298	C299	C300		
G181	U182	G183	C184	A185	G186	G187	C188	G189	U190	U191	G192	C193	G194	G195	A196	U197	U198	C199	C200	A201	A202	G203	U204	A205	C206	A207	A208	C209	G210	C211	A212	A213	A214	C215	C216	C217	G218	U219	U220	C221	C222	U223	C224	A225	A226	C227	U227	U228	C229	A230	U231	U232	U233	A234	C235	A236	A237	C238	C239	C240			



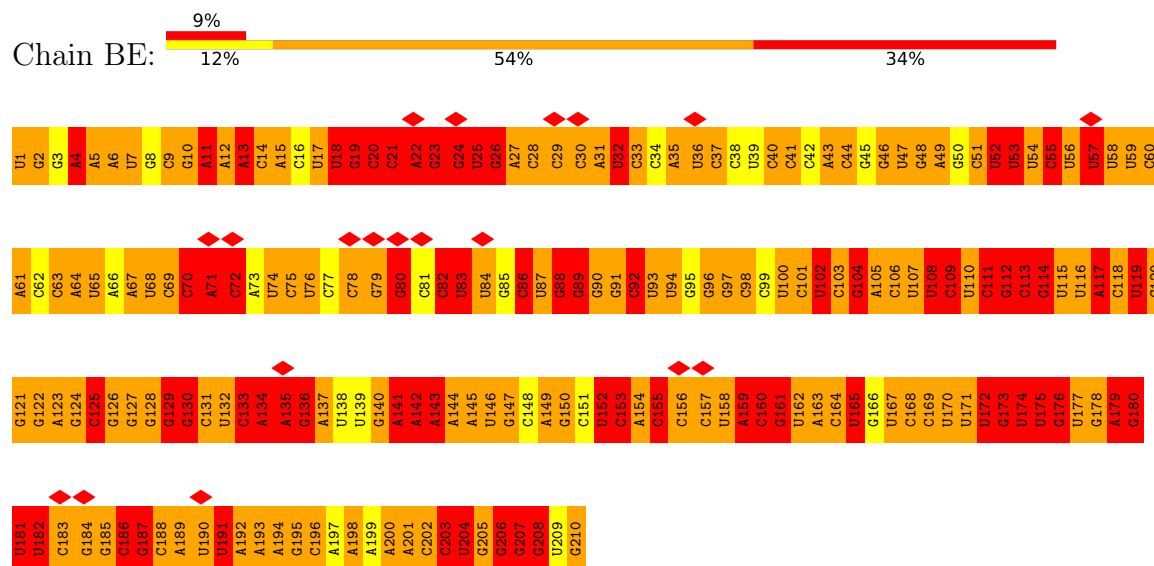




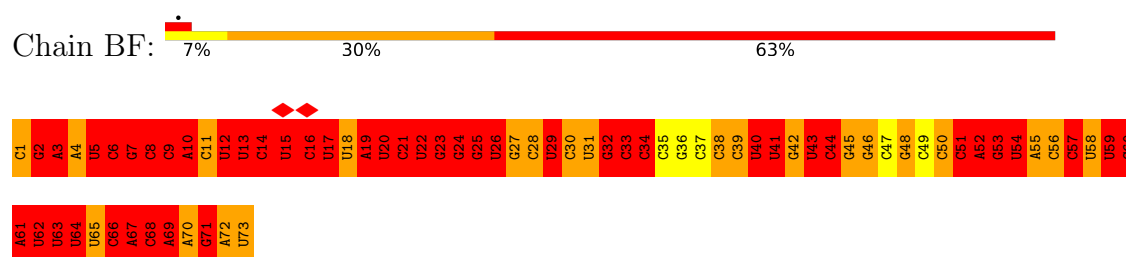
• Molecule 37: 5S RRNA CHAIN OF THE LARGE RIBOSOMAL SUBUNIT



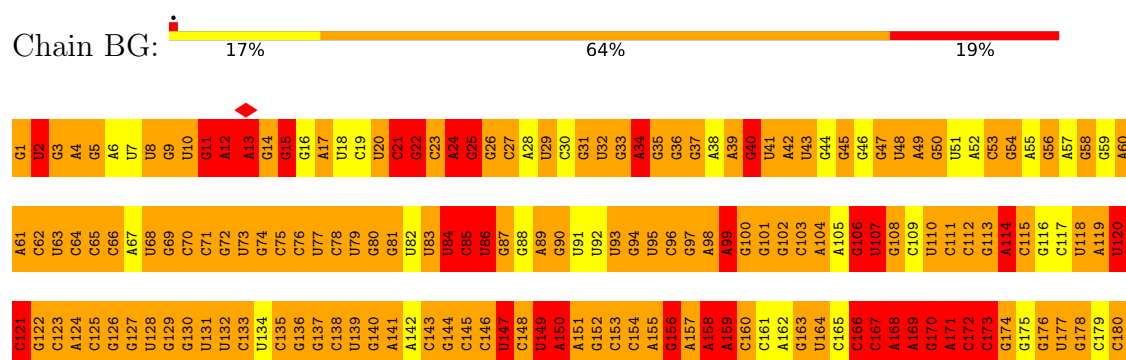
• Molecule 38: SHORT RRNA-I OF THE LARGE RIBOSOMAL SUBUNIT



• Molecule 39: SHORT RRNA-II OF THE LARGE RIBOSOMAL SUBUNIT

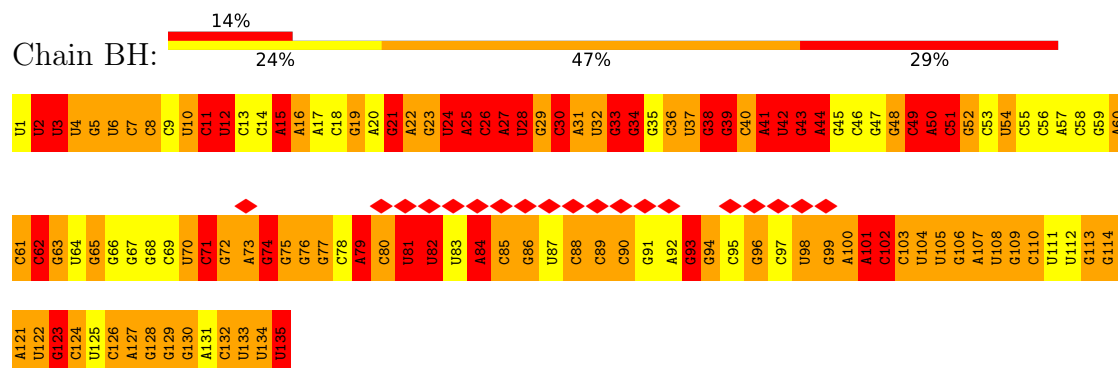


• Molecule 40: SHORT RRNA-III OF THE LARGE RIBOSOMAL SUBUNIT

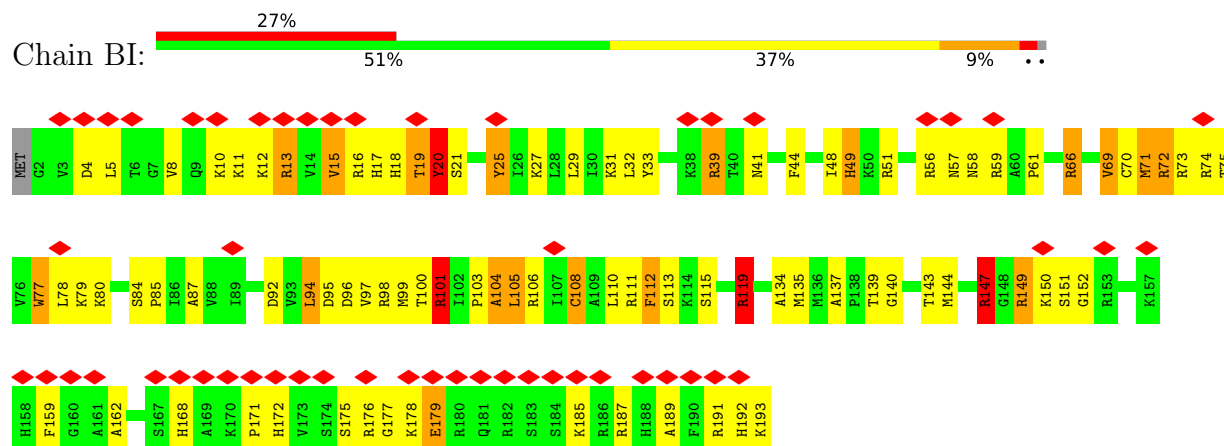




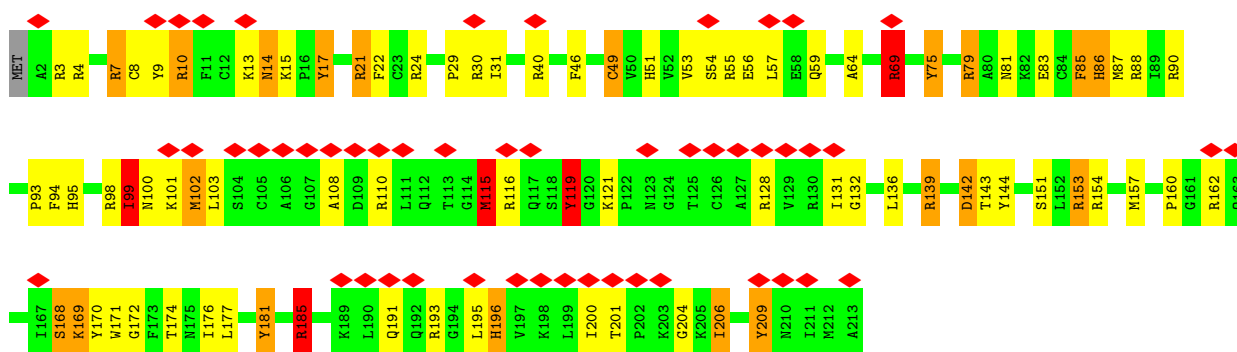
• Molecule 41: SHORT RRNA-IV OF THE LARGE RIBOSOMAL SUBUNIT



• Molecule 42: 60S RIBOSOMAL PROTEIN L18

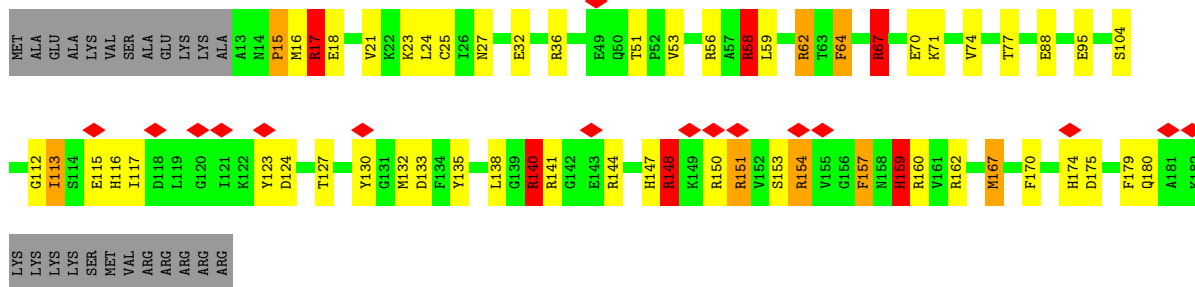


Chain BK: 24% 60% 28% 9%



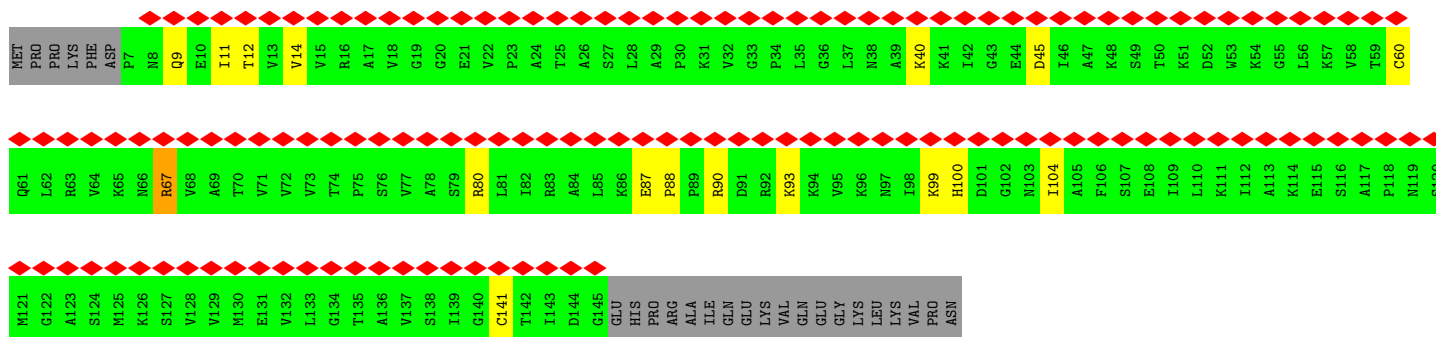
• Molecule 45: 60S RIBOSOMAL PROTEIN L11, PUTATIVE

Chain BL: 8% 58% 23% 12%



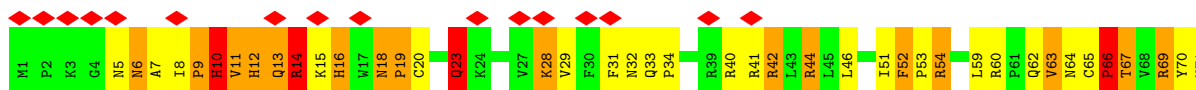
• Molecule 46: 60S RIBOSOMAL PROTEIN L12, PUTATIVE

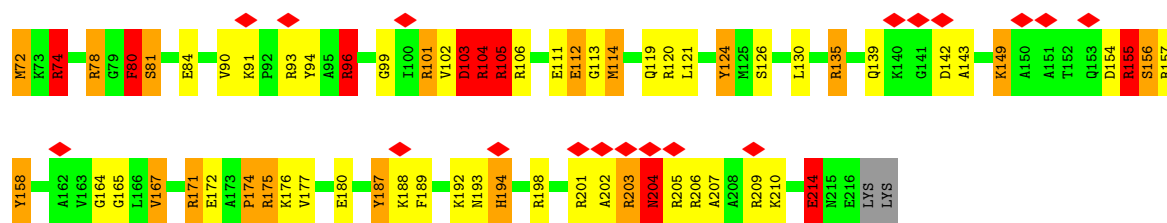
Chain BM: 84% 74% 10% 15%



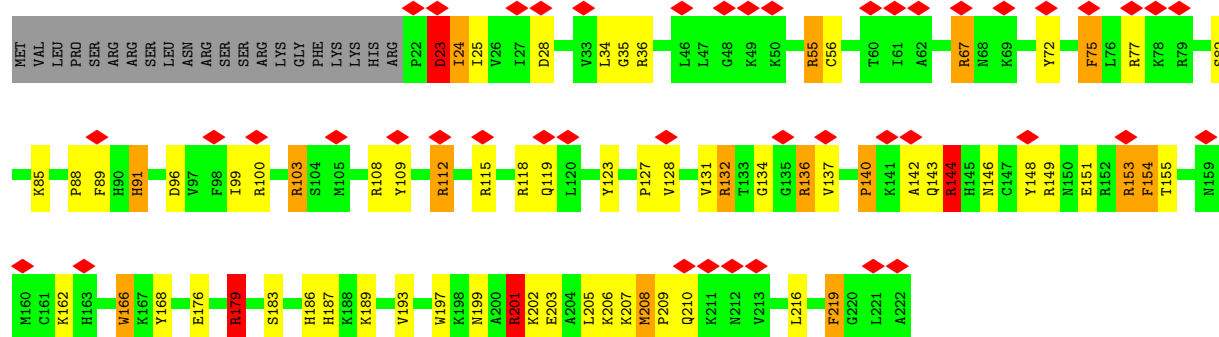
• Molecule 47: 60S RIBOSOMAL PROTEIN L13

Chain BN: 16% 50% 28% 16% 6%

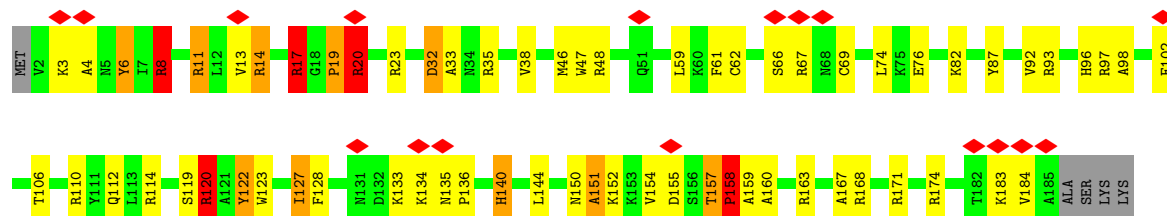




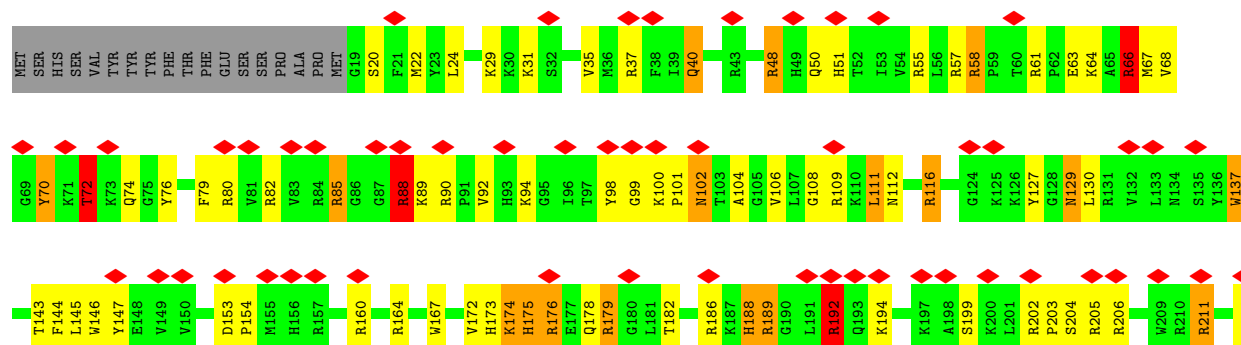
• Molecule 48: 60S RIBOSOMAL PROTEIN L13A, PUTATIVE

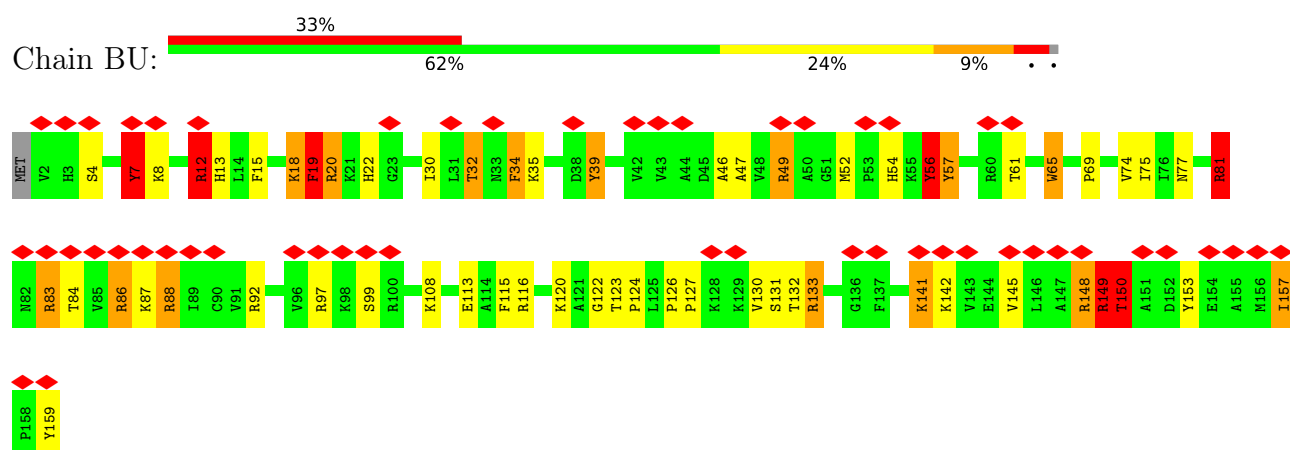


• Molecule 49: PROBABLE 60S RIBOSOMAL PROTEIN L14

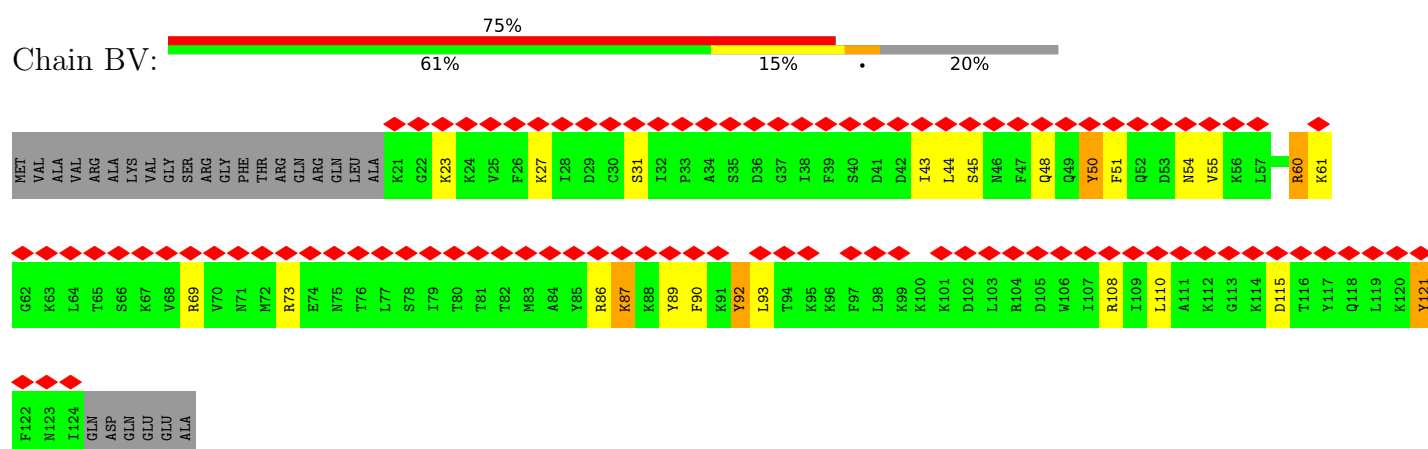


• Molecule 50: RIBOSOMAL PROTEIN L15

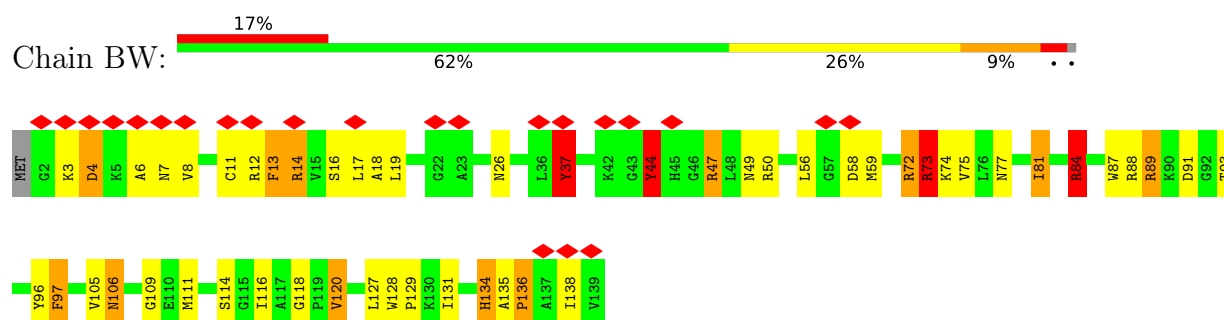




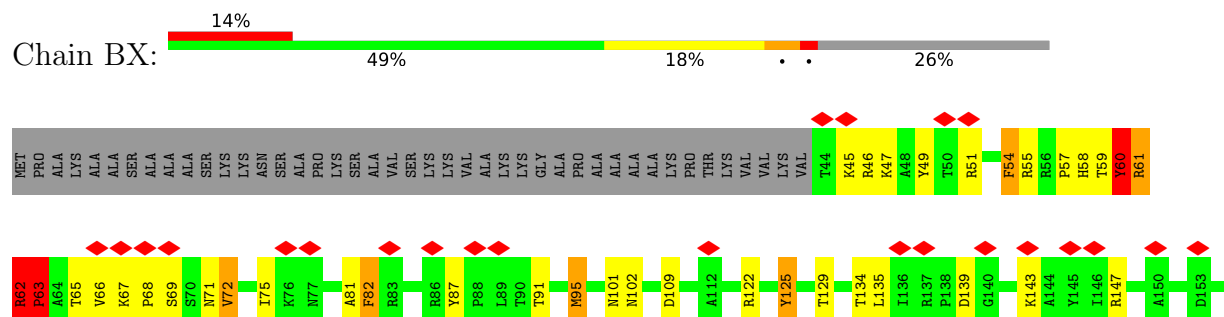
- Molecule 55: 60S RIBOSOMAL PROTEIN L22, PUTATIVE



- Molecule 56: 60S RIBOSOMAL PROTEIN L23, PUTATIVE

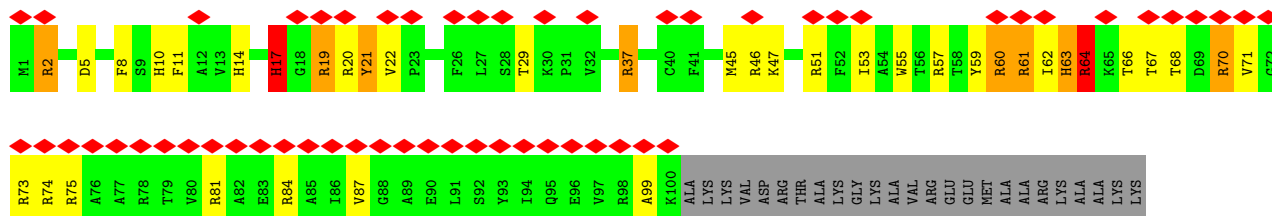


- Molecule 57: 60S RIBOSOMAL PROTEIN L23A

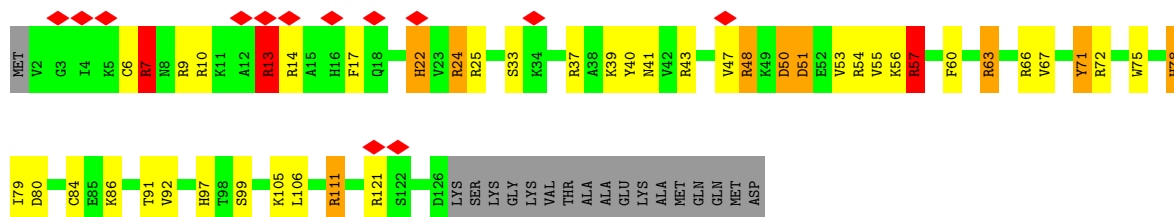




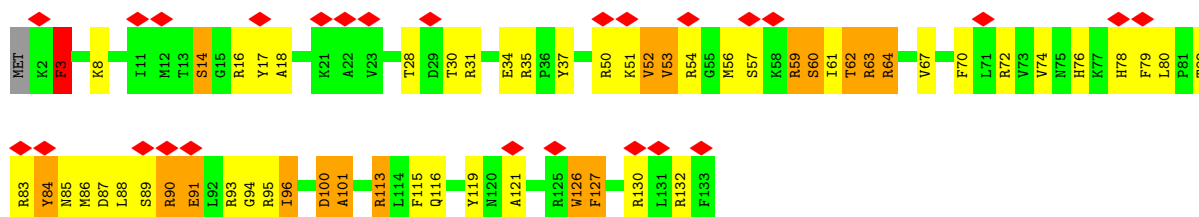
• Molecule 58: 60S RIBOSOMAL PROTEIN L24, PUTATIVE



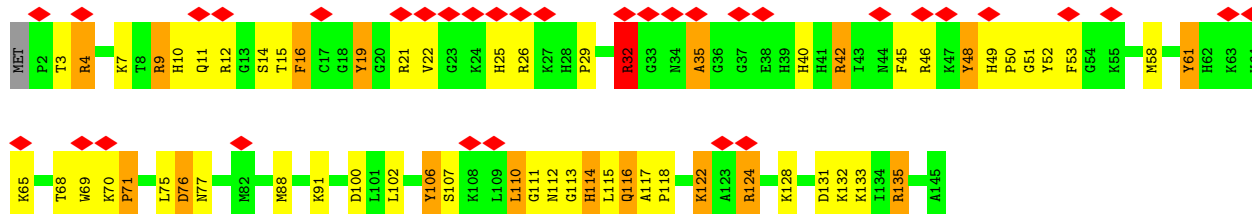
• Molecule 59: 60S RIBOSOMAL PROTEIN L26, PUTATIVE



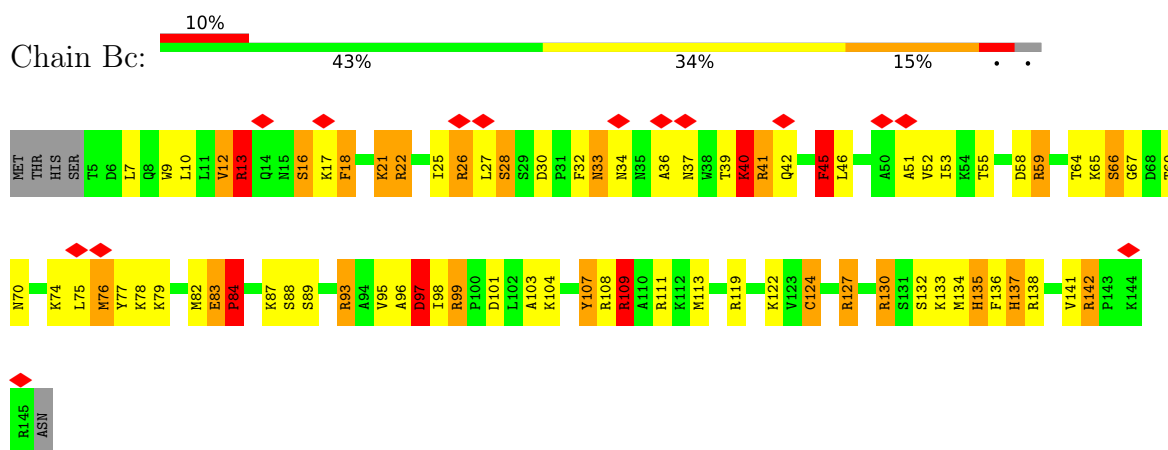
• Molecule 60: 60S RIBOSOMAL PROTEIN L27, PUTATIVE



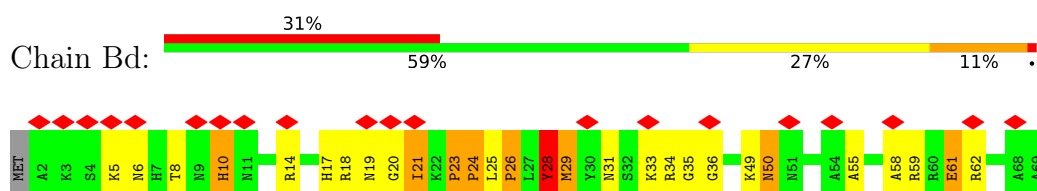
• Molecule 61: 60S RIBOSOMAL PROTEIN L27A



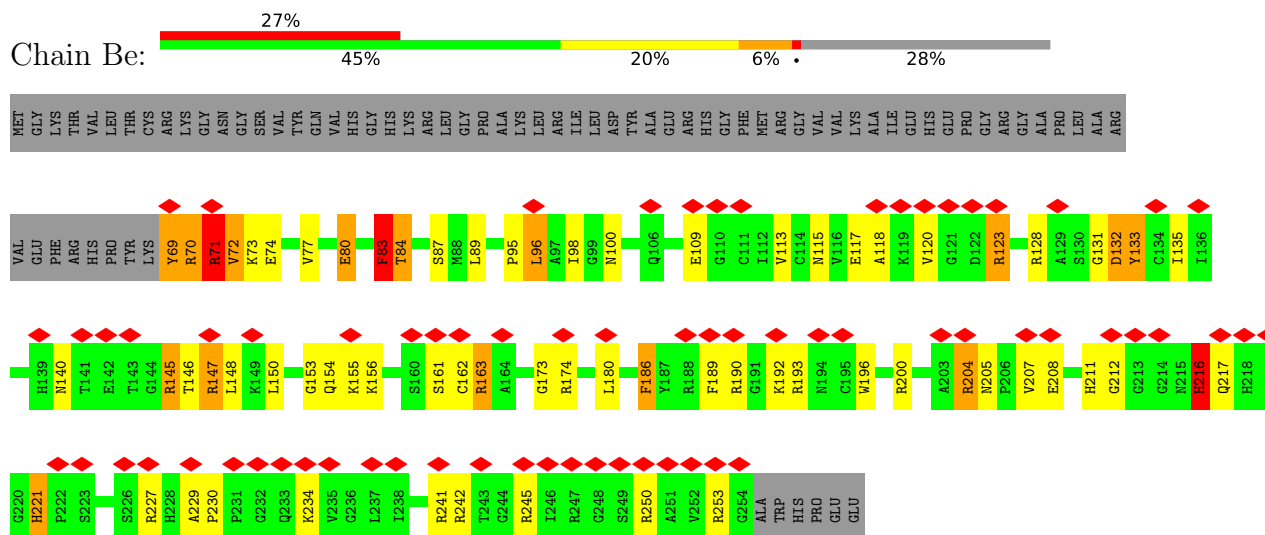
• Molecule 62: 60S RIBOSOMAL PROTEIN L28, PUTATIVE



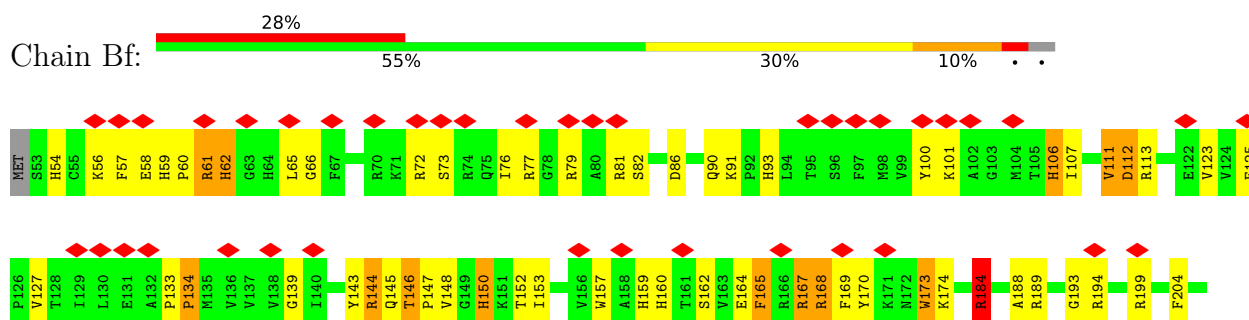
• Molecule 63: 60S RIBOSOMAL PROTEIN L29, PUTATIVE

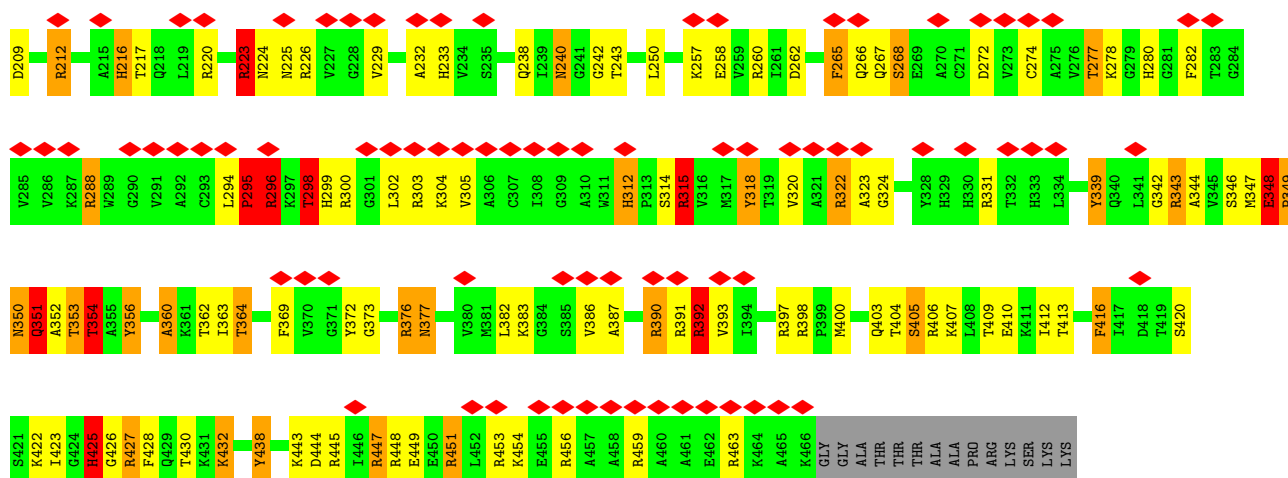


• Molecule 64: 60S RIBOSOMAL PROTEIN L2, PUTATIVE

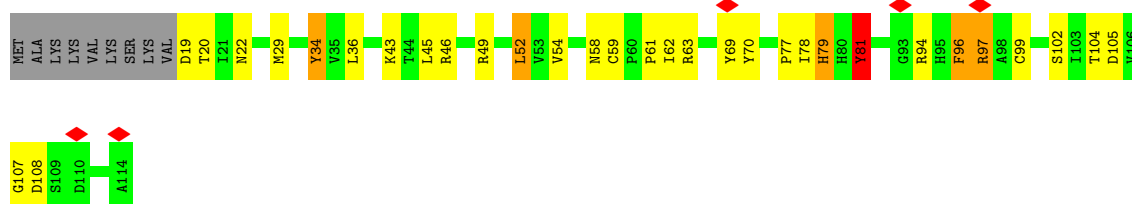


• Molecule 65: RIBOSOMAL PROTEIN L3, MITOCHONDRIAL, PUTATIVE

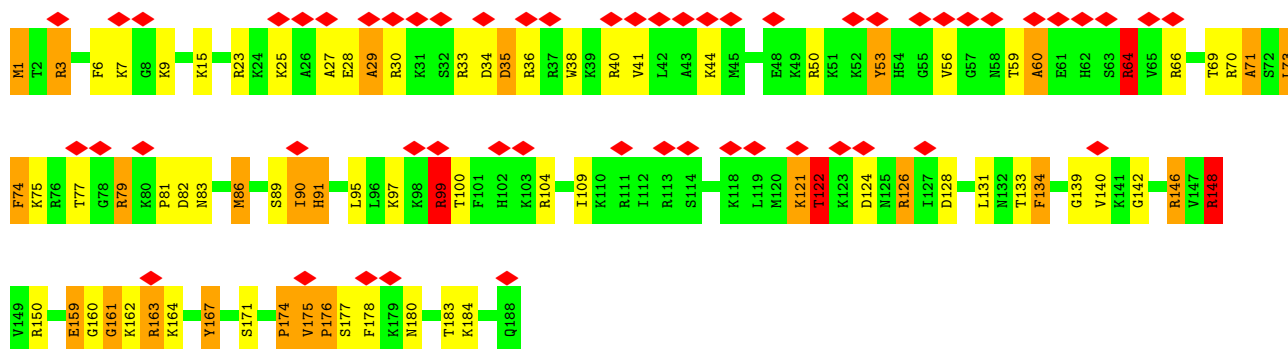




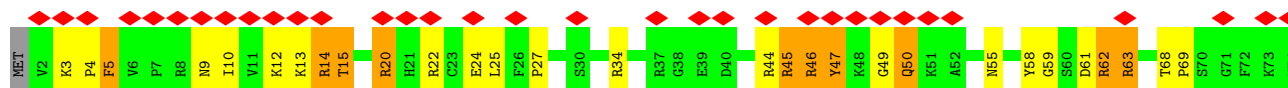
• Molecule 66: 60S RIBOSOMAL PROTEIN L30

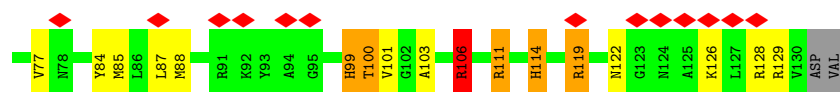


• Molecule 67: 60S RIBOSOMAL SUBUNIT PROTEIN L31, PUTATIVE

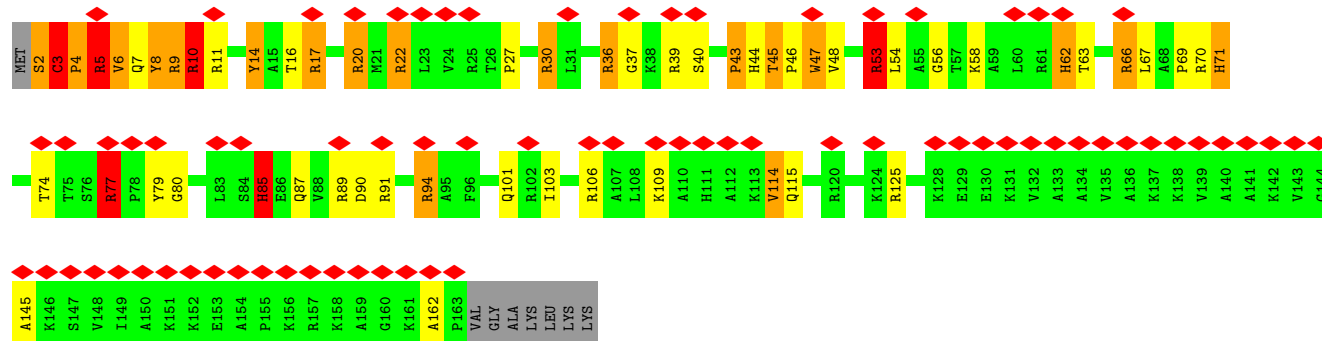


• Molecule 68: 60S RIBOSOMAL PROTEIN L32, PUTATIVE

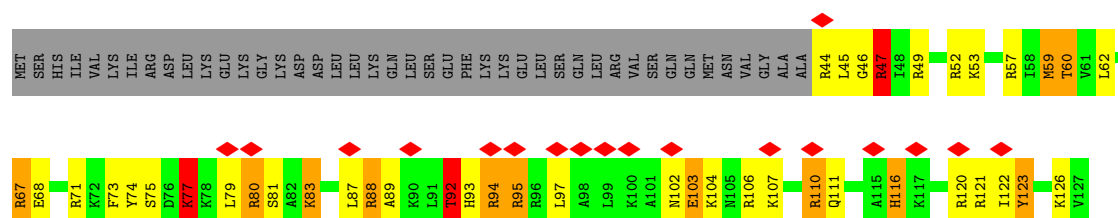
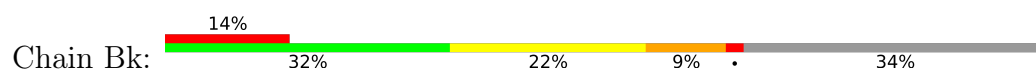




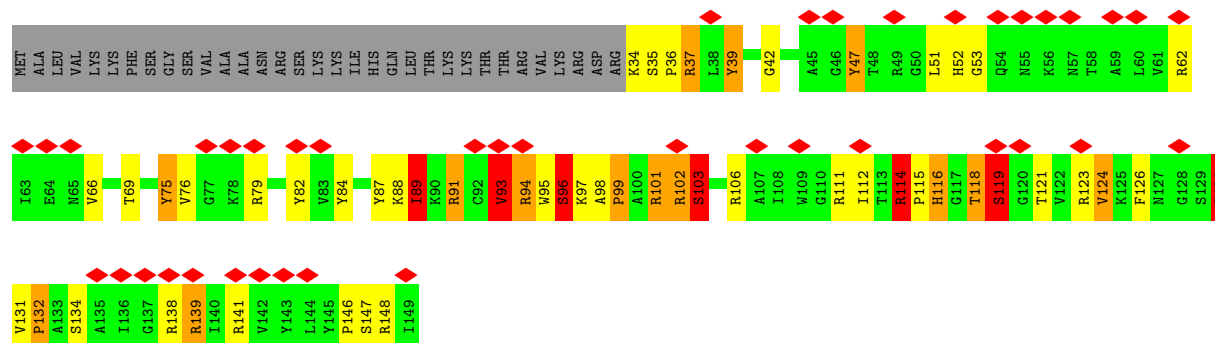
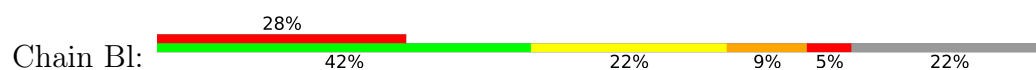
- Molecule 69: 60S RIBOSOMAL PROTEIN L34, PUTATIVE



- Molecule 70: 60S RIBOSOMAL PROTEIN L35, PUTATIVE

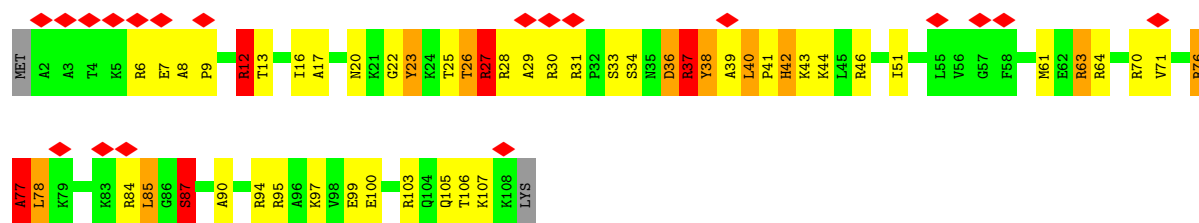


- Molecule 71: 60S RIBOSOMAL PROTEIN L35A, PUTATIVE

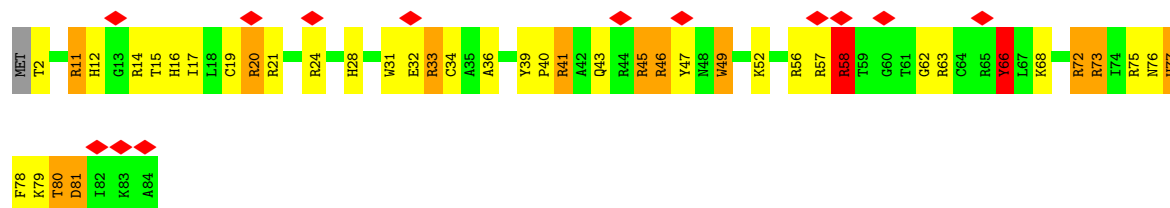


- Molecule 72: RIBOSOMAL PROTEIN L36, PUTATIVE

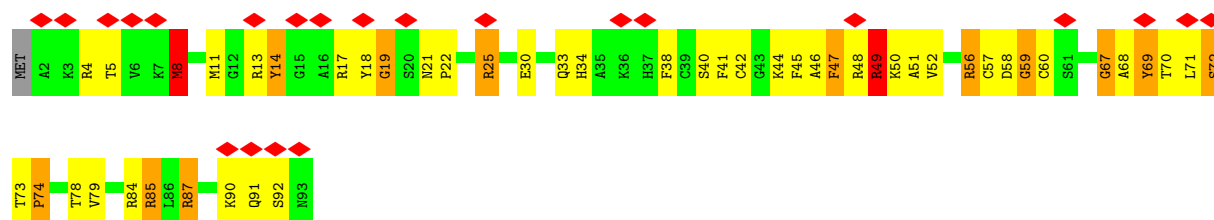




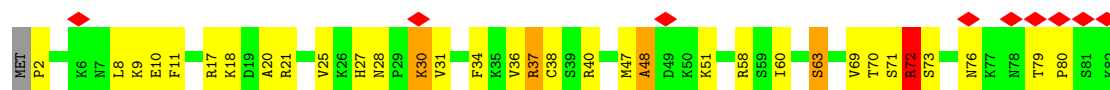
• Molecule 73: RIBOSOMAL PROTEIN L37



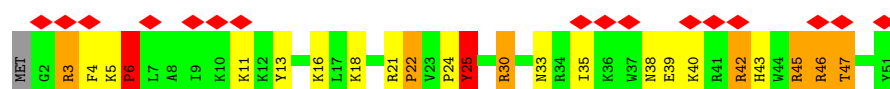
• Molecule 74: 60S RIBOSOMAL PROTEIN L37A, PUTATIVE



• Molecule 75: 60S RIBOSOMAL PROTEIN L38, PUTATIVE

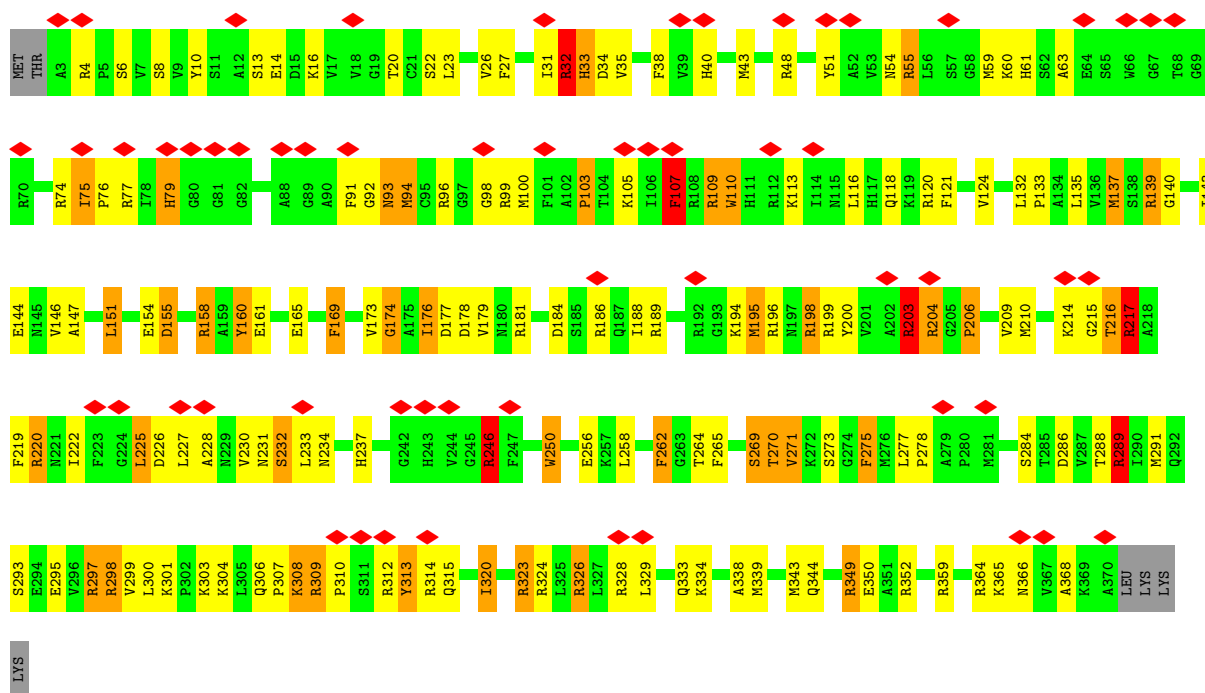


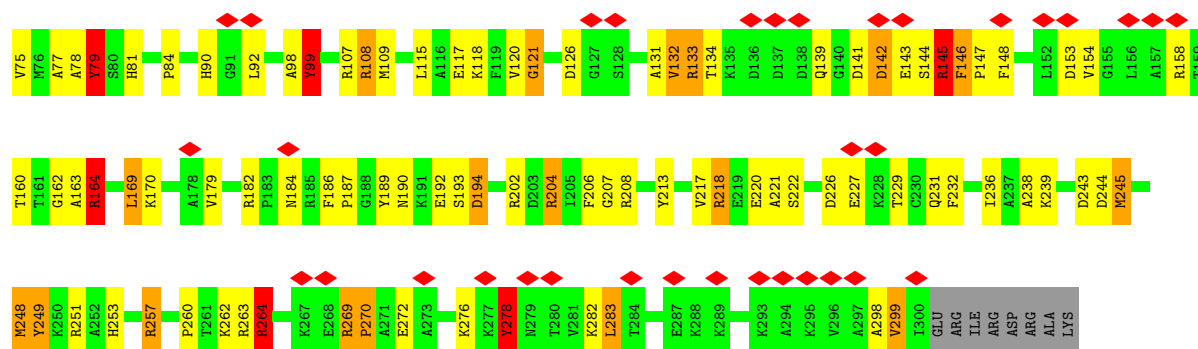
• Molecule 76: 60S RIBOSOMAL PROTEIN L39, PUTATIVE



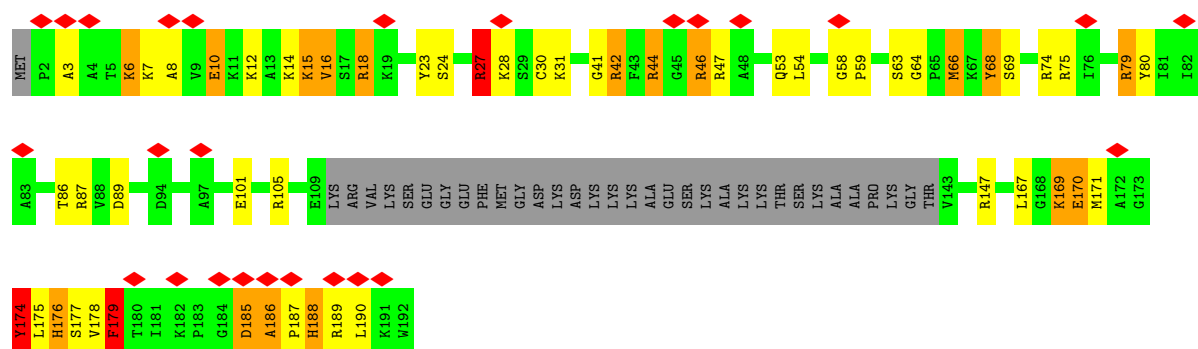
• Molecule 77: 60S RIBOSOMAL PROTEIN L4



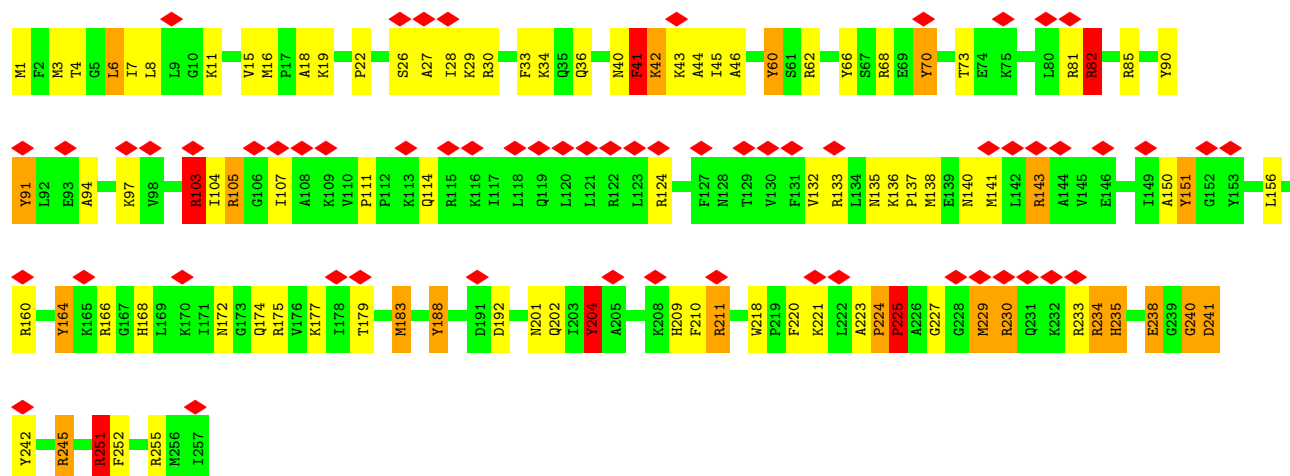




• Molecule 81: 60S RIBOSOMAL PROTEIN L6, PUTATIVE

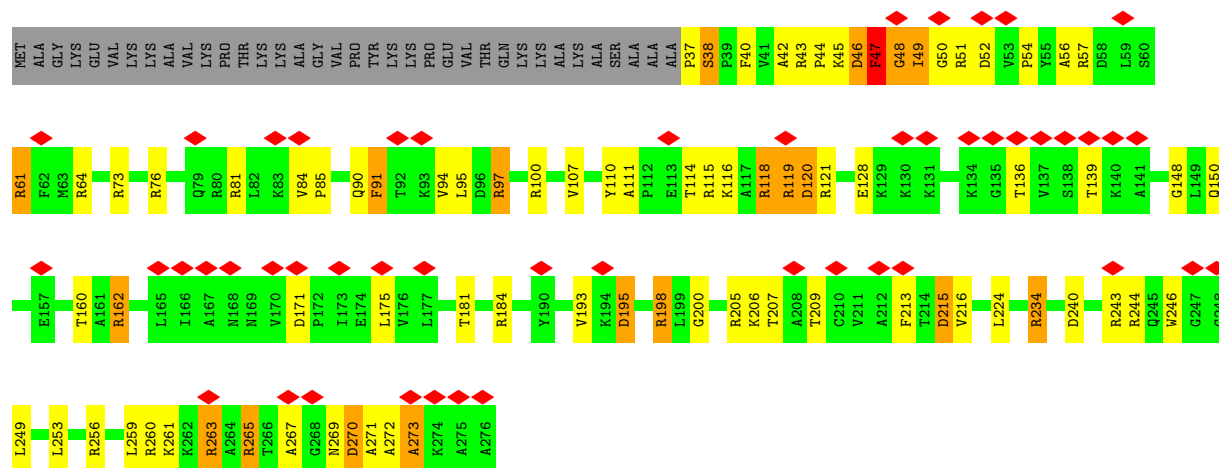


• Molecule 82: 60S RIBOSOMAL PROTEIN L7, PUTATIVE

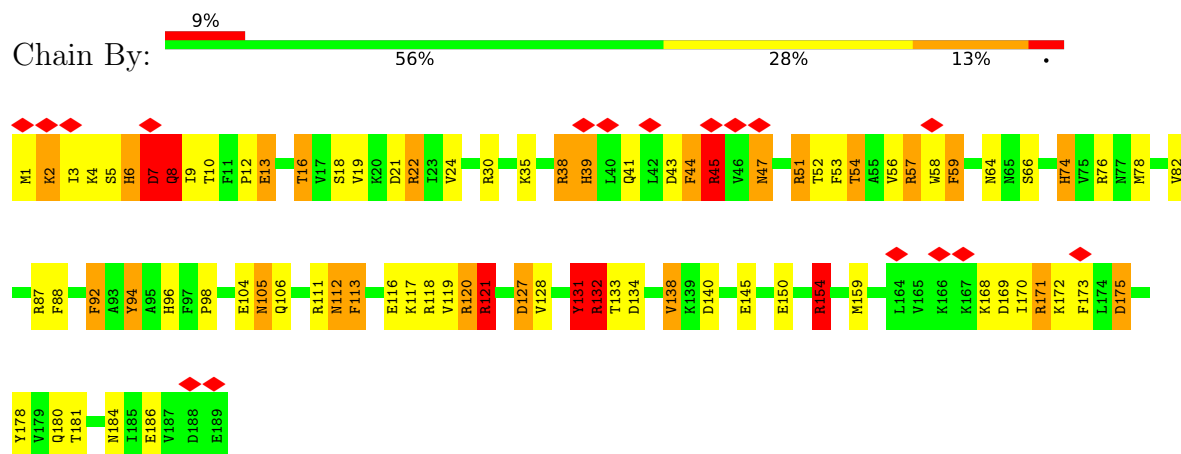


• Molecule 83: 60S RIBOSOMAL PROTEIN L7A, PUTATIVE

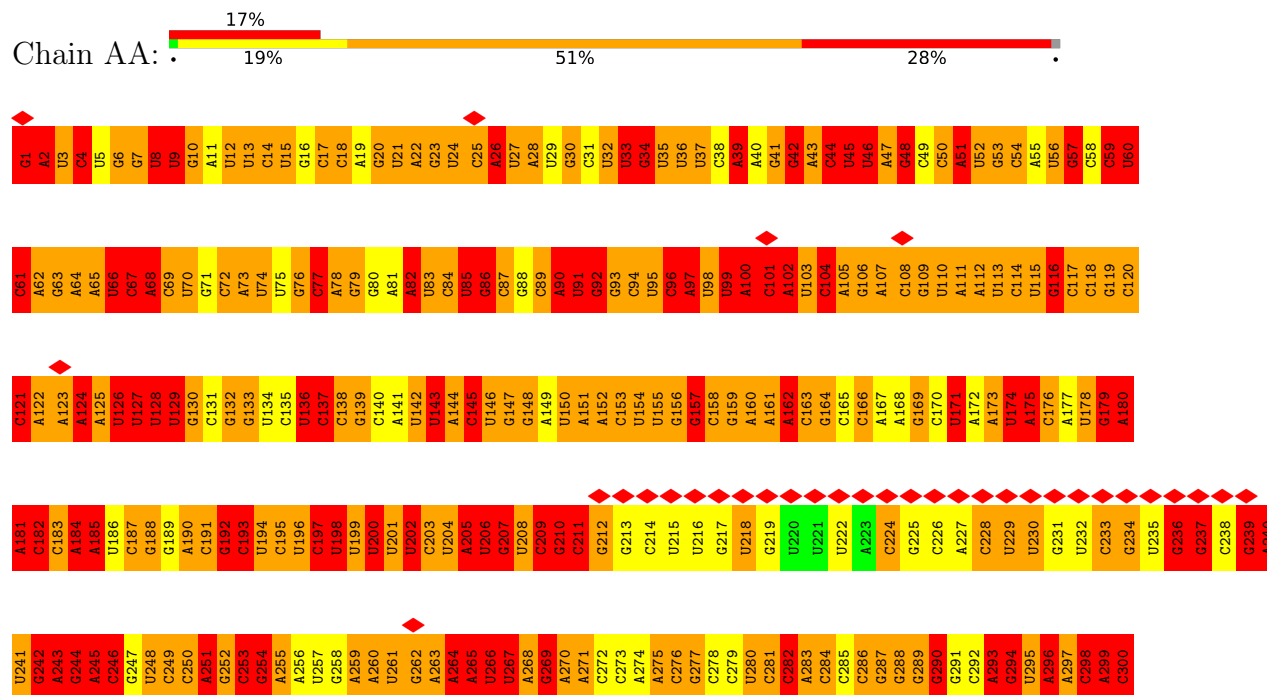




• Molecule 84: 60S RIBOSOMAL PROTEIN L9, PUTATIVE

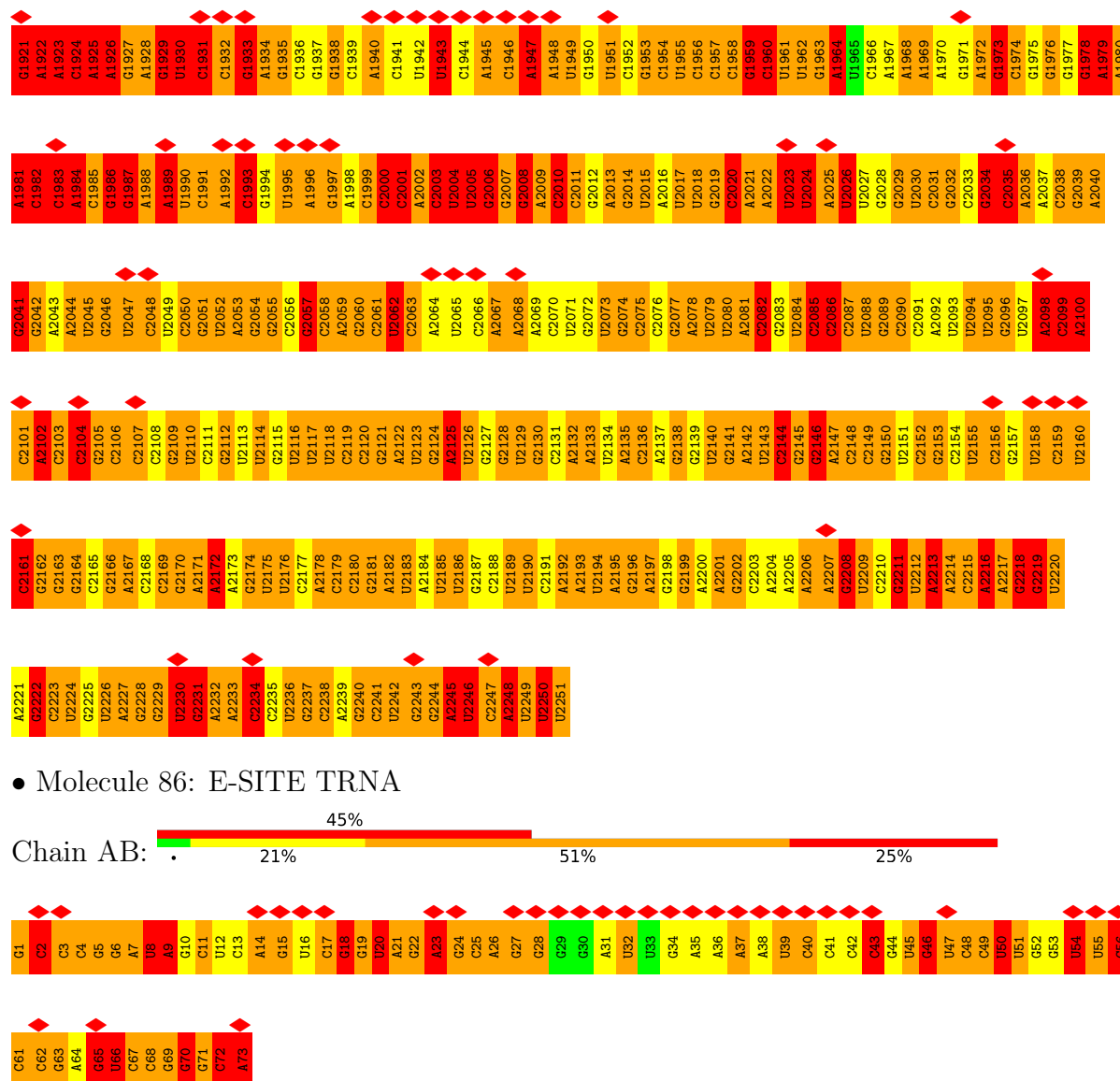


• Molecule 85: 18S RRNA OF THE SMALL RIBOSOMAL SUBUNIT



U1081	U1082	C1083	A1084	U1085	U1086	G1087	U1088	G1089	A1090	C1091	G1092	C1093	C1094	C1095	G1096	G1097	C1098	U1099	U1100	C1101	C1102	A1103	G1104	A1105	A1106	A1107	U1108	G1109	A1110	A1111	G1112	G1113	A1114	G1115	G1116	U1117	U1118	A1119	G1120	U1121	U1122	C1123	G1124	G1125	G1126	G1127	G1128	A1129	G1130	A1131	C1132	C1133	G1134	U1135	A1136	C1137	U1138	G1139	G1140	
G1021	G1022	U1023	G1024	U1025	U1026	U1027	U1028	U1029	U1030	G1031	U1032	C1033	U1034	U1035	A1036	U1037	U1038	U1039	U1040	U1041	U1042	U1043	G1044	U1045	U1046	U1047	U1048	U1049	A1051	C1052	A1053	U1054	U1055	C1056	G1057	U1058	C1059	U1060	C1061	U1062	U1063	C1064	U1065	U1066	C1067	U1068	U1069	G1070	U1071	U1072	U1073	U1074	U1075	U1076	U1077	A1078	C1079	A1080		
U961	U962	U963	C964	G965	G966	C967	U968	C969	U970	U971	G972	U973	U974	G975	G976	U977	U978	U979	U980	A981	G982	A983	A984	G985	U986	C987	U988	U989	U990	G991	C992	G993	A994	A995	A996	U997	U998	A999	U1000	G1001	U1002	G1003	G1004	C1005	C1006	G1007	C1008	G1009	U1010	G1011	C1012	C1013	U1014	C1015	U1016	G1017	C958	G1018	U1019	C1020
C901	A902	G903	U904	C905	U906	G907	C908	C909	G910	U911	C912	U913	U914	G915	A916	A917	U918	U919	A920	C921	A922	C923	A924	G925	C926	A927	U928	G929	C930	G931	A932	U933	A934	A935	C936	G937	A938	A939	C940	C941	A942	U943	C944	A945	G	C	C	C	U	G	G	G	C	C956	A957	C958	C959	C960		
U841	G842	U843	C844	A845	U846	G847	C848	U849	U850	G851	C852	G853	A854	G855	G856	G857	U858	G859	C860	U861	U862	C863	C864	G865	U866	G867	A868	A869	U870	U871	U872	U873	A874	C875	U876	G877	U878	C879	A880	C881	C882	A883	A884	A885	A886	A887	C888	C889	U889	C890	C891	C892	C893	A894	C895	C896	A897	A898	A899	G900
G781	G782	C783	C784	G785	G786	U787	G788	A789	U790	A791	C792	C793	A794	C795	U796	G797	A798	G799	A800	U801	A802	C803	A804	A805	G806	A807	A808	A809	C810	A811	C812	G813	G814	G815	A816	G817	C818	G819	C820	U821	C822	C823	C824	U825	C826	C827	U828	C829	A830	C831	U832	U833	U834	C835	A836	C837	G838	C839	A840	
C721	G722	U723	A724	U725	U726	U727	U728	U729	G730	U731	G732	C733	C734	G735	U736	G737	U738	C739	A740	U741	U742	C743	C744	G745	G746	U747	C748	C749	A750	U751	G752	U753	C754	G755	G756	A757	C758	G759	U760	G761	U762	U763	U764	U765	U766	A767	C768	C769	U770	C771	U772	G773	C774	C775	U776	U777	C778	G779	U780	
C661	U662	C663	C664	A665	A666	A667	A668	G669	C670	G671	U672	A673	U674	A675	U676	U677	A678	A679	U680	G681	C682	U683	G684	U685	U686	G687	C688	U689	G690	U691	U692	A693	A694	A695	G696	G697	G698	U699	U700	C701	G702	U703	A704	G705	U706	U707	G708	A709	U710	C711	U712	G713	U714	G715	U716	G717	C718	C719	A720	
A601	U602	C603	C604	A605	A606	U607	A608	U609	C610	G611	A612	G613	U614	A615	A616	C617	A618	A619	U620	A621	G622	C623	A624	G625	G626	A627	C628	A629	A630	G631	U632	C633	U634	G635	G636	U637	G638	C639	C640	A641	G642	C643	A644	C645	C646	C647	G648	C649	G650	G651	U652	A653	A654	U655	U656	C657	C658	A659	G660	
A541	G542	A543	A544	A545	U546	U547	G548	A549	C550	G551	C552	G553	A554	C555	A556	G557	U558	G559	C560	A561	C562	C563	U564	G565	U566	C567	C568	A569	U570	G571	G572	U573	U574	G575	U576	U577	U578	U579	C580	A581	A582	U583	A584	G585	G586	G587	G588	A589	U590	A591	C592	U593	C594	A595	U596	A597	C598	C599	C600	
U361	G362	A363	G364	G365	A366	G367	C368	A369	U370	C371	U372	G373	C374	C375	G376	U377	A378	U379	C380	A381	C382	G383	C384	A385	G386	U387	U388	A389	U390	G391	C392	A393	C394	G395	U396	G397	U398	A399	U400	U401	G402	G403	A404	C405	U406	U407	C408	C409	U410	U411	G412	G413	C414	G415	U416	U417	G418	A419	C420	
U301	C302	A303	G304	A305	C306	G307	U308	G309	U310	U311	G312	A313	C314	U315	C316	A317	A318	U319	U320	A321	A322	U323	U324	C325	C326	G327	U328	G329	C330	G331	A332	A333	A334	G335	C336	C337	G338	A339	G340	C341	C342	U343	U344	U345	U346	U347	G348	C349	U350	C351	G352	G353	C354	G355	U356	C357	U358	A359	C360	

A1861	U1801	A	G1681	A1561	A1501	G1441	C1381	G1321	U1261	A1201	U1141
C1862	U1802	C	U1682	U1562	A1502	U1442	A1382	C1322	A1262	G1202	C1142
A1863	U1803	A	U1683	U1563	G1503	U1443	C1383	C1323	G1263	G1203	G1143
G1864	U1804	C	U1684	U1564	A1504	C1444	C1384	G1323	U1264	A1204	U1144
C1865	U1805	A	G1685	C1565	G1505	C1445	C1385	G1324	C1265	U1205	U1145
A1866	A1806	G	G1686	A1566	U1506	U1446	C1386	C1325	C1266	A1206	C1146
G1867	C1806	G	U1687	C1567	G1507	U1447	C1386	C1326	A1267	C1207	A1147
U1868	U1807	A	U1688	U1568	A1508	U1448	C1387	U1327	C1268	C1208	G1148
G1869	G1809	C	C1689	C1569	A1509	C1449	C1387	U1327	A1269	U1209	U1149
U1870	C1810	A	A1690	A1570	A1510	U1450	G1388	U1328	C1270	U1210	G1150
U1871	C1811		U1691	A1571	C1511	U1451	G1389	U1329	U1271	C1211	U1151
C1872	C1812		U1692	C1572	U1512	C1452	U1390	U1330	G1272	C1212	U1152
U1873	C1813		C1693	A1573	U1513	U1453	U1391	U1331	C1273	U1213	G1153
G1874	U1814		G1694	C1574	A1514	U1454	U1391	G1331	A1274	A1214	A1154
A1875	U1815		U1695	G1575	A1515	C1455	C1392	U1332	A1275	A1215	A1155
U1876	C1816		U1696	G1576	A1516	A1456	C1393	G1333	A1276	A1216	A1156
U1877	U1817		C1697	G1577	A1517	C1457	C1394	C1334	C1277	C1218	U1157
U1878	C1818		U1698	G1578	A1518	C1458	U1395	C1335	C1278	A1219	U1158
U1879	U1819		A1699	A1579	A1519	C1459	U1396	C1336	A1279	A1220	C1159
U1880	G1820		C1700	A1580	A1520	G1460	C1396	G1337	U1280	G1221	U1161
C1881	C1821		G1701	C1581	U1521	A1461	C1397	A1337	G1281	A1222	A1162
U1882	G1822		G1702	U1582	U1522	A1462	U1398	C1338	A1282	A1223	G1163
C1883	G1823		A1703	U1583	G1523	A1463	U1398	C1338	C1283	C1224	A1164
U1884	G1824		C1704	A1584	A1524	G1464	U1399	C1339	A1284	C1225	C1165
A1885	A1825		G1705	C1585	C1525	C1465	U1400	C1340	C1285	A1226	C1166
U1886	U1826		A1706	C1586	G1526	U1466	G1401	U1341	C1286	G1227	C1167
U1887	U1827		G1707	C1587	G1527	U1467	A1402	C1342	C1287	A1228	C1168
U1888	C1828		A1708	A1588	A1528	G1468	A1402	C1342	A1288	G1229	A1169
U1889	U1829		U1709	G1589	A1529	U1469	G1403	G1343	U1289	U1230	C1170
U1890	C1830		C1710	A1590	U1530	A1470	G1404	G1344	A1291	G1231	A1171
U1891	U1831		U1649	U1591	G1531	G1471	U1405	C1345	U1292	A1172	G1173
C1892	G1832		C1712	C1592	G1532	G1472	U1406	C1346	G1293	U1232	C1174
U1893	C1833		A1713	G1593	C1533	U1473	U1406	C1346	G1297	A1237	G1177
G1894	U1834		G1714	C1594	A1534	U1474	U1407	C1347	A1299	U1238	A1178
C1895	U1835		C1715	G1595	C1535	A1475	C1407	C1348	A1300	C1239	A1179
U1896	U1836		U1716	A1596	C1536	C1476	U1408	A1349	C1301	A1241	U1181
U1897	C1837		C1717	C1597	A1537	A1477	U1409	A1349	A1302	A1242	A1182
C1898	U1838		G1718	A1598	C1538	G1478	C1410	U1350	U1303	G1243	C1183
A1899	G1839		C1719	G1599	A1539	U1479	C1411	U1351	U1304	A1244	G1185
U1900	C1840		C1720	A1600	A1540	C1480	G1412	U1352	A1305	U1245	G1186
G1901	U1841		U1721	G1601	G1541	U1481	G1413	U1353	U1306	G1246	C1187
C1902	C1842		G1722	C1602	C1542	C1482	G1414	U1354	A1307	A1247	A1188
U1903	A1843		U1723	G1603	C1543	A1483	G1415	U1355	U1307	G1248	G1190
C1904	A1844		U1724	A1604	U1544	G1484	G1416	U1356	U1307	A1247	C1191
A1905	G1845		G1725	G1605	U1545	G1485	U1417	U1357	U1307	G1246	G1192
U1906	U1846		C1726	C1606	G1546	G1486	U1418	A1358	G1308	A1247	A1193
U1907	U1847		U1727	A1607	G1547	G1487	U1419	C1359	G1308	U1248	U1194
A1908	G1848		G1728	U1608	A1548	G1488	U1420	C1360	G1309	U1249	U1195
U1909	A1849		C1729	U1609	G1549	G1489	A1422	A1361	G1310	G1250	C1196
A1910	G1850		G1730	U1610	C1550	A1490	U1421	U1362	U1311	A1251	U1197
U1911	A1851		G1731	A1611	U1551	G1491	U1422	U1363	G1312	G1252	U1198
U1912	U1852		G1732	C1612	U1552	U1492	A1423	U1364	G1313	C1253	C1199
U1913	U1853		G1733	G1613	G1553	A1493	G1424	U1365	U1314	U1254	U1900
G1914	U1854		G1734	G1614	G1554	C1494	G1425	C1366	C1315	C1255	
U1915	U1855		U1735	A1615	U1555	G1495	G1426	G1367	G1316	A1257	
C1916	U1856		U1736	U1616	U1556	U1496	A1427	U1368	G1317	U1258	
U1917	G1857		G	U1617	U1557	U1497	A1428	G1369	G1318	U1259	
U1918	U1858		U	G1618	U1558	C1498	U1429	C1370	U1319	G1260	
U1919	G1859		C	A1619	U1559	G1499	U1430	C1371	G1320		
A1920	A1860		C	G1620	A1560	C1500	U1431	U1372			
							U1432	U1373			
							A1433	U1374			
							C1434	U1375			
							U1435	C1376			
							C1436	U1377			
							A1437	U1378			
							U1438	U1379			
							C1439	U1380			
							A1440				



• Molecule 86: E-SITE TRNA



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	164000	Depositor
Resolution determination method	Not provided	
CTF correction method	PHASE-FLIPPING	Depositor
Microscope	FEI TECNAI F30	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	25	Depositor
Minimum defocus (nm)	1500	Depositor
Maximum defocus (nm)	4000	Depositor
Magnification	59000	Depositor
Image detector	KODAK SO-163 FILM	Depositor
Maximum map value	483783.781	Depositor
Minimum map value	-202702.703	Depositor
Average map value	7679.429	Depositor
Map value standard deviation	33729.660	Depositor
Recommended contour level	108000	Depositor
Map size (\AA)	391.31, 391.31, 391.31	wwPDB
Map dimensions	359, 359, 359	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (\AA)	1.09, 1.09, 1.09	Depositor

5 Model quality ⓘ

5.1 Standard geometry ⓘ

The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 5$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# $ Z > 5$	RMSZ	# $ Z > 5$
1	A0	1.19	2/1808 (0.1%)	1.83	45/2432 (1.9%)
2	A1	1.37	5/1973 (0.3%)	1.90	53/2657 (2.0%)
3	A2	1.08	0/1507	1.76	36/2027 (1.8%)
4	A3	1.32	1/2026 (0.0%)	1.92	53/2699 (2.0%)
5	A4	1.33	2/1623 (0.1%)	2.04	60/2185 (2.7%)
6	A5	1.42	6/1574 (0.4%)	1.83	36/2100 (1.7%)
7	A6	1.38	3/1548 (0.2%)	1.95	44/2076 (2.1%)
8	A7	1.06	0/2471	1.64	33/3368 (1.0%)
9	A8	1.09	0/337	1.68	4/445 (0.9%)
10	A9	0.91	0/542	1.55	8/722 (1.1%)
11	AC	1.41	1/1655 (0.1%)	1.76	29/2240 (1.3%)
12	AD	1.02	0/877	1.60	13/1182 (1.1%)
13	AE	1.32	4/1324 (0.3%)	1.83	29/1771 (1.6%)
14	AF	0.76	0/946	1.36	11/1270 (0.9%)
15	AG	1.45	2/1170 (0.2%)	1.93	41/1567 (2.6%)
16	AH	1.24	0/937	1.77	21/1263 (1.7%)
17	AI	0.92	0/1098	1.64	15/1473 (1.0%)
18	AJ	1.29	0/1035	1.78	13/1386 (0.9%)
19	AK	1.06	0/1211	1.77	24/1625 (1.5%)
20	AL	1.24	1/1033 (0.1%)	1.91	27/1380 (2.0%)
21	AM	1.05	0/1247	1.76	29/1666 (1.7%)
22	AO	1.08	0/1206	1.81	23/1613 (1.4%)
23	AP	1.40	1/1766 (0.1%)	1.83	37/2383 (1.6%)
24	AQ	1.11	0/839	1.77	23/1139 (2.0%)
25	AR	1.52	3/612 (0.5%)	1.96	26/835 (3.1%)
26	AS	1.22	0/1137	1.73	27/1520 (1.8%)
27	AT	1.17	2/1065 (0.2%)	1.99	30/1411 (2.1%)
28	AU	1.00	0/681	1.54	6/907 (0.7%)
29	AV	1.17	0/825	1.95	19/1105 (1.7%)
30	AW	1.37	1/648 (0.2%)	1.84	14/868 (1.6%)
31	AX	1.11	2/1649 (0.1%)	1.71	35/2203 (1.6%)
32	AY	1.22	0/521	1.83	9/685 (1.3%)
33	AZ	1.20	1/527 (0.2%)	1.77	13/702 (1.9%)
34	BA	4.08	8852/44057 (20.1%)	3.34	8033/68678 (11.7%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
35	BB	3.66	5797/34826 (16.6%)	2.99	5090/54269 (9.4%)
36	BC	4.32	938/4004 (23.4%)	3.24	674/6235 (10.8%)
37	BD	4.09	601/2830 (21.2%)	3.33	521/4410 (11.8%)
38	BE	3.90	902/4956 (18.2%)	3.64	1067/7716 (13.8%)
39	BF	3.69	277/1691 (16.4%)	3.64	354/2627 (13.5%)
40	BG	4.34	1057/4358 (24.3%)	3.41	840/6797 (12.4%)
41	BH	3.79	548/3201 (17.1%)	3.52	652/4987 (13.1%)
42	BI	1.47	4/1553 (0.3%)	2.04	64/2070 (3.1%)
43	BJ	0.72	0/1743	1.28	12/2339 (0.5%)
44	BK	1.40	3/1760 (0.2%)	1.89	45/2359 (1.9%)
45	BL	1.33	1/1385 (0.1%)	1.80	33/1851 (1.8%)
46	BM	0.67	0/1033	1.21	1/1394 (0.1%)
47	BN	2.40	4/1793 (0.2%)	2.18	80/2392 (3.3%)
48	BO	1.46	2/1655 (0.1%)	1.84	44/2214 (2.0%)
49	BP	1.48	3/1506 (0.2%)	1.97	45/2014 (2.2%)
50	BQ	1.52	1/1755 (0.1%)	1.84	42/2346 (1.8%)
51	BR	1.39	1/1270 (0.1%)	1.99	39/1705 (2.3%)
52	BS	1.36	3/1508 (0.2%)	2.06	54/2028 (2.7%)
53	BT	1.45	1/1689 (0.1%)	1.97	65/2232 (2.9%)
54	BU	1.35	1/1290 (0.1%)	1.89	39/1734 (2.2%)
55	BV	0.96	0/878	1.57	13/1169 (1.1%)
56	BW	1.44	5/1059 (0.5%)	2.00	32/1424 (2.2%)
57	BX	1.56	4/1007 (0.4%)	1.97	35/1353 (2.6%)
58	BY	1.28	1/857 (0.1%)	1.82	26/1150 (2.3%)
59	BZ	1.44	3/1021 (0.3%)	2.07	43/1362 (3.2%)
60	Ba	1.40	0/1111	1.95	31/1479 (2.1%)
61	Bb	1.35	4/1165 (0.3%)	1.92	34/1554 (2.2%)
62	Bc	2.50	3/1145 (0.3%)	2.20	62/1528 (4.1%)
63	Bd	1.21	2/582 (0.3%)	1.98	22/777 (2.8%)
64	Be	1.50	7/1416 (0.5%)	1.91	38/1905 (2.0%)
65	Bf	1.43	8/3387 (0.2%)	1.99	129/4548 (2.8%)
66	Bg	1.75	3/745 (0.4%)	2.09	23/1005 (2.3%)
67	Bh	1.27	4/1551 (0.3%)	2.00	56/2059 (2.7%)
68	Bi	1.39	2/1076 (0.2%)	1.91	30/1439 (2.1%)
69	Bj	1.24	3/1312 (0.2%)	1.85	34/1743 (2.0%)
70	Bk	1.46	0/726	2.06	31/957 (3.2%)
71	Bl	1.44	4/958 (0.4%)	2.05	35/1290 (2.7%)
72	Bm	1.54	1/859 (0.1%)	1.97	29/1141 (2.5%)
73	Bn	1.60	1/713 (0.1%)	2.01	28/949 (3.0%)
74	Bo	1.70	8/727 (1.1%)	1.88	26/968 (2.7%)
75	Bp	1.50	1/666 (0.2%)	1.87	15/885 (1.7%)
76	Bq	1.31	1/471 (0.2%)	1.71	9/626 (1.4%)
77	Br	1.46	5/2937 (0.2%)	1.98	106/3943 (2.7%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
78	Bs	1.40	1/433 (0.2%)	1.86	12/572 (2.1%)
79	Bt	1.36	1/883 (0.1%)	1.87	24/1170 (2.1%)
80	Bu	1.30	1/2397 (0.0%)	1.88	65/3219 (2.0%)
81	Bv	1.30	1/1242 (0.1%)	1.84	29/1667 (1.7%)
82	Bw	1.44	2/2105 (0.1%)	1.91	49/2823 (1.7%)
83	Bx	1.48	5/1936 (0.3%)	1.85	38/2603 (1.5%)
84	By	1.32	1/1561 (0.1%)	1.95	62/2098 (3.0%)
85	AA	3.42	7237/52940 (13.7%)	3.10	8226/82489 (10.0%)
86	AB	2.36	80/1740 (4.6%)	2.66	188/2712 (6.9%)
All	All	3.06	26432/250887 (10.5%)	2.80	28260/369909 (7.6%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A0	0	19
2	A1	0	18
3	A2	0	11
4	A3	0	23
5	A4	0	27
6	A5	0	23
7	A6	0	26
8	A7	0	17
9	A8	0	4
10	A9	0	4
11	AC	0	21
12	AD	0	10
13	AE	0	25
14	AF	0	6
15	AG	0	17
16	AH	0	8
17	AI	0	10
18	AJ	0	11
19	AK	0	20
20	AL	0	6
21	AM	0	18
22	AO	0	17
23	AP	0	14
24	AQ	0	7
25	AR	0	12

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Mol	Chain	#Chirality outliers	#Planarity outliers
26	AS	0	7
27	AT	0	27
28	AU	0	7
29	AV	0	13
30	AW	0	13
31	AX	0	10
32	AY	0	13
33	AZ	0	3
34	BA	0	1404
35	BB	0	967
36	BC	0	128
37	BD	0	89
38	BE	0	169
39	BF	0	61
40	BG	0	144
41	BH	0	87
42	BI	0	19
43	BJ	0	7
44	BK	0	26
45	BL	0	15
47	BN	1	31
48	BO	0	21
49	BP	0	24
50	BQ	0	28
51	BR	0	21
52	BS	0	21
53	BT	0	24
54	BU	0	21
55	BV	0	6
56	BW	0	10
57	BX	0	7
58	BY	0	15
59	BZ	0	12
60	Ba	0	20
61	Bb	0	15
62	Bc	0	27
63	Bd	0	10
64	Be	0	21
65	Bf	0	48
66	Bg	0	8
67	Bh	0	23
68	Bi	0	13

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Mol	Chain	#Chirality outliers	#Planarity outliers
69	Bj	1	23
70	Bk	0	12
71	Bl	0	23
72	Bm	0	16
73	Bn	0	14
74	Bo	0	12
75	Bp	0	6
76	Bq	0	9
77	Br	0	46
78	Bs	0	5
79	Bt	0	13
80	Bu	0	34
81	Bv	0	22
82	Bw	0	32
83	Bx	0	19
84	By	1	27
85	AA	0	1567
86	AB	0	45
All	All	3	5934

All (26432) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	469	G	C5-C4	80.15	1.94	1.38
47	BN	7	ALA	CA-C	79.39	3.59	1.52
34	BA	743	A	N9-C4	74.20	1.82	1.37
62	Bc	78	LYS	CD-CE	69.91	3.26	1.51
34	BA	214	A	C6-N1	57.25	1.75	1.35
34	BA	214	A	C5-C4	55.92	1.77	1.38
85	AA	469	G	C8-N7	52.60	1.62	1.30
34	BA	743	A	N9-C8	46.86	1.75	1.37
34	BA	214	A	C5-C6	42.37	1.79	1.41
34	BA	546	U	O3'-P	42.32	2.12	1.61
34	BA	214	A	C2-N3	40.77	1.70	1.33
34	BA	214	A	N3-C4	40.20	1.58	1.34
34	BA	214	A	N1-C2	39.07	1.69	1.34
85	AA	469	G	N9-C4	33.75	1.65	1.38
85	AA	469	G	N9-C8	32.48	1.60	1.37
34	BA	743	A	N7-C5	-32.37	1.19	1.39
34	BA	743	A	C5-C6	31.22	1.69	1.41
34	BA	481	A	N9-C4	-30.64	1.19	1.37
34	BA	743	A	C5-C4	27.33	1.57	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	62	A	N9-C4	-26.76	1.21	1.37
34	BA	399	G	N9-C4	-23.70	1.19	1.38
34	BA	547	C	C4'-C3'	23.33	1.78	1.53
34	BA	557	U	C3'-C2'	22.13	1.77	1.52
85	AA	1916	A	N9-C4	-22.12	1.24	1.37
34	BA	557	U	O3'-P	21.67	1.87	1.61
38	BE	123	A	N9-C4	-21.08	1.25	1.37
38	BE	139	U	P-O5'	-20.49	1.39	1.59
34	BA	471	U	N1-C2	-20.26	1.20	1.38
40	BG	9	G	O3'-P	-19.82	1.37	1.61
34	BA	1011	G	P-O5'	-19.73	1.40	1.59
41	BH	34	G	P-O5'	-19.71	1.40	1.59
85	AA	1228	A	N9-C4	-19.61	1.26	1.37
34	BA	1556	A	N9-C4	-19.58	1.26	1.37
34	BA	1213	A	N9-C4	-19.42	1.26	1.37
85	AA	2193	A	N9-C4	-19.41	1.26	1.37
40	BG	25	G	P-O5'	-19.30	1.40	1.59
34	BA	49	A	N9-C4	-19.27	1.26	1.37
85	AA	597	A	N9-C4	-19.22	1.26	1.37
34	BA	1321	A	P-O5'	-19.21	1.40	1.59
85	AA	2133	A	N9-C4	-19.21	1.26	1.37
36	BC	39	G	P-O5'	-19.11	1.40	1.59
34	BA	759	A	N9-C4	-19.10	1.26	1.37
40	BG	64	C	P-O5'	-19.07	1.40	1.59
41	BH	72	G	C4'-C3'	18.84	1.73	1.53
40	BG	39	A	N9-C4	-18.75	1.26	1.37
34	BA	1442	A	N9-C4	18.72	1.49	1.37
35	BB	1315	C	P-O5'	-18.65	1.41	1.59
35	BB	822	G	N9-C4	18.62	1.52	1.38
85	AA	970	U	P-O5'	-18.62	1.41	1.59
85	AA	989	U	P-O5'	-18.61	1.41	1.59
34	BA	517	A	P-O5'	-18.60	1.41	1.59
35	BB	1306	G	P-O5'	-18.57	1.41	1.59
34	BA	1608	C	P-O5'	-18.53	1.41	1.59
41	BH	16	A	N9-C4	-18.53	1.26	1.37
35	BB	1132	A	P-O5'	-18.49	1.41	1.59
40	BG	24	A	O3'-P	-18.48	1.39	1.61
85	AA	2125	A	N9-C4	-18.42	1.26	1.37
34	BA	1177	C	P-O5'	-18.39	1.41	1.59
34	BA	1411	C	P-O5'	-18.36	1.41	1.59
34	BA	480	G	N9-C4	-18.33	1.23	1.38
38	BE	149	A	N9-C4	-18.31	1.26	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1832	A	P-O5'	-18.28	1.41	1.59
41	BH	121	A	N9-C4	-18.26	1.26	1.37
34	BA	992	A	N9-C4	-18.25	1.26	1.37
34	BA	757	G	C2'-C1'	-18.21	1.33	1.53
34	BA	1468	U	P-O5'	-18.19	1.41	1.59
85	AA	917	A	N9-C4	-18.15	1.26	1.37
34	BA	1649	A	N9-C4	-18.09	1.26	1.37
38	BE	194	A	N9-C4	-18.07	1.27	1.37
38	BE	117	A	P-O5'	-18.00	1.41	1.59
34	BA	321	G	N7-C5	-17.89	1.28	1.39
34	BA	875	G	P-O5'	-17.88	1.41	1.59
35	BB	109	U	P-O5'	-17.85	1.42	1.59
35	BB	577	U	P-O5'	-17.83	1.42	1.59
85	AA	869	A	N9-C4	-17.80	1.27	1.37
35	BB	99	G	P-O5'	-17.79	1.42	1.59
85	AA	1731	G	P-O5'	-17.69	1.42	1.59
35	BB	993	A	N9-C4	-17.68	1.27	1.37
36	BC	113	G	P-O5'	-17.68	1.42	1.59
35	BB	703	U	P-O5'	-17.66	1.42	1.59
35	BB	404	A	N9-C4	-17.64	1.27	1.37
38	BE	163	A	N9-C4	-17.61	1.27	1.37
34	BA	15	G	N9-C4	-17.54	1.24	1.38
35	BB	435	A	N9-C4	-17.49	1.27	1.37
34	BA	135	G	P-O5'	-17.40	1.42	1.59
85	AA	788	G	P-O5'	-17.39	1.42	1.59
34	BA	111	U	C1'-N1	-17.38	1.22	1.46
34	BA	1017	C	P-O5'	-17.37	1.42	1.59
34	BA	919	A	P-O5'	-17.37	1.42	1.59
36	BC	24	G	P-O5'	-17.35	1.42	1.59
35	BB	415	A	N9-C4	-17.29	1.27	1.37
85	AA	367	A	N9-C4	-17.28	1.27	1.37
85	AA	97	A	N9-C4	-17.20	1.27	1.37
85	AA	2063	C	P-O5'	-17.13	1.42	1.59
34	BA	57	A	N9-C4	-17.11	1.27	1.37
40	BG	33	G	N9-C4	-17.07	1.24	1.38
85	AA	2060	G	P-O5'	-17.07	1.42	1.59
41	BH	72	G	O3'-P	-17.03	1.40	1.61
34	BA	1454	G	N9-C4	-16.99	1.24	1.38
85	AA	679	A	N9-C4	-16.92	1.27	1.37
37	BD	100	A	N9-C4	-16.87	1.27	1.37
34	BA	115	U	P-O5'	-16.87	1.42	1.59
34	BA	436	U	P-O5'	-16.83	1.43	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	19	A	N9-C4	-16.79	1.27	1.37
34	BA	141	G	N9-C4	-16.76	1.24	1.38
35	BB	1129	C	P-O5'	-16.74	1.43	1.59
34	BA	1211	G	N7-C5	-16.73	1.29	1.39
85	AA	2092	A	P-O5'	-16.71	1.43	1.59
41	BH	29	G	C6-N1	-16.70	1.27	1.39
34	BA	383	G	P-O5'	-16.70	1.43	1.59
34	BA	316	G	P-O5'	-16.62	1.43	1.59
85	AA	1501	A	P-O5'	-16.61	1.43	1.59
34	BA	6	C	P-O5'	-16.58	1.43	1.59
34	BA	86	A	N9-C4	-16.58	1.27	1.37
34	BA	422	C	P-O5'	-16.57	1.43	1.59
34	BA	1656	A	N9-C4	-16.57	1.27	1.37
34	BA	20	A	N7-C5	-16.53	1.29	1.39
34	BA	605	G	N9-C4	16.53	1.51	1.38
35	BB	1130	U	P-O5'	-16.48	1.43	1.59
35	BB	33	A	N9-C4	-16.48	1.27	1.37
34	BA	900	A	N9-C4	-16.46	1.27	1.37
85	AA	19	A	N9-C4	-16.44	1.27	1.37
36	BC	41	A	N9-C4	-16.41	1.28	1.37
34	BA	45	A	N9-C4	-16.38	1.28	1.37
85	AA	794	A	N9-C4	-16.37	1.28	1.37
85	AA	1240	A	N9-C4	-16.36	1.28	1.37
36	BC	23	G	P-O5'	-16.36	1.43	1.59
85	AA	1254	A	N9-C4	-16.32	1.28	1.37
38	BE	192	A	N9-C4	-16.32	1.28	1.37
35	BB	1246	C	P-O5'	-16.31	1.43	1.59
85	AA	661	C	P-O5'	-16.31	1.43	1.59
35	BB	1148	U	P-O5'	-16.28	1.43	1.59
35	BB	1399	A	N9-C4	-16.27	1.28	1.37
34	BA	1808	A	P-O5'	-16.23	1.43	1.59
85	AA	483	G	P-O5'	-16.16	1.43	1.59
85	AA	431	G	P-O5'	-16.14	1.43	1.59
34	BA	26	C	P-O5'	-16.11	1.43	1.59
85	AA	316	C	P-O5'	-16.10	1.43	1.59
34	BA	292	C	P-O5'	-16.09	1.43	1.59
36	BC	16	A	N9-C4	-16.09	1.28	1.37
35	BB	399	A	N9-C4	-16.09	1.28	1.37
35	BB	798	A	N7-C5	-16.07	1.29	1.39
34	BA	1260	G	P-O5'	-16.07	1.43	1.59
34	BA	502	U	O3'-P	-16.07	1.41	1.61
34	BA	1542	A	N9-C4	-16.04	1.28	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	81	C	P-O5'	-16.04	1.43	1.59
34	BA	348	U	P-O5'	-16.03	1.43	1.59
35	BB	423	G	P-O5'	-16.02	1.43	1.59
34	BA	1503	U	O3'-P	-16.00	1.42	1.61
85	AA	674	U	P-O5'	-15.96	1.43	1.59
34	BA	494	A	P-O5'	-15.93	1.43	1.59
41	BH	44	A	N9-C4	-15.91	1.28	1.37
34	BA	82	A	N9-C4	-15.89	1.28	1.37
34	BA	294	C	P-O5'	-15.89	1.43	1.59
34	BA	481	A	N3-C4	-15.88	1.25	1.34
85	AA	867	G	O3'-P	-15.82	1.42	1.61
36	BC	6	G	N9-C4	-15.80	1.25	1.38
40	BG	30	C	P-O5'	-15.77	1.44	1.59
34	BA	1585	A	N9-C4	-15.76	1.28	1.37
35	BB	96	A	N7-C5	-15.74	1.29	1.39
35	BB	1514	G	P-O5'	-15.74	1.44	1.59
34	BA	684	G	C5'-C4'	15.73	1.70	1.51
40	BG	142	A	N9-C4	-15.72	1.28	1.37
85	AA	1216	A	P-O5'	-15.72	1.44	1.59
36	BC	18	G	N9-C4	-15.70	1.25	1.38
34	BA	1604	A	N9-C4	-15.69	1.28	1.37
40	BG	157	A	N7-C5	-15.67	1.29	1.39
35	BB	79	U	P-O5'	-15.64	1.44	1.59
35	BB	1220	A	N9-C4	-15.62	1.28	1.37
34	BA	1845	G	N9-C4	-15.61	1.25	1.38
35	BB	468	U	P-O5'	-15.61	1.44	1.59
34	BA	96	G	P-O5'	-15.60	1.44	1.59
41	BH	126	C	P-O5'	-15.59	1.44	1.59
85	AA	114	C	P-O5'	-15.59	1.44	1.59
34	BA	398	G	N9-C4	-15.58	1.25	1.38
85	AA	1495	G	P-O5'	-15.58	1.44	1.59
34	BA	1721	U	C4'-C3'	-15.57	1.36	1.53
35	BB	1226	G	N9-C4	-15.57	1.25	1.38
34	BA	1620	U	P-O5'	-15.55	1.44	1.59
36	BC	25	C	P-O5'	-15.55	1.44	1.59
85	AA	635	G	P-O5'	-15.53	1.44	1.59
34	BA	401	A	N9-C4	-15.50	1.28	1.37
35	BB	657	A	P-O5'	-15.49	1.44	1.59
35	BB	993	A	P-O5'	-15.47	1.44	1.59
41	BH	20	A	P-O5'	-15.47	1.44	1.59
37	BD	75	G	N9-C4	-15.46	1.25	1.38
40	BG	35	G	P-O5'	-15.45	1.44	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	447	U	P-O5'	-15.45	1.44	1.59
40	BG	81	G	P-O5'	-15.42	1.44	1.59
34	BA	335	C	P-O5'	-15.41	1.44	1.59
85	AA	937	G	P-O5'	-15.41	1.44	1.59
35	BB	1196	A	N9-C4	-15.39	1.28	1.37
34	BA	740	A	C2'-C1'	-15.39	1.36	1.53
34	BA	417	A	N9-C4	-15.38	1.28	1.37
35	BB	36	U	P-O5'	-15.37	1.44	1.59
34	BA	881	C	P-O5'	-15.37	1.44	1.59
85	AA	527	A	P-O5'	-15.37	1.44	1.59
34	BA	68	A	N9-C4	-15.34	1.28	1.37
85	AA	363	A	N9-C4	-15.33	1.28	1.37
38	BE	6	A	N9-C4	-15.33	1.28	1.37
35	BB	45	A	N9-C4	-15.32	1.28	1.37
35	BB	634	A	P-O5'	-15.32	1.44	1.59
37	BD	14	C	P-O5'	-15.31	1.44	1.59
34	BA	372	U	P-O5'	-15.31	1.44	1.59
85	AA	514	U	P-O5'	-15.27	1.44	1.59
35	BB	701	U	P-O5'	-15.26	1.44	1.59
85	AA	1459	C	P-O5'	-15.26	1.44	1.59
34	BA	1577	U	P-O5'	-15.25	1.44	1.59
34	BA	462	C	P-O5'	-15.24	1.44	1.59
35	BB	837	A	P-O5'	-15.23	1.44	1.59
35	BB	1360	A	P-O5'	-15.23	1.44	1.59
85	AA	699	U	O3'-P	-15.23	1.42	1.61
35	BB	1138	A	C4'-C3'	-15.23	1.36	1.53
85	AA	521	A	N9-C4	-15.22	1.28	1.37
34	BA	906	A	N9-C4	-15.22	1.28	1.37
35	BB	1486	C	P-O5'	-15.22	1.44	1.59
85	AA	443	A	O3'-P	-15.20	1.43	1.61
85	AA	102	A	N9-C4	-15.20	1.28	1.37
35	BB	483	C	P-O5'	-15.19	1.44	1.59
35	BB	1431	G	P-O5'	-15.19	1.44	1.59
34	BA	121	A	N9-C4	-15.18	1.28	1.37
85	AA	557	G	P-O5'	-15.16	1.44	1.59
36	BC	28	C	N1-C2	-15.16	1.25	1.40
34	BA	761	U	C2-N3	-15.13	1.27	1.37
85	AA	1106	A	C4'-C3'	-15.13	1.36	1.53
35	BB	802	G	P-O5'	-15.09	1.44	1.59
34	BA	481	A	P-O5'	-15.08	1.44	1.59
35	BB	1206	G	P-O5'	-15.08	1.44	1.59
38	BE	128	G	P-O5'	-15.08	1.44	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	29	U	P-O5'	-15.07	1.44	1.59
34	BA	843	G	P-O5'	-15.06	1.44	1.59
85	AA	1840	C	P-O5'	-15.04	1.44	1.59
40	BG	140	G	P-O5'	-15.03	1.44	1.59
34	BA	344	G	O3'-P	-15.02	1.43	1.61
34	BA	1277	G	P-O5'	-15.02	1.44	1.59
34	BA	21	C	P-O5'	-15.02	1.44	1.59
40	BG	28	A	P-O5'	-15.02	1.44	1.59
34	BA	1619	U	P-O5'	-14.99	1.44	1.59
34	BA	130	U	P-O5'	-14.98	1.44	1.59
34	BA	1267	A	P-O5'	-14.98	1.44	1.59
34	BA	1615	A	N9-C4	-14.97	1.28	1.37
34	BA	1720	U	C2-N3	-14.97	1.27	1.37
85	AA	1182	A	N9-C4	-14.96	1.28	1.37
41	BH	111	U	P-O5'	-14.95	1.44	1.59
34	BA	1146	U	P-O5'	-14.95	1.44	1.59
35	BB	971	A	N9-C4	-14.93	1.28	1.37
85	AA	424	A	C3'-C2'	-14.93	1.36	1.52
85	AA	2147	A	N9-C4	-14.93	1.28	1.37
85	AA	706	U	P-O5'	-14.92	1.44	1.59
34	BA	894	G	N9-C4	-14.89	1.26	1.38
41	BH	107	A	N9-C4	-14.89	1.28	1.37
35	BB	1515	C	P-O5'	-14.88	1.44	1.59
35	BB	1209	A	N9-C4	-14.87	1.28	1.37
34	BA	1275	G	O3'-P	-14.87	1.43	1.61
35	BB	456	A	N9-C4	-14.86	1.28	1.37
35	BB	630	A	P-O5'	-14.85	1.44	1.59
85	AA	1521	U	P-O5'	-14.85	1.45	1.59
34	BA	196	A	O3'-P	-14.84	1.43	1.61
35	BB	505	G	P-O5'	-14.83	1.45	1.59
85	AA	986	U	P-O5'	-14.80	1.45	1.59
38	BE	31	A	P-O5'	-14.79	1.45	1.59
34	BA	12	G	N9-C4	-14.79	1.26	1.38
85	AA	644	A	N9-C4	-14.78	1.28	1.37
34	BA	530	A	N9-C4	-14.77	1.28	1.37
34	BA	946	A	N9-C4	-14.75	1.28	1.37
34	BA	212	A	P-O5'	-14.75	1.45	1.59
35	BB	621	C	P-O5'	-14.74	1.45	1.59
38	BE	119	U	P-O5'	-14.72	1.45	1.59
34	BA	67	A	N9-C4	-14.71	1.29	1.37
85	AA	2140	U	P-O5'	-14.71	1.45	1.59
40	BG	22	G	O3'-P	-14.69	1.43	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	806	G	P-O5'	-14.69	1.45	1.59
34	BA	1485	U	P-O5'	-14.68	1.45	1.59
34	BA	1647	G	P-O5'	-14.68	1.45	1.59
85	AA	1219	A	N9-C4	-14.67	1.29	1.37
35	BB	1483	A	P-O5'	-14.65	1.45	1.59
34	BA	897	U	P-O5'	-14.65	1.45	1.59
85	AA	1469	G	O3'-P	-14.64	1.43	1.61
34	BA	759	A	P-O5'	-14.64	1.45	1.59
34	BA	1706	A	N9-C4	-14.64	1.29	1.37
34	BA	30	A	P-O5'	-14.62	1.45	1.59
40	BG	20	U	P-O5'	-14.62	1.45	1.59
34	BA	547	C	C5'-C4'	14.62	1.68	1.51
34	BA	547	C	O3'-P	14.61	1.78	1.61
35	BB	784	C	P-O5'	-14.61	1.45	1.59
85	AA	1996	A	N9-C4	-14.60	1.29	1.37
34	BA	1709	A	O3'-P	-14.59	1.43	1.61
35	BB	1234	G	P-O5'	-14.59	1.45	1.59
34	BA	141	G	N3-C4	-14.57	1.25	1.35
34	BA	241	U	P-O5'	-14.57	1.45	1.59
34	BA	888	G	P-O5'	-14.56	1.45	1.59
35	BB	1052	G	O3'-P	-14.56	1.43	1.61
35	BB	1109	A	P-O5'	-14.55	1.45	1.59
34	BA	207	A	N9-C4	-14.54	1.29	1.37
34	BA	297	A	P-O5'	-14.54	1.45	1.59
40	BG	34	A	N9-C4	-14.54	1.29	1.37
35	BB	835	C	P-O5'	-14.53	1.45	1.59
37	BD	51	G	P-O5'	-14.53	1.45	1.59
34	BA	389	U	P-O5'	-14.53	1.45	1.59
34	BA	1204	U	O3'-P	-14.51	1.43	1.61
85	AA	2120	C	P-O5'	-14.51	1.45	1.59
41	BH	26	C	P-O5'	-14.50	1.45	1.59
34	BA	166	G	P-O5'	-14.50	1.45	1.59
37	BD	19	C	P-O5'	-14.50	1.45	1.59
34	BA	1675	C	P-O5'	-14.49	1.45	1.59
34	BA	1176	C	P-O5'	-14.48	1.45	1.59
35	BB	5	A	C6-N6	-14.48	1.22	1.33
37	BD	35	C	P-O5'	-14.48	1.45	1.59
85	AA	2142	A	N9-C4	-14.48	1.29	1.37
85	AA	2174	G	N9-C4	-14.46	1.26	1.38
34	BA	591	G	P-O5'	-14.46	1.45	1.59
34	BA	1039	G	P-O5'	-14.46	1.45	1.59
34	BA	1148	U	P-O5'	-14.44	1.45	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	169	A	N9-C4	-14.42	1.29	1.37
34	BA	281	C	P-O5'	-14.41	1.45	1.59
34	BA	804	G	C6-N1	-14.41	1.29	1.39
85	AA	731	U	P-O5'	-14.41	1.45	1.59
35	BB	666	A	N9-C4	-14.40	1.29	1.37
85	AA	413	G	P-O5'	-14.39	1.45	1.59
85	AA	862	U	O3'-P	-14.39	1.43	1.61
34	BA	1798	G	P-O5'	-14.38	1.45	1.59
34	BA	299	C	P-O5'	-14.37	1.45	1.59
35	BB	687	C	P-O5'	-14.37	1.45	1.59
34	BA	57	A	N7-C5	-14.37	1.30	1.39
85	AA	887	A	N9-C4	-14.37	1.29	1.37
34	BA	957	A	O3'-P	-14.36	1.44	1.61
34	BA	1143	U	P-O5'	-14.36	1.45	1.59
34	BA	1412	G	C6-N1	-14.36	1.29	1.39
34	BA	500	C	O3'-P	-14.35	1.44	1.61
34	BA	1011	G	N9-C4	-14.35	1.26	1.38
40	BG	51	U	P-O5'	-14.34	1.45	1.59
35	BB	430	A	N9-C4	-14.34	1.29	1.37
34	BA	1563	G	P-O5'	-14.34	1.45	1.59
38	BE	101	C	O3'-P	-14.34	1.44	1.61
85	AA	385	A	N9-C4	-14.33	1.29	1.37
36	BC	117	A	N9-C4	-14.32	1.29	1.37
37	BD	72	U	O3'-P	-14.32	1.44	1.61
34	BA	401	A	P-O5'	-14.31	1.45	1.59
40	BG	135	C	P-O5'	-14.31	1.45	1.59
36	BC	51	A	N9-C4	-14.30	1.29	1.37
34	BA	1592	U	P-O5'	-14.28	1.45	1.59
85	AA	1666	U	P-O5'	-14.28	1.45	1.59
34	BA	1650	G	C1'-N9	-14.26	1.26	1.46
36	BC	148	C	P-O5'	-14.26	1.45	1.59
34	BA	777	C	P-O5'	-14.26	1.45	1.59
35	BB	21	C	O3'-P	-14.24	1.44	1.61
35	BB	1005	A	P-O5'	-14.24	1.45	1.59
38	BE	194	A	N7-C5	-14.24	1.30	1.39
85	AA	2122	A	N7-C5	-14.23	1.30	1.39
85	AA	991	G	N7-C5	-14.22	1.30	1.39
85	AA	870	U	P-O5'	-14.22	1.45	1.59
35	BB	605	C	O3'-P	-14.22	1.44	1.61
85	AA	414	C	P-O5'	-14.19	1.45	1.59
35	BB	93	A	P-O5'	-14.18	1.45	1.59
85	AA	474	C	P-O5'	-14.17	1.45	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1262	A	N9-C4	-14.16	1.29	1.37
34	BA	665	C	P-O5'	-14.15	1.45	1.59
34	BA	1213	A	P-O5'	-14.15	1.45	1.59
85	AA	1256	C	P-O5'	-14.14	1.45	1.59
35	BB	871	C	P-O5'	-14.14	1.45	1.59
34	BA	1415	C	P-O5'	-14.13	1.45	1.59
35	BB	1449	G	P-O5'	-14.13	1.45	1.59
39	BF	14	C	O3'-P	-14.13	1.44	1.61
34	BA	1438	C	P-O5'	-14.12	1.45	1.59
39	BF	4	A	N9-C4	-14.12	1.29	1.37
35	BB	1452	U	P-O5'	-14.12	1.45	1.59
35	BB	593	A	N9-C4	-14.11	1.29	1.37
38	BE	45	G	P-O5'	-14.12	1.45	1.59
39	BF	11	C	C5'-C4'	14.11	1.68	1.51
85	AA	313	A	N9-C4	-14.11	1.29	1.37
35	BB	1355	C	O3'-P	-14.11	1.44	1.61
85	AA	932	A	N9-C4	-14.11	1.29	1.37
85	AA	586	G	P-O5'	-14.10	1.45	1.59
38	BE	21	C	O3'-P	-14.10	1.44	1.61
85	AA	800	A	N9-C4	-14.10	1.29	1.37
34	BA	1697	U	O3'-P	-14.09	1.44	1.61
36	BC	137	C	P-O5'	-14.08	1.45	1.59
34	BA	296	G	C4'-C3'	14.07	1.68	1.53
34	BA	452	A	N9-C4	-14.07	1.29	1.37
85	AA	427	G	P-O5'	-14.06	1.45	1.59
34	BA	720	A	N9-C4	-14.05	1.29	1.37
35	BB	1404	A	N9-C4	-14.05	1.29	1.37
85	AA	5	U	P-O5'	-14.05	1.45	1.59
85	AA	419	A	N9-C4	-14.05	1.29	1.37
34	BA	921	G	P-O5'	-14.04	1.45	1.59
34	BA	1528	U	P-O5'	-14.04	1.45	1.59
34	BA	1790	U	P-O5'	-14.04	1.45	1.59
39	BF	22	U	P-O5'	-14.04	1.45	1.59
35	BB	577	U	C2-N3	-14.03	1.27	1.37
85	AA	2151	U	P-O5'	-14.03	1.45	1.59
85	AA	492	C	P-O5'	-14.02	1.45	1.59
37	BD	111	U	P-O5'	-14.00	1.45	1.59
85	AA	1465	C	P-O5'	-13.99	1.45	1.59
34	BA	458	G	P-O5'	-13.98	1.45	1.59
34	BA	755	G	N9-C4	-13.98	1.26	1.38
85	AA	440	U	P-O5'	-13.98	1.45	1.59
34	BA	480	G	P-O5'	-13.97	1.45	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	431	U	P-O5'	-13.97	1.45	1.59
34	BA	376	U	C2-N3	-13.96	1.27	1.37
85	AA	484	G	C2'-C1'	-13.96	1.38	1.53
34	BA	104	A	N9-C4	-13.96	1.29	1.37
85	AA	1247	A	O3'-P	-13.93	1.44	1.61
35	BB	491	A	N9-C4	-13.93	1.29	1.37
85	AA	1520	A	N9-C4	-13.93	1.29	1.37
35	BB	1164	U	P-O5'	-13.93	1.45	1.59
34	BA	1049	G	N9-C4	-13.92	1.26	1.38
34	BA	1599	A	O3'-P	-13.92	1.44	1.61
34	BA	983	A	N9-C4	-13.90	1.29	1.37
34	BA	792	A	N9-C4	-13.90	1.29	1.37
39	BF	69	A	N9-C4	-13.90	1.29	1.37
35	BB	604	C	P-O5'	-13.88	1.45	1.59
34	BA	605	G	N7-C5	-13.88	1.30	1.39
38	BE	59	U	P-O5'	-13.88	1.45	1.59
85	AA	2071	U	P-O5'	-13.88	1.45	1.59
37	BD	77	A	P-O5'	-13.88	1.45	1.59
34	BA	1619	U	O3'-P	-13.88	1.44	1.61
34	BA	858	C	P-O5'	-13.87	1.45	1.59
35	BB	766	G	P-O5'	-13.87	1.45	1.59
85	AA	1679	U	P-O5'	-13.87	1.45	1.59
34	BA	12	G	P-O5'	-13.86	1.45	1.59
34	BA	1025	A	N9-C4	-13.86	1.29	1.37
35	BB	1334	C	C2'-C1'	-13.86	1.38	1.53
35	BB	1195	A	P-O5'	-13.85	1.46	1.59
35	BB	80	C	O3'-P	-13.84	1.44	1.61
34	BA	254	U	C2-N3	-13.84	1.28	1.37
35	BB	1243	A	N9-C4	-13.84	1.29	1.37
41	BH	127	A	P-O5'	-13.83	1.46	1.59
85	AA	77	C	P-O5'	-13.83	1.46	1.59
34	BA	484	A	N9-C4	-13.82	1.29	1.37
35	BB	1222	A	N9-C4	-13.81	1.29	1.37
34	BA	793	A	P-O5'	-13.80	1.46	1.59
34	BA	138	C	P-O5'	-13.80	1.46	1.59
38	BE	96	G	P-O5'	-13.80	1.46	1.59
38	BE	193	A	N9-C4	-13.79	1.29	1.37
85	AA	130	G	P-O5'	-13.79	1.46	1.59
85	AA	504	U	O3'-P	-13.78	1.44	1.61
85	AA	932	A	P-O5'	-13.78	1.46	1.59
34	BA	257	G	N9-C4	-13.77	1.26	1.38
85	AA	1463	A	P-O5'	-13.77	1.46	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	762	A	N9-C4	-13.76	1.29	1.37
34	BA	64	A	N9-C4	-13.76	1.29	1.37
85	AA	1539	A	N9-C4	-13.76	1.29	1.37
34	BA	62	A	N9-C4	-13.75	1.29	1.37
34	BA	1558	C	P-O5'	-13.75	1.46	1.59
36	BC	88	A	P-O5'	-13.75	1.46	1.59
40	BG	118	U	P-O5'	-13.74	1.46	1.59
34	BA	1556	A	C1'-N9	-13.74	1.27	1.46
36	BC	24	G	O3'-P	-13.74	1.44	1.61
85	AA	1180	C	O3'-P	-13.73	1.44	1.61
34	BA	1561	C	P-O5'	-13.72	1.46	1.59
40	BG	18	U	P-O5'	-13.72	1.46	1.59
35	BB	5	A	C2'-C1'	-13.71	1.38	1.53
85	AA	2176	U	P-O5'	-13.72	1.46	1.59
41	BH	37	U	O3'-P	-13.69	1.44	1.61
35	BB	552	C	P-O5'	-13.67	1.46	1.59
85	AA	28	A	P-O5'	-13.67	1.46	1.59
35	BB	1033	U	P-O5'	-13.67	1.46	1.59
40	BG	28	A	O3'-P	-13.66	1.44	1.61
34	BA	127	U	P-O5'	-13.66	1.46	1.59
34	BA	774	A	N9-C4	-13.66	1.29	1.37
34	BA	802	G	P-O5'	-13.65	1.46	1.59
34	BA	926	A	P-O5'	-13.63	1.46	1.59
38	BE	27	A	C1'-N9	-13.63	1.27	1.46
35	BB	1181	A	N9-C4	-13.63	1.29	1.37
85	AA	168	A	P-O5'	-13.62	1.46	1.59
34	BA	1097	G	O3'-P	-13.62	1.44	1.61
34	BA	290	G	P-O5'	-13.61	1.46	1.59
34	BA	409	A	N9-C4	-13.61	1.29	1.37
34	BA	680	C	O3'-P	-13.61	1.44	1.61
34	BA	1028	A	P-O5'	-13.61	1.46	1.59
85	AA	692	U	P-O5'	-13.61	1.46	1.59
85	AA	484	G	P-O5'	-13.60	1.46	1.59
34	BA	582	U	C2-N3	-13.59	1.28	1.37
36	BC	156	A	N9-C4	-13.59	1.29	1.37
36	BC	102	G	P-O5'	-13.58	1.46	1.59
35	BB	1252	G	N9-C4	-13.56	1.27	1.38
34	BA	557	U	C2'-C1'	-13.54	1.38	1.53
34	BA	1489	U	P-O5'	-13.54	1.46	1.59
85	AA	508	C	P-O5'	-13.54	1.46	1.59
39	BF	28	C	O3'-P	-13.53	1.45	1.61
34	BA	1637	G	P-O5'	-13.53	1.46	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	692	U	N1-C2	-13.52	1.26	1.38
85	AA	668	A	N9-C4	-13.52	1.29	1.37
34	BA	557	U	N1-C6	-13.52	1.25	1.38
35	BB	1063	C	P-O5'	-13.51	1.46	1.59
85	AA	1226	A	P-O5'	-13.51	1.46	1.59
85	AA	435	A	N9-C4	-13.51	1.29	1.37
34	BA	141	G	C2-N3	-13.50	1.22	1.32
34	BA	214	A	N9-C4	-13.49	1.29	1.37
37	BD	82	G	C3'-C2'	-13.49	1.38	1.52
39	BF	23	G	C2'-C1'	-13.49	1.38	1.53
38	BE	32	U	P-O5'	-13.48	1.46	1.59
34	BA	1666	U	P-O5'	-13.46	1.46	1.59
85	AA	115	U	O3'-P	-13.46	1.45	1.61
35	BB	39	C	P-O5'	-13.46	1.46	1.59
35	BB	8	U	P-O5'	-13.46	1.46	1.59
34	BA	894	G	O3'-P	-13.46	1.45	1.61
34	BA	93	A	O3'-P	-13.45	1.45	1.61
34	BA	1160	U	P-O5'	-13.44	1.46	1.59
34	BA	1247	G	P-O5'	-13.44	1.46	1.59
34	BA	1463	U	P-O5'	-13.44	1.46	1.59
35	BB	613	C	P-O5'	-13.44	1.46	1.59
37	BD	61	C	P-O5'	-13.44	1.46	1.59
85	AA	2188	C	O3'-P	-13.43	1.45	1.61
34	BA	1685	C	P-O5'	-13.43	1.46	1.59
85	AA	455	G	N9-C4	-13.42	1.27	1.38
85	AA	152	A	N7-C5	-13.42	1.31	1.39
35	BB	27	C	O3'-P	-13.42	1.45	1.61
37	BD	4	U	P-O5'	-13.42	1.46	1.59
34	BA	74	A	C2'-C1'	-13.41	1.38	1.53
85	AA	2075	C	P-O5'	-13.40	1.46	1.59
34	BA	236	A	O3'-P	-13.40	1.45	1.61
34	BA	1197	U	P-O5'	-13.39	1.46	1.59
38	BE	133	C	O3'-P	-13.39	1.45	1.61
34	BA	741	A	N9-C4	-13.39	1.29	1.37
34	BA	182	U	P-O5'	-13.38	1.46	1.59
34	BA	65	A	C2'-C1'	-13.38	1.38	1.53
34	BA	206	C	C5'-C4'	13.37	1.67	1.51
35	BB	1102	U	O3'-P	-13.38	1.45	1.61
85	AA	521	A	N7-C5	-13.37	1.31	1.39
85	AA	2127	G	N9-C4	-13.37	1.27	1.38
35	BB	1282	G	P-O5'	-13.37	1.46	1.59
34	BA	330	A	N9-C4	-13.36	1.29	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	788	C	C2'-C1'	-13.36	1.38	1.53
85	AA	1456	A	N9-C4	-13.36	1.29	1.37
34	BA	1031	U	P-O5'	-13.35	1.46	1.59
85	AA	365	G	C2'-C1'	-13.35	1.38	1.53
35	BB	999	G	P-O5'	-13.34	1.46	1.59
85	AA	511	A	P-O5'	-13.34	1.46	1.59
34	BA	982	A	N9-C4	-13.33	1.29	1.37
34	BA	1217	A	N9-C4	-13.33	1.29	1.37
35	BB	549	U	O3'-P	-13.33	1.45	1.61
35	BB	1425	A	N9-C4	-13.33	1.29	1.37
85	AA	982	G	N9-C4	-13.33	1.27	1.38
34	BA	1202	G	P-O5'	-13.32	1.46	1.59
35	BB	1327	U	O3'-P	-13.32	1.45	1.61
34	BA	1500	G	N9-C4	-13.32	1.48	1.38
85	AA	1116	G	P-O5'	-13.32	1.46	1.59
34	BA	922	C	O3'-P	-13.32	1.45	1.61
34	BA	1160	U	O3'-P	-13.32	1.45	1.61
34	BA	1796	A	N9-C4	-13.31	1.29	1.37
85	AA	1511	C	P-O5'	-13.31	1.46	1.59
85	AA	1230	U	P-O5'	-13.31	1.46	1.59
34	BA	891	C	P-O5'	-13.31	1.46	1.59
38	BE	162	U	P-O5'	-13.30	1.46	1.59
34	BA	1598	U	P-O5'	-13.30	1.46	1.59
85	AA	1471	G	P-O5'	-13.30	1.46	1.59
85	AA	1708	A	N9-C4	-13.29	1.29	1.37
35	BB	1385	C	O3'-P	-13.29	1.45	1.61
85	AA	2131	C	P-O5'	-13.28	1.46	1.59
34	BA	1550	G	P-O5'	-13.27	1.46	1.59
34	BA	1564	A	P-O5'	-13.27	1.46	1.59
85	AA	53	G	P-O5'	-13.26	1.46	1.59
35	BB	1426	G	C6-N1	-13.26	1.30	1.39
37	BD	106	G	P-O5'	-13.25	1.46	1.59
85	AA	862	U	P-O5'	-13.25	1.46	1.59
34	BA	1466	U	C2-N3	-13.24	1.28	1.37
34	BA	1520	A	O3'-P	-13.24	1.45	1.61
35	BB	697	G	P-O5'	-13.24	1.46	1.59
35	BB	56	U	P-O5'	-13.24	1.46	1.59
85	AA	2196	G	P-O5'	-13.24	1.46	1.59
85	AA	1525	C	P-O5'	-13.24	1.46	1.59
36	BC	31	A	C1'-N9	-13.23	1.28	1.46
35	BB	690	C	P-O5'	-13.22	1.46	1.59
85	AA	755	G	P-O5'	-13.22	1.46	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	525	C	C2'-C1'	-13.22	1.38	1.53
34	BA	4	A	N9-C4	-13.21	1.29	1.37
34	BA	421	G	P-O5'	-13.21	1.46	1.59
34	BA	1636	C	O3'-P	-13.21	1.45	1.61
35	BB	1025	A	C5'-C4'	13.21	1.67	1.51
34	BA	1409	A	N9-C4	-13.18	1.29	1.37
41	BH	33	G	O3'-P	-13.18	1.45	1.61
85	AA	169	G	P-O5'	-13.17	1.46	1.59
35	BB	1123	A	P-O5'	-13.17	1.46	1.59
85	AA	939	A	N7-C5	-13.16	1.31	1.39
85	AA	1831	U	P-O5'	-13.15	1.46	1.59
39	BF	32	G	O3'-P	-13.15	1.45	1.61
35	BB	778	A	O3'-P	-13.15	1.45	1.61
34	BA	1462	U	O3'-P	-13.15	1.45	1.61
85	AA	1669	G	P-O5'	-13.15	1.46	1.59
39	BF	58	U	P-O5'	-13.14	1.46	1.59
35	BB	121	A	P-O5'	-13.13	1.46	1.59
35	BB	648	G	O3'-P	-13.13	1.45	1.61
85	AA	100	A	N9-C4	-13.13	1.29	1.37
34	BA	1422	A	N9-C4	-13.12	1.29	1.37
85	AA	923	A	O3'-P	-13.12	1.45	1.61
35	BB	81	A	P-O5'	-13.12	1.46	1.59
34	BA	590	U	P-O5'	-13.11	1.46	1.59
34	BA	1675	C	O3'-P	-13.11	1.45	1.61
37	BD	93	G	P-O5'	-13.11	1.46	1.59
38	BE	94	U	O3'-P	-13.11	1.45	1.61
34	BA	1658	G	P-O5'	-13.11	1.46	1.59
34	BA	280	A	P-O5'	-13.10	1.46	1.59
35	BB	1334	C	O3'-P	-13.10	1.45	1.61
34	BA	1804	A	N7-C5	-13.09	1.31	1.39
35	BB	1440	A	N9-C4	-13.09	1.29	1.37
34	BA	1809	G	P-O5'	-13.08	1.46	1.59
34	BA	221	G	P-O5'	-13.08	1.46	1.59
40	BG	38	A	N9-C4	-13.07	1.30	1.37
37	BD	101	A	N9-C4	-13.06	1.30	1.37
34	BA	228	A	N9-C4	13.06	1.45	1.37
35	BB	21	C	P-O5'	-13.06	1.46	1.59
85	AA	69	C	C2'-C1'	-13.06	1.39	1.53
41	BH	36	C	P-O5'	-13.06	1.46	1.59
39	BF	31	U	O3'-P	-13.06	1.45	1.61
85	AA	960	G	N7-C5	-13.06	1.31	1.39
35	BB	1445	A	N9-C4	-13.05	1.30	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1699	A	P-O5'	-13.05	1.46	1.59
35	BB	1339	C	P-O5'	-13.04	1.46	1.59
85	AA	1462	A	C2'-C1'	-13.04	1.39	1.53
34	BA	1284	G	C6-N1	-13.04	1.30	1.39
34	BA	1545	C	P-O5'	-13.03	1.46	1.59
35	BB	101	U	P-O5'	-13.03	1.46	1.59
34	BA	1315	C	O3'-P	-13.03	1.45	1.61
35	BB	512	C	P-O5'	-13.02	1.46	1.59
34	BA	1503	U	C2'-C1'	-13.01	1.39	1.53
34	BA	1475	G	O3'-P	-13.01	1.45	1.61
34	BA	1262	A	N9-C4	-13.01	1.30	1.37
36	BC	98	C	P-O5'	-13.01	1.46	1.59
85	AA	100	A	P-O5'	-13.01	1.46	1.59
85	AA	1703	A	P-O5'	-13.00	1.46	1.59
36	BC	108	A	P-O5'	-13.00	1.46	1.59
85	AA	390	U	C2'-C1'	-12.99	1.39	1.53
37	BD	8	A	P-O5'	-12.99	1.46	1.59
39	BF	56	C	P-O5'	-12.99	1.46	1.59
34	BA	969	A	N9-C4	-12.99	1.30	1.37
34	BA	1710	C	P-O5'	-12.99	1.46	1.59
36	BC	124	A	N9-C4	12.98	1.45	1.37
85	AA	1499	G	P-O5'	-12.98	1.46	1.59
34	BA	696	A	O3'-P	-12.97	1.45	1.61
85	AA	699	U	P-O5'	-12.97	1.46	1.59
35	BB	492	U	P-O5'	-12.97	1.46	1.59
85	AA	2107	C	P-O5'	-12.97	1.46	1.59
85	AA	1454	U	P-O5'	-12.96	1.46	1.59
34	BA	1518	A	N9-C4	-12.96	1.30	1.37
35	BB	50	A	P-O5'	-12.95	1.46	1.59
35	BB	458	U	P-O5'	-12.94	1.46	1.59
39	BF	73	U	P-O5'	-12.94	1.46	1.59
40	BG	105	A	P-O5'	-12.94	1.46	1.59
41	BH	45	G	P-O5'	-12.94	1.46	1.59
35	BB	1322	A	N9-C4	-12.93	1.30	1.37
34	BA	798	G	N9-C4	-12.92	1.27	1.38
85	AA	323	U	P-O5'	-12.92	1.46	1.59
35	BB	586	U	P-O5'	-12.91	1.46	1.59
85	AA	49	C	P-O5'	-12.91	1.46	1.59
36	BC	100	U	P-O5'	-12.91	1.46	1.59
85	AA	1471	G	O3'-P	-12.91	1.45	1.61
34	BA	801	U	P-O5'	-12.90	1.46	1.59
85	AA	1221	G	N9-C4	-12.90	1.27	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	98	A	N9-C4	-12.89	1.30	1.37
34	BA	809	U	C2-N3	-12.89	1.28	1.37
34	BA	906	A	P-O5'	-12.89	1.46	1.59
35	BB	1134	G	P-O5'	-12.89	1.46	1.59
34	BA	1728	G	O3'-P	-12.88	1.45	1.61
34	BA	1789	A	O3'-P	-12.88	1.45	1.61
35	BB	638	G	P-O5'	-12.88	1.46	1.59
40	BG	109	C	P-O5'	-12.87	1.46	1.59
34	BA	105	U	P-O5'	-12.87	1.46	1.59
34	BA	1689	U	O3'-P	-12.87	1.45	1.61
34	BA	297	A	O3'-P	-12.86	1.45	1.61
34	BA	369	A	O3'-P	-12.86	1.45	1.61
36	BC	42	G	P-O5'	-12.86	1.46	1.59
85	AA	2127	G	C2'-C1'	-12.86	1.39	1.53
35	BB	1220	A	O3'-P	-12.85	1.45	1.61
35	BB	778	A	P-O5'	-12.85	1.47	1.59
41	BH	23	G	O3'-P	-12.84	1.45	1.61
34	BA	10	G	P-O5'	-12.84	1.47	1.59
85	AA	2132	A	O3'-P	-12.84	1.45	1.61
85	AA	1197	U	P-O5'	-12.83	1.47	1.59
35	BB	1156	U	P-O5'	-12.82	1.47	1.59
38	BE	201	A	N9-C4	-12.82	1.30	1.37
85	AA	889	G	P-O5'	-12.82	1.47	1.59
35	BB	1422	G	N9-C4	-12.82	1.27	1.38
34	BA	326	A	C4'-C3'	-12.82	1.39	1.53
85	AA	1924	C	P-O5'	-12.82	1.47	1.59
85	AA	1124	G	C3'-C2'	-12.81	1.38	1.52
85	AA	1114	A	N9-C4	-12.81	1.30	1.37
34	BA	264	A	N9-C4	-12.81	1.30	1.37
34	BA	1668	C	P-O5'	-12.80	1.47	1.59
34	BA	1609	U	P-O5'	-12.80	1.47	1.59
37	BD	34	C	P-O5'	-12.80	1.47	1.59
34	BA	1329	U	P-O5'	-12.79	1.47	1.59
40	BG	103	C	P-O5'	-12.78	1.47	1.59
85	AA	1538	C	O3'-P	-12.78	1.45	1.61
34	BA	1222	C	O3'-P	-12.77	1.45	1.61
85	AA	2182	A	N9-C4	-12.77	1.30	1.37
85	AA	935	A	P-O5'	-12.73	1.47	1.59
34	BA	1323	G	P-O5'	-12.73	1.47	1.59
40	BG	49	A	O3'-P	-12.73	1.45	1.61
85	AA	1206	A	O3'-P	-12.73	1.45	1.61
40	BG	156	G	N9-C4	-12.72	1.27	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	91	G	P-O5'	-12.72	1.47	1.59
35	BB	832	C	P-O5'	-12.72	1.47	1.59
35	BB	1484	A	N7-C5	-12.72	1.31	1.39
34	BA	1564	A	N9-C4	-12.71	1.30	1.37
36	BC	17	U	P-O5'	-12.71	1.47	1.59
34	BA	896	U	C3'-C2'	12.71	1.67	1.52
35	BB	1254	G	N9-C4	-12.71	1.27	1.38
35	BB	1260	A	N9-C4	-12.71	1.30	1.37
36	BC	26	U	N1-C2	-12.71	1.27	1.38
85	AA	251	A	P-O5'	-12.70	1.47	1.59
34	BA	774	A	C2'-C1'	-12.70	1.39	1.53
35	BB	782	A	P-O5'	-12.70	1.47	1.59
85	AA	152	A	N9-C4	-12.70	1.30	1.37
34	BA	1578	A	N9-C4	-12.69	1.30	1.37
85	AA	2054	G	P-O5'	-12.69	1.47	1.59
34	BA	1556	A	P-O5'	-12.69	1.47	1.59
41	BH	33	G	N9-C4	-12.69	1.27	1.38
36	BC	54	G	C2'-C1'	-12.68	1.39	1.53
38	BE	48	G	P-O5'	-12.68	1.47	1.59
40	BG	139	U	O3'-P	-12.67	1.46	1.61
85	AA	420	C	P-O5'	-12.67	1.47	1.59
85	AA	424	A	N9-C4	-12.67	1.30	1.37
41	BH	6	U	P-O5'	-12.66	1.47	1.59
34	BA	1804	A	N9-C4	-12.65	1.30	1.37
35	BB	580	A	N9-C4	-12.65	1.30	1.37
34	BA	972	C	P-O5'	-12.65	1.47	1.59
37	BD	90	A	N9-C4	-12.65	1.30	1.37
35	BB	1155	U	C2-N3	-12.65	1.28	1.37
34	BA	874	G	O3'-P	-12.64	1.46	1.61
34	BA	1461	A	P-O5'	-12.64	1.47	1.59
36	BC	116	C	P-O5'	-12.64	1.47	1.59
85	AA	967	C	P-O5'	-12.64	1.47	1.59
41	BH	108	U	O3'-P	-12.63	1.46	1.61
34	BA	1499	A	P-O5'	-12.63	1.47	1.59
85	AA	1104	G	P-O5'	-12.63	1.47	1.59
35	BB	451	A	N9-C4	-12.62	1.30	1.37
34	BA	8	G	P-O5'	-12.62	1.47	1.59
37	BD	25	G	P-O5'	-12.61	1.47	1.59
34	BA	748	C	P-O5'	-12.61	1.47	1.59
35	BB	1259	A	N9-C4	-12.61	1.30	1.37
34	BA	1786	C	P-O5'	-12.60	1.47	1.59
85	AA	107	A	N9-C4	-12.60	1.30	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	758	G	O3'-P	-12.60	1.46	1.61
34	BA	1591	G	P-O5'	-12.60	1.47	1.59
34	BA	209	A	N7-C5	-12.59	1.31	1.39
85	AA	479	C	P-O5'	-12.59	1.47	1.59
35	BB	1168	G	P-O5'	-12.59	1.47	1.59
85	AA	2143	U	P-O5'	-12.59	1.47	1.59
34	BA	736	G	N9-C4	-12.59	1.27	1.38
34	BA	1709	A	N9-C4	-12.58	1.30	1.37
35	BB	795	A	N9-C4	-12.58	1.30	1.37
35	BB	1147	G	O3'-P	-12.58	1.46	1.61
85	AA	411	U	O3'-P	-12.58	1.46	1.61
34	BA	1179	U	O3'-P	-12.57	1.46	1.61
38	BE	177	U	C2-N3	-12.57	1.28	1.37
35	BB	1211	C	P-O5'	-12.57	1.47	1.59
37	BD	74	A	N9-C4	-12.56	1.30	1.37
35	BB	1196	A	O3'-P	-12.56	1.46	1.61
34	BA	56	G	C1'-N9	-12.55	1.29	1.46
35	BB	1320	U	P-O5'	-12.55	1.47	1.59
35	BB	622	G	N7-C5	-12.55	1.31	1.39
39	BF	9	C	P-O5'	-12.55	1.47	1.59
35	BB	786	A	N9-C4	-12.55	1.30	1.37
34	BA	1189	A	P-O5'	-12.55	1.47	1.59
34	BA	1064	A	N9-C4	-12.54	1.30	1.37
35	BB	1201	G	C4'-C3'	12.54	1.67	1.53
37	BD	99	G	N9-C4	-12.54	1.27	1.38
35	BB	93	A	N9-C4	-12.54	1.30	1.37
34	BA	1321	A	N9-C4	-12.52	1.30	1.37
35	BB	1376	G	P-O5'	-12.52	1.47	1.59
34	BA	397	A	O3'-P	-12.52	1.46	1.61
37	BD	48	G	C1'-N9	-12.52	1.29	1.46
85	AA	2145	G	C2'-C1'	-12.52	1.39	1.53
34	BA	323	C	C2'-C1'	-12.52	1.39	1.53
40	BG	22	G	P-O5'	-12.51	1.47	1.59
85	AA	687	G	P-O5'	-12.51	1.47	1.59
85	AA	1153	G	N7-C5	-12.51	1.31	1.39
86	AB	68	C	P-O5'	-12.50	1.47	1.59
34	BA	705	C	P-O5'	-12.50	1.47	1.59
35	BB	1489	A	N9-C4	-12.50	1.30	1.37
38	BE	23	G	P-O5'	-12.50	1.47	1.59
38	BE	201	A	P-O5'	-12.50	1.47	1.59
35	BB	788	U	C2-N3	-12.50	1.29	1.37
34	BA	606	G	O3'-P	-12.49	1.46	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	939	C	P-O5'	-12.49	1.47	1.59
35	BB	122	U	C2-N3	-12.49	1.29	1.37
35	BB	1195	A	N9-C4	-12.49	1.30	1.37
36	BC	107	C	P-O5'	-12.49	1.47	1.59
38	BE	18	U	P-O5'	-12.49	1.47	1.59
85	AA	2096	G	P-O5'	-12.49	1.47	1.59
34	BA	1418	G	P-O5'	-12.48	1.47	1.59
34	BA	1639	U	P-O5'	-12.48	1.47	1.59
35	BB	1024	G	P-O5'	-12.47	1.47	1.59
85	AA	1516	A	O3'-P	-12.47	1.46	1.61
85	AA	1871	U	P-O5'	-12.47	1.47	1.59
36	BC	17	U	N1-C2	-12.47	1.27	1.38
34	BA	1003	A	P-O5'	-12.46	1.47	1.59
85	AA	505	U	P-O5'	-12.46	1.47	1.59
34	BA	1599	A	P-O5'	-12.46	1.47	1.59
85	AA	422	G	P-O5'	-12.46	1.47	1.59
34	BA	474	A	N9-C4	-12.46	1.30	1.37
34	BA	1652	G	C1'-N9	-12.46	1.29	1.46
34	BA	93	A	N9-C4	-12.45	1.30	1.37
85	AA	450	A	N9-C4	-12.45	1.30	1.37
85	AA	1235	G	P-O5'	-12.45	1.47	1.59
40	BG	16	G	C2'-C1'	-12.44	1.39	1.53
34	BA	138	C	O3'-P	-12.44	1.46	1.61
35	BB	824	C	P-O5'	-12.44	1.47	1.59
36	BC	3	C	P-O5'	-12.44	1.47	1.59
85	AA	450	A	P-O5'	-12.44	1.47	1.59
34	BA	1656	A	P-O5'	-12.44	1.47	1.59
35	BB	17	U	P-O5'	-12.44	1.47	1.59
85	AA	519	A	N9-C4	-12.44	1.30	1.37
85	AA	99	U	C2'-C1'	-12.43	1.39	1.53
40	BG	123	C	P-O5'	-12.43	1.47	1.59
85	AA	591	A	N9-C4	-12.43	1.30	1.37
34	BA	618	G	P-O5'	-12.43	1.47	1.59
34	BA	988	U	P-O5'	-12.43	1.47	1.59
41	BH	33	G	C2'-C1'	-12.42	1.39	1.53
85	AA	455	G	O3'-P	-12.42	1.46	1.61
35	BB	18	A	N9-C4	-12.42	1.30	1.37
34	BA	1065	U	O3'-P	-12.42	1.46	1.61
34	BA	1095	G	P-O5'	-12.42	1.47	1.59
36	BC	164	G	O3'-P	-12.42	1.46	1.61
38	BE	111	C	O3'-P	-12.41	1.46	1.61
35	BB	649	A	O3'-P	-12.41	1.46	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	238	C	P-O5'	-12.40	1.47	1.59
34	BA	97	A	N7-C5	-12.40	1.31	1.39
34	BA	1429	A	P-O5'	-12.40	1.47	1.59
36	BC	54	G	N9-C4	-12.40	1.28	1.38
40	BG	33	G	C1'-N9	-12.40	1.29	1.46
85	AA	364	C	P-O5'	-12.39	1.47	1.59
85	AA	1516	A	C1'-N9	-12.38	1.29	1.46
35	BB	452	A	P-O5'	-12.37	1.47	1.59
85	AA	76	G	P-O5'	-12.38	1.47	1.59
34	BA	615	A	N9-C4	-12.37	1.30	1.37
34	BA	732	A	N9-C4	-12.37	1.30	1.37
34	BA	1192	A	O3'-P	-12.37	1.46	1.61
34	BA	4	A	O3'-P	-12.37	1.46	1.61
34	BA	1543	A	N9-C4	-12.37	1.30	1.37
85	AA	2179	C	P-O5'	-12.37	1.47	1.59
38	BE	185	G	P-O5'	-12.36	1.47	1.59
85	AA	2008	G	N9-C4	-12.36	1.28	1.38
34	BA	423	G	N9-C4	-12.36	1.28	1.38
35	BB	1026	G	C2'-C1'	-12.36	1.39	1.53
34	BA	1806	A	N9-C4	-12.36	1.30	1.37
85	AA	456	A	C1'-N9	-12.35	1.29	1.46
35	BB	40	C	P-O5'	-12.35	1.47	1.59
85	AA	929	G	P-O5'	-12.35	1.47	1.59
34	BA	426	A	N9-C4	-12.35	1.30	1.37
85	AA	1484	G	O3'-P	-12.35	1.46	1.61
35	BB	386	G	P-O5'	-12.35	1.47	1.59
41	BH	112	U	O3'-P	-12.35	1.46	1.61
34	BA	142	A	O3'-P	-12.34	1.46	1.61
34	BA	234	A	P-O5'	-12.34	1.47	1.59
35	BB	1138	A	C2'-C1'	-12.34	1.39	1.53
40	BG	182	G	N7-C5	-12.34	1.31	1.39
85	AA	1099	U	P-O5'	-12.34	1.47	1.59
34	BA	136	A	P-O5'	-12.34	1.47	1.59
34	BA	20	A	N9-C4	-12.34	1.30	1.37
85	AA	899	A	C2'-C1'	-12.34	1.39	1.53
34	BA	185	A	P-O5'	-12.33	1.47	1.59
34	BA	457	A	N9-C4	-12.33	1.30	1.37
34	BA	609	G	P-O5'	-12.33	1.47	1.59
34	BA	547	C	C3'-O3'	12.33	1.59	1.42
35	BB	1127	A	N9-C4	-12.33	1.30	1.37
36	BC	37	U	O3'-P	-12.33	1.46	1.61
38	BE	27	A	P-O5'	-12.33	1.47	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	132	G	P-O5'	-12.32	1.47	1.59
85	AA	976	G	P-O5'	-12.32	1.47	1.59
35	BB	1169	A	P-O5'	-12.31	1.47	1.59
34	BA	765	U	C5'-C4'	12.31	1.66	1.51
34	BA	1211	G	N9-C4	-12.31	1.28	1.38
39	BF	35	C	P-O5'	-12.31	1.47	1.59
85	AA	400	G	N9-C4	-12.31	1.28	1.38
35	BB	653	G	P-O5'	-12.30	1.47	1.59
35	BB	1401	G	N9-C4	-12.30	1.28	1.38
41	BH	109	G	C2'-C1'	-12.30	1.39	1.53
85	AA	1439	A	P-O5'	-12.30	1.47	1.59
85	AA	1519	A	N9-C4	-12.30	1.30	1.37
85	AA	694	A	O3'-P	-12.30	1.46	1.61
40	BG	27	C	P-O5'	-12.29	1.47	1.59
85	AA	313	A	C1'-N9	-12.29	1.29	1.46
85	AA	373	G	C2'-C1'	-12.29	1.39	1.53
35	BB	606	C	P-O5'	-12.29	1.47	1.59
35	BB	676	G	P-O5'	-12.28	1.47	1.59
85	AA	392	G	P-O5'	-12.28	1.47	1.59
35	BB	1138	A	O3'-P	-12.28	1.46	1.61
85	AA	2075	C	C2'-C1'	-12.28	1.39	1.53
34	BA	17	A	P-O5'	-12.28	1.47	1.59
34	BA	1548	A	O3'-P	-12.28	1.46	1.61
35	BB	1430	G	O3'-P	-12.28	1.46	1.61
35	BB	446	U	O3'-P	-12.27	1.46	1.61
85	AA	1225	C	P-O5'	-12.27	1.47	1.59
85	AA	426	C	C2-N3	-12.27	1.25	1.35
85	AA	43	A	N9-C4	-12.27	1.30	1.37
34	BA	387	A	P-O5'	-12.27	1.47	1.59
40	BG	55	A	N9-C4	-12.27	1.30	1.37
36	BC	58	G	P-O5'	-12.26	1.47	1.59
40	BG	149	U	C1'-N1	-12.26	1.29	1.46
34	BA	408	U	P-O5'	-12.26	1.47	1.59
34	BA	1110	A	O3'-P	-12.26	1.46	1.61
34	BA	540	G	P-O5'	-12.25	1.47	1.59
35	BB	22	A	N9-C4	-12.24	1.30	1.37
34	BA	473	A	O3'-P	-12.24	1.46	1.61
37	BD	108	G	P-O5'	-12.24	1.47	1.59
34	BA	1265	G	P-O5'	-12.24	1.47	1.59
36	BC	154	A	C2'-C1'	-12.24	1.39	1.53
38	BE	31	A	N9-C4	-12.24	1.30	1.37
85	AA	112	A	O3'-P	-12.24	1.46	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	542	A	N9-C4	-12.24	1.30	1.37
85	AA	2036	A	N7-C5	-12.24	1.31	1.39
85	AA	1856	G	P-O5'	-12.23	1.47	1.59
34	BA	804	G	N7-C5	-12.23	1.31	1.39
34	BA	1006	G	P-O5'	-12.23	1.47	1.59
34	BA	184	C	P-O5'	-12.22	1.47	1.59
34	BA	463	A	N9-C4	-12.22	1.30	1.37
85	AA	1107	A	P-O5'	-12.22	1.47	1.59
34	BA	1041	U	P-O5'	-12.22	1.47	1.59
35	BB	479	U	P-O5'	-12.22	1.47	1.59
85	AA	1490	A	O3'-P	-12.22	1.46	1.61
35	BB	83	G	P-O5'	-12.22	1.47	1.59
85	AA	1561	A	N9-C4	-12.21	1.30	1.37
34	BA	381	A	O3'-P	-12.21	1.46	1.61
38	BE	146	U	C2-N3	-12.21	1.29	1.37
34	BA	1058	C	O3'-P	-12.20	1.46	1.61
34	BA	1328	U	C2'-C1'	-12.20	1.40	1.53
34	BA	1737	A	P-O5'	-12.20	1.47	1.59
85	AA	2133	A	O3'-P	-12.20	1.46	1.61
38	BE	108	U	C2'-C1'	-12.19	1.40	1.53
35	BB	1052	G	N9-C4	-12.19	1.28	1.38
37	BD	33	U	O3'-P	-12.18	1.46	1.61
34	BA	395	G	C1'-N9	-12.18	1.29	1.46
35	BB	637	G	O3'-P	-12.18	1.46	1.61
85	AA	657	C	P-O5'	-12.17	1.47	1.59
34	BA	461	A	N9-C4	-12.17	1.30	1.37
38	BE	140	G	P-O5'	-12.17	1.47	1.59
85	AA	264	A	P-O5'	-12.16	1.47	1.59
35	BB	1542	C	P-O5'	-12.16	1.47	1.59
35	BB	398	A	P-O5'	-12.16	1.47	1.59
85	AA	723	U	P-O5'	-12.16	1.47	1.59
34	BA	611	A	O3'-P	-12.15	1.46	1.61
35	BB	667	G	P-O5'	-12.15	1.47	1.59
36	BC	10	C	C2'-C1'	-12.15	1.40	1.53
40	BG	141	A	N9-C4	-12.15	1.30	1.37
36	BC	94	C	O3'-P	-12.14	1.46	1.61
37	BD	4	U	C2-N3	-12.14	1.29	1.37
85	AA	466	A	N9-C4	-12.14	1.30	1.37
34	BA	1618	A	P-O5'	-12.14	1.47	1.59
35	BB	1449	G	N9-C4	-12.14	1.28	1.38
34	BA	1699	A	C2'-C1'	-12.14	1.40	1.53
85	AA	1978	G	N9-C4	-12.14	1.28	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	466	A	C6-N6	12.14	1.43	1.33
85	AA	936	C	O3'-P	-12.13	1.46	1.61
35	BB	101	U	O3'-P	-12.13	1.46	1.61
85	AA	432	A	P-O5'	-12.13	1.47	1.59
34	BA	206	C	C4'-C3'	12.12	1.66	1.53
41	BH	3	U	O3'-P	-12.13	1.46	1.61
85	AA	2181	G	P-O5'	-12.12	1.47	1.59
34	BA	615	A	P-O5'	-12.12	1.47	1.59
36	BC	107	C	C2'-C1'	-12.12	1.40	1.53
85	AA	1521	U	O3'-P	-12.12	1.46	1.61
35	BB	1235	A	N9-C4	-12.11	1.30	1.37
38	BE	5	A	N9-C4	-12.11	1.30	1.37
35	BB	127	U	P-O5'	-12.11	1.47	1.59
35	BB	700	C	P-O5'	-12.11	1.47	1.59
35	BB	1342	C	C4'-C3'	-12.11	1.39	1.53
41	BH	119	U	P-O5'	-12.11	1.47	1.59
36	BC	141	C	O3'-P	-12.11	1.46	1.61
85	AA	385	A	O3'-P	-12.11	1.46	1.61
34	BA	449	G	P-O5'	-12.10	1.47	1.59
41	BH	4	U	P-O5'	-12.10	1.47	1.59
36	BC	135	A	P-O5'	-12.10	1.47	1.59
34	BA	1669	C	P-O5'	-12.09	1.47	1.59
35	BB	14	C	O3'-P	-12.08	1.46	1.61
85	AA	1676	G	P-O5'	-12.08	1.47	1.59
85	AA	2084	U	P-O5'	-12.08	1.47	1.59
85	AA	2148	C	P-O5'	-12.08	1.47	1.59
85	AA	535	G	P-O5'	-12.08	1.47	1.59
34	BA	420	A	N9-C4	-12.07	1.30	1.37
34	BA	947	A	N9-C4	-12.07	1.30	1.37
35	BB	1264	U	P-O5'	-12.07	1.47	1.59
36	BC	53	A	O3'-P	-12.07	1.46	1.61
85	AA	2070	C	P-O5'	-12.06	1.47	1.59
85	AA	811	A	P-O5'	-12.06	1.47	1.59
85	AA	412	G	P-O5'	-12.06	1.47	1.59
85	AA	366	A	P-O5'	-12.05	1.47	1.59
35	BB	1180	G	P-O5'	-12.05	1.47	1.59
34	BA	186	G	P-O5'	-12.05	1.47	1.59
40	BG	8	U	P-O5'	-12.05	1.47	1.59
35	BB	61	A	O3'-P	-12.04	1.46	1.61
85	AA	2185	U	P-O5'	-12.04	1.47	1.59
34	BA	272	A	N9-C4	-12.04	1.30	1.37
37	BD	109	U	P-O5'	-12.04	1.47	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	180	G	N9-C4	-12.03	1.28	1.38
35	BB	451	A	O3'-P	-12.03	1.46	1.61
34	BA	531	C	O3'-P	-12.03	1.46	1.61
34	BA	1215	U	P-O5'	-12.03	1.47	1.59
34	BA	598	G	P-O5'	-12.03	1.47	1.59
35	BB	1079	G	P-O5'	-12.02	1.47	1.59
34	BA	1641	G	C3'-C2'	-12.02	1.39	1.52
85	AA	1157	U	O3'-P	-12.02	1.46	1.61
34	BA	82	A	C1'-N9	-12.01	1.30	1.46
34	BA	1215	U	C2-N3	-12.01	1.29	1.37
85	AA	641	A	P-O5'	-12.01	1.47	1.59
35	BB	1312	U	C2-N3	-12.01	1.29	1.37
35	BB	1435	G	N9-C4	-12.01	1.28	1.38
86	AB	66	U	P-O5'	-12.01	1.47	1.59
85	AA	889	G	N7-C5	-12.00	1.32	1.39
41	BH	36	C	O3'-P	-12.00	1.46	1.61
34	BA	923	C	O3'-P	-11.99	1.46	1.61
34	BA	337	C	P-O5'	-11.99	1.47	1.59
34	BA	473	A	P-O5'	-11.99	1.47	1.59
34	BA	524	G	C6-N1	-11.99	1.31	1.39
34	BA	265	A	P-O5'	-11.99	1.47	1.59
34	BA	1555	G	O3'-P	-11.99	1.46	1.61
36	BC	26	U	O3'-P	-11.99	1.46	1.61
85	AA	1106	A	P-O5'	-11.99	1.47	1.59
85	AA	2102	A	N9-C4	-11.98	1.30	1.37
35	BB	1218	G	C2'-C1'	-11.98	1.40	1.53
85	AA	938	A	N7-C5	-11.98	1.32	1.39
34	BA	1286	C	O3'-P	-11.97	1.46	1.61
37	BD	24	U	P-O5'	-11.97	1.47	1.59
35	BB	648	G	C6-N1	-11.97	1.31	1.39
38	BE	108	U	O3'-P	-11.97	1.46	1.61
34	BA	326	A	O3'-P	-11.96	1.46	1.61
34	BA	1239	G	O3'-P	-11.96	1.46	1.61
35	BB	1440	A	O3'-P	-11.96	1.46	1.61
34	BA	537	C	P-O5'	-11.95	1.47	1.59
40	BG	8	U	C2-N3	-11.95	1.29	1.37
34	BA	1069	U	C3'-C2'	-11.94	1.39	1.52
38	BE	166	G	P-O5'	-11.94	1.47	1.59
34	BA	856	G	C2'-C1'	-11.94	1.40	1.53
34	BA	1016	A	N9-C4	-11.94	1.30	1.37
34	BA	91	C	O3'-P	-11.93	1.46	1.61
85	AA	983	A	N9-C4	-11.93	1.30	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	805	A	P-O5'	-11.93	1.47	1.59
85	AA	283	A	N9-C4	-11.93	1.30	1.37
34	BA	1268	C	P-O5'	-11.93	1.47	1.59
85	AA	249	C	P-O5'	-11.93	1.47	1.59
34	BA	662	U	P-O5'	-11.92	1.47	1.59
34	BA	800	G	O4'-C1'	-11.92	1.26	1.41
34	BA	1232	C	P-O5'	-11.92	1.47	1.59
35	BB	630	A	C4'-C3'	-11.92	1.40	1.53
85	AA	201	U	P-O5'	-11.92	1.47	1.59
85	AA	2147	A	N7-C5	-11.92	1.32	1.39
85	AA	2147	A	O3'-P	-11.91	1.46	1.61
38	BE	132	U	P-O5'	-11.91	1.47	1.59
39	BF	12	U	P-O5'	-11.91	1.47	1.59
34	BA	22	C	P-O5'	-11.90	1.47	1.59
34	BA	440	A	O3'-P	-11.90	1.46	1.61
85	AA	866	U	C2'-C1'	-11.90	1.40	1.53
34	BA	48	C	O3'-P	-11.90	1.46	1.61
38	BE	14	C	P-O5'	-11.90	1.47	1.59
40	BG	112	C	P-O5'	-11.90	1.47	1.59
36	BC	3	C	C2'-C1'	-11.89	1.40	1.53
34	BA	1223	C	P-O5'	-11.89	1.47	1.59
85	AA	1829	C	P-O5'	-11.89	1.47	1.59
36	BC	79	A	N9-C4	-11.89	1.30	1.37
34	BA	513	U	C2-N3	-11.89	1.29	1.37
34	BA	504	A	P-O5'	-11.88	1.47	1.59
34	BA	1335	A	P-O5'	-11.88	1.47	1.59
85	AA	731	U	O3'-P	-11.88	1.46	1.61
34	BA	953	G	P-O5'	-11.88	1.47	1.59
85	AA	407	G	P-O5'	-11.88	1.47	1.59
34	BA	1276	G	P-O5'	-11.88	1.47	1.59
34	BA	958	G	P-O5'	-11.87	1.47	1.59
34	BA	1275	G	P-O5'	-11.87	1.47	1.59
34	BA	1291	A	N9-C4	-11.87	1.30	1.37
85	AA	438	G	P-O5'	-11.87	1.47	1.59
85	AA	1455	C	C2'-C1'	-11.87	1.40	1.53
34	BA	382	G	C6-N1	-11.87	1.31	1.39
34	BA	301	U	C2'-C1'	-11.86	1.40	1.53
85	AA	1518	A	N9-C4	-11.86	1.30	1.37
85	AA	1541	G	P-O5'	-11.86	1.47	1.59
34	BA	1583	A	N9-C4	-11.85	1.30	1.37
36	BC	120	G	N9-C4	-11.85	1.28	1.38
85	AA	515	C	O3'-P	-11.85	1.47	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2130	G	O3'-P	-11.85	1.47	1.61
34	BA	1591	G	C1'-N9	-11.85	1.30	1.46
35	BB	1498	G	N9-C4	-11.85	1.28	1.38
85	AA	1227	A	P-O5'	-11.85	1.48	1.59
34	BA	609	G	C2'-C1'	-11.84	1.40	1.53
34	BA	979	G	O3'-P	-11.84	1.47	1.61
85	AA	936	C	P-O5'	-11.84	1.48	1.59
34	BA	1844	U	P-O5'	-11.84	1.48	1.59
35	BB	59	U	C2-N3	-11.84	1.29	1.37
85	AA	374	C	P-O5'	-11.84	1.48	1.59
34	BA	340	U	P-O5'	-11.84	1.48	1.59
85	AA	299	A	P-O5'	-11.84	1.48	1.59
34	BA	504	A	N9-C4	-11.83	1.30	1.37
34	BA	762	A	P-O5'	-11.83	1.48	1.59
34	BA	2	A	P-O5'	-11.83	1.48	1.59
34	BA	667	U	P-O5'	-11.83	1.48	1.59
34	BA	1641	G	O3'-P	-11.83	1.47	1.61
35	BB	990	G	O3'-P	-11.83	1.47	1.61
34	BA	355	U	P-O5'	-11.82	1.48	1.59
35	BB	805	G	C4'-C3'	-11.82	1.40	1.53
85	AA	1471	G	C5-C6	-11.82	1.30	1.42
85	AA	1661	U	P-O5'	-11.82	1.48	1.59
35	BB	572	G	N7-C5	-11.81	1.32	1.39
85	AA	695	A	N9-C4	-11.81	1.30	1.37
85	AA	2137	A	N9-C4	-11.81	1.30	1.37
34	BA	985	C	O3'-P	-11.81	1.47	1.61
34	BA	1190	A	P-O5'	-11.81	1.48	1.59
34	BA	1220	C	P-O5'	-11.81	1.48	1.59
85	AA	822	U	P-O5'	-11.81	1.48	1.59
35	BB	499	A	N9-C4	-11.80	1.30	1.37
85	AA	902	A	N9-C4	-11.80	1.30	1.37
85	AA	1674	G	N9-C4	-11.81	1.28	1.38
40	BG	95	U	C1'-N1	-11.80	1.30	1.46
38	BE	20	C	C2'-C1'	-11.80	1.40	1.53
38	BE	61	A	P-O5'	-11.80	1.48	1.59
85	AA	2203	C	C2'-C1'	-11.80	1.40	1.53
35	BB	1246	C	O3'-P	-11.79	1.47	1.61
35	BB	1546	C	P-O5'	-11.79	1.48	1.59
34	BA	687	G	O3'-P	-11.79	1.47	1.61
85	AA	397	G	O3'-P	-11.79	1.47	1.61
85	AA	879	G	C2'-C1'	-11.79	1.40	1.53
34	BA	16	C	P-O5'	-11.79	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	758	G	N9-C8	-11.78	1.29	1.37
34	BA	838	U	P-O5'	-11.78	1.48	1.59
85	AA	1191	G	P-O5'	-11.78	1.48	1.59
34	BA	856	G	O3'-P	-11.78	1.47	1.61
34	BA	1720	U	P-O5'	-11.78	1.48	1.59
41	BH	35	G	P-O5'	-11.78	1.48	1.59
41	BH	133	U	P-O5'	-11.78	1.48	1.59
40	BG	88	G	O3'-P	-11.77	1.47	1.61
35	BB	1172	U	O3'-P	-11.77	1.47	1.61
85	AA	99	U	O3'-P	-11.77	1.47	1.61
85	AA	1224	C	O3'-P	-11.77	1.47	1.61
34	BA	162	G	N9-C4	-11.77	1.28	1.38
85	AA	927	A	N9-C4	11.77	1.45	1.37
85	AA	991	G	N9-C8	-11.77	1.29	1.37
38	BE	190	U	O3'-P	-11.76	1.47	1.61
85	AA	684	G	P-O5'	-11.76	1.48	1.59
41	BH	126	C	O3'-P	-11.76	1.47	1.61
34	BA	1604	A	O3'-P	-11.76	1.47	1.61
34	BA	333	A	N9-C4	-11.76	1.30	1.37
34	BA	1223	C	O3'-P	-11.76	1.47	1.61
34	BA	449	G	O3'-P	-11.76	1.47	1.61
35	BB	370	A	P-O5'	-11.75	1.48	1.59
35	BB	1019	C	P-O5'	-11.75	1.48	1.59
35	BB	440	U	P-O5'	-11.75	1.48	1.59
35	BB	1227	G	O3'-P	-11.75	1.47	1.61
34	BA	1177	C	C2'-C1'	-11.75	1.40	1.53
85	AA	2174	G	C2'-C1'	-11.75	1.40	1.53
35	BB	1314	G	O3'-P	-11.74	1.47	1.61
34	BA	1282	G	O3'-P	-11.74	1.47	1.61
40	BG	118	U	C2-N3	-11.74	1.29	1.37
34	BA	938	C	C3'-C2'	-11.74	1.39	1.52
40	BG	23	C	P-O5'	-11.74	1.48	1.59
34	BA	1661	U	O3'-P	-11.74	1.47	1.61
35	BB	409	U	P-O5'	-11.74	1.48	1.59
35	BB	1243	A	O3'-P	-11.73	1.47	1.61
85	AA	464	A	N7-C5	-11.73	1.32	1.39
35	BB	777	C	P-O5'	-11.73	1.48	1.59
35	BB	1103	A	O3'-P	-11.72	1.47	1.61
85	AA	1182	A	C2'-C1'	-11.72	1.40	1.53
41	BH	44	A	N7-C5	-11.72	1.32	1.39
40	BG	70	C	P-O5'	-11.72	1.48	1.59
85	AA	83	U	P-O5'	-11.72	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2125	A	O3'-P	-11.72	1.47	1.61
85	AA	1219	A	N7-C5	-11.72	1.32	1.39
34	BA	909	G	O3'-P	-11.71	1.47	1.61
39	BF	38	C	P-O5'	-11.71	1.48	1.59
40	BG	23	C	O3'-P	-11.71	1.47	1.61
85	AA	809	A	P-O5'	-11.71	1.48	1.59
34	BA	108	A	N9-C4	-11.71	1.30	1.37
35	BB	1367	U	P-O5'	-11.70	1.48	1.59
85	AA	1976	G	P-O5'	-11.70	1.48	1.59
34	BA	382	G	P-O5'	-11.70	1.48	1.59
34	BA	1003	A	C3'-C2'	-11.70	1.39	1.52
85	AA	189	G	P-O5'	-11.70	1.48	1.59
35	BB	1183	U	O3'-P	-11.70	1.47	1.61
85	AA	868	A	P-O5'	-11.70	1.48	1.59
41	BH	105	U	P-O5'	-11.69	1.48	1.59
34	BA	824	C	P-O5'	-11.69	1.48	1.59
85	AA	1510	A	N9-C4	-11.69	1.30	1.37
34	BA	651	U	C2'-C1'	-11.68	1.40	1.53
34	BA	1796	A	C1'-N9	-11.68	1.30	1.46
38	BE	113	C	P-O5'	-11.68	1.48	1.59
38	BE	148	C	P-O5'	-11.68	1.48	1.59
41	BH	113	G	N9-C4	-11.68	1.28	1.38
39	BF	39	C	O3'-P	-11.67	1.47	1.61
85	AA	1228	A	C2'-C1'	-11.67	1.40	1.53
85	AA	1549	G	N9-C4	-11.67	1.28	1.38
34	BA	1341	A	N9-C4	-11.67	1.30	1.37
85	AA	2029	G	P-O5'	-11.67	1.48	1.59
34	BA	400	A	P-O5'	-11.67	1.48	1.59
34	BA	1002	U	C3'-C2'	-11.67	1.40	1.52
38	BE	16	C	C5'-C4'	11.67	1.65	1.51
34	BA	201	A	O3'-P	-11.66	1.47	1.61
34	BA	979	G	N9-C4	-11.66	1.28	1.38
34	BA	1424	G	P-O5'	-11.66	1.48	1.59
85	AA	472	A	P-O5'	-11.66	1.48	1.59
37	BD	48	G	N9-C4	-11.66	1.28	1.38
85	AA	1283	C	C2-N3	-11.66	1.26	1.35
34	BA	93	A	P-O5'	-11.65	1.48	1.59
34	BA	1465	C	O3'-P	-11.65	1.47	1.61
35	BB	1528	U	P-O5'	-11.65	1.48	1.59
41	BH	124	C	O3'-P	-11.65	1.47	1.61
85	AA	2149	C	C2'-C1'	-11.65	1.40	1.53
85	AA	2227	A	N9-C4	-11.65	1.30	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	649	A	P-O5'	-11.65	1.48	1.59
36	BC	43	A	C3'-C2'	-11.65	1.40	1.52
37	BD	73	U	P-O5'	-11.65	1.48	1.59
85	AA	2123	U	P-O5'	-11.65	1.48	1.59
35	BB	1180	G	O3'-P	-11.64	1.47	1.61
36	BC	67	U	P-O5'	-11.64	1.48	1.59
85	AA	488	G	P-O5'	-11.64	1.48	1.59
34	BA	53	G	O3'-P	-11.64	1.47	1.61
34	BA	1732	A	N9-C4	11.64	1.44	1.37
34	BA	484	A	P-O5'	-11.64	1.48	1.59
34	BA	471	U	P-O5'	-11.64	1.48	1.59
35	BB	387	G	N9-C8	-11.64	1.29	1.37
85	AA	2199	G	C2'-C1'	-11.63	1.40	1.53
34	BA	1002	U	P-O5'	-11.63	1.48	1.59
35	BB	615	A	N9-C4	-11.63	1.30	1.37
37	BD	77	A	N9-C4	-11.63	1.30	1.37
41	BH	17	A	P-O5'	-11.63	1.48	1.59
85	AA	605	A	N7-C5	-11.63	1.32	1.39
85	AA	712	U	P-O5'	-11.63	1.48	1.59
34	BA	350	C	O3'-P	-11.62	1.47	1.61
35	BB	1463	A	N9-C4	11.62	1.44	1.37
85	AA	393	C	O3'-P	-11.62	1.47	1.61
34	BA	734	G	P-O5'	-11.62	1.48	1.59
34	BA	896	U	C2'-C1'	11.62	1.66	1.53
34	BA	942	G	P-O5'	-11.61	1.48	1.59
35	BB	1490	G	C2'-C1'	-11.61	1.40	1.53
34	BA	782	C	C2'-C1'	-11.61	1.40	1.53
85	AA	106	G	P-O5'	-11.61	1.48	1.59
35	BB	1075	A	N9-C4	-11.61	1.30	1.37
41	BH	41	A	N9-C4	-11.61	1.30	1.37
85	AA	763	U	C2-N3	-11.61	1.29	1.37
85	AA	1540	A	P-O5'	-11.61	1.48	1.59
34	BA	31	A	N9-C4	-11.60	1.30	1.37
40	BG	27	C	O3'-P	-11.60	1.47	1.61
41	BH	33	G	P-O5'	-11.60	1.48	1.59
85	AA	601	A	O3'-P	-11.60	1.47	1.61
34	BA	89	G	P-O5'	-11.59	1.48	1.59
35	BB	1048	A	N9-C4	-11.59	1.30	1.37
85	AA	383	C	P-O5'	-11.59	1.48	1.59
34	BA	1100	A	P-O5'	-11.59	1.48	1.59
34	BA	1193	A	O3'-P	-11.59	1.47	1.61
35	BB	818	U	P-O5'	-11.59	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	115	A	P-O5'	-11.59	1.48	1.59
40	BG	95	U	C2-N3	-11.59	1.29	1.37
36	BC	78	G	O3'-P	-11.58	1.47	1.61
36	BC	92	C	P-O5'	-11.58	1.48	1.59
34	BA	954	U	P-O5'	-11.58	1.48	1.59
85	AA	630	A	N9-C4	-11.58	1.30	1.37
35	BB	1348	C	O3'-P	-11.58	1.47	1.61
85	AA	1531	G	P-O5'	-11.58	1.48	1.59
34	BA	877	U	C2-N3	-11.57	1.29	1.37
37	BD	68	C	P-O5'	-11.57	1.48	1.59
34	BA	737	U	O3'-P	-11.57	1.47	1.61
36	BC	2	A	N9-C4	-11.57	1.30	1.37
38	BE	200	A	N9-C4	-11.57	1.30	1.37
34	BA	1430	C	O3'-P	-11.57	1.47	1.61
34	BA	1809	G	O3'-P	-11.57	1.47	1.61
34	BA	1697	U	C2-N3	-11.56	1.45	1.37
35	BB	1413	U	P-O5'	-11.56	1.48	1.59
40	BG	78	C	O3'-P	-11.56	1.47	1.61
85	AA	605	A	N9-C4	-11.56	1.30	1.37
85	AA	436	G	P-O5'	-11.56	1.48	1.59
38	BE	128	G	O3'-P	-11.56	1.47	1.61
34	BA	850	C	P-O5'	-11.55	1.48	1.59
85	AA	244	G	P-O5'	-11.56	1.48	1.59
34	BA	984	U	C2-N3	-11.55	1.29	1.37
35	BB	1461	C	C2'-C1'	-11.55	1.40	1.53
85	AA	270	A	P-O5'	-11.55	1.48	1.59
35	BB	644	A	N9-C4	-11.54	1.30	1.37
34	BA	297	A	C4'-C3'	-11.54	1.40	1.53
34	BA	1159	A	P-O5'	-11.54	1.48	1.59
34	BA	1321	A	C2'-C1'	-11.54	1.40	1.53
34	BA	720	A	O3'-P	-11.53	1.47	1.61
85	AA	2105	G	P-O5'	-11.54	1.48	1.59
35	BB	798	A	C2'-C1'	-11.53	1.40	1.53
85	AA	455	G	P-O5'	-11.53	1.48	1.59
35	BB	1242	C	P-O5'	-11.53	1.48	1.59
35	BB	824	C	C2'-C1'	-11.53	1.40	1.53
35	BB	1232	A	P-O5'	-11.52	1.48	1.59
34	BA	912	G	O3'-P	-11.52	1.47	1.61
38	BE	28	C	P-O5'	-11.52	1.48	1.59
34	BA	86	A	P-O5'	-11.52	1.48	1.59
34	BA	810	A	P-O5'	-11.52	1.48	1.59
34	BA	918	U	P-O5'	-11.52	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	482	A	N9-C4	-11.52	1.30	1.37
34	BA	35	U	P-O5'	-11.51	1.48	1.59
34	BA	125	G	P-O5'	-11.51	1.48	1.59
35	BB	1360	A	N9-C4	-11.51	1.30	1.37
34	BA	1549	U	P-O5'	-11.51	1.48	1.59
35	BB	5	A	N7-C5	-11.51	1.32	1.39
34	BA	899	G	O3'-P	-11.51	1.47	1.61
35	BB	1421	C	P-O5'	-11.51	1.48	1.59
37	BD	77	A	C1'-N9	-11.51	1.30	1.46
34	BA	545	U	P-O5'	-11.50	1.48	1.59
85	AA	1676	G	O3'-P	-11.50	1.47	1.61
34	BA	556	A	O3'-P	-11.50	1.47	1.61
34	BA	1322	A	O3'-P	-11.50	1.47	1.61
37	BD	84	U	P-O5'	-11.50	1.48	1.59
85	AA	2228	G	P-O5'	-11.50	1.48	1.59
34	BA	1722	U	P-O5'	-11.50	1.48	1.59
35	BB	1368	A	N9-C4	-11.50	1.30	1.37
41	BH	31	A	C1'-N9	-11.50	1.30	1.46
35	BB	970	C	O3'-P	-11.49	1.47	1.61
36	BC	101	U	C2'-C1'	-11.49	1.40	1.53
85	AA	24	U	P-O5'	-11.49	1.48	1.59
85	AA	436	G	N9-C4	-11.49	1.28	1.38
85	AA	541	A	P-O5'	-11.49	1.48	1.59
34	BA	1709	A	C3'-C2'	-11.49	1.40	1.52
34	BA	44	U	P-O5'	-11.49	1.48	1.59
39	BF	65	U	P-O5'	-11.49	1.48	1.59
35	BB	1415	G	C6-N1	-11.48	1.31	1.39
38	BE	39	U	P-O5'	-11.48	1.48	1.59
35	BB	440	U	C3'-C2'	-11.48	1.40	1.52
35	BB	1293	C	P-O5'	-11.48	1.48	1.59
85	AA	210	G	C2'-C1'	-11.48	1.40	1.53
34	BA	717	U	P-O5'	-11.48	1.48	1.59
34	BA	1099	U	O3'-P	-11.48	1.47	1.61
38	BE	129	G	C6-N1	-11.48	1.31	1.39
35	BB	1199	A	N9-C4	-11.47	1.30	1.37
40	BG	69	G	O3'-P	-11.47	1.47	1.61
85	AA	521	A	C2'-C1'	-11.47	1.40	1.53
85	AA	2075	C	N1-C6	-11.47	1.30	1.37
85	AA	922	A	O3'-P	-11.46	1.47	1.61
34	BA	587	U	O3'-P	-11.46	1.47	1.61
34	BA	1050	A	N9-C4	-11.46	1.30	1.37
85	AA	1470	A	N9-C4	-11.46	1.30	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1631	U	C4'-C3'	11.46	1.65	1.53
34	BA	1190	A	C2'-C1'	-11.46	1.40	1.53
34	BA	1556	A	O3'-P	-11.45	1.47	1.61
38	BE	141	A	P-O5'	-11.45	1.48	1.59
36	BC	162	C	P-O5'	-11.45	1.48	1.59
85	AA	1481	U	P-O5'	-11.45	1.48	1.59
34	BA	1076	U	C2-N3	-11.44	1.29	1.37
37	BD	82	G	O3'-P	-11.45	1.47	1.61
34	BA	10	G	C3'-C2'	-11.44	1.40	1.52
34	BA	1027	C	O3'-P	-11.44	1.47	1.61
34	BA	1693	U	O3'-P	-11.44	1.47	1.61
35	BB	1398	A	N9-C4	-11.44	1.30	1.37
34	BA	1144	A	P-O5'	-11.44	1.48	1.59
34	BA	1529	G	C6-N1	-11.43	1.31	1.39
34	BA	1688	G	P-O5'	-11.43	1.48	1.59
34	BA	1256	A	N9-C4	-11.42	1.30	1.37
34	BA	1259	C	O3'-P	-11.42	1.47	1.61
85	AA	808	A	O3'-P	-11.42	1.47	1.61
85	AA	1357	U	P-O5'	-11.42	1.48	1.59
41	BH	54	U	P-O5'	-11.42	1.48	1.59
85	AA	1499	G	O3'-P	-11.42	1.47	1.61
35	BB	368	C	O3'-P	-11.41	1.47	1.61
38	BE	200	A	C2'-C1'	-11.41	1.40	1.53
85	AA	469	G	N3-C4	-11.41	1.27	1.35
85	AA	1182	A	P-O5'	-11.41	1.48	1.59
34	BA	907	A	O3'-P	-11.41	1.47	1.61
34	BA	986	G	P-O5'	-11.41	1.48	1.59
85	AA	314	C	P-O5'	-11.41	1.48	1.59
85	AA	370	A	P-O5'	-11.41	1.48	1.59
85	AA	441	C	O3'-P	-11.41	1.47	1.61
85	AA	918	U	O3'-P	-11.41	1.47	1.61
34	BA	1322	A	C2'-C1'	-11.40	1.40	1.53
85	AA	119	G	C2'-C1'	-11.40	1.40	1.53
34	BA	900	A	N7-C5	-11.40	1.32	1.39
40	BG	105	A	N9-C4	-11.40	1.31	1.37
35	BB	411	A	N9-C4	-11.40	1.31	1.37
35	BB	818	U	C2-N3	-11.40	1.29	1.37
37	BD	95	G	C6-N1	-11.40	1.31	1.39
34	BA	1028	A	N9-C4	-11.39	1.31	1.37
35	BB	814	A	P-O5'	-11.39	1.48	1.59
38	BE	46	G	P-O5'	-11.39	1.48	1.59
35	BB	1226	G	C1'-N9	-11.39	1.30	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1412	G	O3'-P	-11.39	1.47	1.61
36	BC	168	C	P-O5'	-11.39	1.48	1.59
34	BA	837	U	O3'-P	-11.39	1.47	1.61
85	AA	545	A	P-O5'	-11.39	1.48	1.59
34	BA	606	G	P-O5'	-11.38	1.48	1.59
35	BB	1028	C	O3'-P	-11.38	1.47	1.61
35	BB	1312	U	P-O5'	-11.38	1.48	1.59
35	BB	56	U	O3'-P	-11.38	1.47	1.61
35	BB	1467	A	N7-C5	-11.38	1.32	1.39
34	BA	1254	C	P-O5'	-11.38	1.48	1.59
35	BB	1342	C	O3'-P	-11.38	1.47	1.61
34	BA	358	A	N9-C4	-11.37	1.31	1.37
40	BG	48	U	P-O5'	-11.37	1.48	1.59
85	AA	1909	C	P-O5'	-11.37	1.48	1.59
85	AA	116	G	N9-C4	-11.37	1.28	1.38
34	BA	1147	C	O3'-P	-11.36	1.47	1.61
35	BB	414	C	P-O5'	-11.36	1.48	1.59
85	AA	2176	U	N1-C2	-11.36	1.28	1.38
35	BB	68	G	N9-C4	-11.35	1.28	1.38
85	AA	1127	G	C1'-N9	-11.35	1.30	1.46
85	AA	2136	C	P-O5'	-11.35	1.48	1.59
34	BA	1597	G	N9-C4	-11.35	1.28	1.38
38	BE	34	C	C2'-C1'	-11.35	1.40	1.53
85	AA	1281	G	N9-C4	-11.35	1.28	1.38
34	BA	113	G	C3'-C2'	-11.35	1.40	1.52
34	BA	356	C	O3'-P	-11.35	1.47	1.61
85	AA	4	C	O3'-P	-11.35	1.47	1.61
85	AA	25	C	P-O5'	-11.35	1.48	1.59
34	BA	1282	G	N9-C4	-11.34	1.28	1.38
35	BB	781	U	O3'-P	-11.34	1.47	1.61
36	BC	9	G	C2'-C1'	-11.34	1.40	1.53
36	BC	111	C	P-O5'	-11.34	1.48	1.59
85	AA	2227	A	P-O5'	-11.34	1.48	1.59
34	BA	1494	G	O3'-P	-11.34	1.47	1.61
34	BA	1650	G	O3'-P	-11.33	1.47	1.61
85	AA	2185	U	C2'-C1'	-11.33	1.40	1.53
34	BA	1071	G	O3'-P	-11.32	1.47	1.61
34	BA	262	A	N9-C4	-11.32	1.31	1.37
34	BA	1676	A	P-O5'	-11.32	1.48	1.59
34	BA	961	C	C2'-C1'	-11.32	1.41	1.53
34	BA	1325	G	O3'-P	-11.31	1.47	1.61
35	BB	46	U	P-O5'	-11.31	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	164	G	P-O5'	-11.31	1.48	1.59
34	BA	1055	U	P-O5'	-11.31	1.48	1.59
35	BB	1147	G	P-O5'	-11.31	1.48	1.59
40	BG	46	G	P-O5'	-11.31	1.48	1.59
40	BG	176	G	P-O5'	-11.31	1.48	1.59
85	AA	2008	G	N3-C4	-11.31	1.27	1.35
36	BC	93	C	O3'-P	-11.31	1.47	1.61
85	AA	45	U	O3'-P	-11.31	1.47	1.61
85	AA	1486	G	C2'-C1'	-11.30	1.41	1.53
34	BA	592	G	C4'-C3'	11.30	1.65	1.53
34	BA	6	C	O3'-P	-11.29	1.47	1.61
34	BA	478	G	C2'-C1'	-11.30	1.41	1.53
36	BC	128	U	P-O5'	-11.30	1.48	1.59
36	BC	55	U	C2'-C1'	-11.29	1.41	1.53
35	BB	1466	A	N9-C4	-11.29	1.31	1.37
37	BD	65	G	P-O5'	-11.29	1.48	1.59
34	BA	432	A	N9-C4	-11.29	1.31	1.37
36	BC	20	C	P-O5'	-11.29	1.48	1.59
36	BC	100	U	O3'-P	-11.29	1.47	1.61
40	BG	162	A	P-O5'	-11.28	1.48	1.59
85	AA	663	C	O3'-P	-11.28	1.47	1.61
39	BF	62	U	O3'-P	-11.28	1.47	1.61
34	BA	13	U	P-O5'	-11.28	1.48	1.59
34	BA	59	A	O3'-P	-11.28	1.47	1.61
35	BB	654	C	O3'-P	-11.28	1.47	1.61
34	BA	1013	A	N9-C4	-11.27	1.31	1.37
35	BB	493	U	P-O5'	-11.27	1.48	1.59
34	BA	1739	G	N9-C4	-11.27	1.28	1.38
34	BA	1322	A	N9-C4	-11.26	1.31	1.37
35	BB	47	C	O3'-P	-11.26	1.47	1.61
36	BC	114	C	O3'-P	-11.26	1.47	1.61
39	BF	29	U	O3'-P	-11.26	1.47	1.61
36	BC	153	C	N1-C2	-11.26	1.28	1.40
85	AA	471	U	P-O5'	-11.26	1.48	1.59
34	BA	754	G	C2'-C1'	-11.26	1.41	1.53
85	AA	655	U	P-O5'	-11.26	1.48	1.59
34	BA	933	U	O3'-P	-11.25	1.47	1.61
38	BE	101	C	C3'-C2'	-11.25	1.40	1.52
35	BB	58	G	N9-C4	-11.25	1.28	1.38
35	BB	1484	A	P-O5'	-11.25	1.48	1.59
41	BH	102	C	O3'-P	-11.25	1.47	1.61
35	BB	1170	U	O3'-P	-11.25	1.47	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	549	A	P-O5'	-11.25	1.48	1.59
35	BB	74	U	P-O5'	-11.25	1.48	1.59
37	BD	16	U	P-O5'	-11.25	1.48	1.59
85	AA	757	A	P-O5'	-11.25	1.48	1.59
34	BA	572	G	N9-C4	11.24	1.47	1.38
35	BB	54	U	O3'-P	-11.23	1.47	1.61
85	AA	393	C	C3'-C2'	-11.23	1.40	1.52
34	BA	1092	U	C2-N3	-11.23	1.29	1.37
35	BB	1356	G	C2-N2	-11.23	1.23	1.34
34	BA	1261	G	C2'-C1'	-11.23	1.41	1.53
35	BB	783	U	O3'-P	-11.23	1.47	1.61
37	BD	98	G	O3'-P	-11.23	1.47	1.61
85	AA	924	A	P-O5'	-11.23	1.48	1.59
35	BB	640	A	N9-C4	-11.22	1.31	1.37
85	AA	384	C	P-O5'	-11.22	1.48	1.59
85	AA	596	A	C2'-C1'	-11.22	1.41	1.53
85	AA	1687	U	C2-N3	-11.22	1.29	1.37
35	BB	1070	G	P-O5'	-11.22	1.48	1.59
34	BA	416	A	N9-C4	-11.22	1.31	1.37
85	AA	1228	A	P-O5'	-11.22	1.48	1.59
34	BA	475	A	N9-C4	-11.21	1.31	1.37
85	AA	551	C	P-O5'	-11.21	1.48	1.59
85	AA	1248	U	O3'-P	-11.21	1.47	1.61
34	BA	381	A	N9-C4	-11.21	1.31	1.37
35	BB	1370	G	N7-C5	-11.21	1.32	1.39
34	BA	936	A	N9-C4	-11.21	1.31	1.37
34	BA	1600	G	P-O5'	-11.21	1.48	1.59
34	BA	1010	C	O3'-P	-11.20	1.47	1.61
35	BB	1105	G	P-O5'	-11.20	1.48	1.59
34	BA	652	C	P-O5'	-11.20	1.48	1.59
34	BA	929	A	N9-C4	-11.20	1.31	1.37
34	BA	1460	U	P-O5'	-11.20	1.48	1.59
38	BE	65	U	P-O5'	-11.20	1.48	1.59
34	BA	741	A	C2'-C1'	-11.20	1.41	1.53
85	AA	158	C	P-O5'	-11.20	1.48	1.59
85	AA	485	A	P-O5'	-11.20	1.48	1.59
85	AA	1110	A	P-O5'	-11.20	1.48	1.59
85	AA	1529	A	N9-C4	-11.20	1.31	1.37
35	BB	368	C	P-O5'	-11.19	1.48	1.59
35	BB	1029	U	P-O5'	-11.19	1.48	1.59
85	AA	372	U	C2-N3	-11.19	1.29	1.37
35	BB	1229	A	C2'-C1'	-11.18	1.41	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	702	G	N9-C8	-11.18	1.30	1.37
34	BA	310	C	P-O5'	-11.18	1.48	1.59
36	BC	163	A	P-O5'	-11.18	1.48	1.59
40	BG	106	G	N9-C4	-11.18	1.29	1.38
85	AA	79	G	P-O5'	-11.18	1.48	1.59
35	BB	475	A	N9-C4	-11.18	1.31	1.37
34	BA	96	G	N9-C4	-11.17	1.29	1.38
34	BA	1784	G	P-O5'	-11.17	1.48	1.59
35	BB	1456	G	O3'-P	-11.17	1.47	1.61
36	BC	44	A	O3'-P	-11.17	1.47	1.61
34	BA	593	G	C5'-C4'	11.16	1.64	1.51
35	BB	682	U	C2-N3	-11.16	1.29	1.37
35	BB	1204	C	O3'-P	-11.16	1.47	1.61
35	BB	1444	U	C2-N3	-11.16	1.29	1.37
35	BB	1444	U	P-O5'	-11.16	1.48	1.59
85	AA	496	C	P-O5'	-11.16	1.48	1.59
35	BB	50	A	O3'-P	-11.16	1.47	1.61
34	BA	71	G	O3'-P	-11.16	1.47	1.61
85	AA	2238	C	P-O5'	-11.16	1.48	1.59
36	BC	36	G	N9-C4	-11.15	1.29	1.38
85	AA	4	C	P-O5'	-11.15	1.48	1.59
85	AA	2229	G	C6-N1	-11.15	1.31	1.39
35	BB	795	A	O3'-P	-11.15	1.47	1.61
85	AA	1700	C	P-O5'	-11.15	1.48	1.59
85	AA	2139	G	N9-C4	-11.15	1.29	1.38
34	BA	75	U	C2-N3	-11.14	1.29	1.37
85	AA	603	C	P-O5'	-11.14	1.48	1.59
85	AA	1448	A	P-O5'	-11.14	1.48	1.59
37	BD	87	G	P-O5'	-11.14	1.48	1.59
34	BA	145	U	P-O5'	-11.14	1.48	1.59
34	BA	305	C	P-O5'	-11.14	1.48	1.59
34	BA	1555	G	P-O5'	-11.14	1.48	1.59
38	BE	38	C	P-O5'	-11.13	1.48	1.59
34	BA	428	C	P-O5'	-11.13	1.48	1.59
35	BB	374	A	C2'-C1'	-11.13	1.41	1.53
35	BB	416	U	O3'-P	-11.13	1.47	1.61
37	BD	69	U	P-O5'	-11.13	1.48	1.59
35	BB	1287	U	P-O5'	-11.13	1.48	1.59
34	BA	919	A	N9-C4	-11.12	1.31	1.37
34	BA	902	C	O3'-P	-11.12	1.47	1.61
34	BA	409	A	P-O5'	-11.12	1.48	1.59
36	BC	18	G	C2'-C1'	-11.12	1.41	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	120	G	C6-N1	-11.12	1.31	1.39
85	AA	381	A	C1'-N9	-11.12	1.31	1.46
85	AA	1225	C	O3'-P	-11.12	1.47	1.61
34	BA	498	A	O3'-P	-11.12	1.47	1.61
85	AA	613	G	P-O5'	-11.11	1.48	1.59
36	BC	105	C	P-O5'	-11.11	1.48	1.59
40	BG	17	A	C4'-C3'	-11.11	1.41	1.53
85	AA	696	G	O3'-P	-11.11	1.47	1.61
38	BE	2	G	O3'-P	-11.11	1.47	1.61
85	AA	756	G	O3'-P	-11.11	1.47	1.61
35	BB	1508	G	C2'-C1'	-11.11	1.41	1.53
85	AA	2146	G	C2-N2	-11.11	1.23	1.34
85	AA	1526	G	P-O5'	-11.11	1.48	1.59
85	AA	1289	U	P-O5'	-11.10	1.48	1.59
34	BA	900	A	C2'-C1'	-11.10	1.41	1.53
35	BB	1424	G	O3'-P	-11.10	1.47	1.61
85	AA	1152	U	P-O5'	-11.10	1.48	1.59
85	AA	1263	G	C6-N1	-11.10	1.31	1.39
85	AA	1442	U	C2-N3	-11.10	1.29	1.37
34	BA	244	A	P-O5'	-11.10	1.48	1.59
34	BA	812	A	P-O5'	-11.10	1.48	1.59
41	BH	29	G	N9-C4	-11.10	1.29	1.38
35	BB	1274	G	N7-C5	-11.09	1.32	1.39
85	AA	185	A	P-O5'	-11.09	1.48	1.59
34	BA	1406	U	P-O5'	-11.09	1.48	1.59
35	BB	827	U	P-O5'	-11.09	1.48	1.59
36	BC	7	U	N1-C2	-11.09	1.28	1.38
85	AA	10	G	O3'-P	-11.08	1.47	1.61
85	AA	105	A	N9-C4	-11.08	1.31	1.37
85	AA	442	G	C5-C4	-11.08	1.30	1.38
85	AA	2139	G	P-O5'	-11.08	1.48	1.59
34	BA	96	G	O3'-P	-11.08	1.47	1.61
34	BA	761	U	C1'-N1	-11.07	1.31	1.46
35	BB	584	A	C1'-N9	-11.07	1.31	1.46
34	BA	1659	G	P-O5'	-11.07	1.48	1.59
85	AA	1189	A	N9-C4	-11.07	1.31	1.37
34	BA	36	A	P-O5'	-11.07	1.48	1.59
34	BA	1194	G	C3'-C2'	-11.07	1.40	1.52
34	BA	324	C	C4'-C3'	-11.07	1.41	1.53
34	BA	758	G	N9-C4	-11.06	1.29	1.38
34	BA	1779	U	P-O5'	-11.06	1.48	1.59
35	BB	1492	C	P-O5'	-11.06	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2236	U	C2'-C1'	-11.06	1.41	1.53
34	BA	1726	U	O3'-P	-11.06	1.47	1.61
34	BA	1800	G	N9-C4	-11.06	1.29	1.38
34	BA	1665	G	P-O5'	-11.06	1.48	1.59
34	BA	927	A	P-O5'	-11.06	1.48	1.59
35	BB	87	G	O3'-P	-11.06	1.47	1.61
35	BB	1309	A	N9-C4	-11.06	1.31	1.37
85	AA	1493	A	N9-C4	-11.06	1.31	1.37
34	BA	1098	G	P-O5'	-11.05	1.48	1.59
35	BB	1447	U	O3'-P	-11.05	1.47	1.61
40	BG	37	G	P-O5'	-11.05	1.48	1.59
85	AA	1223	A	N9-C4	-11.05	1.31	1.37
34	BA	531	C	C2'-C1'	-11.05	1.41	1.53
34	BA	12	G	C2-N2	-11.05	1.23	1.34
35	BB	25	A	P-O5'	-11.04	1.48	1.59
85	AA	543	A	N9-C4	-11.04	1.31	1.37
34	BA	1613	G	P-O5'	-11.04	1.48	1.59
35	BB	576	A	C2'-C1'	-11.04	1.41	1.53
35	BB	642	G	O3'-P	-11.04	1.48	1.61
38	BE	96	G	C2-N2	-11.04	1.23	1.34
38	BE	141	A	O3'-P	-11.04	1.47	1.61
34	BA	14	G	N9-C4	-11.04	1.29	1.38
34	BA	1266	A	N9-C4	-11.04	1.31	1.37
85	AA	433	U	P-O5'	-11.04	1.48	1.59
34	BA	248	G	C2'-C1'	-11.04	1.41	1.53
34	BA	262	A	P-O5'	-11.04	1.48	1.59
34	BA	774	A	P-O5'	-11.04	1.48	1.59
34	BA	851	C	P-O5'	-11.04	1.48	1.59
35	BB	502	C	P-O5'	-11.04	1.48	1.59
34	BA	999	G	P-O5'	-11.04	1.48	1.59
34	BA	741	A	C1'-N9	-11.04	1.31	1.46
34	BA	955	G	P-O5'	-11.03	1.48	1.59
34	BA	1736	A	N7-C5	-11.03	1.32	1.39
36	BC	5	U	P-O5'	-11.03	1.48	1.59
85	AA	1119	A	O3'-P	-11.03	1.48	1.61
85	AA	1515	A	C1'-N9	-11.03	1.31	1.46
34	BA	390	A	N9-C4	-11.03	1.31	1.37
34	BA	1224	A	C4'-O4'	-11.03	1.31	1.45
35	BB	429	C	P-O5'	-11.03	1.48	1.59
85	AA	50	C	P-O5'	-11.03	1.48	1.59
85	AA	1136	A	C1'-N9	-11.03	1.31	1.46
35	BB	1145	G	N9-C4	-11.02	1.29	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1199	A	P-O5'	-11.02	1.48	1.59
34	BA	79	C	O3'-P	-11.02	1.48	1.61
34	BA	406	G	P-O5'	-11.02	1.48	1.59
85	AA	383	C	C2'-C1'	-11.02	1.41	1.53
85	AA	681	G	N9-C8	-11.02	1.30	1.37
85	AA	815	G	N9-C4	-11.02	1.46	1.38
34	BA	1673	G	C6-N1	-11.02	1.31	1.39
35	BB	1131	C	O3'-P	-11.02	1.48	1.61
35	BB	1371	G	O3'-P	-11.02	1.48	1.61
35	BB	1377	A	N9-C4	-11.02	1.31	1.37
34	BA	1845	G	C2-N3	-11.01	1.24	1.32
35	BB	1114	A	N9-C4	-11.01	1.31	1.37
85	AA	449	G	C2'-C1'	-11.01	1.41	1.53
34	BA	65	A	N7-C5	-11.01	1.32	1.39
35	BB	481	A	O3'-P	-11.01	1.48	1.61
34	BA	174	A	O3'-P	-11.01	1.48	1.61
34	BA	1052	G	O3'-P	-11.01	1.48	1.61
35	BB	569	G	C6-N1	-11.00	1.31	1.39
35	BB	6	A	P-O5'	-11.00	1.48	1.59
37	BD	75	G	P-O5'	-11.00	1.48	1.59
34	BA	445	C	C2'-C1'	-11.00	1.41	1.53
35	BB	1185	G	C1'-N9	-11.00	1.31	1.46
85	AA	1115	G	P-O5'	-11.00	1.48	1.59
34	BA	844	U	O3'-P	-11.00	1.48	1.61
35	BB	380	G	P-O5'	-11.00	1.48	1.59
85	AA	2195	A	O3'-P	-11.00	1.48	1.61
85	AA	2153	G	P-O5'	-10.99	1.48	1.59
85	AA	1895	C	P-O5'	-10.99	1.48	1.59
38	BE	172	U	O3'-P	-10.99	1.48	1.61
85	AA	464	A	N9-C4	-10.99	1.31	1.37
34	BA	1549	U	O3'-P	-10.99	1.48	1.61
40	BG	181	C	O3'-P	-10.99	1.48	1.61
34	BA	42	A	P-O5'	-10.98	1.48	1.59
34	BA	482	C	P-O5'	-10.98	1.48	1.59
34	BA	10	G	O3'-P	-10.98	1.48	1.61
35	BB	970	C	C2'-C1'	-10.98	1.41	1.53
34	BA	49	A	C1'-N9	-10.98	1.31	1.46
34	BA	924	U	P-O5'	-10.98	1.48	1.59
37	BD	48	G	C8-N7	-10.98	1.24	1.30
34	BA	301	U	O3'-P	-10.98	1.48	1.61
85	AA	1107	A	O3'-P	-10.98	1.48	1.61
35	BB	93	A	C2'-C1'	-10.97	1.41	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	167	U	O3'-P	-10.97	1.48	1.61
85	AA	1879	U	O3'-P	-10.97	1.48	1.61
85	AA	2062	U	O3'-P	-10.97	1.48	1.61
85	AA	2146	G	C2'-C1'	-10.97	1.41	1.53
34	BA	593	G	O3'-P	-10.97	1.48	1.61
35	BB	130	G	O3'-P	-10.97	1.48	1.61
36	BC	127	C	P-O5'	-10.97	1.48	1.59
38	BE	5	A	O3'-P	-10.97	1.48	1.61
38	BE	28	C	O3'-P	-10.97	1.48	1.61
35	BB	1171	U	O3'-P	-10.97	1.48	1.61
37	BD	92	G	P-O5'	-10.97	1.48	1.59
40	BG	122	G	O3'-P	-10.97	1.48	1.61
40	BG	176	G	O3'-P	-10.97	1.48	1.61
34	BA	383	G	C3'-C2'	-10.96	1.40	1.52
35	BB	1540	U	C2'-C1'	-10.96	1.41	1.53
34	BA	460	G	O3'-P	-10.96	1.48	1.61
35	BB	1016	C	C2'-C1'	-10.96	1.41	1.53
85	AA	362	G	C1'-N9	-10.96	1.31	1.46
35	BB	1167	C	O3'-P	-10.96	1.48	1.61
34	BA	1461	A	N9-C4	-10.96	1.31	1.37
35	BB	1089	A	N9-C4	-10.96	1.31	1.37
35	BB	1479	C	P-O5'	-10.96	1.48	1.59
85	AA	1288	A	N3-C4	-10.96	1.28	1.34
34	BA	143	A	P-O5'	-10.95	1.48	1.59
34	BA	1591	G	C5-C4	-10.95	1.30	1.38
35	BB	1378	U	P-O5'	-10.95	1.48	1.59
34	BA	1033	G	N9-C4	-10.95	1.29	1.38
34	BA	1493	U	C5'-C4'	10.95	1.64	1.51
34	BA	95	C	C3'-C2'	-10.95	1.40	1.52
34	BA	243	C	P-O5'	-10.95	1.48	1.59
35	BB	39	C	O3'-P	-10.95	1.48	1.61
40	BG	130	G	C1'-N9	-10.95	1.31	1.46
85	AA	1493	A	C2'-C1'	-10.95	1.41	1.53
34	BA	681	G	N9-C4	-10.94	1.29	1.38
35	BB	1051	U	C2-N3	-10.94	1.30	1.37
85	AA	1196	C	P-O5'	-10.94	1.48	1.59
85	AA	1457	C	C2'-C1'	-10.94	1.41	1.53
34	BA	997	U	C2-N3	-10.94	1.30	1.37
85	AA	1598	A	O3'-P	-10.94	1.48	1.61
34	BA	1614	G	C8-N7	-10.93	1.24	1.30
85	AA	160	A	C2'-C1'	-10.93	1.41	1.53
85	AA	2022	A	N7-C5	-10.93	1.32	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	8	C	N1-C2	-10.93	1.29	1.40
85	AA	1548	A	O3'-P	-10.93	1.48	1.61
34	BA	1612	C	O3'-P	-10.93	1.48	1.61
34	BA	1796	A	O3'-P	-10.93	1.48	1.61
85	AA	1913	G	P-O5'	-10.93	1.48	1.59
40	BG	148	C	P-O5'	-10.93	1.48	1.59
85	AA	790	A	C2'-C1'	-10.93	1.41	1.53
85	AA	1209	U	O3'-P	-10.93	1.48	1.61
35	BB	538	A	P-O5'	-10.92	1.48	1.59
34	BA	793	A	N7-C5	-10.92	1.32	1.39
35	BB	1057	G	P-O5'	-10.92	1.48	1.59
35	BB	1419	G	P-O5'	-10.92	1.48	1.59
34	BA	1252	G	O3'-P	-10.91	1.48	1.61
85	AA	364	C	C2'-C1'	-10.91	1.41	1.53
85	AA	498	C	O3'-P	-10.91	1.48	1.61
34	BA	1410	C	P-O5'	-10.91	1.48	1.59
85	AA	890	U	N1-C2	-10.91	1.28	1.38
35	BB	1350	A	O3'-P	-10.91	1.48	1.61
36	BC	102	G	O3'-P	-10.91	1.48	1.61
85	AA	454	G	O3'-P	-10.90	1.48	1.61
34	BA	911	G	C6-N1	-10.90	1.31	1.39
34	BA	788	C	O3'-P	-10.90	1.48	1.61
35	BB	630	A	O3'-P	-10.90	1.48	1.61
35	BB	1038	G	O3'-P	-10.90	1.48	1.61
85	AA	1115	G	C2'-C1'	-10.90	1.41	1.53
35	BB	1089	A	C1'-N9	-10.90	1.31	1.46
36	BC	97	U	O3'-P	-10.90	1.48	1.61
85	AA	2167	A	P-O5'	-10.90	1.48	1.59
34	BA	297	A	N9-C4	-10.90	1.31	1.37
85	AA	370	A	O3'-P	-10.90	1.48	1.61
85	AA	1266	C	P-O5'	-10.89	1.48	1.59
34	BA	1143	U	O3'-P	-10.89	1.48	1.61
35	BB	1310	C	O3'-P	-10.89	1.48	1.61
85	AA	2132	A	C2'-C1'	-10.89	1.41	1.53
34	BA	1018	U	O3'-P	-10.88	1.48	1.61
34	BA	1269	C	O3'-P	-10.88	1.48	1.61
35	BB	1045	G	C1'-N9	-10.88	1.31	1.46
36	BC	9	G	C1'-N9	-10.89	1.31	1.46
85	AA	1287	C	O3'-P	-10.88	1.48	1.61
34	BA	1518	A	N7-C5	-10.88	1.32	1.39
34	BA	1792	U	P-O5'	-10.88	1.48	1.59
85	AA	2030	U	P-O5'	-10.88	1.48	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1240	A	O3'-P	-10.88	1.48	1.61
34	BA	417	A	O3'-P	-10.88	1.48	1.61
35	BB	1323	U	P-O5'	-10.88	1.48	1.59
34	BA	406	G	N9-C4	-10.88	1.29	1.38
34	BA	440	A	N9-C4	-10.88	1.31	1.37
35	BB	1286	G	O3'-P	-10.88	1.48	1.61
36	BC	95	A	C2'-C1'	-10.87	1.41	1.53
35	BB	1466	A	N9-C8	-10.87	1.29	1.37
35	BB	1121	A	N9-C4	-10.87	1.31	1.37
35	BB	1134	G	N9-C4	-10.87	1.29	1.38
85	AA	2085	C	P-O5'	-10.87	1.48	1.59
34	BA	34	U	O3'-P	-10.86	1.48	1.61
34	BA	927	A	C1'-N9	-10.87	1.31	1.46
34	BA	478	G	C6-N1	-10.86	1.31	1.39
34	BA	775	C	P-O5'	-10.86	1.48	1.59
34	BA	900	A	C1'-N9	-10.86	1.31	1.46
35	BB	413	A	O3'-P	-10.86	1.48	1.61
35	BB	620	G	C2'-C1'	-10.86	1.41	1.53
34	BA	420	A	O3'-P	-10.86	1.48	1.61
34	BA	1324	G	P-O5'	-10.86	1.48	1.59
40	BG	147	U	P-O5'	-10.86	1.48	1.59
34	BA	1156	U	C2-N3	-10.86	1.30	1.37
35	BB	391	G	C6-N1	-10.86	1.31	1.39
35	BB	638	G	N9-C4	-10.86	1.29	1.38
85	AA	1490	A	N9-C4	-10.86	1.31	1.37
34	BA	1845	G	O3'-P	-10.86	1.48	1.61
34	BA	128	C	O3'-P	-10.85	1.48	1.61
34	BA	383	G	O3'-P	-10.85	1.48	1.61
35	BB	1135	U	O3'-P	-10.85	1.48	1.61
35	BB	1334	C	C3'-C2'	-10.85	1.40	1.52
85	AA	859	G	P-O5'	-10.85	1.48	1.59
85	AA	1098	C	O3'-P	-10.85	1.48	1.61
34	BA	218	G	P-O5'	-10.85	1.49	1.59
38	BE	59	U	C2-N3	-10.85	1.30	1.37
35	BB	873	C	P-O5'	-10.85	1.49	1.59
85	AA	315	U	O3'-P	-10.85	1.48	1.61
85	AA	863	C	P-O5'	-10.85	1.49	1.59
34	BA	1641	G	P-O5'	-10.84	1.49	1.59
35	BB	471	U	P-O5'	-10.84	1.49	1.59
34	BA	1274	A	N9-C4	-10.84	1.31	1.37
35	BB	971	A	C2'-C1'	-10.84	1.41	1.53
85	AA	48	G	C6-N1	-10.84	1.31	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	502	U	P-O5'	-10.84	1.49	1.59
34	BA	1163	G	C6-N1	-10.84	1.31	1.39
85	AA	624	A	N9-C4	-10.83	1.31	1.37
34	BA	876	C	O3'-P	-10.83	1.48	1.61
35	BB	85	A	O3'-P	-10.83	1.48	1.61
85	AA	523	U	P-O5'	-10.83	1.49	1.59
35	BB	119	G	O3'-P	-10.83	1.48	1.61
85	AA	595	A	N9-C4	-10.83	1.31	1.37
34	BA	538	G	C2'-C1'	-10.82	1.41	1.53
36	BC	110	A	O3'-P	-10.82	1.48	1.61
85	AA	369	A	P-O5'	-10.82	1.49	1.59
34	BA	1007	G	N9-C4	-10.82	1.29	1.38
34	BA	517	A	O3'-P	-10.81	1.48	1.61
35	BB	830	G	C2'-C1'	-10.81	1.41	1.53
85	AA	1259	U	P-O5'	-10.81	1.49	1.59
85	AA	365	G	C1'-N9	-10.81	1.31	1.46
85	AA	707	U	C4'-C3'	-10.81	1.41	1.53
35	BB	1122	C	P-O5'	-10.80	1.49	1.59
34	BA	1284	G	N1-C2	-10.80	1.29	1.37
34	BA	899	G	P-O5'	-10.80	1.49	1.59
34	BA	1671	A	C2'-C1'	-10.80	1.41	1.53
85	AA	1878	C	P-O5'	-10.80	1.49	1.59
34	BA	19	G	P-O5'	-10.80	1.49	1.59
35	BB	978	C	P-O5'	-10.80	1.49	1.59
34	BA	366	G	P-O5'	-10.79	1.49	1.59
34	BA	1602	A	N9-C4	-10.79	1.31	1.37
34	BA	1680	G	C6-N1	-10.79	1.31	1.39
34	BA	1730	A	C5'-C4'	10.79	1.64	1.51
37	BD	37	G	P-O5'	-10.79	1.49	1.59
85	AA	2138	G	C2'-C1'	-10.79	1.41	1.53
35	BB	561	C	P-O5'	-10.79	1.49	1.59
34	BA	1662	U	N3-C4	-10.79	1.28	1.38
35	BB	994	A	P-O5'	-10.79	1.49	1.59
39	BF	16	C	C5'-C4'	10.79	1.64	1.51
34	BA	1336	U	P-O5'	-10.78	1.49	1.59
85	AA	2146	G	N9-C8	-10.78	1.30	1.37
34	BA	1680	G	O3'-P	-10.78	1.48	1.61
35	BB	464	C	O3'-P	-10.78	1.48	1.61
36	BC	52	A	P-O5'	-10.78	1.49	1.59
34	BA	1220	C	O3'-P	-10.78	1.48	1.61
34	BA	1588	U	P-O5'	-10.78	1.49	1.59
40	BG	38	A	O3'-P	-10.77	1.48	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	758	C	P-O5'	-10.77	1.49	1.59
85	AA	966	G	P-O5'	-10.77	1.49	1.59
34	BA	596	G	N7-C5	-10.77	1.32	1.39
85	AA	447	C	O3'-P	-10.77	1.48	1.61
34	BA	95	C	O3'-P	-10.76	1.48	1.61
34	BA	1039	G	C3'-C2'	-10.76	1.41	1.52
34	BA	490	A	N9-C4	-10.76	1.31	1.37
35	BB	1403	G	O3'-P	-10.76	1.48	1.61
35	BB	1455	A	N7-C5	-10.76	1.32	1.39
34	BA	895	U	C4'-C3'	-10.76	1.41	1.53
85	AA	1198	U	O3'-P	-10.76	1.48	1.61
34	BA	1062	G	C6-N1	-10.76	1.32	1.39
34	BA	1809	G	C1'-N9	-10.76	1.31	1.46
85	AA	676	U	P-O5'	-10.76	1.49	1.59
34	BA	707	C	O3'-P	-10.75	1.48	1.61
34	BA	1001	G	O3'-P	-10.75	1.48	1.61
35	BB	1036	G	P-O5'	-10.75	1.49	1.59
85	AA	802	A	N9-C4	-10.75	1.31	1.37
38	BE	121	G	C2'-C1'	-10.75	1.41	1.53
41	BH	63	G	N7-C5	-10.75	1.32	1.39
85	AA	819	G	N7-C5	-10.75	1.32	1.39
34	BA	756	A	P-O5'	-10.75	1.49	1.59
34	BA	962	U	C2'-C1'	-10.75	1.41	1.53
35	BB	379	U	P-O5'	-10.75	1.49	1.59
35	BB	418	G	P-O5'	-10.75	1.49	1.59
35	BB	1123	A	N9-C4	-10.75	1.31	1.37
34	BA	876	C	C2'-C1'	-10.74	1.41	1.53
35	BB	1235	A	O3'-P	-10.74	1.48	1.61
40	BG	15	G	C2'-C1'	-10.74	1.41	1.53
40	BG	139	U	N3-C4	-10.74	1.28	1.38
35	BB	18	A	O3'-P	-10.74	1.48	1.61
35	BB	38	C	P-O5'	-10.74	1.49	1.59
85	AA	1828	C	P-O5'	-10.74	1.49	1.59
35	BB	650	A	P-O5'	-10.74	1.49	1.59
34	BA	962	U	O3'-P	-10.74	1.48	1.61
85	AA	687	G	C5-C4	-10.74	1.30	1.38
85	AA	1480	C	O3'-P	-10.74	1.48	1.61
85	AA	2191	C	P-O5'	-10.74	1.49	1.59
34	BA	619	U	P-O5'	-10.73	1.49	1.59
34	BA	1087	A	N9-C4	-10.73	1.31	1.37
85	AA	2210	C	P-O5'	-10.73	1.49	1.59
85	AA	1247	A	N9-C4	-10.73	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1527	G	O3'-P	-10.73	1.48	1.61
39	BF	51	C	P-O5'	-10.73	1.49	1.59
85	AA	702	G	C6-N1	-10.73	1.32	1.39
34	BA	1282	G	P-O5'	-10.72	1.49	1.59
85	AA	700	U	O3'-P	-10.72	1.48	1.61
85	AA	1923	A	N7-C5	-10.72	1.32	1.39
35	BB	634	A	N9-C4	-10.72	1.31	1.37
85	AA	456	A	C2'-C1'	-10.72	1.41	1.53
85	AA	1116	G	O3'-P	-10.72	1.48	1.61
34	BA	1094	U	P-O5'	-10.72	1.49	1.59
34	BA	856	G	P-O5'	-10.72	1.49	1.59
36	BC	38	U	O3'-P	-10.72	1.48	1.61
85	AA	1112	G	N7-C5	-10.72	1.32	1.39
34	BA	327	G	C2'-C1'	-10.72	1.41	1.53
85	AA	2038	C	P-O5'	-10.72	1.49	1.59
34	BA	1835	A	P-O5'	-10.71	1.49	1.59
35	BB	1194	A	C2'-C1'	-10.71	1.41	1.53
34	BA	224	G	O3'-P	-10.71	1.48	1.61
35	BB	825	U	C2'-C1'	-10.71	1.41	1.53
85	AA	667	A	N9-C4	-10.71	1.31	1.37
85	AA	2182	A	N7-C5	-10.71	1.32	1.39
35	BB	1424	G	N9-C4	-10.71	1.29	1.38
85	AA	526	G	P-O5'	-10.71	1.49	1.59
40	BG	166	C	P-O5'	-10.71	1.49	1.59
38	BE	118	C	O3'-P	-10.71	1.48	1.61
40	BG	54	G	P-O5'	-10.71	1.49	1.59
34	BA	895	U	O3'-P	-10.70	1.48	1.61
85	AA	1282	A	O3'-P	-10.70	1.48	1.61
34	BA	1504	A	P-O5'	-10.70	1.49	1.59
85	AA	1096	G	P-O5'	-10.70	1.49	1.59
85	AA	21	U	O3'-P	-10.70	1.48	1.61
85	AA	2083	G	O3'-P	-10.70	1.48	1.61
34	BA	999	G	O3'-P	-10.69	1.48	1.61
34	BA	1642	A	P-O5'	-10.69	1.49	1.59
85	AA	1644	G	N7-C5	-10.69	1.32	1.39
34	BA	886	G	O3'-P	-10.69	1.48	1.61
34	BA	910	U	O3'-P	-10.69	1.48	1.61
34	BA	934	G	O3'-P	-10.69	1.48	1.61
34	BA	321	G	N9-C8	-10.68	1.30	1.37
34	BA	341	U	C2-N3	-10.68	1.30	1.37
34	BA	384	U	P-O5'	-10.68	1.49	1.59
35	BB	837	A	N9-C4	-10.68	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1369	A	O3'-P	-10.68	1.48	1.61
85	AA	1291	A	N9-C4	-10.68	1.31	1.37
34	BA	1839	G	P-O5'	-10.67	1.49	1.59
35	BB	1045	G	C2'-C1'	-10.67	1.41	1.53
37	BD	36	C	C2'-C1'	-10.67	1.41	1.53
85	AA	2189	U	C2'-C1'	-10.67	1.41	1.53
34	BA	513	U	N3-C4	-10.67	1.28	1.38
34	BA	735	A	N9-C4	-10.67	1.31	1.37
85	AA	1255	C	P-O5'	-10.67	1.49	1.59
35	BB	2	C	P-O5'	-10.67	1.49	1.59
34	BA	722	A	N9-C4	-10.66	1.31	1.37
34	BA	972	C	O3'-P	-10.66	1.48	1.61
34	BA	1631	U	C5'-C4'	10.66	1.64	1.51
35	BB	441	G	O3'-P	-10.66	1.48	1.61
35	BB	606	C	O3'-P	-10.66	1.48	1.61
35	BB	1220	A	P-O5'	-10.66	1.49	1.59
37	BD	75	G	C2-N3	-10.66	1.24	1.32
40	BG	11	G	P-O5'	-10.66	1.49	1.59
34	BA	720	A	N7-C5	-10.66	1.32	1.39
85	AA	864	C	C2'-C1'	-10.66	1.41	1.53
34	BA	803	U	P-O5'	-10.66	1.49	1.59
35	BB	556	U	C2-N3	-10.66	1.30	1.37
38	BE	175	U	O3'-P	-10.66	1.48	1.61
40	BG	32	U	C2-N3	-10.66	1.30	1.37
34	BA	291	C	P-O5'	-10.65	1.49	1.59
34	BA	1211	G	N3-C4	-10.65	1.27	1.35
85	AA	57	G	P-O5'	-10.65	1.49	1.59
34	BA	1286	C	C2-N3	-10.65	1.27	1.35
35	BB	372	U	C2-N3	-10.65	1.30	1.37
41	BH	30	C	P-O5'	-10.65	1.49	1.59
34	BA	743	A	P-O5'	-10.65	1.49	1.59
35	BB	120	C	O3'-P	-10.65	1.48	1.61
34	BA	1143	U	C2-N3	-10.65	1.30	1.37
35	BB	69	A	N9-C4	-10.65	1.31	1.37
38	BE	141	A	C4'-C3'	-10.65	1.41	1.53
85	AA	471	U	O3'-P	-10.65	1.48	1.61
85	AA	47	A	O3'-P	-10.65	1.48	1.61
34	BA	1485	U	C2-N3	-10.64	1.30	1.37
35	BB	110	U	P-O5'	-10.64	1.49	1.59
34	BA	997	U	O3'-P	-10.64	1.48	1.61
85	AA	902	A	O4'-C1'	-10.64	1.27	1.41
85	AA	1466	U	N1-C2	-10.64	1.28	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	58	A	N9-C4	-10.64	1.31	1.37
85	AA	1146	C	P-O5'	-10.64	1.49	1.59
34	BA	1832	A	O3'-P	-10.64	1.48	1.61
35	BB	426	A	P-O5'	-10.64	1.49	1.59
40	BG	41	U	P-O5'	-10.64	1.49	1.59
85	AA	341	C	P-O5'	-10.64	1.49	1.59
34	BA	437	G	O3'-P	-10.64	1.48	1.61
34	BA	718	U	C2'-C1'	-10.63	1.41	1.53
85	AA	1911	A	O3'-P	-10.64	1.48	1.61
34	BA	1541	G	N9-C4	-10.63	1.29	1.38
34	BA	1617	U	C2-N3	-10.63	1.30	1.37
34	BA	1708	A	C3'-C2'	-10.63	1.41	1.52
35	BB	1307	C	O3'-P	-10.63	1.48	1.61
34	BA	1724	G	C2'-C1'	-10.63	1.41	1.53
35	BB	1117	G	C3'-C2'	-10.63	1.41	1.52
35	BB	119	G	C3'-C2'	-10.62	1.41	1.52
36	BC	113	G	N9-C4	-10.62	1.29	1.38
85	AA	1655	G	C2'-C1'	-10.62	1.41	1.53
85	AA	2139	G	O3'-P	-10.62	1.48	1.61
34	BA	1587	C	O3'-P	-10.62	1.48	1.61
35	BB	5	A	N9-C4	-10.62	1.31	1.37
36	BC	119	G	N9-C4	-10.61	1.29	1.38
85	AA	1283	C	P-O5'	-10.61	1.49	1.59
34	BA	253	U	C2-N3	-10.61	1.30	1.37
35	BB	28	G	P-O5'	-10.61	1.49	1.59
85	AA	2119	C	C3'-C2'	-10.61	1.41	1.52
35	BB	1462	G	C6-N1	-10.61	1.32	1.39
39	BF	25	G	O3'-P	-10.61	1.48	1.61
85	AA	1489	G	O3'-P	-10.61	1.48	1.61
34	BA	276	C	C2'-C1'	-10.61	1.41	1.53
34	BA	1502	G	C2'-C1'	-10.61	1.41	1.53
34	BA	926	A	C1'-N9	-10.60	1.32	1.46
34	BA	1272	U	O3'-P	-10.60	1.48	1.61
34	BA	798	G	N7-C5	-10.60	1.32	1.39
85	AA	352	G	C1'-N9	-10.60	1.32	1.46
85	AA	1123	C	P-O5'	-10.60	1.49	1.59
85	AA	1867	G	O3'-P	-10.60	1.48	1.61
35	BB	1331	U	O3'-P	-10.60	1.48	1.61
40	BG	165	C	P-O5'	-10.60	1.49	1.59
34	BA	1407	C	P-O5'	-10.59	1.49	1.59
85	AA	1819	U	P-O5'	-10.59	1.49	1.59
34	BA	370	U	P-O5'	-10.59	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	715	U	C2-N3	-10.59	1.30	1.37
35	BB	391	G	P-O5'	-10.59	1.49	1.59
35	BB	1370	G	P-O5'	-10.59	1.49	1.59
34	BA	190	U	P-O5'	-10.59	1.49	1.59
38	BE	192	A	P-O5'	-10.59	1.49	1.59
85	AA	273	C	P-O5'	-10.59	1.49	1.59
85	AA	1223	A	P-O5'	-10.59	1.49	1.59
34	BA	1270	G	O3'-P	-10.58	1.48	1.61
40	BG	139	U	C2-N3	-10.58	1.30	1.37
85	AA	689	U	P-O5'	-10.58	1.49	1.59
34	BA	191	G	O3'-P	-10.58	1.48	1.61
34	BA	930	A	P-O5'	-10.58	1.49	1.59
35	BB	135	C	C3'-C2'	-10.58	1.41	1.52
35	BB	491	A	O3'-P	-10.58	1.48	1.61
34	BA	1711	G	N9-C4	-10.58	1.29	1.38
34	BA	1069	U	C2'-C1'	-10.57	1.41	1.53
34	BA	1696	G	N9-C8	-10.57	1.30	1.37
41	BH	41	A	C1'-N9	-10.57	1.32	1.46
85	AA	1703	A	N9-C4	-10.57	1.31	1.37
34	BA	1175	G	C6-N1	-10.57	1.32	1.39
35	BB	1075	A	C2'-C1'	-10.57	1.41	1.53
34	BA	1173	C	P-O5'	-10.57	1.49	1.59
35	BB	1453	G	C2'-C1'	-10.57	1.41	1.53
85	AA	618	A	N9-C4	-10.57	1.31	1.37
34	BA	27	G	C6-N1	-10.57	1.32	1.39
34	BA	1540	C	C2-N3	10.57	1.44	1.35
35	BB	1033	U	C2-N3	-10.57	1.30	1.37
35	BB	1353	G	P-O5'	-10.57	1.49	1.59
85	AA	537	G	P-O5'	-10.57	1.49	1.59
85	AA	1560	A	N7-C5	-10.57	1.32	1.39
34	BA	1509	U	C2-N3	-10.56	1.30	1.37
34	BA	1542	A	N3-C4	-10.56	1.28	1.34
34	BA	1711	G	P-O5'	-10.56	1.49	1.59
41	BH	135	U	P-O5'	-10.56	1.49	1.59
85	AA	118	C	C2'-C1'	-10.56	1.41	1.53
85	AA	939	A	N9-C4	-10.56	1.31	1.37
34	BA	1694	C	O3'-P	-10.56	1.48	1.61
35	BB	1153	G	O3'-P	-10.56	1.48	1.61
85	AA	941	C	O3'-P	-10.56	1.48	1.61
35	BB	34	G	C6-N1	-10.55	1.32	1.39
85	AA	1209	U	P-O5'	-10.56	1.49	1.59
85	AA	1858	G	P-O5'	-10.56	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1080	U	O3'-P	-10.55	1.48	1.61
85	AA	880	A	C2'-C1'	-10.55	1.41	1.53
85	AA	1832	G	P-O5'	-10.55	1.49	1.59
85	AA	2199	G	C3'-C2'	-10.55	1.41	1.52
34	BA	695	A	C2'-C1'	-10.54	1.41	1.53
34	BA	1107	A	N9-C4	-10.54	1.31	1.37
85	AA	2174	G	C1'-N9	-10.54	1.32	1.46
40	BG	142	A	C1'-N9	-10.54	1.32	1.46
85	AA	366	A	N9-C4	-10.54	1.31	1.37
37	BD	81	C	C3'-C2'	-10.54	1.41	1.52
85	AA	725	G	P-O5'	-10.54	1.49	1.59
34	BA	438	A	N9-C4	-10.53	1.31	1.37
35	BB	9	G	P-O5'	-10.53	1.49	1.59
85	AA	180	A	N9-C4	-10.54	1.31	1.37
85	AA	1292	A	N9-C4	-10.53	1.31	1.37
35	BB	516	G	P-O5'	-10.53	1.49	1.59
85	AA	446	C	O3'-P	-10.53	1.48	1.61
85	AA	1167	G	C6-N1	-10.53	1.32	1.39
34	BA	1476	G	P-O5'	-10.53	1.49	1.59
34	BA	1603	A	N9-C4	-10.53	1.31	1.37
85	AA	1900	C	P-O5'	-10.53	1.49	1.59
35	BB	1534	U	P-O5'	-10.53	1.49	1.59
85	AA	766	G	O3'-P	-10.53	1.48	1.61
34	BA	465	A	N9-C4	-10.52	1.31	1.37
34	BA	529	A	N9-C4	-10.52	1.31	1.37
34	BA	933	U	P-O5'	-10.52	1.49	1.59
39	BF	15	U	O3'-P	-10.52	1.48	1.61
34	BA	3	G	C3'-C2'	-10.52	1.41	1.52
35	BB	1063	C	O3'-P	-10.52	1.48	1.61
35	BB	1520	C	P-O5'	-10.52	1.49	1.59
85	AA	160	A	P-O5'	-10.52	1.49	1.59
85	AA	974	U	C5'-C4'	10.52	1.64	1.51
36	BC	120	G	N7-C5	-10.51	1.32	1.39
40	BG	8	U	C2'-C1'	-10.51	1.41	1.53
34	BA	696	A	C2'-C1'	-10.51	1.41	1.53
34	BA	777	C	O3'-P	-10.51	1.48	1.61
37	BD	95	G	N9-C4	10.51	1.46	1.38
85	AA	418	G	P-O5'	-10.51	1.49	1.59
34	BA	1166	A	O3'-P	-10.51	1.48	1.61
35	BB	1326	U	C2'-C1'	-10.51	1.41	1.53
41	BH	29	G	C1'-N9	-10.51	1.32	1.46
85	AA	1292	A	O3'-P	-10.51	1.48	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	992	A	N7-C5	-10.50	1.32	1.39
34	BA	246	G	O3'-P	-10.50	1.48	1.61
85	AA	2056	C	P-O5'	-10.50	1.49	1.59
34	BA	1166	A	C3'-C2'	-10.50	1.41	1.52
34	BA	1203	G	N7-C5	-10.50	1.32	1.39
35	BB	541	U	C2-N3	-10.50	1.30	1.37
38	BE	26	G	O3'-P	-10.50	1.48	1.61
85	AA	789	A	C5'-C4'	10.50	1.64	1.51
38	BE	129	G	N7-C5	-10.50	1.32	1.39
34	BA	1262	A	P-O5'	-10.49	1.49	1.59
35	BB	52	G	P-O5'	-10.49	1.49	1.59
35	BB	709	G	C2'-C1'	-10.49	1.41	1.53
38	BE	124	G	N9-C4	-10.49	1.29	1.38
40	BG	181	C	C2'-C1'	-10.49	1.41	1.53
85	AA	878	U	P-O5'	-10.49	1.49	1.59
34	BA	1003	A	C1'-N9	-10.49	1.32	1.46
40	BG	178	G	O3'-P	-10.49	1.48	1.61
39	BF	16	C	C4'-C3'	10.49	1.64	1.53
40	BG	100	G	N7-C5	-10.49	1.32	1.39
85	AA	2187	G	C4'-C3'	-10.49	1.41	1.53
34	BA	196	A	C5-C4	-10.49	1.31	1.38
34	BA	796	G	N7-C5	-10.49	1.32	1.39
35	BB	1367	U	O3'-P	-10.49	1.48	1.61
35	BB	1441	C	O3'-P	-10.49	1.48	1.61
85	AA	34	G	O3'-P	-10.49	1.48	1.61
35	BB	66	G	O3'-P	-10.48	1.48	1.61
35	BB	1036	G	O3'-P	-10.48	1.48	1.61
85	AA	2172	A	O3'-P	-10.48	1.48	1.61
85	AA	2218	G	N7-C5	-10.48	1.32	1.39
34	BA	427	G	P-O5'	-10.48	1.49	1.59
34	BA	711	C	C5'-C4'	10.48	1.64	1.51
35	BB	1207	C	O3'-P	-10.48	1.48	1.61
38	BE	97	G	C3'-C2'	-10.48	1.41	1.52
85	AA	1264	U	O3'-P	-10.48	1.48	1.61
85	AA	2007	G	C2'-C1'	-10.48	1.41	1.53
34	BA	616	G	P-O5'	-10.48	1.49	1.59
34	BA	1046	G	P-O5'	-10.48	1.49	1.59
34	BA	308	C	P-O5'	-10.47	1.49	1.59
35	BB	647	U	C2-N3	-10.47	1.30	1.37
34	BA	1018	U	C4'-C3'	-10.47	1.41	1.53
35	BB	1039	A	N9-C4	-10.47	1.31	1.37
39	BF	53	G	O3'-P	-10.47	1.48	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	771	A	O3'-P	-10.47	1.48	1.61
85	AA	938	A	P-O5'	-10.47	1.49	1.59
34	BA	82	A	P-O5'	-10.47	1.49	1.59
34	BA	1218	G	N9-C4	10.47	1.46	1.38
35	BB	880	G	P-O5'	-10.47	1.49	1.59
38	BE	40	C	O3'-P	-10.47	1.48	1.61
85	AA	645	C	O3'-P	-10.47	1.48	1.61
34	BA	1175	G	P-O5'	-10.46	1.49	1.59
38	BE	60	C	O3'-P	-10.46	1.48	1.61
40	BG	89	A	O3'-P	-10.46	1.48	1.61
85	AA	428	G	N9-C4	-10.46	1.29	1.38
85	AA	960	G	N9-C8	-10.46	1.30	1.37
34	BA	326	A	N9-C4	-10.46	1.31	1.37
85	AA	768	C	O3'-P	-10.46	1.48	1.61
34	BA	504	A	O3'-P	-10.46	1.48	1.61
35	BB	1363	A	O3'-P	-10.46	1.48	1.61
36	BC	125	A	P-O5'	-10.46	1.49	1.59
38	BE	192	A	N7-C5	-10.46	1.32	1.39
34	BA	894	G	C1'-N9	-10.46	1.32	1.46
35	BB	1316	U	O3'-P	-10.46	1.48	1.61
40	BG	140	G	O3'-P	-10.46	1.48	1.61
85	AA	463	G	P-O5'	-10.46	1.49	1.59
85	AA	1288	A	P-O5'	-10.45	1.49	1.59
34	BA	370	U	O3'-P	-10.45	1.48	1.61
34	BA	1674	G	O3'-P	-10.45	1.48	1.61
35	BB	651	G	N9-C8	-10.45	1.30	1.37
35	BB	1190	U	P-O5'	-10.45	1.49	1.59
36	BC	37	U	C2'-C1'	-10.45	1.41	1.53
85	AA	96	C	P-O5'	-10.45	1.49	1.59
34	BA	352	G	O3'-P	-10.45	1.48	1.61
85	AA	396	U	C2-N3	-10.45	1.30	1.37
85	AA	496	C	C2-N3	-10.45	1.27	1.35
34	BA	414	A	N9-C4	-10.45	1.31	1.37
35	BB	633	C	O3'-P	-10.44	1.48	1.61
40	BG	33	G	C2'-C1'	-10.45	1.41	1.53
34	BA	1034	U	C2-N3	-10.44	1.30	1.37
34	BA	1795	A	N9-C4	-10.44	1.31	1.37
35	BB	1372	G	P-O5'	-10.44	1.49	1.59
37	BD	47	U	O3'-P	-10.44	1.48	1.61
85	AA	2086	C	O3'-P	-10.44	1.48	1.61
34	BA	57	A	N3-C4	-10.44	1.28	1.34
36	BC	141	C	P-O5'	-10.44	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	649	C	P-O5'	-10.44	1.49	1.59
40	BG	9	G	C3'-C2'	-10.44	1.41	1.52
35	BB	1512	C	P-O5'	-10.44	1.49	1.59
35	BB	77	A	N9-C4	-10.43	1.31	1.37
85	AA	860	C	C3'-C2'	-10.43	1.41	1.52
35	BB	120	C	C3'-C2'	-10.43	1.41	1.52
34	BA	78	U	C2'-C1'	-10.43	1.41	1.53
85	AA	63	G	O3'-P	-10.43	1.48	1.61
85	AA	424	A	C2'-C1'	-10.43	1.41	1.53
85	AA	743	C	C2'-C1'	-10.43	1.41	1.53
85	AA	1160	U	P-O5'	-10.43	1.49	1.59
85	AA	2174	G	P-O5'	-10.43	1.49	1.59
34	BA	1485	U	O3'-P	-10.43	1.48	1.61
40	BG	15	G	O3'-P	-10.43	1.48	1.61
85	AA	385	A	C1'-N9	-10.43	1.32	1.46
35	BB	1054	G	O3'-P	-10.42	1.48	1.61
85	AA	1263	G	N9-C4	-10.42	1.29	1.38
34	BA	949	C	O3'-P	-10.42	1.48	1.61
36	BC	27	U	P-O5'	-10.42	1.49	1.59
34	BA	1801	G	C1'-N9	-10.42	1.32	1.46
34	BA	535	G	N9-C4	10.42	1.46	1.38
34	BA	1680	G	N7-C5	-10.42	1.32	1.39
85	AA	1730	C	O3'-P	-10.42	1.48	1.61
34	BA	80	U	O3'-P	-10.41	1.48	1.61
35	BB	1427	A	O3'-P	-10.41	1.48	1.61
41	BH	104	U	C1'-N1	-10.41	1.32	1.46
40	BG	89	A	P-O5'	-10.41	1.49	1.59
34	BA	238	C	N1-C6	-10.41	1.30	1.37
34	BA	800	G	N7-C5	-10.41	1.33	1.39
34	BA	996	U	C2-N3	-10.41	1.30	1.37
35	BB	375	G	C1'-N9	-10.41	1.32	1.46
34	BA	146	G	C2'-C1'	-10.41	1.42	1.53
38	BE	56	U	P-O5'	-10.41	1.49	1.59
34	BA	894	G	C6-N1	-10.40	1.32	1.39
35	BB	404	A	P-O5'	-10.40	1.49	1.59
35	BB	1017	U	C2'-C1'	-10.40	1.42	1.53
35	BB	1450	G	P-O5'	-10.40	1.49	1.59
85	AA	2058	C	C2'-C1'	-10.40	1.42	1.53
34	BA	28	C	O3'-P	-10.40	1.48	1.61
34	BA	502	U	C3'-C2'	-10.40	1.41	1.52
35	BB	1474	A	C4'-C3'	10.40	1.64	1.53
36	BC	44	A	P-O5'	-10.40	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	750	A	N9-C4	-10.40	1.31	1.37
85	AA	1695	G	N9-C4	-10.40	1.29	1.38
34	BA	1335	A	O3'-P	-10.40	1.48	1.61
34	BA	1639	U	O3'-P	-10.40	1.48	1.61
35	BB	15	C	C2-N3	-10.40	1.27	1.35
35	BB	434	A	P-O5'	-10.40	1.49	1.59
35	BB	1254	G	O4'-C1'	-10.40	1.28	1.41
85	AA	420	C	C3'-C2'	-10.40	1.41	1.52
34	BA	1656	A	C5-C4	-10.39	1.31	1.38
85	AA	381	A	P-O5'	-10.39	1.49	1.59
85	AA	1483	A	C2'-C1'	-10.39	1.42	1.53
34	BA	331	G	C6-N1	-10.39	1.32	1.39
37	BD	91	U	C3'-C2'	-10.39	1.41	1.52
85	AA	384	C	O3'-P	-10.39	1.48	1.61
34	BA	247	U	C3'-C2'	10.39	1.64	1.52
34	BA	1714	A	N9-C4	-10.39	1.31	1.37
34	BA	1782	C	C2'-C1'	-10.39	1.42	1.53
41	BH	29	G	C8-N7	-10.39	1.24	1.30
85	AA	1516	A	C3'-C2'	-10.39	1.41	1.52
35	BB	568	A	N9-C4	-10.39	1.31	1.37
38	BE	18	U	C2-N3	-10.39	1.30	1.37
85	AA	507	C	C2-N3	-10.39	1.27	1.35
85	AA	1472	G	P-O5'	-10.39	1.49	1.59
85	AA	1955	U	P-O5'	-10.39	1.49	1.59
35	BB	1423	U	P-O5'	-10.38	1.49	1.59
38	BE	177	U	P-O5'	-10.38	1.49	1.59
85	AA	1483	A	N9-C4	-10.38	1.31	1.37
40	BG	38	A	P-O5'	-10.38	1.49	1.59
85	AA	12	U	P-O5'	-10.38	1.49	1.59
35	BB	1191	G	P-O5'	-10.38	1.49	1.59
36	BC	4	G	O3'-P	-10.38	1.48	1.61
86	AB	69	G	N9-C4	-10.38	1.29	1.38
85	AA	882	C	O3'-P	-10.38	1.48	1.61
85	AA	1797	U	P-O5'	-10.38	1.49	1.59
34	BA	513	U	P-O5'	-10.38	1.49	1.59
85	AA	411	U	C2-N3	-10.38	1.30	1.37
34	BA	1204	U	C2-N3	-10.38	1.30	1.37
35	BB	1237	C	C2-N3	-10.38	1.27	1.35
34	BA	336	A	O3'-P	-10.37	1.48	1.61
35	BB	1239	A	P-O5'	-10.37	1.49	1.59
85	AA	1257	A	C1'-N9	-10.38	1.32	1.46
85	AA	1682	U	O3'-P	-10.38	1.48	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	111	U	C3'-C2'	-10.37	1.41	1.52
34	BA	1591	G	N9-C8	-10.37	1.30	1.37
34	BA	1733	G	P-O5'	-10.37	1.49	1.59
35	BB	121	A	C3'-C2'	-10.37	1.41	1.52
85	AA	81	A	P-O5'	-10.37	1.49	1.59
85	AA	2173	A	P-O5'	-10.37	1.49	1.59
34	BA	780	U	C5'-C4'	10.37	1.63	1.51
85	AA	1215	A	P-O5'	-10.37	1.49	1.59
34	BA	679	U	C2-N3	-10.37	1.30	1.37
34	BA	1542	A	C1'-N9	-10.37	1.32	1.46
36	BC	2	A	P-O5'	-10.37	1.49	1.59
35	BB	688	U	P-O5'	-10.36	1.49	1.59
36	BC	101	U	P-O5'	-10.37	1.49	1.59
34	BA	1411	C	O3'-P	-10.36	1.48	1.61
37	BD	23	A	C2'-C1'	-10.36	1.42	1.53
38	BE	6	A	P-O5'	-10.36	1.49	1.59
34	BA	1599	A	C2'-C1'	-10.36	1.42	1.53
35	BB	119	G	C1'-N9	-10.36	1.32	1.46
35	BB	1432	U	C2-N3	-10.36	1.30	1.37
85	AA	1252	A	O3'-P	-10.36	1.48	1.61
34	BA	1439	C	P-O5'	-10.36	1.49	1.59
40	BG	149	U	O3'-P	-10.36	1.48	1.61
35	BB	787	A	P-O5'	-10.36	1.49	1.59
35	BB	1415	G	N7-C5	-10.36	1.33	1.39
40	BG	30	C	O3'-P	-10.36	1.48	1.61
85	AA	31	C	C2'-C1'	-10.36	1.42	1.53
34	BA	1501	U	C2'-C1'	-10.35	1.42	1.53
35	BB	1188	A	P-O5'	-10.35	1.49	1.59
35	BB	1370	G	C6-N1	-10.35	1.32	1.39
35	BB	1397	G	P-O5'	-10.35	1.49	1.59
36	BC	55	U	O3'-P	-10.35	1.48	1.61
40	BG	17	A	C2'-C1'	-10.35	1.42	1.53
34	BA	426	A	O3'-P	-10.35	1.48	1.61
34	BA	538	G	C2-N2	-10.35	1.24	1.34
85	AA	54	C	P-O5'	-10.35	1.49	1.59
35	BB	479	U	O3'-P	-10.35	1.48	1.61
36	BC	113	G	N7-C5	-10.35	1.33	1.39
85	AA	70	U	O3'-P	-10.35	1.48	1.61
85	AA	1490	A	N3-C4	-10.35	1.28	1.34
40	BG	163	G	N7-C5	-10.35	1.33	1.39
85	AA	766	G	P-O5'	-10.35	1.49	1.59
34	BA	909	G	N9-C4	-10.34	1.29	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	977	G	P-O5'	-10.34	1.49	1.59
35	BB	608	A	N9-C4	-10.34	1.31	1.37
35	BB	1023	G	N9-C4	-10.34	1.29	1.38
85	AA	404	A	O3'-P	-10.34	1.48	1.61
35	BB	1159	U	C2-N3	-10.34	1.30	1.37
35	BB	1377	A	P-O5'	-10.34	1.49	1.59
40	BG	158	A	N9-C4	-10.34	1.31	1.37
85	AA	362	G	O3'-P	-10.34	1.48	1.61
34	BA	663	U	P-O5'	-10.34	1.49	1.59
34	BA	979	G	N7-C5	-10.34	1.33	1.39
85	AA	363	A	O3'-P	-10.34	1.48	1.61
34	BA	1152	A	N9-C4	-10.34	1.31	1.37
34	BA	1807	G	C6-N1	-10.34	1.32	1.39
35	BB	1039	A	P-O5'	-10.34	1.49	1.59
85	AA	377	U	P-O5'	-10.34	1.49	1.59
85	AA	764	U	P-O5'	-10.34	1.49	1.59
34	BA	214	A	C1'-N9	-10.33	1.32	1.46
35	BB	778	A	C3'-C2'	-10.33	1.41	1.52
85	AA	363	A	C3'-C2'	-10.33	1.41	1.52
35	BB	416	U	N1-C2	-10.33	1.29	1.38
34	BA	859	G	N9-C4	-10.33	1.29	1.38
35	BB	836	U	O3'-P	-10.33	1.48	1.61
36	BC	15	G	N9-C4	-10.33	1.29	1.38
36	BC	55	U	P-O5'	-10.33	1.49	1.59
41	BH	110	C	C3'-C2'	-10.33	1.41	1.52
34	BA	596	G	C5'-C4'	10.32	1.63	1.51
34	BA	604	G	O3'-P	-10.32	1.48	1.61
34	BA	1706	A	C2'-C1'	-10.32	1.42	1.53
85	AA	529	G	O3'-P	-10.32	1.48	1.61
85	AA	985	G	P-O5'	-10.32	1.49	1.59
85	AA	362	G	C2'-C1'	-10.32	1.42	1.53
85	AA	1253	G	C2'-C1'	-10.32	1.42	1.53
85	AA	1670	U	P-O5'	-10.32	1.49	1.59
85	AA	2174	G	O3'-P	-10.32	1.48	1.61
34	BA	1338	G	N9-C4	-10.32	1.29	1.38
35	BB	610	U	P-O5'	-10.32	1.49	1.59
40	BG	94	G	N9-C4	-10.32	1.29	1.38
85	AA	259	A	N9-C4	-10.32	1.31	1.37
85	AA	466	A	N3-C4	-10.32	1.28	1.34
34	BA	1721	U	O4'-C1'	-10.32	1.28	1.41
36	BC	122	A	O3'-P	-10.32	1.48	1.61
34	BA	763	U	P-O5'	-10.32	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	151	A	O3'-P	-10.32	1.48	1.61
34	BA	992	A	C1'-N9	-10.31	1.32	1.46
34	BA	1065	U	C3'-C2'	-10.31	1.41	1.52
35	BB	1060	U	O3'-P	-10.31	1.48	1.61
85	AA	1515	A	C2'-C1'	-10.31	1.42	1.53
85	AA	2126	U	C2'-C1'	-10.31	1.42	1.53
34	BA	1711	G	C3'-C2'	-10.31	1.41	1.52
34	BA	1068	C	P-O5'	-10.31	1.49	1.59
85	AA	879	G	N9-C4	-10.31	1.29	1.38
85	AA	1578	G	C5'-C4'	10.31	1.63	1.51
34	BA	1103	G	O3'-P	-10.30	1.48	1.61
34	BA	1610	A	P-O5'	-10.30	1.49	1.59
85	AA	1892	G	P-O5'	-10.30	1.49	1.59
34	BA	1807	G	N7-C5	-10.30	1.33	1.39
40	BG	39	A	O3'-P	-10.30	1.48	1.61
40	BG	65	C	O3'-P	-10.30	1.48	1.61
40	BG	102	G	C6-N1	-10.30	1.32	1.39
34	BA	1464	C	P-O5'	-10.30	1.49	1.59
35	BB	572	G	N9-C8	-10.30	1.30	1.37
34	BA	33	C	O3'-P	-10.30	1.48	1.61
34	BA	98	A	C1'-N9	-10.30	1.32	1.46
39	BF	57	C	N1-C6	-10.30	1.30	1.37
34	BA	13	U	O3'-P	-10.30	1.48	1.61
34	BA	237	A	N9-C4	-10.29	1.31	1.37
34	BA	1006	G	C6-N1	-10.29	1.32	1.39
85	AA	531	G	C2'-C1'	-10.29	1.42	1.53
34	BA	965	A	O3'-P	-10.29	1.48	1.61
35	BB	505	G	C6-N1	-10.29	1.32	1.39
40	BG	14	G	C2'-C1'	-10.29	1.42	1.53
40	BG	137	G	O3'-P	-10.29	1.48	1.61
85	AA	17	C	P-O5'	-10.29	1.49	1.59
36	BC	149	A	O3'-P	-10.29	1.48	1.61
35	BB	1426	G	N1-C2	-10.29	1.29	1.37
37	BD	67	C	C2'-C1'	-10.29	1.42	1.53
41	BH	45	G	N9-C4	-10.29	1.29	1.38
85	AA	641	A	O3'-P	-10.28	1.48	1.61
34	BA	413	A	C2'-C1'	-10.28	1.42	1.53
85	AA	8	U	O3'-P	-10.28	1.48	1.61
85	AA	1471	G	N9-C4	-10.28	1.29	1.38
34	BA	1060	C	O3'-P	-10.28	1.48	1.61
85	AA	2174	G	C3'-C2'	-10.28	1.41	1.52
35	BB	802	G	C6-N1	-10.28	1.32	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1531	G	P-O5'	-10.28	1.49	1.59
85	AA	1265	C	P-O5'	-10.28	1.49	1.59
36	BC	69	U	O3'-P	-10.28	1.48	1.61
85	AA	497	G	O3'-P	-10.28	1.48	1.61
85	AA	1503	G	C1'-N9	-10.28	1.32	1.46
85	AA	1814	U	C2'-C1'	-10.28	1.42	1.53
35	BB	93	A	O3'-P	-10.27	1.48	1.61
85	AA	1494	C	P-O5'	-10.27	1.49	1.59
34	BA	139	U	C2-N3	-10.27	1.30	1.37
34	BA	440	A	C1'-N9	-10.27	1.32	1.46
34	BA	543	A	N9-C4	-10.27	1.31	1.37
34	BA	948	C	P-O5'	-10.27	1.49	1.59
34	BA	1432	C	P-O5'	-10.27	1.49	1.59
85	AA	32	U	P-O5'	-10.27	1.49	1.59
35	BB	828	G	C2'-C1'	-10.27	1.42	1.53
34	BA	875	G	C1'-N9	-10.27	1.32	1.46
34	BA	1546	C	C2'-C1'	-10.27	1.42	1.53
34	BA	1566	G	P-O5'	-10.27	1.49	1.59
34	BA	1609	U	C2-N3	-10.27	1.30	1.37
35	BB	10	C	P-O5'	-10.27	1.49	1.59
40	BG	83	U	O3'-P	-10.27	1.48	1.61
34	BA	384	U	C3'-C2'	-10.27	1.41	1.52
85	AA	1682	U	P-O5'	-10.27	1.49	1.59
34	BA	452	A	C1'-N9	-10.27	1.32	1.46
38	BE	63	C	O3'-P	-10.27	1.48	1.61
85	AA	292	C	C2'-C1'	-10.26	1.42	1.53
34	BA	56	G	O3'-P	-10.26	1.48	1.61
34	BA	372	U	C3'-C2'	-10.26	1.41	1.52
41	BH	127	A	N9-C4	-10.26	1.31	1.37
34	BA	65	A	C1'-N9	-10.26	1.32	1.46
34	BA	108	A	O3'-P	-10.26	1.48	1.61
34	BA	20	A	C2'-C1'	-10.26	1.42	1.53
34	BA	128	C	P-O5'	-10.26	1.49	1.59
35	BB	996	G	C1'-N9	-10.26	1.32	1.46
35	BB	1202	G	N9-C4	10.26	1.46	1.38
34	BA	1803	A	N9-C4	-10.26	1.31	1.37
35	BB	1178	A	N9-C4	-10.26	1.31	1.37
34	BA	38	G	C2'-C1'	-10.26	1.42	1.53
34	BA	975	A	C3'-C2'	-10.26	1.41	1.52
40	BG	14	G	C5-C4	-10.26	1.31	1.38
85	AA	596	A	P-O5'	-10.26	1.49	1.59
85	AA	1227	A	N9-C4	-10.26	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
39	BF	17	U	P-O5'	-10.25	1.49	1.59
41	BH	29	G	C2'-C1'	-10.25	1.42	1.53
35	BB	629	C	P-O5'	-10.25	1.49	1.59
35	BB	1117	G	O3'-P	-10.25	1.48	1.61
35	BB	1266	A	O3'-P	-10.25	1.48	1.61
34	BA	1211	G	C8-N7	-10.25	1.24	1.30
34	BA	514	U	C2'-C1'	-10.25	1.42	1.53
34	BA	950	C	O3'-P	-10.24	1.48	1.61
34	BA	1702	G	P-O5'	-10.24	1.49	1.59
85	AA	1536	C	C5'-C4'	10.24	1.63	1.51
85	AA	2209	U	P-O5'	-10.24	1.49	1.59
34	BA	266	G	C6-N1	-10.24	1.32	1.39
34	BA	1283	U	C2'-C1'	-10.24	1.42	1.53
85	AA	1244	A	N9-C4	-10.24	1.31	1.37
85	AA	1371	C	O3'-P	-10.24	1.48	1.61
35	BB	1383	C	N1-C6	-10.24	1.31	1.37
36	BC	49	G	C6-N1	-10.24	1.32	1.39
37	BD	71	G	C1'-N9	-10.24	1.32	1.46
35	BB	428	G	P-O5'	-10.23	1.49	1.59
35	BB	1157	G	O3'-P	-10.23	1.48	1.61
38	BE	180	G	P-O5'	-10.23	1.49	1.59
38	BE	116	U	O3'-P	-10.23	1.48	1.61
34	BA	788	C	C3'-C2'	-10.23	1.41	1.52
34	BA	1435	A	C1'-N9	-10.23	1.32	1.46
34	BA	1695	G	O3'-P	-10.23	1.48	1.61
38	BE	70	C	P-O5'	-10.23	1.49	1.59
39	BF	47	C	P-O5'	-10.23	1.49	1.59
85	AA	11	A	P-O5'	-10.23	1.49	1.59
85	AA	1502	A	N7-C5	-10.23	1.33	1.39
34	BA	24	C	O3'-P	-10.22	1.48	1.61
34	BA	259	C	P-O5'	-10.22	1.49	1.59
34	BA	1573	C	C2-N3	-10.22	1.27	1.35
85	AA	1102	C	O3'-P	-10.22	1.48	1.61
86	AB	16	U	C5'-C4'	10.22	1.63	1.51
34	BA	1845	G	N3-C4	-10.22	1.28	1.35
35	BB	465	C	O3'-P	-10.22	1.48	1.61
85	AA	1441	G	P-O5'	-10.22	1.49	1.59
34	BA	386	A	N9-C4	-10.22	1.31	1.37
35	BB	417	A	N9-C4	-10.22	1.31	1.37
34	BA	890	G	C1'-N9	-10.22	1.32	1.46
35	BB	995	C	O3'-P	-10.22	1.48	1.61
35	BB	1271	A	P-O5'	-10.22	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	131	C	P-O5'	-10.22	1.49	1.59
85	AA	508	C	C2'-C1'	-10.22	1.42	1.53
35	BB	443	A	N9-C4	-10.21	1.31	1.37
85	AA	1197	U	C2-N3	-10.22	1.30	1.37
35	BB	688	U	O3'-P	-10.21	1.48	1.61
85	AA	2133	A	C4'-C3'	-10.21	1.42	1.53
34	BA	1154	U	P-O5'	-10.21	1.49	1.59
85	AA	1148	G	C2'-C1'	-10.21	1.42	1.53
34	BA	1071	G	C3'-C2'	-10.21	1.41	1.52
39	BF	6	C	C3'-C2'	-10.21	1.41	1.52
40	BG	138	C	P-O5'	-10.21	1.49	1.59
85	AA	157	G	C1'-N9	-10.21	1.32	1.46
85	AA	1736	U	P-O5'	-10.21	1.49	1.59
35	BB	1399	A	N7-C5	-10.21	1.33	1.39
34	BA	389	U	C2-N3	-10.21	1.30	1.37
34	BA	896	U	C2-N3	10.21	1.44	1.37
34	BA	1025	A	C5-C4	-10.21	1.31	1.38
34	BA	269	G	C2-N2	-10.21	1.24	1.34
34	BA	1314	A	N9-C4	-10.21	1.31	1.37
35	BB	103	C	P-O5'	-10.20	1.49	1.59
34	BA	1434	U	P-O5'	-10.20	1.49	1.59
85	AA	659	A	C2'-C1'	-10.20	1.42	1.53
34	BA	1724	G	C3'-C2'	-10.20	1.41	1.52
34	BA	1739	G	C2-N3	-10.20	1.24	1.32
35	BB	816	U	N1-C2	10.20	1.47	1.38
85	AA	509	C	P-O5'	-10.20	1.49	1.59
34	BA	330	A	O3'-P	-10.20	1.49	1.61
38	BE	21	C	N1-C6	-10.20	1.31	1.37
85	AA	2082	C	C4-C5	-10.20	1.34	1.43
34	BA	897	U	O3'-P	-10.19	1.49	1.61
34	BA	1167	A	O3'-P	-10.19	1.49	1.61
85	AA	1140	G	C2'-C1'	-10.19	1.42	1.53
37	BD	10	C	O3'-P	-10.19	1.49	1.61
35	BB	8	U	O3'-P	-10.19	1.49	1.61
36	BC	53	A	C1'-N9	-10.19	1.32	1.46
37	BD	60	C	O3'-P	-10.19	1.49	1.61
35	BB	648	G	C4'-C3'	-10.19	1.42	1.53
36	BC	145	G	C6-N1	-10.19	1.32	1.39
40	BG	109	C	O3'-P	-10.19	1.49	1.61
85	AA	475	A	C1'-N9	-10.19	1.32	1.46
35	BB	100	A	N9-C4	-10.18	1.31	1.37
35	BB	572	G	N9-C4	-10.18	1.29	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	665	A	C2'-C1'	-10.18	1.42	1.53
85	AA	1220	A	O3'-P	-10.18	1.49	1.61
85	AA	1892	G	O3'-P	-10.18	1.49	1.61
34	BA	321	G	C8-N7	-10.18	1.24	1.30
34	BA	608	G	O4'-C1'	-10.18	1.28	1.41
41	BH	36	C	N1-C6	-10.18	1.31	1.37
34	BA	905	A	N9-C4	-10.18	1.31	1.37
85	AA	760	U	C3'-C2'	-10.18	1.41	1.52
85	AA	1991	C	C4-C5	-10.18	1.34	1.43
37	BD	81	C	P-O5'	-10.18	1.49	1.59
85	AA	640	C	P-O5'	-10.18	1.49	1.59
85	AA	2239	A	N9-C4	-10.18	1.31	1.37
37	BD	74	A	O3'-P	-10.17	1.49	1.61
34	BA	882	G	P-O5'	-10.17	1.49	1.59
34	BA	973	U	O3'-P	-10.17	1.49	1.61
34	BA	1142	C	O3'-P	-10.17	1.49	1.61
34	BA	475	A	C1'-N9	-10.17	1.32	1.46
34	BA	1682	A	O3'-P	-10.17	1.49	1.61
35	BB	1061	G	C6-N1	-10.17	1.32	1.39
37	BD	86	A	C3'-C2'	-10.17	1.41	1.52
35	BB	43	G	C2'-C1'	-10.16	1.42	1.53
35	BB	1019	C	C2'-C1'	-10.16	1.42	1.53
38	BE	200	A	O3'-P	-10.16	1.49	1.61
41	BH	112	U	P-O5'	-10.16	1.49	1.59
34	BA	1220	C	C2-N3	-10.16	1.27	1.35
35	BB	1064	U	P-O5'	-10.16	1.49	1.59
35	BB	1199	A	O3'-P	-10.16	1.49	1.61
35	BB	1211	C	O3'-P	-10.16	1.49	1.61
41	BH	19	G	P-O5'	-10.16	1.49	1.59
85	AA	2069	A	C2'-C1'	-10.16	1.42	1.53
34	BA	125	G	C6-N1	-10.16	1.32	1.39
34	BA	807	U	C2'-C1'	-10.16	1.42	1.53
35	BB	625	A	P-O5'	-10.16	1.49	1.59
35	BB	1384	A	C2'-C1'	-10.16	1.42	1.53
85	AA	117	C	C3'-C2'	-10.16	1.41	1.52
85	AA	1491	G	N9-C4	-10.16	1.29	1.38
85	AA	1671	G	P-O5'	-10.16	1.49	1.59
36	BC	107	C	C2-N3	-10.15	1.27	1.35
36	BC	149	A	P-O5'	-10.15	1.49	1.59
34	BA	1010	C	C3'-C2'	-10.15	1.41	1.52
35	BB	441	G	N9-C4	-10.15	1.29	1.38
35	BB	805	G	O3'-P	-10.15	1.49	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1146	C	P-O5'	-10.15	1.49	1.59
85	AA	1926	A	C2'-C1'	-10.15	1.42	1.53
34	BA	724	A	O3'-P	-10.15	1.49	1.61
38	BE	25	U	C3'-C2'	-10.14	1.41	1.52
38	BE	101	C	C2'-C1'	-10.14	1.42	1.53
38	BE	192	A	C2'-C1'	-10.14	1.42	1.53
38	BE	26	G	C4'-O4'	-10.14	1.32	1.45
39	BF	6	C	O3'-P	-10.14	1.49	1.61
35	BB	74	U	O3'-P	-10.14	1.49	1.61
85	AA	532	G	N9-C4	-10.14	1.29	1.38
85	AA	1973	G	O3'-P	-10.14	1.49	1.61
35	BB	431	U	O3'-P	-10.14	1.49	1.61
35	BB	1340	U	C2'-C1'	-10.14	1.42	1.53
35	BB	1392	A	P-O5'	-10.14	1.49	1.59
40	BG	134	U	O3'-P	-10.14	1.49	1.61
34	BA	461	A	N7-C5	-10.14	1.33	1.39
38	BE	176	G	O3'-P	-10.13	1.49	1.61
85	AA	1218	C	P-O5'	-10.13	1.49	1.59
34	BA	430	A	N9-C4	-10.13	1.31	1.37
34	BA	1022	C	O3'-P	-10.13	1.49	1.61
35	BB	1292	G	P-O5'	-10.13	1.49	1.59
85	AA	1553	G	P-O5'	-10.13	1.49	1.59
34	BA	479	U	O3'-P	-10.13	1.49	1.61
34	BA	1696	G	C6-N1	-10.13	1.32	1.39
36	BC	104	A	P-O5'	-10.13	1.49	1.59
38	BE	110	U	P-O5'	-10.13	1.49	1.59
34	BA	1663	U	O3'-P	-10.13	1.49	1.61
35	BB	1217	C	O3'-P	-10.13	1.49	1.61
36	BC	52	A	O3'-P	-10.13	1.49	1.61
38	BE	158	U	O3'-P	-10.12	1.49	1.61
34	BA	1696	G	C3'-C2'	-10.12	1.41	1.52
37	BD	6	C	O3'-P	-10.12	1.49	1.61
37	BD	78	C	C2-N3	-10.12	1.27	1.35
85	AA	95	U	C2-N3	-10.12	1.30	1.37
34	BA	322	U	O3'-P	-10.12	1.49	1.61
34	BA	415	C	P-O5'	-10.12	1.49	1.59
34	BA	1694	C	P-O5'	-10.12	1.49	1.59
34	BA	1706	A	N3-C4	-10.12	1.28	1.34
35	BB	1151	A	O3'-P	-10.12	1.49	1.61
40	BG	25	G	C2'-C1'	-10.12	1.42	1.53
41	BH	33	G	C1'-N9	-10.12	1.32	1.46
85	AA	618	A	C2'-C1'	-10.12	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1979	A	O3'-P	-10.12	1.49	1.61
34	BA	42	A	O3'-P	-10.12	1.49	1.61
38	BE	60	C	C2-N3	-10.12	1.27	1.35
85	AA	535	G	N9-C4	-10.12	1.29	1.38
85	AA	572	G	C2'-C1'	-10.12	1.42	1.53
34	BA	772	G	N9-C8	-10.11	1.30	1.37
40	BG	59	G	P-O5'	-10.11	1.49	1.59
85	AA	675	A	O3'-P	-10.11	1.49	1.61
85	AA	1671	G	O3'-P	-10.11	1.49	1.61
85	AA	396	U	O3'-P	-10.11	1.49	1.61
85	AA	2228	G	C2'-C1'	-10.11	1.42	1.53
40	BG	58	G	C4'-C3'	-10.11	1.42	1.53
85	AA	373	G	O3'-P	-10.11	1.49	1.61
85	AA	2129	U	C2'-C1'	-10.11	1.42	1.53
36	BC	17	U	C3'-C2'	-10.11	1.41	1.52
85	AA	1673	A	P-O5'	-10.11	1.49	1.59
85	AA	2213	A	C2'-C1'	-10.11	1.42	1.53
34	BA	1088	G	N9-C4	-10.11	1.29	1.38
85	AA	1890	C	C2'-C1'	-10.11	1.42	1.53
35	BB	434	A	N9-C4	-10.11	1.31	1.37
38	BE	130	G	P-O5'	-10.11	1.49	1.59
34	BA	49	A	O3'-P	-10.10	1.49	1.61
34	BA	543	A	C2'-C1'	-10.10	1.42	1.53
34	BA	912	G	P-O5'	-10.10	1.49	1.59
34	BA	1441	C	O3'-P	-10.10	1.49	1.61
34	BA	1594	G	C2'-C1'	-10.10	1.42	1.53
36	BC	130	U	P-O5'	-10.10	1.49	1.59
34	BA	62	A	O3'-P	-10.10	1.49	1.61
34	BA	894	G	C2-N2	-10.10	1.24	1.34
34	BA	1092	U	P-O5'	-10.10	1.49	1.59
35	BB	1443	C	C2'-C1'	-10.10	1.42	1.53
38	BE	116	U	P-O5'	-10.10	1.49	1.59
40	BG	70	C	C2-N3	-10.10	1.27	1.35
85	AA	1125	G	C1'-N9	-10.10	1.32	1.46
34	BA	277	A	P-O5'	-10.10	1.49	1.59
34	BA	1518	A	C1'-N9	-10.10	1.32	1.46
35	BB	642	G	C6-N1	-10.10	1.32	1.39
35	BB	1401	G	P-O5'	-10.10	1.49	1.59
35	BB	587	A	C2'-C1'	-10.09	1.42	1.53
85	AA	1217	U	C2'-C1'	-10.09	1.42	1.53
35	BB	1170	U	P-O5'	-10.09	1.49	1.59
38	BE	94	U	C3'-C2'	-10.09	1.41	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	83	G	N7-C5	-10.09	1.33	1.39
34	BA	1462	U	C3'-C2'	-10.09	1.41	1.52
85	AA	1506	U	O3'-P	-10.09	1.49	1.61
34	BA	366	G	C6-N1	-10.09	1.32	1.39
34	BA	909	G	C4'-C3'	-10.09	1.42	1.53
85	AA	1586	C	C3'-C2'	-10.09	1.41	1.52
35	BB	1015	U	C2'-C1'	-10.09	1.42	1.53
38	BE	27	A	C2'-C1'	-10.09	1.42	1.53
34	BA	1011	G	C2-N3	-10.08	1.24	1.32
35	BB	1507	U	O3'-P	-10.08	1.49	1.61
34	BA	386	A	N7-C5	-10.08	1.33	1.39
34	BA	98	A	C2'-C1'	-10.08	1.42	1.53
34	BA	1253	G	C6-N1	-10.08	1.32	1.39
34	BA	1676	A	C2'-C1'	-10.08	1.42	1.53
85	AA	131	C	P-O5'	-10.08	1.49	1.59
85	AA	579	U	C2'-C1'	-10.08	1.42	1.53
85	AA	817	G	O3'-P	-10.07	1.49	1.61
34	BA	84	U	P-O5'	-10.07	1.49	1.59
36	BC	166	G	P-O5'	-10.07	1.49	1.59
85	AA	1001	G	P-O5'	-10.07	1.49	1.59
34	BA	742	C	C2-N3	-10.07	1.27	1.35
34	BA	1837	U	P-O5'	-10.07	1.49	1.59
38	BE	87	U	P-O5'	-10.07	1.49	1.59
85	AA	2037	A	N9-C4	-10.07	1.31	1.37
34	BA	106	U	O3'-P	-10.07	1.49	1.61
34	BA	269	G	N3-C4	-10.07	1.28	1.35
34	BA	860	G	O3'-P	-10.07	1.49	1.61
35	BB	489	A	N9-C4	-10.07	1.31	1.37
35	BB	1342	C	C2'-C1'	-10.07	1.42	1.53
35	BB	698	C	O3'-P	-10.07	1.49	1.61
85	AA	1211	C	C2'-C1'	-10.07	1.42	1.53
34	BA	132	U	C2'-C1'	-10.06	1.42	1.53
34	BA	748	C	C3'-C2'	-10.06	1.41	1.52
34	BA	1789	A	N9-C4	-10.06	1.31	1.37
39	BF	34	C	C2'-C1'	-10.06	1.42	1.53
85	AA	107	A	N9-C8	-10.06	1.29	1.37
34	BA	1463	U	O3'-P	-10.06	1.49	1.61
36	BC	31	A	N9-C4	-10.06	1.31	1.37
85	AA	2204	A	C2'-C1'	-10.06	1.42	1.53
85	AA	2149	C	P-O5'	-10.06	1.49	1.59
34	BA	1572	G	O3'-P	-10.05	1.49	1.61
35	BB	85	A	N9-C4	-10.06	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2108	C	C2'-C1'	-10.06	1.42	1.53
34	BA	1164	C	C2'-C1'	-10.05	1.42	1.53
34	BA	1519	G	P-O5'	-10.05	1.49	1.59
35	BB	416	U	C3'-C2'	-10.05	1.41	1.52
85	AA	1461	A	P-O5'	-10.05	1.49	1.59
34	BA	784	C	C2'-C1'	-10.05	1.42	1.53
35	BB	1214	U	O3'-P	-10.05	1.49	1.61
34	BA	1501	U	C2-N3	-10.05	1.30	1.37
35	BB	680	A	C3'-C2'	-10.05	1.41	1.52
35	BB	1470	G	C5'-C4'	10.05	1.63	1.51
34	BA	926	A	C3'-C2'	-10.05	1.41	1.52
41	BH	118	U	O3'-P	-10.05	1.49	1.61
85	AA	366	A	O3'-P	-10.05	1.49	1.61
85	AA	1495	G	C3'-C2'	-10.05	1.41	1.52
34	BA	436	U	O3'-P	-10.05	1.49	1.61
85	AA	705	G	C3'-C2'	-10.05	1.41	1.52
34	BA	420	A	P-O5'	-10.04	1.49	1.59
34	BA	790	G	O3'-P	-10.04	1.49	1.61
35	BB	1205	A	N7-C5	-10.04	1.33	1.39
35	BB	542	A	O3'-P	-10.04	1.49	1.61
35	BB	1181	A	C1'-N9	-10.04	1.32	1.46
85	AA	398	U	P-O5'	-10.04	1.49	1.59
85	AA	690	G	P-O5'	-10.04	1.49	1.59
85	AA	1921	G	N9-C4	10.04	1.46	1.38
34	BA	1817	G	O3'-P	-10.03	1.49	1.61
35	BB	15	C	P-O5'	-10.03	1.49	1.59
85	AA	2180	C	O3'-P	-10.03	1.49	1.61
34	BA	292	C	C2'-C1'	-10.03	1.42	1.53
34	BA	1454	G	C2'-C1'	-10.03	1.42	1.53
37	BD	11	A	O3'-P	-10.03	1.49	1.61
85	AA	90	A	N9-C4	-10.03	1.31	1.37
34	BA	357	A	C2'-C1'	-10.03	1.42	1.53
34	BA	400	A	C3'-C2'	-10.03	1.41	1.52
34	BA	532	C	C2'-C1'	-10.03	1.42	1.53
34	BA	605	G	C6-N1	-10.03	1.32	1.39
34	BA	744	G	C2'-C1'	-10.03	1.42	1.53
35	BB	1099	U	C2-N3	-10.03	1.30	1.37
36	BC	123	G	N9-C4	-10.03	1.29	1.38
35	BB	999	G	C2'-C1'	-10.03	1.42	1.53
85	AA	84	C	P-O5'	-10.03	1.49	1.59
85	AA	196	U	C2-N3	-10.03	1.30	1.37
34	BA	681	G	P-O5'	-10.02	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	944	G	O3'-P	-10.02	1.49	1.61
36	BC	168	C	O3'-P	-10.02	1.49	1.61
34	BA	1585	A	O3'-P	-10.02	1.49	1.61
35	BB	137	A	N9-C4	-10.02	1.31	1.37
85	AA	1118	U	C2-N3	-10.02	1.30	1.37
34	BA	366	G	O3'-P	-10.02	1.49	1.61
34	BA	917	C	P-O5'	-10.02	1.49	1.59
34	BA	245	U	P-O5'	-10.02	1.49	1.59
34	BA	1174	A	O3'-P	-10.02	1.49	1.61
34	BA	1259	C	P-O5'	-10.02	1.49	1.59
34	BA	1676	A	C1'-N9	-10.02	1.32	1.46
35	BB	813	C	P-O5'	-10.02	1.49	1.59
39	BF	32	G	N7-C5	-10.02	1.33	1.39
34	BA	1690	U	P-O5'	-10.01	1.49	1.59
35	BB	489	A	N3-C4	-10.01	1.28	1.34
85	AA	737	G	P-O5'	-10.01	1.49	1.59
85	AA	1699	A	O3'-P	-10.01	1.49	1.61
34	BA	1585	A	C1'-N9	-10.01	1.32	1.46
40	BG	26	G	P-O5'	-10.01	1.49	1.59
34	BA	370	U	C2-N3	-10.00	1.30	1.37
35	BB	64	U	O3'-P	-10.00	1.49	1.61
36	BC	14	G	P-O5'	-10.00	1.49	1.59
36	BC	152	C	N1-C2	-10.00	1.30	1.40
37	BD	98	G	N9-C4	-10.00	1.29	1.38
85	AA	2176	U	O3'-P	-10.00	1.49	1.61
36	BC	153	C	P-O5'	-10.00	1.49	1.59
38	BE	104	G	C6-N1	-10.00	1.32	1.39
85	AA	120	C	P-O5'	-10.00	1.49	1.59
34	BA	9	A	O3'-P	-10.00	1.49	1.61
34	BA	691	A	N9-C4	-10.00	1.31	1.37
85	AA	1563	U	P-O5'	-10.00	1.49	1.59
35	BB	1423	U	C2'-C1'	-9.99	1.42	1.53
85	AA	2218	G	C1'-N9	-9.99	1.32	1.46
34	BA	1293	A	P-O5'	-9.99	1.49	1.59
85	AA	474	C	C3'-C2'	-9.99	1.41	1.52
34	BA	471	U	C4'-C3'	-9.99	1.42	1.53
34	BA	1820	G	N9-C4	-9.99	1.29	1.38
34	BA	1506	C	C3'-C2'	-9.99	1.41	1.52
40	BG	94	G	P-O5'	-9.99	1.49	1.59
41	BH	11	C	O3'-P	-9.99	1.49	1.61
85	AA	1125	G	C2'-C1'	-9.99	1.42	1.53
34	BA	963	G	C2'-C1'	-9.98	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1310	C	P-O5'	-9.98	1.49	1.59
34	BA	1808	A	N9-C4	-9.98	1.31	1.37
35	BB	491	A	C3'-C2'	-9.98	1.41	1.52
35	BB	428	G	C2'-C1'	-9.98	1.42	1.53
34	BA	1795	A	C1'-N9	-9.98	1.32	1.46
85	AA	568	C	P-O5'	-9.98	1.49	1.59
35	BB	118	A	P-O5'	-9.98	1.49	1.59
37	BD	74	A	P-O5'	-9.98	1.49	1.59
85	AA	1216	A	C3'-C2'	-9.98	1.41	1.52
34	BA	1700	C	O3'-P	-9.98	1.49	1.61
85	AA	1136	A	C2'-C1'	-9.98	1.42	1.53
34	BA	1106	A	N9-C4	-9.97	1.31	1.37
34	BA	981	A	O3'-P	-9.97	1.49	1.61
34	BA	1559	C	O3'-P	-9.97	1.49	1.61
35	BB	136	A	O3'-P	-9.97	1.49	1.61
35	BB	662	G	C1'-N9	-9.97	1.32	1.46
85	AA	481	A	N9-C4	-9.97	1.31	1.37
85	AA	1676	G	N9-C4	-9.97	1.29	1.38
86	AB	66	U	O3'-P	-9.97	1.49	1.61
34	BA	89	G	N9-C4	-9.97	1.29	1.38
85	AA	2100	A	C2'-C1'	-9.97	1.42	1.53
34	BA	1165	A	P-O5'	-9.97	1.49	1.59
35	BB	1229	A	N9-C4	-9.97	1.31	1.37
85	AA	542	G	C2'-C1'	-9.97	1.42	1.53
85	AA	644	A	P-O5'	-9.97	1.49	1.59
85	AA	1104	G	O3'-P	-9.97	1.49	1.61
34	BA	1529	G	P-O5'	-9.97	1.49	1.59
35	BB	5	A	C6-N1	-9.97	1.28	1.35
35	BB	6	A	C2'-C1'	-9.97	1.42	1.53
85	AA	517	A	N9-C4	-9.97	1.31	1.37
35	BB	1445	A	O3'-P	-9.97	1.49	1.61
38	BE	26	G	N9-C4	-9.97	1.29	1.38
34	BA	340	U	C2-N3	-9.97	1.30	1.37
34	BA	889	U	P-O5'	-9.96	1.49	1.59
34	BA	1609	U	C3'-C2'	-9.97	1.41	1.52
34	BA	1568	A	N9-C4	-9.96	1.31	1.37
34	BA	1592	U	C2-N3	-9.96	1.30	1.37
36	BC	18	G	N3-C4	-9.97	1.28	1.35
41	BH	22	A	O3'-P	-9.97	1.49	1.61
37	BD	16	U	C2-N3	-9.96	1.30	1.37
85	AA	902	A	P-O5'	-9.96	1.49	1.59
85	AA	1458	G	C6-N1	-9.96	1.32	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2022	A	C6-N1	-9.96	1.28	1.35
34	BA	35	U	C3'-C2'	-9.96	1.41	1.52
35	BB	564	U	P-O5'	-9.96	1.49	1.59
38	BE	171	U	O3'-P	-9.96	1.49	1.61
85	AA	447	C	C2'-C1'	-9.96	1.42	1.53
85	AA	1967	A	O3'-P	-9.96	1.49	1.61
35	BB	504	C	O3'-P	-9.96	1.49	1.61
35	BB	436	G	P-O5'	-9.95	1.49	1.59
41	BH	19	G	O3'-P	-9.96	1.49	1.61
34	BA	281	C	C2-N3	-9.95	1.27	1.35
85	AA	921	C	P-O5'	-9.95	1.49	1.59
34	BA	371	U	P-O5'	-9.95	1.49	1.59
34	BA	1475	G	C2'-C1'	-9.95	1.42	1.53
85	AA	2170	G	P-O5'	-9.95	1.49	1.59
34	BA	60	A	P-O5'	-9.95	1.49	1.59
35	BB	368	C	C2'-C1'	-9.95	1.42	1.53
34	BA	888	G	O3'-P	-9.95	1.49	1.61
41	BH	21	G	C2'-C1'	-9.95	1.42	1.53
34	BA	1256	A	C1'-N9	-9.94	1.32	1.46
85	AA	161	A	O3'-P	-9.95	1.49	1.61
34	BA	141	G	C2-N2	-9.94	1.24	1.34
34	BA	1451	A	C4'-C3'	-9.94	1.42	1.53
34	BA	1583	A	C1'-N9	-9.94	1.32	1.46
35	BB	490	G	N9-C4	-9.94	1.29	1.38
40	BG	79	U	C2'-C1'	-9.94	1.42	1.53
41	BH	104	U	O3'-P	-9.94	1.49	1.61
85	AA	1496	U	C2-N3	-9.94	1.30	1.37
85	AA	2039	G	O3'-P	-9.94	1.49	1.61
34	BA	1169	A	P-O5'	-9.94	1.49	1.59
41	BH	18	C	P-O5'	-9.94	1.49	1.59
34	BA	1245	C	O3'-P	-9.94	1.49	1.61
41	BH	109	G	C4'-C3'	-9.94	1.42	1.53
35	BB	13	A	P-O5'	-9.93	1.49	1.59
85	AA	456	A	C5-C4	-9.93	1.31	1.38
36	BC	3	C	O3'-P	-9.93	1.49	1.61
85	AA	972	G	C5'-C4'	9.93	1.63	1.51
38	BE	67	A	P-O5'	-9.93	1.49	1.59
34	BA	965	A	N9-C4	-9.93	1.31	1.37
34	BA	1278	A	O3'-P	-9.93	1.49	1.61
35	BB	397	C	O3'-P	-9.93	1.49	1.61
35	BB	574	G	P-O5'	-9.93	1.49	1.59
35	BB	1186	A	C3'-C2'	-9.93	1.41	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	42	A	P-O5'	-9.93	1.49	1.59
38	BE	124	G	C1'-N9	-9.93	1.32	1.46
34	BA	49	A	N7-C5	-9.93	1.33	1.39
34	BA	158	U	P-O5'	-9.93	1.49	1.59
34	BA	1557	G	C6-N1	-9.93	1.32	1.39
85	AA	161	A	C2'-C1'	-9.93	1.42	1.53
85	AA	1544	G	C1'-N9	-9.93	1.32	1.46
34	BA	480	G	O3'-P	-9.92	1.49	1.61
85	AA	1821	C	P-O5'	-9.92	1.49	1.59
34	BA	799	A	N9-C4	-9.92	1.31	1.37
34	BA	900	A	C5-C4	-9.92	1.31	1.38
34	BA	1808	A	O3'-P	-9.92	1.49	1.61
34	BA	882	G	C2'-C1'	-9.92	1.42	1.53
34	BA	1801	G	C2'-C1'	-9.92	1.42	1.53
35	BB	50	A	N9-C4	-9.92	1.31	1.37
85	AA	913	U	P-O5'	-9.92	1.49	1.59
34	BA	971	G	C1'-N9	-9.91	1.32	1.46
35	BB	1062	G	P-O5'	-9.91	1.49	1.59
34	BA	82	A	O3'-P	-9.91	1.49	1.61
34	BA	1532	G	C6-N1	-9.91	1.32	1.39
35	BB	618	U	C3'-C2'	-9.91	1.41	1.52
35	BB	1321	G	O3'-P	-9.91	1.49	1.61
85	AA	64	A	P-O5'	-9.91	1.49	1.59
85	AA	2232	A	P-O5'	-9.91	1.49	1.59
36	BC	138	C	P-O5'	-9.91	1.49	1.59
38	BE	123	A	C8-N7	-9.91	1.24	1.31
35	BB	43	G	N9-C4	-9.91	1.30	1.38
41	BH	13	C	C2'-C1'	-9.91	1.42	1.53
34	BA	397	A	C1'-N9	-9.91	1.32	1.46
35	BB	2	C	C4'-C3'	-9.91	1.42	1.53
34	BA	718	U	P-O5'	-9.90	1.49	1.59
34	BA	760	G	P-O5'	-9.90	1.49	1.59
35	BB	673	C	P-O5'	-9.90	1.49	1.59
35	BB	1097	U	P-O5'	-9.90	1.49	1.59
85	AA	1430	A	P-O5'	-9.90	1.49	1.59
34	BA	904	G	C1'-N9	-9.90	1.32	1.46
34	BA	1578	A	C1'-N9	-9.90	1.32	1.46
35	BB	1396	G	N9-C4	-9.90	1.30	1.38
35	BB	1409	G	P-O5'	-9.90	1.49	1.59
40	BG	33	G	C3'-C2'	-9.90	1.41	1.52
36	BC	106	G	C6-N1	-9.90	1.32	1.39
41	BH	37	U	P-O5'	-9.90	1.49	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1252	A	C2'-C1'	-9.90	1.42	1.53
34	BA	428	C	O3'-P	-9.90	1.49	1.61
34	BA	669	U	P-O5'	-9.90	1.49	1.59
37	BD	57	C	P-O5'	-9.90	1.49	1.59
40	BG	112	C	O3'-P	-9.90	1.49	1.61
38	BE	13	A	O3'-P	-9.90	1.49	1.61
85	AA	702	G	N9-C4	-9.90	1.30	1.38
85	AA	1212	C	P-O5'	-9.90	1.49	1.59
34	BA	504	A	C2'-C1'	-9.89	1.42	1.53
35	BB	40	C	O3'-P	-9.89	1.49	1.61
35	BB	1364	C	P-O5'	-9.89	1.49	1.59
34	BA	658	C	P-O5'	-9.89	1.49	1.59
35	BB	778	A	C1'-N9	-9.89	1.32	1.46
85	AA	487	G	P-O5'	-9.89	1.49	1.59
36	BC	12	A	P-O5'	-9.89	1.49	1.59
40	BG	67	A	O3'-P	-9.89	1.49	1.61
40	BG	128	U	C1'-N1	-9.89	1.33	1.46
85	AA	995	G	C2'-C1'	-9.89	1.42	1.53
35	BB	465	C	P-O5'	-9.89	1.49	1.59
38	BE	49	A	O3'-P	-9.89	1.49	1.61
40	BG	75	C	P-O5'	-9.89	1.49	1.59
35	BB	782	A	O3'-P	-9.88	1.49	1.61
85	AA	480	U	O3'-P	-9.89	1.49	1.61
34	BA	348	U	O3'-P	-9.88	1.49	1.61
41	BH	103	C	C2'-C1'	-9.88	1.42	1.53
34	BA	1698	C	O3'-P	-9.88	1.49	1.61
35	BB	387	G	N7-C5	-9.88	1.33	1.39
85	AA	2113	U	P-O5'	-9.88	1.49	1.59
85	AA	554	A	P-O5'	-9.88	1.49	1.59
85	AA	1368	G	P-O5'	-9.88	1.49	1.59
34	BA	1061	A	N9-C4	-9.88	1.31	1.37
34	BA	1478	G	O3'-P	-9.88	1.49	1.61
34	BA	1597	G	C2'-C1'	-9.88	1.42	1.53
35	BB	1401	G	C8-N7	-9.88	1.25	1.30
35	BB	1430	G	C1'-N9	-9.88	1.33	1.46
34	BA	726	G	C6-N1	-9.88	1.32	1.39
34	BA	1607	U	C3'-C2'	-9.88	1.41	1.52
35	BB	1034	U	O3'-P	-9.87	1.49	1.61
85	AA	1243	G	C2'-C1'	-9.88	1.42	1.53
34	BA	749	G	O3'-P	-9.87	1.49	1.61
37	BD	75	G	N3-C4	-9.87	1.28	1.35
35	BB	1087	A	N9-C4	-9.87	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	800	A	N3-C4	-9.87	1.28	1.34
34	BA	180	G	N9-C4	9.87	1.45	1.38
35	BB	1418	C	P-O5'	-9.87	1.49	1.59
41	BH	31	A	N9-C4	-9.87	1.31	1.37
34	BA	1030	C	O3'-P	-9.87	1.49	1.61
85	AA	623	G	P-O5'	-9.87	1.49	1.59
85	AA	1105	G	C2-N2	-9.87	1.24	1.34
34	BA	675	C	P-O5'	-9.86	1.49	1.59
35	BB	1179	C	O3'-P	-9.87	1.49	1.61
40	BG	9	G	P-O5'	-9.86	1.49	1.59
34	BA	83	G	O3'-P	-9.86	1.49	1.61
34	BA	1189	A	N7-C5	-9.86	1.33	1.39
35	BB	117	A	N9-C4	-9.86	1.31	1.37
35	BB	779	C	O3'-P	-9.86	1.49	1.61
85	AA	303	A	C5'-C4'	9.86	1.63	1.51
85	AA	830	A	N9-C4	-9.86	1.31	1.37
34	BA	100	A	N9-C4	-9.86	1.31	1.37
34	BA	1544	G	P-O5'	-9.86	1.49	1.59
35	BB	1161	G	N9-C4	-9.86	1.30	1.38
34	BA	1557	G	N7-C5	-9.86	1.33	1.39
34	BA	1814	U	O3'-P	-9.86	1.49	1.61
40	BG	161	C	O3'-P	-9.86	1.49	1.61
34	BA	518	C	O3'-P	-9.85	1.49	1.61
34	BA	1672	C	P-O5'	-9.85	1.49	1.59
35	BB	557	C	P-O5'	-9.85	1.49	1.59
35	BB	1139	A	O3'-P	-9.85	1.49	1.61
38	BE	173	G	N9-C4	-9.85	1.30	1.38
85	AA	93	G	P-O5'	-9.85	1.49	1.59
85	AA	1275	A	C2'-C1'	-9.85	1.42	1.53
85	AA	270	A	C2'-C1'	-9.85	1.42	1.53
34	BA	487	A	N9-C4	-9.85	1.31	1.37
34	BA	1228	G	P-O5'	-9.85	1.50	1.59
35	BB	1157	G	P-O5'	-9.85	1.50	1.59
85	AA	1663	U	P-O5'	-9.85	1.50	1.59
34	BA	99	G	P-O5'	-9.85	1.50	1.59
34	BA	1003	A	N9-C4	-9.85	1.31	1.37
34	BA	76	U	O3'-P	-9.85	1.49	1.61
34	BA	335	C	O3'-P	-9.85	1.49	1.61
35	BB	1412	U	P-O5'	-9.85	1.50	1.59
85	AA	1867	G	P-O5'	-9.85	1.50	1.59
34	BA	1681	U	O3'-P	-9.84	1.49	1.61
35	BB	1036	G	C3'-C2'	-9.84	1.42	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	91	G	O3'-P	-9.84	1.49	1.61
41	BH	127	A	C2'-C1'	-9.84	1.42	1.53
85	AA	363	A	C2'-C1'	-9.84	1.42	1.53
85	AA	1464	G	P-O5'	-9.84	1.50	1.59
34	BA	173	U	P-O5'	-9.84	1.50	1.59
34	BA	467	A	P-O5'	-9.84	1.50	1.59
38	BE	20	C	C4-N4	-9.84	1.25	1.33
37	BD	72	U	P-O5'	-9.84	1.50	1.59
34	BA	84	U	O3'-P	-9.84	1.49	1.61
34	BA	1526	C	P-O5'	-9.84	1.50	1.59
38	BE	62	C	P-O5'	-9.84	1.50	1.59
38	BE	136	G	C4'-O4'	-9.84	1.32	1.45
85	AA	744	C	C2-N3	-9.84	1.27	1.35
85	AA	769	C	O3'-P	-9.84	1.49	1.61
34	BA	800	G	C1'-N9	-9.83	1.33	1.46
34	BA	826	C	P-O5'	-9.83	1.50	1.59
34	BA	1688	G	N9-C4	-9.83	1.30	1.38
85	AA	152	A	P-O5'	-9.83	1.50	1.59
85	AA	1963	G	N7-C5	-9.83	1.33	1.39
36	BC	9	G	P-O5'	-9.83	1.50	1.59
85	AA	2142	A	C2'-C1'	-9.83	1.42	1.53
34	BA	1582	C	P-O5'	-9.82	1.50	1.59
34	BA	992	A	C2'-C1'	-9.82	1.42	1.53
35	BB	112	G	P-O5'	-9.82	1.50	1.59
35	BB	1060	U	P-O5'	-9.82	1.50	1.59
85	AA	485	A	C2'-C1'	-9.82	1.42	1.53
85	AA	1093	C	P-O5'	-9.82	1.50	1.59
85	AA	1950	G	P-O5'	-9.82	1.50	1.59
34	BA	1252	G	C1'-N9	-9.82	1.33	1.46
34	BA	1283	U	P-O5'	-9.82	1.50	1.59
35	BB	1154	C	O3'-P	-9.82	1.49	1.61
40	BG	99	A	N9-C4	-9.82	1.31	1.37
34	BA	954	U	C2-N3	-9.81	1.30	1.37
38	BE	134	A	P-O5'	-9.81	1.50	1.59
40	BG	94	G	C2'-C1'	-9.81	1.42	1.53
40	BG	52	A	O3'-P	-9.81	1.49	1.61
85	AA	387	U	O3'-P	-9.81	1.49	1.61
85	AA	420	C	C2-N3	-9.81	1.27	1.35
85	AA	486	G	P-O5'	-9.81	1.50	1.59
34	BA	1475	G	C1'-N9	-9.81	1.33	1.46
34	BA	1590	G	C6-N1	-9.81	1.32	1.39
85	AA	506	G	N9-C4	-9.81	1.30	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1562	G	O3'-P	-9.80	1.49	1.61
34	BA	1158	A	N7-C5	-9.80	1.33	1.39
34	BA	1429	A	O3'-P	-9.80	1.49	1.61
35	BB	578	G	P-O5'	-9.80	1.50	1.59
41	BH	41	A	P-O5'	-9.80	1.50	1.59
85	AA	674	U	C3'-C2'	-9.80	1.42	1.52
85	AA	916	A	O3'-P	-9.80	1.49	1.61
85	AA	1139	G	C2'-C1'	-9.80	1.42	1.53
34	BA	1581	G	C6-N1	-9.80	1.32	1.39
34	BA	1455	C	O3'-P	-9.80	1.49	1.61
85	AA	584	G	N7-C5	-9.80	1.33	1.39
85	AA	1828	C	O3'-P	-9.80	1.49	1.61
34	BA	1135	U	P-O5'	-9.80	1.50	1.59
34	BA	440	A	P-O5'	-9.80	1.50	1.59
34	BA	1443	U	C2-N3	-9.80	1.30	1.37
35	BB	991	C	P-O5'	-9.80	1.50	1.59
40	BG	84	U	P-O5'	-9.80	1.50	1.59
85	AA	898	A	P-O5'	-9.80	1.50	1.59
85	AA	1197	U	N3-C4	-9.80	1.29	1.38
85	AA	2184	A	O3'-P	-9.80	1.49	1.61
34	BA	272	A	C1'-N9	-9.80	1.33	1.46
34	BA	327	G	O3'-P	-9.80	1.49	1.61
85	AA	90	A	P-O5'	-9.79	1.50	1.59
85	AA	411	U	P-O5'	-9.79	1.50	1.59
34	BA	440	A	N7-C5	-9.79	1.33	1.39
34	BA	670	U	C2'-C1'	-9.79	1.42	1.53
40	BG	41	U	C2-N3	-9.79	1.30	1.37
85	AA	1434	U	P-O5'	-9.79	1.50	1.59
34	BA	478	G	N7-C5	-9.79	1.33	1.39
35	BB	135	C	O3'-P	-9.79	1.49	1.61
85	AA	397	G	N9-C4	-9.79	1.30	1.38
34	BA	1646	U	O3'-P	-9.79	1.49	1.61
35	BB	73	G	N7-C5	-9.79	1.33	1.39
34	BA	1191	C	C2'-C1'	-9.79	1.42	1.53
35	BB	868	C	C2'-C1'	-9.78	1.42	1.53
40	BG	112	C	C3'-C2'	-9.79	1.42	1.52
85	AA	20	G	P-O5'	-9.78	1.50	1.59
85	AA	1565	G	P-O5'	-9.79	1.50	1.59
34	BA	449	G	C3'-C2'	-9.78	1.42	1.52
34	BA	749	G	C3'-C2'	-9.78	1.42	1.52
35	BB	1203	C	C2'-C1'	-9.78	1.42	1.53
85	AA	425	G	O3'-P	-9.78	1.49	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	792	A	N7-C5	-9.78	1.33	1.39
34	BA	30	A	C3'-C2'	-9.78	1.42	1.52
85	AA	1214	C	O3'-P	-9.78	1.49	1.61
34	BA	1826	C	P-O5'	-9.78	1.50	1.59
35	BB	1202	G	C2'-C1'	-9.78	1.42	1.53
36	BC	150	U	O3'-P	-9.78	1.49	1.61
36	BC	8	C	P-O5'	-9.78	1.50	1.59
34	BA	1164	C	P-O5'	-9.78	1.50	1.59
85	AA	385	A	C2'-C1'	-9.77	1.42	1.53
34	BA	1162	U	P-O5'	-9.77	1.50	1.59
85	AA	2139	G	N7-C5	-9.77	1.33	1.39
34	BA	939	C	C3'-C2'	-9.77	1.42	1.52
37	BD	64	A	P-O5'	-9.77	1.50	1.59
85	AA	2177	C	O3'-P	-9.77	1.49	1.61
85	AA	1541	G	C4'-C3'	-9.77	1.42	1.53
35	BB	95	A	N7-C5	-9.77	1.33	1.39
35	BB	1098	G	P-O5'	-9.77	1.50	1.59
37	BD	44	U	P-O5'	-9.77	1.50	1.59
38	BE	27	A	N9-C4	-9.77	1.31	1.37
34	BA	374	U	C2-N3	-9.76	1.30	1.37
34	BA	1559	C	C2-N3	-9.76	1.27	1.35
35	BB	1405	G	C1'-N9	-9.76	1.33	1.46
37	BD	67	C	O3'-P	-9.76	1.49	1.61
85	AA	644	A	O3'-P	-9.76	1.49	1.61
35	BB	1291	G	C1'-N9	-9.76	1.33	1.46
35	BB	1466	A	C2'-C1'	-9.76	1.42	1.53
37	BD	105	G	N7-C5	-9.76	1.33	1.39
38	BE	146	U	P-O5'	-9.76	1.50	1.59
34	BA	266	G	N1-C2	-9.76	1.29	1.37
34	BA	409	A	O3'-P	-9.76	1.49	1.61
35	BB	1070	G	N9-C8	-9.76	1.31	1.37
38	BE	204	U	O3'-P	-9.76	1.49	1.61
41	BH	109	G	N9-C4	-9.76	1.30	1.38
34	BA	407	A	P-O5'	-9.75	1.50	1.59
40	BG	126	G	O3'-P	-9.75	1.49	1.61
34	BA	232	U	C5'-C4'	9.75	1.63	1.51
85	AA	474	C	O3'-P	-9.75	1.49	1.61
34	BA	322	U	C2-N3	-9.75	1.30	1.37
34	BA	427	G	O3'-P	-9.75	1.49	1.61
34	BA	1016	A	O3'-P	-9.75	1.49	1.61
34	BA	1534	U	C2-N3	-9.75	1.30	1.37
34	BA	88	C	O3'-P	-9.75	1.49	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1107	C	O3'-P	-9.75	1.49	1.61
35	BB	1310	C	P-O5'	-9.75	1.50	1.59
40	BG	72	G	C6-N1	-9.75	1.32	1.39
40	BG	140	G	C3'-C2'	-9.75	1.42	1.52
35	BB	639	A	O3'-P	-9.74	1.49	1.61
34	BA	1332	U	C2'-C1'	-9.74	1.42	1.53
34	BA	1451	A	C2'-C1'	-9.74	1.42	1.53
35	BB	101	U	C2-N3	-9.74	1.30	1.37
40	BG	23	C	C5'-C4'	-9.74	1.39	1.51
85	AA	2036	A	C8-N7	-9.74	1.24	1.31
34	BA	754	G	P-O5'	-9.74	1.50	1.59
34	BA	1519	G	C6-N1	-9.74	1.32	1.39
34	BA	1676	A	N9-C4	-9.74	1.32	1.37
35	BB	1339	C	O3'-P	-9.74	1.49	1.61
35	BB	1487	G	P-O5'	-9.74	1.50	1.59
85	AA	681	G	O3'-P	-9.74	1.49	1.61
34	BA	331	G	O3'-P	-9.74	1.49	1.61
34	BA	1094	U	C2-N3	-9.74	1.30	1.37
35	BB	86	A	P-O5'	-9.74	1.50	1.59
35	BB	1100	C	O3'-P	-9.74	1.49	1.61
40	BG	9	G	C1'-N9	-9.74	1.33	1.46
34	BA	702	G	C3'-C2'	-9.73	1.42	1.52
38	BE	8	G	C6-N1	-9.73	1.32	1.39
41	BH	36	C	C3'-C2'	-9.73	1.42	1.52
85	AA	577	U	P-O5'	-9.73	1.50	1.59
85	AA	2075	C	O3'-P	-9.73	1.49	1.61
34	BA	69	C	O3'-P	-9.73	1.49	1.61
34	BA	431	A	O3'-P	-9.73	1.49	1.61
34	BA	880	G	O3'-P	-9.73	1.49	1.61
36	BC	69	U	C3'-C2'	-9.73	1.42	1.52
34	BA	185	A	C1'-N9	-9.73	1.33	1.46
34	BA	1843	G	C2'-C1'	-9.73	1.42	1.53
35	BB	1403	G	N9-C4	-9.73	1.30	1.38
38	BE	183	C	N1-C6	9.73	1.43	1.37
85	AA	626	G	P-O5'	-9.72	1.50	1.59
85	AA	2054	G	C2'-C1'	-9.72	1.42	1.53
34	BA	787	A	N9-C4	-9.72	1.32	1.37
34	BA	974	G	C1'-N9	-9.72	1.33	1.46
85	AA	1122	U	O3'-P	-9.72	1.49	1.61
34	BA	1554	C	P-O5'	-9.72	1.50	1.59
35	BB	1192	C	P-O5'	-9.72	1.50	1.59
34	BA	1196	C	C4'-C3'	-9.72	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1533	G	C6-N1	-9.72	1.32	1.39
34	BA	1668	C	N1-C6	-9.72	1.31	1.37
35	BB	64	U	C4'-C3'	-9.72	1.42	1.53
34	BA	783	U	P-O5'	-9.72	1.50	1.59
38	BE	66	A	N9-C4	-9.72	1.32	1.37
35	BB	27	C	C2'-C1'	-9.71	1.42	1.53
35	BB	680	A	N9-C4	-9.71	1.32	1.37
35	BB	1241	U	C2-N3	-9.71	1.30	1.37
85	AA	538	A	O3'-P	-9.71	1.49	1.61
85	AA	646	C	P-O5'	-9.71	1.50	1.59
34	BA	795	G	N9-C4	-9.71	1.30	1.38
85	AA	533	C	C2-N3	-9.71	1.27	1.35
85	AA	2085	C	O3'-P	-9.71	1.49	1.61
38	BE	204	U	C2-N3	-9.71	1.30	1.37
85	AA	156	G	C6-N1	-9.71	1.32	1.39
85	AA	994	A	C2'-C1'	-9.71	1.42	1.53
35	BB	58	G	P-O5'	-9.71	1.50	1.59
39	BF	36	G	O3'-P	-9.71	1.49	1.61
85	AA	1661	U	O3'-P	-9.71	1.49	1.61
34	BA	714	G	N9-C4	-9.71	1.30	1.38
36	BC	4	G	N9-C4	-9.71	1.30	1.38
34	BA	1064	A	P-O5'	-9.70	1.50	1.59
35	BB	434	A	O3'-P	-9.70	1.49	1.61
85	AA	185	A	N9-C4	-9.70	1.32	1.37
85	AA	509	C	O3'-P	-9.70	1.49	1.61
34	BA	135	G	C6-N1	-9.70	1.32	1.39
34	BA	168	U	P-O5'	-9.70	1.50	1.59
40	BG	73	U	C2-N3	-9.70	1.30	1.37
85	AA	2098	A	N9-C4	-9.70	1.32	1.37
34	BA	894	G	C2'-C1'	-9.70	1.42	1.53
34	BA	1811	A	N9-C4	-9.70	1.32	1.37
85	AA	317	A	N9-C4	-9.70	1.32	1.37
35	BB	72	G	C6-N1	-9.70	1.32	1.39
85	AA	928	U	C2'-C1'	-9.70	1.42	1.53
35	BB	827	U	O3'-P	-9.69	1.49	1.61
85	AA	398	U	O3'-P	-9.70	1.49	1.61
85	AA	2244	G	P-O5'	-9.69	1.50	1.59
34	BA	1192	A	P-O5'	-9.69	1.50	1.59
35	BB	1283	C	P-O5'	-9.69	1.50	1.59
85	AA	788	G	O3'-P	-9.69	1.49	1.61
34	BA	701	G	C3'-C2'	-9.69	1.42	1.52
34	BA	758	G	N7-C5	-9.69	1.33	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	946	A	C1'-N9	-9.69	1.33	1.46
34	BA	1608	C	O3'-P	-9.69	1.49	1.61
35	BB	412	A	N9-C4	-9.68	1.32	1.37
85	AA	102	A	O3'-P	-9.68	1.49	1.61
35	BB	1359	G	C2'-C1'	-9.68	1.42	1.53
40	BG	28	A	C2'-C1'	-9.68	1.42	1.53
85	AA	111	A	O3'-P	-9.68	1.49	1.61
85	AA	423	G	P-O5'	-9.68	1.50	1.59
85	AA	1124	G	P-O5'	-9.68	1.50	1.59
85	AA	1678	U	C2'-C1'	-9.68	1.42	1.53
34	BA	131	A	O3'-P	-9.68	1.49	1.61
34	BA	859	G	P-O5'	-9.68	1.50	1.59
34	BA	1701	U	P-O5'	-9.68	1.50	1.59
35	BB	1330	A	N9-C4	-9.68	1.32	1.37
35	BB	439	G	O3'-P	-9.68	1.49	1.61
85	AA	1248	U	C2'-C1'	-9.68	1.42	1.53
36	BC	55	U	C3'-C2'	-9.68	1.42	1.52
38	BE	197	A	C1'-N9	-9.68	1.33	1.46
34	BA	580	U	P-O5'	-9.67	1.50	1.59
34	BA	1289	C	O3'-P	-9.67	1.49	1.61
35	BB	691	A	N9-C4	-9.67	1.32	1.37
35	BB	968	C	C2'-C1'	-9.67	1.42	1.53
34	BA	599	U	O3'-P	-9.67	1.49	1.61
85	AA	2154	C	C2'-C1'	-9.67	1.42	1.53
35	BB	382	U	P-O5'	-9.67	1.50	1.59
35	BB	450	A	N9-C4	-9.67	1.32	1.37
85	AA	658	C	P-O5'	-9.67	1.50	1.59
35	BB	1483	A	N9-C4	-9.67	1.32	1.37
40	BG	52	A	C3'-C2'	-9.67	1.42	1.52
85	AA	1142	G	P-O5'	-9.67	1.50	1.59
85	AA	1471	G	C1'-N9	-9.67	1.33	1.46
34	BA	1287	G	C2'-C1'	-9.67	1.42	1.53
35	BB	1467	A	N9-C4	-9.67	1.32	1.37
36	BC	153	C	O3'-P	-9.66	1.49	1.61
85	AA	429	G	P-O5'	-9.66	1.50	1.59
85	AA	1263	G	N7-C5	-9.66	1.33	1.39
34	BA	1208	U	O3'-P	-9.66	1.49	1.61
34	BA	1236	U	C2-N3	-9.66	1.30	1.37
34	BA	1421	A	N9-C4	-9.66	1.32	1.37
34	BA	1510	C	C3'-C2'	-9.66	1.42	1.52
85	AA	1126	G	C6-N1	-9.66	1.32	1.39
35	BB	1292	G	C6-N1	-9.66	1.32	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1645	G	P-O5'	-9.66	1.50	1.59
34	BA	154	A	P-O5'	-9.66	1.50	1.59
35	BB	1018	U	P-O5'	-9.66	1.50	1.59
34	BA	255	G	O3'-P	-9.66	1.49	1.61
34	BA	1281	U	C3'-C2'	-9.66	1.42	1.52
35	BB	867	C	C2'-C1'	-9.66	1.42	1.53
35	BB	1256	C	O3'-P	-9.66	1.49	1.61
35	BB	1480	G	C3'-C2'	-9.66	1.42	1.52
34	BA	1583	A	C2'-C1'	-9.66	1.42	1.53
35	BB	134	G	C6-N1	-9.66	1.32	1.39
35	BB	823	G	C1'-N9	-9.66	1.33	1.46
35	BB	1133	C	O3'-P	-9.66	1.49	1.61
85	AA	466	A	P-O5'	-9.66	1.50	1.59
85	AA	2218	G	C8-N7	-9.66	1.25	1.30
34	BA	388	A	N9-C4	-9.65	1.32	1.37
35	BB	484	G	O3'-P	-9.65	1.49	1.61
85	AA	250	C	N1-C6	-9.65	1.31	1.37
85	AA	547	A	P-O5'	-9.65	1.50	1.59
34	BA	932	G	C6-N1	-9.65	1.32	1.39
39	BF	41	U	P-O5'	-9.65	1.50	1.59
85	AA	707	U	O3'-P	-9.65	1.49	1.61
34	BA	1599	A	C3'-C2'	-9.65	1.42	1.52
38	BE	127	G	N1-C2	-9.65	1.30	1.37
85	AA	2177	C	C2'-C1'	-9.65	1.42	1.53
38	BE	174	U	C3'-C2'	-9.65	1.42	1.52
35	BB	778	A	N9-C4	-9.64	1.32	1.37
35	BB	1343	C	O3'-P	-9.64	1.49	1.61
85	AA	605	A	O3'-P	-9.64	1.49	1.61
85	AA	780	U	P-O5'	-9.64	1.50	1.59
34	BA	6	C	C3'-C2'	-9.64	1.42	1.52
34	BA	349	G	C2'-C1'	-9.64	1.42	1.53
34	BA	783	U	O3'-P	-9.64	1.49	1.61
35	BB	1152	U	O3'-P	-9.64	1.49	1.61
35	BB	1215	U	P-O5'	-9.64	1.50	1.59
35	BB	1308	G	P-O5'	-9.64	1.50	1.59
85	AA	1542	A	P-O5'	-9.64	1.50	1.59
35	BB	1368	A	P-O5'	-9.64	1.50	1.59
36	BC	52	A	N9-C4	-9.64	1.32	1.37
34	BA	893	U	C2'-C1'	-9.63	1.42	1.53
85	AA	477	U	C3'-C2'	-9.64	1.42	1.52
35	BB	68	G	P-O5'	-9.63	1.50	1.59
35	BB	1299	G	N9-C4	-9.63	1.30	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	103	C	O3'-P	-9.63	1.49	1.61
85	AA	160	A	C1'-N9	-9.63	1.33	1.46
34	BA	1419	A	P-O5'	-9.63	1.50	1.59
35	BB	1231	U	O3'-P	-9.63	1.49	1.61
41	BH	53	C	P-O5'	-9.63	1.50	1.59
34	BA	349	G	O3'-P	-9.63	1.49	1.61
34	BA	680	C	C3'-C2'	-9.63	1.42	1.52
34	BA	1578	A	C2'-C1'	-9.63	1.42	1.53
34	BA	1812	C	O3'-P	-9.63	1.49	1.61
85	AA	1133	C	O3'-P	-9.63	1.49	1.61
34	BA	1287	G	P-O5'	-9.63	1.50	1.59
35	BB	1210	U	C2-N3	-9.63	1.31	1.37
85	AA	443	A	N9-C4	-9.63	1.32	1.37
34	BA	1333	G	C1'-N9	-9.62	1.33	1.46
35	BB	415	A	P-O5'	-9.62	1.50	1.59
34	BA	565	U	P-O5'	-9.62	1.50	1.59
34	BA	658	C	C2'-C1'	-9.62	1.42	1.53
38	BE	88	G	P-O5'	-9.62	1.50	1.59
85	AA	52	U	O3'-P	-9.62	1.49	1.61
85	AA	241	U	O3'-P	-9.62	1.49	1.61
34	BA	926	A	C5-C4	-9.62	1.32	1.38
35	BB	654	C	C1'-N1	-9.62	1.33	1.46
85	AA	1978	G	C1'-N9	-9.62	1.33	1.46
85	AA	2183	U	P-O5'	-9.62	1.50	1.59
34	BA	61	G	O3'-P	-9.62	1.49	1.61
34	BA	817	U	C3'-C2'	-9.62	1.42	1.52
35	BB	1136	G	P-O5'	-9.62	1.50	1.59
34	BA	1037	C	O3'-P	-9.61	1.49	1.61
35	BB	118	A	C1'-N9	-9.62	1.33	1.46
37	BD	80	G	P-O5'	-9.62	1.50	1.59
40	BG	166	C	O3'-P	-9.62	1.49	1.61
85	AA	726	U	P-O5'	-9.62	1.50	1.59
85	AA	2139	G	C2'-C1'	-9.62	1.42	1.53
34	BA	19	G	C1'-N9	-9.61	1.33	1.46
34	BA	1645	C	O3'-P	-9.61	1.49	1.61
35	BB	86	A	N9-C4	-9.61	1.32	1.37
36	BC	95	A	O3'-P	-9.61	1.49	1.61
37	BD	82	G	C6-N1	-9.61	1.32	1.39
34	BA	994	G	O3'-P	-9.61	1.49	1.61
35	BB	482	A	O3'-P	-9.61	1.49	1.61
41	BH	40	C	C2'-C1'	-9.61	1.42	1.53
36	BC	43	A	C2'-C1'	-9.61	1.42	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	4	U	O3'-P	-9.61	1.49	1.61
39	BF	10	A	N7-C5	-9.61	1.33	1.39
34	BA	99	G	N3-C4	-9.61	1.28	1.35
34	BA	902	C	P-O5'	-9.61	1.50	1.59
34	BA	1073	G	N7-C5	-9.61	1.33	1.39
34	BA	1588	U	O3'-P	-9.61	1.49	1.61
35	BB	18	A	P-O5'	-9.61	1.50	1.59
35	BB	1248	A	C1'-N9	-9.61	1.33	1.46
85	AA	357	C	P-O5'	-9.61	1.50	1.59
85	AA	763	U	O3'-P	-9.61	1.49	1.61
85	AA	1506	U	P-O5'	-9.61	1.50	1.59
85	AA	1557	U	P-O5'	-9.61	1.50	1.59
34	BA	478	G	N9-C8	-9.60	1.31	1.37
36	BC	94	C	P-O5'	-9.60	1.50	1.59
85	AA	2169	C	P-O5'	-9.60	1.50	1.59
34	BA	1597	G	C5-C4	-9.60	1.31	1.38
35	BB	1519	U	C2'-C1'	-9.60	1.42	1.53
35	BB	605	C	P-O5'	-9.60	1.50	1.59
35	BB	1301	U	P-O5'	-9.60	1.50	1.59
38	BE	194	A	C8-N7	-9.60	1.24	1.31
85	AA	1498	C	P-O5'	-9.60	1.50	1.59
39	BF	71	G	O3'-P	-9.60	1.49	1.61
34	BA	1502	G	N9-C8	-9.60	1.31	1.37
35	BB	1155	U	C2'-C1'	-9.60	1.42	1.53
35	BB	1367	U	C2-N3	-9.60	1.31	1.37
36	BC	116	C	C2-N3	-9.60	1.28	1.35
36	BC	151	G	P-O5'	-9.60	1.50	1.59
41	BH	114	G	N9-C4	-9.60	1.30	1.38
85	AA	367	A	N7-C5	-9.60	1.33	1.39
35	BB	1178	A	O3'-P	-9.59	1.49	1.61
85	AA	402	G	N7-C5	-9.59	1.33	1.39
35	BB	1140	C	P-O5'	-9.59	1.50	1.59
35	BB	693	U	P-O5'	-9.59	1.50	1.59
85	AA	397	G	P-O5'	-9.59	1.50	1.59
85	AA	1217	U	N3-C4	-9.59	1.29	1.38
35	BB	1249	G	C1'-N9	-9.59	1.33	1.46
38	BE	12	A	N9-C4	-9.59	1.32	1.37
34	BA	103	G	O3'-P	-9.59	1.49	1.61
34	BA	598	G	O3'-P	-9.59	1.49	1.61
34	BA	1250	C	C2-N3	-9.59	1.28	1.35
34	BA	1558	C	O3'-P	-9.59	1.49	1.61
35	BB	786	A	C1'-N9	-9.59	1.33	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1510	G	P-O5'	-9.59	1.50	1.59
36	BC	99	U	P-O5'	-9.59	1.50	1.59
85	AA	2033	C	C4'-C3'	-9.59	1.42	1.53
34	BA	1418	G	N9-C8	-9.58	1.31	1.37
85	AA	1823	G	O3'-P	-9.58	1.49	1.61
35	BB	1420	U	O3'-P	-9.58	1.49	1.61
85	AA	302	C	C5'-C4'	9.58	1.62	1.51
85	AA	437	G	O3'-P	-9.58	1.49	1.61
34	BA	183	G	P-O5'	-9.58	1.50	1.59
34	BA	539	C	P-O5'	-9.58	1.50	1.59
34	BA	616	G	C2'-C1'	-9.58	1.42	1.53
85	AA	248	U	O3'-P	-9.58	1.49	1.61
85	AA	627	A	N9-C4	-9.58	1.32	1.37
35	BB	663	G	P-O5'	-9.58	1.50	1.59
41	BH	132	C	P-O5'	-9.58	1.50	1.59
34	BA	967	C	O3'-P	-9.57	1.49	1.61
34	BA	1327	G	O3'-P	-9.57	1.49	1.61
35	BB	611	U	O3'-P	-9.57	1.49	1.61
35	BB	1027	U	O3'-P	-9.57	1.49	1.61
37	BD	108	G	N7-C5	-9.57	1.33	1.39
34	BA	798	G	P-O5'	-9.57	1.50	1.59
35	BB	675	U	O3'-P	-9.57	1.49	1.61
37	BD	32	A	O3'-P	-9.57	1.49	1.61
34	BA	320	G	N7-C5	-9.57	1.33	1.39
35	BB	662	G	C2'-C1'	-9.57	1.42	1.53
35	BB	852	G	P-O5'	-9.57	1.50	1.59
35	BB	481	A	C1'-N9	-9.57	1.33	1.46
35	BB	1340	U	C3'-C2'	-9.57	1.42	1.52
85	AA	2010	C	P-O5'	-9.57	1.50	1.59
41	BH	15	A	P-O5'	-9.56	1.50	1.59
85	AA	1885	A	C2'-C1'	-9.56	1.42	1.53
85	AA	1254	A	C2'-C1'	-9.56	1.42	1.53
34	BA	908	G	P-O5'	-9.56	1.50	1.59
34	BA	1719	G	C6-N1	-9.56	1.32	1.39
36	BC	91	G	P-O5'	-9.56	1.50	1.59
85	AA	1471	G	C6-N1	-9.56	1.32	1.39
38	BE	178	G	P-O5'	-9.56	1.50	1.59
85	AA	455	G	C1'-N9	-9.56	1.33	1.46
85	AA	2058	C	C3'-C2'	-9.56	1.42	1.52
34	BA	1091	U	C2-N3	-9.56	1.31	1.37
38	BE	117	A	C4'-O4'	-9.56	1.33	1.45
39	BF	65	U	O3'-P	-9.56	1.49	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	86	U	C2'-C1'	-9.56	1.42	1.53
85	AA	1692	U	C2-N3	-9.56	1.31	1.37
85	AA	2211	G	P-O5'	-9.56	1.50	1.59
34	BA	557	U	C4'-C3'	9.56	1.63	1.53
34	BA	297	A	N3-C4	-9.55	1.29	1.34
35	BB	442	U	O3'-P	-9.55	1.49	1.61
38	BE	195	G	N9-C4	9.55	1.45	1.38
85	AA	769	C	P-O5'	-9.55	1.50	1.59
85	AA	2119	C	O3'-P	-9.55	1.49	1.61
34	BA	1802	C	C2'-C1'	-9.55	1.42	1.53
35	BB	654	C	P-O5'	-9.55	1.50	1.59
41	BH	110	C	O3'-P	-9.55	1.49	1.61
85	AA	1487	G	C1'-N9	-9.55	1.33	1.46
35	BB	1284	U	C2-N3	-9.55	1.31	1.37
34	BA	150	C	C4'-C3'	-9.55	1.42	1.53
34	BA	1052	G	N9-C4	-9.54	1.30	1.38
34	BA	1598	U	O3'-P	-9.54	1.49	1.61
34	BA	1599	A	N9-C4	-9.54	1.32	1.37
35	BB	374	A	C1'-N9	-9.55	1.33	1.46
35	BB	485	U	P-O5'	-9.54	1.50	1.59
85	AA	1238	U	O3'-P	-9.54	1.49	1.61
34	BA	705	C	C3'-C2'	-9.54	1.42	1.52
34	BA	1193	A	P-O5'	-9.54	1.50	1.59
36	BC	24	G	N9-C4	-9.54	1.30	1.38
85	AA	2129	U	P-O5'	-9.54	1.50	1.59
34	BA	365	A	P-O5'	-9.54	1.50	1.59
35	BB	704	G	P-O5'	-9.54	1.50	1.59
38	BE	139	U	O3'-P	-9.54	1.49	1.61
39	BF	25	G	N7-C5	-9.54	1.33	1.39
41	BH	119	U	O4'-C1'	-9.54	1.29	1.41
85	AA	1566	A	P-O5'	-9.54	1.50	1.59
34	BA	133	A	N9-C4	-9.54	1.32	1.37
34	BA	1687	A	O3'-P	-9.54	1.49	1.61
35	BB	608	A	O3'-P	-9.54	1.49	1.61
85	AA	2011	C	C2'-C1'	-9.54	1.42	1.53
85	AA	2223	C	P-O5'	-9.53	1.50	1.59
34	BA	1260	G	C5-C4	-9.53	1.31	1.38
35	BB	375	G	O3'-P	-9.53	1.49	1.61
35	BB	1421	C	C2-N3	-9.53	1.28	1.35
41	BH	38	G	C2'-C1'	-9.53	1.42	1.53
34	BA	74	A	C1'-N9	-9.53	1.33	1.46
34	BA	733	G	N9-C4	-9.53	1.30	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	556	U	C3'-C2'	-9.53	1.42	1.52
34	BA	1559	C	C3'-C2'	-9.53	1.42	1.52
35	BB	703	U	O3'-P	-9.53	1.49	1.61
34	BA	893	U	O3'-P	-9.53	1.49	1.61
34	BA	1307	U	P-O5'	-9.53	1.50	1.59
35	BB	1386	C	P-O5'	-9.53	1.50	1.59
85	AA	736	U	C2'-C1'	-9.53	1.42	1.53
36	BC	29	C	N1-C2	-9.53	1.30	1.40
34	BA	135	G	C3'-C2'	-9.52	1.42	1.52
35	BB	667	G	O3'-P	-9.52	1.49	1.61
85	AA	307	G	C2'-C1'	-9.52	1.42	1.53
85	AA	1282	A	N9-C4	-9.52	1.32	1.37
85	AA	1650	G	O3'-P	-9.52	1.49	1.61
85	AA	1271	U	P-O5'	-9.52	1.50	1.59
34	BA	975	A	O3'-P	-9.52	1.49	1.61
34	BA	1675	C	C3'-C2'	-9.52	1.42	1.52
37	BD	83	A	P-O5'	-9.52	1.50	1.59
34	BA	1108	U	P-O5'	-9.52	1.50	1.59
35	BB	35	G	C5-C4	-9.52	1.31	1.38
34	BA	809	U	P-O5'	-9.52	1.50	1.59
35	BB	688	U	C2-N3	-9.52	1.31	1.37
85	AA	449	G	N9-C4	-9.52	1.30	1.38
34	BA	45	A	N7-C5	-9.51	1.33	1.39
34	BA	790	G	N9-C4	-9.51	1.30	1.38
85	AA	119	G	N9-C4	-9.51	1.30	1.38
35	BB	131	A	P-O5'	-9.51	1.50	1.59
34	BA	372	U	C2-N3	-9.51	1.31	1.37
34	BA	1486	U	C2'-C1'	-9.51	1.42	1.53
34	BA	1635	A	P-O5'	-9.51	1.50	1.59
34	BA	423	G	O3'-P	-9.51	1.49	1.61
34	BA	1328	U	C3'-C2'	-9.51	1.42	1.52
34	BA	1560	U	C2-N3	-9.51	1.31	1.37
35	BB	651	G	N7-C5	-9.51	1.33	1.39
34	BA	1816	G	C5-C4	-9.51	1.31	1.38
35	BB	661	G	O3'-P	-9.51	1.49	1.61
38	BE	46	G	O3'-P	-9.51	1.49	1.61
85	AA	194	U	P-O5'	-9.51	1.50	1.59
85	AA	369	A	N9-C4	-9.51	1.32	1.37
34	BA	125	G	O3'-P	-9.50	1.49	1.61
34	BA	213	A	N9-C4	-9.50	1.32	1.37
34	BA	932	G	N7-C5	-9.50	1.33	1.39
34	BA	1290	A	N9-C4	-9.50	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2039	G	C4'-C3'	-9.50	1.42	1.53
34	BA	713	C	C2'-C1'	-9.50	1.42	1.53
34	BA	1522	G	C1'-N9	-9.50	1.33	1.46
35	BB	1517	G	N7-C5	-9.50	1.33	1.39
38	BE	108	U	C3'-C2'	-9.50	1.42	1.52
40	BG	136	G	C2'-C1'	-9.50	1.43	1.53
85	AA	1137	C	P-O5'	-9.50	1.50	1.59
40	BG	168	A	N9-C4	-9.50	1.32	1.37
34	BA	300	C	C2'-C1'	-9.50	1.43	1.53
85	AA	745	C	P-O5'	-9.50	1.50	1.59
34	BA	800	G	C2'-C1'	-9.50	1.43	1.53
34	BA	1329	U	O3'-P	-9.50	1.49	1.61
34	BA	670	U	C2-N3	-9.49	1.31	1.37
35	BB	1114	A	P-O5'	-9.49	1.50	1.59
34	BA	390	A	O3'-P	-9.49	1.49	1.61
34	BA	1279	U	C2-N3	-9.49	1.31	1.37
36	BC	125	A	C2'-C1'	-9.49	1.43	1.53
37	BD	31	U	P-O5'	-9.49	1.50	1.59
38	BE	13	A	C2'-C1'	-9.49	1.43	1.53
38	BE	47	U	O3'-P	-9.49	1.49	1.61
41	BH	39	G	C2'-C1'	-9.49	1.43	1.53
85	AA	85	U	P-O5'	-9.49	1.50	1.59
85	AA	604	C	O3'-P	-9.49	1.49	1.61
34	BA	323	C	C3'-C2'	-9.49	1.42	1.52
34	BA	452	A	P-O5'	-9.49	1.50	1.59
34	BA	1157	A	N9-C4	-9.49	1.32	1.37
34	BA	1555	G	C2'-C1'	-9.49	1.43	1.53
35	BB	992	C	O3'-P	-9.49	1.49	1.61
85	AA	1262	A	C2'-C1'	-9.49	1.43	1.53
85	AA	1502	A	C2'-C1'	-9.49	1.43	1.53
34	BA	851	C	O3'-P	-9.48	1.49	1.61
34	BA	1018	U	N3-C4	-9.48	1.29	1.38
34	BA	1531	G	C2'-C1'	-9.48	1.43	1.53
35	BB	881	G	P-O5'	-9.48	1.50	1.59
36	BC	61	A	N9-C4	-9.48	1.32	1.37
37	BD	32	A	P-O5'	-9.48	1.50	1.59
40	BG	40	G	C6-N1	-9.48	1.32	1.39
85	AA	259	A	P-O5'	-9.48	1.50	1.59
38	BE	105	A	P-O5'	-9.48	1.50	1.59
34	BA	649	A	P-O5'	-9.48	1.50	1.59
34	BA	906	A	N3-C4	-9.48	1.29	1.34
35	BB	1389	C	P-O5'	-9.48	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	59	G	N9-C4	-9.48	1.30	1.38
41	BH	38	G	N9-C4	-9.48	1.30	1.38
85	AA	994	A	P-O5'	-9.48	1.50	1.59
34	BA	557	U	C4'-O4'	9.48	1.57	1.45
85	AA	2103	C	O3'-P	-9.48	1.49	1.61
37	BD	60	C	P-O5'	-9.48	1.50	1.59
85	AA	465	A	P-O5'	-9.48	1.50	1.59
85	AA	806	G	C2'-C1'	-9.48	1.43	1.53
34	BA	1510	C	O3'-P	-9.47	1.49	1.61
34	BA	1840	C	O3'-P	-9.47	1.49	1.61
35	BB	49	A	O3'-P	-9.47	1.49	1.61
35	BB	456	A	O3'-P	-9.47	1.49	1.61
35	BB	588	A	N9-C4	-9.47	1.32	1.37
36	BC	109	A	N9-C4	-9.47	1.32	1.37
40	BG	1	G	N9-C4	-9.47	1.30	1.38
34	BA	1152	A	O3'-P	-9.47	1.49	1.61
34	BA	138	C	C3'-C2'	-9.47	1.42	1.52
35	BB	1062	G	C2'-C1'	-9.47	1.43	1.53
36	BC	73	U	C3'-C2'	-9.47	1.42	1.52
36	BC	75	G	C6-N1	-9.47	1.32	1.39
85	AA	1460	G	N9-C4	-9.47	1.30	1.38
34	BA	654	C	P-O5'	-9.47	1.50	1.59
34	BA	1573	C	O3'-P	-9.47	1.49	1.61
35	BB	1385	C	P-O5'	-9.47	1.50	1.59
36	BC	3	C	N1-C6	-9.47	1.31	1.37
39	BF	65	U	C2-N3	-9.47	1.31	1.37
34	BA	892	C	N3-C4	-9.47	1.27	1.33
35	BB	1427	A	P-O5'	-9.47	1.50	1.59
35	BB	1482	A	O3'-P	-9.47	1.49	1.61
38	BE	13	A	C5'-C4'	9.47	1.62	1.51
40	BG	163	G	N9-C4	-9.47	1.30	1.38
85	AA	2124	G	O3'-P	-9.47	1.49	1.61
34	BA	21	C	O3'-P	-9.46	1.49	1.61
34	BA	1210	A	C4'-C3'	-9.46	1.42	1.53
35	BB	487	A	N9-C4	-9.46	1.32	1.37
35	BB	491	A	C2'-C1'	-9.46	1.43	1.53
38	BE	114	G	C4'-C3'	-9.46	1.42	1.53
85	AA	605	A	C8-N7	-9.46	1.25	1.31
34	BA	1213	A	C1'-N9	-9.46	1.33	1.46
34	BA	1672	C	O3'-P	-9.46	1.49	1.61
35	BB	1407	U	O3'-P	-9.46	1.49	1.61
36	BC	4	G	C2'-C1'	-9.46	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	131	A	N9-C4	-9.46	1.32	1.37
34	BA	1171	C	P-O5'	-9.46	1.50	1.59
34	BA	1474	G	O3'-P	-9.46	1.49	1.61
34	BA	790	G	C2-N2	-9.46	1.25	1.34
34	BA	1054	U	P-O5'	-9.46	1.50	1.59
39	BF	49	C	O3'-P	-9.45	1.49	1.61
40	BG	179	C	O3'-P	-9.45	1.49	1.61
85	AA	718	C	P-O5'	-9.46	1.50	1.59
85	AA	2182	A	P-O5'	-9.45	1.50	1.59
34	BA	216	C	C2'-C1'	-9.45	1.43	1.53
34	BA	1077	G	P-O5'	-9.45	1.50	1.59
34	BA	1283	U	O3'-P	-9.45	1.49	1.61
34	BA	1556	A	C2'-C1'	-9.45	1.43	1.53
34	BA	1668	C	O3'-P	-9.45	1.49	1.61
34	BA	1678	U	P-O5'	-9.45	1.50	1.59
36	BC	31	A	P-O5'	-9.45	1.50	1.59
85	AA	1677	A	N9-C4	-9.45	1.32	1.37
34	BA	94	G	C6-N1	-9.45	1.32	1.39
36	BC	16	A	C6-N6	-9.45	1.26	1.33
38	BE	95	G	P-O5'	-9.45	1.50	1.59
40	BG	171	A	N7-C5	-9.45	1.33	1.39
85	AA	534	A	N9-C4	-9.45	1.32	1.37
85	AA	637	U	O3'-P	-9.45	1.49	1.61
34	BA	881	C	O3'-P	-9.45	1.49	1.61
85	AA	16	G	C5-C4	-9.45	1.31	1.38
35	BB	1164	U	O3'-P	-9.45	1.49	1.61
85	AA	495	G	O3'-P	-9.45	1.49	1.61
34	BA	682	A	N9-C4	-9.44	1.32	1.37
40	BG	120	U	P-O5'	-9.44	1.50	1.59
85	AA	710	A	N7-C5	-9.44	1.33	1.39
85	AA	926	C	O3'-P	-9.44	1.49	1.61
85	AA	710	A	O3'-P	-9.44	1.49	1.61
85	AA	1124	G	O3'-P	-9.44	1.49	1.61
34	BA	892	C	O3'-P	-9.44	1.49	1.61
37	BD	27	A	P-O5'	-9.44	1.50	1.59
38	BE	23	G	C6-N1	-9.44	1.32	1.39
85	AA	695	A	O3'-P	-9.44	1.49	1.61
85	AA	288	G	P-O5'	-9.44	1.50	1.59
35	BB	82	G	C2'-C1'	-9.43	1.43	1.53
35	BB	647	U	O3'-P	-9.43	1.49	1.61
35	BB	1377	A	O3'-P	-9.43	1.49	1.61
36	BC	102	G	N9-C4	-9.43	1.30	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	143	C	O3'-P	-9.43	1.49	1.61
85	AA	547	A	N9-C4	-9.43	1.32	1.37
34	BA	177	G	C2-N2	-9.43	1.25	1.34
34	BA	1283	U	C3'-C2'	-9.43	1.42	1.52
35	BB	485	U	C2-N3	-9.43	1.31	1.37
35	BB	1114	A	N7-C5	-9.43	1.33	1.39
35	BB	1285	U	O3'-P	-9.43	1.49	1.61
36	BC	74	U	P-O5'	-9.43	1.50	1.59
85	AA	687	G	N7-C5	-9.43	1.33	1.39
34	BA	871	G	N7-C5	-9.43	1.33	1.39
85	AA	173	A	O3'-P	-9.43	1.49	1.61
34	BA	31	A	O3'-P	-9.43	1.49	1.61
34	BA	399	G	N3-C4	-9.43	1.28	1.35
34	BA	940	C	C3'-C2'	-9.43	1.42	1.52
35	BB	389	G	O3'-P	-9.43	1.49	1.61
39	BF	11	C	O3'-P	-9.43	1.49	1.61
85	AA	174	U	O3'-P	-9.43	1.49	1.61
34	BA	1204	U	P-O5'	-9.42	1.50	1.59
37	BD	89	G	C2'-C1'	-9.42	1.43	1.53
40	BG	133	C	P-O5'	-9.42	1.50	1.59
34	BA	203	U	P-O5'	-9.42	1.50	1.59
34	BA	907	A	N9-C4	-9.42	1.32	1.37
34	BA	1511	C	C3'-C2'	-9.42	1.42	1.52
35	BB	1300	U	P-O5'	-9.42	1.50	1.59
85	AA	331	G	P-O5'	-9.42	1.50	1.59
34	BA	272	A	O3'-P	-9.42	1.49	1.61
34	BA	520	G	N7-C5	-9.42	1.33	1.39
35	BB	694	C	C2-N3	-9.42	1.28	1.35
35	BB	1040	C	C2'-C1'	-9.42	1.43	1.53
85	AA	1486	G	N9-C8	-9.42	1.31	1.37
34	BA	1704	G	P-O5'	-9.42	1.50	1.59
34	BA	1713	U	O3'-P	-9.42	1.49	1.61
34	BA	1732	A	N7-C5	-9.42	1.33	1.39
34	BA	983	A	C2'-C1'	-9.42	1.43	1.53
34	BA	1037	C	C3'-C2'	-9.42	1.42	1.52
34	BA	1335	A	C3'-C2'	-9.42	1.42	1.52
34	BA	1719	G	N9-C8	-9.42	1.31	1.37
35	BB	501	G	C6-N1	-9.42	1.32	1.39
37	BD	114	U	P-O5'	-9.42	1.50	1.59
86	AB	63	G	C2'-C1'	-9.42	1.43	1.53
35	BB	901	U	P-O5'	-9.41	1.50	1.59
35	BB	1237	C	C2'-C1'	-9.41	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1296	A	N7-C5	-9.41	1.33	1.39
38	BE	141	A	N7-C5	-9.41	1.33	1.39
40	BG	74	G	C6-N1	-9.41	1.32	1.39
85	AA	572	G	C2-N2	-9.41	1.25	1.34
85	AA	1474	U	O3'-P	-9.41	1.49	1.61
85	AA	1549	G	O3'-P	-9.41	1.49	1.61
35	BB	1401	G	N7-C5	-9.41	1.33	1.39
85	AA	425	G	C2'-C1'	-9.41	1.43	1.53
85	AA	1493	A	N7-C5	-9.41	1.33	1.39
85	AA	690	G	N1-C2	-9.41	1.30	1.37
85	AA	2054	G	O3'-P	-9.41	1.49	1.61
34	BA	961	C	O3'-P	-9.41	1.49	1.61
34	BA	1615	A	O3'-P	-9.41	1.49	1.61
40	BG	177	U	C2-N3	-9.41	1.31	1.37
35	BB	1115	G	P-O5'	-9.41	1.50	1.59
36	BC	90	U	N1-C2	-9.41	1.30	1.38
85	AA	505	U	C4'-C3'	-9.41	1.42	1.53
85	AA	1182	A	O3'-P	-9.41	1.49	1.61
35	BB	72	G	C3'-C2'	-9.40	1.42	1.52
35	BB	705	C	P-O5'	-9.40	1.50	1.59
41	BH	72	G	P-O5'	9.40	1.69	1.59
85	AA	627	A	C2'-C1'	-9.40	1.43	1.53
85	AA	1879	U	P-O5'	-9.40	1.50	1.59
85	AA	2094	U	O3'-P	-9.40	1.49	1.61
85	AA	1923	A	N9-C4	-9.40	1.32	1.37
34	BA	1841	A	P-O5'	-9.40	1.50	1.59
35	BB	474	G	N9-C4	-9.40	1.30	1.38
35	BB	1441	C	C2'-C1'	-9.40	1.43	1.53
85	AA	1481	U	C2-N3	-9.40	1.31	1.37
85	AA	995	G	N9-C4	-9.40	1.30	1.38
37	BD	41	G	P-O5'	-9.40	1.50	1.59
34	BA	58	A	P-O5'	-9.40	1.50	1.59
34	BA	1609	U	O3'-P	-9.40	1.49	1.61
40	BG	175	G	C2'-C1'	-9.40	1.43	1.53
34	BA	391	U	O3'-P	-9.40	1.49	1.61
35	BB	488	G	C2'-C1'	-9.40	1.43	1.53
35	BB	1192	C	O3'-P	-9.40	1.49	1.61
35	BB	1306	G	C6-N1	-9.40	1.32	1.39
35	BB	1378	U	C2-N3	-9.40	1.31	1.37
36	BC	126	G	P-O5'	-9.40	1.50	1.59
40	BG	119	A	P-O5'	-9.40	1.50	1.59
85	AA	338	G	P-O5'	-9.40	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	596	A	N9-C4	-9.40	1.32	1.37
85	AA	709	A	N9-C4	-9.40	1.32	1.37
35	BB	1019	C	O3'-P	-9.39	1.49	1.61
85	AA	707	U	P-O5'	-9.39	1.50	1.59
34	BA	1567	G	P-O5'	-9.39	1.50	1.59
35	BB	494	C	C2'-C1'	-9.39	1.43	1.53
35	BB	97	U	O3'-P	-9.39	1.49	1.61
35	BB	661	G	N3-C4	-9.39	1.28	1.35
85	AA	156	G	N7-C5	-9.39	1.33	1.39
85	AA	682	C	O3'-P	-9.39	1.49	1.61
85	AA	2184	A	P-O5'	-9.39	1.50	1.59
85	AA	2214	A	C2'-C1'	-9.39	1.43	1.53
34	BA	159	U	C2'-C1'	-9.39	1.43	1.53
34	BA	825	G	N7-C5	-9.39	1.33	1.39
34	BA	1711	G	C5-C4	-9.39	1.31	1.38
35	BB	426	A	O3'-P	-9.39	1.49	1.61
34	BA	1816	G	C2'-C1'	-9.39	1.43	1.53
37	BD	95	G	N1-C2	-9.39	1.30	1.37
85	AA	2139	G	N9-C8	-9.39	1.31	1.37
35	BB	506	G	C3'-C2'	-9.39	1.42	1.52
35	BB	1486	C	O3'-P	-9.39	1.49	1.61
85	AA	629	A	O3'-P	-9.39	1.49	1.61
34	BA	328	A	C2'-C1'	-9.38	1.43	1.53
34	BA	1468	U	O3'-P	-9.38	1.49	1.61
85	AA	2191	C	O3'-P	-9.38	1.49	1.61
34	BA	708	C	O3'-P	-9.38	1.49	1.61
36	BC	144	C	C3'-C2'	-9.38	1.42	1.52
40	BG	49	A	C1'-N9	-9.38	1.33	1.46
85	AA	2182	A	O3'-P	-9.38	1.49	1.61
85	AA	1194	U	C2-N3	-9.38	1.31	1.37
34	BA	338	U	P-O5'	-9.38	1.50	1.59
85	AA	2175	U	O3'-P	-9.38	1.49	1.61
39	BF	34	C	C2-N3	-9.37	1.28	1.35
35	BB	79	U	O3'-P	-9.37	1.50	1.61
34	BA	727	G	N7-C5	-9.37	1.33	1.39
36	BC	41	A	C2'-C1'	-9.37	1.43	1.53
40	BG	73	U	O3'-P	-9.37	1.50	1.61
85	AA	706	U	C2-N3	-9.37	1.31	1.37
85	AA	2037	A	O3'-P	-9.37	1.50	1.61
34	BA	117	C	C2'-C1'	-9.37	1.43	1.53
34	BA	1550	G	C1'-N9	-9.37	1.33	1.46
35	BB	1311	G	O3'-P	-9.37	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	443	A	C2'-C1'	-9.37	1.43	1.53
34	BA	82	A	C2'-C1'	-9.36	1.43	1.53
85	AA	486	G	O3'-P	-9.36	1.50	1.61
35	BB	1449	G	O3'-P	-9.36	1.50	1.61
35	BB	1470	G	C4'-C3'	9.36	1.63	1.53
85	AA	97	A	C2'-C1'	-9.36	1.43	1.53
34	BA	193	C	O3'-P	-9.36	1.50	1.61
37	BD	32	A	C3'-C2'	-9.36	1.42	1.52
38	BE	116	U	C2-N3	-9.36	1.31	1.37
85	AA	352	G	N9-C4	-9.36	1.30	1.38
34	BA	23	A	C1'-N9	-9.36	1.33	1.46
34	BA	503	C	C2'-C1'	-9.36	1.43	1.53
34	BA	517	A	C5-C4	-9.36	1.32	1.38
34	BA	1070	G	N9-C4	-9.36	1.30	1.38
35	BB	391	G	N1-C2	-9.36	1.30	1.37
35	BB	1270	C	P-O5'	-9.36	1.50	1.59
85	AA	94	C	P-O5'	-9.36	1.50	1.59
39	BF	69	A	C2'-C1'	-9.35	1.43	1.53
34	BA	52	G	C2'-C1'	-9.35	1.43	1.53
34	BA	476	U	P-O5'	-9.35	1.50	1.59
34	BA	1657	A	N7-C5	-9.35	1.33	1.39
35	BB	832	C	C3'-C2'	-9.35	1.42	1.52
35	BB	1369	A	P-O5'	-9.35	1.50	1.59
85	AA	1541	G	O3'-P	-9.35	1.50	1.61
85	AA	2135	A	P-O5'	-9.35	1.50	1.59
40	BG	16	G	P-O5'	-9.35	1.50	1.59
34	BA	400	A	C1'-N9	-9.35	1.33	1.46
34	BA	1210	A	N9-C4	-9.35	1.32	1.37
38	BE	147	G	C1'-N9	-9.35	1.33	1.46
34	BA	857	C	C3'-C2'	-9.35	1.42	1.52
38	BE	124	G	P-O5'	-9.35	1.50	1.59
35	BB	552	C	O3'-P	-9.34	1.50	1.61
35	BB	1145	G	N7-C5	-9.34	1.33	1.39
39	BF	52	A	O3'-P	-9.34	1.50	1.61
34	BA	1282	G	C3'-C2'	-9.34	1.42	1.52
34	BA	1426	A	N7-C5	-9.34	1.33	1.39
36	BC	160	C	C3'-C2'	-9.34	1.42	1.52
34	BA	195	G	C6-N1	-9.34	1.33	1.39
34	BA	1257	U	O3'-P	-9.34	1.50	1.61
34	BA	453	A	P-O5'	-9.34	1.50	1.59
36	BC	43	A	C5-C4	-9.34	1.32	1.38
37	BD	89	G	N7-C5	-9.34	1.33	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
39	BF	43	U	O3'-P	-9.34	1.50	1.61
85	AA	1667	C	O3'-P	-9.34	1.50	1.61
34	BA	211	C	C3'-C2'	-9.33	1.42	1.52
34	BA	712	C	O3'-P	-9.33	1.50	1.61
34	BA	860	G	P-O5'	-9.33	1.50	1.59
35	BB	620	G	O3'-P	-9.33	1.50	1.61
34	BA	262	A	N3-C4	-9.33	1.29	1.34
34	BA	759	A	O3'-P	-9.33	1.50	1.61
34	BA	996	U	C3'-C2'	-9.33	1.42	1.52
34	BA	1312	A	O3'-P	-9.33	1.50	1.61
34	BA	768	G	P-O5'	-9.33	1.50	1.59
35	BB	1033	U	C2'-C1'	-9.33	1.43	1.53
35	BB	1281	G	N9-C4	-9.33	1.30	1.38
85	AA	463	G	N9-C4	-9.33	1.30	1.38
85	AA	1366	A	P-O5'	-9.33	1.50	1.59
34	BA	1486	U	O3'-P	-9.33	1.50	1.61
35	BB	384	A	N3-C4	-9.33	1.29	1.34
35	BB	454	U	P-O5'	-9.33	1.50	1.59
36	BC	113	G	C2'-C1'	-9.33	1.43	1.53
38	BE	123	A	O3'-P	-9.33	1.50	1.61
85	AA	179	G	P-O5'	-9.33	1.50	1.59
85	AA	768	C	P-O5'	-9.33	1.50	1.59
85	AA	1165	C	O3'-P	-9.33	1.50	1.61
85	AA	2007	G	N9-C4	-9.33	1.30	1.38
38	BE	146	U	N3-C4	-9.32	1.30	1.38
85	AA	470	C	C2-N3	-9.32	1.28	1.35
34	BA	1049	G	P-O5'	-9.32	1.50	1.59
34	BA	1695	G	C3'-C2'	-9.32	1.42	1.52
35	BB	1116	U	P-O5'	-9.32	1.50	1.59
38	BE	20	C	C2-N3	-9.32	1.28	1.35
40	BG	134	U	P-O5'	-9.32	1.50	1.59
85	AA	70	U	C2-N3	-9.32	1.31	1.37
34	BA	1725	U	C4'-C3'	-9.32	1.42	1.53
35	BB	1286	G	C2'-C1'	-9.32	1.43	1.53
38	BE	25	U	P-O5'	-9.32	1.50	1.59
85	AA	241	U	P-O5'	-9.32	1.50	1.59
85	AA	1463	A	O3'-P	-9.32	1.50	1.61
34	BA	220	U	P-O5'	-9.32	1.50	1.59
34	BA	419	U	C2-N3	-9.32	1.31	1.37
34	BA	506	U	C2'-C1'	-9.32	1.43	1.53
34	BA	1260	G	N1-C2	-9.32	1.30	1.37
38	BE	199	A	C2'-C1'	-9.32	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	452	A	O3'-P	-9.32	1.50	1.61
34	BA	670	U	O3'-P	-9.32	1.50	1.61
35	BB	52	G	C1'-N9	-9.32	1.33	1.46
35	BB	582	G	C4'-C3'	-9.32	1.43	1.53
35	BB	866	A	P-O5'	-9.32	1.50	1.59
41	BH	129	G	C2'-C1'	-9.32	1.43	1.53
34	BA	685	C	C5'-C4'	9.31	1.62	1.51
34	BA	1239	G	C6-N1	-9.31	1.33	1.39
85	AA	785	C	P-O5'	-9.31	1.50	1.59
36	BC	134	G	C2'-C1'	-9.31	1.43	1.53
85	AA	530	A	P-O5'	-9.31	1.50	1.59
85	AA	1571	A	C2'-C1'	-9.31	1.43	1.53
34	BA	51	C	C2-N3	-9.31	1.28	1.35
34	BA	88	C	C4'-O4'	-9.31	1.33	1.45
35	BB	1535	G	O3'-P	-9.31	1.50	1.61
35	BB	594	U	C2-N3	-9.31	1.31	1.37
35	BB	1014	U	P-O5'	-9.31	1.50	1.59
35	BB	1222	A	N7-C5	-9.31	1.33	1.39
83	Bx	48	GLY	CA-C	9.31	1.66	1.51
85	AA	463	G	C1'-N9	-9.31	1.33	1.46
85	AA	1575	G	P-O5'	-9.31	1.50	1.59
35	BB	118	A	N9-C4	-9.31	1.32	1.37
34	BA	1598	U	N1-C2	-9.31	1.30	1.38
40	BG	41	U	O3'-P	-9.31	1.50	1.61
34	BA	161	U	C4-C5	-9.30	1.35	1.43
34	BA	407	A	O3'-P	-9.31	1.50	1.61
34	BA	691	A	C2'-C1'	-9.31	1.43	1.53
34	BA	1178	U	P-O5'	-9.31	1.50	1.59
34	BA	1742	G	C2'-C1'	-9.30	1.43	1.53
85	AA	1122	U	P-O5'	-9.31	1.50	1.59
34	BA	909	G	C5-C4	-9.30	1.31	1.38
34	BA	925	G	N3-C4	-9.30	1.28	1.35
34	BA	1648	G	O3'-P	-9.30	1.50	1.61
34	BA	703	U	O3'-P	-9.30	1.50	1.61
34	BA	1796	A	C2'-C1'	-9.30	1.43	1.53
34	BA	1094	U	O3'-P	-9.30	1.50	1.61
35	BB	1145	G	O3'-P	-9.30	1.50	1.61
85	AA	351	C	P-O5'	-9.30	1.50	1.59
86	AB	6	G	P-O5'	-9.30	1.50	1.59
38	BE	16	C	O3'-P	-9.30	1.50	1.61
85	AA	548	G	C2'-C1'	-9.30	1.43	1.53
85	AA	2190	U	C2-N3	-9.30	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	349	G	C6-N1	-9.30	1.33	1.39
35	BB	576	A	O3'-P	-9.30	1.50	1.61
85	AA	1585	A	N9-C4	-9.30	1.32	1.37
35	BB	1145	G	P-O5'	-9.30	1.50	1.59
34	BA	321	G	C5-C4	-9.29	1.31	1.38
34	BA	843	G	O3'-P	-9.29	1.50	1.61
34	BA	930	A	N9-C4	-9.30	1.32	1.37
35	BB	806	U	C4-C5	-9.29	1.35	1.43
41	BH	41	A	C2'-C1'	-9.29	1.43	1.53
85	AA	1258	U	P-O5'	-9.29	1.50	1.59
85	AA	2031	C	P-O5'	-9.29	1.50	1.59
34	BA	936	A	C3'-C2'	-9.29	1.42	1.52
38	BE	195	G	C2-N3	9.29	1.40	1.32
85	AA	190	A	C3'-C2'	-9.29	1.42	1.52
85	AA	190	A	C5-C4	-9.29	1.32	1.38
34	BA	1783	C	C2'-C1'	-9.29	1.43	1.53
85	AA	318	A	N9-C4	-9.29	1.32	1.37
85	AA	1928	A	C2'-C1'	-9.29	1.43	1.53
85	AA	1522	U	P-O5'	-9.29	1.50	1.59
34	BA	17	A	C2'-C1'	-9.29	1.43	1.53
35	BB	1130	U	O3'-P	-9.29	1.50	1.61
34	BA	27	G	O3'-P	-9.28	1.50	1.61
34	BA	1260	G	O3'-P	-9.28	1.50	1.61
85	AA	81	A	O3'-P	-9.29	1.50	1.61
85	AA	1356	U	O3'-P	-9.28	1.50	1.61
34	BA	753	G	O3'-P	-9.28	1.50	1.61
34	BA	350	C	P-O5'	-9.28	1.50	1.59
34	BA	825	G	C3'-C2'	-9.28	1.42	1.52
35	BB	94	A	N9-C4	-9.28	1.32	1.37
35	BB	709	G	C1'-N9	-9.28	1.33	1.46
85	AA	598	C	O3'-P	-9.28	1.50	1.61
85	AA	1528	A	P-O5'	-9.28	1.50	1.59
85	AA	1471	G	N7-C5	-9.28	1.33	1.39
85	AA	2175	U	P-O5'	-9.28	1.50	1.59
34	BA	1715	C	P-O5'	-9.28	1.50	1.59
35	BB	586	U	O3'-P	-9.28	1.50	1.61
35	BB	1292	G	N3-C4	-9.28	1.28	1.35
38	BE	89	G	C2'-C1'	-9.28	1.43	1.53
38	BE	105	A	C2'-C1'	-9.28	1.43	1.53
85	AA	1809	G	P-O5'	-9.28	1.50	1.59
34	BA	815	C	O3'-P	-9.28	1.50	1.61
85	AA	1018	G	P-O5'	-9.28	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1200	A	O3'-P	-9.28	1.50	1.61
34	BA	1690	U	O3'-P	-9.28	1.50	1.61
35	BB	107	A	N9-C4	-9.28	1.32	1.37
35	BB	692	G	C3'-C2'	-9.28	1.42	1.52
39	BF	18	U	C2-N3	-9.28	1.31	1.37
40	BG	176	G	C3'-C2'	-9.28	1.42	1.52
85	AA	497	G	P-O5'	-9.28	1.50	1.59
34	BA	29	U	O3'-P	-9.27	1.50	1.61
34	BA	1240	G	N9-C4	-9.27	1.30	1.38
35	BB	441	G	C2'-C1'	-9.27	1.43	1.53
35	BB	786	A	C2'-C1'	-9.27	1.43	1.53
35	BB	1293	C	O3'-P	-9.27	1.50	1.61
36	BC	45	C	C2'-C1'	-9.27	1.43	1.53
85	AA	1196	C	C2-N3	-9.27	1.28	1.35
34	BA	1107	A	P-O5'	-9.27	1.50	1.59
41	BH	114	G	O3'-P	-9.27	1.50	1.61
85	AA	112	A	P-O5'	-9.27	1.50	1.59
85	AA	469	G	C1'-N9	-9.27	1.33	1.46
34	BA	1311	G	C6-N1	-9.27	1.33	1.39
85	AA	774	C	P-O5'	-9.27	1.50	1.59
34	BA	495	A	C3'-C2'	-9.26	1.42	1.52
41	BH	34	G	C6-N1	-9.26	1.33	1.39
34	BA	1628	A	O3'-P	-9.26	1.50	1.61
36	BC	43	A	O3'-P	-9.26	1.50	1.61
40	BG	79	U	C2-N3	-9.26	1.31	1.37
35	BB	1375	G	C6-N1	-9.26	1.33	1.39
35	BB	27	C	P-O5'	-9.26	1.50	1.59
85	AA	71	G	C6-N1	-9.26	1.33	1.39
35	BB	1341	U	P-O5'	-9.26	1.50	1.59
34	BA	114	U	P-O5'	-9.25	1.50	1.59
34	BA	1093	G	C6-N1	-9.25	1.33	1.39
36	BC	119	G	O3'-P	-9.25	1.50	1.61
85	AA	5	U	C2'-C1'	-9.25	1.43	1.53
85	AA	1507	G	N9-C4	-9.25	1.30	1.38
85	AA	1830	U	C2-N3	-9.25	1.31	1.37
34	BA	1217	A	O3'-P	-9.25	1.50	1.61
34	BA	574	U	O3'-P	-9.25	1.50	1.61
34	BA	1265	G	O3'-P	-9.25	1.50	1.61
35	BB	447	C	O3'-P	-9.25	1.50	1.61
85	AA	519	A	C2'-C1'	-9.25	1.43	1.53
85	AA	1147	A	C1'-N9	-9.25	1.33	1.46
34	BA	137	C	P-O5'	-9.25	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	380	G	C2-N2	-9.25	1.25	1.34
40	BG	37	G	C3'-C2'	-9.25	1.42	1.52
40	BG	128	U	C2-N3	-9.25	1.31	1.37
40	BG	133	C	O3'-P	-9.25	1.50	1.61
85	AA	1452	C	O3'-P	-9.25	1.50	1.61
85	AA	1106	A	C2'-C1'	-9.25	1.43	1.53
34	BA	314	A	P-O5'	-9.24	1.50	1.59
34	BA	813	C	P-O5'	-9.24	1.50	1.59
35	BB	1400	C	O3'-P	-9.24	1.50	1.61
36	BC	7	U	P-O5'	-9.24	1.50	1.59
39	BF	48	G	O3'-P	-9.24	1.50	1.61
40	BG	9	G	C3'-O3'	-9.24	1.29	1.42
40	BG	79	U	O3'-P	-9.24	1.50	1.61
34	BA	103	G	C2'-C1'	-9.24	1.43	1.53
34	BA	605	G	P-O5'	-9.24	1.50	1.59
34	BA	1664	C	P-O5'	-9.24	1.50	1.59
40	BG	132	U	P-O5'	-9.24	1.50	1.59
41	BH	72	G	C5'-C4'	9.24	1.62	1.51
85	AA	150	U	O3'-P	-9.24	1.50	1.61
34	BA	1467	U	O3'-P	-9.24	1.50	1.61
35	BB	608	A	P-O5'	-9.24	1.50	1.59
85	AA	1151	G	N9-C4	-9.24	1.30	1.38
34	BA	517	A	C1'-N9	-9.24	1.33	1.46
34	BA	631	G	P-O5'	-9.24	1.50	1.59
85	AA	2174	G	N7-C5	-9.24	1.33	1.39
34	BA	356	C	C2'-C1'	-9.24	1.43	1.53
34	BA	1685	C	O3'-P	-9.24	1.50	1.61
35	BB	674	C	C2'-C1'	-9.24	1.43	1.53
36	BC	69	U	P-O5'	-9.24	1.50	1.59
40	BG	28	A	N9-C4	-9.24	1.32	1.37
40	BG	49	A	N9-C4	-9.24	1.32	1.37
85	AA	161	A	N9-C4	-9.24	1.32	1.37
85	AA	1149	A	O3'-P	-9.24	1.50	1.61
34	BA	1595	G	O3'-P	-9.23	1.50	1.61
34	BA	1684	A	P-O5'	-9.23	1.50	1.59
35	BB	77	A	C1'-N9	-9.23	1.33	1.46
35	BB	404	A	C3'-C2'	-9.23	1.42	1.52
85	AA	1172	A	P-O5'	-9.23	1.50	1.59
85	AA	1515	A	P-O5'	-9.23	1.50	1.59
34	BA	776	U	O3'-P	-9.23	1.50	1.61
35	BB	634	A	O3'-P	-9.23	1.50	1.61
85	AA	903	G	C4'-C3'	-9.23	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	386	G	N9-C4	-9.23	1.30	1.38
35	BB	1106	G	C6-N1	-9.23	1.33	1.39
85	AA	1650	G	N7-C5	-9.23	1.33	1.39
85	AA	1839	G	P-O5'	-9.23	1.50	1.59
85	AA	2001	C	O3'-P	-9.23	1.50	1.61
35	BB	1147	G	C3'-C2'	-9.23	1.42	1.52
85	AA	1302	A	C5'-C4'	9.23	1.62	1.51
34	BA	994	G	N9-C4	-9.23	1.30	1.38
34	BA	1017	C	O3'-P	-9.23	1.50	1.61
34	BA	1734	U	P-O5'	-9.23	1.50	1.59
85	AA	1492	U	C3'-C2'	-9.23	1.42	1.52
41	BH	41	A	C5-C4	-9.23	1.32	1.38
35	BB	421	U	C2-N3	-9.22	1.31	1.37
34	BA	937	G	C3'-C2'	-9.22	1.42	1.52
35	BB	121	A	C5-C4	-9.22	1.32	1.38
36	BC	126	G	C2'-C1'	-9.22	1.43	1.53
85	AA	2174	G	N9-C8	-9.22	1.31	1.37
35	BB	1444	U	N3-C4	-9.22	1.30	1.38
41	BH	131	A	O3'-P	-9.22	1.50	1.61
85	AA	604	C	C2'-C1'	-9.22	1.43	1.53
85	AA	1000	U	O4'-C1'	-9.22	1.29	1.41
34	BA	1457	C	P-O5'	-9.22	1.50	1.59
35	BB	1323	U	C3'-C2'	-9.22	1.42	1.52
35	BB	665	A	N7-C5	-9.22	1.33	1.39
35	BB	1109	A	O3'-P	-9.22	1.50	1.61
34	BA	614	A	P-O5'	-9.21	1.50	1.59
35	BB	129	U	C2'-C1'	-9.21	1.43	1.53
34	BA	60	A	N9-C4	-9.21	1.32	1.37
38	BE	122	G	C2'-C1'	-9.21	1.43	1.53
40	BG	24	A	N9-C4	-9.21	1.32	1.37
85	AA	1721	A	P-O5'	-9.21	1.50	1.59
34	BA	741	A	N7-C5	-9.21	1.33	1.39
35	BB	374	A	C5-C4	-9.21	1.32	1.38
85	AA	928	U	O3'-P	-9.21	1.50	1.61
85	AA	2008	G	C2-N3	-9.21	1.25	1.32
34	BA	655	U	C2-N3	-9.21	1.31	1.37
40	BG	16	G	C3'-C2'	-9.21	1.42	1.52
85	AA	1240	A	C1'-N9	-9.21	1.33	1.46
34	BA	752	A	N9-C4	-9.21	1.32	1.37
34	BA	809	U	C2'-C1'	-9.21	1.43	1.53
34	BA	1069	U	P-O5'	-9.21	1.50	1.59
34	BA	1436	A	P-O5'	-9.21	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1696	G	N1-C2	-9.21	1.30	1.37
35	BB	1118	G	P-O5'	-9.20	1.50	1.59
85	AA	1221	G	C2'-C1'	-9.20	1.43	1.53
34	BA	5	C	C3'-C2'	-9.20	1.42	1.52
34	BA	121	A	C2'-C1'	-9.20	1.43	1.53
34	BA	1284	G	C1'-N9	-9.20	1.33	1.46
35	BB	60	A	C1'-N9	-9.20	1.33	1.46
35	BB	451	A	C5-C4	-9.20	1.32	1.38
35	BB	1087	A	C3'-C2'	-9.20	1.42	1.52
41	BH	105	U	C2-N3	-9.20	1.31	1.37
85	AA	1701	G	C2-N3	-9.20	1.25	1.32
34	BA	1686	G	C2'-C1'	-9.20	1.43	1.53
85	AA	533	C	C2'-C1'	-9.20	1.43	1.53
35	BB	84	G	P-O5'	-9.20	1.50	1.59
85	AA	696	G	C2-N3	-9.20	1.25	1.32
85	AA	982	G	C2'-C1'	-9.20	1.43	1.53
34	BA	1234	U	C3'-C2'	-9.20	1.42	1.52
34	BA	1255	G	N9-C4	-9.20	1.30	1.38
34	BA	1687	A	C2'-C1'	-9.20	1.43	1.53
36	BC	12	A	C5-C4	-9.20	1.32	1.38
37	BD	107	G	N7-C5	-9.20	1.33	1.39
38	BE	145	A	O3'-P	-9.20	1.50	1.61
85	AA	545	A	C2'-C1'	-9.20	1.43	1.53
34	BA	1098	G	O3'-P	-9.20	1.50	1.61
35	BB	627	G	P-O5'	-9.20	1.50	1.59
35	BB	1129	C	O3'-P	-9.20	1.50	1.61
35	BB	1146	C	O3'-P	-9.20	1.50	1.61
35	BB	1294	C	P-O5'	-9.20	1.50	1.59
40	BG	124	A	O3'-P	-9.20	1.50	1.61
40	BG	164	U	O3'-P	-9.20	1.50	1.61
85	AA	681	G	N9-C4	-9.20	1.30	1.38
85	AA	2193	A	O3'-P	-9.20	1.50	1.61
85	AA	2241	C	P-O5'	-9.20	1.50	1.59
35	BB	1361	A	P-O5'	-9.19	1.50	1.59
37	BD	65	G	C5-C4	-9.19	1.31	1.38
40	BG	39	A	C1'-N9	-9.20	1.33	1.46
85	AA	1190	G	N9-C4	-9.20	1.30	1.38
85	AA	2137	A	N3-C4	-9.20	1.29	1.34
34	BA	12	G	C2-N3	-9.19	1.25	1.32
34	BA	395	G	N3-C4	-9.19	1.29	1.35
34	BA	915	A	O3'-P	-9.19	1.50	1.61
35	BB	415	A	N3-C4	-9.19	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1464	G	C3'-C2'	-9.19	1.42	1.52
86	AB	68	C	C2-N3	-9.19	1.28	1.35
34	BA	76	U	C2-N3	-9.19	1.31	1.37
85	AA	2074	G	P-O5'	-9.19	1.50	1.59
41	BH	127	A	C1'-N9	-9.19	1.33	1.46
85	AA	854	A	C5'-C4'	9.19	1.62	1.51
34	BA	556	A	C3'-O3'	9.19	1.55	1.42
40	BG	20	U	C2-N3	-9.19	1.31	1.37
85	AA	925	G	O3'-P	-9.19	1.50	1.61
34	BA	736	G	O3'-P	-9.18	1.50	1.61
41	BH	133	U	C4'-C3'	-9.18	1.43	1.53
34	BA	232	U	C2'-C1'	-9.18	1.43	1.53
34	BA	1644	A	N9-C4	-9.18	1.32	1.37
35	BB	1032	U	O3'-P	-9.18	1.50	1.61
85	AA	1927	G	O3'-P	-9.18	1.50	1.61
85	AA	2106	C	P-O5'	-9.18	1.50	1.59
34	BA	1208	U	C2-N3	-9.18	1.31	1.37
36	BC	114	C	P-O5'	-9.18	1.50	1.59
85	AA	1542	A	O3'-P	-9.18	1.50	1.61
34	BA	1570	C	O3'-P	-9.18	1.50	1.61
35	BB	679	G	C6-N1	-9.18	1.33	1.39
34	BA	1808	A	N7-C5	-9.18	1.33	1.39
35	BB	390	G	C6-N1	-9.18	1.33	1.39
34	BA	757	G	N9-C4	-9.18	1.30	1.38
34	BA	917	C	C3'-C2'	-9.18	1.42	1.52
34	BA	1167	A	P-O5'	-9.18	1.50	1.59
34	BA	1707	C	C3'-C2'	-9.18	1.42	1.52
38	BE	48	G	C6-N1	-9.18	1.33	1.39
85	AA	887	A	C1'-N9	-9.18	1.34	1.46
85	AA	1565	G	O3'-P	-9.18	1.50	1.61
85	AA	2186	U	C2-N3	-9.18	1.31	1.37
34	BA	293	A	O3'-P	-9.17	1.50	1.61
34	BA	1019	C	C3'-C2'	-9.17	1.42	1.52
35	BB	1430	G	C4'-C3'	-9.17	1.43	1.53
34	BA	1164	C	O3'-P	-9.17	1.50	1.61
34	BA	1221	A	C5'-C4'	-9.17	1.32	1.38
35	BB	103	C	C2-N3	-9.17	1.28	1.35
35	BB	1408	G	N7-C5	-9.17	1.33	1.39
85	AA	1862	C	P-O5'	-9.17	1.50	1.59
34	BA	353	U	O3'-P	-9.17	1.50	1.61
35	BB	1070	G	C2'-C1'	-9.17	1.43	1.53
35	BB	1085	C	C2-N3	-9.17	1.28	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	198	A	O3'-P	-9.17	1.50	1.61
34	BA	1024	A	N9-C4	-9.17	1.32	1.37
34	BA	1409	A	C2'-C1'	-9.17	1.43	1.53
35	BB	1485	G	C2'-C1'	-9.17	1.43	1.53
35	BB	1198	C	O3'-P	-9.16	1.50	1.61
36	BC	81	U	O3'-P	-9.16	1.50	1.61
34	BA	73	G	C2-N2	-9.16	1.25	1.34
34	BA	800	G	C4'-O4'	-9.16	1.33	1.45
35	BB	570	A	O3'-P	-9.16	1.50	1.61
36	BC	134	G	C6-N1	-9.16	1.33	1.39
85	AA	1670	U	C2-N3	-9.16	1.31	1.37
34	BA	617	G	O4'-C1'	-9.16	1.29	1.41
37	BD	87	G	C2-N3	-9.16	1.25	1.32
35	BB	463	C	P-O5'	-9.16	1.50	1.59
35	BB	611	U	P-O5'	-9.16	1.50	1.59
38	BE	7	U	O3'-P	-9.16	1.50	1.61
85	AA	495	G	C3'-C2'	-9.16	1.42	1.52
85	AA	2238	C	N1-C6	-9.16	1.31	1.37
34	BA	672	G	N9-C4	-9.15	1.30	1.38
34	BA	18	G	P-O5'	-9.15	1.50	1.59
34	BA	1502	G	C5-C4	-9.15	1.31	1.38
35	BB	429	C	C2'-C1'	-9.15	1.43	1.53
35	BB	578	G	N9-C4	-9.15	1.30	1.38
41	BH	52	G	C2'-C1'	-9.15	1.43	1.53
85	AA	917	A	C2'-C1'	-9.15	1.43	1.53
85	AA	177	A	O3'-P	-9.15	1.50	1.61
85	AA	939	A	C5-C4	-9.15	1.32	1.38
85	AA	1896	G	N7-C5	-9.15	1.33	1.39
35	BB	131	A	O3'-P	-9.15	1.50	1.61
34	BA	1559	C	C2'-C1'	-9.15	1.43	1.53
34	BA	1707	C	P-O5'	-9.15	1.50	1.59
41	BH	112	U	C2-N3	-9.15	1.31	1.37
34	BA	126	G	N7-C5	-9.15	1.33	1.39
34	BA	341	U	O3'-P	-9.15	1.50	1.61
34	BA	391	U	P-O5'	-9.15	1.50	1.59
34	BA	1816	G	C1'-N9	-9.15	1.34	1.46
35	BB	1311	G	P-O5'	-9.15	1.50	1.59
85	AA	376	C	O3'-P	-9.15	1.50	1.61
85	AA	440	U	C2'-C1'	-9.15	1.43	1.53
85	AA	869	A	P-O5'	-9.15	1.50	1.59
34	BA	333	A	N7-C5	-9.14	1.33	1.39
34	BA	995	A	P-O5'	-9.14	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	401	U	P-O5'	-9.14	1.50	1.59
34	BA	515	U	C5'-C4'	9.14	1.62	1.51
34	BA	587	U	C4-C5	-9.14	1.35	1.43
34	BA	1511	C	O3'-P	-9.14	1.50	1.61
34	BA	1223	C	C3'-C2'	-9.14	1.42	1.52
40	BG	104	A	P-O5'	-9.14	1.50	1.59
85	AA	2203	C	C3'-C2'	-9.14	1.42	1.52
34	BA	1229	G	N9-C4	-9.14	1.30	1.38
35	BB	686	A	O3'-P	-9.14	1.50	1.61
85	AA	1518	A	P-O5'	-9.14	1.50	1.59
35	BB	959	C	P-O5'	-9.14	1.50	1.59
36	BC	14	G	N9-C8	-9.14	1.31	1.37
38	BE	147	G	O3'-P	-9.14	1.50	1.61
39	BF	62	U	C3'-C2'	-9.14	1.42	1.52
85	AA	740	A	C2'-C1'	-9.14	1.43	1.53
35	BB	1445	A	P-O5'	-9.14	1.50	1.59
34	BA	291	C	O3'-P	-9.13	1.50	1.61
34	BA	426	A	P-O5'	-9.13	1.50	1.59
34	BA	1503	U	C3'-C2'	-9.13	1.42	1.52
35	BB	553	U	P-O5'	-9.13	1.50	1.59
36	BC	65	G	O3'-P	-9.13	1.50	1.61
85	AA	1247	A	C2'-C1'	-9.13	1.43	1.53
34	BA	1563	G	C6-N1	-9.13	1.33	1.39
34	BA	1822	U	O3'-P	-9.13	1.50	1.61
35	BB	640	A	P-O5'	-9.13	1.50	1.59
35	BB	1342	C	C2-N3	-9.13	1.28	1.35
40	BG	5	G	O3'-P	-9.13	1.50	1.61
85	AA	177	A	C1'-N9	-9.13	1.34	1.46
85	AA	2206	A	P-O5'	-9.13	1.50	1.59
34	BA	694	G	N7-C5	-9.13	1.33	1.39
40	BG	76	C	O3'-P	-9.13	1.50	1.61
85	AA	857	G	P-O5'	-9.13	1.50	1.59
34	BA	669	U	C2'-C1'	-9.13	1.43	1.53
34	BA	1224	A	N9-C4	-9.13	1.32	1.37
34	BA	1424	G	N9-C4	-9.13	1.30	1.38
35	BB	1294	C	C4'-C3'	-9.13	1.43	1.53
57	BX	87	TYR	CB-CG	9.13	1.65	1.51
85	AA	1664	G	P-O5'	-9.13	1.50	1.59
35	BB	551	C	P-O5'	-9.13	1.50	1.59
40	BG	4	A	P-O5'	-9.13	1.50	1.59
85	AA	1187	G	O3'-P	-9.13	1.50	1.61
85	AA	1504	A	C1'-N9	-9.13	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1864	G	N9-C4	-9.13	1.30	1.38
34	BA	1067	G	P-O5'	-9.12	1.50	1.59
35	BB	1130	U	C2-N3	-9.12	1.31	1.37
35	BB	1505	U	C2'-C1'	-9.12	1.43	1.53
34	BA	330	A	C2'-C1'	-9.12	1.43	1.53
34	BA	399	G	C2-N3	-9.12	1.25	1.32
35	BB	1071	G	O3'-P	-9.12	1.50	1.61
36	BC	35	C	C3'-C2'	-9.12	1.42	1.52
85	AA	509	C	C2'-C1'	-9.12	1.43	1.53
35	BB	1161	G	O3'-P	-9.12	1.50	1.61
38	BE	205	G	P-O5'	-9.12	1.50	1.59
85	AA	31	C	O3'-P	-9.12	1.50	1.61
34	BA	338	U	O3'-P	-9.12	1.50	1.61
35	BB	612	A	O3'-P	-9.12	1.50	1.61
85	AA	1551	G	C6-N1	-9.12	1.33	1.39
85	AA	2138	G	N7-C5	-9.12	1.33	1.39
85	AA	511	A	O3'-P	-9.12	1.50	1.61
85	AA	2091	C	C2'-C1'	-9.12	1.43	1.53
34	BA	45	A	O3'-P	-9.12	1.50	1.61
34	BA	1109	G	C1'-N9	-9.12	1.34	1.46
35	BB	23	U	P-O5'	-9.12	1.50	1.59
35	BB	71	A	P-O5'	-9.12	1.50	1.59
35	BB	461	U	C2-N3	-9.12	1.31	1.37
35	BB	1252	G	O3'-P	-9.12	1.50	1.61
85	AA	96	C	C2-N3	-9.12	1.28	1.35
85	AA	831	C	C2'-C1'	-9.12	1.43	1.53
85	AA	960	G	C2'-C1'	-9.12	1.43	1.53
85	AA	997	U	P-O5'	-9.12	1.50	1.59
85	AA	1851	A	N9-C4	-9.12	1.32	1.37
40	BG	167	C	O3'-P	-9.11	1.50	1.61
34	BA	1475	G	N9-C4	-9.11	1.30	1.38
85	AA	440	U	O3'-P	-9.11	1.50	1.61
36	BC	49	G	O3'-P	-9.11	1.50	1.61
36	BC	124	A	C5'-C4'	9.11	1.62	1.51
85	AA	545	A	N9-C4	-9.11	1.32	1.37
85	AA	1140	G	O3'-P	-9.11	1.50	1.61
34	BA	28	C	P-O5'	-9.11	1.50	1.59
34	BA	1000	G	O3'-P	-9.11	1.50	1.61
34	BA	1066	A	P-O5'	-9.11	1.50	1.59
34	BA	1312	A	C2'-C1'	-9.11	1.43	1.53
34	BA	1612	C	P-O5'	-9.11	1.50	1.59
35	BB	1289	G	P-O5'	-9.11	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	512	U	P-O5'	-9.11	1.50	1.59
85	AA	1220	A	N7-C5	-9.11	1.33	1.39
34	BA	134	U	C3'-C2'	-9.10	1.42	1.52
34	BA	349	G	C1'-N9	-9.10	1.34	1.46
34	BA	721	A	N7-C5	-9.10	1.33	1.39
36	BC	91	G	C1'-N9	-9.10	1.34	1.46
40	BG	10	U	O4'-C1'	-9.10	1.29	1.41
34	BA	1603	A	N3-C4	-9.10	1.29	1.34
35	BB	42	A	N9-C4	-9.10	1.32	1.37
34	BA	17	A	C3'-C2'	-9.10	1.42	1.52
34	BA	757	G	P-O5'	-9.10	1.50	1.59
34	BA	1234	U	O3'-P	-9.10	1.50	1.61
35	BB	1246	C	C3'-C2'	-9.10	1.42	1.52
38	BE	129	G	P-O5'	-9.10	1.50	1.59
38	BE	203	C	P-O5'	-9.10	1.50	1.59
40	BG	35	G	C2'-C1'	-9.10	1.43	1.53
85	AA	969	U	O3'-P	-9.09	1.50	1.61
85	AA	1221	G	C1'-N9	-9.09	1.34	1.46
34	BA	128	C	C3'-C2'	-9.09	1.42	1.52
34	BA	460	G	C2'-C1'	-9.09	1.43	1.53
34	BA	481	A	O3'-P	-9.09	1.50	1.61
34	BA	1803	A	C1'-N9	-9.09	1.34	1.46
35	BB	632	U	C2'-C1'	-9.09	1.43	1.53
35	BB	410	A	N9-C4	-9.09	1.32	1.37
36	BC	46	G	P-O5'	-9.09	1.50	1.59
85	AA	1518	A	O3'-P	-9.09	1.50	1.61
40	BG	81	G	O3'-P	-9.09	1.50	1.61
85	AA	1458	G	N1-C2	-9.09	1.30	1.37
85	AA	2043	A	N9-C4	-9.09	1.32	1.37
34	BA	123	C	P-O5'	-9.09	1.50	1.59
34	BA	513	U	O3'-P	-9.09	1.50	1.61
34	BA	723	C	P-O5'	-9.09	1.50	1.59
34	BA	783	U	C2-N3	-9.09	1.31	1.37
35	BB	460	C	C2'-C1'	-9.09	1.43	1.53
37	BD	23	A	N9-C4	-9.09	1.32	1.37
85	AA	686	U	P-O5'	-9.09	1.50	1.59
85	AA	2146	G	C5-C4	-9.09	1.31	1.38
85	AA	2183	U	O3'-P	-9.09	1.50	1.61
36	BC	154	A	O3'-P	-9.09	1.50	1.61
38	BE	23	G	O3'-P	-9.09	1.50	1.61
40	BG	16	G	C1'-N9	-9.09	1.34	1.46
36	BC	129	C	P-O5'	-9.08	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	28	U	O3'-P	-9.08	1.50	1.61
41	BH	35	G	O3'-P	-9.08	1.50	1.61
34	BA	889	U	C3'-C2'	-9.08	1.42	1.52
34	BA	1510	C	C2-N3	-9.08	1.28	1.35
35	BB	425	G	C6-N1	-9.08	1.33	1.39
85	AA	1677	A	O3'-P	-9.08	1.50	1.61
37	BD	69	U	C2-N3	-9.08	1.31	1.37
34	BA	1685	C	C3'-C2'	-9.08	1.42	1.52
38	BE	68	U	C2-N3	-9.08	1.31	1.37
85	AA	1168	C	P-O5'	-9.08	1.50	1.59
34	BA	610	A	C4'-C3'	-9.07	1.43	1.53
34	BA	931	G	C1'-N9	-9.07	1.34	1.46
35	BB	775	U	C3'-C2'	-9.07	1.42	1.52
85	AA	817	G	C2'-C1'	-9.07	1.43	1.53
34	BA	478	G	C1'-N9	-9.07	1.34	1.46
85	AA	714	U	P-O5'	-9.07	1.50	1.59
35	BB	649	A	C4'-C3'	-9.07	1.43	1.53
35	BB	781	U	C2'-C1'	-9.07	1.43	1.53
34	BA	17	A	N9-C4	-9.07	1.32	1.37
34	BA	222	C	P-O5'	-9.07	1.50	1.59
35	BB	450	A	P-O5'	-9.07	1.50	1.59
34	BA	306	G	P-O5'	-9.07	1.50	1.59
34	BA	1103	G	P-O5'	-9.07	1.50	1.59
34	BA	1612	C	C3'-C2'	-9.07	1.42	1.52
35	BB	1359	G	P-O5'	-9.07	1.50	1.59
37	BD	43	U	O3'-P	-9.07	1.50	1.61
38	BE	64	A	P-O5'	-9.07	1.50	1.59
41	BH	67	G	P-O5'	-9.07	1.50	1.59
85	AA	620	U	C2-N3	-9.07	1.31	1.37
34	BA	1166	A	C2'-C1'	-9.06	1.43	1.53
85	AA	555	C	P-O5'	-9.06	1.50	1.59
34	BA	147	U	P-O5'	-9.06	1.50	1.59
34	BA	1639	U	C2-N3	-9.06	1.31	1.37
34	BA	341	U	P-O5'	-9.06	1.50	1.59
34	BA	375	C	P-O5'	-9.06	1.50	1.59
85	AA	627	A	O3'-P	-9.06	1.50	1.61
85	AA	1495	G	C1'-N9	-9.06	1.34	1.46
36	BC	160	C	O3'-P	-9.06	1.50	1.61
34	BA	1050	A	N7-C5	-9.06	1.33	1.39
34	BA	28	C	C2-N3	-9.06	1.28	1.35
36	BC	147	G	P-O5'	-9.06	1.50	1.59
37	BD	59	G	P-O5'	-9.06	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	111	U	C2-N3	-9.06	1.31	1.37
85	AA	1453	U	P-O5'	-9.06	1.50	1.59
34	BA	1276	G	C1'-N9	-9.05	1.34	1.46
35	BB	379	U	C3'-C2'	-9.05	1.42	1.52
38	BE	1	U	C2-N3	-9.06	1.31	1.37
39	BF	63	U	P-O5'	-9.06	1.50	1.59
40	BG	26	G	O3'-P	-9.05	1.50	1.61
40	BG	40	G	C1'-N9	-9.05	1.34	1.46
85	AA	1281	G	C1'-N9	-9.05	1.34	1.46
85	AA	2005	U	C4'-C3'	-9.05	1.43	1.53
34	BA	378	C	O3'-P	-9.05	1.50	1.61
34	BA	430	A	C2'-C1'	-9.05	1.43	1.53
34	BA	728	A	O3'-P	-9.05	1.50	1.61
36	BC	16	A	O3'-P	-9.05	1.50	1.61
38	BE	100	U	O3'-P	-9.05	1.50	1.61
38	BE	192	A	O3'-P	-9.05	1.50	1.61
34	BA	1262	A	C1'-N9	-9.05	1.34	1.46
40	BG	89	A	N9-C4	-9.05	1.32	1.37
40	BG	174	G	O3'-P	-9.05	1.50	1.61
34	BA	455	A	O3'-P	-9.05	1.50	1.61
85	AA	188	G	C2'-C1'	-9.05	1.43	1.53
85	AA	1580	A	O3'-P	-9.05	1.50	1.61
34	BA	413	A	N7-C5	-9.05	1.33	1.39
34	BA	890	G	C2'-C1'	-9.05	1.43	1.53
35	BB	547	A	O3'-P	-9.05	1.50	1.61
34	BA	258	C	O3'-P	-9.04	1.50	1.61
34	BA	1210	A	N7-C5	-9.05	1.33	1.39
35	BB	1371	G	C2'-C1'	-9.05	1.43	1.53
36	BC	163	A	O3'-P	-9.05	1.50	1.61
85	AA	1248	U	C4'-C3'	-9.04	1.43	1.53
34	BA	492	G	C1'-N9	-9.04	1.34	1.46
85	AA	1249	U	C2'-C1'	-9.04	1.43	1.53
85	AA	620	U	C2'-C1'	-9.04	1.43	1.53
34	BA	369	A	C3'-C2'	-9.04	1.42	1.52
34	BA	401	A	C4'-C3'	-9.04	1.43	1.53
35	BB	86	A	C4'-C3'	-9.04	1.43	1.53
35	BB	1163	U	P-O5'	-9.04	1.50	1.59
35	BB	1523	U	C2'-C1'	-9.04	1.43	1.53
85	AA	1473	U	C2'-C1'	-9.04	1.43	1.53
34	BA	714	G	C2'-C1'	-9.04	1.43	1.53
35	BB	1081	U	O3'-P	-9.04	1.50	1.61
35	BB	1161	G	C2'-C1'	-9.04	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	162	A	P-O5'	-9.04	1.50	1.59
85	AA	1248	U	C3'-C2'	-9.04	1.42	1.52
34	BA	27	G	C4'-C3'	-9.04	1.43	1.53
35	BB	805	G	C2'-C1'	-9.04	1.43	1.53
35	BB	597	C	C2'-C1'	-9.03	1.43	1.53
39	BF	42	G	C6-N1	-9.04	1.33	1.39
85	AA	868	A	O3'-P	-9.03	1.50	1.61
34	BA	275	C	O3'-P	-9.03	1.50	1.61
35	BB	1034	U	C2-N3	-9.03	1.31	1.37
85	AA	1935	G	P-O5'	-9.03	1.50	1.59
85	AA	2187	G	C2'-C1'	-9.03	1.43	1.53
34	BA	1830	A	C4'-C3'	-9.03	1.43	1.53
85	AA	1539	A	P-O5'	-9.03	1.50	1.59
34	BA	1482	A	C2'-C1'	-9.02	1.43	1.53
38	BE	115	U	P-O5'	-9.02	1.50	1.59
38	BE	23	G	N9-C8	-9.02	1.31	1.37
85	AA	592	C	O3'-P	-9.02	1.50	1.61
85	AA	838	G	C2'-C1'	-9.02	1.43	1.53
85	AA	1977	G	P-O5'	-9.02	1.50	1.59
34	BA	774	A	O3'-P	-9.02	1.50	1.61
35	BB	997	G	P-O5'	-9.02	1.50	1.59
36	BC	95	A	P-O5'	-9.02	1.50	1.59
41	BH	22	A	N9-C4	-9.02	1.32	1.37
34	BA	610	A	C2'-C1'	-9.02	1.43	1.53
34	BA	1826	C	O3'-P	-9.02	1.50	1.61
35	BB	559	U	P-O5'	-9.02	1.50	1.59
36	BC	55	U	C2-N3	-9.02	1.31	1.37
85	AA	766	G	N7-C5	-9.02	1.33	1.39
85	AA	1095	C	P-O5'	-9.02	1.50	1.59
85	AA	1829	C	C2'-C1'	-9.02	1.43	1.53
34	BA	1418	G	N9-C4	-9.01	1.30	1.38
35	BB	2	C	C1'-N1	-9.01	1.34	1.46
35	BB	679	G	N7-C5	-9.01	1.33	1.39
38	BE	106	C	O3'-P	-9.01	1.50	1.61
41	BH	106	G	O3'-P	-9.01	1.50	1.61
85	AA	527	A	C1'-N9	-9.01	1.34	1.46
85	AA	2022	A	C6-N6	-9.01	1.26	1.33
85	AA	2184	A	C3'-C2'	-9.01	1.42	1.52
34	BA	580	U	C2'-C1'	-9.01	1.43	1.53
34	BA	692	U	P-O5'	-9.01	1.50	1.59
34	BA	693	G	C2-N2	-9.01	1.25	1.34
35	BB	500	C	O3'-P	-9.01	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	601	U	O3'-P	-9.01	1.50	1.61
35	BB	1249	G	C5-C4	-9.01	1.32	1.38
35	BB	1428	C	O3'-P	-9.01	1.50	1.61
38	BE	40	C	P-O5'	-9.01	1.50	1.59
85	AA	1282	A	C3'-C2'	-9.01	1.42	1.52
86	AB	72	C	C2'-C1'	-9.01	1.43	1.53
34	BA	355	U	C2-N3	-9.01	1.31	1.37
37	BD	3	G	O3'-P	-9.01	1.50	1.61
38	BE	107	U	C4'-C3'	-9.01	1.43	1.53
85	AA	1535	C	O3'-P	-9.01	1.50	1.61
34	BA	1587	C	C2'-C1'	-9.01	1.43	1.53
40	BG	139	U	C3'-C2'	-9.01	1.42	1.52
34	BA	562	C	P-O5'	-9.00	1.50	1.59
34	BA	911	G	C1'-N9	-9.00	1.34	1.46
34	BA	1342	C	P-O5'	-9.00	1.50	1.59
34	BA	1433	U	C2'-C1'	-9.00	1.43	1.53
35	BB	32	C	N1-C6	-9.00	1.31	1.37
38	BE	25	U	O3'-P	-9.00	1.50	1.61
85	AA	396	U	P-O5'	-9.00	1.50	1.59
34	BA	8	G	C2'-C1'	-9.00	1.43	1.53
34	BA	1312	A	P-O5'	-9.00	1.50	1.59
34	BA	1443	U	O4'-C1'	-9.00	1.29	1.41
35	BB	77	A	C5-C4	-9.00	1.32	1.38
35	BB	1065	G	N9-C4	-9.00	1.30	1.38
37	BD	25	G	C2'-C1'	-9.00	1.43	1.53
37	BD	48	G	O3'-P	-9.00	1.50	1.61
85	AA	2002	A	N9-C4	9.00	1.43	1.37
85	AA	883	A	O3'-P	-9.00	1.50	1.61
85	AA	1183	C	O3'-P	-9.00	1.50	1.61
85	AA	1830	U	P-O5'	-9.00	1.50	1.59
85	AA	2172	A	P-O5'	-9.00	1.50	1.59
34	BA	747	G	C2'-C1'	-8.99	1.43	1.53
35	BB	763	U	P-O5'	-8.99	1.50	1.59
34	BA	272	A	N7-C5	-8.99	1.33	1.39
34	BA	346	A	N9-C4	-8.99	1.32	1.37
34	BA	784	C	C1'-N1	-8.99	1.34	1.46
34	BA	1576	C	O3'-P	-8.99	1.50	1.61
37	BD	75	G	C2-N2	-8.99	1.25	1.34
85	AA	485	A	N9-C4	-8.99	1.32	1.37
85	AA	2186	U	O3'-P	-8.99	1.50	1.61
34	BA	57	A	C2'-C1'	-8.99	1.43	1.53
34	BA	658	C	O3'-P	-8.99	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1704	G	C3'-C2'	-8.99	1.42	1.52
35	BB	452	A	N9-C4	-8.99	1.32	1.37
35	BB	1119	G	C3'-C2'	-8.99	1.42	1.52
38	BE	192	A	C1'-N9	-8.99	1.34	1.46
85	AA	686	U	O3'-P	-8.99	1.50	1.61
35	BB	1194	A	P-O5'	-8.99	1.50	1.59
34	BA	412	G	N9-C4	-8.99	1.30	1.38
41	BH	10	U	N3-C4	-8.99	1.30	1.38
85	AA	155	U	C2-N3	-8.99	1.31	1.37
34	BA	154	A	N9-C4	-8.98	1.32	1.37
34	BA	1281	U	C2-N3	-8.98	1.31	1.37
35	BB	1086	G	O3'-P	-8.98	1.50	1.61
35	BB	1149	A	N9-C4	-8.98	1.32	1.37
35	BB	1338	U	O3'-P	-8.98	1.50	1.61
35	BB	571	C	O3'-P	-8.98	1.50	1.61
38	BE	108	U	C2-N3	-8.98	1.31	1.37
34	BA	326	A	C2'-C1'	-8.98	1.43	1.53
35	BB	419	G	C1'-N9	-8.98	1.34	1.46
36	BC	101	U	C2-N3	-8.98	1.31	1.37
40	BG	142	A	N7-C5	-8.98	1.33	1.39
34	BA	78	U	O3'-P	-8.98	1.50	1.61
34	BA	290	G	C3'-C2'	-8.98	1.42	1.52
34	BA	1720	U	N3-C4	-8.98	1.30	1.38
34	BA	437	G	N7-C5	-8.98	1.33	1.39
34	BA	495	A	C2'-C1'	-8.98	1.43	1.53
34	BA	1180	A	P-O5'	-8.98	1.50	1.59
34	BA	1564	A	N7-C5	-8.98	1.33	1.39
35	BB	579	A	O3'-P	-8.98	1.50	1.61
85	AA	421	G	C6-N1	-8.98	1.33	1.39
35	BB	1094	A	P-O5'	-8.98	1.50	1.59
35	BB	1371	G	N9-C4	-8.98	1.30	1.38
85	AA	170	C	O3'-P	-8.98	1.50	1.61
85	AA	2151	U	C2-N3	-8.97	1.31	1.37
34	BA	441	A	C2'-C1'	-8.97	1.43	1.53
34	BA	1341	A	P-O5'	-8.97	1.50	1.59
34	BA	1579	G	C6-N1	-8.97	1.33	1.39
35	BB	433	C	P-O5'	-8.97	1.50	1.59
35	BB	1362	G	N7-C5	-8.97	1.33	1.39
41	BH	43	G	N9-C4	-8.97	1.30	1.38
34	BA	1808	A	C3'-C2'	-8.97	1.42	1.52
35	BB	423	G	C6-N1	-8.97	1.33	1.39
35	BB	615	A	C2'-C1'	-8.97	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	26	G	C1'-N9	-8.97	1.34	1.46
35	BB	1073	A	O3'-P	-8.97	1.50	1.61
36	BC	14	G	C2'-C1'	-8.97	1.43	1.53
37	BD	26	C	P-O5'	-8.97	1.50	1.59
85	AA	456	A	O3'-P	-8.97	1.50	1.61
34	BA	238	C	C4-C5	-8.97	1.35	1.43
36	BC	106	G	O3'-P	-8.97	1.50	1.61
36	BC	117	A	O3'-P	-8.97	1.50	1.61
41	BH	44	A	C2'-C1'	-8.97	1.43	1.53
85	AA	1963	G	O3'-P	-8.97	1.50	1.61
34	BA	341	U	C4'-C3'	-8.96	1.43	1.53
34	BA	1327	G	C2'-C1'	-8.97	1.43	1.53
85	AA	860	C	C2-N3	-8.97	1.28	1.35
34	BA	1496	G	N9-C4	-8.96	1.30	1.38
40	BG	137	G	C1'-N9	-8.96	1.34	1.46
85	AA	1651	C	C2'-C1'	-8.96	1.43	1.53
85	AA	2203	C	O3'-P	-8.96	1.50	1.61
85	AA	2251	U	C5'-C4'	8.96	1.62	1.51
34	BA	1405	A	C2'-C1'	-8.96	1.43	1.53
38	BE	167	U	C2-N3	-8.96	1.31	1.37
40	BG	5	G	P-O5'	-8.96	1.50	1.59
85	AA	128	U	O3'-P	-8.96	1.50	1.61
85	AA	268	A	P-O5'	-8.96	1.50	1.59
34	BA	11	U	C2'-C1'	-8.96	1.43	1.53
34	BA	742	C	N1-C2	-8.96	1.31	1.40
34	BA	1241	U	C2'-C1'	-8.96	1.43	1.53
35	BB	1047	C	O3'-P	-8.96	1.50	1.61
35	BB	1406	C	O3'-P	-8.96	1.50	1.61
41	BH	29	G	C5-C4	-8.96	1.32	1.38
41	BH	110	C	P-O5'	-8.96	1.50	1.59
85	AA	1756	C	P-O5'	-8.96	1.50	1.59
36	BC	96	A	C1'-N9	-8.96	1.34	1.46
85	AA	52	U	P-O5'	-8.96	1.50	1.59
34	BA	518	C	C2'-C1'	-8.96	1.43	1.53
35	BB	847	U	P-O5'	-8.96	1.50	1.59
35	BB	1542	C	C2'-C1'	-8.96	1.43	1.53
85	AA	707	U	C2-N3	-8.96	1.31	1.37
34	BA	1223	C	C2'-C1'	-8.95	1.43	1.53
40	BG	68	U	C4'-C3'	-8.96	1.43	1.53
85	AA	397	G	C5-C4	-8.95	1.32	1.38
35	BB	687	C	O3'-P	-8.95	1.50	1.61
35	BB	1387	C	C2-N3	-8.95	1.28	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	722	G	P-O5'	-8.95	1.50	1.59
34	BA	484	A	C5'-C4'	-8.95	1.40	1.51
34	BA	1211	G	C2-N2	-8.95	1.25	1.34
35	BB	551	C	O3'-P	-8.95	1.50	1.61
35	BB	490	G	C2'-C1'	-8.95	1.43	1.53
37	BD	74	A	C2'-C1'	-8.95	1.43	1.53
85	AA	402	G	N9-C4	-8.95	1.30	1.38
35	BB	117	A	O3'-P	-8.95	1.50	1.61
34	BA	102	G	C6-N1	-8.95	1.33	1.39
34	BA	205	G	P-O5'	-8.95	1.50	1.59
34	BA	322	U	N3-C4	-8.94	1.30	1.38
34	BA	1570	C	C4'-C3'	-8.95	1.43	1.53
34	BA	1657	A	N9-C4	-8.95	1.32	1.37
35	BB	787	A	C3'-C2'	-8.95	1.43	1.52
35	BB	977	G	N7-C5	-8.95	1.33	1.39
35	BB	1138	A	P-O5'	-8.95	1.50	1.59
38	BE	21	C	C4'-C3'	-8.95	1.43	1.53
85	AA	2224	U	P-O5'	-8.95	1.50	1.59
40	BG	82	U	O3'-P	-8.94	1.50	1.61
85	AA	299	A	O3'-P	-8.94	1.50	1.61
85	AA	757	A	O3'-P	-8.95	1.50	1.61
34	BA	59	A	N9-C4	-8.94	1.32	1.37
34	BA	994	G	P-O5'	-8.94	1.50	1.59
35	BB	130	G	C1'-N9	-8.94	1.34	1.46
37	BD	54	A	N7-C5	-8.94	1.33	1.39
40	BG	95	U	O4'-C1'	-8.94	1.30	1.41
34	BA	111	U	C2'-C1'	-8.94	1.43	1.53
34	BA	1107	A	C2'-C1'	-8.94	1.43	1.53
34	BA	1542	A	O3'-P	-8.94	1.50	1.61
38	BE	170	U	C4'-C3'	-8.94	1.43	1.53
85	AA	1471	G	N1-C2	-8.94	1.30	1.37
85	AA	1670	U	C2'-C1'	-8.94	1.43	1.53
35	BB	1314	G	C3'-C2'	-8.94	1.43	1.52
85	AA	1659	C	P-O5'	-8.94	1.50	1.59
86	AB	70	G	O3'-P	-8.94	1.50	1.61
34	BA	985	C	C3'-C2'	-8.94	1.43	1.52
34	BA	1597	G	C1'-N9	-8.94	1.34	1.46
36	BC	106	G	C2'-C1'	-8.94	1.43	1.53
40	BG	28	A	C5-C4	-8.94	1.32	1.38
34	BA	1641	G	N9-C4	-8.93	1.30	1.38
34	BA	1652	G	C2'-C1'	-8.93	1.43	1.53
40	BG	8	U	C1'-N1	-8.93	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	116	G	P-O5'	-8.93	1.50	1.59
35	BB	109	U	O3'-P	-8.93	1.50	1.61
34	BA	17	A	O3'-P	-8.93	1.50	1.61
34	BA	151	A	C1'-N9	8.93	1.62	1.48
34	BA	856	G	C4'-C3'	-8.93	1.43	1.53
34	BA	996	U	O3'-P	-8.93	1.50	1.61
34	BA	1176	C	C4'-C3'	-8.93	1.43	1.53
34	BA	1221	A	P-O5'	-8.93	1.50	1.59
34	BA	1250	C	O3'-P	-8.93	1.50	1.61
35	BB	569	G	C2'-C1'	-8.93	1.43	1.53
35	BB	669	A	N9-C4	-8.93	1.32	1.37
35	BB	1051	U	C3'-C2'	-8.93	1.43	1.52
35	BB	826	G	O3'-P	-8.93	1.50	1.61
38	BE	25	U	C2-N3	-8.93	1.31	1.37
34	BA	708	C	C2'-C1'	-8.92	1.43	1.53
34	BA	1590	G	O3'-P	-8.92	1.50	1.61
35	BB	662	G	C3'-C2'	-8.92	1.43	1.52
36	BC	6	G	O3'-P	-8.92	1.50	1.61
85	AA	860	C	O3'-P	-8.92	1.50	1.61
85	AA	1196	C	C2'-C1'	-8.92	1.43	1.53
34	BA	399	G	C5-C6	-8.92	1.33	1.42
35	BB	620	G	C3'-C2'	-8.92	1.43	1.52
37	BD	50	A	C1'-N9	-8.92	1.34	1.46
39	BF	4	A	P-O5'	-8.92	1.50	1.59
34	BA	875	G	O3'-P	-8.92	1.50	1.61
34	BA	844	U	P-O5'	-8.92	1.50	1.59
35	BB	373	C	O3'-P	-8.92	1.50	1.61
36	BC	18	G	O3'-P	-8.92	1.50	1.61
35	BB	566	A	P-O5'	-8.92	1.50	1.59
36	BC	65	G	P-O5'	-8.92	1.50	1.59
38	BE	6	A	C1'-N9	-8.92	1.34	1.46
40	BG	73	U	P-O5'	-8.92	1.50	1.59
85	AA	30	G	C2'-C1'	-8.92	1.43	1.53
85	AA	2008	G	C3'-C2'	-8.92	1.43	1.52
34	BA	134	U	P-O5'	-8.91	1.50	1.59
34	BA	1442	A	C1'-N9	-8.91	1.34	1.46
34	BA	1547	G	P-O5'	-8.91	1.50	1.59
35	BB	605	C	C4'-C3'	-8.91	1.43	1.53
35	BB	1103	A	C1'-N9	-8.91	1.34	1.46
85	AA	403	G	C2'-C1'	-8.91	1.43	1.53
35	BB	598	C	O3'-P	-8.91	1.50	1.61
36	BC	6	G	P-O5'	-8.91	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	182	G	P-O5'	-8.91	1.50	1.59
85	AA	999	A	C5'-C4'	8.91	1.62	1.51
34	BA	267	G	O3'-P	-8.91	1.50	1.61
36	BC	33	U	O3'-P	-8.91	1.50	1.61
40	BG	95	U	C2'-C1'	-8.91	1.43	1.53
85	AA	701	C	O3'-P	-8.91	1.50	1.61
85	AA	995	G	C1'-N9	-8.91	1.34	1.46
85	AA	1005	C	P-O5'	-8.91	1.50	1.59
85	AA	1483	A	P-O5'	-8.91	1.50	1.59
85	AA	1823	G	C3'-C2'	-8.91	1.43	1.52
34	BA	474	A	C2'-C1'	-8.91	1.43	1.53
35	BB	1321	G	C4'-C3'	-8.91	1.43	1.53
85	AA	448	G	C6-N1	-8.91	1.33	1.39
85	AA	1162	A	C2'-C1'	-8.91	1.43	1.53
85	AA	2041	G	C2-N2	-8.91	1.25	1.34
85	AA	2148	C	C3'-C2'	-8.91	1.43	1.52
34	BA	136	A	O3'-P	-8.90	1.50	1.61
34	BA	191	G	C2'-C1'	-8.90	1.43	1.53
34	BA	1292	A	N9-C4	-8.90	1.32	1.37
35	BB	994	A	C1'-N9	8.90	1.62	1.48
37	BD	112	U	P-O5'	-8.90	1.50	1.59
85	AA	454	G	N9-C8	-8.90	1.31	1.37
85	AA	1829	C	O3'-P	-8.90	1.50	1.61
85	AA	639	C	O3'-P	-8.90	1.50	1.61
85	AA	939	A	C8-N7	-8.90	1.25	1.31
34	BA	740	A	C1'-N9	-8.90	1.34	1.46
34	BA	149	G	P-O5'	-8.90	1.50	1.59
35	BB	799	A	C4'-C3'	-8.90	1.43	1.53
40	BG	111	C	P-O5'	-8.90	1.50	1.59
85	AA	589	A	N7-C5	-8.90	1.33	1.39
85	AA	1143	C	O3'-P	-8.90	1.50	1.61
34	BA	203	U	O3'-P	-8.90	1.50	1.61
34	BA	216	C	P-O5'	-8.90	1.50	1.59
34	BA	297	A	O4'-C1'	-8.90	1.30	1.41
34	BA	594	G	C2'-C1'	-8.90	1.43	1.53
85	AA	1524	A	N9-C4	-8.90	1.32	1.37
34	BA	451	A	O3'-P	-8.90	1.50	1.61
34	BA	1234	U	P-O5'	-8.90	1.50	1.59
35	BB	66	G	C4'-C3'	-8.90	1.43	1.53
35	BB	132	G	P-O5'	-8.90	1.50	1.59
35	BB	737	C	P-O5'	-8.90	1.50	1.59
41	BH	19	G	C1'-N9	-8.90	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	97	A	C1'-N9	-8.90	1.34	1.46
34	BA	688	G	P-O5'	-8.89	1.50	1.59
34	BA	856	G	C1'-N9	-8.89	1.34	1.46
34	BA	1538	G	N1-C2	-8.89	1.30	1.37
35	BB	1425	A	C2'-C1'	-8.89	1.43	1.53
35	BB	1440	A	C2'-C1'	-8.89	1.43	1.53
85	AA	2037	A	P-O5'	-8.89	1.50	1.59
85	AA	702	G	C8-N7	-8.89	1.25	1.30
85	AA	879	G	C1'-N9	-8.89	1.34	1.46
85	AA	1226	A	C2'-C1'	-8.89	1.43	1.53
34	BA	751	A	O3'-P	-8.89	1.50	1.61
34	BA	1293	A	C5-C4	-8.89	1.32	1.38
34	BA	1614	G	C1'-N9	-8.89	1.34	1.46
41	BH	14	C	O3'-P	-8.89	1.50	1.61
85	AA	1526	G	O3'-P	-8.89	1.50	1.61
37	BD	93	G	C6-N1	-8.89	1.33	1.39
85	AA	422	G	C5-C4	-8.89	1.32	1.38
34	BA	10	G	C2'-C1'	-8.89	1.43	1.53
34	BA	53	G	C5-C4	-8.89	1.32	1.38
35	BB	392	G	C6-N1	-8.89	1.33	1.39
35	BB	631	G	O3'-P	-8.89	1.50	1.61
35	BB	1220	A	C8-N7	-8.89	1.25	1.31
35	BB	1366	C	O3'-P	-8.89	1.50	1.61
37	BD	18	G	O3'-P	-8.89	1.50	1.61
85	AA	1107	A	C6-N6	-8.89	1.26	1.33
85	AA	887	A	C2'-C1'	-8.89	1.43	1.53
85	AA	2147	A	N9-C8	-8.89	1.30	1.37
34	BA	755	G	C2-N2	-8.88	1.25	1.34
35	BB	645	C	C3'-C2'	-8.89	1.43	1.52
85	AA	1502	A	O3'-P	-8.89	1.50	1.61
85	AA	1676	G	N3-C4	-8.89	1.29	1.35
35	BB	1282	G	O3'-P	-8.88	1.50	1.61
85	AA	437	G	C1'-N9	-8.88	1.34	1.46
85	AA	654	A	C6-N1	-8.88	1.29	1.35
34	BA	49	A	C3'-C2'	-8.88	1.43	1.52
34	BA	480	G	C4'-C3'	-8.88	1.43	1.53
34	BA	1289	C	C4'-C3'	-8.88	1.43	1.53
35	BB	1299	G	P-O5'	-8.88	1.50	1.59
40	BG	98	A	N9-C4	-8.88	1.32	1.37
85	AA	1966	C	P-O5'	-8.88	1.50	1.59
85	AA	2003	C	O3'-P	-8.88	1.50	1.61
85	AA	2044	A	C2'-C1'	-8.88	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	786	A	C3'-C2'	-8.88	1.43	1.52
34	BA	916	A	N3-C4	-8.88	1.29	1.34
35	BB	49	A	N9-C4	-8.88	1.32	1.37
40	BG	92	U	C2'-C1'	-8.88	1.43	1.53
85	AA	1217	U	O3'-P	-8.88	1.50	1.61
36	BC	25	C	O3'-P	-8.88	1.50	1.61
34	BA	1001	G	C1'-N9	-8.87	1.34	1.46
35	BB	1183	U	C3'-C2'	-8.87	1.43	1.52
35	BB	1225	A	O3'-P	-8.87	1.50	1.61
41	BH	38	G	O3'-P	-8.87	1.50	1.61
36	BC	17	U	O3'-P	-8.87	1.50	1.61
38	BE	139	U	C2-N3	-8.87	1.31	1.37
34	BA	954	U	O3'-P	-8.87	1.50	1.61
34	BA	1507	C	P-O5'	-8.87	1.50	1.59
35	BB	1388	A	C5-C4	-8.87	1.32	1.38
85	AA	173	A	C4'-C3'	-8.87	1.43	1.53
34	BA	1592	U	O3'-P	-8.87	1.50	1.61
85	AA	1281	G	C2'-C1'	-8.87	1.43	1.53
85	AA	2178	A	C2'-C1'	-8.87	1.43	1.53
34	BA	7	U	C3'-C2'	-8.87	1.43	1.52
34	BA	780	U	C1'-N1	8.87	1.62	1.48
34	BA	798	G	C6-N1	-8.87	1.33	1.39
34	BA	336	A	C2'-C1'	-8.86	1.43	1.53
35	BB	491	A	N7-C5	-8.86	1.33	1.39
34	BA	568	G	P-O5'	-8.86	1.50	1.59
35	BB	1422	G	O3'-P	-8.86	1.50	1.61
38	BE	105	A	N9-C4	-8.86	1.32	1.37
34	BA	483	A	P-O5'	-8.86	1.50	1.59
35	BB	430	A	C2'-C1'	-8.86	1.43	1.53
35	BB	446	U	C4'-O4'	-8.86	1.34	1.45
35	BB	634	A	C2'-C1'	-8.86	1.43	1.53
35	BB	1241	U	C3'-C2'	-8.86	1.43	1.52
37	BD	12	U	O3'-P	-8.86	1.50	1.61
38	BE	23	G	C4'-C3'	-8.86	1.43	1.53
85	AA	937	G	C2'-C1'	-8.86	1.43	1.53
85	AA	1204	A	O3'-P	-8.86	1.50	1.61
85	AA	1233	G	N9-C4	-8.86	1.30	1.38
85	AA	1801	U	P-O5'	-8.86	1.50	1.59
85	AA	2031	C	O3'-P	-8.86	1.50	1.61
35	BB	591	A	P-O5'	-8.86	1.50	1.59
85	AA	43	A	O3'-P	-8.86	1.50	1.61
85	AA	70	U	C4'-C3'	-8.86	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	162	A	C2'-C1'	-8.86	1.43	1.53
85	AA	415	G	P-O5'	-8.86	1.50	1.59
85	AA	897	A	N9-C4	-8.86	1.32	1.37
85	AA	1221	G	O3'-P	-8.86	1.50	1.61
34	BA	596	G	C5-C6	-8.86	1.33	1.42
35	BB	1114	A	C8-N7	-8.86	1.25	1.31
38	BE	176	G	C2'-C1'	-8.86	1.43	1.53
34	BA	557	U	P-OP1	8.85	1.64	1.49
37	BD	103	C	P-O5'	-8.85	1.50	1.59
34	BA	620	C	C2'-C1'	-8.85	1.43	1.53
35	BB	622	G	C8-N7	-8.85	1.25	1.30
85	AA	1955	U	C2-N3	-8.85	1.31	1.37
34	BA	923	C	C3'-C2'	-8.85	1.43	1.52
41	BH	31	A	O3'-P	-8.85	1.50	1.61
85	AA	336	C	O3'-P	-8.85	1.50	1.61
85	AA	1105	G	N9-C8	-8.85	1.31	1.37
85	AA	1822	G	O3'-P	-8.85	1.50	1.61
34	BA	1835	A	O3'-P	-8.85	1.50	1.61
35	BB	572	G	P-O5'	-8.85	1.50	1.59
85	AA	28	A	O3'-P	-8.85	1.50	1.61
85	AA	477	U	P-O5'	-8.85	1.50	1.59
34	BA	165	C	O3'-P	-8.85	1.50	1.61
34	BA	1007	G	O3'-P	-8.85	1.50	1.61
34	BA	1099	U	P-O5'	-8.85	1.50	1.59
35	BB	86	A	O3'-P	-8.85	1.50	1.61
35	BB	1083	C	O3'-P	-8.85	1.50	1.61
35	BB	775	U	P-O5'	-8.85	1.50	1.59
36	BC	31	A	C5-C4	-8.85	1.32	1.38
39	BF	72	A	P-O5'	-8.85	1.50	1.59
85	AA	662	U	O3'-P	-8.85	1.50	1.61
34	BA	126	G	C2'-C1'	-8.84	1.43	1.53
34	BA	266	G	C1'-N9	-8.84	1.34	1.46
34	BA	371	U	C2-N3	-8.84	1.31	1.37
35	BB	1341	U	C2'-C1'	-8.84	1.43	1.53
36	BC	96	A	N9-C4	-8.84	1.32	1.37
37	BD	81	C	O3'-P	-8.84	1.50	1.61
39	BF	21	C	C4'-C3'	-8.84	1.43	1.53
85	AA	366	A	C2'-C1'	-8.84	1.43	1.53
35	BB	1415	G	N3-C4	-8.84	1.29	1.35
36	BC	148	C	C2-N3	-8.84	1.28	1.35
34	BA	1741	G	C2'-C1'	-8.84	1.43	1.53
36	BC	74	U	O3'-P	-8.84	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	399	A	O3'-P	-8.84	1.50	1.61
85	AA	929	G	O3'-P	-8.84	1.50	1.61
34	BA	1555	G	C2-N2	-8.84	1.25	1.34
34	BA	1838	U	O3'-P	-8.84	1.50	1.61
35	BB	504	C	P-O5'	-8.84	1.50	1.59
35	BB	1382	U	O3'-P	-8.84	1.50	1.61
37	BD	57	C	O3'-P	-8.84	1.50	1.61
34	BA	1108	U	O3'-P	-8.84	1.50	1.61
34	BA	1652	G	N9-C8	-8.84	1.31	1.37
35	BB	613	C	O3'-P	-8.84	1.50	1.61
35	BB	981	A	P-O5'	-8.84	1.50	1.59
38	BE	144	A	P-O5'	-8.84	1.50	1.59
85	AA	1503	G	O3'-P	-8.84	1.50	1.61
34	BA	660	C	C3'-C2'	-8.83	1.43	1.52
36	BC	68	A	N7-C5	-8.83	1.33	1.39
85	AA	305	A	N9-C4	-8.83	1.32	1.37
85	AA	887	A	P-O5'	-8.83	1.50	1.59
85	AA	1573	A	N9-C4	-8.83	1.32	1.37
85	AA	1637	C	P-O5'	-8.83	1.50	1.59
85	AA	2099	C	O3'-P	-8.83	1.50	1.61
85	AA	2218	G	P-O5'	-8.83	1.50	1.59
35	BB	1383	C	C2'-C1'	-8.83	1.43	1.53
34	BA	933	U	C1'-N1	-8.83	1.34	1.46
85	AA	1483	A	C5-C4	-8.83	1.32	1.38
34	BA	493	G	C2'-C1'	-8.83	1.43	1.53
34	BA	1508	C	P-O5'	-8.83	1.50	1.59
34	BA	1797	A	P-O5'	-8.83	1.50	1.59
35	BB	483	C	N1-C6	-8.83	1.31	1.37
35	BB	1334	C	C4'-C3'	-8.83	1.43	1.53
36	BC	53	A	C2'-C1'	-8.83	1.43	1.53
36	BC	55	U	N1-C2	-8.83	1.30	1.38
85	AA	391	G	O3'-P	-8.83	1.50	1.61
85	AA	2148	C	C2'-C1'	-8.83	1.43	1.53
34	BA	409	A	C2'-C1'	-8.82	1.43	1.53
34	BA	672	G	C2'-C1'	-8.82	1.43	1.53
41	BH	24	U	C2'-C1'	-8.82	1.43	1.53
35	BB	1296	A	N9-C4	-8.82	1.32	1.37
38	BE	6	A	C5-C4	-8.82	1.32	1.38
85	AA	1144	G	C6-N1	-8.82	1.33	1.39
85	AA	1206	A	C3'-C2'	-8.82	1.43	1.52
34	BA	347	A	N9-C4	-8.82	1.32	1.37
34	BA	477	C	P-O5'	-8.82	1.50	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	613	A	N9-C4	-8.82	1.32	1.37
34	BA	1807	G	C2-N2	-8.82	1.25	1.34
35	BB	131	A	N9-C4	-8.82	1.32	1.37
35	BB	516	G	O3'-P	-8.82	1.50	1.61
35	BB	1062	G	N9-C4	-8.82	1.30	1.38
34	BA	516	U	C2'-C1'	-8.82	1.43	1.53
40	BG	142	A	C2'-C1'	-8.82	1.43	1.53
34	BA	849	G	P-O5'	-8.82	1.50	1.59
34	BA	1686	G	O3'-P	-8.82	1.50	1.61
85	AA	410	A	N9-C4	-8.82	1.32	1.37
85	AA	1534	A	O3'-P	-8.82	1.50	1.61
85	AA	2240	G	N9-C4	-8.82	1.30	1.38
35	BB	1425	A	O3'-P	-8.82	1.50	1.61
34	BA	433	G	C5'-C4'	8.81	1.61	1.51
34	BA	333	A	C1'-N9	-8.81	1.34	1.46
35	BB	565	U	O3'-P	-8.81	1.50	1.61
35	BB	1264	U	C2-N3	-8.81	1.31	1.37
40	BG	130	G	C5-C4	-8.81	1.32	1.38
41	BH	29	G	N7-C5	-8.81	1.33	1.39
85	AA	1928	A	N9-C4	-8.81	1.32	1.37
85	AA	622	G	C1'-N9	-8.81	1.34	1.46
85	AA	1702	G	P-O5'	-8.81	1.50	1.59
85	AA	2180	C	C2'-C1'	-8.81	1.43	1.53
34	BA	1039	G	C1'-N9	-8.81	1.34	1.46
35	BB	999	G	C1'-N9	-8.81	1.34	1.46
35	BB	1390	U	O3'-P	-8.81	1.50	1.61
34	BA	1437	G	C2'-C1'	-8.81	1.43	1.53
34	BA	1613	G	N9-C4	-8.81	1.30	1.38
85	AA	876	U	P-O5'	-8.81	1.50	1.59
35	BB	1053	G	C1'-N9	-8.81	1.34	1.46
38	BE	142	A	P-O5'	-8.81	1.50	1.59
85	AA	620	U	C3'-C2'	-8.81	1.43	1.52
85	AA	1460	G	C5-C4	-8.81	1.32	1.38
85	AA	242	G	C2-N2	-8.81	1.25	1.34
85	AA	1911	A	C2'-C1'	-8.81	1.43	1.53
34	BA	236	A	P-O5'	-8.81	1.50	1.59
34	BA	1269	C	C2-N3	-8.81	1.28	1.35
34	BA	1556	A	C5-C4	-8.81	1.32	1.38
35	BB	7	C	O3'-P	-8.81	1.50	1.61
40	BG	10	U	C4'-O4'	-8.81	1.34	1.45
37	BD	29	C	P-O5'	-8.81	1.50	1.59
34	BA	765	U	C1'-N1	8.80	1.61	1.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1575	U	P-O5'	-8.81	1.50	1.59
85	AA	7	G	N9-C4	-8.81	1.30	1.38
34	BA	465	A	O3'-P	-8.80	1.50	1.61
35	BB	778	A	C5-C4	-8.80	1.32	1.38
37	BD	98	G	N1-C2	-8.80	1.30	1.37
37	BD	108	G	C3'-C2'	-8.80	1.43	1.52
35	BB	513	G	P-O5'	-8.80	1.50	1.59
35	BB	1244	U	O3'-P	-8.80	1.50	1.61
35	BB	1305	A	N7-C5	-8.80	1.33	1.39
37	BD	80	G	O3'-P	-8.80	1.50	1.61
40	BG	89	A	C5-C4	-8.80	1.32	1.38
85	AA	447	C	C2-N3	-8.80	1.28	1.35
85	AA	1735	U	C2'-C1'	-8.80	1.43	1.53
85	AA	597	A	C1'-N9	-8.80	1.34	1.46
35	BB	1326	U	P-O5'	-8.80	1.50	1.59
85	AA	471	U	C2-N3	-8.80	1.31	1.37
85	AA	1234	G	O3'-P	-8.80	1.50	1.61
34	BA	388	A	O3'-P	-8.79	1.50	1.61
34	BA	1555	G	N9-C4	-8.79	1.30	1.38
35	BB	776	U	O3'-P	-8.80	1.50	1.61
34	BA	1653	G	C1'-N9	-8.79	1.34	1.46
35	BB	1303	A	O3'-P	-8.79	1.50	1.61
38	BE	127	G	C3'-C2'	-8.79	1.43	1.52
40	BG	37	G	O3'-P	-8.79	1.50	1.61
85	AA	165	C	O3'-P	-8.79	1.50	1.61
85	AA	1181	U	O3'-P	-8.79	1.50	1.61
85	AA	1793	A	C4'-C3'	-8.79	1.43	1.53
85	AA	2203	C	P-O5'	-8.79	1.50	1.59
35	BB	407	A	P-O5'	-8.79	1.50	1.59
35	BB	470	C	O3'-P	-8.79	1.50	1.61
35	BB	831	C	C2'-C1'	-8.79	1.43	1.53
40	BG	178	G	P-O5'	-8.79	1.50	1.59
35	BB	869	G	C2'-C1'	-8.79	1.43	1.53
34	BA	1355	G	N7-C5	-8.79	1.33	1.39
35	BB	135	C	P-O5'	-8.79	1.50	1.59
35	BB	1251	G	P-O5'	-8.79	1.50	1.59
35	BB	1539	C	O3'-P	-8.79	1.50	1.61
34	BA	480	G	C8-N7	-8.79	1.25	1.30
35	BB	971	A	O3'-P	-8.79	1.50	1.61
35	BB	1084	A	N9-C4	-8.79	1.32	1.37
38	BE	145	A	C1'-N9	-8.79	1.34	1.46
40	BG	52	A	N9-C4	-8.79	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	501	A	P-O5'	-8.79	1.50	1.59
34	BA	1590	G	C1'-N9	-8.79	1.34	1.46
37	BD	95	G	N7-C5	-8.79	1.33	1.39
34	BA	1249	G	N1-C2	-8.78	1.30	1.37
35	BB	482	A	C2'-C1'	-8.78	1.43	1.53
85	AA	1538	C	C2'-C1'	-8.78	1.43	1.53
34	BA	1418	G	N7-C5	-8.78	1.33	1.39
35	BB	1095	G	P-O5'	-8.78	1.50	1.59
85	AA	1490	A	C1'-N9	-8.78	1.34	1.46
34	BA	1095	G	O3'-P	-8.78	1.50	1.61
34	BA	500	C	C3'-C2'	-8.78	1.43	1.52
34	BA	794	G	O3'-P	-8.78	1.50	1.61
34	BA	1814	U	N3-C4	-8.78	1.30	1.38
35	BB	76	C	P-O5'	-8.78	1.50	1.59
34	BA	52	G	C1'-N9	-8.78	1.34	1.46
34	BA	901	C	C2'-C1'	-8.78	1.43	1.53
34	BA	429	G	N3-C4	-8.78	1.29	1.35
85	AA	454	G	P-O5'	-8.78	1.50	1.59
34	BA	1023	G	O3'-P	-8.77	1.50	1.61
35	BB	583	G	C2'-C1'	-8.77	1.43	1.53
35	BB	768	A	C2'-C1'	-8.77	1.43	1.53
35	BB	1110	G	O3'-P	-8.77	1.50	1.61
34	BA	760	G	C2'-C1'	-8.77	1.43	1.53
35	BB	7	C	C2'-C1'	-8.77	1.43	1.53
35	BB	487	A	C1'-N9	-8.77	1.34	1.46
36	BC	10	C	O3'-P	-8.77	1.50	1.61
37	BD	74	A	C4'-C3'	-8.77	1.43	1.53
85	AA	456	A	N9-C4	-8.77	1.32	1.37
85	AA	2147	A	C2'-C1'	-8.77	1.43	1.53
34	BA	1721	U	N3-C4	-8.77	1.30	1.38
35	BB	593	A	O3'-P	-8.77	1.50	1.61
34	BA	482	C	C4'-O4'	-8.77	1.34	1.45
34	BA	524	G	C2-N2	-8.77	1.25	1.34
35	BB	1178	A	C2'-C1'	-8.77	1.43	1.53
41	BH	42	U	P-O5'	-8.77	1.50	1.59
85	AA	2217	A	C2'-C1'	-8.77	1.43	1.53
34	BA	981	A	N9-C4	-8.77	1.32	1.37
35	BB	115	A	P-O5'	-8.77	1.50	1.59
36	BC	42	G	C1'-N9	-8.77	1.34	1.46
85	AA	627	A	P-O5'	-8.77	1.50	1.59
35	BB	786	A	P-O5'	-8.77	1.50	1.59
40	BG	56	G	O3'-P	-8.77	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	339	G	P-O5'	-8.76	1.50	1.59
34	BA	413	A	N9-C4	-8.76	1.32	1.37
34	BA	1278	A	C3'-C2'	-8.76	1.43	1.52
35	BB	1077	C	P-O5'	-8.76	1.50	1.59
85	AA	601	A	C2'-C1'	-8.76	1.43	1.53
85	AA	688	C	C3'-C2'	-8.76	1.43	1.52
35	BB	1444	U	O3'-P	-8.76	1.50	1.61
40	BG	10	U	C1'-N1	-8.76	1.34	1.46
85	AA	36	U	O3'-P	-8.76	1.50	1.61
85	AA	516	G	O3'-P	-8.76	1.50	1.61
85	AA	1801	U	O3'-P	-8.76	1.50	1.61
34	BA	557	U	O4'-C1'	8.76	1.53	1.41
34	BA	1803	A	O3'-P	-8.76	1.50	1.61
35	BB	1051	U	P-O5'	-8.76	1.50	1.59
35	BB	1053	G	P-O5'	-8.76	1.50	1.59
85	AA	374	C	C2'-C1'	-8.76	1.43	1.53
85	AA	1107	A	C4'-C3'	-8.76	1.43	1.53
35	BB	503	G	C2'-C1'	-8.76	1.43	1.53
35	BB	1189	C	O3'-P	-8.76	1.50	1.61
35	BB	1292	G	C4'-C3'	-8.76	1.43	1.53
85	AA	927	A	P-O5'	-8.76	1.50	1.59
40	BG	65	C	P-O5'	-8.76	1.50	1.59
85	AA	1496	U	P-O5'	-8.76	1.50	1.59
85	AA	2190	U	P-O5'	-8.76	1.50	1.59
34	BA	825	G	C5-C4	-8.76	1.32	1.38
37	BD	64	A	O3'-P	-8.76	1.50	1.61
34	BA	938	C	C2'-C1'	-8.76	1.43	1.53
35	BB	127	U	C2'-C1'	-8.76	1.43	1.53
35	BB	568	A	C5-C4	-8.76	1.32	1.38
35	BB	1103	A	N7-C5	-8.76	1.33	1.39
35	BB	1399	A	C8-N7	-8.76	1.25	1.31
36	BC	76	C	P-O5'	-8.76	1.50	1.59
34	BA	515	U	C3'-C2'	-8.75	1.43	1.52
35	BB	465	C	C4-N4	-8.75	1.26	1.33
35	BB	1448	U	P-O5'	-8.75	1.50	1.59
37	BD	67	C	C3'-C2'	-8.75	1.43	1.52
40	BG	3	G	C2-N2	-8.75	1.25	1.34
34	BA	701	G	P-O5'	-8.75	1.50	1.59
34	BA	1176	C	O3'-P	-8.75	1.50	1.61
85	AA	1656	C	O3'-P	-8.75	1.50	1.61
85	AA	2146	G	C4'-O4'	-8.75	1.34	1.45
34	BA	219	U	C2'-C1'	-8.75	1.43	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1142	C	O3'-P	-8.75	1.50	1.61
34	BA	504	A	C1'-N9	-8.75	1.34	1.46
34	BA	864	G	C8-N7	8.75	1.36	1.30
34	BA	1092	U	C3'-C2'	-8.75	1.43	1.52
34	BA	1208	U	C1'-N1	-8.75	1.34	1.46
37	BD	41	G	C4'-O4'	-8.75	1.34	1.45
38	BE	144	A	C3'-C2'	-8.75	1.43	1.52
85	AA	1244	A	P-O5'	-8.75	1.51	1.59
34	BA	20	A	C8-N7	-8.75	1.25	1.31
34	BA	240	C	C2'-C1'	-8.75	1.43	1.53
34	BA	579	U	C2-N3	-8.75	1.31	1.37
85	AA	877	G	C2'-C1'	-8.75	1.43	1.53
34	BA	500	C	P-O5'	-8.75	1.51	1.59
35	BB	451	A	P-O5'	-8.75	1.51	1.59
35	BB	646	U	P-O5'	-8.75	1.51	1.59
34	BA	43	U	O3'-P	-8.74	1.50	1.61
34	BA	269	G	C2'-C1'	-8.74	1.43	1.53
34	BA	503	C	C2-N3	-8.74	1.28	1.35
34	BA	700	G	C3'-C2'	-8.74	1.43	1.52
34	BA	1064	A	C5-C4	-8.74	1.32	1.38
34	BA	1671	A	O3'-P	-8.74	1.50	1.61
35	BB	670	G	O3'-P	-8.74	1.50	1.61
35	BB	825	U	O3'-P	-8.74	1.50	1.61
35	BB	1144	A	C2'-C1'	-8.74	1.43	1.53
36	BC	95	A	C1'-N9	-8.74	1.34	1.46
85	AA	364	C	C3'-C2'	-8.74	1.43	1.52
85	AA	1486	G	C6-N1	-8.74	1.33	1.39
36	BC	53	A	P-O5'	-8.74	1.51	1.59
85	AA	1858	G	C3'-C2'	-8.74	1.43	1.52
85	AA	2126	U	C2-N3	-8.74	1.31	1.37
40	BG	113	G	C6-N1	-8.74	1.33	1.39
34	BA	1009	G	O3'-P	-8.74	1.50	1.61
34	BA	1745	G	P-O5'	-8.74	1.51	1.59
34	BA	306	G	C3'-C2'	-8.74	1.43	1.52
34	BA	457	A	C3'-C2'	-8.74	1.43	1.52
34	BA	1222	C	C4'-C3'	-8.74	1.43	1.53
35	BB	609	G	P-O5'	-8.74	1.51	1.59
35	BB	467	G	P-O5'	-8.74	1.51	1.59
40	BG	8	U	C3'-C2'	-8.74	1.43	1.52
85	AA	1268	C	O3'-P	-8.74	1.50	1.61
85	AA	1717	G	C4'-C3'	8.74	1.62	1.53
85	AA	1984	A	O3'-P	-8.74	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	441	A	N9-C4	-8.73	1.32	1.37
34	BA	1483	U	C2-N3	-8.73	1.31	1.37
85	AA	397	G	N9-C8	-8.73	1.31	1.37
35	BB	496	C	O3'-P	-8.73	1.50	1.61
35	BB	1376	G	O3'-P	-8.73	1.50	1.61
85	AA	730	G	N9-C4	8.73	1.45	1.38
85	AA	2147	A	N3-C4	-8.73	1.29	1.34
38	BE	75	C	P-O5'	-8.73	1.51	1.59
85	AA	2081	A	P-O5'	-8.73	1.51	1.59
34	BA	664	C	P-O5'	-8.73	1.51	1.59
35	BB	783	U	P-O5'	-8.73	1.51	1.59
36	BC	89	U	P-O5'	-8.73	1.51	1.59
37	BD	75	G	O3'-P	-8.73	1.50	1.61
34	BA	94	G	N1-C2	-8.73	1.30	1.37
34	BA	542	A	P-O5'	-8.73	1.51	1.59
40	BG	105	A	C5-C4	-8.73	1.32	1.38
40	BG	137	G	P-O5'	-8.73	1.51	1.59
85	AA	47	A	N9-C4	8.73	1.43	1.37
85	AA	1178	A	N9-C4	-8.73	1.32	1.37
34	BA	1107	A	C3'-C2'	-8.73	1.43	1.52
34	BA	1800	G	C2'-C1'	-8.73	1.43	1.53
35	BB	469	G	O3'-P	-8.73	1.50	1.61
35	BB	1026	G	C5'-C4'	8.73	1.61	1.51
40	BG	108	G	C1'-N9	-8.73	1.34	1.46
85	AA	1283	C	O3'-P	-8.73	1.50	1.61
85	AA	474	C	C2'-C1'	-8.73	1.43	1.53
85	AA	1936	C	P-O5'	-8.73	1.51	1.59
85	AA	1678	U	C2-N3	-8.73	1.31	1.37
35	BB	1359	G	N9-C4	-8.72	1.30	1.38
34	BA	517	A	N9-C4	-8.72	1.32	1.37
35	BB	1078	U	O3'-P	-8.72	1.50	1.61
38	BE	97	G	P-O5'	-8.72	1.51	1.59
85	AA	2060	G	O3'-P	-8.72	1.50	1.61
34	BA	1841	A	O3'-P	-8.72	1.50	1.61
35	BB	1090	A	N9-C4	-8.72	1.32	1.37
38	BE	23	G	N7-C5	-8.72	1.34	1.39
34	BA	459	U	C3'-C2'	-8.72	1.43	1.52
85	AA	454	G	C1'-N9	-8.72	1.34	1.46
85	AA	1366	A	N9-C4	-8.72	1.32	1.37
35	BB	363	A	N7-C5	-8.72	1.34	1.39
85	AA	57	G	C6-N1	-8.72	1.33	1.39
85	AA	2185	U	C2-N3	-8.72	1.31	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	538	G	C6-N1	-8.71	1.33	1.39
34	BA	1034	U	O3'-P	-8.72	1.50	1.61
34	BA	1404	A	P-O5'	-8.72	1.51	1.59
41	BH	16	A	N3-C4	-8.71	1.29	1.34
85	AA	1464	G	N9-C8	-8.72	1.31	1.37
85	AA	611	G	P-O5'	-8.71	1.51	1.59
85	AA	764	U	C2-N3	-8.71	1.31	1.37
85	AA	1206	A	N9-C4	-8.71	1.32	1.37
85	AA	1476	C	O3'-P	-8.71	1.50	1.61
85	AA	1923	A	C2'-C1'	-8.71	1.43	1.53
34	BA	248	G	N7-C5	-8.71	1.34	1.39
39	BF	46	G	P-O5'	-8.71	1.51	1.59
34	BA	446	U	O3'-P	-8.71	1.50	1.61
34	BA	739	A	N3-C4	-8.71	1.29	1.34
34	BA	796	G	P-O5'	-8.71	1.51	1.59
38	BE	140	G	O3'-P	-8.71	1.50	1.61
40	BG	52	A	P-O5'	-8.71	1.51	1.59
35	BB	1067	G	C2'-C1'	-8.71	1.43	1.53
36	BC	31	A	C2'-C1'	-8.71	1.43	1.53
85	AA	741	G	C6-N1	-8.71	1.33	1.39
85	AA	1682	U	C2'-C1'	-8.71	1.43	1.53
85	AA	2146	G	N9-C4	-8.71	1.30	1.38
85	AA	1759	U	O3'-P	-8.71	1.50	1.61
34	BA	687	G	C6-N1	-8.71	1.33	1.39
35	BB	131	A	C5-C4	-8.71	1.32	1.38
35	BB	1194	A	O3'-P	-8.71	1.50	1.61
85	AA	1179	A	P-O5'	-8.71	1.51	1.59
35	BB	436	G	N9-C4	-8.71	1.30	1.38
41	BH	41	A	C3'-C2'	-8.71	1.43	1.52
34	BA	1431	G	C6-N1	-8.70	1.33	1.39
35	BB	106	A	P-O5'	-8.71	1.51	1.59
35	BB	441	G	C1'-N9	-8.71	1.34	1.46
35	BB	812	G	P-O5'	-8.70	1.51	1.59
35	BB	1057	G	O3'-P	-8.71	1.50	1.61
36	BC	54	G	N3-C4	-8.71	1.29	1.35
40	BG	47	G	P-O5'	-8.71	1.51	1.59
85	AA	1709	U	O3'-P	-8.70	1.50	1.61
34	BA	1696	G	C5-C4	-8.70	1.32	1.38
85	AA	514	U	O3'-P	-8.70	1.50	1.61
34	BA	694	G	C6-N1	-8.70	1.33	1.39
34	BA	1177	C	O3'-P	-8.70	1.50	1.61
34	BA	1285	G	C6-N1	-8.70	1.33	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1144	A	O3'-P	-8.70	1.50	1.61
36	BC	18	G	C6-N1	-8.70	1.33	1.39
34	BA	104	A	O3'-P	-8.70	1.50	1.61
34	BA	807	U	O3'-P	-8.70	1.50	1.61
34	BA	881	C	C3'-C2'	-8.70	1.43	1.52
34	BA	1037	C	C2'-C1'	-8.70	1.43	1.53
35	BB	567	G	C6-N1	-8.70	1.33	1.39
85	AA	1140	G	C4'-C3'	-8.70	1.43	1.53
34	BA	971	G	P-O5'	-8.69	1.51	1.59
34	BA	1344	G	P-O5'	-8.69	1.51	1.59
34	BA	1716	A	C5'-C4'	8.69	1.61	1.51
35	BB	126	C	C3'-C2'	-8.69	1.43	1.52
35	BB	546	A	N9-C4	-8.69	1.32	1.37
35	BB	622	G	C2'-C1'	-8.69	1.43	1.53
35	BB	1134	G	O3'-P	-8.69	1.50	1.61
35	BB	1427	A	C5-C4	-8.69	1.32	1.38
38	BE	27	A	O3'-P	-8.69	1.50	1.61
41	BH	47	G	C6-N1	-8.69	1.33	1.39
85	AA	642	G	P-O5'	-8.69	1.51	1.59
85	AA	704	A	C1'-N9	-8.69	1.34	1.46
85	AA	2081	A	N7-C5	-8.69	1.34	1.39
34	BA	961	C	C3'-C2'	-8.69	1.43	1.52
34	BA	1409	A	P-O5'	-8.69	1.51	1.59
35	BB	100	A	O3'-P	-8.69	1.50	1.61
85	AA	1477	A	O3'-P	-8.69	1.50	1.61
35	BB	1119	G	P-O5'	-8.69	1.51	1.59
41	BH	107	A	N7-C5	-8.69	1.34	1.39
85	AA	2141	G	C3'-C2'	-8.69	1.43	1.52
34	BA	574	U	N1-C2	8.69	1.46	1.38
34	BA	1252	G	C6-N1	-8.69	1.33	1.39
35	BB	1122	C	O3'-P	-8.69	1.50	1.61
34	BA	188	C	P-O5'	-8.68	1.51	1.59
34	BA	680	C	C4'-C3'	-8.68	1.43	1.53
85	AA	983	A	P-O5'	-8.68	1.51	1.59
35	BB	1210	U	P-O5'	-8.68	1.51	1.59
40	BG	16	G	O3'-P	-8.68	1.50	1.61
41	BH	58	C	P-O5'	-8.68	1.51	1.59
85	AA	2110	U	O3'-P	-8.68	1.50	1.61
34	BA	5	C	C2-N3	-8.68	1.28	1.35
34	BA	1105	A	N9-C4	-8.68	1.32	1.37
34	BA	1279	U	P-O5'	-8.68	1.51	1.59
34	BA	1535	G	C1'-N9	-8.68	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1149	A	C3'-C2'	-8.68	1.43	1.52
34	BA	1192	A	C2'-C1'	-8.68	1.43	1.53
34	BA	1332	U	P-O5'	-8.68	1.51	1.59
34	BA	422	C	O3'-P	-8.68	1.50	1.61
34	BA	1537	G	P-O5'	-8.68	1.51	1.59
35	BB	694	C	O3'-P	-8.68	1.50	1.61
35	BB	1385	C	C3'-C2'	-8.68	1.43	1.52
38	BE	64	A	O3'-P	-8.68	1.50	1.61
85	AA	1851	A	C1'-N9	-8.68	1.34	1.46
85	AA	1904	C	P-O5'	-8.68	1.51	1.59
34	BA	92	G	C2'-C1'	-8.67	1.43	1.53
34	BA	730	C	C2'-C1'	-8.67	1.43	1.53
34	BA	1657	A	C1'-N9	-8.67	1.34	1.46
35	BB	1062	G	O3'-P	-8.67	1.50	1.61
37	BD	101	A	P-O5'	-8.67	1.51	1.59
37	BD	111	U	C2-N3	-8.67	1.31	1.37
38	BE	182	U	C4'-C3'	-8.67	1.43	1.53
34	BA	296	G	C5'-C4'	8.67	1.61	1.51
40	BG	97	G	O3'-P	-8.67	1.50	1.61
40	BG	120	U	O3'-P	-8.67	1.50	1.61
35	BB	1221	G	N9-C4	-8.67	1.31	1.38
85	AA	352	G	C2'-C1'	-8.67	1.43	1.53
85	AA	1100	U	P-O5'	-8.67	1.51	1.59
35	BB	1286	G	N7-C5	-8.67	1.34	1.39
85	AA	372	U	N3-C4	-8.67	1.30	1.38
34	BA	792	A	C8-N7	-8.67	1.25	1.31
35	BB	33	A	C5-C4	-8.67	1.32	1.38
35	BB	648	G	N1-C2	-8.67	1.30	1.37
85	AA	2137	A	P-O5'	-8.67	1.51	1.59
34	BA	1289	C	P-O5'	-8.66	1.51	1.59
34	BA	1290	A	P-O5'	-8.66	1.51	1.59
35	BB	51	U	C2'-C1'	-8.66	1.43	1.53
37	BD	110	G	C5-C4	-8.66	1.32	1.38
85	AA	419	A	C1'-N9	-8.66	1.34	1.46
85	AA	2121	G	N7-C5	-8.66	1.34	1.39
34	BA	112	C	C3'-C2'	-8.66	1.43	1.52
34	BA	1163	G	C2'-C1'	-8.66	1.43	1.53
34	BA	1477	C	C4'-C3'	-8.66	1.43	1.53
34	BA	1823	A	N9-C4	-8.66	1.32	1.37
36	BC	26	U	C4'-O4'	-8.66	1.34	1.45
85	AA	832	U	P-O5'	-8.66	1.51	1.59
85	AA	1678	U	O3'-P	-8.66	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2145	G	O3'-P	-8.66	1.50	1.61
36	BC	30	U	O3'-P	-8.66	1.50	1.61
38	BE	170	U	C2-N3	-8.66	1.31	1.37
85	AA	705	G	O3'-P	-8.66	1.50	1.61
85	AA	1590	A	P-O5'	-8.66	1.51	1.59
34	BA	106	U	C3'-C2'	-8.66	1.43	1.52
34	BA	1045	C	P-O5'	-8.66	1.51	1.59
35	BB	1076	U	O3'-P	-8.66	1.50	1.61
40	BG	87	G	P-O5'	-8.66	1.51	1.59
85	AA	1877	G	O3'-P	-8.66	1.50	1.61
35	BB	478	G	P-O5'	-8.66	1.51	1.59
35	BB	1136	G	C6-N1	-8.66	1.33	1.39
85	AA	388	G	C2'-C1'	-8.66	1.43	1.53
85	AA	447	C	C3'-C2'	-8.66	1.43	1.52
85	AA	1466	U	C1'-N1	-8.66	1.34	1.46
34	BA	13	U	N1-C2	-8.65	1.30	1.38
34	BA	541	C	P-O5'	-8.65	1.51	1.59
34	BA	567	U	C2'-C1'	-8.65	1.43	1.53
35	BB	971	A	C1'-N9	-8.65	1.34	1.46
36	BC	156	A	C4'-C3'	-8.65	1.43	1.53
37	BD	3	G	C1'-N9	-8.65	1.34	1.46
40	BG	33	G	O3'-P	-8.65	1.50	1.61
85	AA	1250	A	N9-C4	-8.65	1.32	1.37
85	AA	1655	G	N9-C4	-8.65	1.31	1.38
35	BB	1344	U	C2-N3	-8.65	1.31	1.37
36	BC	147	G	O3'-P	-8.65	1.50	1.61
38	BE	3	G	N9-C4	-8.65	1.31	1.38
34	BA	882	G	O3'-P	-8.65	1.50	1.61
34	BA	1544	G	N9-C4	-8.65	1.31	1.38
35	BB	697	G	O3'-P	-8.65	1.50	1.61
85	AA	71	G	P-O5'	-8.65	1.51	1.59
85	AA	917	A	C1'-N9	-8.65	1.34	1.46
34	BA	1458	A	C5'-C4'	8.65	1.61	1.51
35	BB	586	U	C2-N3	-8.65	1.31	1.37
35	BB	4	C	P-O5'	-8.65	1.51	1.59
36	BC	52	A	C4'-C3'	-8.65	1.43	1.53
40	BG	16	G	N9-C4	8.65	1.44	1.38
85	AA	5	U	C2-N3	-8.65	1.31	1.37
85	AA	1228	A	C3'-C2'	-8.65	1.43	1.52
34	BA	370	U	C3'-C2'	-8.64	1.43	1.52
34	BA	939	C	O3'-P	-8.64	1.50	1.61
85	AA	2013	A	N9-C4	-8.64	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1430	C	C2'-C1'	-8.64	1.43	1.53
34	BA	917	C	C2-N3	-8.64	1.28	1.35
35	BB	1045	G	C5-C4	-8.64	1.32	1.38
38	BE	183	C	C5'-C4'	8.64	1.61	1.51
85	AA	629	A	C5-C4	-8.64	1.32	1.38
34	BA	818	G	C3'-C2'	-8.64	1.43	1.52
34	BA	896	U	P-O5'	-8.64	1.51	1.59
34	BA	942	G	O3'-P	-8.64	1.50	1.61
34	BA	1107	A	O3'-P	-8.64	1.50	1.61
35	BB	839	G	C4'-C3'	8.64	1.62	1.53
40	BG	85	C	O3'-P	-8.64	1.50	1.61
85	AA	248	U	C2-N3	-8.64	1.31	1.37
85	AA	321	C	C2'-C1'	-8.64	1.43	1.53
85	AA	1515	A	O3'-P	-8.64	1.50	1.61
85	AA	1171	C	C2'-C1'	-8.64	1.43	1.53
85	AA	1688	U	O3'-P	-8.64	1.50	1.61
34	BA	457	A	P-O5'	-8.64	1.51	1.59
35	BB	1259	A	C5'-C4'	8.64	1.61	1.51
34	BA	1028	A	C2'-C1'	-8.63	1.43	1.53
34	BA	1743	U	C2'-C1'	-8.63	1.43	1.53
35	BB	80	C	P-O5'	-8.64	1.51	1.59
35	BB	811	C	C2'-C1'	-8.63	1.43	1.53
37	BD	4	U	C1'-N1	-8.64	1.34	1.46
34	BA	103	G	C1'-N9	-8.63	1.34	1.46
34	BA	307	C	P-O5'	-8.63	1.51	1.59
35	BB	490	G	N3-C4	-8.63	1.29	1.35
35	BB	1057	G	N9-C4	-8.63	1.31	1.38
35	BB	1440	A	C4'-C3'	-8.63	1.43	1.53
85	AA	1024	G	P-O5'	-8.63	1.51	1.59
34	BA	757	G	N7-C5	-8.63	1.34	1.39
34	BA	903	C	C2'-C1'	-8.63	1.43	1.53
34	BA	970	U	P-O5'	-8.63	1.51	1.59
35	BB	596	C	C2'-C1'	-8.63	1.43	1.53
37	BD	89	G	N9-C4	-8.63	1.31	1.38
38	BE	89	G	C3'-C2'	-8.63	1.43	1.52
34	BA	260	A	C1'-N9	-8.63	1.34	1.46
34	BA	366	G	C1'-N9	-8.63	1.34	1.46
34	BA	1712	U	O3'-P	-8.63	1.50	1.61
35	BB	1191	G	O3'-P	-8.63	1.50	1.61
40	BG	34	A	C3'-C2'	-8.63	1.43	1.52
34	BA	1072	U	C3'-C2'	-8.63	1.43	1.52
34	BA	1197	U	O3'-P	-8.63	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	269	A	N9-C4	-8.63	1.32	1.37
36	BC	121	G	N9-C4	-8.63	1.31	1.38
38	BE	31	A	N7-C5	-8.63	1.34	1.39
85	AA	2117	U	P-O5'	-8.63	1.51	1.59
34	BA	503	C	O3'-P	-8.62	1.50	1.61
34	BA	1543	A	N7-C5	-8.62	1.34	1.39
35	BB	879	G	C2'-C1'	-8.63	1.43	1.53
85	AA	1550	C	P-O5'	-8.62	1.51	1.59
34	BA	1256	A	C2'-C1'	-8.62	1.43	1.53
34	BA	1642	A	C5-C4	-8.62	1.32	1.38
35	BB	1311	G	C3'-C2'	-8.62	1.43	1.52
85	AA	2196	G	N7-C5	-8.62	1.34	1.39
34	BA	1070	G	N1-C2	-8.62	1.30	1.37
34	BA	1295	U	C2-N3	-8.62	1.31	1.37
34	BA	1301	G	C5-C4	-8.62	1.32	1.38
35	BB	579	A	N9-C4	-8.62	1.32	1.37
36	BC	9	G	C6-N1	-8.62	1.33	1.39
38	BE	121	G	C1'-N9	-8.62	1.34	1.46
85	AA	368	C	C3'-C2'	-8.62	1.43	1.52
85	AA	459	C	C2'-C1'	-8.62	1.43	1.53
85	AA	913	U	C2'-C1'	-8.62	1.43	1.53
85	AA	1752	C	P-O5'	-8.62	1.51	1.59
34	BA	1590	G	P-O5'	-8.62	1.51	1.59
34	BA	1435	A	N7-C5	-8.61	1.34	1.39
40	BG	2	U	P-O5'	-8.62	1.51	1.59
40	BG	35	G	C1'-N9	-8.61	1.34	1.46
85	AA	542	G	P-O5'	-8.62	1.51	1.59
34	BA	1107	A	N3-C4	-8.61	1.29	1.34
35	BB	1254	G	C4'-O4'	-8.61	1.34	1.45
34	BA	81	C	O3'-P	-8.61	1.50	1.61
34	BA	940	C	P-O5'	-8.61	1.51	1.59
34	BA	1000	G	C6-N1	-8.61	1.33	1.39
34	BA	1249	G	C6-N1	-8.61	1.33	1.39
35	BB	430	A	P-O5'	-8.61	1.51	1.59
35	BB	679	G	C2'-C1'	-8.61	1.43	1.53
35	BB	972	C	P-O5'	-8.61	1.51	1.59
35	BB	1190	U	C2-N3	-8.61	1.31	1.37
35	BB	1430	G	C2'-C1'	-8.61	1.43	1.53
36	BC	14	G	N9-C4	-8.61	1.31	1.38
38	BE	124	G	O3'-P	-8.61	1.50	1.61
40	BG	35	G	N9-C8	-8.61	1.31	1.37
85	AA	1248	U	P-O5'	-8.61	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	839	U	C3'-C2'	-8.61	1.43	1.52
34	BA	1472	G	N1-C2	-8.61	1.30	1.37
34	BA	977	G	O3'-P	-8.61	1.50	1.61
35	BB	1021	C	C2'-C1'	-8.61	1.43	1.53
35	BB	1079	G	O3'-P	-8.61	1.50	1.61
37	BD	51	G	N9-C4	-8.61	1.31	1.38
39	BF	58	U	C2'-C1'	-8.61	1.43	1.53
34	BA	755	G	O3'-P	-8.61	1.50	1.61
35	BB	570	A	C3'-C2'	-8.61	1.43	1.52
39	BF	19	A	O3'-P	-8.61	1.50	1.61
34	BA	713	C	C3'-C2'	-8.61	1.43	1.52
34	BA	732	A	C2'-C1'	-8.61	1.43	1.53
36	BC	112	G	N7-C5	-8.61	1.34	1.39
41	BH	47	G	O3'-P	-8.61	1.50	1.61
85	AA	520	A	C5'-C4'	8.61	1.61	1.51
34	BA	689	C	C2-N3	-8.60	1.28	1.35
36	BC	43	A	C4'-C3'	-8.60	1.43	1.53
34	BA	1012	A	P-O5'	-8.60	1.51	1.59
34	BA	1836	A	O3'-P	-8.60	1.50	1.61
36	BC	2	A	C2'-C1'	-8.60	1.43	1.53
37	BD	73	U	O3'-P	-8.60	1.50	1.61
38	BE	189	A	N9-C4	-8.60	1.32	1.37
85	AA	621	U	O3'-P	-8.60	1.50	1.61
85	AA	1969	A	O3'-P	-8.60	1.50	1.61
38	BE	35	A	P-O5'	-8.60	1.51	1.59
40	BG	174	G	N9-C4	-8.60	1.31	1.38
85	AA	243	A	C5'-C4'	8.60	1.61	1.51
85	AA	276	C	C2'-C1'	-8.60	1.43	1.53
34	BA	99	G	C3'-C2'	-8.60	1.43	1.52
34	BA	395	G	C6-N1	-8.60	1.33	1.39
34	BA	930	A	O3'-P	-8.60	1.50	1.61
34	BA	1710	C	C2-N3	-8.60	1.28	1.35
35	BB	899	C	P-O5'	-8.60	1.51	1.59
85	AA	1466	U	C4'-O4'	-8.60	1.34	1.45
34	BA	941	G	C1'-N9	-8.60	1.34	1.46
35	BB	1265	U	P-O5'	-8.60	1.51	1.59
85	AA	1485	G	O3'-P	-8.60	1.50	1.61
85	AA	92	G	P-O5'	-8.60	1.51	1.59
34	BA	1099	U	C3'-C2'	-8.60	1.43	1.52
34	BA	1447	C	C2'-C1'	-8.60	1.43	1.53
37	BD	54	A	N9-C4	-8.60	1.32	1.37
38	BE	114	G	C1'-N9	-8.60	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	137	G	C5-C4	-8.60	1.32	1.38
85	AA	105	A	C3'-C2'	-8.60	1.43	1.52
85	AA	490	A	O3'-P	-8.60	1.50	1.61
85	AA	578	U	C5'-C4'	-8.60	1.41	1.51
85	AA	866	U	C3'-C2'	-8.60	1.43	1.52
85	AA	1199	C	C2'-C1'	-8.60	1.43	1.53
85	AA	1958	C	P-O5'	-8.60	1.51	1.59
34	BA	78	U	C4'-C3'	-8.59	1.43	1.53
34	BA	978	U	O3'-P	-8.59	1.50	1.61
85	AA	441	C	C2'-C1'	-8.59	1.43	1.53
34	BA	79	C	C2-N3	-8.59	1.28	1.35
34	BA	1070	G	P-O5'	-8.59	1.51	1.59
35	BB	1373	U	C2-N3	-8.59	1.31	1.37
34	BA	1519	G	C1'-N9	-8.59	1.34	1.46
35	BB	1087	A	O3'-P	-8.59	1.50	1.61
41	BH	21	G	P-O5'	-8.59	1.51	1.59
85	AA	690	G	C6-N1	-8.59	1.33	1.39
85	AA	1063	U	P-O5'	-8.59	1.51	1.59
34	BA	1040	G	C5-C4	-8.59	1.32	1.38
35	BB	399	A	O3'-P	-8.59	1.50	1.61
34	BA	63	A	O3'-P	-8.59	1.50	1.61
34	BA	307	C	C2'-C1'	-8.59	1.44	1.53
34	BA	593	G	C1'-N9	8.59	1.61	1.48
35	BB	1119	G	O3'-P	-8.59	1.50	1.61
85	AA	158	C	O3'-P	-8.59	1.50	1.61
85	AA	1516	A	C4'-C3'	-8.59	1.43	1.53
85	AA	2045	U	O3'-P	-8.59	1.50	1.61
34	BA	766	A	N1-C2	-8.59	1.26	1.34
34	BA	816	G	C2'-C1'	-8.59	1.44	1.53
34	BA	1651	C	O3'-P	-8.59	1.50	1.61
35	BB	1208	G	O3'-P	-8.59	1.50	1.61
85	AA	411	U	C2'-C1'	-8.59	1.44	1.53
85	AA	429	G	O3'-P	-8.59	1.50	1.61
85	AA	1167	G	N7-C5	-8.59	1.34	1.39
34	BA	975	A	N9-C4	-8.59	1.32	1.37
35	BB	436	G	N7-C5	-8.59	1.34	1.39
39	BF	57	C	C3'-C2'	-8.59	1.43	1.52
85	AA	399	A	N9-C4	-8.59	1.32	1.37
85	AA	452	A	C2'-C1'	-8.59	1.44	1.53
85	AA	322	A	N9-C4	-8.59	1.32	1.37
85	AA	493	A	C2'-C1'	-8.59	1.44	1.53
85	AA	527	A	C2'-C1'	-8.58	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	113	G	O3'-P	-8.58	1.50	1.61
34	BA	316	G	C1'-N9	-8.58	1.34	1.46
34	BA	1846	G	C3'-C2'	-8.58	1.43	1.52
34	BA	376	U	O3'-P	-8.58	1.50	1.61
34	BA	817	U	C2-N3	-8.58	1.31	1.37
34	BA	1262	A	C2'-C1'	-8.58	1.44	1.53
34	BA	1830	A	P-O5'	-8.58	1.51	1.59
35	BB	794	G	C6-N1	-8.58	1.33	1.39
41	BH	113	G	C2'-C1'	-8.58	1.44	1.53
35	BB	1145	G	C8-N7	-8.58	1.25	1.30
34	BA	1804	A	P-O5'	-8.58	1.51	1.59
35	BB	555	G	P-O5'	-8.58	1.51	1.59
35	BB	1294	C	O3'-P	-8.58	1.50	1.61
41	BH	123	G	P-O5'	-8.58	1.51	1.59
34	BA	805	A	C2'-C1'	-8.57	1.44	1.53
35	BB	484	G	C6-N1	-8.57	1.33	1.39
85	AA	2087	C	P-O5'	-8.57	1.51	1.59
34	BA	10	G	C1'-N9	-8.57	1.34	1.46
34	BA	260	A	N9-C4	-8.57	1.32	1.37
34	BA	1245	C	C2-N3	-8.57	1.28	1.35
85	AA	330	C	O3'-P	-8.57	1.50	1.61
85	AA	1540	A	N9-C4	-8.57	1.32	1.37
34	BA	189	G	N3-C4	-8.57	1.29	1.35
34	BA	1196	C	O3'-P	-8.57	1.50	1.61
34	BA	1671	A	N9-C4	-8.57	1.32	1.37
35	BB	1377	A	C1'-N9	-8.57	1.34	1.46
35	BB	1388	A	O3'-P	-8.57	1.50	1.61
35	BB	1459	U	N1-C2	-8.57	1.30	1.38
40	BG	96	C	C2-N3	-8.57	1.28	1.35
85	AA	187	C	P-O5'	-8.57	1.51	1.59
85	AA	441	C	P-O5'	-8.57	1.51	1.59
34	BA	1166	A	N9-C4	-8.57	1.32	1.37
34	BA	1523	U	C2-N3	-8.57	1.31	1.37
35	BB	854	G	P-O5'	-8.57	1.51	1.59
85	AA	485	A	C1'-N9	-8.57	1.34	1.46
85	AA	1216	A	C2'-C1'	-8.57	1.44	1.53
85	AA	1855	U	O3'-P	-8.57	1.50	1.61
34	BA	448	U	C2-N3	-8.57	1.31	1.37
34	BA	49	A	P-O5'	-8.57	1.51	1.59
34	BA	1632	G	P-O5'	-8.57	1.51	1.59
35	BB	392	G	C1'-N9	-8.57	1.34	1.46
35	BB	1049	G	O3'-P	-8.57	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	575	G	C1'-N9	-8.57	1.34	1.46
35	BB	603	U	O3'-P	-8.57	1.50	1.61
38	BE	193	A	O3'-P	-8.57	1.50	1.61
34	BA	699	G	P-O5'	-8.56	1.51	1.59
34	BA	1331	G	C2'-C1'	-8.56	1.44	1.53
35	BB	1511	U	C2-N3	-8.56	1.31	1.37
85	AA	360	C	P-O5'	-8.56	1.51	1.59
85	AA	804	A	N9-C4	-8.56	1.32	1.37
85	AA	878	U	C2'-C1'	-8.56	1.44	1.53
34	BA	95	C	C2-N3	-8.56	1.28	1.35
34	BA	973	U	C4'-C3'	-8.56	1.43	1.53
35	BB	1546	C	C3'-C2'	-8.56	1.43	1.52
34	BA	463	A	N3-C4	-8.56	1.29	1.34
85	AA	1812	C	P-O5'	-8.56	1.51	1.59
34	BA	325	A	C2'-C1'	-8.56	1.44	1.53
34	BA	1103	G	N9-C4	-8.56	1.31	1.38
34	BA	1318	G	C2'-C1'	-8.56	1.44	1.53
35	BB	104	G	C2'-C1'	-8.56	1.44	1.53
35	BB	1406	C	C3'-C2'	-8.56	1.43	1.52
40	BG	88	G	C6-N1	-8.56	1.33	1.39
85	AA	1586	C	C2'-C1'	-8.56	1.44	1.53
85	AA	2123	U	C3'-C2'	-8.56	1.43	1.52
34	BA	178	C	C2-N3	-8.55	1.28	1.35
34	BA	792	A	N7-C5	-8.55	1.34	1.39
34	BA	887	U	C3'-C2'	-8.56	1.43	1.52
34	BA	757	G	C4'-O4'	-8.55	1.34	1.45
34	BA	983	A	O3'-P	-8.55	1.50	1.61
34	BA	1838	U	P-O5'	-8.55	1.51	1.59
35	BB	614	U	O3'-P	-8.55	1.50	1.61
35	BB	1199	A	C3'-C2'	-8.55	1.43	1.52
35	BB	1326	U	O3'-P	-8.55	1.50	1.61
35	BB	1523	U	P-O5'	-8.55	1.51	1.59
36	BC	77	A	N9-C4	-8.55	1.32	1.37
40	BG	126	G	P-O5'	-8.55	1.51	1.59
34	BA	41	U	C2-N3	-8.55	1.31	1.37
34	BA	726	G	O3'-P	-8.55	1.50	1.61
34	BA	815	C	C2'-C1'	-8.55	1.44	1.53
34	BA	1579	G	O3'-P	-8.55	1.50	1.61
35	BB	1153	G	N7-C5	-8.55	1.34	1.39
36	BC	144	C	P-O5'	-8.55	1.51	1.59
38	BE	66	A	O3'-P	-8.55	1.50	1.61
34	BA	764	G	C5'-C4'	8.55	1.61	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1053	G	C2-N2	-8.55	1.26	1.34
34	BA	1262	A	C5-C4	-8.55	1.32	1.38
35	BB	39	C	C3'-C2'	-8.55	1.43	1.52
35	BB	393	A	P-O5'	-8.55	1.51	1.59
35	BB	844	G	P-O5'	-8.55	1.51	1.59
40	BG	105	A	C1'-N9	-8.55	1.34	1.46
85	AA	515	C	C2-N3	-8.55	1.28	1.35
85	AA	1707	G	N7-C5	-8.55	1.34	1.39
34	BA	114	U	C2'-C1'	-8.55	1.44	1.53
34	BA	218	G	C2'-C1'	-8.54	1.44	1.53
34	BA	474	A	C1'-N9	-8.54	1.34	1.46
34	BA	501	U	C1'-N1	8.55	1.61	1.48
34	BA	1163	G	O3'-P	-8.55	1.50	1.61
34	BA	1821	A	P-O5'	-8.55	1.51	1.59
37	BD	87	G	C5-C4	-8.55	1.32	1.38
34	BA	1670	A	O3'-P	-8.54	1.50	1.61
35	BB	67	A	P-O5'	-8.54	1.51	1.59
85	AA	989	U	O3'-P	-8.54	1.50	1.61
85	AA	1272	G	C2'-C1'	-8.55	1.44	1.53
85	AA	1509	A	C2'-C1'	-8.54	1.44	1.53
34	BA	400	A	N9-C4	-8.54	1.32	1.37
34	BA	681	G	O3'-P	-8.54	1.50	1.61
34	BA	804	G	P-O5'	-8.54	1.51	1.59
34	BA	1014	A	P-O5'	-8.54	1.51	1.59
34	BA	1550	G	C5-C4	-8.54	1.32	1.38
34	BA	1565	U	C2-N3	-8.54	1.31	1.37
34	BA	1706	A	C5'-C4'	8.54	1.61	1.51
35	BB	1183	U	C2'-C1'	-8.54	1.44	1.53
85	AA	1885	A	C1'-N9	-8.54	1.34	1.46
34	BA	1614	G	N9-C4	-8.54	1.31	1.38
35	BB	587	A	C1'-N9	-8.54	1.34	1.46
34	BA	20	A	C1'-N9	-8.54	1.34	1.46
34	BA	760	G	N9-C4	-8.54	1.31	1.38
34	BA	1845	G	P-O5'	-8.54	1.51	1.59
40	BG	152	G	P-O5'	-8.54	1.51	1.59
34	BA	161	U	N1-C2	-8.54	1.30	1.38
34	BA	1292	A	P-O5'	-8.54	1.51	1.59
34	BA	1295	U	C1'-N1	-8.54	1.34	1.46
34	BA	1513	G	C6-N1	-8.54	1.33	1.39
85	AA	119	G	N3-C4	-8.54	1.29	1.35
85	AA	912	C	P-O5'	-8.54	1.51	1.59
35	BB	64	U	C2'-C1'	-8.54	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1509	G	C2'-C1'	-8.54	1.44	1.53
41	BH	20	A	O3'-P	-8.54	1.50	1.61
85	AA	890	U	O3'-P	-8.54	1.50	1.61
34	BA	200	C	O3'-P	-8.54	1.50	1.61
35	BB	793	A	P-O5'	-8.54	1.51	1.59
34	BA	819	G	C1'-N9	-8.53	1.34	1.46
34	BA	1474	G	C6-N1	-8.54	1.33	1.39
35	BB	1404	A	N3-C4	-8.54	1.29	1.34
35	BB	1003	G	C2-N2	-8.53	1.26	1.34
34	BA	53	G	P-O5'	-8.53	1.51	1.59
35	BB	366	G	C2'-C1'	-8.53	1.44	1.53
35	BB	683	U	O3'-P	-8.53	1.50	1.61
35	BB	967	G	P-O5'	-8.53	1.51	1.59
85	AA	373	G	P-O5'	-8.53	1.51	1.59
34	BA	470	C	N1-C2	-8.53	1.31	1.40
34	BA	507	U	C5'-C4'	8.53	1.61	1.51
34	BA	1225	A	O3'-P	-8.53	1.50	1.61
34	BA	1243	A	N9-C4	-8.53	1.32	1.37
34	BA	1032	A	N9-C4	-8.53	1.32	1.37
35	BB	548	A	N7-C5	-8.53	1.34	1.39
35	BB	1067	G	N7-C5	-8.53	1.34	1.39
85	AA	181	A	N7-C5	-8.53	1.34	1.39
85	AA	1822	G	C2'-C1'	-8.53	1.44	1.53
34	BA	1549	U	C2-N3	-8.53	1.31	1.37
34	BA	276	C	C2-N3	-8.53	1.28	1.35
34	BA	398	G	C2-N3	-8.53	1.25	1.32
34	BA	617	G	C2'-C1'	-8.53	1.44	1.53
35	BB	546	A	C5-C4	-8.53	1.32	1.38
38	BE	39	U	C3'-C2'	-8.53	1.43	1.52
35	BB	778	A	C5'-C4'	-8.53	1.41	1.51
38	BE	90	G	O3'-P	-8.53	1.50	1.61
38	BE	117	A	N7-C5	-8.53	1.34	1.39
34	BA	484	A	O3'-P	-8.52	1.50	1.61
34	BA	915	A	N9-C4	-8.52	1.32	1.37
34	BA	1114	G	C6-N1	-8.52	1.33	1.39
34	BA	1417	C	C3'-C2'	-8.52	1.43	1.52
37	BD	35	C	C2'-C1'	-8.52	1.44	1.53
40	BG	38	A	C3'-C2'	-8.52	1.43	1.52
41	BH	69	C	O3'-P	-8.52	1.50	1.61
35	BB	133	G	C1'-N9	-8.52	1.34	1.46
35	BB	1148	U	O3'-P	-8.52	1.50	1.61
40	BG	31	G	C5-C4	-8.52	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	16	A	N7-C5	-8.52	1.34	1.39
41	BH	19	G	C3'-C2'	-8.52	1.43	1.52
85	AA	1557	U	O3'-P	-8.52	1.50	1.61
34	BA	74	A	O3'-P	-8.52	1.50	1.61
35	BB	637	G	N3-C4	-8.52	1.29	1.35
34	BA	1800	G	N3-C4	-8.52	1.29	1.35
35	BB	1074	U	O3'-P	-8.52	1.50	1.61
38	BE	37	C	O3'-P	-8.52	1.50	1.61
85	AA	269	G	P-O5'	-8.52	1.51	1.59
85	AA	1699	A	N7-C5	-8.52	1.34	1.39
85	AA	1865	C	O3'-P	-8.52	1.50	1.61
34	BA	52	G	C6-N1	-8.52	1.33	1.39
34	BA	763	U	C4'-C3'	-8.52	1.43	1.53
34	BA	1513	G	P-O5'	-8.52	1.51	1.59
35	BB	493	U	C2'-C1'	-8.52	1.44	1.53
35	BB	378	C	P-O5'	-8.52	1.51	1.59
35	BB	662	G	N9-C4	-8.52	1.31	1.38
35	BB	1052	G	C1'-N9	-8.52	1.34	1.46
36	BC	60	U	O3'-P	-8.52	1.50	1.61
34	BA	43	U	P-O5'	-8.51	1.51	1.59
34	BA	110	C	C2'-C1'	-8.51	1.44	1.53
34	BA	406	G	C2'-C1'	-8.51	1.44	1.53
34	BA	1287	G	C1'-N9	-8.51	1.34	1.46
38	BE	77	C	O3'-P	-8.51	1.50	1.61
37	BD	82	G	C2'-C1'	-8.51	1.44	1.53
39	BF	58	U	O3'-P	-8.51	1.50	1.61
40	BG	75	C	O3'-P	-8.51	1.50	1.61
85	AA	1584	U	O3'-P	-8.51	1.50	1.61
85	AA	2082	C	P-O5'	-8.51	1.51	1.59
40	BG	165	C	O3'-P	-8.51	1.50	1.61
34	BA	308	C	O3'-P	-8.51	1.50	1.61
35	BB	638	G	O3'-P	-8.51	1.50	1.61
35	BB	1185	G	C2'-C1'	-8.51	1.44	1.53
85	AA	429	G	N9-C4	-8.51	1.31	1.38
85	AA	1700	C	C2'-C1'	-8.51	1.44	1.53
40	BG	95	U	P-O5'	-8.51	1.51	1.59
85	AA	922	A	N9-C4	-8.51	1.32	1.37
85	AA	2127	G	C5-C6	-8.51	1.33	1.42
34	BA	260	A	C2'-C1'	-8.51	1.44	1.53
34	BA	1149	C	C2-N3	-8.51	1.28	1.35
34	BA	1189	A	O3'-P	-8.51	1.50	1.61
34	BA	1324	G	O3'-P	-8.51	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1840	C	C3'-C2'	-8.51	1.43	1.52
35	BB	422	U	C2-N3	-8.51	1.31	1.37
35	BB	810	G	C2'-C1'	-8.51	1.44	1.53
36	BC	66	G	C2'-C1'	-8.51	1.44	1.53
36	BC	136	G	C2'-C1'	-8.51	1.44	1.53
39	BF	3	A	O3'-P	-8.51	1.50	1.61
85	AA	286	C	P-O5'	-8.51	1.51	1.59
85	AA	1672	G	P-O5'	-8.51	1.51	1.59
85	AA	62	A	N9-C4	-8.51	1.32	1.37
85	AA	703	U	P-O5'	-8.51	1.51	1.59
34	BA	280	A	N7-C5	-8.50	1.34	1.39
34	BA	498	A	C3'-C2'	-8.50	1.43	1.52
35	BB	772	U	P-O5'	-8.50	1.51	1.59
35	BB	1378	U	O3'-P	-8.50	1.50	1.61
36	BC	115	G	O3'-P	-8.50	1.50	1.61
85	AA	702	G	P-O5'	-8.50	1.51	1.59
85	AA	856	G	C2'-C1'	-8.50	1.44	1.53
85	AA	917	A	P-O5'	-8.50	1.51	1.59
85	AA	1583	U	P-O5'	-8.50	1.51	1.59
34	BA	494	A	C2'-C1'	-8.50	1.44	1.53
34	BA	1722	U	O4'-C1'	-8.50	1.30	1.41
35	BB	66	G	N9-C4	-8.50	1.31	1.38
36	BC	127	C	C2'-C1'	-8.50	1.44	1.53
85	AA	97	A	C5-C4	-8.50	1.32	1.38
85	AA	1242	A	P-O5'	-8.50	1.51	1.59
85	AA	2072	G	P-O5'	-8.50	1.51	1.59
34	BA	410	G	C2'-C1'	-8.50	1.44	1.53
34	BA	706	C	C2'-C1'	-8.50	1.44	1.53
35	BB	1488	G	C6-N1	-8.50	1.33	1.39
85	AA	1717	G	C5'-C4'	8.50	1.61	1.51
34	BA	1721	U	C2-N3	-8.50	1.31	1.37
35	BB	435	A	C5-C4	-8.50	1.32	1.38
85	AA	372	U	O3'-P	-8.50	1.50	1.61
85	AA	1132	A	N9-C4	-8.50	1.32	1.37
85	AA	1860	A	P-O5'	-8.50	1.51	1.59
34	BA	324	C	O3'-P	-8.49	1.50	1.61
34	BA	341	U	O4'-C1'	-8.49	1.30	1.41
85	AA	1882	U	P-O5'	-8.49	1.51	1.59
34	BA	1275	G	C6-N1	-8.49	1.33	1.39
35	BB	614	U	C3'-C2'	-8.49	1.43	1.52
34	BA	1408	C	C2'-C1'	-8.49	1.44	1.53
41	BH	17	A	O3'-P	-8.49	1.50	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	88	G	C6-N1	-8.49	1.33	1.39
85	AA	1057	G	C2'-C1'	-8.49	1.44	1.53
85	AA	2090	C	P-O5'	-8.49	1.51	1.59
34	BA	185	A	C2'-C1'	-8.49	1.44	1.53
35	BB	52	G	C3'-C2'	-8.49	1.43	1.52
35	BB	505	G	C1'-N9	-8.49	1.34	1.46
35	BB	644	A	P-O5'	-8.49	1.51	1.59
35	BB	1047	C	C2-N3	-8.49	1.28	1.35
35	BB	1025	A	C4'-C3'	8.49	1.62	1.53
85	AA	442	G	O3'-P	-8.49	1.50	1.61
34	BA	576	C	O3'-P	-8.49	1.50	1.61
40	BG	28	A	C3'-C2'	-8.49	1.43	1.52
34	BA	1724	G	N9-C4	-8.49	1.31	1.38
38	BE	193	A	N7-C5	-8.49	1.34	1.39
85	AA	62	A	C1'-N9	-8.49	1.34	1.46
85	AA	1264	U	P-O5'	-8.49	1.51	1.59
34	BA	20	A	N3-C4	-8.48	1.29	1.34
34	BA	1168	C	O3'-P	-8.48	1.50	1.61
34	BA	1227	U	C3'-C2'	-8.48	1.43	1.52
34	BA	1240	G	N7-C5	-8.48	1.34	1.39
34	BA	1475	G	C5-C4	-8.48	1.32	1.38
35	BB	624	A	O3'-P	-8.48	1.50	1.61
85	AA	1214	C	C4'-C3'	-8.48	1.43	1.53
41	BH	32	U	P-O5'	-8.48	1.51	1.59
85	AA	442	G	N9-C8	-8.48	1.31	1.37
85	AA	851	G	O3'-P	-8.48	1.50	1.61
34	BA	1025	A	C1'-N9	-8.48	1.34	1.46
34	BA	1686	G	C1'-N9	-8.48	1.34	1.46
35	BB	108	G	N9-C4	-8.48	1.31	1.38
35	BB	1470	G	N7-C5	-8.48	1.34	1.39
41	BH	35	G	N7-C5	-8.48	1.34	1.39
85	AA	163	C	P-O5'	-8.48	1.51	1.59
85	AA	289	G	C2'-C1'	-8.48	1.44	1.53
85	AA	2213	A	N9-C4	-8.48	1.32	1.37
85	AA	1105	G	C6-N1	-8.48	1.33	1.39
35	BB	1051	U	O3'-P	-8.48	1.50	1.61
37	BD	9	C	C2-N3	-8.48	1.28	1.35
85	AA	595	A	C2'-C1'	-8.48	1.44	1.53
35	BB	540	G	N7-C5	-8.48	1.34	1.39
38	BE	106	C	C4'-C3'	-8.48	1.43	1.53
40	BG	131	U	C2-N3	-8.48	1.31	1.37
85	AA	1868	G	C2'-C1'	-8.48	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2228	G	O3'-P	-8.48	1.50	1.61
34	BA	38	G	C6-N1	-8.47	1.33	1.39
85	AA	887	A	C4'-C3'	-8.47	1.43	1.53
34	BA	987	C	C2'-C1'	-8.47	1.44	1.53
35	BB	662	G	N1-C2	-8.47	1.30	1.37
35	BB	1058	U	C2-N3	-8.47	1.31	1.37
36	BC	29	C	O3'-P	-8.47	1.50	1.61
36	BC	113	G	C6-N1	-8.47	1.33	1.39
37	BD	68	C	C2'-C1'	-8.47	1.44	1.53
40	BG	99	A	O3'-P	-8.47	1.50	1.61
85	AA	704	A	N9-C4	-8.47	1.32	1.37
85	AA	838	G	N9-C4	8.47	1.44	1.38
85	AA	930	G	C6-N1	8.47	1.45	1.39
85	AA	2022	A	N9-C4	-8.47	1.32	1.37
34	BA	180	G	C2'-C1'	-8.47	1.44	1.53
34	BA	446	U	C3'-C2'	-8.47	1.43	1.52
35	BB	509	A	N9-C4	-8.47	1.32	1.37
35	BB	1335	G	N9-C4	-8.47	1.31	1.38
34	BA	1181	G	O3'-P	-8.47	1.50	1.61
36	BC	75	G	P-O5'	-8.47	1.51	1.59
36	BC	134	G	P-O5'	-8.47	1.51	1.59
40	BG	66	C	C2'-C1'	-8.47	1.44	1.53
85	AA	1294	U	O4'-C1'	-8.47	1.30	1.41
85	AA	1928	A	P-O5'	-8.47	1.51	1.59
34	BA	111	U	O3'-P	-8.47	1.50	1.61
37	BD	42	A	N9-C4	-8.47	1.32	1.37
40	BG	92	U	C2-N3	-8.47	1.31	1.37
40	BG	169	A	P-O5'	-8.47	1.51	1.59
85	AA	677	U	O3'-P	-8.47	1.50	1.61
85	AA	1694	C	O3'-P	-8.47	1.50	1.61
85	AA	2234	C	P-O5'	-8.47	1.51	1.59
34	BA	174	A	P-O5'	-8.46	1.51	1.59
34	BA	474	A	O3'-P	-8.46	1.50	1.61
34	BA	888	G	C1'-N9	-8.47	1.34	1.46
34	BA	1807	G	O3'-P	-8.47	1.50	1.61
35	BB	59	U	P-O5'	-8.46	1.51	1.59
40	BG	78	C	C3'-C2'	-8.47	1.43	1.52
40	BG	123	C	O3'-P	-8.47	1.50	1.61
35	BB	1259	A	C1'-N9	-8.46	1.35	1.46
38	BE	44	C	P-O5'	-8.46	1.51	1.59
85	AA	450	A	C1'-N9	-8.46	1.35	1.46
85	AA	1149	A	C1'-N9	-8.46	1.34	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1240	A	O3'-P	-8.46	1.50	1.61
85	AA	1877	G	C2'-C1'	-8.46	1.44	1.53
34	BA	782	C	O3'-P	-8.46	1.50	1.61
34	BA	1549	U	C4'-C3'	-8.46	1.43	1.53
34	BA	1700	C	C3'-C2'	-8.46	1.43	1.52
35	BB	1250	A	P-O5'	-8.46	1.51	1.59
38	BE	186	C	P-O5'	-8.46	1.51	1.59
41	BH	124	C	C3'-C2'	-8.46	1.43	1.52
85	AA	963	U	P-O5'	-8.46	1.51	1.59
85	AA	1502	A	C1'-N9	-8.46	1.35	1.46
85	AA	1650	G	C2'-C1'	-8.46	1.44	1.53
34	BA	481	A	N7-C5	-8.46	1.34	1.39
34	BA	1301	G	O3'-P	-8.46	1.50	1.61
34	BA	1836	A	C1'-N9	-8.46	1.35	1.46
35	BB	407	A	O3'-P	-8.46	1.50	1.61
35	BB	450	A	O3'-P	-8.46	1.50	1.61
35	BB	799	A	N7-C5	-8.46	1.34	1.39
40	BG	117	C	C2'-C1'	-8.46	1.44	1.53
40	BG	155	A	P-O5'	-8.46	1.51	1.59
85	AA	415	G	N9-C4	-8.46	1.31	1.38
85	AA	719	C	P-O5'	-8.46	1.51	1.59
34	BA	594	G	C6-N1	-8.46	1.33	1.39
34	BA	1221	A	C1'-N9	-8.46	1.35	1.46
35	BB	1460	G	C2'-C1'	-8.46	1.44	1.53
35	BB	102	G	O3'-P	-8.46	1.51	1.61
35	BB	1053	G	C3'-C2'	-8.46	1.43	1.52
85	AA	504	U	C1'-N1	-8.46	1.35	1.46
35	BB	1097	U	O3'-P	-8.46	1.51	1.61
85	AA	288	G	N9-C4	-8.46	1.31	1.38
85	AA	326	C	O3'-P	-8.46	1.50	1.61
85	AA	395	G	O3'-P	-8.46	1.51	1.61
34	BA	1776	G	C1'-N9	-8.46	1.35	1.46
37	BD	5	A	O3'-P	-8.45	1.51	1.61
39	BF	70	A	P-O5'	-8.45	1.51	1.59
85	AA	463	G	C2-N2	-8.46	1.26	1.34
85	AA	1466	U	O4'-C1'	-8.46	1.30	1.41
34	BA	790	G	N7-C5	-8.45	1.34	1.39
34	BA	1000	G	C1'-N9	-8.45	1.35	1.46
34	BA	1153	C	P-O5'	-8.45	1.51	1.59
35	BB	550	G	C2-N3	-8.45	1.25	1.32
35	BB	1272	G	O3'-P	-8.45	1.51	1.61
38	BE	27	A	C4'-O4'	-8.45	1.34	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	151	C	P-O5'	-8.45	1.51	1.59
85	AA	1127	G	C5-C4	-8.45	1.32	1.38
85	AA	1976	G	O3'-P	-8.45	1.51	1.61
34	BA	604	G	P-O5'	-8.45	1.51	1.59
40	BG	27	C	C3'-C2'	-8.45	1.43	1.52
34	BA	1568	A	C2'-C1'	-8.45	1.44	1.53
35	BB	98	A	O3'-P	-8.45	1.51	1.61
35	BB	696	G	O3'-P	-8.45	1.51	1.61
35	BB	704	G	C6-N1	-8.45	1.33	1.39
85	AA	879	G	O3'-P	-8.45	1.51	1.61
85	AA	1195	U	O3'-P	-8.45	1.51	1.61
35	BB	1271	A	O3'-P	-8.45	1.51	1.61
85	AA	1562	U	O3'-P	-8.45	1.51	1.61
85	AA	2074	G	O3'-P	-8.45	1.51	1.61
85	AA	2077	G	N7-C5	-8.45	1.34	1.39
34	BA	472	G	N7-C5	-8.45	1.34	1.39
34	BA	269	G	N9-C4	-8.45	1.31	1.38
34	BA	1676	A	O3'-P	-8.45	1.51	1.61
34	BA	1695	G	P-O5'	-8.45	1.51	1.59
35	BB	684	U	P-O5'	-8.45	1.51	1.59
35	BB	1021	C	N1-C6	-8.45	1.32	1.37
35	BB	1278	A	C1'-N9	-8.45	1.35	1.46
37	BD	10	C	N1-C6	-8.45	1.32	1.37
85	AA	300	C	P-O5'	-8.45	1.51	1.59
34	BA	886	G	C2'-C1'	-8.44	1.44	1.53
34	BA	1119	A	N9-C4	-8.44	1.32	1.37
34	BA	1459	U	P-O5'	-8.45	1.51	1.59
35	BB	1202	G	P-O5'	-8.45	1.51	1.59
41	BH	104	U	C2-N3	-8.44	1.31	1.37
85	AA	605	A	C2'-C1'	-8.45	1.44	1.53
85	AA	1127	G	O3'-P	-8.45	1.51	1.61
85	AA	2075	C	C3'-C2'	-8.45	1.43	1.52
85	AA	2076	C	O3'-P	-8.44	1.51	1.61
34	BA	662	U	C2-N3	-8.44	1.31	1.37
34	BA	676	G	C6-N1	-8.44	1.33	1.39
34	BA	882	G	C3'-C2'	-8.44	1.43	1.52
85	AA	505	U	O3'-P	-8.44	1.51	1.61
36	BC	141	C	C3'-C2'	-8.44	1.43	1.52
85	AA	6	G	P-O5'	-8.44	1.51	1.59
85	AA	1290	G	C2-N2	-8.44	1.26	1.34
34	BA	237	A	N9-C8	-8.44	1.30	1.37
34	BA	1200	U	O3'-P	-8.44	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1328	U	O3'-P	-8.44	1.51	1.61
35	BB	1537	C	P-O5'	-8.44	1.51	1.59
36	BC	115	G	C6-N1	-8.44	1.33	1.39
85	AA	1004	G	C2'-C1'	-8.44	1.44	1.53
34	BA	1066	A	O3'-P	-8.44	1.51	1.61
34	BA	32	A	N9-C4	-8.44	1.32	1.37
34	BA	115	U	O4'-C1'	-8.44	1.30	1.41
34	BA	390	A	C5-C4	-8.44	1.32	1.38
34	BA	481	A	C8-N7	-8.44	1.25	1.31
34	BA	1270	G	C3'-C2'	-8.44	1.43	1.52
34	BA	1796	A	P-O5'	-8.44	1.51	1.59
36	BC	148	C	O3'-P	-8.44	1.51	1.61
40	BG	58	G	P-O5'	-8.44	1.51	1.59
40	BG	72	G	C5-C4	-8.44	1.32	1.38
40	BG	88	G	C2'-C1'	-8.44	1.44	1.53
35	BB	379	U	O3'-P	-8.43	1.51	1.61
38	BE	123	A	N7-C5	-8.43	1.34	1.39
35	BB	1494	G	P-O5'	-8.43	1.51	1.59
40	BG	178	G	N9-C4	-8.43	1.31	1.38
85	AA	13	U	P-O5'	-8.43	1.51	1.59
85	AA	2133	A	C3'-C2'	-8.43	1.43	1.52
34	BA	1467	U	C3'-C2'	-8.43	1.43	1.52
35	BB	569	G	O3'-P	-8.43	1.51	1.61
37	BD	61	C	O3'-P	-8.43	1.51	1.61
40	BG	99	A	C3'-C2'	-8.43	1.43	1.52
85	AA	30	G	P-O5'	-8.43	1.51	1.59
85	AA	742	U	P-O5'	-8.43	1.51	1.59
85	AA	1713	A	P-O5'	-8.43	1.51	1.59
85	AA	2142	A	C1'-N9	-8.43	1.35	1.46
85	AA	1796	C	C2'-C1'	-8.43	1.44	1.53
34	BA	525	A	N9-C4	-8.43	1.32	1.37
34	BA	1565	U	N3-C4	-8.43	1.30	1.38
35	BB	1094	A	C1'-N9	-8.43	1.35	1.46
37	BD	21	G	P-O5'	-8.43	1.51	1.59
85	AA	496	C	C3'-C2'	-8.43	1.43	1.52
85	AA	2081	A	O3'-P	-8.43	1.51	1.61
85	AA	1704	C	P-O5'	-8.43	1.51	1.59
34	BA	181	G	C2'-C1'	-8.42	1.44	1.53
38	BE	89	G	C1'-N9	-8.42	1.35	1.46
34	BA	103	G	C4'-C3'	-8.42	1.43	1.53
34	BA	1311	G	O3'-P	-8.42	1.51	1.61
36	BC	77	A	O3'-P	-8.42	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	105	A	O3'-P	-8.42	1.51	1.61
34	BA	776	U	C2'-C1'	-8.42	1.44	1.53
34	BA	1099	U	C2-N3	-8.42	1.31	1.37
34	BA	1418	G	O3'-P	-8.42	1.51	1.61
34	BA	1456	C	O3'-P	-8.42	1.51	1.61
35	BB	7	C	C2-N3	-8.42	1.29	1.35
35	BB	72	G	O3'-P	-8.42	1.51	1.61
35	BB	557	C	O3'-P	-8.42	1.51	1.61
35	BB	1272	G	P-O5'	-8.42	1.51	1.59
38	BE	181	U	O3'-P	-8.42	1.51	1.61
40	BG	71	C	C1'-N1	-8.42	1.35	1.46
41	BH	125	U	C2-N3	-8.42	1.31	1.37
85	AA	98	U	P-O5'	-8.42	1.51	1.59
85	AA	1486	G	N7-C5	-8.42	1.34	1.39
34	BA	956	G	N9-C4	-8.42	1.31	1.38
85	AA	2176	U	C2-N3	-8.42	1.31	1.37
34	BA	959	G	P-O5'	-8.42	1.51	1.59
34	BA	1225	A	N7-C5	-8.42	1.34	1.39
35	BB	372	U	C2'-C1'	-8.42	1.44	1.53
36	BC	154	A	C4'-C3'	-8.42	1.43	1.53
34	BA	66	C	P-O5'	-8.41	1.51	1.59
34	BA	520	G	N9-C4	-8.41	1.31	1.38
34	BA	1088	G	C1'-N9	-8.41	1.35	1.46
34	BA	1562	G	C6-N1	-8.41	1.33	1.39
34	BA	1696	G	P-O5'	-8.41	1.51	1.59
37	BD	96	C	P-O5'	-8.41	1.51	1.59
39	BF	48	G	P-O5'	-8.41	1.51	1.59
34	BA	761	U	P-O5'	-8.41	1.51	1.59
35	BB	434	A	C2'-C1'	-8.41	1.44	1.53
85	AA	419	A	O3'-P	-8.41	1.51	1.61
35	BB	1305	A	N9-C4	-8.41	1.32	1.37
37	BD	71	G	C2'-C1'	-8.41	1.44	1.53
85	AA	308	U	C2'-C1'	-8.41	1.44	1.53
85	AA	1260	G	C2'-C1'	-8.41	1.44	1.53
35	BB	3	C	O3'-P	-8.41	1.51	1.61
34	BA	1088	G	C5-C4	-8.41	1.32	1.38
35	BB	495	A	O3'-P	-8.41	1.51	1.61
35	BB	605	C	C3'-C2'	-8.41	1.43	1.52
35	BB	1450	G	O3'-P	-8.41	1.51	1.61
41	BH	38	G	C6-N1	-8.41	1.33	1.39
85	AA	640	C	O3'-P	-8.41	1.51	1.61
34	BA	88	C	O4'-C1'	-8.40	1.30	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1331	G	C1'-N9	-8.40	1.35	1.46
35	BB	645	C	O3'-P	-8.40	1.51	1.61
85	AA	1170	C	P-O5'	-8.40	1.51	1.59
34	BA	1580	U	O3'-P	-8.40	1.51	1.61
35	BB	828	G	C1'-N9	-8.40	1.35	1.46
40	BG	93	U	C2-N3	-8.40	1.31	1.37
85	AA	715	G	C2'-C1'	-8.40	1.44	1.53
85	AA	830	A	O3'-P	-8.40	1.51	1.61
85	AA	861	G	O3'-P	-8.40	1.51	1.61
34	BA	996	U	C1'-N1	-8.40	1.35	1.46
34	BA	1052	G	C5-C4	-8.40	1.32	1.38
34	BA	1239	G	P-O5'	-8.40	1.51	1.59
36	BC	154	A	C3'-C2'	-8.40	1.43	1.52
85	AA	1090	A	C2'-C1'	-8.40	1.44	1.53
40	BG	137	G	C4'-C3'	-8.40	1.44	1.53
85	AA	368	C	P-O5'	-8.40	1.51	1.59
85	AA	469	G	O3'-P	-8.40	1.51	1.61
85	AA	2021	A	P-O5'	-8.40	1.51	1.59
85	AA	2087	C	O3'-P	-8.40	1.51	1.61
34	BA	798	G	C2'-C1'	-8.40	1.44	1.53
34	BA	316	G	O3'-P	-8.40	1.51	1.61
34	BA	1541	G	C2'-C1'	-8.40	1.44	1.53
36	BC	53	A	C3'-C2'	-8.40	1.43	1.52
40	BG	77	U	P-O5'	-8.40	1.51	1.59
34	BA	946	A	N7-C5	-8.40	1.34	1.39
34	BA	1003	A	O3'-P	-8.40	1.51	1.61
34	BA	1806	A	C1'-N9	-8.40	1.35	1.46
85	AA	2130	G	C6-N1	-8.40	1.33	1.39
85	AA	2144	C	C2'-C1'	-8.40	1.44	1.53
34	BA	268	U	O3'-P	-8.39	1.51	1.61
34	BA	1531	G	O3'-P	-8.39	1.51	1.61
35	BB	1185	G	O3'-P	-8.39	1.51	1.61
39	BF	39	C	P-O5'	-8.39	1.51	1.59
85	AA	1952	C	P-O5'	-8.39	1.51	1.59
35	BB	1226	G	O3'-P	-8.39	1.51	1.61
85	AA	2019	G	P-O5'	-8.39	1.51	1.59
34	BA	523	A	C1'-N9	-8.39	1.35	1.46
34	BA	1203	G	O3'-P	-8.39	1.51	1.61
34	BA	1275	G	C2'-C1'	-8.39	1.44	1.53
38	BE	16	C	C2'-C1'	-8.39	1.44	1.53
35	BB	611	U	C2-N3	-8.39	1.31	1.37
37	BD	72	U	N1-C2	-8.39	1.30	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	137	G	C2'-C1'	-8.39	1.44	1.53
85	AA	2145	G	C3'-C2'	-8.39	1.43	1.52
38	BE	50	G	C1'-N9	-8.39	1.35	1.46
85	AA	962	U	C2'-C1'	-8.39	1.44	1.53
85	AA	1215	A	N9-C4	-8.39	1.32	1.37
37	BD	50	A	O3'-P	-8.39	1.51	1.61
34	BA	887	U	C2'-C1'	-8.39	1.44	1.53
35	BB	5	A	C5-C6	-8.39	1.33	1.41
35	BB	505	G	C3'-C2'	-8.39	1.43	1.52
35	BB	584	A	C5-C4	-8.39	1.32	1.38
41	BH	106	G	C6-N1	-8.39	1.33	1.39
85	AA	2066	C	C2'-C1'	-8.39	1.44	1.53
39	BF	11	C	C4'-C3'	8.39	1.62	1.53
85	AA	2153	G	N9-C4	-8.39	1.31	1.38
85	AA	2184	A	N9-C4	-8.39	1.32	1.37
34	BA	819	G	C2'-C1'	-8.39	1.44	1.53
34	BA	909	G	C1'-N9	-8.39	1.35	1.46
34	BA	461	A	P-O5'	-8.38	1.51	1.59
35	BB	619	A	C3'-C2'	-8.38	1.43	1.52
35	BB	1346	A	P-O5'	-8.38	1.51	1.59
35	BB	1445	A	N7-C5	-8.38	1.34	1.39
36	BC	152	C	O3'-P	-8.38	1.51	1.61
40	BG	80	G	O3'-P	-8.39	1.51	1.61
40	BG	124	A	C2'-C1'	-8.39	1.44	1.53
38	BE	22	A	O3'-P	-8.38	1.51	1.61
85	AA	449	G	O3'-P	-8.38	1.51	1.61
85	AA	960	G	N9-C4	-8.38	1.31	1.38
85	AA	1471	G	C5-C4	-8.38	1.32	1.38
85	AA	2188	C	C2'-C1'	-8.38	1.44	1.53
34	BA	526	C	C2'-C1'	-8.38	1.44	1.53
34	BA	696	A	C5-C4	-8.38	1.32	1.38
85	AA	1571	A	N9-C4	-8.38	1.32	1.37
34	BA	414	A	C2'-C1'	-8.38	1.44	1.53
34	BA	1441	C	C2-N3	-8.38	1.29	1.35
35	BB	462	G	P-O5'	-8.38	1.51	1.59
34	BA	1614	G	C5-C4	-8.38	1.32	1.38
36	BC	10	C	C4'-C3'	-8.38	1.44	1.53
38	BE	190	U	P-O5'	-8.38	1.51	1.59
85	AA	2127	G	C5-C4	-8.38	1.32	1.38
34	BA	1073	G	N9-C8	-8.38	1.31	1.37
34	BA	1100	A	N9-C4	-8.38	1.32	1.37
34	BA	1835	A	C2'-C1'	-8.38	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	780	U	O3'-P	-8.38	1.51	1.61
35	BB	1233	U	C2'-C1'	-8.38	1.44	1.53
36	BC	78	G	C2'-C1'	-8.38	1.44	1.53
41	BH	2	U	P-O5'	-8.38	1.51	1.59
35	BB	449	C	C2'-C1'	-8.38	1.44	1.53
85	AA	1588	A	P-O5'	-8.38	1.51	1.59
34	BA	152	C	C5'-C4'	8.38	1.61	1.51
34	BA	241	U	O3'-P	-8.38	1.51	1.61
34	BA	1244	G	C2'-C1'	-8.38	1.44	1.53
35	BB	804	U	O3'-P	-8.37	1.51	1.61
35	BB	816	U	P-O5'	-8.38	1.51	1.59
35	BB	1388	A	P-O5'	-8.38	1.51	1.59
41	BH	127	A	O3'-P	-8.38	1.51	1.61
36	BC	144	C	C2'-C1'	-8.37	1.44	1.53
40	BG	129	G	C1'-N9	-8.37	1.35	1.46
85	AA	79	G	O3'-P	-8.37	1.51	1.61
85	AA	257	U	C2-N3	-8.37	1.31	1.37
34	BA	1433	U	C1'-N1	-8.37	1.35	1.46
35	BB	1155	U	P-O5'	-8.37	1.51	1.59
35	BB	1160	U	P-O5'	-8.37	1.51	1.59
34	BA	516	U	C3'-C2'	-8.37	1.43	1.52
34	BA	1563	G	N7-C5	-8.37	1.34	1.39
34	BA	1637	G	O3'-P	-8.37	1.51	1.61
37	BD	94	C	C4-C5	-8.37	1.36	1.43
85	AA	1483	A	C1'-N9	-8.37	1.35	1.46
35	BB	486	G	C3'-C2'	-8.37	1.43	1.52
85	AA	105	A	O3'-P	-8.37	1.51	1.61
40	BG	108	G	C2'-C1'	-8.37	1.44	1.53
41	BH	4	U	C3'-C2'	-8.37	1.43	1.52
85	AA	629	A	N9-C4	-8.37	1.32	1.37
85	AA	687	G	C2'-C1'	-8.37	1.44	1.53
35	BB	815	G	C1'-N9	-8.37	1.35	1.46
38	BE	98	C	C2'-C1'	-8.37	1.44	1.53
34	BA	906	A	O3'-P	-8.36	1.51	1.61
34	BA	1470	G	P-O5'	-8.36	1.51	1.59
34	BA	1744	C	C3'-C2'	-8.37	1.43	1.52
35	BB	1076	U	P-O5'	-8.37	1.51	1.59
40	BG	98	A	P-O5'	-8.37	1.51	1.59
85	AA	437	G	C2'-C1'	-8.37	1.44	1.53
34	BA	736	G	N7-C5	-8.36	1.34	1.39
34	BA	1251	A	C1'-N9	-8.36	1.35	1.46
85	AA	932	A	O3'-P	-8.36	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	100	A	O3'-P	-8.36	1.51	1.61
34	BA	425	G	N3-C4	-8.36	1.29	1.35
34	BA	520	G	C5-C6	-8.36	1.33	1.42
35	BB	1142	C	C1'-N1	-8.36	1.35	1.46
39	BF	32	G	C1'-N9	-8.36	1.35	1.46
40	BG	87	G	N7-C5	-8.36	1.34	1.39
85	AA	1279	A	N9-C4	-8.36	1.32	1.37
34	BA	62	A	P-O5'	-8.36	1.51	1.59
34	BA	589	A	O3'-P	-8.36	1.51	1.61
34	BA	962	U	C2-N3	-8.36	1.31	1.37
34	BA	1522	G	C6-N1	-8.36	1.33	1.39
40	BG	146	C	C3'-C2'	-8.36	1.43	1.52
36	BC	125	A	N7-C5	-8.36	1.34	1.39
38	BE	93	U	C2'-C1'	-8.36	1.44	1.53
85	AA	1551	G	O3'-P	-8.36	1.51	1.61
34	BA	572	G	C5'-C4'	8.36	1.61	1.51
34	BA	1044	A	C2'-C1'	-8.36	1.44	1.53
34	BA	1226	G	C5-C4	-8.36	1.32	1.38
35	BB	75	A	O3'-P	-8.36	1.51	1.61
35	BB	1080	U	C2-N3	-8.36	1.31	1.37
35	BB	1520	C	C2'-C1'	-8.36	1.44	1.53
35	BB	1369	A	N9-C4	-8.36	1.32	1.37
85	AA	1163	G	P-O5'	-8.36	1.51	1.59
34	BA	1172	C	C2-N3	-8.35	1.29	1.35
34	BA	1195	G	C2'-C1'	-8.35	1.44	1.53
35	BB	464	C	P-O5'	-8.35	1.51	1.59
35	BB	495	A	N9-C4	-8.35	1.32	1.37
34	BA	263	G	P-O5'	-8.35	1.51	1.59
34	BA	782	C	C1'-N1	-8.35	1.35	1.46
34	BA	1305	A	O3'-P	-8.35	1.51	1.61
40	BG	55	A	P-O5'	-8.35	1.51	1.59
34	BA	650	C	C2'-C1'	-8.35	1.44	1.53
34	BA	1007	G	C1'-N9	-8.35	1.35	1.46
35	BB	662	G	O3'-P	-8.35	1.51	1.61
35	BB	1266	A	P-O5'	-8.35	1.51	1.59
36	BC	9	G	O3'-P	-8.35	1.51	1.61
34	BA	899	G	N7-C5	-8.35	1.34	1.39
34	BA	994	G	N7-C5	-8.35	1.34	1.39
35	BB	479	U	C3'-C2'	-8.35	1.43	1.52
35	BB	779	C	C2-N3	-8.35	1.29	1.35
36	BC	147	G	C6-N1	-8.35	1.33	1.39
85	AA	2227	A	C5-C4	-8.35	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1166	A	C1'-N9	-8.34	1.35	1.46
35	BB	798	A	O3'-P	-8.34	1.51	1.61
35	BB	1082	A	P-O5'	-8.34	1.51	1.59
35	BB	1372	G	O3'-P	-8.34	1.51	1.61
35	BB	1530	U	P-O5'	-8.34	1.51	1.59
37	BD	112	U	C2-N3	-8.34	1.31	1.37
38	BE	23	G	C1'-N9	-8.34	1.35	1.46
34	BA	1410	C	O3'-P	-8.34	1.51	1.61
34	BA	1575	U	O3'-P	-8.34	1.51	1.61
39	BF	71	G	N9-C4	-8.34	1.31	1.38
85	AA	1504	A	C2'-C1'	-8.34	1.44	1.53
35	BB	1194	A	N9-C4	-8.34	1.32	1.37
35	BB	1521	G	P-O5'	-8.34	1.51	1.59
36	BC	56	G	O3'-P	-8.34	1.51	1.61
38	BE	104	G	C4'-C3'	8.34	1.62	1.53
85	AA	995	G	C5-C4	-8.34	1.32	1.38
85	AA	1000	U	C4'-C3'	-8.34	1.44	1.53
85	AA	1912	U	P-O5'	-8.34	1.51	1.59
34	BA	1719	G	N1-C2	-8.34	1.31	1.37
34	BA	57	A	O3'-P	-8.34	1.51	1.61
34	BA	306	G	C6-N1	-8.34	1.33	1.39
35	BB	643	G	C6-N1	-8.34	1.33	1.39
85	AA	1484	G	C2'-C1'	-8.34	1.44	1.53
85	AA	2122	A	N9-C8	-8.34	1.31	1.37
34	BA	991	U	C3'-C2'	-8.34	1.43	1.52
35	BB	829	C	P-O5'	-8.34	1.51	1.59
36	BC	121	G	O3'-P	-8.34	1.51	1.61
40	BG	87	G	C6-N1	-8.34	1.33	1.39
74	Bo	74	PRO	CA-C	-8.34	1.36	1.52
85	AA	340	G	O3'-P	-8.34	1.51	1.61
85	AA	588	G	P-O5'	-8.34	1.51	1.59
85	AA	2127	G	C1'-N9	-8.34	1.35	1.46
34	BA	463	A	O3'-P	-8.33	1.51	1.61
35	BB	666	A	C2'-C1'	-8.33	1.44	1.53
38	BE	198	A	N9-C4	-8.33	1.32	1.37
40	BG	132	U	C2-N3	-8.33	1.31	1.37
37	BD	64	A	N9-C4	-8.33	1.32	1.37
40	BG	110	U	C2'-C1'	-8.33	1.44	1.53
85	AA	1516	A	C2'-C1'	-8.33	1.44	1.53
85	AA	2007	G	C3'-C2'	-8.33	1.43	1.52
34	BA	34	U	C2-N3	-8.33	1.31	1.37
34	BA	427	G	C6-N1	-8.33	1.33	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	729	C	C3'-C2'	-8.33	1.43	1.52
35	BB	33	A	N7-C5	-8.33	1.34	1.39
35	BB	658	G	C6-N1	-8.33	1.33	1.39
85	AA	2038	C	O3'-P	-8.33	1.51	1.61
85	AA	2130	G	N9-C4	-8.33	1.31	1.38
85	AA	703	U	C2-N3	-8.33	1.31	1.37
34	BA	706	C	C3'-C2'	-8.33	1.43	1.52
35	BB	599	U	P-O5'	-8.33	1.51	1.59
41	BH	130	G	C5'-C4'	-8.33	1.41	1.51
40	BG	11	G	N7-C5	-8.33	1.34	1.39
41	BH	119	U	C1'-N1	-8.33	1.35	1.46
85	AA	519	A	C1'-N9	-8.33	1.35	1.46
85	AA	1288	A	O3'-P	-8.33	1.51	1.61
34	BA	761	U	O3'-P	-8.32	1.51	1.61
34	BA	999	G	C2'-C1'	-8.32	1.44	1.53
35	BB	30	A	C2'-C1'	-8.32	1.44	1.53
35	BB	81	A	O3'-P	-8.32	1.51	1.61
35	BB	506	G	C2'-C1'	-8.32	1.44	1.53
35	BB	647	U	C3'-C2'	-8.32	1.43	1.52
35	BB	797	C	C2'-C1'	-8.32	1.44	1.53
35	BB	1136	G	N9-C4	-8.32	1.31	1.38
35	BB	1182	A	C1'-N9	-8.32	1.35	1.46
35	BB	1188	A	N9-C4	-8.32	1.32	1.37
85	AA	1290	G	O3'-P	-8.32	1.51	1.61
85	AA	1292	A	C4'-C3'	-8.32	1.44	1.53
85	AA	2133	A	C2'-C1'	-8.32	1.44	1.53
34	BA	1413	G	C5-C4	-8.32	1.32	1.38
35	BB	377	A	O3'-P	-8.32	1.51	1.61
35	BB	964	G	P-O5'	-8.32	1.51	1.59
36	BC	113	G	O3'-P	-8.32	1.51	1.61
40	BG	55	A	C1'-N9	-8.32	1.35	1.46
40	BG	144	G	P-O5'	-8.32	1.51	1.59
85	AA	1194	U	C2'-C1'	-8.32	1.44	1.53
34	BA	661	C	C2'-C1'	-8.32	1.44	1.53
37	BD	62	A	N9-C4	-8.32	1.32	1.37
85	AA	2200	A	O3'-P	-8.32	1.51	1.61
85	AA	841	U	P-O5'	-8.32	1.51	1.59
34	BA	566	G	C2'-C1'	-8.32	1.44	1.53
34	BA	613	A	P-O5'	-8.32	1.51	1.59
35	BB	1045	G	N9-C8	-8.32	1.32	1.37
85	AA	65	A	C2'-C1'	-8.32	1.44	1.53
85	AA	1106	A	N9-C4	-8.32	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1546	G	P-O5'	-8.32	1.51	1.59
34	BA	696	A	C1'-N9	-8.31	1.35	1.46
34	BA	747	G	N7-C5	-8.31	1.34	1.39
34	BA	794	G	P-O5'	-8.31	1.51	1.59
35	BB	888	U	P-O5'	-8.31	1.51	1.59
85	AA	37	U	C2'-C1'	-8.31	1.44	1.53
34	BA	1322	A	C3'-C2'	-8.31	1.43	1.52
35	BB	1473	U	P-O5'	-8.31	1.51	1.59
41	BH	38	G	C5-C4	-8.31	1.32	1.38
85	AA	1004	G	P-O5'	-8.31	1.51	1.59
85	AA	1136	A	O3'-P	-8.31	1.51	1.61
34	BA	123	C	O3'-P	-8.31	1.51	1.61
34	BA	140	C	N3-C4	8.31	1.39	1.33
34	BA	966	G	C1'-N9	-8.31	1.35	1.46
35	BB	440	U	C2'-C1'	-8.31	1.44	1.53
85	AA	717	G	P-O5'	-8.31	1.51	1.59
40	BG	174	G	N7-C5	-8.31	1.34	1.39
85	AA	800	A	O3'-P	-8.31	1.51	1.61
85	AA	933	U	O3'-P	-8.31	1.51	1.61
85	AA	1204	A	C3'-C2'	-8.31	1.43	1.52
34	BA	1226	G	O3'-P	-8.31	1.51	1.61
35	BB	562	A	N9-C4	-8.31	1.32	1.37
35	BB	810	G	C1'-N9	-8.31	1.35	1.46
34	BA	237	A	C1'-N9	-8.31	1.35	1.46
34	BA	1799	G	O3'-P	-8.31	1.51	1.61
35	BB	1206	G	C6-N1	-8.31	1.33	1.39
35	BB	401	U	C3'-C2'	-8.31	1.43	1.52
35	BB	694	C	N3-C4	-8.31	1.28	1.33
38	BE	8	G	N7-C5	-8.31	1.34	1.39
85	AA	772	C	P-O5'	-8.31	1.51	1.59
34	BA	1258	G	C2'-C1'	-8.30	1.44	1.53
85	AA	823	C	C2'-C1'	-8.30	1.44	1.53
85	AA	1190	G	O3'-P	-8.30	1.51	1.61
36	BC	70	C	C2'-C1'	-8.30	1.44	1.53
34	BA	45	A	C2'-C1'	-8.30	1.44	1.53
34	BA	471	U	O3'-P	-8.30	1.51	1.61
34	BA	490	A	C1'-N9	-8.30	1.35	1.46
34	BA	623	U	C4'-C3'	-8.30	1.44	1.53
34	BA	878	G	C2'-C1'	-8.30	1.44	1.53
34	BA	922	C	C2'-C1'	-8.30	1.44	1.53
34	BA	1288	U	C2'-C1'	-8.30	1.44	1.53
35	BB	816	U	O3'-P	-8.30	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	376	C	P-O5'	-8.30	1.51	1.59
40	BG	43	U	C2'-C1'	-8.30	1.44	1.53
85	AA	456	A	C4'-C3'	-8.30	1.44	1.53
34	BA	401	A	C5'-C4'	-8.30	1.41	1.51
34	BA	418	G	C2-N2	-8.30	1.26	1.34
34	BA	915	A	C2'-C1'	-8.30	1.44	1.53
34	BA	1011	G	C2-N2	-8.30	1.26	1.34
34	BA	1086	A	C2'-C1'	-8.30	1.44	1.53
34	BA	1580	U	C4'-C3'	-8.30	1.44	1.53
40	BG	45	G	C2'-C1'	-8.30	1.44	1.53
38	BE	148	C	C2'-C1'	-8.30	1.44	1.53
40	BG	95	U	C4'-O4'	-8.30	1.34	1.45
85	AA	438	G	C1'-N9	-8.30	1.35	1.46
85	AA	1522	U	C2'-C1'	-8.30	1.44	1.53
85	AA	2172	A	C3'-C2'	-8.30	1.43	1.52
37	BD	60	C	C2'-C1'	-8.30	1.44	1.53
85	AA	1229	G	O3'-P	-8.30	1.51	1.61
34	BA	193	C	C2-N3	-8.29	1.29	1.35
34	BA	1332	U	C2-N3	-8.29	1.31	1.37
35	BB	687	C	C2'-C1'	-8.29	1.44	1.53
35	BB	1509	G	P-O5'	-8.29	1.51	1.59
37	BD	52	U	P-O5'	-8.30	1.51	1.59
38	BE	144	A	N9-C4	-8.29	1.32	1.37
41	BH	31	A	N7-C5	-8.30	1.34	1.39
85	AA	1091	C	P-O5'	-8.29	1.51	1.59
34	BA	1194	G	P-O5'	-8.29	1.51	1.59
34	BA	1323	G	C3'-C2'	-8.29	1.43	1.52
34	BA	1431	G	C2'-C1'	-8.29	1.44	1.53
85	AA	1261	U	C3'-C2'	-8.29	1.43	1.52
34	BA	1225	A	N9-C4	-8.29	1.32	1.37
35	BB	779	C	P-O5'	-8.29	1.51	1.59
41	BH	123	G	O3'-P	-8.29	1.51	1.61
85	AA	2209	U	C3'-C2'	-8.29	1.43	1.52
35	BB	1025	A	C4'-O4'	8.29	1.56	1.45
37	BD	105	G	C3'-C2'	-8.29	1.43	1.52
40	BG	46	G	N9-C4	-8.29	1.31	1.38
41	BH	51	C	C2-N3	-8.29	1.29	1.35
85	AA	1967	A	N9-C4	-8.29	1.32	1.37
85	AA	2072	G	C1'-N9	-8.29	1.35	1.46
85	AA	2215	C	C2'-C1'	-8.29	1.44	1.53
85	AA	2172	A	N7-C5	-8.29	1.34	1.39
34	BA	1003	A	C5-C4	-8.29	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1259	C	C3'-C2'	-8.29	1.43	1.52
35	BB	16	G	O3'-P	-8.29	1.51	1.61
35	BB	799	A	C6-N6	-8.29	1.27	1.33
35	BB	808	U	C3'-C2'	-8.29	1.43	1.52
35	BB	967	G	C2'-C1'	-8.29	1.44	1.53
36	BC	37	U	C3'-C2'	-8.29	1.43	1.52
36	BC	132	U	P-O5'	-8.29	1.51	1.59
37	BD	1	G	C2'-C1'	-8.29	1.44	1.53
85	AA	366	A	N7-C5	-8.28	1.34	1.39
34	BA	333	A	C8-N7	-8.28	1.25	1.31
34	BA	726	G	N1-C2	-8.28	1.31	1.37
34	BA	801	U	C2'-C1'	-8.28	1.44	1.53
35	BB	400	C	O3'-P	-8.28	1.51	1.61
85	AA	166	C	C2'-C1'	-8.28	1.44	1.53
38	BE	207	G	O3'-P	-8.28	1.51	1.61
34	BA	763	U	C3'-C2'	8.28	1.62	1.52
34	BA	1838	U	C2-N3	-8.28	1.31	1.37
35	BB	525	U	P-O5'	-8.28	1.51	1.59
35	BB	535	U	C2-N3	-8.28	1.31	1.37
35	BB	1406	C	C2'-C1'	-8.28	1.44	1.53
38	BE	5	A	N3-C4	-8.28	1.29	1.34
85	AA	1681	G	P-O5'	-8.28	1.51	1.59
85	AA	1964	A	P-O5'	-8.28	1.51	1.59
34	BA	399	G	O3'-P	-8.28	1.51	1.61
34	BA	794	G	C1'-N9	-8.28	1.35	1.46
34	BA	1291	A	O3'-P	-8.28	1.51	1.61
35	BB	26	C	O3'-P	-8.28	1.51	1.61
35	BB	430	A	C1'-N9	-8.28	1.35	1.46
35	BB	574	G	C1'-N9	-8.28	1.35	1.46
35	BB	879	G	O3'-P	-8.28	1.51	1.61
34	BA	98	A	C3'-C2'	-8.27	1.43	1.52
34	BA	1555	G	N3-C4	-8.27	1.29	1.35
34	BA	1203	G	C2'-C1'	-8.27	1.44	1.53
34	BA	1224	A	C4'-C3'	-8.27	1.44	1.53
34	BA	1442	A	P-O5'	-8.27	1.51	1.59
34	BA	1611	A	N7-C5	-8.27	1.34	1.39
34	BA	1641	G	C5-C4	-8.27	1.32	1.38
35	BB	610	U	C2-N3	-8.27	1.31	1.37
35	BB	647	U	P-O5'	-8.27	1.51	1.59
85	AA	699	U	C3'-C2'	-8.27	1.43	1.52
38	BE	101	C	C1'-N1	-8.27	1.35	1.46
38	BE	117	A	C5'-C4'	-8.27	1.41	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	141	A	P-O5'	-8.27	1.51	1.59
34	BA	50	G	N1-C2	-8.27	1.31	1.37
35	BB	817	C	O3'-P	-8.27	1.51	1.61
35	BB	1370	G	O3'-P	-8.27	1.51	1.61
38	BE	116	U	C4'-C3'	-8.27	1.44	1.53
85	AA	2222	G	C2'-C1'	-8.27	1.44	1.53
34	BA	334	G	C3'-C2'	-8.27	1.43	1.52
34	BA	1496	G	C1'-N9	-8.27	1.35	1.46
35	BB	490	G	N7-C5	-8.27	1.34	1.39
85	AA	82	A	O3'-P	-8.27	1.51	1.61
85	AA	582	A	O3'-P	-8.27	1.51	1.61
35	BB	461	U	P-O5'	-8.27	1.51	1.59
36	BC	152	C	P-O5'	-8.27	1.51	1.59
85	AA	1124	G	C2'-C1'	-8.27	1.44	1.53
85	AA	1724	A	P-O5'	-8.27	1.51	1.59
34	BA	54	A	O3'-P	-8.26	1.51	1.61
34	BA	142	A	P-O5'	-8.26	1.51	1.59
40	BG	68	U	P-O5'	-8.26	1.51	1.59
41	BH	39	G	C1'-N9	-8.26	1.35	1.46
34	BA	385	U	O3'-P	-8.26	1.51	1.61
34	BA	960	C	C3'-C2'	-8.26	1.43	1.52
34	BA	1275	G	C3'-C2'	-8.26	1.43	1.52
34	BA	1702	G	C2'-C1'	-8.26	1.44	1.53
35	BB	628	A	O3'-P	-8.26	1.51	1.61
40	BG	164	U	P-O5'	-8.26	1.51	1.59
85	AA	2130	G	N7-C5	-8.26	1.34	1.39
34	BA	936	A	O3'-P	-8.26	1.51	1.61
34	BA	1109	G	O3'-P	-8.26	1.51	1.61
34	BA	1506	C	C2'-C1'	-8.26	1.44	1.53
35	BB	868	C	O3'-P	-8.26	1.51	1.61
35	BB	1388	A	C3'-C2'	-8.26	1.43	1.52
85	AA	432	A	N9-C4	-8.26	1.32	1.37
85	AA	465	A	N9-C4	-8.26	1.32	1.37
34	BA	566	G	N9-C4	-8.26	1.31	1.38
34	BA	1027	C	C3'-C2'	-8.26	1.43	1.52
35	BB	465	C	C2'-C1'	-8.26	1.44	1.53
34	BA	1011	G	C1'-N9	-8.26	1.35	1.46
34	BA	1821	A	N9-C4	-8.26	1.32	1.37
35	BB	555	G	C3'-C2'	-8.26	1.43	1.52
35	BB	814	A	C3'-C2'	-8.26	1.43	1.52
35	BB	1267	C	O3'-P	-8.26	1.51	1.61
85	AA	403	G	C1'-N9	-8.26	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	704	A	O3'-P	-8.26	1.51	1.61
85	AA	877	G	O3'-P	-8.26	1.51	1.61
85	AA	1501	A	O3'-P	-8.26	1.51	1.61
85	AA	1111	A	N9-C4	-8.26	1.32	1.37
34	BA	67	A	O3'-P	-8.25	1.51	1.61
34	BA	209	A	C3'-C2'	-8.25	1.43	1.52
34	BA	253	U	P-O5'	-8.25	1.51	1.59
34	BA	626	G	C2'-C1'	-8.25	1.44	1.53
36	BC	30	U	P-O5'	-8.25	1.51	1.59
34	BA	1730	A	C2'-C1'	-8.25	1.44	1.53
35	BB	387	G	C6-N1	-8.25	1.33	1.39
85	AA	373	G	C6-N1	-8.25	1.33	1.39
85	AA	461	G	N7-C5	-8.25	1.34	1.39
85	AA	640	C	C2'-C1'	-8.25	1.44	1.53
85	AA	867	G	N9-C4	-8.25	1.31	1.38
85	AA	1532	G	P-O5'	-8.25	1.51	1.59
34	BA	398	G	C3'-C2'	-8.25	1.43	1.52
34	BA	588	C	P-O5'	-8.25	1.51	1.59
34	BA	708	C	C3'-C2'	-8.25	1.43	1.52
34	BA	406	G	C5-C4	-8.25	1.32	1.38
34	BA	955	G	N9-C4	-8.25	1.31	1.38
34	BA	1213	A	C5-C4	-8.25	1.32	1.38
35	BB	58	G	C5-C4	-8.25	1.32	1.38
35	BB	1142	C	C3'-C2'	-8.25	1.43	1.52
35	BB	570	A	P-O5'	-8.25	1.51	1.59
35	BB	1315	C	O3'-P	-8.25	1.51	1.61
35	BB	1387	C	P-O5'	-8.25	1.51	1.59
85	AA	1237	A	O3'-P	-8.25	1.51	1.61
85	AA	1610	G	P-O5'	-8.25	1.51	1.59
34	BA	19	G	C6-N1	-8.25	1.33	1.39
37	BD	104	C	C2'-C1'	-8.25	1.44	1.53
85	AA	1205	U	P-O5'	-8.25	1.51	1.59
34	BA	974	G	C2'-C1'	-8.25	1.44	1.53
34	BA	1233	U	P-O5'	-8.25	1.51	1.59
34	BA	1261	G	C6-N1	-8.25	1.33	1.39
34	BA	1525	G	C6-N1	-8.25	1.33	1.39
34	BA	1846	G	C2'-C1'	-8.25	1.44	1.53
35	BB	1050	A	O3'-P	-8.25	1.51	1.61
36	BC	65	G	C1'-N9	-8.25	1.35	1.46
36	BC	114	C	C2-N3	-8.25	1.29	1.35
40	BG	50	G	C2-N2	-8.25	1.26	1.34
85	AA	2233	A	N9-C4	-8.25	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	13	U	C2-N3	-8.24	1.31	1.37
35	BB	979	G	O3'-P	-8.24	1.51	1.61
34	BA	444	A	N9-C4	-8.24	1.32	1.37
34	BA	1459	U	O3'-P	-8.24	1.51	1.61
35	BB	975	G	N9-C8	-8.24	1.32	1.37
36	BC	51	A	N7-C5	-8.24	1.34	1.39
85	AA	156	G	P-O5'	-8.24	1.51	1.59
85	AA	1916	A	N3-C4	-8.24	1.29	1.34
85	AA	2143	U	C2-N3	-8.24	1.31	1.37
35	BB	700	C	O3'-P	-8.24	1.51	1.61
85	AA	1505	G	P-O5'	-8.24	1.51	1.59
34	BA	194	G	C2'-C1'	-8.24	1.44	1.53
34	BA	417	A	C1'-N9	-8.24	1.35	1.46
34	BA	529	A	N7-C5	-8.24	1.34	1.39
34	BA	892	C	C3'-C2'	-8.24	1.43	1.52
85	AA	544	A	O3'-P	-8.24	1.51	1.61
34	BA	754	G	O3'-P	-8.24	1.51	1.61
34	BA	1235	C	C2-N3	-8.24	1.29	1.35
34	BA	1654	G	C1'-N9	-8.24	1.35	1.46
34	BA	1720	U	O3'-P	-8.24	1.51	1.61
35	BB	539	G	O3'-P	-8.24	1.51	1.61
38	BE	142	A	C2'-C1'	-8.24	1.44	1.53
41	BH	2	U	O3'-P	-8.24	1.51	1.61
34	BA	80	U	P-O5'	-8.24	1.51	1.59
34	BA	1587	C	P-O5'	-8.24	1.51	1.59
36	BC	82	C	O3'-P	-8.24	1.51	1.61
85	AA	996	A	N9-C4	-8.24	1.32	1.37
37	BD	30	A	N9-C4	-8.24	1.32	1.37
40	BG	70	C	C2'-C1'	-8.24	1.44	1.53
85	AA	664	C	O3'-P	-8.24	1.51	1.61
85	AA	1286	C	C2-N3	-8.24	1.29	1.35
34	BA	1640	G	C3'-C2'	-8.23	1.43	1.52
36	BC	151	G	O3'-P	-8.23	1.51	1.61
85	AA	39	A	N9-C4	-8.23	1.32	1.37
85	AA	601	A	N9-C4	-8.23	1.32	1.37
85	AA	2187	G	O3'-P	-8.23	1.51	1.61
34	BA	12	G	O3'-P	-8.23	1.51	1.61
34	BA	789	U	O3'-P	-8.23	1.51	1.61
34	BA	924	U	N3-C4	-8.23	1.31	1.38
34	BA	940	C	O3'-P	-8.23	1.51	1.61
35	BB	1252	G	N7-C5	-8.23	1.34	1.39
34	BA	1049	G	N7-C5	-8.23	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1195	A	N7-C5	-8.23	1.34	1.39
35	BB	1258	G	N7-C5	-8.23	1.34	1.39
36	BC	151	G	N3-C4	-8.23	1.29	1.35
34	BA	1231	C	P-O5'	-8.23	1.51	1.59
34	BA	1591	G	O3'-P	-8.23	1.51	1.61
34	BA	1726	U	P-O5'	-8.23	1.51	1.59
35	BB	1312	U	C1'-N1	-8.23	1.35	1.46
36	BC	121	G	C2'-C1'	-8.23	1.44	1.53
34	BA	29	U	C2'-C1'	-8.23	1.44	1.53
34	BA	139	U	O3'-P	-8.23	1.51	1.61
34	BA	290	G	C1'-N9	-8.23	1.35	1.46
34	BA	890	G	N1-C2	-8.23	1.31	1.37
35	BB	1127	A	N7-C5	-8.23	1.34	1.39
35	BB	1311	G	C6-N1	-8.23	1.33	1.39
85	AA	420	C	O3'-P	-8.23	1.51	1.61
85	AA	584	G	C8-N7	-8.23	1.26	1.30
85	AA	616	A	C2'-C1'	-8.23	1.44	1.53
34	BA	807	U	C4'-O4'	-8.22	1.34	1.45
34	BA	1333	G	O3'-P	-8.22	1.51	1.61
34	BA	1465	C	C2'-C1'	-8.22	1.44	1.53
34	BA	1708	A	C5-C4	-8.22	1.32	1.38
35	BB	650	A	N9-C4	-8.22	1.32	1.37
35	BB	1185	G	P-O5'	-8.22	1.51	1.59
34	BA	1802	C	O3'-P	-8.22	1.51	1.61
37	BD	81	C	C2'-C1'	-8.22	1.44	1.53
85	AA	1449	C	P-O5'	-8.22	1.51	1.59
85	AA	1517	G	P-O5'	-8.22	1.51	1.59
34	BA	447	U	C3'-C2'	-8.22	1.43	1.52
34	BA	826	C	C2-N3	-8.22	1.29	1.35
34	BA	1607	U	O3'-P	-8.22	1.51	1.61
35	BB	94	A	N7-C5	-8.22	1.34	1.39
37	BD	99	G	N7-C5	-8.22	1.34	1.39
34	BA	1119	A	C3'-C2'	-8.22	1.43	1.52
34	BA	1684	A	C3'-C2'	-8.22	1.43	1.52
35	BB	353	G	N9-C4	-8.22	1.31	1.38
35	BB	1228	A	C5'-C4'	8.22	1.61	1.51
41	BH	29	G	C3'-C2'	-8.22	1.43	1.52
85	AA	157	G	N9-C4	-8.22	1.31	1.38
34	BA	267	G	C1'-N9	-8.22	1.35	1.46
34	BA	956	G	P-O5'	-8.22	1.51	1.59
35	BB	1404	A	N7-C5	-8.22	1.34	1.39
34	BA	322	U	C3'-C2'	-8.22	1.43	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	432	A	O3'-P	-8.22	1.51	1.61
34	BA	1532	G	N1-C2	-8.22	1.31	1.37
34	BA	1839	G	O3'-P	-8.22	1.51	1.61
35	BB	616	U	O3'-P	-8.22	1.51	1.61
37	BD	41	G	C1'-N9	-8.22	1.35	1.46
85	AA	962	U	C3'-C2'	-8.22	1.43	1.52
34	BA	1023	G	C2'-C1'	-8.22	1.44	1.53
34	BA	1474	G	N7-C5	-8.21	1.34	1.39
35	BB	398	A	C1'-N9	-8.22	1.35	1.46
35	BB	1342	C	C1'-N1	-8.22	1.35	1.46
35	BB	672	C	P-O5'	-8.21	1.51	1.59
35	BB	1278	A	O3'-P	-8.21	1.51	1.61
85	AA	195	C	P-O5'	-8.21	1.51	1.59
85	AA	491	G	C2'-C1'	-8.21	1.44	1.53
85	AA	820	G	N9-C4	8.21	1.44	1.38
85	AA	962	U	C2-N3	-8.21	1.31	1.37
85	AA	674	U	C2'-C1'	-8.21	1.44	1.53
34	BA	8	G	C1'-N9	-8.21	1.35	1.46
35	BB	370	A	O3'-P	-8.21	1.51	1.61
35	BB	612	A	C1'-N9	-8.21	1.35	1.46
38	BE	8	G	N9-C8	-8.21	1.32	1.37
38	BE	124	G	C2'-C1'	-8.21	1.44	1.53
40	BG	2	U	C3'-C2'	-8.21	1.43	1.52
40	BG	51	U	C2'-C1'	-8.21	1.44	1.53
40	BG	86	U	O3'-P	-8.21	1.51	1.61
40	BG	154	C	C2'-C1'	-8.21	1.44	1.53
85	AA	267	U	C1'-N1	8.21	1.61	1.48
85	AA	336	C	C3'-C2'	-8.21	1.43	1.52
85	AA	672	U	O3'-P	-8.21	1.51	1.61
85	AA	765	U	P-O5'	-8.21	1.51	1.59
34	BA	117	C	N1-C6	-8.21	1.32	1.37
34	BA	239	C	C3'-C2'	-8.21	1.43	1.52
34	BA	347	A	P-O5'	-8.21	1.51	1.59
34	BA	1465	C	C3'-C2'	-8.21	1.43	1.52
34	BA	1689	U	P-O5'	-8.21	1.51	1.59
35	BB	1046	C	O3'-P	-8.21	1.51	1.61
35	BB	1525	G	C2'-C1'	-8.21	1.44	1.53
36	BC	100	U	C2-N3	-8.21	1.32	1.37
85	AA	7	G	O3'-P	-8.21	1.51	1.61
85	AA	352	G	C6-N1	-8.21	1.33	1.39
85	AA	383	C	C2-N3	-8.21	1.29	1.35
85	AA	1729	C	O3'-P	-8.21	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2213	A	C3'-C2'	-8.21	1.43	1.52
35	BB	1431	G	C5-C4	-8.21	1.32	1.38
34	BA	175	G	P-O5'	-8.20	1.51	1.59
34	BA	948	C	O3'-P	-8.21	1.51	1.61
35	BB	1522	G	C3'-C2'	-8.21	1.43	1.52
36	BC	140	U	P-O5'	-8.21	1.51	1.59
85	AA	1992	A	N7-C5	-8.21	1.34	1.39
34	BA	1316	G	P-O5'	-8.20	1.51	1.59
35	BB	127	U	C3'-C2'	-8.20	1.43	1.52
35	BB	1176	G	O3'-P	-8.20	1.51	1.61
35	BB	1259	A	C2'-C1'	-8.20	1.44	1.53
34	BA	726	G	P-O5'	-8.20	1.51	1.59
34	BA	1223	C	C4'-C3'	-8.20	1.44	1.53
85	AA	978	U	P-O5'	-8.20	1.51	1.59
85	AA	1891	U	C2-N3	-8.20	1.32	1.37
34	BA	72	U	O3'-P	-8.20	1.51	1.61
34	BA	323	C	P-O5'	-8.20	1.51	1.59
35	BB	1260	A	C1'-N9	-8.20	1.35	1.46
34	BA	300	C	C2-N3	-8.20	1.29	1.35
34	BA	412	G	C2'-C1'	-8.20	1.44	1.53
34	BA	1597	G	N9-C8	-8.20	1.32	1.37
34	BA	1792	U	C3'-C2'	-8.20	1.43	1.52
38	BE	125	C	C3'-C2'	-8.20	1.43	1.52
40	BG	31	G	O3'-P	-8.20	1.51	1.61
85	AA	397	G	C3'-C2'	-8.20	1.43	1.52
34	BA	367	G	N7-C5	-8.20	1.34	1.39
85	AA	2004	U	O3'-P	-8.20	1.51	1.61
38	BE	111	C	C4'-C3'	-8.19	1.44	1.53
39	BF	61	A	C2'-C1'	-8.20	1.44	1.53
85	AA	2219	G	O3'-P	-8.19	1.51	1.61
34	BA	1068	C	N3-C4	-8.19	1.28	1.33
34	BA	1663	U	P-O5'	-8.19	1.51	1.59
85	AA	478	U	O3'-P	-8.19	1.51	1.61
34	BA	4	A	C1'-N9	-8.19	1.35	1.46
34	BA	68	A	C2'-C1'	-8.19	1.44	1.53
38	BE	139	U	C5'-C4'	-8.19	1.41	1.51
34	BA	282	A	N3-C4	-8.19	1.29	1.34
34	BA	1702	G	C3'-C2'	-8.19	1.43	1.52
35	BB	662	G	C5-C4	-8.19	1.32	1.38
35	BB	619	A	N9-C4	-8.19	1.32	1.37
34	BA	506	U	O3'-P	-8.19	1.51	1.61
34	BA	812	A	O3'-P	-8.19	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1492	G	C2-N2	-8.19	1.26	1.34
35	BB	53	C	C2'-C1'	-8.19	1.44	1.53
35	BB	392	G	O3'-P	-8.19	1.51	1.61
35	BB	423	G	O3'-P	-8.19	1.51	1.61
85	AA	996	A	O3'-P	-8.19	1.51	1.61
85	AA	1500	C	C2'-C1'	-8.19	1.44	1.53
34	BA	727	G	C6-N1	-8.19	1.33	1.39
35	BB	578	G	O3'-P	-8.19	1.51	1.61
36	BC	38	U	C2'-C1'	-8.19	1.44	1.53
85	AA	413	G	C6-N1	-8.19	1.33	1.39
85	AA	1889	U	P-O5'	-8.19	1.51	1.59
34	BA	701	G	O3'-P	-8.19	1.51	1.61
34	BA	1687	A	C1'-N9	-8.19	1.35	1.46
35	BB	687	C	C1'-N1	-8.19	1.35	1.46
35	BB	1082	A	C2'-C1'	-8.19	1.44	1.53
34	BA	395	G	C2-N2	-8.18	1.26	1.34
34	BA	1823	A	P-O5'	-8.18	1.51	1.59
37	BD	73	U	N1-C2	-8.18	1.31	1.38
35	BB	1143	A	O3'-P	-8.18	1.51	1.61
39	BF	52	A	P-O5'	-8.18	1.51	1.59
40	BG	173	C	C2'-C1'	-8.18	1.44	1.53
85	AA	424	A	N7-C5	-8.18	1.34	1.39
85	AA	199	U	O3'-P	-8.18	1.51	1.61
85	AA	373	G	N9-C4	-8.18	1.31	1.38
85	AA	629	A	C2'-C1'	-8.18	1.44	1.53
85	AA	910	G	C2-N2	-8.18	1.26	1.34
85	AA	1835	U	C2-N3	-8.18	1.32	1.37
34	BA	1022	C	C2'-C1'	-8.18	1.44	1.53
35	BB	459	U	C3'-C2'	-8.18	1.43	1.52
85	AA	887	A	C5-C4	-8.18	1.33	1.38
85	AA	1491	G	P-O5'	-8.18	1.51	1.59
34	BA	420	A	C3'-C2'	-8.18	1.43	1.52
34	BA	696	A	C3'-C2'	-8.18	1.43	1.52
34	BA	1017	C	C2-N3	-8.18	1.29	1.35
34	BA	1495	A	C4'-C3'	-8.18	1.44	1.53
34	BA	1640	G	C1'-N9	-8.18	1.35	1.46
34	BA	1732	A	C8-N7	-8.18	1.25	1.31
35	BB	28	G	N7-C5	-8.18	1.34	1.39
35	BB	501	G	C1'-N9	-8.18	1.35	1.46
38	BE	96	G	C3'-C2'	-8.18	1.43	1.52
85	AA	376	C	C3'-C2'	-8.18	1.43	1.52
85	AA	386	G	N7-C5	-8.18	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	680	U	O3'-P	-8.18	1.51	1.61
34	BA	732	A	C5-C4	-8.18	1.33	1.38
35	BB	428	G	C2-N2	-8.18	1.26	1.34
38	BE	140	G	C6-N1	-8.18	1.33	1.39
35	BB	1238	A	O3'-P	-8.18	1.51	1.61
40	BG	17	A	O3'-P	-8.18	1.51	1.61
85	AA	1484	G	P-O5'	-8.18	1.51	1.59
40	BG	103	C	C2-N3	-8.18	1.29	1.35
35	BB	125	G	N7-C5	-8.17	1.34	1.39
34	BA	439	A	N9-C4	-8.17	1.32	1.37
35	BB	1128	U	O3'-P	-8.17	1.51	1.61
37	BD	80	G	C3'-C2'	-8.17	1.43	1.52
39	BF	72	A	O3'-P	-8.17	1.51	1.61
41	BH	15	A	N9-C4	-8.17	1.32	1.37
85	AA	1098	C	C4'-C3'	-8.17	1.44	1.53
34	BA	1427	U	O3'-P	-8.17	1.51	1.61
34	BA	1637	G	N7-C5	-8.17	1.34	1.39
85	AA	419	A	N7-C5	-8.17	1.34	1.39
85	AA	2183	U	C3'-C2'	-8.17	1.43	1.52
40	BG	172	C	O3'-P	-8.17	1.51	1.61
34	BA	810	A	C1'-N9	-8.17	1.35	1.46
34	BA	956	G	N7-C5	-8.17	1.34	1.39
35	BB	1096	G	C2'-C1'	-8.17	1.44	1.53
35	BB	1243	A	C1'-N9	-8.17	1.35	1.46
85	AA	313	A	C5-C4	-8.17	1.33	1.38
85	AA	1114	A	N3-C4	-8.17	1.29	1.34
85	AA	1120	G	C2'-C1'	-8.17	1.44	1.53
34	BA	195	G	O3'-P	-8.16	1.51	1.61
34	BA	1110	A	C1'-N9	-8.16	1.35	1.46
85	AA	1199	C	P-O5'	-8.16	1.51	1.59
34	BA	215	C	C2'-C1'	-8.16	1.44	1.53
34	BA	920	U	C2-N3	-8.16	1.32	1.37
35	BB	1467	A	C5'-C4'	8.16	1.61	1.51
36	BC	123	G	N3-C4	-8.16	1.29	1.35
85	AA	138	C	C2'-C1'	-8.16	1.44	1.53
85	AA	2196	G	C6-N1	-8.16	1.33	1.39
85	AA	2072	G	C2-N2	-8.16	1.26	1.34
34	BA	99	G	O3'-P	-8.16	1.51	1.61
34	BA	362	G	C1'-N9	-8.16	1.35	1.46
34	BA	575	U	C2'-C1'	-8.16	1.44	1.53
34	BA	575	U	P-O5'	-8.16	1.51	1.59
35	BB	813	C	C2'-C1'	-8.16	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1361	A	C2'-C1'	-8.16	1.44	1.53
40	BG	31	G	P-O5'	-8.16	1.51	1.59
35	BB	1391	G	O3'-P	-8.16	1.51	1.61
37	BD	41	G	O3'-P	-8.16	1.51	1.61
85	AA	340	G	C3'-C2'	-8.16	1.43	1.52
85	AA	476	C	C3'-C2'	-8.16	1.43	1.52
34	BA	281	C	O3'-P	-8.16	1.51	1.61
34	BA	711	C	C2-N3	-8.16	1.29	1.35
34	BA	1040	G	O3'-P	-8.16	1.51	1.61
35	BB	17	U	O3'-P	-8.16	1.51	1.61
34	BA	1603	A	C2'-C1'	-8.16	1.44	1.53
35	BB	1258	G	C2'-C1'	-8.16	1.44	1.53
38	BE	179	A	O3'-P	-8.16	1.51	1.61
85	AA	457	G	C1'-N9	-8.16	1.35	1.46
34	BA	1002	U	O3'-P	-8.15	1.51	1.61
35	BB	102	G	C2'-C1'	-8.15	1.44	1.53
35	BB	506	G	C1'-N9	-8.15	1.35	1.46
35	BB	563	A	C2'-C1'	-8.15	1.44	1.53
41	BH	16	A	C3'-C2'	-8.15	1.43	1.52
85	AA	1153	G	C8-N7	-8.15	1.26	1.30
34	BA	115	U	C2-N3	-8.15	1.32	1.37
34	BA	1563	G	C2-N2	-8.15	1.26	1.34
35	BB	792	G	N9-C4	-8.15	1.31	1.38
85	AA	1655	G	O3'-P	-8.15	1.51	1.61
34	BA	746	C	P-O5'	-8.15	1.51	1.59
34	BA	1694	C	C3'-C2'	-8.15	1.43	1.52
38	BE	52	U	P-O5'	-8.15	1.51	1.59
34	BA	112	C	O3'-P	-8.15	1.51	1.61
34	BA	944	G	C3'-C2'	-8.15	1.43	1.52
35	BB	502	C	C2-N3	-8.15	1.29	1.35
35	BB	648	G	N7-C5	-8.15	1.34	1.39
35	BB	976	U	P-O5'	-8.15	1.51	1.59
41	BH	6	U	C3'-C2'	-8.15	1.43	1.52
36	BC	11	G	O3'-P	-8.15	1.51	1.61
36	BC	44	A	C1'-N9	-8.15	1.35	1.46
36	BC	141	C	C2'-C1'	-8.15	1.44	1.53
38	BE	50	G	O3'-P	-8.15	1.51	1.61
85	AA	190	A	C1'-N9	-8.15	1.35	1.46
85	AA	207	G	P-O5'	-8.15	1.51	1.59
85	AA	591	A	P-O5'	-8.15	1.51	1.59
85	AA	2219	G	P-O5'	-8.15	1.51	1.59
34	BA	697	A	N7-C5	-8.15	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1034	U	C3'-C2'	-8.15	1.43	1.52
35	BB	33	A	C1'-N9	-8.15	1.35	1.46
35	BB	1049	G	C6-N1	-8.15	1.33	1.39
35	BB	1398	A	C1'-N9	-8.15	1.35	1.46
40	BG	155	A	N9-C4	-8.15	1.32	1.37
85	AA	367	A	N3-C4	-8.15	1.29	1.34
41	BH	46	C	C3'-C2'	-8.15	1.43	1.52
85	AA	474	C	C2-N3	-8.15	1.29	1.35
85	AA	1130	G	C6-N1	-8.15	1.33	1.39
85	AA	1272	G	N7-C5	-8.15	1.34	1.39
85	AA	1421	U	O3'-P	-8.15	1.51	1.61
34	BA	1194	G	C2-N3	-8.14	1.26	1.32
35	BB	123	U	P-O5'	-8.14	1.51	1.59
85	AA	86	G	N9-C4	-8.14	1.31	1.38
85	AA	107	A	O3'-P	-8.14	1.51	1.61
85	AA	2171	A	C2'-C1'	-8.14	1.44	1.53
34	BA	1026	C	C3'-C2'	-8.14	1.43	1.52
35	BB	25	A	N9-C4	-8.14	1.32	1.37
34	BA	1693	U	P-O5'	-8.14	1.51	1.59
35	BB	54	U	C4'-C3'	-8.14	1.44	1.53
38	BE	8	G	C2'-C1'	-8.14	1.44	1.53
40	BG	24	A	C3'-O3'	-8.14	1.30	1.42
40	BG	59	G	O3'-P	-8.14	1.51	1.61
85	AA	152	A	C8-N7	-8.14	1.25	1.31
85	AA	469	G	N7-C5	8.14	1.44	1.39
34	BA	327	G	P-O5'	-8.14	1.51	1.59
35	BB	652	G	C2'-C1'	-8.14	1.44	1.53
34	BA	1101	A	C2'-C1'	-8.14	1.44	1.53
34	BA	1298	U	P-O5'	-8.14	1.51	1.59
38	BE	38	C	O3'-P	-8.14	1.51	1.61
85	AA	172	A	C1'-N9	-8.14	1.35	1.46
85	AA	457	G	P-O5'	-8.14	1.51	1.59
34	BA	79	C	C3'-C2'	-8.13	1.43	1.52
34	BA	273	G	C1'-N9	-8.13	1.35	1.46
34	BA	1240	G	O3'-P	-8.13	1.51	1.61
35	BB	677	U	C2-N3	-8.13	1.32	1.37
35	BB	1306	G	N1-C2	-8.13	1.31	1.37
35	BB	1313	C	O3'-P	-8.13	1.51	1.61
37	BD	74	A	C5-C4	-8.13	1.33	1.38
38	BE	93	U	O3'-P	-8.14	1.51	1.61
85	AA	16	G	C3'-C2'	-8.13	1.43	1.52
85	AA	101	C	P-O5'	-8.13	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	794	A	O3'-P	-8.13	1.51	1.61
85	AA	1631	C	P-O5'	-8.13	1.51	1.59
34	BA	259	C	O3'-P	-8.13	1.51	1.61
34	BA	1072	U	C2-N3	-8.13	1.32	1.37
34	BA	1490	U	C4'-C3'	8.13	1.62	1.53
35	BB	633	C	C2-N3	-8.13	1.29	1.35
34	BA	203	U	C2-N3	-8.13	1.32	1.37
34	BA	395	G	C4'-C3'	-8.13	1.44	1.53
34	BA	910	U	C3'-C2'	-8.13	1.43	1.52
34	BA	1543	A	O3'-P	-8.13	1.51	1.61
35	BB	375	G	C4'-C3'	-8.13	1.44	1.53
35	BB	416	U	P-O5'	-8.13	1.51	1.59
35	BB	1098	G	O3'-P	-8.13	1.51	1.61
35	BB	1170	U	C2-N3	-8.13	1.32	1.37
36	BC	40	A	C2'-C1'	-8.13	1.44	1.53
35	BB	1042	U	C5'-C4'	8.13	1.61	1.51
35	BB	1141	A	O3'-P	-8.13	1.51	1.61
37	BD	11	A	N9-C4	-8.13	1.32	1.37
85	AA	1235	G	N3-C4	-8.13	1.29	1.35
85	AA	1275	A	C1'-N9	-8.13	1.35	1.46
34	BA	97	A	N9-C4	-8.13	1.32	1.37
36	BC	76	C	C2'-C1'	-8.13	1.44	1.53
34	BA	291	C	C3'-C2'	-8.13	1.43	1.52
35	BB	96	A	P-O5'	-8.13	1.51	1.59
35	BB	1387	C	C3'-C2'	-8.13	1.43	1.52
85	AA	975	G	N9-C4	8.13	1.44	1.38
34	BA	786	U	P-O5'	-8.13	1.51	1.59
34	BA	1154	U	C2-N3	-8.13	1.32	1.37
35	BB	786	A	O3'-P	-8.13	1.51	1.61
35	BB	95	A	N9-C4	-8.12	1.32	1.37
35	BB	126	C	P-O5'	-8.12	1.51	1.59
35	BB	1185	G	C5-C4	-8.12	1.32	1.38
37	BD	47	U	C2-N3	-8.12	1.32	1.37
37	BD	79	G	O3'-P	-8.12	1.51	1.61
85	AA	100	A	O4'-C1'	-8.12	1.31	1.41
85	AA	543	A	C2'-C1'	-8.12	1.44	1.53
85	AA	1195	U	P-O5'	-8.12	1.51	1.59
34	BA	95	C	C2'-C1'	-8.12	1.44	1.53
34	BA	928	C	P-O5'	-8.12	1.51	1.59
34	BA	1837	U	C2-N3	-8.12	1.32	1.37
35	BB	1519	U	P-O5'	-8.12	1.51	1.59
36	BC	50	C	P-O5'	-8.12	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	147	G	C5-C4	-8.12	1.32	1.38
85	AA	36	U	P-O5'	-8.12	1.51	1.59
85	AA	399	A	P-O5'	-8.12	1.51	1.59
85	AA	461	G	P-O5'	-8.12	1.51	1.59
85	AA	556	C	C2'-C1'	-8.12	1.44	1.53
85	AA	689	U	O3'-P	-8.12	1.51	1.61
34	BA	1165	A	C5-C4	-8.12	1.33	1.38
35	BB	375	G	C2'-C1'	-8.12	1.44	1.53
35	BB	420	U	P-O5'	-8.12	1.51	1.59
35	BB	1288	G	C2'-C1'	-8.12	1.44	1.53
40	BG	130	G	O3'-P	-8.12	1.51	1.61
85	AA	473	C	C3'-C2'	-8.12	1.43	1.52
85	AA	1799	C	P-O5'	-8.12	1.51	1.59
34	BA	292	C	O3'-P	-8.12	1.51	1.61
34	BA	870	C	N3-C4	8.12	1.39	1.33
34	BA	1618	A	C2'-C1'	-8.12	1.44	1.53
35	BB	1079	G	C1'-N9	-8.12	1.35	1.46
35	BB	1443	C	O3'-P	-8.12	1.51	1.61
37	BD	32	A	C1'-N9	-8.12	1.35	1.46
38	BE	41	C	C2'-C1'	-8.12	1.44	1.53
39	BF	5	U	C2-N3	-8.12	1.32	1.37
85	AA	993	G	O3'-P	-8.12	1.51	1.61
86	AB	73	A	N9-C4	-8.12	1.32	1.37
85	AA	1660	U	C3'-C2'	-8.12	1.43	1.52
85	AA	1681	G	O3'-P	-8.12	1.51	1.61
34	BA	1647	G	C2'-C1'	-8.12	1.44	1.53
34	BA	344	G	C2'-C1'	-8.11	1.44	1.53
34	BA	855	C	O3'-P	-8.11	1.51	1.61
34	BA	897	U	C4'-C3'	-8.11	1.44	1.53
38	BE	21	C	C2'-C1'	-8.11	1.44	1.53
38	BE	180	G	C2-N3	-8.12	1.26	1.32
85	AA	1109	G	O3'-P	-8.12	1.51	1.61
85	AA	1861	A	O3'-P	-8.12	1.51	1.61
34	BA	101	G	C6-N1	-8.11	1.33	1.39
34	BA	1125	G	P-O5'	-8.11	1.51	1.59
36	BC	66	G	N9-C4	-8.11	1.31	1.38
85	AA	889	G	C2-N3	-8.11	1.26	1.32
85	AA	2146	G	O3'-P	-8.11	1.51	1.61
35	BB	1139	A	C2'-C1'	-8.11	1.44	1.53
34	BA	262	A	N7-C5	-8.11	1.34	1.39
34	BA	293	A	C2'-C1'	-8.11	1.44	1.53
34	BA	467	A	N9-C4	-8.11	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	506	U	C3'-C2'	-8.11	1.43	1.52
34	BA	1103	G	C1'-N9	-8.11	1.35	1.46
85	AA	489	C	C2'-C1'	-8.11	1.44	1.53
34	BA	1594	G	O3'-P	-8.11	1.51	1.61
34	BA	1814	U	C2'-C1'	-8.11	1.44	1.53
35	BB	1291	G	C6-N1	-8.11	1.33	1.39
85	AA	1970	A	C2'-C1'	-8.11	1.44	1.53
34	BA	19	G	C2'-C1'	-8.11	1.44	1.53
34	BA	313	C	C2'-C1'	-8.11	1.44	1.53
34	BA	1030	C	C2-N3	-8.11	1.29	1.35
34	BA	1437	G	O3'-P	-8.11	1.51	1.61
35	BB	114	A	N9-C4	-8.11	1.32	1.37
34	BA	1743	U	P-O5'	-8.11	1.51	1.59
35	BB	563	A	N9-C4	-8.11	1.32	1.37
35	BB	1149	A	O3'-P	-8.11	1.51	1.61
36	BC	151	G	C2-N2	-8.11	1.26	1.34
85	AA	2221	A	N9-C4	-8.11	1.32	1.37
35	BB	598	C	C3'-C2'	-8.11	1.43	1.52
85	AA	1259	U	C4'-C3'	-8.11	1.44	1.53
85	AA	2174	G	C8-N7	-8.11	1.26	1.30
34	BA	239	C	C2'-C1'	-8.10	1.44	1.53
34	BA	1235	C	O3'-P	-8.10	1.51	1.61
34	BA	1502	G	C1'-N9	-8.10	1.35	1.46
35	BB	1003	G	C2'-C1'	-8.10	1.44	1.53
35	BB	1166	A	C1'-N9	-8.10	1.35	1.46
34	BA	431	A	C3'-C2'	-8.10	1.43	1.52
34	BA	454	G	O3'-P	-8.10	1.51	1.61
35	BB	1063	C	O4'-C1'	-8.10	1.31	1.41
85	AA	7	G	C3'-C2'	-8.10	1.43	1.52
85	AA	1470	A	C5-C4	-8.10	1.33	1.38
34	BA	1708	A	C2'-C1'	-8.10	1.44	1.53
35	BB	1045	G	N7-C5	-8.10	1.34	1.39
85	AA	53	G	O3'-P	-8.10	1.51	1.61
85	AA	693	A	N7-C5	-8.10	1.34	1.39
85	AA	1211	C	C4'-C3'	-8.10	1.44	1.53
85	AA	1478	G	O3'-P	-8.10	1.51	1.61
85	AA	1539	A	C1'-N9	-8.10	1.35	1.46
34	BA	320	G	O3'-P	-8.10	1.51	1.61
34	BA	608	G	C4'-O4'	-8.10	1.35	1.45
34	BA	734	G	C5-C4	-8.10	1.32	1.38
39	BF	12	U	O3'-P	-8.10	1.51	1.61
34	BA	1024	A	P-O5'	-8.10	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1102	A	O3'-P	-8.10	1.51	1.61
34	BA	1466	U	C2'-C1'	-8.10	1.44	1.53
85	AA	1450	U	C2-N3	-8.10	1.32	1.37
35	BB	120	C	C2-N3	-8.10	1.29	1.35
35	BB	590	G	N1-C2	-8.10	1.31	1.37
35	BB	669	A	O3'-P	-8.10	1.51	1.61
85	AA	2122	A	O3'-P	-8.10	1.51	1.61
34	BA	313	C	O3'-P	-8.10	1.51	1.61
34	BA	772	G	N7-C5	-8.10	1.34	1.39
34	BA	909	G	C3'-C2'	-8.10	1.43	1.52
34	BA	1143	U	C3'-C2'	-8.10	1.43	1.52
35	BB	1325	C	P-O5'	-8.10	1.51	1.59
85	AA	1475	A	N9-C4	-8.10	1.32	1.37
34	BA	286	C	C2'-C1'	-8.09	1.44	1.53
34	BA	296	G	N7-C5	-8.09	1.34	1.39
85	AA	1486	G	N3-C4	-8.09	1.29	1.35
34	BA	967	C	C3'-C2'	-8.09	1.43	1.52
34	BA	1840	C	P-O5'	-8.09	1.51	1.59
34	BA	1845	G	C2-N2	-8.09	1.26	1.34
35	BB	459	U	C2'-C1'	-8.09	1.44	1.53
41	BH	129	G	C2-N3	-8.09	1.26	1.32
85	AA	35	U	P-O5'	-8.09	1.51	1.59
35	BB	692	G	C6-N1	-8.09	1.33	1.39
35	BB	986	C	N1-C6	-8.09	1.32	1.37
36	BC	41	A	C1'-N9	-8.09	1.35	1.46
37	BD	46	G	O3'-P	-8.09	1.51	1.61
38	BE	141	A	N9-C4	-8.09	1.32	1.37
34	BA	38	G	O3'-P	-8.09	1.51	1.61
34	BA	617	G	O3'-P	-8.09	1.51	1.61
34	BA	1559	C	P-O5'	-8.09	1.51	1.59
35	BB	125	G	O3'-P	-8.09	1.51	1.61
35	BB	556	U	O3'-P	-8.09	1.51	1.61
85	AA	392	G	O3'-P	-8.09	1.51	1.61
85	AA	2175	U	C2-N3	-8.09	1.32	1.37
34	BA	899	G	N9-C4	-8.09	1.31	1.38
34	BA	905	A	C2'-C1'	-8.09	1.44	1.53
34	BA	1249	G	O3'-P	-8.09	1.51	1.61
34	BA	1303	U	P-O5'	-8.09	1.51	1.59
35	BB	472	C	P-O5'	-8.09	1.51	1.59
34	BA	424	U	C2'-C1'	-8.09	1.44	1.53
34	BA	861	C	N1-C6	-8.09	1.32	1.37
35	BB	127	U	C1'-N1	-8.09	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	365	U	O3'-P	-8.09	1.51	1.61
35	BB	1168	G	O3'-P	-8.09	1.51	1.61
35	BB	1284	U	O3'-P	-8.09	1.51	1.61
37	BD	79	G	C5-C4	-8.09	1.32	1.38
37	BD	25	G	N9-C4	-8.09	1.31	1.38
85	AA	1288	A	C2'-C1'	-8.09	1.44	1.53
41	BH	41	A	N9-C8	-8.09	1.31	1.37
35	BB	456	A	C1'-N9	-8.08	1.35	1.46
85	AA	254	G	P-O5'	-8.08	1.51	1.59
85	AA	869	A	N7-C5	-8.08	1.34	1.39
34	BA	449	G	C5-C4	-8.08	1.32	1.38
34	BA	875	G	C2'-C1'	-8.08	1.44	1.53
34	BA	944	G	N9-C4	-8.08	1.31	1.38
34	BA	1294	C	C2-N3	-8.08	1.29	1.35
34	BA	1563	G	N1-C2	-8.08	1.31	1.37
35	BB	136	A	N7-C5	-8.08	1.34	1.39
35	BB	1458	U	N1-C2	-8.08	1.31	1.38
37	BD	110	G	C2'-C1'	-8.08	1.44	1.53
38	BE	201	A	C2'-C1'	-8.08	1.44	1.53
41	BH	45	G	C1'-N9	-8.08	1.35	1.46
85	AA	354	C	C2'-C1'	-8.08	1.44	1.53
85	AA	473	C	C2'-C1'	-8.08	1.44	1.53
85	AA	650	G	P-O5'	-8.08	1.51	1.59
34	BA	301	U	C2-N3	-8.08	1.32	1.37
34	BA	1250	C	C1'-N1	-8.08	1.35	1.46
35	BB	457	U	O3'-P	-8.08	1.51	1.61
34	BA	538	G	N1-C2	-8.08	1.31	1.37
34	BA	1022	C	C3'-C2'	-8.08	1.43	1.52
34	BA	1165	A	C1'-N9	-8.08	1.35	1.46
35	BB	702	G	C1'-N9	-8.08	1.35	1.46
35	BB	1489	A	P-O5'	-8.08	1.51	1.59
34	BA	122	U	O3'-P	-8.07	1.51	1.61
85	AA	644	A	C1'-N9	-8.07	1.35	1.46
34	BA	493	G	O3'-P	-8.07	1.51	1.61
34	BA	1613	G	C8-N7	-8.07	1.26	1.30
35	BB	1150	A	O3'-P	-8.07	1.51	1.61
38	BE	32	U	O4'-C1'	-8.07	1.31	1.41
85	AA	119	G	C5-C4	-8.07	1.32	1.38
34	BA	104	A	C5-C4	-8.07	1.33	1.38
34	BA	1527	G	C6-N1	-8.07	1.33	1.39
34	BA	1638	U	C2-N3	-8.07	1.32	1.37
35	BB	651	G	C8-N7	-8.07	1.26	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1120	A	P-O5'	-8.07	1.51	1.59
36	BC	12	A	O3'-P	-8.07	1.51	1.61
36	BC	19	A	N3-C4	-8.07	1.30	1.34
85	AA	902	A	C4'-C3'	-8.07	1.44	1.53
34	BA	926	A	O3'-P	-8.07	1.51	1.61
85	AA	190	A	O3'-P	-8.07	1.51	1.61
85	AA	390	U	P-O5'	-8.07	1.51	1.59
85	AA	1532	G	O3'-P	-8.07	1.51	1.61
85	AA	450	A	O3'-P	-8.07	1.51	1.61
35	BB	1103	A	C4'-O4'	-8.07	1.35	1.45
34	BA	353	U	C2-N3	-8.07	1.32	1.37
35	BB	794	G	N9-C4	-8.07	1.31	1.38
85	AA	106	G	C6-N1	-8.07	1.33	1.39
85	AA	925	G	N1-C2	-8.07	1.31	1.37
85	AA	1242	A	N7-C5	-8.07	1.34	1.39
34	BA	195	G	C2'-C1'	-8.06	1.44	1.53
34	BA	52	G	C5-C4	-8.06	1.32	1.38
34	BA	494	A	C3'-C2'	-8.06	1.43	1.52
34	BA	710	A	C2'-C1'	-8.06	1.44	1.53
34	BA	1288	U	C1'-N1	-8.06	1.35	1.46
35	BB	1094	A	C5-C4	-8.06	1.33	1.38
85	AA	462	A	N9-C4	-8.06	1.33	1.37
85	AA	535	G	C2-N2	-8.06	1.26	1.34
34	BA	1695	G	C6-N1	-8.06	1.33	1.39
37	BD	20	C	P-O5'	-8.06	1.51	1.59
85	AA	113	U	C3'-C2'	-8.06	1.43	1.52
85	AA	553	G	P-O5'	-8.06	1.51	1.59
85	AA	718	C	C2'-C1'	-8.06	1.44	1.53
85	AA	1894	G	C5-C4	-8.06	1.32	1.38
34	BA	387	A	N7-C5	-8.06	1.34	1.39
34	BA	498	A	C1'-N9	-8.06	1.35	1.46
34	BA	911	G	N1-C2	-8.06	1.31	1.37
34	BA	929	A	N7-C5	-8.06	1.34	1.39
35	BB	1183	U	C2-N3	-8.06	1.32	1.37
85	AA	125	A	O3'-P	-8.06	1.51	1.61
35	BB	1049	G	N1-C2	-8.06	1.31	1.37
34	BA	871	G	N3-C4	8.06	1.41	1.35
34	BA	1482	A	C1'-N9	-8.06	1.35	1.46
35	BB	93	A	N7-C5	-8.06	1.34	1.39
85	AA	1554	C	C2'-C1'	-8.06	1.44	1.53
85	AA	2072	G	C6-N1	-8.06	1.33	1.39
34	BA	53	G	C3'-C2'	-8.05	1.43	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	401	A	C6-N1	-8.05	1.29	1.35
34	BA	875	G	C5-C4	-8.05	1.32	1.38
36	BC	47	C	P-O5'	-8.06	1.51	1.59
34	BA	670	U	C4'-C3'	-8.05	1.44	1.53
34	BA	1108	U	C2-N3	-8.05	1.32	1.37
34	BA	1294	C	C2'-C1'	-8.05	1.44	1.53
34	BA	1337	A	O3'-P	-8.05	1.51	1.61
35	BB	1386	C	O3'-P	-8.05	1.51	1.61
85	AA	290	G	C2'-C1'	-8.05	1.44	1.53
34	BA	1001	G	C5-C4	-8.05	1.32	1.38
34	BA	1070	G	N7-C5	-8.05	1.34	1.39
34	BA	1026	C	O3'-P	-8.05	1.51	1.61
34	BA	1508	C	C2-N3	-8.05	1.29	1.35
35	BB	418	G	C1'-N9	-8.05	1.35	1.46
85	AA	925	G	C3'-C2'	-8.05	1.43	1.52
35	BB	823	G	N9-C4	8.05	1.44	1.38
35	BB	990	G	P-O5'	-8.05	1.51	1.59
39	BF	37	C	P-O5'	-8.05	1.51	1.59
85	AA	177	A	N9-C4	-8.05	1.33	1.37
40	BG	34	A	P-O5'	-8.05	1.51	1.59
85	AA	194	U	C2-N3	-8.05	1.32	1.37
85	AA	400	G	P-O5'	-8.05	1.51	1.59
85	AA	1105	G	P-O5'	-8.05	1.51	1.59
34	BA	876	C	C3'-C2'	-8.05	1.43	1.52
34	BA	888	G	C2'-C1'	-8.05	1.44	1.53
34	BA	1411	C	C2'-C1'	-8.05	1.44	1.53
34	BA	1522	G	C2'-C1'	-8.05	1.44	1.53
34	BA	1703	A	N9-C4	-8.05	1.33	1.37
34	BA	1717	C	O3'-P	-8.05	1.51	1.61
35	BB	560	C	C2'-C1'	-8.05	1.44	1.53
85	AA	1732	G	P-O5'	-8.05	1.51	1.59
85	AA	1974	C	C4-C5	-8.05	1.36	1.43
34	BA	196	A	C1'-N9	-8.05	1.35	1.46
34	BA	214	A	C2'-C1'	-8.04	1.44	1.53
34	BA	531	C	P-O5'	-8.04	1.51	1.59
35	BB	1098	G	C1'-N9	-8.05	1.35	1.46
36	BC	87	C	O3'-P	-8.05	1.51	1.61
85	AA	1233	G	N7-C5	-8.05	1.34	1.39
35	BB	1149	A	P-O5'	-8.04	1.51	1.59
36	BC	48	A	O3'-P	-8.04	1.51	1.61
37	BD	68	C	O3'-P	-8.04	1.51	1.61
85	AA	1244	A	O3'-P	-8.04	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	418	G	P-O5'	-8.04	1.51	1.59
34	BA	420	A	C1'-N9	-8.04	1.35	1.46
34	BA	459	U	O3'-P	-8.04	1.51	1.61
34	BA	1673	G	O3'-P	-8.04	1.51	1.61
35	BB	1138	A	C1'-N9	-8.04	1.35	1.46
85	AA	394	C	P-O5'	-8.04	1.51	1.59
35	BB	1545	U	C2'-C1'	-8.04	1.44	1.53
37	BD	3	G	C4'-C3'	-8.04	1.44	1.53
38	BE	97	G	O3'-P	-8.04	1.51	1.61
40	BG	174	G	C2-N2	-8.04	1.26	1.34
85	AA	1119	A	N9-C4	-8.04	1.33	1.37
40	BG	25	G	C3'-C2'	-8.04	1.44	1.52
85	AA	92	G	O3'-P	-8.04	1.51	1.61
35	BB	522	A	C2'-C1'	-8.04	1.44	1.53
36	BC	11	G	C6-N1	-8.04	1.33	1.39
39	BF	70	A	N9-C4	-8.04	1.33	1.37
34	BA	413	A	C3'-C2'	-8.04	1.44	1.52
35	BB	1296	A	O3'-P	-8.04	1.51	1.61
37	BD	69	U	C3'-C2'	-8.04	1.44	1.52
85	AA	794	A	N7-C5	-8.04	1.34	1.39
85	AA	2177	C	P-O5'	-8.04	1.51	1.59
41	BH	39	G	O3'-P	-8.04	1.51	1.61
85	AA	1134	G	C1'-N9	-8.04	1.35	1.46
35	BB	1070	G	N7-C5	-8.04	1.34	1.39
34	BA	702	G	C2'-C1'	-8.03	1.44	1.53
34	BA	1158	A	C3'-C2'	-8.04	1.44	1.52
35	BB	1268	C	C2'-C1'	-8.04	1.44	1.53
85	AA	178	U	C2-N3	-8.04	1.32	1.37
85	AA	612	A	O3'-P	-8.03	1.51	1.61
85	AA	1196	C	O3'-P	-8.04	1.51	1.61
34	BA	2	A	N9-C4	-8.03	1.33	1.37
34	BA	146	G	N9-C4	-8.03	1.31	1.38
35	BB	1076	U	C3'-C2'	-8.03	1.44	1.52
85	AA	470	C	C1'-N1	-8.03	1.35	1.46
34	BA	1281	U	C2'-C1'	-8.03	1.44	1.53
34	BA	1528	U	C2-N3	-8.03	1.32	1.37
35	BB	1310	C	C2'-C1'	-8.03	1.44	1.53
36	BC	168	C	C2-N3	-8.03	1.29	1.35
38	BE	61	A	O3'-P	-8.03	1.51	1.61
38	BE	177	U	N3-C4	-8.03	1.31	1.38
38	BE	204	U	C3'-C2'	-8.03	1.44	1.52
85	AA	469	G	C6-N1	-8.03	1.33	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	291	C	C2'-C1'	-8.03	1.44	1.53
34	BA	1511	C	P-O5'	-8.03	1.51	1.59
35	BB	557	C	C2-N3	-8.03	1.29	1.35
35	BB	1102	U	C2-N3	-8.03	1.32	1.37
34	BA	1016	A	C3'-C2'	-8.03	1.44	1.52
35	BB	434	A	C1'-N9	-8.03	1.35	1.46
35	BB	563	A	P-O5'	-8.03	1.51	1.59
35	BB	1298	C	O3'-P	-8.03	1.51	1.61
38	BE	96	G	N9-C4	-8.03	1.31	1.38
38	BE	189	A	C2'-C1'	-8.03	1.44	1.53
41	BH	34	G	N7-C5	-8.03	1.34	1.39
34	BA	15	G	C3'-C2'	-8.03	1.44	1.52
34	BA	44	U	O3'-P	-8.03	1.51	1.61
34	BA	233	U	C2-N3	-8.03	1.32	1.37
35	BB	69	A	O3'-P	-8.03	1.51	1.61
85	AA	167	A	O3'-P	-8.03	1.51	1.61
34	BA	531	C	C3'-O3'	-8.03	1.30	1.42
34	BA	799	A	N7-C5	-8.03	1.34	1.39
34	BA	1047	U	O3'-P	-8.03	1.51	1.61
35	BB	115	A	N9-C4	-8.03	1.33	1.37
40	BG	74	G	C2-N2	-8.03	1.26	1.34
85	AA	2188	C	C3'-C2'	-8.03	1.44	1.52
34	BA	933	U	C2-N3	-8.02	1.32	1.37
34	BA	1614	G	P-O5'	-8.02	1.51	1.59
35	BB	438	G	P-O5'	-8.02	1.51	1.59
35	BB	607	G	N9-C4	-8.02	1.31	1.38
36	BC	27	U	O3'-P	-8.02	1.51	1.61
85	AA	1185	G	C5'-C4'	8.02	1.60	1.51
85	AA	2095	U	O3'-P	-8.02	1.51	1.61
34	BA	454	G	P-O5'	-8.02	1.51	1.59
35	BB	425	G	N9-C4	-8.02	1.31	1.38
34	BA	1653	G	O3'-P	-8.02	1.51	1.61
35	BB	572	G	C1'-N9	-8.02	1.35	1.46
38	BE	39	U	C2'-C1'	-8.02	1.44	1.53
38	BE	158	U	C2'-C1'	-8.02	1.44	1.53
39	BF	57	C	P-O5'	-8.02	1.51	1.59
41	BH	24	U	C2-N3	-8.02	1.32	1.37
85	AA	1284	A	C4'-C3'	-8.02	1.44	1.53
34	BA	690	G	C1'-N9	-8.02	1.35	1.46
34	BA	1091	U	O3'-P	-8.02	1.51	1.61
35	BB	620	G	C1'-N9	-8.02	1.35	1.46
85	AA	120	C	C3'-C2'	-8.02	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1817	U	O3'-P	-8.02	1.51	1.61
34	BA	1491	U	C5'-C4'	8.02	1.60	1.51
35	BB	515	C	O3'-P	-8.02	1.51	1.61
36	BC	160	C	P-O5'	-8.02	1.51	1.59
85	AA	543	A	P-O5'	-8.02	1.51	1.59
85	AA	1925	A	N9-C4	-8.02	1.33	1.37
34	BA	325	A	N9-C4	-8.01	1.33	1.37
35	BB	131	A	C3'-C2'	-8.01	1.44	1.52
35	BB	1052	G	C3'-C2'	-8.01	1.44	1.52
34	BA	753	G	C2'-C1'	-8.01	1.44	1.53
34	BA	1103	G	C5-C4	-8.01	1.32	1.38
34	BA	1165	A	O3'-P	-8.01	1.51	1.61
35	BB	425	G	O3'-P	-8.01	1.51	1.61
35	BB	1115	G	N1-C2	-8.01	1.31	1.37
85	AA	34	G	C2'-C1'	-8.01	1.44	1.53
34	BA	611	A	P-O5'	-8.01	1.51	1.59
34	BA	1167	A	C2'-C1'	-8.01	1.44	1.53
34	BA	1406	U	C2'-C1'	-8.01	1.44	1.53
35	BB	85	A	C1'-N9	-8.01	1.35	1.46
36	BC	24	G	C3'-C2'	-8.01	1.44	1.52
35	BB	787	A	O3'-P	-8.01	1.51	1.61
85	AA	2181	G	N9-C4	-8.01	1.31	1.38
36	BC	53	A	N9-C4	-8.01	1.33	1.37
34	BA	895	U	C2'-C1'	-8.01	1.44	1.53
34	BA	1057	C	P-O5'	-8.01	1.51	1.59
34	BA	1426	A	N9-C4	-8.01	1.33	1.37
34	BA	1691	G	N9-C4	-8.01	1.31	1.38
34	BA	1809	G	C4'-C3'	8.01	1.61	1.53
34	BA	1815	G	C6-N1	-8.01	1.33	1.39
35	BB	1041	A	N7-C5	-8.01	1.34	1.39
39	BF	49	C	C3'-C2'	-8.01	1.44	1.52
85	AA	820	G	P-O5'	-8.01	1.51	1.59
85	AA	2138	G	C6-N1	-8.01	1.33	1.39
34	BA	625	U	O3'-P	-8.00	1.51	1.61
85	AA	1847	U	P-O5'	-8.00	1.51	1.59
34	BA	893	U	C2-N3	-8.00	1.32	1.37
34	BA	1313	U	O3'-P	-8.00	1.51	1.61
40	BG	26	G	C3'-C2'	-8.00	1.44	1.52
85	AA	418	G	C5-C4	-8.00	1.32	1.38
85	AA	1167	G	P-O5'	-8.00	1.51	1.59
85	AA	2238	C	C3'-C2'	-8.00	1.44	1.52
34	BA	1845	G	N7-C5	-8.00	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
39	BF	10	A	C3'-O3'	8.00	1.53	1.42
40	BG	31	G	C2-N2	-8.00	1.26	1.34
34	BA	423	G	C2'-C1'	-8.00	1.44	1.53
34	BA	690	G	O3'-P	-8.00	1.51	1.61
34	BA	919	A	C5-C4	-8.00	1.33	1.38
34	BA	1070	G	C2'-C1'	-8.00	1.44	1.53
34	BA	1172	C	C3'-C2'	-8.00	1.44	1.52
35	BB	1416	A	C5'-C4'	8.00	1.60	1.51
85	AA	52	U	C2'-C1'	-8.00	1.44	1.53
85	AA	1731	G	N3-C4	-8.00	1.29	1.35
85	AA	1810	C	O3'-P	-8.00	1.51	1.61
34	BA	20	A	O3'-P	-8.00	1.51	1.61
34	BA	58	A	O3'-P	-8.00	1.51	1.61
34	BA	1518	A	C8-N7	-8.00	1.25	1.31
34	BA	1661	U	P-O5'	-8.00	1.51	1.59
35	BB	129	U	C2-N3	-8.00	1.32	1.37
37	BD	95	G	C2-N2	-8.00	1.26	1.34
85	AA	1860	A	N9-C4	-8.00	1.33	1.37
40	BG	50	G	O3'-P	-8.00	1.51	1.61
85	AA	396	U	C3'-C2'	-8.00	1.44	1.52
85	AA	431	G	O3'-P	-8.00	1.51	1.61
34	BA	599	U	C2-N3	-7.99	1.32	1.37
34	BA	1226	G	P-O5'	-7.99	1.51	1.59
35	BB	575	C	P-O5'	-7.99	1.51	1.59
35	BB	617	C	C2'-C1'	-7.99	1.44	1.53
35	BB	628	A	C2'-C1'	-7.99	1.44	1.53
35	BB	1457	A	C4'-C3'	-7.99	1.44	1.53
36	BC	42	G	N9-C8	-7.99	1.32	1.37
85	AA	597	A	P-O5'	-7.99	1.51	1.59
85	AA	1916	A	N7-C5	-7.99	1.34	1.39
85	AA	1928	A	C4'-O4'	-7.99	1.35	1.45
36	BC	106	G	N7-C5	-7.99	1.34	1.39
85	AA	54	C	C2-N3	-7.99	1.29	1.35
34	BA	788	C	C4'-C3'	-7.99	1.44	1.53
34	BA	889	U	C2-N3	-7.99	1.32	1.37
34	BA	1080	U	C4'-C3'	7.99	1.61	1.53
34	BA	1814	U	C3'-C2'	-7.99	1.44	1.52
35	BB	89	C	P-O5'	-7.99	1.51	1.59
35	BB	828	G	P-O5'	-7.99	1.51	1.59
36	BC	78	G	P-O5'	-7.99	1.51	1.59
85	AA	48	G	N1-C2	-7.99	1.31	1.37
85	AA	373	G	C1'-N9	-7.99	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1962	U	P-O5'	-7.99	1.51	1.59
86	AB	68	C	O3'-P	-7.99	1.51	1.61
34	BA	1170	A	C2'-C1'	-7.99	1.44	1.53
34	BA	1720	U	C2'-C1'	-7.99	1.44	1.53
35	BB	597	C	C2-N3	-7.99	1.29	1.35
35	BB	1046	C	C2'-C1'	-7.99	1.44	1.53
35	BB	1262	A	C5-C4	-7.99	1.33	1.38
40	BG	109	C	C3'-C2'	-7.99	1.44	1.52
40	BG	139	U	C2'-C1'	-7.99	1.44	1.53
40	BG	163	G	C5-C6	-7.99	1.34	1.42
86	AB	67	C	O3'-P	-7.99	1.51	1.61
34	BA	1666	U	C2-N3	-7.99	1.32	1.37
40	BG	26	G	N9-C4	-7.99	1.31	1.38
41	BH	13	C	O3'-P	-7.99	1.51	1.61
85	AA	895	C	O3'-P	-7.99	1.51	1.61
34	BA	961	C	C4-C5	-7.99	1.36	1.43
34	BA	1065	U	P-O5'	-7.99	1.51	1.59
34	BA	1300	G	C2-N2	-7.99	1.26	1.34
35	BB	72	G	N1-C2	-7.99	1.31	1.37
35	BB	129	U	C3'-C2'	-7.99	1.44	1.52
85	AA	378	A	C2'-C1'	-7.99	1.44	1.53
85	AA	1144	G	P-O5'	-7.99	1.51	1.59
85	AA	1484	G	C4'-C3'	-7.99	1.44	1.53
85	AA	1711	C	C2'-C1'	-7.99	1.44	1.53
34	BA	251	U	O3'-P	-7.98	1.51	1.61
34	BA	258	C	P-O5'	-7.98	1.51	1.59
35	BB	1543	C	C2'-C1'	-7.98	1.44	1.53
85	AA	531	G	C6-N1	-7.98	1.33	1.39
34	BA	566	G	N7-C5	-7.98	1.34	1.39
34	BA	582	U	N3-C4	-7.98	1.31	1.38
34	BA	744	G	N7-C5	-7.98	1.34	1.39
35	BB	679	G	O3'-P	-7.98	1.51	1.61
35	BB	689	C	C3'-C2'	-7.98	1.44	1.52
85	AA	790	A	N9-C4	-7.98	1.33	1.37
85	AA	1197	U	O3'-P	-7.98	1.51	1.61
34	BA	1115	A	C3'-C2'	-7.98	1.44	1.52
35	BB	1037	A	C2'-C1'	-7.98	1.44	1.53
35	BB	1078	U	P-O5'	-7.98	1.51	1.59
35	BB	1163	U	C2-N3	-7.98	1.32	1.37
36	BC	117	A	N7-C5	-7.98	1.34	1.39
37	BD	27	A	O3'-P	-7.98	1.51	1.61
37	BD	71	G	P-O5'	-7.98	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2201	A	C2'-C1'	-7.98	1.44	1.53
35	BB	1254	G	P-O5'	-7.98	1.51	1.59
35	BB	1482	A	C2'-C1'	-7.98	1.44	1.53
35	BB	1484	A	C8-N7	-7.98	1.25	1.31
37	BD	8	A	N9-C4	-7.98	1.33	1.37
85	AA	455	G	C4'-C3'	-7.98	1.44	1.53
85	AA	483	G	O3'-P	-7.98	1.51	1.61
85	AA	547	A	O3'-P	-7.98	1.51	1.61
34	BA	690	G	O4'-C1'	-7.98	1.31	1.41
34	BA	1137	U	P-O5'	-7.98	1.51	1.59
35	BB	667	G	N9-C4	-7.98	1.31	1.38
35	BB	1200	A	C2'-C1'	-7.98	1.44	1.53
35	BB	1441	C	C1'-N1	-7.98	1.35	1.46
37	BD	10	C	P-O5'	-7.98	1.51	1.59
85	AA	628	C	O3'-P	-7.98	1.51	1.61
34	BA	1718	C	C2'-C1'	-7.97	1.44	1.53
35	BB	427	U	O3'-P	-7.97	1.51	1.61
35	BB	509	A	O3'-P	-7.97	1.51	1.61
35	BB	544	C	N1-C6	-7.97	1.32	1.37
35	BB	659	C	P-O5'	-7.97	1.51	1.59
35	BB	1290	C	C2-N3	-7.97	1.29	1.35
35	BB	1314	G	C5'-C4'	-7.97	1.41	1.51
36	BC	31	A	N9-C8	-7.97	1.31	1.37
36	BC	49	G	C2-N2	-7.97	1.26	1.34
85	AA	491	G	O3'-P	-7.97	1.51	1.61
85	AA	767	A	C1'-N9	-7.97	1.35	1.46
85	AA	1947	A	P-O5'	-7.97	1.51	1.59
85	AA	2060	G	N9-C4	-7.97	1.31	1.38
34	BA	513	U	C3'-C2'	-7.97	1.44	1.52
34	BA	1171	C	O3'-P	-7.97	1.51	1.61
34	BA	1450	G	O3'-P	-7.97	1.51	1.61
34	BA	1594	G	C3'-C2'	-7.97	1.44	1.52
35	BB	22	A	C6-N1	-7.97	1.29	1.35
38	BE	148	C	C4'-C3'	-7.97	1.44	1.53
85	AA	1525	C	C2-N3	-7.97	1.29	1.35
85	AA	1890	C	P-O5'	-7.97	1.51	1.59
85	AA	2029	G	O3'-P	-7.97	1.51	1.61
34	BA	71	G	P-O5'	-7.97	1.51	1.59
34	BA	1469	G	C5-C6	-7.97	1.34	1.42
36	BC	36	G	C2'-C1'	-7.97	1.44	1.53
34	BA	224	G	C2-N2	-7.97	1.26	1.34
35	BB	1398	A	N7-C5	-7.97	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1445	A	C1'-N9	-7.97	1.35	1.46
36	BC	113	G	C5-C6	-7.97	1.34	1.42
85	AA	16	G	N9-C8	-7.97	1.32	1.37
85	AA	382	G	O4'-C1'	-7.97	1.31	1.41
35	BB	1235	A	C2'-C1'	-7.97	1.44	1.53
85	AA	619	A	O3'-P	-7.97	1.51	1.61
34	BA	676	G	N7-C5	-7.97	1.34	1.39
34	BA	1300	G	N9-C4	-7.97	1.31	1.38
35	BB	1325	C	O3'-P	-7.97	1.51	1.61
40	BG	33	G	N7-C5	-7.97	1.34	1.39
40	BG	122	G	C5-C4	-7.97	1.32	1.38
85	AA	186	U	O3'-P	-7.97	1.51	1.61
85	AA	1148	G	P-O5'	-7.97	1.51	1.59
86	AB	66	U	C2'-C1'	-7.97	1.44	1.53
34	BA	713	C	C4'-C3'	-7.96	1.44	1.53
41	BH	36	C	C2'-C1'	-7.96	1.44	1.53
85	AA	361	U	C2'-C1'	-7.96	1.44	1.53
85	AA	2115	G	P-O5'	-7.96	1.51	1.59
34	BA	1252	G	C2-N2	-7.96	1.26	1.34
35	BB	1001	G	N9-C4	7.96	1.44	1.38
37	BD	48	G	C5-C4	-7.96	1.32	1.38
40	BG	154	C	O3'-P	-7.96	1.51	1.61
36	BC	145	G	N7-C5	-7.96	1.34	1.39
85	AA	2057	G	P-O5'	-7.96	1.51	1.59
85	AA	2170	G	O3'-P	-7.96	1.51	1.61
34	BA	320	G	N9-C8	-7.96	1.32	1.37
34	BA	1441	C	C2'-C1'	-7.96	1.44	1.53
34	BA	727	G	O3'-P	-7.96	1.51	1.61
35	BB	452	A	O4'-C1'	-7.96	1.31	1.41
38	BE	9	C	C2-N3	-7.96	1.29	1.35
85	AA	24	U	O3'-P	-7.96	1.51	1.61
85	AA	2100	A	C1'-N9	-7.96	1.35	1.46
34	BA	293	A	C4'-C3'	-7.96	1.44	1.53
34	BA	429	G	N7-C5	-7.96	1.34	1.39
34	BA	656	U	O3'-P	-7.96	1.51	1.61
34	BA	841	G	N7-C5	-7.96	1.34	1.39
34	BA	1290	A	N3-C4	-7.96	1.30	1.34
34	BA	1613	G	C6-N1	-7.96	1.33	1.39
35	BB	387	G	N9-C4	-7.96	1.31	1.38
35	BB	1291	G	N1-C2	-7.96	1.31	1.37
35	BB	1401	G	N3-C4	-7.96	1.29	1.35
34	BA	177	G	C1'-N9	-7.96	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1744	C	C2'-C1'	-7.96	1.44	1.53
35	BB	1019	C	C1'-N1	-7.96	1.35	1.46
35	BB	1143	A	C4'-C3'	-7.96	1.44	1.53
36	BC	62	A	N3-C4	-7.96	1.30	1.34
34	BA	1008	A	P-O5'	-7.95	1.51	1.59
35	BB	85	A	C2'-C1'	-7.95	1.44	1.53
35	BB	1035	C	P-O5'	-7.95	1.51	1.59
35	BB	1097	U	C3'-C2'	-7.95	1.44	1.52
35	BB	1254	G	C4'-C3'	-7.95	1.44	1.53
35	BB	1428	C	P-O5'	-7.95	1.51	1.59
36	BC	51	A	O3'-P	-7.95	1.51	1.61
39	BF	26	U	O3'-P	-7.95	1.51	1.61
40	BG	80	G	C3'-C2'	-7.95	1.44	1.52
85	AA	129	U	P-O5'	-7.95	1.51	1.59
85	AA	513	G	O3'-P	-7.95	1.51	1.61
34	BA	236	A	C1'-N9	-7.95	1.35	1.46
34	BA	1240	G	C5-C6	-7.95	1.34	1.42
34	BA	1640	G	C2'-C1'	-7.95	1.44	1.53
35	BB	543	G	P-O5'	-7.95	1.51	1.59
35	BB	1088	C	O3'-P	-7.95	1.51	1.61
35	BB	1144	A	N9-C4	-7.95	1.33	1.37
36	BC	106	G	P-O5'	-7.95	1.51	1.59
39	BF	32	G	C2'-C1'	-7.95	1.44	1.53
41	BH	43	G	C2'-C1'	-7.95	1.44	1.53
85	AA	679	A	O3'-P	-7.95	1.51	1.61
85	AA	741	G	C2'-C1'	-7.95	1.44	1.53
34	BA	147	U	C2'-C1'	-7.95	1.44	1.53
34	BA	269	G	N7-C5	-7.95	1.34	1.39
34	BA	533	U	C2'-C1'	-7.95	1.44	1.53
35	BB	1395	G	O3'-P	-7.95	1.51	1.61
36	BC	102	G	C2'-C1'	-7.95	1.44	1.53
85	AA	1558	U	C2-N3	-7.95	1.32	1.37
34	BA	465	A	C2'-C1'	-7.95	1.44	1.53
34	BA	845	U	O3'-P	-7.95	1.51	1.61
35	BB	57	G	C1'-N9	-7.95	1.35	1.46
35	BB	1222	A	C3'-C2'	-7.95	1.44	1.52
37	BD	36	C	C1'-N1	-7.95	1.35	1.46
85	AA	119	G	C3'-C2'	-7.95	1.44	1.52
85	AA	1723	U	P-O5'	-7.95	1.51	1.59
35	BB	806	U	O3'-P	-7.95	1.51	1.61
40	BG	123	C	C3'-C2'	-7.95	1.44	1.52
85	AA	372	U	P-O5'	-7.95	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	445	U	O3'-P	-7.95	1.51	1.61
85	AA	1125	G	N9-C4	-7.95	1.31	1.38
85	AA	1487	G	C2-N2	-7.95	1.26	1.34
34	BA	894	G	N7-C5	-7.94	1.34	1.39
38	BE	89	G	C6-N1	-7.94	1.33	1.39
34	BA	1071	G	C2'-C1'	-7.94	1.44	1.53
34	BA	1192	A	C3'-C2'	-7.94	1.44	1.52
40	BG	70	C	C1'-N1	-7.94	1.35	1.46
34	BA	30	A	C5-C4	-7.94	1.33	1.38
34	BA	123	C	N1-C2	-7.94	1.32	1.40
35	BB	1196	A	C3'-C2'	-7.94	1.44	1.52
36	BC	99	U	O3'-P	-7.94	1.51	1.61
40	BG	39	A	C3'-C2'	-7.94	1.44	1.52
41	BH	2	U	C2'-C1'	-7.94	1.44	1.53
85	AA	389	A	C1'-N9	-7.94	1.35	1.46
85	AA	410	A	N7-C5	-7.94	1.34	1.39
34	BA	580	U	C4'-C3'	-7.94	1.44	1.53
34	BA	1214	U	C4'-C3'	-7.94	1.44	1.53
85	AA	930	G	P-O5'	-7.94	1.51	1.59
85	AA	1528	A	N9-C4	-7.94	1.33	1.37
85	AA	2124	G	C2'-C1'	-7.94	1.44	1.53
35	BB	1258	G	P-O5'	-7.94	1.51	1.59
35	BB	497	C	O3'-P	-7.94	1.51	1.61
38	BE	132	U	C2-N3	-7.94	1.32	1.37
85	AA	1471	G	C3'-C2'	-7.94	1.44	1.52
34	BA	610	A	O3'-P	-7.93	1.51	1.61
34	BA	1405	A	O3'-P	-7.93	1.51	1.61
34	BA	1711	G	N1-C2	-7.93	1.31	1.37
35	BB	133	G	O3'-P	-7.93	1.51	1.61
35	BB	584	A	O3'-P	-7.93	1.51	1.61
38	BE	180	G	N3-C4	-7.93	1.29	1.35
85	AA	532	G	P-O5'	-7.93	1.51	1.59
34	BA	386	A	N3-C4	-7.93	1.30	1.34
34	BA	1214	U	O3'-P	-7.93	1.51	1.61
34	BA	1543	A	C1'-N9	-7.93	1.35	1.46
34	BA	1567	G	N9-C8	-7.93	1.32	1.37
34	BA	1719	G	C2'-C1'	-7.93	1.44	1.53
36	BC	135	A	O3'-P	-7.93	1.51	1.61
39	BF	15	U	P-O5'	-7.93	1.51	1.59
40	BG	19	C	C2'-C1'	-7.93	1.44	1.53
85	AA	245	A	C2'-C1'	-7.93	1.44	1.53
85	AA	494	G	P-O5'	-7.93	1.51	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1622	G	P-O5'	-7.93	1.51	1.59
85	AA	2240	G	C2'-C1'	-7.93	1.44	1.53
34	BA	1795	A	C2'-C1'	-7.93	1.44	1.53
85	AA	683	U	C2'-C1'	-7.93	1.44	1.53
35	BB	1259	A	C5-C4	-7.93	1.33	1.38
85	AA	409	C	O3'-P	-7.93	1.51	1.61
85	AA	462	A	C2'-C1'	-7.93	1.44	1.53
85	AA	1259	U	O3'-P	-7.93	1.51	1.61
35	BB	631	G	P-O5'	-7.93	1.51	1.59
40	BG	72	G	N1-C2	-7.93	1.31	1.37
85	AA	534	A	C3'-C2'	-7.93	1.44	1.52
85	AA	965	G	P-O5'	-7.93	1.51	1.59
34	BA	50	G	C1'-N9	-7.93	1.35	1.46
34	BA	386	A	C5'-C4'	7.93	1.60	1.51
34	BA	395	G	P-O5'	-7.93	1.51	1.59
34	BA	909	G	C2'-C1'	-7.93	1.44	1.53
34	BA	1178	U	C2-N3	-7.93	1.32	1.37
34	BA	1595	G	P-O5'	-7.93	1.51	1.59
35	BB	119	G	C5-C4	-7.93	1.32	1.38
35	BB	505	G	C2-N2	-7.93	1.26	1.34
35	BB	1086	G	N9-C8	-7.93	1.32	1.37
39	BF	34	C	C1'-N1	-7.93	1.35	1.46
34	BA	364	C	O3'-P	-7.92	1.51	1.61
34	BA	1057	C	O3'-P	-7.92	1.51	1.61
34	BA	1196	C	C2-N3	-7.92	1.29	1.35
34	BA	1417	C	C2'-C1'	-7.92	1.44	1.53
35	BB	51	U	N1-C2	-7.92	1.31	1.38
35	BB	131	A	C1'-N9	-7.92	1.35	1.46
35	BB	406	A	O3'-P	-7.92	1.51	1.61
35	BB	1299	G	O3'-P	-7.92	1.51	1.61
36	BC	54	G	C1'-N9	-7.92	1.35	1.46
85	AA	172	A	C2'-C1'	-7.92	1.44	1.53
34	BA	761	U	O4'-C1'	-7.92	1.31	1.41
34	BA	1498	A	N7-C5	-7.92	1.34	1.39
35	BB	438	G	C3'-C2'	-7.92	1.44	1.52
35	BB	570	A	C5-C4	-7.92	1.33	1.38
35	BB	787	A	C1'-N9	-7.92	1.35	1.46
35	BB	1068	G	N9-C4	-7.92	1.31	1.38
36	BC	81	U	P-O5'	-7.92	1.51	1.59
34	BA	446	U	C2'-C1'	-7.92	1.44	1.53
35	BB	87	G	P-O5'	-7.92	1.51	1.59
85	AA	539	A	O3'-P	-7.92	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1181	A	O3'-P	-7.92	1.51	1.61
36	BC	42	G	N9-C4	-7.92	1.31	1.38
38	BE	176	G	N7-C5	-7.92	1.34	1.39
40	BG	152	G	C5'-C4'	-7.92	1.41	1.51
85	AA	655	U	C4'-C3'	-7.92	1.44	1.53
85	AA	1482	C	O3'-P	-7.92	1.51	1.61
85	AA	1647	G	N9-C4	7.92	1.44	1.38
34	BA	1747	C	P-O5'	-7.92	1.51	1.59
35	BB	435	A	N3-C4	-7.92	1.30	1.34
35	BB	615	A	N9-C8	-7.92	1.31	1.37
85	AA	185	A	C2'-C1'	-7.92	1.44	1.53
34	BA	1472	G	P-O5'	-7.92	1.51	1.59
35	BB	837	A	C1'-N9	-7.92	1.35	1.46
35	BB	1275	A	N9-C4	-7.92	1.33	1.37
85	AA	1220	A	P-O5'	-7.92	1.51	1.59
85	AA	1444	U	C2-N3	-7.92	1.32	1.37
85	AA	2052	U	P-O5'	-7.92	1.51	1.59
34	BA	521	C	C2-N3	-7.91	1.29	1.35
34	BA	999	G	N3-C4	-7.91	1.29	1.35
35	BB	6	A	C3'-C2'	-7.91	1.44	1.52
85	AA	1864	G	C3'-C2'	-7.91	1.44	1.52
34	BA	356	C	C3'-C2'	-7.91	1.44	1.52
85	AA	118	C	C3'-C2'	-7.91	1.44	1.52
34	BA	519	G	C4'-C3'	-7.91	1.44	1.53
34	BA	542	A	N9-C4	-7.91	1.33	1.37
34	BA	1528	U	C3'-C2'	-7.91	1.44	1.52
37	BD	77	A	C5-C4	-7.91	1.33	1.38
40	BG	170	G	C2'-C1'	-7.91	1.44	1.53
85	AA	784	C	P-O5'	-7.91	1.51	1.59
34	BA	57	A	C5-C4	-7.91	1.33	1.38
34	BA	1363	A	P-O5'	-7.91	1.51	1.59
34	BA	1603	A	N7-C5	-7.91	1.34	1.39
34	BA	1496	G	N9-C8	-7.91	1.32	1.37
35	BB	1476	C	C2'-C1'	-7.91	1.44	1.53
34	BA	1557	G	O3'-P	-7.91	1.51	1.61
35	BB	1329	G	P-O5'	-7.91	1.51	1.59
36	BC	68	A	O3'-P	-7.91	1.51	1.61
37	BD	99	G	O3'-P	-7.91	1.51	1.61
41	BH	113	G	O3'-P	-7.91	1.51	1.61
85	AA	423	G	C1'-N9	-7.91	1.35	1.46
85	AA	932	A	C1'-N9	-7.91	1.35	1.46
85	AA	1617	G	C2'-C1'	-7.91	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	793	A	N9-C4	-7.90	1.33	1.37
34	BA	1442	A	O3'-P	-7.90	1.51	1.61
35	BB	116	G	N7-C5	-7.90	1.34	1.39
35	BB	526	A	N7-C5	-7.90	1.34	1.39
37	BD	108	G	O3'-P	-7.90	1.51	1.61
85	AA	539	A	N9-C4	-7.90	1.33	1.37
34	BA	1030	C	C2'-C1'	-7.90	1.44	1.53
35	BB	134	G	N1-C2	-7.90	1.31	1.37
37	BD	13	A	C5'-C4'	-7.90	1.41	1.51
85	AA	182	C	P-O5'	-7.90	1.51	1.59
34	BA	399	G	C3'-C2'	-7.90	1.44	1.52
34	BA	494	A	N9-C4	-7.90	1.33	1.37
34	BA	819	G	C6-N1	-7.90	1.34	1.39
34	BA	1202	G	C1'-N9	-7.90	1.35	1.46
34	BA	1533	G	O3'-P	-7.90	1.51	1.61
35	BB	1117	G	C1'-N9	-7.90	1.35	1.46
36	BC	138	C	C3'-C2'	-7.90	1.44	1.52
39	BF	32	G	C3'-C2'	-7.90	1.44	1.52
35	BB	102	G	N9-C4	-7.90	1.31	1.38
34	BA	1071	G	C2-N2	-7.90	1.26	1.34
34	BA	1614	G	C5-C6	-7.90	1.34	1.42
35	BB	1477	C	P-O5'	-7.90	1.51	1.59
85	AA	504	U	C2'-C1'	-7.90	1.44	1.53
85	AA	1584	U	P-O5'	-7.90	1.51	1.59
34	BA	157	U	O3'-P	-7.90	1.51	1.61
34	BA	934	G	P-O5'	-7.90	1.51	1.59
85	AA	441	C	C3'-C2'	-7.90	1.44	1.52
85	AA	1660	U	C2-N3	-7.90	1.32	1.37
34	BA	268	U	P-O5'	-7.89	1.51	1.59
34	BA	274	C	C2'-C1'	-7.89	1.44	1.53
34	BA	937	G	C2-N2	-7.89	1.26	1.34
34	BA	953	G	N1-C2	-7.89	1.31	1.37
35	BB	534	C	C2'-C1'	-7.89	1.44	1.53
40	BG	17	A	N9-C4	-7.89	1.33	1.37
85	AA	1116	G	C3'-C2'	-7.89	1.44	1.52
85	AA	1174	G	P-O5'	-7.89	1.51	1.59
85	AA	2128	G	O3'-P	-7.89	1.51	1.61
34	BA	845	U	C2'-C1'	-7.89	1.44	1.53
34	BA	1602	A	O3'-P	-7.89	1.51	1.61
36	BC	142	C	C2'-C1'	-7.89	1.44	1.53
37	BD	33	U	C3'-C2'	-7.89	1.44	1.52
40	BG	72	G	O3'-P	-7.89	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1617	G	O3'-P	-7.89	1.51	1.61
34	BA	1287	G	C3'-C2'	-7.89	1.44	1.52
34	BA	1313	U	P-O5'	-7.89	1.51	1.59
34	BA	1438	C	O3'-P	-7.89	1.51	1.61
40	BG	157	A	O3'-P	-7.89	1.51	1.61
85	AA	9	U	N1-C2	-7.89	1.31	1.38
85	AA	608	A	C1'-N9	-7.89	1.35	1.46
34	BA	388	A	P-O5'	-7.89	1.51	1.59
34	BA	1650	G	C6-N1	-7.89	1.34	1.39
34	BA	1789	A	N3-C4	-7.89	1.30	1.34
36	BC	101	U	N3-C4	-7.89	1.31	1.38
36	BC	146	U	C2-N3	-7.89	1.32	1.37
85	AA	1031	G	P-O5'	-7.89	1.51	1.59
85	AA	1496	U	C1'-N1	-7.89	1.35	1.46
34	BA	22	C	N1-C6	-7.89	1.32	1.37
34	BA	1801	G	N9-C8	-7.89	1.32	1.37
34	BA	965	A	C2'-C1'	-7.89	1.44	1.53
35	BB	597	C	O3'-P	-7.89	1.51	1.61
85	AA	817	G	C1'-N9	-7.89	1.35	1.46
85	AA	1290	G	N3-C4	-7.89	1.29	1.35
34	BA	344	G	N9-C4	-7.88	1.31	1.38
34	BA	481	A	C2-N3	-7.88	1.26	1.33
34	BA	587	U	N1-C2	-7.88	1.31	1.38
34	BA	1013	A	O3'-P	-7.88	1.51	1.61
85	AA	1511	C	C3'-C2'	-7.88	1.44	1.52
85	AA	1885	A	P-O5'	-7.88	1.51	1.59
85	AA	2099	C	P-O5'	-7.88	1.51	1.59
85	AA	2240	G	C1'-N9	-7.88	1.35	1.46
34	BA	578	C	C2'-C1'	-7.88	1.44	1.53
34	BA	1246	G	O3'-P	-7.88	1.51	1.61
35	BB	1297	G	O3'-P	-7.88	1.51	1.61
85	AA	1589	G	P-O5'	-7.88	1.51	1.59
34	BA	334	G	C6-N1	-7.88	1.34	1.39
34	BA	798	G	C8-N7	-7.88	1.26	1.30
34	BA	912	G	C5-C4	-7.88	1.32	1.38
34	BA	1158	A	P-O5'	-7.88	1.51	1.59
35	BB	615	A	C1'-N9	-7.88	1.35	1.46
35	BB	1049	G	N9-C4	-7.88	1.31	1.38
35	BB	1432	U	O3'-P	-7.88	1.51	1.61
85	AA	160	A	C4'-C3'	-7.88	1.44	1.53
85	AA	464	A	N3-C4	-7.88	1.30	1.34
85	AA	881	C	C2-N3	-7.88	1.29	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1824	G	N9-C8	-7.88	1.32	1.37
85	AA	2128	G	N9-C4	-7.88	1.31	1.38
34	BA	1019	C	O3'-P	-7.88	1.51	1.61
34	BA	1036	G	N9-C4	-7.88	1.31	1.38
35	BB	1015	U	C2-N3	-7.88	1.32	1.37
85	AA	150	U	P-O5'	-7.88	1.51	1.59
34	BA	758	G	C4'-C3'	-7.88	1.44	1.53
35	BB	589	U	C3'-C2'	-7.88	1.44	1.52
36	BC	146	U	C3'-C2'	-7.88	1.44	1.52
85	AA	976	G	C2'-C1'	-7.88	1.44	1.53
85	AA	1810	C	C2'-C1'	-7.88	1.44	1.53
34	BA	1557	G	C1'-N9	-7.88	1.35	1.46
41	BH	6	U	C2-N3	-7.88	1.32	1.37
85	AA	428	G	O3'-P	-7.88	1.51	1.61
85	AA	859	G	C3'-C2'	-7.88	1.44	1.52
85	AA	1155	A	C1'-N9	-7.88	1.35	1.46
34	BA	1531	G	C1'-N9	-7.88	1.35	1.46
35	BB	426	A	N9-C4	-7.88	1.33	1.37
85	AA	1509	A	C6-N6	-7.88	1.27	1.33
34	BA	1396	A	C1'-N9	-7.87	1.35	1.46
34	BA	1744	C	P-O5'	-7.87	1.51	1.59
35	BB	661	G	C2-N3	-7.87	1.26	1.32
35	BB	1169	A	C3'-C2'	-7.87	1.44	1.52
34	BA	843	G	C6-N1	-7.87	1.34	1.39
34	BA	1642	A	O3'-P	-7.87	1.51	1.61
35	BB	7	C	C3'-C2'	-7.87	1.44	1.52
35	BB	806	U	N1-C6	-7.87	1.30	1.38
85	AA	247	G	C1'-N9	-7.87	1.35	1.46
85	AA	748	C	P-O5'	-7.87	1.51	1.59
85	AA	865	G	C2'-C1'	-7.87	1.44	1.53
85	AA	1166	C	P-O5'	-7.87	1.51	1.59
85	AA	1269	A	C1'-N9	-7.87	1.35	1.46
85	AA	2141	G	C1'-N9	-7.87	1.35	1.46
34	BA	946	A	N9-C8	-7.87	1.31	1.37
41	BH	128	G	N9-C4	-7.87	1.31	1.38
34	BA	473	A	N9-C4	-7.87	1.33	1.37
34	BA	495	A	O3'-P	-7.87	1.51	1.61
34	BA	583	G	C1'-N9	-7.87	1.35	1.46
35	BB	1083	C	P-O5'	-7.87	1.51	1.59
35	BB	1298	C	C2-N3	-7.87	1.29	1.35
35	BB	1377	A	C2'-C1'	-7.87	1.44	1.53
85	AA	2041	G	O3'-P	-7.87	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
86	AB	15	G	N7-C5	-7.87	1.34	1.39
34	BA	1036	G	N7-C5	-7.87	1.34	1.39
35	BB	507	G	C1'-N9	-7.87	1.35	1.46
35	BB	1448	U	C3'-C2'	-7.87	1.44	1.52
38	BE	132	U	O4'-C1'	-7.87	1.31	1.41
39	BF	50	C	P-O5'	-7.87	1.51	1.59
85	AA	252	G	P-O5'	-7.87	1.51	1.59
34	BA	975	A	C5-C4	-7.87	1.33	1.38
34	BA	1577	U	C2-N3	-7.87	1.32	1.37
34	BA	1847	G	P-O5'	-7.87	1.51	1.59
35	BB	125	G	C6-N1	-7.87	1.34	1.39
35	BB	482	A	C3'-C2'	-7.87	1.44	1.52
85	AA	633	C	O3'-P	-7.87	1.51	1.61
85	AA	1263	G	C2'-C1'	-7.87	1.44	1.53
34	BA	88	C	C2-N3	-7.86	1.29	1.35
36	BC	139	A	N7-C5	-7.86	1.34	1.39
41	BH	25	A	O3'-P	-7.86	1.51	1.61
41	BH	56	C	P-O5'	-7.86	1.51	1.59
85	AA	2194	U	O3'-P	-7.86	1.51	1.61
34	BA	448	U	P-O5'	-7.86	1.51	1.59
40	BG	20	U	N3-C4	-7.86	1.31	1.38
85	AA	522	A	P-O5'	-7.86	1.51	1.59
85	AA	1975	G	P-O5'	-7.86	1.51	1.59
34	BA	90	G	P-O5'	-7.86	1.51	1.59
34	BA	530	A	N3-C4	-7.86	1.30	1.34
34	BA	589	A	C3'-C2'	-7.86	1.44	1.52
34	BA	1510	C	C2'-C1'	-7.86	1.44	1.53
34	BA	1547	G	C6-N1	-7.86	1.34	1.39
35	BB	419	G	O3'-P	-7.86	1.51	1.61
35	BB	635	A	O3'-P	-7.86	1.51	1.61
40	BG	14	G	C1'-N9	-7.86	1.35	1.46
85	AA	756	G	C4'-C3'	-7.86	1.44	1.53
34	BA	1581	G	C2-N2	-7.86	1.26	1.34
35	BB	1016	C	C3'-C2'	-7.86	1.44	1.52
36	BC	2	A	C1'-N9	-7.86	1.35	1.46
85	AA	2026	U	P-O5'	-7.86	1.51	1.59
85	AA	2143	U	C3'-C2'	-7.86	1.44	1.52
35	BB	424	U	C2-N3	-7.86	1.32	1.37
35	BB	456	A	N7-C5	-7.86	1.34	1.39
35	BB	787	A	N9-C4	-7.86	1.33	1.37
35	BB	1308	G	C5-C4	-7.86	1.32	1.38
36	BC	9	G	C4'-C3'	-7.86	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	19	A	N7-C5	-7.86	1.34	1.39
85	AA	464	A	P-O5'	-7.86	1.51	1.59
34	BA	289	A	N9-C4	-7.86	1.33	1.37
34	BA	372	U	O3'-P	-7.86	1.51	1.61
34	BA	1028	A	C5-C4	-7.86	1.33	1.38
36	BC	23	G	C6-N1	-7.86	1.34	1.39
85	AA	44	C	P-O5'	-7.86	1.51	1.59
85	AA	1696	U	P-O5'	-7.86	1.51	1.59
37	BD	78	C	O3'-P	-7.85	1.51	1.61
85	AA	2123	U	O4'-C1'	-7.85	1.31	1.41
85	AA	2232	A	N9-C4	-7.85	1.33	1.37
35	BB	802	G	N1-C2	-7.85	1.31	1.37
35	BB	1233	U	C2-N3	-7.85	1.32	1.37
37	BD	95	G	P-O5'	-7.85	1.51	1.59
41	BH	66	G	P-O5'	-7.85	1.51	1.59
85	AA	995	G	O3'-P	-7.85	1.51	1.61
85	AA	1863	A	N9-C4	-7.85	1.33	1.37
35	BB	584	A	C2'-C1'	-7.85	1.44	1.53
40	BG	5	G	N1-C2	-7.85	1.31	1.37
40	BG	151	A	P-O5'	-7.85	1.51	1.59
34	BA	1019	C	C1'-N1	-7.85	1.35	1.46
34	BA	1814	U	C2-N3	-7.85	1.32	1.37
35	BB	1104	A	P-O5'	-7.85	1.51	1.59
35	BB	1396	G	C3'-C2'	-7.85	1.44	1.52
85	AA	35	U	O3'-P	-7.85	1.51	1.61
85	AA	210	G	O3'-P	-7.85	1.51	1.61
85	AA	1150	G	P-O5'	-7.85	1.51	1.59
34	BA	1030	C	C3'-C2'	-7.85	1.44	1.52
35	BB	389	G	C2'-C1'	-7.85	1.44	1.53
35	BB	557	C	C3'-C2'	-7.85	1.44	1.52
38	BE	66	A	C2'-C1'	-7.85	1.44	1.53
39	BF	36	G	P-O5'	-7.85	1.51	1.59
40	BG	153	C	O3'-P	-7.85	1.51	1.61
85	AA	938	A	C2'-C1'	-7.85	1.44	1.53
35	BB	44	C	C1'-N1	-7.85	1.35	1.46
35	BB	1371	G	C1'-N9	-7.85	1.35	1.46
38	BE	139	U	C3'-C2'	-7.85	1.44	1.52
85	AA	55	A	N9-C4	-7.85	1.33	1.37
85	AA	1488	G	P-O5'	-7.85	1.51	1.59
85	AA	1503	G	P-O5'	-7.85	1.51	1.59
85	AA	1518	A	C5-C4	-7.85	1.33	1.38
85	AA	2147	A	C3'-C2'	-7.85	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	929	A	O3'-P	-7.84	1.51	1.61
35	BB	807	U	P-O5'	-7.84	1.51	1.59
85	AA	487	G	O3'-P	-7.84	1.51	1.61
85	AA	877	G	P-O5'	-7.84	1.51	1.59
34	BA	858	C	C3'-C2'	-7.84	1.44	1.52
85	AA	1445	C	O3'-P	-7.84	1.51	1.61
85	AA	1700	C	O3'-P	-7.84	1.51	1.61
85	AA	2084	U	C4'-C3'	-7.84	1.44	1.53
34	BA	546	U	P-O5'	-7.84	1.51	1.59
34	BA	1640	G	O3'-P	-7.84	1.51	1.61
35	BB	680	A	C2'-C1'	-7.84	1.44	1.53
35	BB	730	G	P-O5'	-7.84	1.51	1.59
34	BA	90	G	O3'-P	-7.84	1.51	1.61
34	BA	1088	G	C2'-C1'	-7.84	1.44	1.53
35	BB	1052	G	C8-N7	-7.84	1.26	1.30
35	BB	1126	A	N9-C4	-7.84	1.33	1.37
35	BB	1401	G	N9-C8	-7.84	1.32	1.37
36	BC	7	U	C4'-C3'	-7.84	1.44	1.53
37	BD	97	U	P-O5'	-7.84	1.51	1.59
34	BA	405	C	N1-C6	-7.84	1.32	1.37
34	BA	736	G	P-O5'	-7.84	1.51	1.59
40	BG	81	G	C3'-C2'	-7.84	1.44	1.52
34	BA	1210	A	N9-C8	-7.84	1.31	1.37
35	BB	126	C	N1-C6	-7.84	1.32	1.37
35	BB	1418	C	C3'-C2'	-7.84	1.44	1.52
34	BA	195	G	C1'-N9	-7.83	1.35	1.46
34	BA	483	A	N9-C4	-7.83	1.33	1.37
34	BA	1068	C	C2-N3	-7.83	1.29	1.35
34	BA	1686	G	P-O5'	-7.83	1.51	1.59
35	BB	83	G	O3'-P	-7.83	1.51	1.61
35	BB	631	G	N9-C4	-7.83	1.31	1.38
38	BE	59	U	O3'-P	-7.83	1.51	1.61
34	BA	330	A	C1'-N9	-7.83	1.35	1.46
34	BA	431	A	C5-C4	-7.83	1.33	1.38
34	BA	984	U	C1'-N1	-7.83	1.35	1.46
34	BA	1435	A	C2'-C1'	-7.83	1.44	1.53
35	BB	6	A	C1'-N9	-7.83	1.35	1.46
35	BB	124	G	P-O5'	-7.83	1.51	1.59
35	BB	130	G	C3'-C2'	-7.83	1.44	1.52
35	BB	489	A	N7-C5	-7.83	1.34	1.39
35	BB	569	G	N1-C2	-7.83	1.31	1.37
35	BB	632	U	C2-N3	-7.83	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	53	U	O3'-P	-7.83	1.51	1.61
39	BF	28	C	C4'-C3'	-7.83	1.44	1.53
85	AA	1090	A	O3'-P	-7.83	1.51	1.61
85	AA	2235	C	O3'-P	-7.83	1.51	1.61
86	AB	69	G	C2'-C1'	-7.83	1.44	1.53
34	BA	487	A	C2'-C1'	-7.83	1.44	1.53
34	BA	901	C	C1'-N1	-7.83	1.35	1.46
34	BA	923	C	C2'-C1'	-7.83	1.44	1.53
35	BB	1026	G	N9-C4	-7.83	1.31	1.38
35	BB	1418	C	O3'-P	-7.83	1.51	1.61
85	AA	2182	A	C4'-C3'	-7.83	1.44	1.53
35	BB	1044	U	P-O5'	-7.83	1.51	1.59
41	BH	111	U	N3-C4	-7.83	1.31	1.38
85	AA	716	G	C2'-C1'	-7.83	1.44	1.53
85	AA	2189	U	P-O5'	-7.83	1.51	1.59
34	BA	108	A	C1'-N9	-7.83	1.35	1.46
35	BB	71	A	O3'-P	-7.83	1.51	1.61
35	BB	1396	G	N1-C2	-7.83	1.31	1.37
37	BD	105	G	O3'-P	-7.83	1.51	1.61
38	BE	6	A	C4'-C3'	-7.83	1.44	1.53
40	BG	67	A	P-O5'	-7.83	1.51	1.59
85	AA	1286	C	C2'-C1'	-7.83	1.44	1.53
85	AA	1541	G	C2'-C1'	-7.83	1.44	1.53
85	AA	2186	U	P-O5'	-7.83	1.51	1.59
35	BB	30	A	O3'-P	-7.83	1.51	1.61
35	BB	451	A	C1'-N9	-7.83	1.35	1.46
35	BB	526	A	N9-C4	-7.83	1.33	1.37
35	BB	1242	C	C2-N3	-7.83	1.29	1.35
35	BB	1393	C	O3'-P	-7.83	1.51	1.61
34	BA	1097	G	P-O5'	-7.83	1.51	1.59
35	BB	1058	U	C2'-C1'	-7.83	1.44	1.53
36	BC	166	G	C2-N2	-7.83	1.26	1.34
85	AA	289	G	N9-C4	-7.83	1.31	1.38
85	AA	2039	G	P-O5'	-7.83	1.51	1.59
34	BA	334	G	O3'-P	-7.82	1.51	1.61
34	BA	1119	A	C5'-C4'	-7.82	1.42	1.51
34	BA	1501	U	C3'-C2'	-7.82	1.44	1.52
35	BB	547	A	C1'-N9	-7.82	1.35	1.46
35	BB	1060	U	C2-N3	-7.82	1.32	1.37
35	BB	1103	A	C3'-C2'	-7.82	1.44	1.52
40	BG	164	U	C3'-C2'	-7.82	1.44	1.52
41	BH	129	G	N9-C4	-7.82	1.31	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	354	C	C2-N3	-7.82	1.29	1.35
85	AA	1002	G	C2'-C1'	-7.82	1.44	1.53
85	AA	1575	G	C2-N2	-7.82	1.26	1.34
85	AA	2141	G	N9-C4	-7.82	1.31	1.38
85	AA	2187	G	P-O5'	-7.82	1.51	1.59
34	BA	136	A	C3'-C2'	-7.82	1.44	1.52
34	BA	776	U	C4'-C3'	-7.82	1.44	1.53
36	BC	15	G	C2'-C1'	-7.82	1.44	1.53
40	BG	46	G	C2'-C1'	-7.82	1.44	1.53
34	BA	679	U	O3'-P	-7.82	1.51	1.61
35	BB	43	G	C1'-N9	-7.82	1.35	1.46
85	AA	434	U	P-O5'	-7.82	1.51	1.59
85	AA	663	C	C3'-C2'	-7.82	1.44	1.52
85	AA	976	G	C3'-C2'	-7.82	1.44	1.52
85	AA	1211	C	C3'-C2'	-7.82	1.44	1.52
34	BA	499	C	C2-N3	-7.82	1.29	1.35
34	BA	1573	C	C1'-N1	-7.82	1.35	1.46
85	AA	835	C	O3'-P	-7.82	1.51	1.61
85	AA	2236	U	O3'-P	-7.82	1.51	1.61
34	BA	92	G	N9-C4	-7.82	1.31	1.38
34	BA	98	A	C5-C4	-7.82	1.33	1.38
34	BA	695	A	O3'-P	-7.82	1.51	1.61
34	BA	1664	C	O3'-P	-7.82	1.51	1.61
35	BB	414	C	C2'-C1'	-7.82	1.44	1.53
85	AA	763	U	C3'-C2'	-7.82	1.44	1.52
34	BA	103	G	C5-C4	-7.82	1.32	1.38
34	BA	766	A	N9-C4	-7.82	1.33	1.37
34	BA	1525	G	N7-C5	-7.82	1.34	1.39
35	BB	534	C	P-O5'	-7.82	1.51	1.59
35	BB	661	G	N7-C5	-7.82	1.34	1.39
35	BB	1038	G	C4'-C3'	-7.82	1.44	1.53
35	BB	1341	U	C2-N3	-7.82	1.32	1.37
38	BE	94	U	C2-N3	-7.82	1.32	1.37
40	BG	87	G	C1'-N9	-7.82	1.35	1.46
85	AA	977	U	C2'-C1'	-7.82	1.44	1.53
85	AA	1181	U	P-O5'	-7.82	1.51	1.59
85	AA	1690	A	O3'-P	-7.82	1.51	1.61
34	BA	449	G	C2-N3	-7.81	1.26	1.32
34	BA	461	A	N3-C4	-7.81	1.30	1.34
34	BA	1092	U	C2'-C1'	-7.81	1.44	1.53
34	BA	1472	G	C6-N1	-7.81	1.34	1.39
34	BA	1557	G	N9-C4	-7.81	1.31	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	27	C	C4'-C3'	-7.81	1.44	1.53
37	BD	62	A	O3'-P	-7.81	1.51	1.61
36	BC	2	A	N7-C5	-7.81	1.34	1.39
36	BC	125	A	O3'-P	-7.81	1.51	1.61
40	BG	68	U	O3'-P	-7.81	1.51	1.61
85	AA	914	U	C2-N3	-7.81	1.32	1.37
34	BA	1255	G	N3-C4	-7.81	1.29	1.35
35	BB	400	C	P-O5'	-7.81	1.51	1.59
36	BC	92	C	C2-N3	-7.81	1.29	1.35
39	BF	12	U	C4'-C3'	-7.81	1.44	1.53
35	BB	40	C	C2-N3	-7.81	1.29	1.35
35	BB	690	C	O3'-P	-7.81	1.51	1.61
40	BG	152	G	C6-N1	-7.81	1.34	1.39
85	AA	407	G	C6-N1	-7.81	1.34	1.39
85	AA	816	A	C1'-N9	7.81	1.60	1.48
86	AB	70	G	C4'-C3'	-7.81	1.44	1.53
34	BA	408	U	N1-C2	-7.81	1.31	1.38
34	BA	1284	G	C2'-C1'	-7.81	1.44	1.53
34	BA	1782	C	C2-N3	-7.81	1.29	1.35
35	BB	1261	U	O3'-P	-7.81	1.51	1.61
35	BB	1313	C	C2'-C1'	-7.81	1.44	1.53
39	BF	7	G	O3'-P	-7.81	1.51	1.61
41	BH	111	U	C2'-C1'	-7.81	1.44	1.53
85	AA	448	G	N1-C2	-7.81	1.31	1.37
85	AA	457	G	C3'-C2'	-7.81	1.44	1.52
34	BA	211	C	P-O5'	-7.80	1.51	1.59
34	BA	713	C	O3'-P	-7.80	1.51	1.61
34	BA	1425	G	O3'-P	-7.80	1.51	1.61
35	BB	78	C	C3'-C2'	-7.80	1.44	1.52
35	BB	1175	A	N9-C4	-7.80	1.33	1.37
85	AA	661	C	O3'-P	-7.80	1.51	1.61
85	AA	867	G	C8-N7	-7.80	1.26	1.30
34	BA	167	U	C2'-C1'	-7.80	1.44	1.53
34	BA	931	G	P-O5'	-7.80	1.51	1.59
34	BA	991	U	C2-N3	-7.80	1.32	1.37
34	BA	1657	A	C2'-C1'	-7.80	1.44	1.53
35	BB	379	U	C2'-C1'	-7.80	1.44	1.53
37	BD	7	G	O3'-P	-7.80	1.51	1.61
85	AA	98	U	O3'-P	-7.80	1.51	1.61
85	AA	161	A	P-O5'	-7.80	1.51	1.59
34	BA	248	G	C5'-C4'	7.80	1.60	1.51
34	BA	1522	G	C5-C4	-7.80	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1150	A	C4'-C3'	-7.80	1.44	1.53
85	AA	112	A	N9-C4	-7.80	1.33	1.37
85	AA	1127	G	N9-C8	-7.80	1.32	1.37
34	BA	267	G	C2-N2	-7.80	1.26	1.34
34	BA	839	U	O3'-P	-7.80	1.51	1.61
34	BA	1681	U	P-O5'	-7.80	1.51	1.59
35	BB	130	G	C2'-C1'	-7.80	1.44	1.53
38	BE	195	G	P-O5'	-7.80	1.51	1.59
41	BH	38	G	C1'-N9	-7.80	1.35	1.46
41	BH	57	A	P-O5'	-7.80	1.51	1.59
85	AA	1112	G	P-O5'	-7.80	1.51	1.59
85	AA	2145	G	C5-C4	-7.80	1.32	1.38
35	BB	1072	C	C1'-N1	-7.80	1.35	1.46
35	BB	1483	A	N7-C5	-7.80	1.34	1.39
85	AA	430	G	P-O5'	-7.80	1.51	1.59
34	BA	34	U	N3-C4	-7.80	1.31	1.38
34	BA	842	U	C2'-C1'	-7.80	1.44	1.53
34	BA	857	C	C2-N3	-7.80	1.29	1.35
34	BA	1704	G	C5-C4	-7.80	1.32	1.38
34	BA	1726	U	C5'-C4'	-7.80	1.42	1.51
35	BB	402	G	O3'-P	-7.80	1.51	1.61
35	BB	1118	G	C2'-C1'	-7.80	1.44	1.53
36	BC	18	G	C2-N3	-7.80	1.26	1.32
37	BD	21	G	O3'-P	-7.80	1.51	1.61
41	BH	131	A	P-O5'	-7.80	1.51	1.59
85	AA	166	C	O3'-P	-7.80	1.51	1.61
85	AA	370	A	C2'-C1'	-7.80	1.44	1.53
34	BA	351	A	P-O5'	-7.79	1.51	1.59
34	BA	933	U	O4'-C1'	-7.79	1.31	1.41
34	BA	399	G	C2-N2	-7.79	1.26	1.34
34	BA	425	G	C2-N2	-7.79	1.26	1.34
35	BB	1121	A	N7-C5	-7.79	1.34	1.39
85	AA	167	A	N9-C4	-7.79	1.33	1.37
85	AA	355	G	O3'-P	-7.79	1.51	1.61
85	AA	2072	G	N7-C5	-7.79	1.34	1.39
34	BA	527	C	P-O5'	-7.79	1.51	1.59
35	BB	104	G	C3'-C2'	-7.79	1.44	1.52
35	BB	474	G	C1'-N9	-7.79	1.35	1.46
35	BB	647	U	N3-C4	-7.79	1.31	1.38
35	BB	1274	G	N9-C4	-7.79	1.31	1.38
85	AA	42	G	O3'-P	-7.79	1.51	1.61
34	BA	24	C	C3'-C2'	-7.79	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	38	G	N7-C5	-7.79	1.34	1.39
34	BA	196	A	C3'-C2'	-7.79	1.44	1.52
34	BA	1425	G	N7-C5	-7.79	1.34	1.39
34	BA	1611	A	O3'-P	-7.79	1.51	1.61
37	BD	79	G	C2'-C1'	-7.79	1.44	1.53
85	AA	679	A	C1'-N9	-7.79	1.35	1.46
34	BA	737	U	C2'-C1'	-7.79	1.44	1.53
34	BA	796	G	N9-C8	-7.79	1.32	1.37
34	BA	1230	G	N9-C4	-7.79	1.31	1.38
34	BA	1820	G	N3-C4	-7.79	1.29	1.35
35	BB	14	C	C4'-C3'	-7.79	1.44	1.53
35	BB	665	A	C6-N1	-7.79	1.30	1.35
35	BB	1189	C	P-O5'	-7.79	1.51	1.59
37	BD	116	C	O3'-P	-7.79	1.51	1.61
34	BA	290	G	N1-C2	-7.79	1.31	1.37
34	BA	746	C	O3'-P	-7.79	1.51	1.61
34	BA	1608	C	C2-N3	-7.79	1.29	1.35
35	BB	28	G	N9-C4	-7.79	1.31	1.38
35	BB	1047	C	C2'-C1'	-7.79	1.44	1.53
34	BA	1539	A	N9-C4	-7.79	1.33	1.37
35	BB	392	G	C5-C4	-7.79	1.32	1.38
35	BB	1363	A	N9-C4	-7.79	1.33	1.37
85	AA	19	A	O3'-P	-7.79	1.51	1.61
85	AA	197	C	C4'-C3'	7.79	1.61	1.53
34	BA	524	G	N1-C2	-7.78	1.31	1.37
35	BB	615	A	P-O5'	-7.78	1.51	1.59
35	BB	674	C	O3'-P	-7.78	1.51	1.61
85	AA	248	U	P-O5'	-7.78	1.51	1.59
34	BA	1810	A	O3'-P	-7.78	1.51	1.61
35	BB	1029	U	C2-N3	-7.78	1.32	1.37
85	AA	452	A	C1'-N9	-7.78	1.35	1.46
85	AA	800	A	C2'-C1'	-7.78	1.44	1.53
34	BA	141	G	O3'-P	-7.78	1.51	1.61
35	BB	144	G	P-O5'	-7.78	1.51	1.59
35	BB	1263	A	P-O5'	-7.78	1.51	1.59
40	BG	94	G	C1'-N9	-7.78	1.35	1.46
85	AA	679	A	C3'-C2'	-7.78	1.44	1.52
85	AA	760	U	C2'-C1'	-7.78	1.44	1.53
85	AA	1147	A	C2'-C1'	-7.78	1.44	1.53
85	AA	1486	G	O3'-P	-7.78	1.51	1.61
34	BA	700	G	O3'-P	-7.78	1.51	1.61
35	BB	607	G	C2-N2	-7.78	1.26	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	7	G	N7-C5	-7.78	1.34	1.39
37	BD	110	G	C3'-C2'	-7.78	1.44	1.52
85	AA	866	U	O3'-P	-7.78	1.51	1.61
85	AA	1165	C	C3'-C2'	-7.78	1.44	1.52
34	BA	522	C	P-O5'	-7.78	1.51	1.59
35	BB	100	A	C2'-C1'	-7.78	1.44	1.53
35	BB	1128	U	C2'-C1'	-7.78	1.44	1.53
36	BC	54	G	C8-N7	-7.78	1.26	1.30
40	BG	157	A	C6-N1	-7.78	1.30	1.35
85	AA	773	G	C3'-C2'	-7.78	1.44	1.52
34	BA	727	G	P-O5'	-7.77	1.51	1.59
34	BA	1495	A	O3'-P	-7.77	1.51	1.61
35	BB	65	A	C1'-N9	-7.77	1.35	1.46
35	BB	1269	A	O3'-P	-7.77	1.51	1.61
37	BD	97	U	O3'-P	-7.77	1.51	1.61
34	BA	49	A	C4'-O4'	-7.77	1.35	1.45
34	BA	323	C	C2-N3	-7.77	1.29	1.35
35	BB	826	G	C2'-C1'	-7.77	1.44	1.53
35	BB	977	G	P-O5'	-7.77	1.51	1.59
35	BB	1110	G	C6-N1	-7.77	1.34	1.39
36	BC	44	A	C5-C4	-7.77	1.33	1.38
41	BH	123	G	C1'-N9	-7.77	1.35	1.46
85	AA	970	U	C1'-N1	7.77	1.60	1.48
38	BE	107	U	C2'-C1'	-7.77	1.44	1.53
85	AA	1491	G	C2-N2	-7.77	1.26	1.34
34	BA	176	G	C2'-C1'	-7.77	1.44	1.53
34	BA	494	A	O3'-P	-7.77	1.51	1.61
34	BA	548	G	N7-C5	-7.77	1.34	1.39
34	BA	902	C	C1'-N1	-7.77	1.35	1.46
38	BE	136	G	O4'-C1'	-7.77	1.31	1.41
39	BF	51	C	O3'-P	-7.77	1.51	1.61
40	BG	92	U	C3'-C2'	-7.77	1.44	1.52
85	AA	2164	G	C2'-C1'	-7.77	1.44	1.53
34	BA	482	C	N1-C2	-7.77	1.32	1.40
34	BA	1736	A	P-O5'	-7.77	1.51	1.59
35	BB	1293	C	C2'-C1'	-7.77	1.44	1.53
40	BG	102	G	N1-C2	-7.77	1.31	1.37
85	AA	92	G	N9-C4	-7.77	1.31	1.38
85	AA	1224	C	C3'-C2'	-7.77	1.44	1.52
85	AA	1490	A	N7-C5	-7.77	1.34	1.39
34	BA	504	A	C5-C4	-7.77	1.33	1.38
34	BA	1011	G	C5-C4	-7.77	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1040	G	C2'-C1'	-7.77	1.44	1.53
85	AA	422	G	C3'-C2'	-7.77	1.44	1.52
85	AA	2006	G	N7-C5	-7.77	1.34	1.39
34	BA	245	U	O3'-P	-7.76	1.51	1.61
35	BB	1178	A	C1'-N9	-7.76	1.35	1.46
35	BB	1232	A	C3'-C2'	-7.76	1.44	1.52
35	BB	1455	A	C2'-C1'	-7.76	1.44	1.53
35	BB	1505	U	O3'-P	-7.76	1.51	1.61
37	BD	11	A	C4'-C3'	-7.76	1.44	1.53
41	BH	38	G	C5-C6	-7.76	1.34	1.42
41	BH	109	G	O3'-P	-7.76	1.51	1.61
85	AA	1597	C	C2'-C1'	-7.76	1.44	1.53
85	AA	1662	U	P-O5'	-7.76	1.51	1.59
34	BA	397	A	C2'-C1'	-7.76	1.44	1.53
34	BA	1823	A	N7-C5	-7.76	1.34	1.39
35	BB	798	A	C8-N7	-7.76	1.26	1.31
38	BE	92	C	C3'-C2'	-7.76	1.44	1.52
85	AA	1473	U	C1'-N1	-7.76	1.35	1.46
34	BA	53	G	C1'-N9	-7.76	1.35	1.46
34	BA	189	G	N9-C8	-7.76	1.32	1.37
34	BA	246	G	P-O5'	-7.76	1.51	1.59
34	BA	761	U	C3'-C2'	-7.76	1.44	1.52
34	BA	1674	G	C2'-C1'	-7.76	1.44	1.53
34	BA	1731	A	P-O5'	-7.76	1.51	1.59
35	BB	547	A	P-O5'	-7.76	1.51	1.59
35	BB	1105	G	O3'-P	-7.76	1.51	1.61
35	BB	1423	U	C3'-C2'	-7.76	1.44	1.52
35	BB	1478	G	O3'-P	-7.76	1.51	1.61
36	BC	75	G	C3'-C2'	-7.76	1.44	1.52
85	AA	702	G	N7-C5	-7.76	1.34	1.39
85	AA	708	G	N9-C4	-7.76	1.31	1.38
85	AA	713	G	O3'-P	-7.76	1.51	1.61
34	BA	167	U	C2-N3	-7.76	1.32	1.37
34	BA	500	C	C2'-C1'	-7.76	1.44	1.53
34	BA	997	U	C3'-C2'	-7.76	1.44	1.52
34	BA	1493	U	C4'-C3'	7.76	1.61	1.53
35	BB	666	A	P-O5'	-7.76	1.51	1.59
40	BG	65	C	C4'-C3'	-7.76	1.44	1.53
35	BB	773	G	O3'-P	-7.76	1.51	1.61
38	BE	113	C	O3'-P	-7.76	1.51	1.61
34	BA	131	A	P-O5'	-7.76	1.51	1.59
34	BA	669	U	C2-N3	-7.76	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	683	C	C2-N3	7.76	1.42	1.35
34	BA	747	G	O3'-P	-7.76	1.51	1.61
34	BA	782	C	C3'-C2'	-7.76	1.44	1.52
34	BA	1029	C	P-O5'	-7.76	1.51	1.59
34	BA	1260	G	C1'-N9	-7.76	1.35	1.46
34	BA	1521	C	C3'-C2'	-7.76	1.44	1.52
35	BB	829	C	O3'-P	-7.76	1.51	1.61
35	BB	1027	U	P-O5'	-7.76	1.51	1.59
34	BA	142	A	C1'-N9	-7.75	1.35	1.46
34	BA	937	G	N9-C4	-7.75	1.31	1.38
34	BA	1127	U	P-O5'	-7.75	1.51	1.59
85	AA	247	G	O3'-P	-7.75	1.51	1.61
85	AA	501	A	O3'-P	-7.75	1.51	1.61
34	BA	300	C	P-O5'	-7.75	1.51	1.59
34	BA	973	U	P-O5'	-7.75	1.51	1.59
34	BA	1025	A	C8-N7	-7.75	1.26	1.31
34	BA	1059	U	O3'-P	-7.75	1.51	1.61
34	BA	1682	A	P-O5'	-7.75	1.51	1.59
35	BB	1199	A	C1'-N9	-7.75	1.35	1.46
35	BB	1517	G	N9-C4	-7.75	1.31	1.38
38	BE	96	G	C2-N3	-7.75	1.26	1.32
64	Be	131	GLY	CA-C	-7.75	1.39	1.51
85	AA	313	A	O3'-P	-7.75	1.51	1.61
85	AA	419	A	C5-C4	-7.75	1.33	1.38
85	AA	1520	A	N3-C4	-7.75	1.30	1.34
85	AA	2167	A	O3'-P	-7.75	1.51	1.61
34	BA	76	U	P-O5'	-7.75	1.51	1.59
34	BA	1790	U	C1'-N1	-7.75	1.35	1.46
35	BB	1399	A	N9-C8	-7.75	1.31	1.37
39	BF	23	G	C5'-C4'	7.75	1.60	1.51
85	AA	15	U	P-O5'	-7.75	1.51	1.59
34	BA	780	U	C4'-O4'	7.75	1.55	1.45
34	BA	1194	G	C2'-C1'	-7.75	1.44	1.53
35	BB	114	A	P-O5'	-7.75	1.51	1.59
85	AA	882	C	C2-N3	-7.75	1.29	1.35
85	AA	1235	G	N9-C4	-7.75	1.31	1.38
34	BA	381	A	C5-C4	-7.75	1.33	1.38
34	BA	727	G	N9-C8	-7.75	1.32	1.37
34	BA	776	U	C3'-C2'	-7.75	1.44	1.52
34	BA	904	G	N9-C8	-7.75	1.32	1.37
34	BA	925	G	O3'-P	-7.75	1.51	1.61
34	BA	1073	G	C6-N1	-7.75	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1716	A	P-O5'	7.75	1.67	1.59
35	BB	661	G	P-O5'	-7.75	1.52	1.59
38	BE	122	G	C1'-N9	-7.75	1.36	1.46
39	BF	60	C	C2'-C1'	-7.75	1.44	1.53
40	BG	38	A	C5-C4	-7.75	1.33	1.38
41	BH	38	G	P-O5'	-7.75	1.52	1.59
85	AA	507	C	C2'-C1'	-7.75	1.44	1.53
85	AA	652	U	P-O5'	-7.75	1.52	1.59
85	AA	2200	A	C1'-N9	-7.75	1.36	1.46
34	BA	407	A	N7-C5	-7.75	1.34	1.39
34	BA	1154	U	N3-C4	-7.75	1.31	1.38
34	BA	1423	U	O3'-P	-7.75	1.51	1.61
36	BC	16	A	C5-C4	-7.75	1.33	1.38
37	BD	36	C	O3'-P	-7.75	1.51	1.61
85	AA	683	U	O3'-P	-7.75	1.51	1.61
85	AA	2088	U	P-O5'	-7.75	1.52	1.59
85	AA	2208	G	N7-C5	-7.75	1.34	1.39
34	BA	611	A	N7-C5	-7.75	1.34	1.39
34	BA	1551	G	C2'-C1'	-7.75	1.44	1.53
35	BB	377	A	P-O5'	-7.75	1.52	1.59
35	BB	1216	G	C3'-C2'	-7.75	1.44	1.52
38	BE	121	G	O3'-P	-7.75	1.51	1.61
85	AA	25	C	O3'-P	-7.75	1.51	1.61
34	BA	89	G	C5-C4	-7.74	1.32	1.38
34	BA	1220	C	C1'-N1	-7.74	1.36	1.46
34	BA	1254	C	O3'-P	-7.74	1.51	1.61
34	BA	1505	G	C1'-N9	-7.74	1.36	1.46
34	BA	1544	G	C2'-C1'	-7.74	1.44	1.53
35	BB	41	A	O3'-P	-7.74	1.51	1.61
35	BB	433	C	C2-N3	-7.74	1.29	1.35
35	BB	1045	G	O3'-P	-7.74	1.51	1.61
36	BC	68	A	N9-C4	-7.74	1.33	1.37
38	BE	148	C	C2-N3	-7.74	1.29	1.35
39	BF	50	C	C3'-C2'	-7.74	1.44	1.52
39	BF	62	U	C2'-C1'	-7.74	1.44	1.53
85	AA	402	G	C6-N1	-7.74	1.34	1.39
85	AA	1119	A	C3'-C2'	-7.74	1.44	1.52
40	BG	90	G	C6-N1	-7.74	1.34	1.39
34	BA	545	U	C2-N3	-7.74	1.32	1.37
34	BA	614	A	C2'-C1'	-7.74	1.44	1.53
34	BA	813	C	C2-N3	-7.74	1.29	1.35
34	BA	907	A	N7-C5	-7.74	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	30	A	P-O5'	-7.74	1.52	1.59
35	BB	442	U	C1'-N1	-7.74	1.36	1.46
85	AA	1469	G	C4'-C3'	-7.74	1.44	1.53
85	AA	1544	G	O3'-P	-7.74	1.51	1.61
86	AB	22	G	N9-C4	-7.74	1.31	1.38
34	BA	69	C	N3-C4	-7.74	1.28	1.33
34	BA	1062	G	N7-C5	-7.74	1.34	1.39
35	BB	633	C	C3'-C2'	-7.74	1.44	1.52
34	BA	962	U	P-O5'	-7.74	1.52	1.59
40	BG	53	C	P-O5'	-7.74	1.52	1.59
34	BA	112	C	C2-N3	-7.74	1.29	1.35
34	BA	1224	A	C2'-C1'	-7.74	1.44	1.53
35	BB	1081	U	C2'-C1'	-7.74	1.44	1.53
35	BB	1400	C	P-O5'	-7.74	1.52	1.59
65	Bf	324	GLY	CA-C	-7.74	1.39	1.51
85	AA	1681	G	C4'-C3'	-7.74	1.44	1.53
85	AA	1831	U	C2'-C1'	-7.74	1.44	1.53
35	BB	611	U	C3'-C2'	-7.73	1.44	1.52
34	BA	257	G	C6-N1	-7.73	1.34	1.39
34	BA	380	A	O3'-P	-7.73	1.51	1.61
34	BA	1673	G	N1-C2	-7.73	1.31	1.37
35	BB	1210	U	C2'-C1'	-7.73	1.44	1.53
39	BF	46	G	C3'-C2'	-7.73	1.44	1.52
40	BG	127	G	C2-N2	-7.73	1.26	1.34
85	AA	452	A	P-O5'	-7.73	1.52	1.59
34	BA	482	C	C2-N3	-7.73	1.29	1.35
35	BB	697	G	C2'-C1'	-7.73	1.44	1.53
40	BG	176	G	C5-C4	-7.73	1.32	1.38
41	BH	23	G	N7-C5	-7.73	1.34	1.39
85	AA	274	A	C3'-C2'	-7.73	1.44	1.52
85	AA	685	U	C4'-C3'	-7.73	1.44	1.53
85	AA	1257	A	C5-C4	-7.73	1.33	1.38
85	AA	1560	A	C8-N7	-7.73	1.26	1.31
35	BB	1148	U	O4'-C1'	-7.73	1.31	1.41
35	BB	1368	A	O3'-P	-7.73	1.51	1.61
35	BB	1437	U	P-O5'	-7.73	1.52	1.59
85	AA	804	A	O3'-P	-7.73	1.51	1.61
85	AA	1464	G	C5-C4	-7.73	1.32	1.38
34	BA	993	C	O3'-P	-7.73	1.51	1.61
34	BA	1102	A	P-O5'	-7.73	1.52	1.59
34	BA	1260	G	C2'-C1'	-7.73	1.44	1.53
34	BA	1451	A	O3'-P	-7.73	1.51	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1658	G	C3'-C2'	-7.73	1.44	1.52
35	BB	132	G	C1'-N9	-7.73	1.36	1.46
35	BB	136	A	N9-C4	-7.73	1.33	1.37
35	BB	1232	A	N9-C4	-7.73	1.33	1.37
35	BB	1464	G	N9-C4	-7.73	1.31	1.38
38	BE	26	G	O4'-C1'	-7.73	1.31	1.41
85	AA	1707	G	N9-C4	-7.73	1.31	1.38
35	BB	678	U	O3'-P	-7.73	1.51	1.61
40	BG	15	G	C6-N1	-7.73	1.34	1.39
40	BG	116	G	C2'-C1'	-7.73	1.44	1.53
85	AA	2128	G	C5-C4	-7.73	1.32	1.38
35	BB	582	G	P-O5'	-7.72	1.52	1.59
35	BB	1005	A	C1'-N9	-7.72	1.36	1.46
36	BC	62	A	P-O5'	-7.72	1.52	1.59
85	AA	1229	G	C3'-C2'	-7.72	1.44	1.52
34	BA	102	G	C2'-C1'	-7.72	1.44	1.53
34	BA	914	G	O3'-P	-7.72	1.51	1.61
34	BA	1179	U	P-O5'	-7.72	1.52	1.59
34	BA	1234	U	C2-N3	-7.72	1.32	1.37
34	BA	1803	A	C5-C4	-7.72	1.33	1.38
35	BB	393	A	C5-C4	-7.72	1.33	1.38
36	BC	44	A	C2'-C1'	-7.72	1.44	1.53
85	AA	1481	U	O3'-P	-7.72	1.51	1.61
34	BA	338	U	C2-N3	-7.72	1.32	1.37
34	BA	702	G	C6-N1	-7.72	1.34	1.39
34	BA	739	A	N7-C5	-7.72	1.34	1.39
35	BB	975	G	O3'-P	-7.72	1.51	1.61
37	BD	3	G	C5-C4	-7.72	1.32	1.38
41	BH	108	U	P-O5'	-7.72	1.52	1.59
34	BA	1325	G	C3'-C2'	-7.72	1.44	1.52
34	BA	1431	G	C1'-N9	-7.72	1.36	1.46
34	BA	1649	A	P-O5'	-7.72	1.52	1.59
35	BB	680	A	C1'-N9	-7.72	1.36	1.46
85	AA	705	G	C1'-N9	-7.72	1.36	1.46
34	BA	968	G	P-O5'	-7.72	1.52	1.59
34	BA	1195	G	C1'-N9	-7.72	1.36	1.46
35	BB	403	U	C2-N3	-7.72	1.32	1.37
35	BB	1051	U	N3-C4	-7.72	1.31	1.38
34	BA	23	A	C5-C4	-7.72	1.33	1.38
34	BA	47	U	P-O5'	-7.72	1.52	1.59
34	BA	369	A	P-O5'	-7.72	1.52	1.59
34	BA	920	U	C2'-C1'	-7.72	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	944	G	P-O5'	-7.72	1.52	1.59
34	BA	1243	A	P-O5'	-7.72	1.52	1.59
34	BA	1302	C	P-O5'	-7.72	1.52	1.59
35	BB	65	A	P-O5'	-7.72	1.52	1.59
35	BB	380	G	O3'-P	-7.72	1.51	1.61
35	BB	624	A	C2'-C1'	-7.72	1.44	1.53
85	AA	160	A	N7-C5	-7.72	1.34	1.39
85	AA	427	G	N1-C2	-7.72	1.31	1.37
85	AA	660	G	P-O5'	-7.72	1.52	1.59
85	AA	891	G	O3'-P	-7.72	1.51	1.61
85	AA	944	C	C4-N4	-7.72	1.27	1.33
85	AA	1103	A	C5'-C4'	7.72	1.60	1.51
85	AA	1575	G	O3'-P	-7.72	1.51	1.61
85	AA	2151	U	C2'-C1'	-7.72	1.44	1.53
34	BA	6	C	C2'-C1'	-7.71	1.44	1.53
34	BA	1113	A	O3'-P	-7.71	1.51	1.61
35	BB	879	G	C6-N1	-7.71	1.34	1.39
36	BC	63	G	P-O5'	-7.71	1.52	1.59
85	AA	1932	C	C5'-C4'	7.71	1.60	1.51
34	BA	434	U	C5'-C4'	7.71	1.60	1.51
35	BB	41	A	O4'-C1'	-7.71	1.31	1.41
35	BB	93	A	N9-C8	-7.71	1.31	1.37
35	BB	1053	G	O3'-P	-7.71	1.51	1.61
36	BC	45	C	O3'-P	-7.71	1.51	1.61
38	BE	50	G	C5-C4	-7.71	1.32	1.38
40	BG	142	A	O3'-P	-7.71	1.51	1.61
85	AA	2195	A	C2'-C1'	-7.71	1.44	1.53
34	BA	1551	G	C1'-N9	-7.71	1.36	1.46
34	BA	1641	G	C4'-O4'	-7.71	1.35	1.45
35	BB	387	G	C2'-C1'	-7.71	1.44	1.53
35	BB	496	C	C2'-C1'	-7.71	1.44	1.53
35	BB	1058	U	O3'-P	-7.71	1.51	1.61
85	AA	960	G	O3'-P	-7.71	1.51	1.61
85	AA	2234	C	C2'-C1'	-7.71	1.44	1.53
34	BA	27	G	N1-C2	-7.71	1.31	1.37
34	BA	322	U	P-O5'	-7.71	1.52	1.59
34	BA	611	A	C2'-C1'	-7.71	1.44	1.53
35	BB	1314	G	N1-C2	-7.71	1.31	1.37
85	AA	803	C	C2'-C1'	-7.71	1.44	1.53
85	AA	2202	G	C2'-C1'	-7.71	1.44	1.53
34	BA	36	A	N9-C4	-7.71	1.33	1.37
34	BA	734	G	N9-C4	-7.71	1.31	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	492	U	O3'-P	-7.71	1.51	1.61
35	BB	596	C	P-O5'	-7.71	1.52	1.59
35	BB	986	C	P-O5'	-7.71	1.52	1.59
85	AA	289	G	C3'-C2'	-7.71	1.44	1.52
34	BA	63	A	C3'-C2'	-7.71	1.44	1.52
34	BA	135	G	O3'-P	-7.71	1.51	1.61
34	BA	1511	C	C2'-C1'	-7.71	1.44	1.53
35	BB	1249	G	O3'-P	-7.71	1.51	1.61
35	BB	1425	A	C5-C4	-7.71	1.33	1.38
40	BG	175	G	C6-N1	-7.71	1.34	1.39
85	AA	538	A	N7-C5	-7.71	1.34	1.39
85	AA	852	C	P-O5'	-7.71	1.52	1.59
85	AA	2117	U	O3'-P	-7.71	1.51	1.61
34	BA	476	U	O3'-P	-7.71	1.51	1.61
34	BA	1023	G	N7-C5	-7.71	1.34	1.39
38	BE	130	G	N9-C4	7.71	1.44	1.38
40	BG	151	A	N9-C4	-7.71	1.33	1.37
85	AA	432	A	C5-C4	-7.71	1.33	1.38
34	BA	1458	A	N9-C4	-7.70	1.33	1.37
35	BB	1025	A	P-O5'	7.70	1.67	1.59
35	BB	1258	G	O3'-P	-7.70	1.51	1.61
35	BB	1400	C	C2'-C1'	-7.70	1.44	1.53
35	BB	1537	C	C2'-C1'	-7.70	1.44	1.53
85	AA	366	A	N3-C4	-7.70	1.30	1.34
85	AA	489	C	O3'-P	-7.70	1.51	1.61
85	AA	2237	G	N1-C2	-7.70	1.31	1.37
34	BA	475	A	P-O5'	-7.70	1.52	1.59
85	AA	85	U	O3'-P	-7.70	1.51	1.61
34	BA	495	A	N9-C4	-7.70	1.33	1.37
34	BA	1091	U	N3-C4	-7.70	1.31	1.38
34	BA	1097	G	N7-C5	-7.70	1.34	1.39
34	BA	1428	G	C6-N1	-7.70	1.34	1.39
34	BA	1683	C	C2-N3	-7.70	1.29	1.35
35	BB	89	C	O3'-P	-7.70	1.51	1.61
35	BB	988	G	C2'-C1'	-7.70	1.44	1.53
35	BB	1021	C	O3'-P	-7.70	1.51	1.61
35	BB	1283	C	C3'-C2'	-7.70	1.44	1.52
36	BC	120	G	C8-N7	-7.70	1.26	1.30
85	AA	1814	U	C2-N3	-7.70	1.32	1.37
34	BA	1435	A	N9-C8	-7.70	1.31	1.37
34	BA	1632	G	N7-C5	-7.70	1.34	1.39
35	BB	1161	G	C1'-N9	-7.70	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	186	U	C2-N3	-7.70	1.32	1.37
85	AA	1016	G	O3'-P	-7.70	1.51	1.61
34	BA	1110	A	C5-C4	-7.70	1.33	1.38
34	BA	1518	A	C5-C6	-7.70	1.34	1.41
35	BB	434	A	C5-C4	-7.70	1.33	1.38
34	BA	234	A	O3'-P	-7.70	1.51	1.61
34	BA	893	U	C3'-C2'	-7.70	1.44	1.52
34	BA	912	G	C4'-C3'	-7.70	1.44	1.53
35	BB	467	G	O3'-P	-7.70	1.51	1.61
36	BC	4	G	C4'-C3'	-7.70	1.44	1.53
38	BE	120	C	O3'-P	-7.70	1.51	1.61
85	AA	96	C	O3'-P	-7.70	1.51	1.61
85	AA	977	U	O3'-P	-7.70	1.51	1.61
85	AA	1109	G	C2'-C1'	-7.70	1.44	1.53
34	BA	1193	A	C2'-C1'	-7.69	1.44	1.53
34	BA	1830	A	O3'-P	-7.69	1.51	1.61
35	BB	829	C	C2'-C1'	-7.69	1.44	1.53
35	BB	1014	U	C2-N3	-7.69	1.32	1.37
35	BB	1470	G	N9-C4	-7.69	1.31	1.38
40	BG	179	C	C3'-C2'	-7.69	1.44	1.52
85	AA	177	A	C2'-C1'	-7.69	1.44	1.53
85	AA	2064	A	N9-C4	-7.69	1.33	1.37
34	BA	189	G	C2'-C1'	-7.69	1.44	1.53
34	BA	399	G	C2'-C1'	-7.69	1.44	1.53
34	BA	707	C	C2'-C1'	-7.69	1.44	1.53
34	BA	1115	A	P-O5'	-7.69	1.52	1.59
34	BA	1696	G	N7-C5	-7.69	1.34	1.39
34	BA	1703	A	P-O5'	-7.69	1.52	1.59
35	BB	1419	G	O3'-P	-7.69	1.51	1.61
40	BG	130	G	N9-C8	-7.69	1.32	1.37
85	AA	491	G	C6-N1	-7.69	1.34	1.39
85	AA	1460	G	N7-C5	-7.69	1.34	1.39
85	AA	1641	A	O3'-P	-7.69	1.51	1.61
34	BA	382	G	N1-C2	-7.69	1.31	1.37
34	BA	426	A	N7-C5	-7.69	1.34	1.39
35	BB	41	A	C5-C4	-7.69	1.33	1.38
35	BB	375	G	C5-C4	-7.69	1.32	1.38
35	BB	1030	U	P-O5'	-7.69	1.52	1.59
35	BB	1046	C	P-O5'	-7.69	1.52	1.59
37	BD	71	G	C6-N1	-7.69	1.34	1.39
85	AA	450	A	C3'-C2'	-7.69	1.44	1.52
85	AA	1959	G	C2'-C1'	-7.69	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2141	G	O3'-P	-7.69	1.51	1.61
34	BA	1139	G	O3'-P	-7.69	1.51	1.61
35	BB	14	C	C3'-C2'	-7.69	1.44	1.52
35	BB	1086	G	N9-C4	-7.69	1.31	1.38
35	BB	1177	U	C2'-C1'	-7.69	1.44	1.53
41	BH	16	A	N9-C8	-7.69	1.31	1.37
85	AA	56	U	C5'-C4'	7.69	1.60	1.51
85	AA	246	C	P-O5'	-7.69	1.52	1.59
34	BA	93	A	N7-C5	-7.69	1.34	1.39
34	BA	752	A	C4'-O4'	-7.69	1.35	1.45
35	BB	642	G	P-O5'	-7.69	1.52	1.59
85	AA	2193	A	C5-C6	-7.69	1.34	1.41
34	BA	934	G	N9-C8	-7.69	1.32	1.37
35	BB	490	G	N9-C8	-7.69	1.32	1.37
35	BB	543	G	O3'-P	-7.69	1.51	1.61
85	AA	498	C	P-O5'	-7.69	1.52	1.59
34	BA	934	G	C5-C4	-7.68	1.32	1.38
35	BB	584	A	P-O5'	-7.68	1.52	1.59
35	BB	661	G	C6-N1	-7.68	1.34	1.39
35	BB	795	A	C3'-C2'	-7.68	1.44	1.52
36	BC	9	G	N7-C5	-7.68	1.34	1.39
41	BH	29	G	C2-N2	-7.68	1.26	1.34
41	BH	128	G	C1'-N9	-7.68	1.36	1.46
85	AA	363	A	N3-C4	-7.68	1.30	1.34
85	AA	1479	U	C3'-C2'	-7.68	1.44	1.52
34	BA	791	A	N9-C4	-7.68	1.33	1.37
34	BA	965	A	N7-C5	-7.68	1.34	1.39
34	BA	1021	U	P-O5'	-7.68	1.52	1.59
34	BA	1474	G	N1-C2	-7.68	1.31	1.37
35	BB	622	G	N9-C4	-7.68	1.31	1.38
35	BB	661	G	C2'-C1'	-7.68	1.44	1.53
37	BD	110	G	O3'-P	-7.68	1.51	1.61
85	AA	349	C	P-O5'	-7.68	1.52	1.59
85	AA	648	G	O3'-P	-7.68	1.51	1.61
86	AB	50	U	P-O5'	-7.68	1.52	1.59
34	BA	381	A	P-O5'	-7.68	1.52	1.59
35	BB	496	C	C3'-C2'	-7.68	1.44	1.52
85	AA	51	A	N7-C5	-7.68	1.34	1.39
85	AA	1357	U	C4'-C3'	-7.68	1.44	1.53
34	BA	674	G	C1'-N9	-7.68	1.36	1.46
34	BA	1008	A	N9-C4	-7.68	1.33	1.37
34	BA	1075	U	C3'-C2'	-7.68	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1232	C	C3'-C2'	-7.68	1.44	1.52
34	BA	1330	G	P-O5'	-7.68	1.52	1.59
35	BB	275	A	N9-C4	-7.68	1.33	1.37
36	BC	135	A	C2'-C1'	-7.68	1.45	1.53
38	BE	23	G	C2'-C1'	-7.68	1.45	1.53
85	AA	896	C	C2'-C1'	-7.68	1.45	1.53
85	AA	1618	G	P-O5'	-7.68	1.52	1.59
34	BA	412	G	N7-C5	-7.68	1.34	1.39
35	BB	506	G	O3'-P	-7.68	1.51	1.61
35	BB	1469	A	C1'-N9	-7.68	1.36	1.46
37	BD	7	G	C2-N2	-7.68	1.26	1.34
34	BA	425	G	P-O5'	-7.68	1.52	1.59
34	BA	763	U	O4'-C1'	-7.68	1.31	1.41
34	BA	1203	G	N9-C8	-7.68	1.32	1.37
34	BA	1390	C	O3'-P	-7.68	1.51	1.61
35	BB	19	C	C2'-C1'	-7.68	1.45	1.53
35	BB	442	U	O4'-C1'	-7.68	1.31	1.41
35	BB	1141	A	C2'-C1'	-7.68	1.45	1.53
35	BB	1210	U	C3'-C2'	-7.68	1.44	1.52
35	BB	1252	G	C6-N1	-7.68	1.34	1.39
37	BD	96	C	C2'-C1'	-7.68	1.45	1.53
40	BG	113	G	N9-C4	-7.68	1.31	1.38
41	BH	128	G	C2-N2	-7.68	1.26	1.34
85	AA	892	C	O3'-P	-7.68	1.51	1.61
85	AA	1242	A	C2'-C1'	-7.68	1.45	1.53
34	BA	15	G	C1'-N9	-7.67	1.36	1.46
34	BA	615	A	C2'-C1'	-7.67	1.45	1.53
35	BB	709	G	C3'-C2'	-7.67	1.44	1.52
35	BB	1292	G	O3'-P	-7.67	1.51	1.61
35	BB	1351	G	C3'-C2'	-7.67	1.44	1.52
38	BE	28	C	C1'-N1	-7.67	1.36	1.46
40	BG	182	G	N9-C4	-7.67	1.31	1.38
85	AA	159	G	C1'-N9	-7.67	1.36	1.46
85	AA	161	A	C3'-C2'	-7.67	1.44	1.52
34	BA	1104	C	O3'-P	-7.67	1.51	1.61
35	BB	1054	G	N1-C2	-7.67	1.31	1.37
41	BH	38	G	C2-N2	-7.67	1.26	1.34
34	BA	491	U	C2-N3	-7.67	1.32	1.37
34	BA	547	C	P-O5'	7.67	1.67	1.59
35	BB	28	G	O3'-P	-7.67	1.51	1.61
35	BB	666	A	O3'-P	-7.67	1.51	1.61
35	BB	1430	G	C3'-C2'	-7.67	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	897	A	C3'-C2'	-7.67	1.44	1.52
85	AA	939	A	C2'-C1'	-7.67	1.45	1.53
85	AA	1165	C	C2'-C1'	-7.67	1.45	1.53
85	AA	1262	A	O3'-P	-7.67	1.51	1.61
34	BA	177	G	P-O5'	-7.67	1.52	1.59
34	BA	907	A	C1'-N9	-7.67	1.36	1.46
36	BC	116	C	O3'-P	-7.67	1.51	1.61
85	AA	2019	G	N9-C4	-7.67	1.31	1.38
34	BA	735	A	O3'-P	-7.67	1.51	1.61
34	BA	900	A	O3'-P	-7.67	1.51	1.61
34	BA	965	A	P-O5'	-7.67	1.52	1.59
34	BA	1435	A	O3'-P	-7.67	1.51	1.61
35	BB	782	A	C2'-C1'	-7.67	1.45	1.53
35	BB	1270	C	O3'-P	-7.67	1.51	1.61
40	BG	153	C	P-O5'	-7.67	1.52	1.59
85	AA	813	G	C2'-C1'	-7.67	1.45	1.53
85	AA	1261	U	P-O5'	-7.67	1.52	1.59
85	AA	2191	C	C2-N3	-7.67	1.29	1.35
34	BA	567	U	P-O5'	-7.67	1.52	1.59
34	BA	816	G	C5-C6	-7.67	1.34	1.42
34	BA	1641	G	O4'-C1'	-7.67	1.31	1.41
35	BB	391	G	C5-C4	-7.67	1.32	1.38
35	BB	426	A	C2'-C1'	-7.67	1.45	1.53
35	BB	441	G	C3'-C2'	-7.67	1.44	1.52
35	BB	449	C	C2-N3	-7.67	1.29	1.35
35	BB	1462	G	C3'-O3'	7.67	1.52	1.42
37	BD	24	U	N1-C2	-7.67	1.31	1.38
38	BE	112	G	O3'-P	-7.67	1.51	1.61
85	AA	190	A	P-O5'	-7.67	1.52	1.59
85	AA	502	A	P-O5'	-7.67	1.52	1.59
85	AA	538	A	C6-N6	-7.67	1.27	1.33
85	AA	638	G	C4'-C3'	-7.67	1.44	1.53
85	AA	2028	G	N9-C4	-7.67	1.31	1.38
35	BB	693	U	O3'-P	-7.66	1.51	1.61
35	BB	1360	A	C8-N7	-7.66	1.26	1.31
40	BG	97	G	C2'-C1'	-7.66	1.45	1.53
34	BA	937	G	C2'-C1'	-7.66	1.45	1.53
38	BE	123	A	C1'-N9	-7.66	1.36	1.46
85	AA	1217	U	C4'-C3'	-7.66	1.44	1.53
34	BA	465	A	C4'-C3'	-7.66	1.44	1.53
34	BA	846	U	O3'-P	-7.66	1.51	1.61
34	BA	1073	G	C5-C4	-7.66	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1667	G	C2'-C1'	-7.66	1.45	1.53
34	BA	1673	G	C2'-C1'	-7.66	1.45	1.53
35	BB	780	U	C2'-C1'	-7.66	1.45	1.53
35	BB	1316	U	P-O5'	-7.66	1.52	1.59
35	BB	1476	C	O3'-P	-7.66	1.51	1.61
38	BE	94	U	C2'-C1'	-7.66	1.45	1.53
39	BF	39	C	C4'-C3'	-7.66	1.44	1.53
40	BG	59	G	N9-C8	-7.66	1.32	1.37
40	BG	105	A	O3'-P	-7.66	1.51	1.61
85	AA	1155	A	P-O5'	-7.66	1.52	1.59
85	AA	1562	U	C2'-C1'	-7.66	1.45	1.53
85	AA	1690	A	N9-C4	-7.66	1.33	1.37
85	AA	2037	A	N3-C4	-7.66	1.30	1.34
34	BA	56	G	C2'-C1'	-7.66	1.45	1.53
34	BA	99	G	C2'-C1'	-7.66	1.45	1.53
34	BA	503	C	C3'-C2'	-7.66	1.44	1.52
34	BA	800	G	C4'-C3'	-7.66	1.44	1.53
34	BA	1327	G	C6-N1	-7.66	1.34	1.39
35	BB	653	G	N7-C5	-7.66	1.34	1.39
35	BB	776	U	C4'-C3'	-7.66	1.44	1.53
37	BD	93	G	C2'-C1'	-7.66	1.45	1.53
85	AA	1800	U	P-O5'	-7.66	1.52	1.59
35	BB	1106	G	P-O5'	-7.66	1.52	1.59
34	BA	374	U	C3'-C2'	-7.66	1.44	1.52
34	BA	1047	U	P-O5'	-7.66	1.52	1.59
34	BA	1527	G	C1'-N9	-7.66	1.36	1.46
34	BA	1589	U	N3-C4	-7.66	1.31	1.38
34	BA	1804	A	C5-C4	-7.66	1.33	1.38
35	BB	1433	U	C2-N3	-7.66	1.32	1.37
34	BA	759	A	O4'-C1'	-7.65	1.31	1.41
34	BA	999	G	C3'-C2'	-7.65	1.44	1.52
35	BB	1022	C	C2-N3	-7.65	1.29	1.35
35	BB	1366	C	C4'-C3'	-7.65	1.44	1.53
40	BG	67	A	C2'-C1'	-7.65	1.45	1.53
34	BA	108	A	N7-C5	-7.65	1.34	1.39
34	BA	317	U	P-O5'	-7.65	1.52	1.59
34	BA	1219	G	P-O5'	-7.65	1.52	1.59
34	BA	1402	C	P-O5'	-7.65	1.52	1.59
35	BB	371	C	O3'-P	-7.65	1.51	1.61
35	BB	985	A	P-O5'	-7.65	1.52	1.59
35	BB	1206	G	N1-C2	-7.65	1.31	1.37
35	BB	1471	A	P-O5'	-7.65	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	96	C	C2-N3	-7.65	1.29	1.35
85	AA	159	G	C3'-C2'	-7.65	1.44	1.52
85	AA	1128	G	P-O5'	-7.65	1.52	1.59
85	AA	1431	U	P-O5'	-7.65	1.52	1.59
85	AA	2204	A	C1'-N9	-7.65	1.36	1.46
6	A5	200	GLY	CA-C	-7.65	1.39	1.51
34	BA	165	C	C4'-C3'	-7.65	1.44	1.53
34	BA	1068	C	O3'-P	-7.65	1.51	1.61
35	BB	373	C	C2-N3	-7.65	1.29	1.35
35	BB	458	U	C2'-C1'	-7.65	1.45	1.53
35	BB	843	G	C1'-N9	-7.65	1.36	1.46
38	BE	114	G	C2'-C1'	-7.65	1.45	1.53
85	AA	460	U	C2-N3	-7.65	1.32	1.37
35	BB	1281	G	C5-C4	-7.65	1.32	1.38
36	BC	136	G	O3'-P	-7.65	1.51	1.61
36	BC	146	U	O3'-P	-7.65	1.51	1.61
36	BC	155	C	C2'-C1'	-7.65	1.45	1.53
85	AA	1493	A	N3-C4	-7.65	1.30	1.34
85	AA	1795	C	O3'-P	-7.65	1.51	1.61
85	AA	2222	G	P-O5'	-7.65	1.52	1.59
34	BA	64	A	N7-C5	-7.65	1.34	1.39
34	BA	661	C	C3'-C2'	-7.65	1.44	1.52
34	BA	745	A	P-O5'	-7.65	1.52	1.59
34	BA	1526	C	C2'-C1'	-7.65	1.45	1.53
34	BA	1800	G	N9-C8	-7.65	1.32	1.37
35	BB	12	G	O3'-P	-7.65	1.51	1.61
85	AA	790	A	N3-C4	-7.65	1.30	1.34
35	BB	1079	G	C2'-C1'	-7.65	1.45	1.53
41	BH	44	A	C8-N7	-7.65	1.26	1.31
34	BA	592	G	C6-N1	-7.64	1.34	1.39
34	BA	1454	G	P-O5'	-7.64	1.52	1.59
35	BB	1180	G	N3-C4	-7.64	1.30	1.35
38	BE	131	C	O3'-P	-7.64	1.51	1.61
40	BG	102	G	N9-C4	7.64	1.44	1.38
85	AA	2113	U	C2-N3	-7.64	1.32	1.37
34	BA	445	C	C3'-C2'	-7.64	1.44	1.52
34	BA	1029	C	O3'-P	-7.64	1.51	1.61
34	BA	1239	G	C3'-C2'	-7.64	1.44	1.52
34	BA	1617	U	P-O5'	-7.64	1.52	1.59
35	BB	548	A	N9-C4	-7.64	1.33	1.37
39	BF	36	G	C6-N1	-7.64	1.34	1.39
40	BG	40	G	N9-C4	-7.64	1.31	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	109	G	O3'-P	-7.64	1.51	1.61
85	AA	207	G	N7-C5	-7.64	1.34	1.39
85	AA	423	G	O3'-P	-7.64	1.51	1.61
85	AA	646	C	C2'-C1'	-7.64	1.45	1.53
85	AA	938	A	N9-C8	-7.64	1.31	1.37
85	AA	1886	U	O3'-P	-7.64	1.51	1.61
34	BA	201	A	N3-C4	-7.64	1.30	1.34
34	BA	1599	A	N7-C5	-7.64	1.34	1.39
34	BA	1802	C	N1-C6	-7.64	1.32	1.37
35	BB	649	A	N9-C4	-7.64	1.33	1.37
40	BG	79	U	C1'-N1	-7.64	1.36	1.46
85	AA	687	G	C8-N7	-7.64	1.26	1.30
85	AA	880	A	C3'-C2'	-7.64	1.44	1.52
85	AA	1161	U	O3'-P	-7.64	1.51	1.61
34	BA	724	A	N7-C5	-7.64	1.34	1.39
34	BA	776	U	P-O5'	-7.64	1.52	1.59
34	BA	1523	U	O3'-P	-7.64	1.51	1.61
35	BB	435	A	N7-C5	-7.64	1.34	1.39
35	BB	562	A	C2'-C1'	-7.64	1.45	1.53
36	BC	123	G	P-O5'	-7.64	1.52	1.59
38	BE	15	A	P-O5'	-7.64	1.52	1.59
85	AA	870	U	O3'-P	-7.64	1.51	1.61
34	BA	69	C	C3'-C2'	-7.64	1.44	1.52
34	BA	365	A	O3'-P	-7.64	1.51	1.61
34	BA	414	A	N7-C5	-7.64	1.34	1.39
34	BA	924	U	O3'-P	-7.64	1.51	1.61
34	BA	1518	A	O3'-P	-7.64	1.51	1.61
34	BA	1601	C	C3'-C2'	-7.64	1.44	1.52
35	BB	412	A	N7-C5	-7.64	1.34	1.39
35	BB	442	U	C3'-C2'	-7.64	1.44	1.52
35	BB	1165	A	C1'-N9	-7.64	1.36	1.46
35	BB	1414	A	N9-C4	-7.64	1.33	1.37
35	BB	1423	U	O3'-P	-7.64	1.51	1.61
37	BD	96	C	O3'-P	-7.64	1.51	1.61
38	BE	27	A	C3'-C2'	-7.64	1.44	1.52
38	BE	186	C	O3'-P	-7.64	1.51	1.61
40	BG	152	G	C3'-C2'	-7.64	1.44	1.52
85	AA	1259	U	C2'-C1'	-7.64	1.45	1.53
85	AA	1989	A	P-O5'	-7.64	1.52	1.59
34	BA	68	A	P-O5'	-7.63	1.52	1.59
34	BA	971	G	O3'-P	-7.63	1.51	1.61
35	BB	587	A	N9-C4	-7.63	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1029	U	C1'-N1	-7.63	1.36	1.46
35	BB	1349	U	O3'-P	-7.63	1.51	1.61
39	BF	39	C	C3'-C2'	-7.63	1.44	1.52
85	AA	1839	G	O3'-P	-7.63	1.51	1.61
34	BA	566	G	C8-N7	-7.63	1.26	1.30
34	BA	889	U	O3'-P	-7.63	1.51	1.61
34	BA	935	A	N9-C4	-7.63	1.33	1.37
34	BA	1013	A	C5-C4	-7.63	1.33	1.38
34	BA	1585	A	C4'-O4'	-7.63	1.35	1.45
35	BB	1396	G	O3'-P	-7.63	1.51	1.61
37	BD	102	C	O3'-P	-7.63	1.51	1.61
34	BA	1215	U	O3'-P	-7.63	1.51	1.61
34	BA	1454	G	C8-N7	-7.63	1.26	1.30
34	BA	1461	A	O3'-P	-7.63	1.51	1.61
34	BA	1545	C	C3'-C2'	-7.63	1.44	1.52
38	BE	191	U	C3'-C2'	-7.63	1.44	1.52
85	AA	679	A	C2'-C1'	-7.63	1.45	1.53
85	AA	973	U	C3'-O3'	7.63	1.52	1.42
85	AA	1921	G	C5-C6	7.63	1.50	1.42
35	BB	1039	A	N3-C4	-7.63	1.30	1.34
85	AA	20	G	C2'-C1'	-7.63	1.45	1.53
35	BB	1326	U	C3'-C2'	-7.63	1.44	1.52
37	BD	110	G	C1'-N9	-7.63	1.36	1.46
85	AA	419	A	C2'-C1'	-7.63	1.45	1.53
34	BA	23	A	N7-C5	-7.63	1.34	1.39
35	BB	1278	A	N9-C4	-7.63	1.33	1.37
40	BG	164	U	C2'-C1'	-7.63	1.45	1.53
85	AA	436	G	O3'-P	-7.63	1.51	1.61
85	AA	1526	G	N7-C5	-7.63	1.34	1.39
34	BA	413	A	C8-N7	-7.62	1.26	1.31
34	BA	755	G	C2'-C1'	-7.62	1.45	1.53
34	BA	1258	G	N7-C5	-7.62	1.34	1.39
34	BA	1300	G	O3'-P	-7.62	1.51	1.61
34	BA	1594	G	N7-C5	-7.62	1.34	1.39
36	BC	91	G	N9-C8	-7.62	1.32	1.37
34	BA	1454	G	C1'-N9	-7.62	1.36	1.46
35	BB	8	U	C4'-C3'	-7.62	1.44	1.53
35	BB	546	A	C6-N1	7.62	1.40	1.35
85	AA	118	C	O3'-P	-7.62	1.52	1.61
85	AA	162	A	N9-C4	-7.62	1.33	1.37
85	AA	1855	U	C2-N3	-7.62	1.32	1.37
85	AA	2065	U	P-O5'	-7.62	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1473	A	O3'-P	-7.62	1.52	1.61
34	BA	1709	A	C4'-O4'	-7.62	1.35	1.45
35	BB	615	A	O3'-P	-7.62	1.52	1.61
35	BB	830	G	C3'-C2'	-7.62	1.44	1.52
35	BB	1514	G	N7-C5	-7.62	1.34	1.39
41	BH	134	U	C2'-C1'	-7.62	1.45	1.53
85	AA	365	G	C5-C4	-7.62	1.33	1.38
85	AA	448	G	P-O5'	-7.62	1.52	1.59
85	AA	1276	A	O3'-P	-7.62	1.52	1.61
85	AA	2182	A	C5-C6	-7.62	1.34	1.41
34	BA	446	U	C2-N3	-7.62	1.32	1.37
34	BA	1606	A	C2'-C1'	-7.62	1.45	1.53
34	BA	1661	U	C5'-C4'	-7.62	1.42	1.51
85	AA	672	U	N1-C2	-7.62	1.31	1.38
85	AA	1538	C	C4'-C3'	-7.62	1.44	1.53
34	BA	222	C	O3'-P	-7.62	1.52	1.61
35	BB	617	C	C2-N3	-7.62	1.29	1.35
85	AA	758	C	O3'-P	-7.62	1.52	1.61
34	BA	1202	G	C6-N1	-7.62	1.34	1.39
35	BB	425	G	C1'-N9	-7.62	1.36	1.46
35	BB	1440	A	P-O5'	-7.62	1.52	1.59
40	BG	142	A	C8-N7	-7.62	1.26	1.31
85	AA	2098	A	C1'-N9	-7.62	1.36	1.46
34	BA	1067	G	C3'-C2'	-7.62	1.44	1.52
34	BA	1255	G	P-O5'	-7.62	1.52	1.59
35	BB	1398	A	N9-C8	-7.62	1.31	1.37
38	BE	69	C	O3'-P	-7.62	1.52	1.61
40	BG	66	C	O3'-P	-7.62	1.52	1.61
85	AA	436	G	C3'-C2'	-7.62	1.44	1.52
85	AA	1216	A	O3'-P	-7.62	1.52	1.61
35	BB	1268	C	P-O5'	-7.61	1.52	1.59
36	BC	101	U	O3'-P	-7.61	1.52	1.61
85	AA	107	A	C1'-N9	-7.61	1.36	1.46
85	AA	184	A	N9-C4	-7.61	1.33	1.37
34	BA	1555	G	C3'-C2'	-7.61	1.44	1.52
34	BA	1574	C	O3'-P	-7.61	1.52	1.61
35	BB	127	U	C2-N3	-7.61	1.32	1.37
35	BB	1368	A	C1'-N9	-7.61	1.36	1.46
37	BD	108	G	N9-C4	-7.61	1.31	1.38
34	BA	487	A	C5-C4	-7.61	1.33	1.38
85	AA	75	U	P-O5'	-7.61	1.52	1.59
85	AA	1221	G	N9-C8	-7.61	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	33	C	C2'-C1'	-7.61	1.45	1.53
34	BA	75	U	O3'-P	-7.61	1.52	1.61
34	BA	419	U	C1'-N1	-7.61	1.36	1.46
34	BA	427	G	N1-C2	-7.61	1.31	1.37
35	BB	399	A	N3-C4	-7.61	1.30	1.34
35	BB	450	A	C5-C4	-7.61	1.33	1.38
35	BB	543	G	N3-C4	-7.61	1.30	1.35
85	AA	274	A	C2'-C1'	-7.61	1.45	1.53
85	AA	713	G	C2'-C1'	-7.61	1.45	1.53
35	BB	435	A	C1'-N9	-7.61	1.36	1.46
35	BB	1251	G	C6-N1	-7.61	1.34	1.39
35	BB	1386	C	C3'-C2'	-7.61	1.44	1.52
37	BD	105	G	C2'-C1'	-7.61	1.45	1.53
85	AA	263	A	O3'-P	-7.61	1.52	1.61
85	AA	442	G	C3'-C2'	-7.61	1.44	1.52
85	AA	1451	U	O3'-P	-7.61	1.52	1.61
35	BB	401	U	C2'-C1'	-7.60	1.45	1.53
35	BB	90	G	P-O5'	-7.60	1.52	1.59
35	BB	112	G	C4'-C3'	-7.60	1.44	1.53
35	BB	528	G	P-O5'	-7.60	1.52	1.59
35	BB	562	A	O3'-P	-7.60	1.52	1.61
38	BE	176	G	C1'-N9	-7.60	1.36	1.46
41	BH	20	A	C5-C4	-7.60	1.33	1.38
41	BH	37	U	C4'-C3'	-7.60	1.44	1.53
85	AA	1573	A	P-O5'	-7.60	1.52	1.59
35	BB	43	G	O3'-P	-7.60	1.52	1.61
34	BA	799	A	C4'-C3'	7.60	1.61	1.53
34	BA	843	G	N1-C2	-7.60	1.31	1.37
34	BA	1532	G	P-O5'	-7.60	1.52	1.59
35	BB	376	A	N9-C4	-7.60	1.33	1.37
35	BB	972	C	C2'-C1'	-7.60	1.45	1.53
35	BB	1433	U	P-O5'	-7.60	1.52	1.59
35	BB	1453	G	C4'-C3'	-7.60	1.44	1.53
37	BD	55	A	C2'-C1'	-7.60	1.45	1.53
85	AA	178	U	N3-C4	-7.60	1.31	1.38
34	BA	15	G	C8-N7	-7.60	1.26	1.30
34	BA	710	A	N9-C4	-7.60	1.33	1.37
34	BA	927	A	C2'-C1'	-7.60	1.45	1.53
34	BA	1064	A	O3'-P	-7.60	1.52	1.61
34	BA	1425	G	P-O5'	-7.60	1.52	1.59
34	BA	1701	U	C4'-C3'	-7.60	1.44	1.53
34	BA	1793	G	C2-N2	-7.60	1.26	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	90	G	O3'-P	-7.60	1.52	1.61
85	AA	106	G	C2'-C1'	-7.60	1.45	1.53
85	AA	1155	A	C2'-C1'	-7.60	1.45	1.53
85	AA	1462	A	C4'-C3'	-7.60	1.44	1.53
85	AA	2052	U	O3'-P	-7.60	1.52	1.61
34	BA	303	C	O4'-C1'	-7.60	1.31	1.41
34	BA	856	G	C5-C4	-7.60	1.33	1.38
34	BA	290	G	C5-C4	-7.59	1.33	1.38
34	BA	807	U	P-O5'	-7.59	1.52	1.59
35	BB	1178	A	C4'-C3'	-7.59	1.44	1.53
85	AA	696	G	C2'-C1'	-7.59	1.45	1.53
34	BA	919	A	O3'-P	-7.59	1.52	1.61
35	BB	501	G	N7-C5	-7.59	1.34	1.39
34	BA	210	G	C3'-C2'	-7.59	1.44	1.52
34	BA	1150	A	N7-C5	-7.59	1.34	1.39
34	BA	1210	A	C1'-N9	-7.59	1.36	1.46
34	BA	1474	G	P-O5'	-7.59	1.52	1.59
40	BG	30	C	C2-N3	-7.59	1.29	1.35
85	AA	1198	U	C3'-C2'	-7.59	1.44	1.52
85	AA	2073	U	C2'-C1'	-7.59	1.45	1.53
34	BA	583	G	P-O5'	-7.59	1.52	1.59
34	BA	1834	A	O3'-P	-7.59	1.52	1.61
35	BB	1441	C	C3'-C2'	-7.59	1.44	1.52
85	AA	699	U	C2-N3	-7.59	1.32	1.37
85	AA	790	A	O3'-P	-7.59	1.52	1.61
85	AA	1135	U	C3'-C2'	-7.59	1.44	1.52
85	AA	2214	A	O3'-P	-7.59	1.52	1.61
34	BA	761	U	C2'-C1'	-7.59	1.45	1.53
34	BA	1588	U	C2-N3	-7.59	1.32	1.37
35	BB	771	U	O3'-P	-7.59	1.52	1.61
35	BB	1086	G	P-O5'	-7.59	1.52	1.59
85	AA	1196	C	C1'-N1	-7.59	1.36	1.46
35	BB	389	G	N7-C5	-7.59	1.34	1.39
35	BB	692	G	C2'-C1'	-7.59	1.45	1.53
35	BB	810	G	C5-C4	-7.59	1.33	1.38
36	BC	9	G	C5-C4	-7.59	1.33	1.38
85	AA	99	U	C4'-C3'	-7.59	1.44	1.53
85	AA	1130	G	P-O5'	-7.59	1.52	1.59
85	AA	2121	G	P-O5'	-7.59	1.52	1.59
34	BA	734	G	N9-C8	-7.58	1.32	1.37
36	BC	3	C	C3'-C2'	-7.58	1.44	1.52
38	BE	43	A	N9-C4	-7.58	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	272	C	C2'-C1'	-7.58	1.45	1.53
34	BA	418	G	O3'-P	-7.58	1.52	1.61
34	BA	480	G	N3-C4	-7.58	1.30	1.35
34	BA	947	A	C1'-N9	-7.58	1.36	1.46
34	BA	1440	C	O3'-P	-7.58	1.52	1.61
34	BA	1511	C	C1'-N1	-7.58	1.36	1.46
34	BA	1695	G	N1-C2	-7.58	1.31	1.37
35	BB	131	A	C2'-C1'	-7.58	1.45	1.53
35	BB	799	A	C8-N7	-7.58	1.26	1.31
35	BB	1096	G	C1'-N9	-7.58	1.36	1.46
40	BG	93	U	O3'-P	-7.58	1.52	1.61
34	BA	412	G	O3'-P	-7.58	1.52	1.61
34	BA	452	A	C8-N7	-7.58	1.26	1.31
34	BA	785	G	C2-N2	-7.58	1.26	1.34
35	BB	1045	G	C6-N1	-7.58	1.34	1.39
35	BB	1359	G	C5-C4	-7.58	1.33	1.38
40	BG	61	A	N9-C4	-7.58	1.33	1.37
40	BG	169	A	O3'-P	-7.58	1.52	1.61
85	AA	63	G	C3'-C2'	-7.58	1.44	1.52
85	AA	117	C	P-O5'	-7.58	1.52	1.59
85	AA	513	G	P-O5'	-7.58	1.52	1.59
85	AA	2181	G	O3'-P	-7.58	1.52	1.61
86	AB	8	U	C2'-C1'	-7.58	1.45	1.53
34	BA	498	A	C2'-C1'	-7.58	1.45	1.53
34	BA	943	G	O3'-P	-7.58	1.52	1.61
34	BA	1310	C	O3'-P	-7.58	1.52	1.61
35	BB	589	U	C2'-C1'	-7.58	1.45	1.53
35	BB	1290	C	C3'-C2'	-7.58	1.44	1.52
85	AA	623	G	O3'-P	-7.58	1.52	1.61
34	BA	289	A	C8-N7	-7.58	1.26	1.31
34	BA	706	C	P-O5'	-7.58	1.52	1.59
34	BA	1323	G	N9-C4	-7.58	1.31	1.38
34	BA	1552	C	P-O5'	-7.58	1.52	1.59
35	BB	1431	G	O3'-P	-7.58	1.52	1.61
40	BG	174	G	C4'-O4'	-7.58	1.35	1.45
85	AA	867	G	P-O5'	-7.58	1.52	1.59
85	AA	1567	C	P-O5'	-7.58	1.52	1.59
34	BA	422	C	C3'-C2'	-7.58	1.44	1.52
34	BA	1685	C	C2'-C1'	-7.58	1.45	1.53
35	BB	752	A	N9-C4	-7.58	1.33	1.37
34	BA	1119	A	C1'-N9	-7.58	1.36	1.46
35	BB	880	G	N7-C5	-7.58	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1193	G	N9-C4	-7.58	1.31	1.38
35	BB	1460	G	O3'-P	-7.58	1.52	1.61
36	BC	23	G	C3'-C2'	-7.58	1.44	1.52
38	BE	59	U	C2'-C1'	-7.58	1.45	1.53
85	AA	12	U	O3'-P	-7.58	1.52	1.61
85	AA	557	G	C2'-C1'	-7.58	1.45	1.53
85	AA	1291	A	N7-C5	-7.58	1.34	1.39
85	AA	1503	G	C2'-C1'	-7.58	1.45	1.53
34	BA	839	U	C2'-C1'	-7.57	1.45	1.53
35	BB	374	A	O3'-P	-7.57	1.52	1.61
35	BB	645	C	C2'-C1'	-7.57	1.45	1.53
37	BD	82	G	N1-C2	-7.57	1.31	1.37
85	AA	2182	A	C8-N7	-7.57	1.26	1.31
34	BA	922	C	C4'-C3'	-7.57	1.44	1.53
85	AA	1281	G	C3'-C2'	-7.57	1.44	1.52
85	AA	1440	C	O3'-P	-7.57	1.52	1.61
34	BA	827	A	N9-C4	-7.57	1.33	1.37
34	BA	1095	G	C4'-C3'	-7.57	1.44	1.53
34	BA	1840	C	C2'-C1'	-7.57	1.45	1.53
85	AA	528	U	O3'-P	-7.57	1.52	1.61
85	AA	1869	U	O3'-P	-7.57	1.52	1.61
34	BA	325	A	O3'-P	-7.57	1.52	1.61
35	BB	779	C	C1'-N1	-7.57	1.36	1.46
35	BB	1245	A	P-O5'	-7.57	1.52	1.59
85	AA	711	C	O3'-P	-7.57	1.52	1.61
34	BA	1160	U	C2'-C1'	-7.57	1.45	1.53
35	BB	134	G	C2'-C1'	-7.57	1.45	1.53
36	BC	9	G	O4'-C1'	-7.57	1.31	1.41
37	BD	45	U	O3'-P	-7.57	1.52	1.61
85	AA	1252	A	N9-C4	-7.57	1.33	1.37
85	AA	1520	A	N7-C5	-7.57	1.34	1.39
85	AA	1637	C	O3'-P	-7.57	1.52	1.61
85	AA	1907	U	O3'-P	-7.57	1.52	1.61
85	AA	2237	G	C3'-C2'	-7.57	1.44	1.52
34	BA	709	C	P-O5'	-7.57	1.52	1.59
34	BA	1331	G	N7-C5	-7.57	1.34	1.39
34	BA	1450	G	N9-C4	-7.57	1.31	1.38
34	BA	1621	U	O3'-P	-7.57	1.52	1.61
35	BB	101	U	C2'-C1'	-7.57	1.45	1.53
35	BB	375	G	C6-N1	-7.57	1.34	1.39
35	BB	1360	A	N9-C8	-7.57	1.31	1.37
41	BH	44	A	N3-C4	-7.57	1.30	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	470	C	C2-N3	-7.56	1.29	1.35
85	AA	1695	G	C2'-C1'	-7.56	1.45	1.53
34	BA	1332	U	O3'-P	-7.56	1.52	1.61
35	BB	460	C	C2-N3	-7.56	1.29	1.35
35	BB	590	G	C6-N1	-7.56	1.34	1.39
85	AA	392	G	C1'-N9	-7.56	1.36	1.46
85	AA	497	G	C3'-C2'	-7.56	1.44	1.52
85	AA	705	G	P-O5'	-7.56	1.52	1.59
34	BA	50	G	C3'-C2'	-7.56	1.44	1.52
34	BA	995	A	C2'-C1'	-7.56	1.45	1.53
38	BE	121	G	N9-C4	-7.56	1.31	1.38
40	BG	136	G	O3'-P	-7.56	1.52	1.61
85	AA	836	A	O3'-P	-7.56	1.52	1.61
85	AA	1166	C	O3'-P	-7.56	1.52	1.61
34	BA	465	A	C1'-N9	-7.56	1.36	1.46
34	BA	1657	A	N3-C4	-7.56	1.30	1.34
34	BA	1684	A	C1'-N9	-7.56	1.36	1.46
35	BB	399	A	C1'-N9	-7.56	1.36	1.46
35	BB	788	U	N3-C4	-7.56	1.31	1.38
38	BE	19	G	O3'-P	-7.56	1.52	1.61
41	BH	46	C	O3'-P	-7.56	1.52	1.61
85	AA	542	G	O3'-P	-7.56	1.52	1.61
34	BA	1297	G	O3'-P	-7.56	1.52	1.61
35	BB	1054	G	C3'-C2'	-7.56	1.44	1.52
36	BC	35	C	C2'-C1'	-7.56	1.45	1.53
38	BE	128	G	C6-N1	-7.56	1.34	1.39
85	AA	1144	G	C3'-C2'	-7.56	1.44	1.52
85	AA	1206	A	N7-C5	-7.56	1.34	1.39
34	BA	1275	G	N1-C2	-7.56	1.31	1.37
34	BA	1294	C	O4'-C1'	-7.56	1.31	1.41
85	AA	1296	G	P-O5'	-7.56	1.52	1.59
35	BB	35	G	N3-C4	-7.55	1.30	1.35
35	BB	1408	G	C2'-C1'	-7.55	1.45	1.53
36	BC	115	G	C3'-C2'	-7.55	1.44	1.52
37	BD	102	C	C3'-C2'	-7.55	1.44	1.52
40	BG	110	U	C4'-O4'	-7.55	1.35	1.45
85	AA	620	U	O3'-P	-7.55	1.52	1.61
34	BA	1843	G	O3'-P	-7.55	1.52	1.61
35	BB	462	G	N9-C4	-7.55	1.31	1.38
35	BB	495	A	C2'-C1'	-7.55	1.45	1.53
35	BB	1254	G	N3-C4	-7.55	1.30	1.35
85	AA	478	U	P-O5'	-7.55	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1149	A	C2'-C1'	-7.55	1.45	1.53
34	BA	716	C	O3'-P	-7.55	1.52	1.61
35	BB	962	U	C2'-C1'	-7.55	1.45	1.53
36	BC	6	G	C8-N7	-7.55	1.26	1.30
57	BX	82	PHE	CB-CG	7.55	1.64	1.51
85	AA	1897	A	O3'-P	-7.55	1.52	1.61
85	AA	2020	C	P-O5'	-7.55	1.52	1.59
34	BA	606	G	N9-C4	7.55	1.44	1.38
34	BA	937	G	C1'-N9	-7.55	1.36	1.46
34	BA	990	G	O3'-P	-7.55	1.52	1.61
34	BA	1398	C	O3'-P	-7.55	1.52	1.61
34	BA	1547	G	N1-C2	-7.55	1.31	1.37
35	BB	708	C	O3'-P	-7.55	1.52	1.61
35	BB	810	G	P-O5'	-7.55	1.52	1.59
35	BB	1341	U	O3'-P	-7.55	1.52	1.61
35	BB	1348	C	C3'-C2'	-7.55	1.44	1.52
85	AA	590	U	O3'-P	-7.55	1.52	1.61
85	AA	1997	G	C5'-C4'	7.55	1.60	1.51
85	AA	2124	G	P-O5'	-7.55	1.52	1.59
34	BA	47	U	C3'-C2'	-7.55	1.44	1.52
34	BA	507	U	C4'-C3'	7.55	1.61	1.53
34	BA	1034	U	N3-C4	-7.55	1.31	1.38
34	BA	1119	A	P-O5'	-7.55	1.52	1.59
35	BB	38	C	O3'-P	-7.55	1.52	1.61
39	BF	40	U	P-O5'	-7.55	1.52	1.59
40	BG	131	U	P-O5'	-7.55	1.52	1.59
85	AA	425	G	C1'-N9	-7.55	1.36	1.46
85	AA	681	G	C4'-O4'	-7.55	1.35	1.45
34	BA	333	A	O3'-P	-7.54	1.52	1.61
85	AA	437	G	N9-C4	-7.54	1.31	1.38
85	AA	475	A	C3'-C2'	-7.54	1.44	1.52
34	BA	465	A	N7-C5	-7.54	1.34	1.39
34	BA	577	U	P-O5'	-7.54	1.52	1.59
34	BA	906	A	C4'-C3'	-7.54	1.44	1.53
34	BA	1107	A	C4'-C3'	-7.54	1.44	1.53
34	BA	1111	U	P-O5'	-7.54	1.52	1.59
34	BA	1273	U	C2'-C1'	-7.54	1.45	1.53
35	BB	1133	C	C2-N3	-7.54	1.29	1.35
35	BB	1398	A	P-O5'	-7.54	1.52	1.59
40	BG	159	A	C1'-N9	-7.54	1.36	1.46
85	AA	605	A	P-O5'	-7.54	1.52	1.59
34	BA	418	G	N9-C8	-7.54	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	651	U	C2-N3	-7.54	1.32	1.37
34	BA	1354	G	O3'-P	-7.54	1.52	1.61
34	BA	1733	G	C2'-C1'	-7.54	1.45	1.53
35	BB	37	C	C3'-C2'	-7.54	1.44	1.52
35	BB	509	A	C1'-N9	-7.54	1.36	1.46
35	BB	625	A	C3'-C2'	-7.54	1.44	1.52
35	BB	1006	C	C2'-C1'	-7.54	1.45	1.53
40	BG	150	A	O3'-P	-7.54	1.52	1.61
41	BH	14	C	P-O5'	-7.54	1.52	1.59
85	AA	103	U	O3'-P	-7.54	1.52	1.61
85	AA	1921	G	C8-N7	7.54	1.35	1.30
34	BA	236	A	N9-C4	-7.54	1.33	1.37
34	BA	824	C	C2'-C1'	-7.54	1.45	1.53
35	BB	1527	A	N9-C4	-7.54	1.33	1.37
35	BB	387	G	N3-C4	-7.54	1.30	1.35
35	BB	399	A	P-O5'	-7.54	1.52	1.59
85	AA	407	G	C2'-C1'	-7.54	1.45	1.53
85	AA	651	G	P-O5'	-7.54	1.52	1.59
35	BB	1453	G	O3'-P	-7.54	1.52	1.61
85	AA	31	C	C2-N3	-7.54	1.29	1.35
85	AA	802	A	N7-C5	-7.54	1.34	1.39
34	BA	93	A	C1'-N9	-7.54	1.36	1.46
34	BA	95	C	C1'-N1	-7.54	1.36	1.46
34	BA	357	A	C3'-C2'	-7.54	1.44	1.52
34	BA	501	U	P-O5'	-7.54	1.52	1.59
34	BA	727	G	C5-C4	-7.54	1.33	1.38
34	BA	765	U	C3'-C2'	7.54	1.61	1.52
34	BA	796	G	N9-C4	-7.54	1.31	1.38
34	BA	1613	G	N7-C5	-7.54	1.34	1.39
35	BB	88	U	C2'-C1'	-7.54	1.45	1.53
35	BB	401	U	N1-C2	-7.54	1.31	1.38
35	BB	448	G	C2'-C1'	-7.54	1.45	1.53
35	BB	794	G	N7-C5	-7.54	1.34	1.39
35	BB	1302	C	C2-N3	-7.54	1.29	1.35
85	AA	1282	A	C5-C4	-7.54	1.33	1.38
34	BA	10	G	C4'-O4'	-7.53	1.35	1.45
34	BA	101	G	O3'-P	-7.53	1.52	1.61
35	BB	501	G	O3'-P	-7.53	1.52	1.61
35	BB	815	G	C2'-C1'	-7.53	1.45	1.53
35	BB	839	G	C2'-C1'	-7.53	1.45	1.53
35	BB	1014	U	C2'-C1'	-7.53	1.45	1.53
35	BB	1522	G	C2'-C1'	-7.53	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	46	G	C3'-C2'	-7.53	1.44	1.52
85	AA	260	A	P-O5'	-7.53	1.52	1.59
85	AA	428	G	N3-C4	-7.53	1.30	1.35
34	BA	736	G	C5-C6	-7.53	1.34	1.42
35	BB	1001	G	C2'-C1'	-7.53	1.45	1.53
85	AA	1485	G	C2-N3	-7.53	1.26	1.32
34	BA	125	G	C2-N2	-7.53	1.27	1.34
34	BA	629	G	P-O5'	-7.53	1.52	1.59
34	BA	788	C	C1'-N1	-7.53	1.36	1.46
34	BA	857	C	P-O5'	-7.53	1.52	1.59
34	BA	917	C	O3'-P	-7.53	1.52	1.61
35	BB	84	G	O3'-P	-7.53	1.52	1.61
35	BB	622	G	C5-C6	-7.53	1.34	1.42
35	BB	672	C	C2'-C1'	-7.53	1.45	1.53
36	BC	126	G	C6-N1	-7.53	1.34	1.39
40	BG	142	A	O4'-C1'	-7.53	1.31	1.41
85	AA	1537	A	N7-C5	-7.53	1.34	1.39
34	BA	505	U	C2'-C1'	-7.53	1.45	1.53
34	BA	1723	U	C5'-C4'	7.53	1.60	1.51
38	BE	68	U	C3'-C2'	-7.53	1.44	1.52
85	AA	491	G	C4'-C3'	-7.53	1.44	1.53
85	AA	1251	G	P-O5'	-7.53	1.52	1.59
34	BA	23	A	P-O5'	-7.53	1.52	1.59
34	BA	295	G	P-O5'	-7.53	1.52	1.59
34	BA	1092	U	O3'-P	-7.53	1.52	1.61
35	BB	45	A	O3'-P	-7.53	1.52	1.61
35	BB	425	G	N9-C8	-7.53	1.32	1.37
85	AA	486	G	C5-C4	-7.53	1.33	1.38
85	AA	835	C	C3'-C2'	-7.53	1.44	1.52
86	AB	5	G	P-O5'	-7.53	1.52	1.59
34	BA	52	G	O3'-P	-7.53	1.52	1.61
34	BA	117	C	C4'-C3'	-7.53	1.44	1.53
34	BA	1095	G	C2-N2	-7.53	1.27	1.34
34	BA	1435	A	N9-C4	-7.53	1.33	1.37
35	BB	776	U	C2'-C1'	-7.53	1.45	1.53
35	BB	1199	A	C5-C4	-7.53	1.33	1.38
35	BB	1471	A	O3'-P	-7.53	1.52	1.61
37	BD	65	G	O3'-P	-7.53	1.52	1.61
38	BE	197	A	C2'-C1'	-7.53	1.45	1.53
85	AA	36	U	C2-N3	-7.53	1.32	1.37
85	AA	597	A	O3'-P	-7.53	1.52	1.61
85	AA	678	A	C2'-C1'	-7.53	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2107	C	O3'-P	-7.53	1.52	1.61
34	BA	849	G	N9-C4	-7.52	1.31	1.38
35	BB	619	A	O3'-P	-7.52	1.52	1.61
40	BG	111	C	C2'-C1'	-7.52	1.45	1.53
34	BA	1666	U	C2'-C1'	-7.52	1.45	1.53
35	BB	96	A	C5-C4	-7.52	1.33	1.38
52	BS	127	GLY	CA-C	-7.52	1.39	1.51
85	AA	157	G	C2'-C1'	-7.52	1.45	1.53
85	AA	2032	G	C2'-C1'	-7.52	1.45	1.53
34	BA	958	G	C1'-N9	-7.52	1.36	1.46
34	BA	1299	G	C5-C4	-7.52	1.33	1.38
35	BB	1395	G	C6-N1	-7.52	1.34	1.39
85	AA	1151	G	N7-C5	-7.52	1.34	1.39
34	BA	508	C	O3'-P	-7.52	1.52	1.61
34	BA	1284	G	O3'-P	-7.52	1.52	1.61
35	BB	136	A	C3'-C2'	-7.52	1.44	1.52
35	BB	415	A	N7-C5	-7.52	1.34	1.39
35	BB	584	A	N9-C8	-7.52	1.31	1.37
35	BB	1246	C	C2'-C1'	-7.52	1.45	1.53
85	AA	414	C	C2-N3	-7.52	1.29	1.35
85	AA	801	U	P-O5'	-7.52	1.52	1.59
34	BA	1162	U	C3'-C2'	-7.52	1.44	1.52
35	BB	1208	G	C6-N1	-7.52	1.34	1.39
35	BB	1221	G	P-O5'	-7.52	1.52	1.59
85	AA	739	C	C2'-C1'	-7.52	1.45	1.53
85	AA	1229	G	N9-C4	7.52	1.44	1.38
34	BA	202	A	P-O5'	-7.51	1.52	1.59
34	BA	257	G	O3'-P	-7.51	1.52	1.61
34	BA	398	G	C2'-C1'	-7.51	1.45	1.53
35	BB	608	A	C1'-N9	-7.51	1.36	1.46
35	BB	653	G	C2'-C1'	-7.51	1.45	1.53
35	BB	1165	A	C5-C4	-7.51	1.33	1.38
85	AA	584	G	C6-N1	-7.51	1.34	1.39
35	BB	1262	A	C1'-N9	-7.51	1.36	1.46
35	BB	1290	C	C2'-C1'	-7.51	1.45	1.53
35	BB	1521	G	O3'-P	-7.51	1.52	1.61
85	AA	978	U	C2'-C1'	-7.51	1.45	1.53
34	BA	222	C	C3'-C2'	-7.51	1.44	1.52
34	BA	354	G	C2'-C1'	-7.51	1.45	1.53
34	BA	607	C	O3'-P	-7.51	1.52	1.61
34	BA	719	G	C2'-C1'	-7.51	1.45	1.53
34	BA	1421	A	C2'-C1'	-7.51	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1721	U	O3'-P	-7.51	1.52	1.61
40	BG	128	U	O3'-P	-7.51	1.52	1.61
41	BH	105	U	O3'-P	-7.51	1.52	1.61
85	AA	259	A	C5-C4	-7.51	1.33	1.38
34	BA	888	G	C2-N2	-7.51	1.27	1.34
34	BA	1249	G	C5-C4	-7.51	1.33	1.38
35	BB	96	A	N9-C8	-7.51	1.31	1.37
35	BB	696	G	P-O5'	-7.51	1.52	1.59
35	BB	1019	C	C2-N3	-7.51	1.29	1.35
40	BG	90	G	P-O5'	-7.51	1.52	1.59
85	AA	381	A	C6-N1	-7.51	1.30	1.35
85	AA	514	U	C2-N3	-7.51	1.32	1.37
85	AA	976	G	C6-N1	-7.51	1.34	1.39
85	AA	1491	G	C2'-C1'	-7.51	1.45	1.53
85	AA	1657	C	C2'-C1'	-7.51	1.45	1.53
34	BA	1505	G	C3'-C2'	-7.51	1.44	1.52
38	BE	192	A	C4'-C3'	-7.51	1.44	1.53
85	AA	1287	C	C4'-C3'	-7.51	1.44	1.53
85	AA	1904	C	C2-N3	-7.51	1.29	1.35
34	BA	754	G	C1'-N9	-7.51	1.36	1.46
34	BA	932	G	P-O5'	-7.51	1.52	1.59
35	BB	386	G	C2-N3	-7.51	1.26	1.32
35	BB	1426	G	P-O5'	-7.51	1.52	1.59
85	AA	2215	C	C3'-C2'	-7.51	1.44	1.52
34	BA	878	G	C1'-N9	-7.50	1.36	1.46
34	BA	1003	A	C2'-C1'	-7.50	1.45	1.53
34	BA	1380	G	P-O5'	-7.50	1.52	1.59
35	BB	585	U	C2-N3	-7.50	1.32	1.37
35	BB	1363	A	P-O5'	-7.50	1.52	1.59
37	BD	98	G	C6-N1	-7.50	1.34	1.39
38	BE	12	A	C1'-N9	-7.50	1.36	1.46
85	AA	504	U	P-O5'	-7.50	1.52	1.59
34	BA	212	A	C1'-N9	-7.50	1.36	1.46
34	BA	421	G	O3'-P	-7.50	1.52	1.61
34	BA	1039	G	O3'-P	-7.50	1.52	1.61
34	BA	1818	A	N9-C4	-7.50	1.33	1.37
35	BB	1205	A	N9-C4	-7.50	1.33	1.37
40	BG	57	A	P-O5'	-7.50	1.52	1.59
41	BH	113	G	C3'-C2'	-7.50	1.44	1.52
85	AA	130	G	C2'-C1'	-7.50	1.45	1.53
85	AA	2173	A	N9-C4	-7.50	1.33	1.37
34	BA	742	C	C4'-C3'	-7.50	1.44	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1441	C	C3'-C2'	-7.50	1.44	1.52
35	BB	1415	G	C2-N2	-7.50	1.27	1.34
39	BF	41	U	C2-N3	-7.50	1.32	1.37
40	BG	15	G	C1'-N9	-7.50	1.36	1.46
85	AA	41	G	O3'-P	-7.50	1.52	1.61
85	AA	73	A	O3'-P	-7.50	1.52	1.61
85	AA	521	A	N9-C8	-7.50	1.31	1.37
85	AA	2074	G	C1'-N9	-7.50	1.36	1.46
34	BA	75	U	C2'-C1'	-7.50	1.45	1.53
35	BB	1424	G	P-O5'	-7.50	1.52	1.59
85	AA	1701	G	N3-C4	-7.50	1.30	1.35
34	BA	229	C	P-O5'	-7.50	1.52	1.59
34	BA	932	G	N9-C4	-7.50	1.31	1.38
36	BC	66	G	O3'-P	-7.50	1.52	1.61
40	BG	31	G	N9-C4	-7.50	1.31	1.38
34	BA	395	G	N9-C8	-7.50	1.32	1.37
34	BA	564	C	N3-C4	-7.50	1.28	1.33
34	BA	993	C	C2'-C1'	-7.50	1.45	1.53
35	BB	57	G	P-O5'	-7.50	1.52	1.59
35	BB	381	C	C2'-C1'	-7.50	1.45	1.53
35	BB	387	G	P-O5'	-7.50	1.52	1.59
35	BB	678	U	P-O5'	-7.50	1.52	1.59
35	BB	1530	U	C2-N3	-7.50	1.32	1.37
37	BD	98	G	P-O5'	-7.50	1.52	1.59
41	BH	69	C	C2'-C1'	-7.50	1.45	1.53
85	AA	1701	G	C2-N2	-7.50	1.27	1.34
85	AA	1818	C	C2'-C1'	-7.50	1.45	1.53
34	BA	703	U	C2-N3	-7.50	1.32	1.37
34	BA	150	C	C2'-C1'	-7.49	1.45	1.53
34	BA	492	G	C6-N1	-7.49	1.34	1.39
34	BA	1574	C	C2'-C1'	-7.49	1.45	1.53
35	BB	618	U	P-O5'	-7.49	1.52	1.59
35	BB	736	G	O3'-P	-7.49	1.52	1.61
35	BB	1294	C	N1-C6	-7.49	1.32	1.37
37	BD	70	C	P-O5'	-7.49	1.52	1.59
85	AA	481	A	C2'-C1'	-7.49	1.45	1.53
85	AA	902	A	O3'-P	-7.49	1.52	1.61
34	BA	1732	A	O3'-P	-7.49	1.52	1.61
35	BB	654	C	C3'-C2'	-7.49	1.44	1.52
35	BB	1404	A	C2'-C1'	-7.49	1.45	1.53
34	BA	51	C	O3'-P	-7.49	1.52	1.61
34	BA	207	A	C1'-N9	-7.49	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	574	G	C2'-C1'	-7.49	1.45	1.53
36	BC	102	G	C3'-C2'	-7.49	1.44	1.52
38	BE	2	G	C3'-C2'	-7.49	1.44	1.52
38	BE	28	C	C2'-C1'	-7.49	1.45	1.53
85	AA	649	C	C5'-C4'	-7.49	1.42	1.51
34	BA	348	U	C3'-C2'	-7.49	1.44	1.52
34	BA	949	C	C3'-C2'	-7.49	1.44	1.52
34	BA	998	U	C2-N3	-7.49	1.32	1.37
35	BB	713	U	P-O5'	-7.49	1.52	1.59
35	BB	1135	U	C2'-C1'	-7.49	1.45	1.53
36	BC	150	U	P-O5'	-7.49	1.52	1.59
85	AA	1201	A	C1'-N9	-7.49	1.36	1.46
85	AA	1213	U	P-O5'	-7.49	1.52	1.59
34	BA	147	U	C2-N3	-7.49	1.32	1.37
35	BB	96	A	C8-N7	-7.49	1.26	1.31
35	BB	818	U	C3'-C2'	-7.49	1.44	1.52
35	BB	1521	G	C2'-C1'	-7.49	1.45	1.53
85	AA	1925	A	C1'-N9	-7.49	1.36	1.46
34	BA	193	C	C2'-C1'	-7.49	1.45	1.53
34	BA	581	U	P-O5'	-7.49	1.52	1.59
34	BA	887	U	C2-N3	-7.49	1.32	1.37
34	BA	1553	G	P-O5'	-7.49	1.52	1.59
34	BA	1730	A	C4'-C3'	7.49	1.61	1.53
34	BA	1829	A	C2'-C1'	-7.49	1.45	1.53
35	BB	817	C	C2'-C1'	-7.49	1.45	1.53
35	BB	1359	G	N7-C5	-7.49	1.34	1.39
85	AA	178	U	C2'-C1'	-7.49	1.45	1.53
85	AA	1127	G	C4'-C3'	-7.49	1.45	1.53
85	AA	1312	G	P-O5'	-7.49	1.52	1.59
34	BA	450	G	O3'-P	-7.48	1.52	1.61
34	BA	1184	A	O3'-P	-7.48	1.52	1.61
34	BA	1258	G	N9-C4	-7.48	1.31	1.38
35	BB	1417	C	C3'-C2'	-7.48	1.44	1.52
85	AA	692	U	O3'-P	-7.48	1.52	1.61
85	AA	716	G	O3'-P	-7.48	1.52	1.61
85	AA	1519	A	N7-C5	-7.48	1.34	1.39
34	BA	56	G	P-O5'	-7.48	1.52	1.59
34	BA	812	A	C3'-C2'	-7.48	1.44	1.52
34	BA	929	A	C2'-C1'	-7.48	1.45	1.53
34	BA	1085	G	N9-C4	7.48	1.44	1.38
34	BA	1194	G	C5-C4	-7.48	1.33	1.38
34	BA	1496	G	C2'-C1'	-7.48	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1383	C	C3'-C2'	-7.48	1.44	1.52
37	BD	80	G	C5-C4	-7.48	1.33	1.38
38	BE	132	U	N3-C4	-7.48	1.31	1.38
38	BE	207	G	C6-N1	7.48	1.44	1.39
40	BG	57	A	C1'-N9	-7.48	1.36	1.46
40	BG	116	G	C1'-N9	-7.48	1.36	1.46
85	AA	475	A	O3'-P	-7.48	1.52	1.61
85	AA	1286	C	C4-N4	-7.48	1.27	1.33
85	AA	1882	U	O3'-P	-7.48	1.52	1.61
86	AB	71	G	O3'-P	-7.48	1.52	1.61
34	BA	431	A	C2'-C1'	-7.48	1.45	1.53
34	BA	1532	G	C2'-C1'	-7.48	1.45	1.53
35	BB	692	G	C1'-N9	-7.48	1.36	1.46
35	BB	792	G	P-O5'	-7.48	1.52	1.59
35	BB	1226	G	C2'-C1'	-7.48	1.45	1.53
85	AA	1656	C	C2'-C1'	-7.48	1.45	1.53
85	AA	1668	G	P-O5'	-7.48	1.52	1.59
34	BA	1553	G	N9-C4	-7.48	1.31	1.38
35	BB	1376	G	C4'-C3'	-7.48	1.45	1.53
35	BB	1397	G	C6-N1	-7.48	1.34	1.39
85	AA	2113	U	C2'-C1'	-7.48	1.45	1.53
34	BA	72	U	P-O5'	-7.48	1.52	1.59
34	BA	729	C	P-O5'	-7.48	1.52	1.59
34	BA	1505	G	O3'-P	-7.48	1.52	1.61
34	BA	1702	G	C4'-C3'	-7.48	1.45	1.53
35	BB	1193	G	O3'-P	-7.48	1.52	1.61
85	AA	1470	A	N7-C5	-7.48	1.34	1.39
34	BA	971	G	N9-C8	-7.48	1.32	1.37
34	BA	1648	G	P-O5'	-7.48	1.52	1.59
39	BF	44	C	C2-N3	-7.48	1.29	1.35
85	AA	397	G	C4'-O4'	-7.48	1.35	1.45
34	BA	54	A	C3'-C2'	-7.47	1.44	1.52
34	BA	755	G	C3'-C2'	-7.47	1.44	1.52
35	BB	77	A	O3'-P	-7.47	1.52	1.61
35	BB	386	G	N3-C4	-7.47	1.30	1.35
35	BB	505	G	O3'-P	-7.47	1.52	1.61
35	BB	1488	G	N1-C2	-7.47	1.31	1.37
36	BC	68	A	C1'-N9	-7.47	1.36	1.46
85	AA	245	A	O3'-P	-7.47	1.52	1.61
34	BA	821	G	N3-C4	-7.47	1.30	1.35
34	BA	992	A	C8-N7	-7.47	1.26	1.31
34	BA	1487	U	C2-N3	-7.47	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	148	C	O3'-P	-7.47	1.52	1.61
85	AA	1845	G	C2'-C1'	-7.47	1.45	1.53
85	AA	1904	C	O3'-P	-7.47	1.52	1.61
34	BA	1045	C	O3'-P	-7.47	1.52	1.61
36	BC	43	A	N9-C4	-7.47	1.33	1.37
38	BE	182	U	O3'-P	-7.47	1.52	1.61
40	BG	14	G	N9-C8	-7.47	1.32	1.37
85	AA	480	U	C4'-C3'	-7.47	1.45	1.53
85	AA	1257	A	C2'-C1'	-7.47	1.45	1.53
85	AA	2192	A	C2'-C1'	-7.47	1.45	1.53
35	BB	552	C	C1'-N1	-7.47	1.36	1.46
35	BB	638	G	C1'-N9	-7.47	1.36	1.46
35	BB	1137	G	P-O5'	-7.47	1.52	1.59
40	BG	97	G	N7-C5	-7.47	1.34	1.39
34	BA	780	U	O4'-C1'	7.47	1.51	1.41
35	BB	429	C	N1-C6	-7.47	1.32	1.37
35	BB	1470	G	N3-C4	-7.47	1.30	1.35
35	BB	1531	G	O3'-P	-7.47	1.52	1.61
36	BC	147	G	C8-N7	-7.47	1.26	1.30
37	BD	45	U	N3-C4	-7.47	1.31	1.38
40	BG	128	U	C3'-C2'	-7.47	1.44	1.52
85	AA	396	U	N3-C4	-7.47	1.31	1.38
85	AA	1275	A	N9-C4	-7.47	1.33	1.37
34	BA	198	U	C2'-C1'	-7.46	1.45	1.53
34	BA	583	G	N1-C2	-7.46	1.31	1.37
34	BA	1070	G	C8-N7	-7.46	1.26	1.30
34	BA	1634	A	N7-C5	-7.46	1.34	1.39
34	BA	1684	A	N9-C4	-7.46	1.33	1.37
35	BB	1044	U	O3'-P	-7.46	1.52	1.61
35	BB	1365	G	P-O5'	-7.46	1.52	1.59
35	BB	1530	U	O3'-P	-7.46	1.52	1.61
85	AA	887	A	N3-C4	-7.46	1.30	1.34
38	BE	21	C	C1'-N1	-7.46	1.36	1.46
39	BF	36	G	N1-C2	-7.46	1.31	1.37
85	AA	1162	A	N9-C4	-7.46	1.33	1.37
85	AA	2130	G	C3'-C2'	-7.46	1.44	1.52
34	BA	1091	U	P-O5'	-7.46	1.52	1.59
35	BB	679	G	N9-C8	-7.46	1.32	1.37
35	BB	1203	C	C4'-C3'	-7.46	1.45	1.53
35	BB	1252	G	C5-C6	-7.46	1.34	1.42
35	BB	1448	U	O3'-P	-7.46	1.52	1.61
40	BG	40	G	P-O5'	-7.46	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1348	C	P-O5'	-7.46	1.52	1.59
85	AA	1489	G	C3'-C2'	-7.46	1.44	1.52
85	AA	2177	C	C3'-C2'	-7.46	1.44	1.52
34	BA	1656	A	C6-N6	-7.46	1.27	1.33
35	BB	1093	C	P-O5'	-7.46	1.52	1.59
36	BC	87	C	P-O5'	-7.46	1.52	1.59
38	BE	129	G	C2-N2	-7.46	1.27	1.34
85	AA	547	A	C3'-C2'	-7.46	1.44	1.52
34	BA	117	C	C2-N3	-7.46	1.29	1.35
34	BA	606	G	N1-C2	-7.46	1.31	1.37
34	BA	1327	G	N7-C5	-7.46	1.34	1.39
34	BA	1828	A	N9-C4	-7.46	1.33	1.37
35	BB	24	C	C2'-C1'	-7.46	1.45	1.53
35	BB	68	G	O3'-P	-7.46	1.52	1.61
85	AA	1299	A	C2'-C1'	-7.46	1.45	1.53
34	BA	1178	U	O3'-P	-7.46	1.52	1.61
34	BA	1303	U	O3'-P	-7.46	1.52	1.61
35	BB	432	C	C2-N3	-7.46	1.29	1.35
35	BB	1327	U	C4'-C3'	-7.46	1.45	1.53
36	BC	148	C	C1'-N1	-7.46	1.36	1.46
85	AA	54	C	C3'-C2'	-7.46	1.44	1.52
85	AA	144	A	O3'-P	-7.46	1.52	1.61
85	AA	414	C	N3-C4	-7.46	1.28	1.33
85	AA	709	A	C2'-C1'	-7.46	1.45	1.53
34	BA	1000	G	C2'-C1'	-7.46	1.45	1.53
34	BA	1177	C	C1'-N1	-7.46	1.36	1.46
85	AA	407	G	N7-C5	-7.46	1.34	1.39
85	AA	898	A	C2'-C1'	-7.46	1.45	1.53
85	AA	1721	A	N7-C5	-7.46	1.34	1.39
34	BA	433	G	C1'-N9	-7.45	1.36	1.46
34	BA	749	G	C1'-N9	-7.45	1.36	1.46
34	BA	912	G	N7-C5	-7.45	1.34	1.39
35	BB	68	G	O4'-C1'	-7.45	1.31	1.41
35	BB	422	U	C2'-C1'	-7.45	1.45	1.53
35	BB	800	U	C5'-C4'	7.45	1.60	1.51
40	BG	139	U	C1'-N1	-7.45	1.36	1.46
85	AA	1358	A	O3'-P	-7.45	1.52	1.61
34	BA	1329	U	C2'-C1'	-7.45	1.45	1.53
85	AA	1355	U	C5'-C4'	7.45	1.60	1.51
85	AA	1517	G	O3'-P	-7.45	1.52	1.61
34	BA	119	G	C5'-C4'	7.45	1.60	1.51
34	BA	148	G	C2-N2	-7.45	1.27	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1109	G	C2'-C1'	-7.45	1.45	1.53
35	BB	1245	A	O3'-P	-7.45	1.52	1.61
35	BB	1360	A	N7-C5	-7.45	1.34	1.39
41	BH	47	G	P-O5'	-7.45	1.52	1.59
85	AA	612	A	N9-C4	-7.45	1.33	1.37
85	AA	1171	C	O3'-P	-7.45	1.52	1.61
85	AA	1444	U	P-O5'	-7.45	1.52	1.59
85	AA	2229	G	P-O5'	-7.45	1.52	1.59
34	BA	113	G	N7-C5	-7.45	1.34	1.39
34	BA	655	U	C2'-C1'	-7.45	1.45	1.53
34	BA	969	A	N7-C5	-7.45	1.34	1.39
34	BA	1172	C	C2'-C1'	-7.45	1.45	1.53
35	BB	14	C	C2'-C1'	-7.45	1.45	1.53
35	BB	768	A	C4'-O4'	7.45	1.55	1.45
35	BB	1093	C	C3'-C2'	-7.45	1.44	1.52
35	BB	1311	G	C1'-N9	-7.45	1.36	1.46
35	BB	1409	G	O3'-P	-7.45	1.52	1.61
36	BC	140	U	C3'-C2'	-7.45	1.44	1.52
41	BH	39	G	C6-N1	-7.45	1.34	1.39
85	AA	365	G	N9-C8	-7.45	1.32	1.37
85	AA	2098	A	O3'-P	-7.45	1.52	1.61
34	BA	698	U	C2-N3	-7.45	1.32	1.37
35	BB	713	U	O3'-P	-7.45	1.52	1.61
38	BE	147	G	P-O5'	-7.45	1.52	1.59
34	BA	413	A	O3'-P	-7.45	1.52	1.61
34	BA	569	C	O3'-P	-7.45	1.52	1.61
34	BA	888	G	C6-N1	-7.45	1.34	1.39
34	BA	1272	U	P-O5'	-7.45	1.52	1.59
34	BA	1311	G	C1'-N9	-7.45	1.36	1.46
34	BA	1544	G	C3'-C2'	-7.45	1.44	1.52
34	BA	1713	U	C2'-C1'	-7.45	1.45	1.53
36	BC	115	G	P-O5'	-7.45	1.52	1.59
38	BE	150	G	C1'-N9	-7.45	1.36	1.46
39	BF	47	C	C2'-C1'	-7.45	1.45	1.53
85	AA	475	A	P-O5'	-7.45	1.52	1.59
85	AA	635	G	C3'-C2'	-7.45	1.44	1.52
85	AA	1973	G	C3'-C2'	-7.45	1.44	1.52
86	AB	57	G	N9-C4	-7.45	1.31	1.38
34	BA	92	G	C3'-C2'	-7.44	1.44	1.52
40	BG	16	G	N9-C8	-7.44	1.32	1.37
85	AA	1244	A	C2'-C1'	-7.44	1.45	1.53
85	AA	2216	A	O3'-P	-7.44	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	434	A	C4'-C3'	-7.44	1.45	1.53
35	BB	816	U	C3'-C2'	-7.44	1.44	1.52
35	BB	827	U	C2'-C1'	-7.44	1.45	1.53
35	BB	1036	G	C4'-C3'	-7.44	1.45	1.53
37	BD	22	A	C2'-C1'	-7.44	1.45	1.53
85	AA	1215	A	O3'-P	-7.44	1.52	1.61
85	AA	1465	C	C2'-C1'	-7.44	1.45	1.53
85	AA	2243	G	C5'-C4'	7.44	1.60	1.51
34	BA	289	A	C5-C4	-7.44	1.33	1.38
34	BA	979	G	C2'-C1'	-7.44	1.45	1.53
34	BA	1009	G	C3'-C2'	-7.44	1.44	1.52
35	BB	503	G	C1'-N9	-7.44	1.36	1.46
35	BB	669	A	N3-C4	-7.44	1.30	1.34
35	BB	845	C	O3'-P	-7.44	1.52	1.61
85	AA	1114	A	C2'-C1'	-7.44	1.45	1.53
85	AA	1141	U	O3'-P	-7.44	1.52	1.61
85	AA	1928	A	C1'-N9	-7.44	1.36	1.46
85	AA	2204	A	O3'-P	-7.44	1.52	1.61
34	BA	19	G	N9-C8	-7.44	1.32	1.37
34	BA	73	G	C2-N3	-7.44	1.26	1.32
34	BA	377	G	C6-N1	-7.44	1.34	1.39
34	BA	1575	U	C2-N3	-7.44	1.32	1.37
34	BA	688	G	O3'-P	-7.44	1.52	1.61
34	BA	939	C	N1-C6	-7.44	1.32	1.37
34	BA	1809	G	N9-C8	-7.44	1.32	1.37
35	BB	843	G	P-O5'	-7.44	1.52	1.59
40	BG	167	C	C3'-C2'	-7.44	1.44	1.52
85	AA	1249	U	P-O5'	-7.44	1.52	1.59
85	AA	2201	A	C1'-N9	-7.44	1.36	1.46
34	BA	61	G	N9-C4	-7.44	1.32	1.38
34	BA	744	G	C3'-C2'	-7.44	1.44	1.52
85	AA	811	A	O3'-P	-7.44	1.52	1.61
85	AA	2143	U	C2'-C1'	-7.44	1.45	1.53
34	BA	337	C	O3'-P	-7.43	1.52	1.61
35	BB	130	G	P-O5'	-7.43	1.52	1.59
35	BB	1222	A	P-O5'	-7.43	1.52	1.59
85	AA	461	G	C5-C6	-7.43	1.34	1.42
85	AA	688	C	C2'-C1'	-7.43	1.45	1.53
85	AA	1200	A	N9-C4	-7.43	1.33	1.37
34	BA	52	G	C3'-C2'	-7.43	1.44	1.52
34	BA	99	G	C2-N2	-7.43	1.27	1.34
34	BA	860	G	C1'-N9	-7.43	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1395	G	C1'-N9	-7.43	1.36	1.46
35	BB	1432	U	P-O5'	-7.43	1.52	1.59
38	BE	170	U	P-O5'	-7.43	1.52	1.59
38	BE	191	U	O3'-P	-7.43	1.52	1.61
34	BA	450	G	N9-C4	-7.43	1.32	1.38
35	BB	1048	A	C2'-C1'	-7.43	1.45	1.53
41	BH	114	G	C1'-N9	-7.43	1.36	1.46
85	AA	298	C	C2'-C1'	-7.43	1.45	1.53
85	AA	792	A	N9-C4	-7.43	1.33	1.37
85	AA	1260	G	P-O5'	-7.43	1.52	1.59
34	BA	200	C	P-O5'	-7.43	1.52	1.59
34	BA	823	G	O3'-P	-7.43	1.52	1.61
34	BA	841	G	P-O5'	-7.43	1.52	1.59
35	BB	654	C	C4'-C3'	-7.43	1.45	1.53
35	BB	825	U	C4'-C3'	-7.43	1.45	1.53
38	BE	96	G	N3-C4	-7.43	1.30	1.35
40	BG	95	U	C3'-C2'	-7.43	1.44	1.52
85	AA	139	G	C2'-C1'	-7.43	1.45	1.53
34	BA	1233	U	C3'-C2'	-7.43	1.44	1.52
35	BB	1137	G	C6-N1	-7.43	1.34	1.39
35	BB	1187	G	C2-N2	-7.43	1.27	1.34
85	AA	131	C	C2'-C1'	-7.43	1.45	1.53
85	AA	1258	U	C2-N3	-7.43	1.32	1.37
34	BA	1710	C	C3'-C2'	-7.43	1.44	1.52
36	BC	110	A	C2'-C1'	-7.43	1.45	1.53
38	BE	52	U	O3'-P	-7.43	1.52	1.61
40	BG	41	U	C3'-C2'	-7.43	1.44	1.52
85	AA	1112	G	N9-C4	-7.43	1.32	1.38
85	AA	1487	G	O3'-P	-7.43	1.52	1.61
85	AA	1505	G	C2-N2	-7.43	1.27	1.34
85	AA	2046	G	P-O5'	-7.43	1.52	1.59
34	BA	1648	G	C8-N7	-7.42	1.26	1.30
36	BC	66	G	C3'-C2'	-7.42	1.44	1.52
40	BG	36	G	O3'-P	-7.42	1.52	1.61
85	AA	1497	U	C2-N3	-7.42	1.32	1.37
85	AA	1636	C	P-O5'	-7.42	1.52	1.59
35	BB	381	C	O3'-P	-7.42	1.52	1.61
35	BB	778	A	C2'-C1'	-7.42	1.45	1.53
34	BA	921	G	O3'-P	-7.42	1.52	1.61
34	BA	969	A	C2'-C1'	-7.42	1.45	1.53
34	BA	1264	U	C4-C5	-7.42	1.36	1.43
35	BB	659	C	C1'-N1	-7.42	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1241	U	O3'-P	-7.42	1.52	1.61
41	BH	38	G	C3'-C2'	-7.42	1.44	1.52
85	AA	129	U	C2'-C1'	-7.42	1.45	1.53
85	AA	2130	G	C5-C4	-7.42	1.33	1.38
35	BB	1336	G	N9-C4	-7.42	1.32	1.38
35	BB	1431	G	N9-C4	-7.42	1.32	1.38
35	BB	1445	A	C5-C4	-7.42	1.33	1.38
85	AA	386	G	C6-N1	-7.42	1.34	1.39
85	AA	1657	C	C3'-C2'	-7.42	1.44	1.52
34	BA	97	A	O3'-P	-7.42	1.52	1.61
35	BB	1288	G	O3'-P	-7.42	1.52	1.61
40	BG	71	C	C2-N3	-7.42	1.29	1.35
85	AA	1003	G	P-O5'	-7.42	1.52	1.59
34	BA	49	A	O4'-C1'	-7.42	1.32	1.41
34	BA	892	C	C2-N3	-7.42	1.29	1.35
34	BA	1520	A	C2'-C1'	-7.42	1.45	1.53
85	AA	381	A	C4'-O4'	-7.42	1.35	1.45
85	AA	879	G	C6-N1	-7.42	1.34	1.39
85	AA	1162	A	P-O5'	-7.42	1.52	1.59
34	BA	142	A	C2'-C1'	-7.42	1.45	1.53
35	BB	1307	C	C2'-C1'	-7.42	1.45	1.53
85	AA	460	U	O3'-P	-7.42	1.52	1.61
85	AA	498	C	C2-N3	-7.42	1.29	1.35
85	AA	1483	A	C3'-C2'	-7.42	1.44	1.52
34	BA	733	G	C5-C4	-7.41	1.33	1.38
34	BA	1269	C	C3'-C2'	-7.41	1.44	1.52
34	BA	1311	G	C2'-C1'	-7.41	1.45	1.53
34	BA	1406	U	C2-N3	-7.41	1.32	1.37
34	BA	1673	G	C4'-C3'	-7.41	1.45	1.53
35	BB	69	A	P-O5'	-7.41	1.52	1.59
35	BB	522	A	O3'-P	-7.41	1.52	1.61
35	BB	610	U	N3-C4	-7.41	1.31	1.38
35	BB	992	C	C4-C5	-7.41	1.37	1.43
36	BC	166	G	N9-C4	-7.41	1.32	1.38
85	AA	82	A	C4'-C3'	-7.41	1.45	1.53
85	AA	296	A	P-O5'	-7.41	1.52	1.59
85	AA	1480	C	C4'-C3'	-7.41	1.45	1.53
35	BB	618	U	C2-N3	-7.41	1.32	1.37
85	AA	99	U	P-O5'	-7.41	1.52	1.59
85	AA	661	C	C2-N3	-7.41	1.29	1.35
85	AA	1214	C	C4'-O4'	-7.41	1.35	1.45
34	BA	911	G	P-O5'	-7.41	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1121	A	C4'-C3'	-7.41	1.45	1.53
35	BB	1124	G	P-O5'	-7.41	1.52	1.59
35	BB	1422	G	N3-C4	-7.41	1.30	1.35
40	BG	176	G	C1'-N9	-7.41	1.36	1.46
85	AA	761	G	C2-N2	-7.41	1.27	1.34
85	AA	787	U	P-O5'	-7.41	1.52	1.59
35	BB	22	A	N7-C5	-7.41	1.34	1.39
35	BB	1541	G	C6-N1	-7.41	1.34	1.39
36	BC	64	U	O3'-P	-7.41	1.52	1.61
85	AA	755	G	N1-C2	-7.41	1.31	1.37
35	BB	43	G	P-O5'	-7.41	1.52	1.59
35	BB	1176	G	N9-C4	-7.41	1.32	1.38
34	BA	240	C	P-O5'	-7.41	1.52	1.59
34	BA	544	U	C2'-C1'	-7.41	1.45	1.53
34	BA	554	A	N7-C5	-7.41	1.34	1.39
34	BA	885	A	O3'-P	-7.41	1.52	1.61
35	BB	91	G	N7-C5	-7.41	1.34	1.39
35	BB	404	A	C2'-C1'	-7.41	1.45	1.53
85	AA	412	G	O3'-P	-7.41	1.52	1.61
85	AA	517	A	C2'-C1'	-7.41	1.45	1.53
85	AA	579	U	C2-N3	-7.41	1.32	1.37
85	AA	674	U	O3'-P	-7.41	1.52	1.61
85	AA	1701	G	O3'-P	-7.41	1.52	1.61
34	BA	433	G	N7-C5	-7.40	1.34	1.39
34	BA	888	G	C5-C4	-7.40	1.33	1.38
34	BA	1521	C	O3'-P	-7.40	1.52	1.61
36	BC	114	C	C2'-C1'	-7.40	1.45	1.53
41	BH	107	A	C5-C6	-7.40	1.34	1.41
85	AA	771	A	N3-C4	-7.40	1.30	1.34
85	AA	1855	U	P-O5'	-7.40	1.52	1.59
34	BA	3	G	P-O5'	-7.40	1.52	1.59
34	BA	195	G	C3'-C2'	-7.40	1.44	1.52
34	BA	343	G	O3'-P	-7.40	1.52	1.61
34	BA	1324	G	C5-C4	-7.40	1.33	1.38
35	BB	574	G	O4'-C1'	-7.40	1.32	1.41
38	BE	119	U	O3'-P	-7.40	1.52	1.61
85	AA	70	U	P-O5'	-7.40	1.52	1.59
85	AA	858	G	C2'-C1'	-7.40	1.45	1.53
85	AA	1524	A	O3'-P	-7.40	1.52	1.61
34	BA	171	U	C3'-C2'	-7.40	1.44	1.52
34	BA	238	C	C3'-C2'	-7.40	1.44	1.52
34	BA	700	G	P-O5'	-7.40	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	110	U	O3'-P	-7.40	1.52	1.61
39	BF	71	G	N7-C5	-7.40	1.34	1.39
85	AA	268	A	C4'-C3'	-7.40	1.45	1.53
85	AA	765	U	C4'-C3'	-7.40	1.45	1.53
34	BA	1255	G	O3'-P	-7.40	1.52	1.61
85	AA	9	U	C2-N3	-7.40	1.32	1.37
34	BA	425	G	N9-C4	-7.40	1.32	1.38
34	BA	697	A	C5-C4	-7.40	1.33	1.38
34	BA	1149	C	N3-C4	-7.40	1.28	1.33
40	BG	94	G	C6-N1	-7.40	1.34	1.39
85	AA	1664	G	O3'-P	-7.40	1.52	1.61
34	BA	132	U	N1-C2	-7.40	1.31	1.38
34	BA	257	G	C5-C4	-7.40	1.33	1.38
34	BA	895	U	C3'-C2'	-7.40	1.44	1.52
34	BA	1260	G	N9-C4	-7.40	1.32	1.38
34	BA	1603	A	C5-C4	-7.40	1.33	1.38
34	BA	6	C	C2-N3	-7.39	1.29	1.35
34	BA	237	A	C5-C4	-7.39	1.33	1.38
34	BA	1323	G	O3'-P	-7.39	1.52	1.61
34	BA	1522	G	C3'-C2'	-7.39	1.44	1.52
34	BA	1609	U	O4'-C1'	-7.39	1.32	1.41
35	BB	393	A	O3'-P	-7.39	1.52	1.61
35	BB	1286	G	C4'-C3'	-7.39	1.45	1.53
36	BC	144	C	O3'-P	-7.39	1.52	1.61
40	BG	118	U	C4'-C3'	-7.39	1.45	1.53
85	AA	291	G	C2'-C1'	-7.39	1.45	1.53
85	AA	934	A	P-O5'	-7.39	1.52	1.59
34	BA	142	A	N9-C4	-7.39	1.33	1.37
35	BB	403	U	O3'-P	-7.39	1.52	1.61
40	BG	110	U	C3'-C2'	-7.39	1.44	1.52
85	AA	38	C	C2'-C1'	-7.39	1.45	1.53
85	AA	1150	G	N9-C4	-7.39	1.32	1.38
85	AA	1260	G	C3'-C2'	-7.39	1.44	1.52
85	AA	1860	A	O3'-P	-7.39	1.52	1.61
85	AA	1872	G	N9-C4	-7.39	1.32	1.38
34	BA	663	U	C2'-C1'	-7.39	1.45	1.53
34	BA	852	C	O3'-P	-7.39	1.52	1.61
34	BA	969	A	C5-C4	-7.39	1.33	1.38
34	BA	1036	G	C2'-C1'	-7.39	1.45	1.53
34	BA	1507	C	C2'-C1'	-7.39	1.45	1.53
35	BB	59	U	O3'-P	-7.39	1.52	1.61
85	AA	984	A	P-O5'	-7.39	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	938	C	P-O5'	-7.39	1.52	1.59
35	BB	28	G	C5-C4	-7.39	1.33	1.38
35	BB	1384	A	C3'-C2'	-7.39	1.44	1.52
35	BB	1490	G	O4'-C1'	7.39	1.51	1.41
40	BG	66	C	C1'-N1	-7.39	1.36	1.46
85	AA	172	A	N9-C4	-7.39	1.33	1.37
86	AB	15	G	C2-N2	-7.39	1.27	1.34
34	BA	50	G	C5-C4	-7.39	1.33	1.38
34	BA	453	A	O3'-P	-7.39	1.52	1.61
35	BB	561	C	C4-C5	-7.39	1.37	1.43
35	BB	1038	G	P-O5'	-7.39	1.52	1.59
35	BB	1211	C	N1-C6	-7.39	1.32	1.37
35	BB	1543	C	P-O5'	-7.39	1.52	1.59
85	AA	1907	U	C2'-C1'	-7.39	1.45	1.53
34	BA	39	C	P-O5'	-7.39	1.52	1.59
34	BA	260	A	P-O5'	-7.39	1.52	1.59
34	BA	1042	U	O3'-P	-7.39	1.52	1.61
34	BA	1349	A	N9-C4	-7.39	1.33	1.37
35	BB	540	G	N9-C4	-7.39	1.32	1.38
35	BB	1406	C	N1-C6	-7.39	1.32	1.37
38	BE	63	C	C2'-C1'	-7.39	1.45	1.53
40	BG	20	U	O3'-P	-7.39	1.52	1.61
34	BA	346	A	C1'-N9	-7.38	1.36	1.46
34	BA	710	A	C4'-C3'	-7.38	1.45	1.53
35	BB	664	A	C1'-N9	-7.38	1.36	1.46
35	BB	1282	G	C3'-C2'	-7.38	1.44	1.52
85	AA	271	A	P-O5'	-7.38	1.52	1.59
85	AA	559	G	P-O5'	-7.38	1.52	1.59
85	AA	587	G	C2-N2	-7.38	1.27	1.34
85	AA	1057	G	O3'-P	-7.38	1.52	1.61
85	AA	1177	G	N7-C5	-7.38	1.34	1.39
85	AA	1199	C	O3'-P	-7.38	1.52	1.61
85	AA	1357	U	O3'-P	-7.38	1.52	1.61
34	BA	892	C	C1'-N1	-7.38	1.36	1.46
85	AA	1145	U	C4'-C3'	-7.38	1.45	1.53
85	AA	2017	U	P-O5'	-7.38	1.52	1.59
34	BA	1016	A	C2'-C1'	-7.38	1.45	1.53
34	BA	1530	G	O3'-P	-7.38	1.52	1.61
35	BB	362	A	C2'-C1'	-7.38	1.45	1.53
35	BB	513	G	O4'-C1'	-7.38	1.32	1.41
35	BB	1018	U	C2'-C1'	-7.38	1.45	1.53
35	BB	1308	G	N9-C4	-7.38	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	98	G	C4'-C3'	-7.38	1.45	1.53
38	BE	127	G	P-O5'	-7.38	1.52	1.59
41	BH	38	G	N3-C4	-7.38	1.30	1.35
85	AA	1135	U	C2-N3	-7.38	1.32	1.37
85	AA	2149	C	C3'-C2'	-7.38	1.44	1.52
34	BA	236	A	C3'-C2'	-7.38	1.44	1.52
34	BA	758	G	O4'-C1'	-7.38	1.32	1.41
85	AA	1135	U	P-O5'	-7.38	1.52	1.59
85	AA	1284	A	O3'-P	-7.38	1.52	1.61
34	BA	610	A	C5-C4	-7.38	1.33	1.38
35	BB	124	G	O3'-P	-7.38	1.52	1.61
35	BB	431	U	C4'-C3'	-7.38	1.45	1.53
36	BC	131	C	O3'-P	-7.38	1.52	1.61
85	AA	419	A	N3-C4	-7.38	1.30	1.34
85	AA	591	A	C1'-N9	-7.38	1.36	1.46
85	AA	678	A	C1'-N9	-7.38	1.36	1.46
85	AA	716	G	P-O5'	-7.38	1.52	1.59
85	AA	2199	G	O3'-P	-7.38	1.52	1.61
34	BA	905	A	P-O5'	-7.38	1.52	1.59
34	BA	1073	G	C1'-N9	-7.38	1.36	1.46
34	BA	1221	A	O4'-C1'	-7.38	1.32	1.41
35	BB	78	C	O3'-P	-7.38	1.52	1.61
35	BB	1456	G	C4'-C3'	-7.38	1.45	1.53
38	BE	141	A	C5-C6	-7.38	1.34	1.41
85	AA	1346	C	P-O5'	-7.38	1.52	1.59
85	AA	1512	U	O3'-P	-7.38	1.52	1.61
34	BA	217	C	P-O5'	-7.38	1.52	1.59
34	BA	293	A	P-O5'	-7.38	1.52	1.59
34	BA	399	G	C8-N7	-7.37	1.26	1.30
34	BA	809	U	O3'-P	-7.37	1.52	1.61
85	AA	1678	U	P-O5'	-7.37	1.52	1.59
85	AA	1877	G	N7-C5	-7.37	1.34	1.39
85	AA	2105	G	O3'-P	-7.37	1.52	1.61
34	BA	1293	A	N7-C5	-7.37	1.34	1.39
34	BA	1645	C	C2'-C1'	-7.37	1.45	1.53
35	BB	1182	A	C2'-C1'	-7.37	1.45	1.53
35	BB	1523	U	C2-N3	-7.37	1.32	1.37
40	BG	6	A	C2'-C1'	-7.37	1.45	1.53
41	BH	53	C	C2'-C1'	-7.37	1.45	1.53
85	AA	759	G	C6-N1	-7.37	1.34	1.39
85	AA	1900	C	O3'-P	-7.37	1.52	1.61
34	BA	1014	A	O3'-P	-7.37	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1338	G	O3'-P	-7.37	1.52	1.61
35	BB	1112	U	P-O5'	-7.37	1.52	1.59
38	BE	64	A	C2'-C1'	-7.37	1.45	1.53
85	AA	857	G	C3'-C2'	-7.37	1.44	1.52
35	BB	1423	U	C2-N3	-7.37	1.32	1.37
40	BG	71	C	C4'-C3'	-7.37	1.45	1.53
34	BA	396	U	C2'-C1'	-7.37	1.45	1.53
34	BA	728	A	N3-C4	-7.37	1.30	1.34
35	BB	1369	A	C3'-C2'	-7.37	1.44	1.52
85	AA	406	U	P-O5'	-7.37	1.52	1.59
85	AA	810	C	C2-N3	-7.37	1.29	1.35
34	BA	260	A	C6-N1	-7.37	1.30	1.35
34	BA	1292	A	N7-C5	-7.37	1.34	1.39
35	BB	584	A	N7-C5	-7.37	1.34	1.39
35	BB	1108	G	C6-N1	-7.37	1.34	1.39
36	BC	157	U	O3'-P	-7.37	1.52	1.61
38	BE	63	C	C1'-N1	-7.37	1.36	1.46
38	BE	110	U	C2'-C1'	-7.37	1.45	1.53
40	BG	182	G	N9-C8	-7.37	1.32	1.37
34	BA	101	G	N1-C2	-7.36	1.31	1.37
34	BA	498	A	C5-C4	-7.36	1.33	1.38
34	BA	1083	A	P-O5'	-7.36	1.52	1.59
34	BA	1479	G	O3'-P	-7.36	1.52	1.61
34	BA	1568	A	C6-N1	7.36	1.40	1.35
35	BB	73	G	C2-N2	-7.36	1.27	1.34
35	BB	112	G	C2-N3	-7.36	1.26	1.32
35	BB	410	A	P-O5'	-7.36	1.52	1.59
35	BB	1182	A	N9-C4	-7.36	1.33	1.37
35	BB	1438	U	C2'-C1'	-7.36	1.45	1.53
38	BE	86	C	C2'-C1'	-7.36	1.45	1.53
85	AA	931	G	N9-C4	-7.36	1.32	1.38
34	BA	1256	A	O3'-P	-7.36	1.52	1.61
34	BA	1629	A	O3'-P	-7.36	1.52	1.61
85	AA	1793	A	O4'-C1'	-7.36	1.32	1.41
85	AA	2217	A	C4'-C3'	-7.36	1.45	1.53
34	BA	900	A	C8-N7	-7.36	1.26	1.31
34	BA	1437	G	C6-N1	-7.36	1.34	1.39
35	BB	1087	A	N7-C5	-7.36	1.34	1.39
35	BB	1547	U	P-O5'	-7.36	1.52	1.59
85	AA	318	A	O3'-P	-7.36	1.52	1.61
34	BA	1000	G	P-O5'	-7.36	1.52	1.59
35	BB	13	A	C2'-C1'	-7.36	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1195	A	C2'-C1'	-7.36	1.45	1.53
85	AA	2	A	N9-C4	-7.36	1.33	1.37
85	AA	779	G	P-O5'	-7.36	1.52	1.59
85	AA	1238	U	C2'-C1'	-7.36	1.45	1.53
34	BA	970	U	C2'-C1'	-7.36	1.45	1.53
34	BA	1256	A	N7-C5	-7.36	1.34	1.39
35	BB	557	C	N3-C4	-7.36	1.28	1.33
36	BC	14	G	C3'-C2'	-7.36	1.44	1.52
85	AA	1176	C	O3'-P	-7.36	1.52	1.61
34	BA	974	G	C2-N2	-7.36	1.27	1.34
34	BA	1379	G	O3'-P	-7.36	1.52	1.61
34	BA	1422	A	C1'-N9	-7.36	1.36	1.46
35	BB	95	A	C2'-C1'	-7.36	1.45	1.53
35	BB	784	C	C3'-C2'	-7.36	1.44	1.52
38	BE	11	A	O3'-P	-7.36	1.52	1.61
85	AA	589	A	C5'-C4'	7.36	1.60	1.51
34	BA	563	A	P-O5'	-7.35	1.52	1.59
35	BB	804	U	C4'-C3'	-7.35	1.45	1.53
36	BC	106	G	N1-C2	-7.35	1.31	1.37
85	AA	1175	A	O3'-P	-7.35	1.52	1.61
85	AA	1524	A	N3-C4	-7.35	1.30	1.34
34	BA	11	U	P-O5'	-7.35	1.52	1.59
34	BA	1153	C	O3'-P	-7.35	1.52	1.61
34	BA	1230	G	N3-C4	-7.35	1.30	1.35
34	BA	1685	C	C2-N3	-7.35	1.29	1.35
34	BA	1829	A	P-O5'	-7.35	1.52	1.59
35	BB	29	C	C2'-C1'	-7.35	1.45	1.53
35	BB	534	C	C3'-C2'	-7.35	1.44	1.52
35	BB	877	A	C2'-C1'	-7.35	1.45	1.53
37	BD	90	A	N7-C5	-7.35	1.34	1.39
85	AA	298	C	O3'-P	-7.35	1.52	1.61
85	AA	329	G	C1'-N9	-7.35	1.36	1.46
85	AA	925	G	P-O5'	-7.35	1.52	1.59
35	BB	111	C	O3'-P	-7.35	1.52	1.61
35	BB	1125	A	N9-C4	-7.35	1.33	1.37
85	AA	1613	A	P-O5'	-7.35	1.52	1.59
34	BA	1422	A	N7-C5	-7.35	1.34	1.39
35	BB	51	U	C3'-C2'	-7.35	1.44	1.52
35	BB	677	U	P-O5'	-7.35	1.52	1.59
35	BB	1487	G	C6-N1	-7.35	1.34	1.39
85	AA	134	U	P-O5'	-7.35	1.52	1.59
85	AA	722	G	N7-C5	-7.35	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	883	A	N9-C4	-7.35	1.33	1.37
85	AA	2134	U	P-O5'	-7.35	1.52	1.59
85	AA	2239	A	P-O5'	-7.35	1.52	1.59
34	BA	1312	A	C4'-C3'	-7.35	1.45	1.53
35	BB	989	C	P-O5'	-7.35	1.52	1.59
37	BD	15	U	C2'-C1'	-7.35	1.45	1.53
85	AA	457	G	C5-C4	-7.35	1.33	1.38
34	BA	1123	G	N7-C5	-7.35	1.34	1.39
34	BA	1534	U	N3-C4	-7.35	1.31	1.38
35	BB	71	A	N9-C4	-7.35	1.33	1.37
35	BB	1198	C	C3'-C2'	-7.35	1.44	1.52
34	BA	108	A	C2'-C1'	-7.34	1.45	1.53
34	BA	340	U	O3'-P	-7.34	1.52	1.61
34	BA	494	A	C1'-N9	-7.34	1.36	1.46
34	BA	544	U	P-O5'	-7.34	1.52	1.59
35	BB	896	C	O3'-P	-7.34	1.52	1.61
36	BC	23	G	O3'-P	-7.34	1.52	1.61
38	BE	89	G	N1-C2	-7.34	1.31	1.37
40	BG	149	U	O4'-C1'	-7.34	1.32	1.41
41	BH	115	A	P-O5'	-7.34	1.52	1.59
85	AA	1000	U	C2-N3	-7.34	1.32	1.37
34	BA	1605	G	O3'-P	-7.34	1.52	1.61
40	BG	156	G	N3-C4	-7.34	1.30	1.35
34	BA	960	C	C2-N3	-7.34	1.29	1.35
34	BA	1202	G	N9-C8	-7.34	1.32	1.37
34	BA	1483	U	C1'-N1	-7.34	1.36	1.46
34	BA	1599	A	C4'-C3'	-7.34	1.45	1.53
34	BA	1843	G	C1'-N9	-7.34	1.36	1.46
35	BB	1418	C	C2-N3	-7.34	1.29	1.35
40	BG	54	G	O3'-P	-7.34	1.52	1.61
40	BG	78	C	P-O5'	-7.34	1.52	1.59
85	AA	923	A	C4'-C3'	-7.34	1.45	1.53
85	AA	1247	A	C4'-C3'	-7.34	1.45	1.53
85	AA	1549	G	P-O5'	-7.34	1.52	1.59
85	AA	1730	C	P-O5'	-7.34	1.52	1.59
34	BA	385	U	P-O5'	-7.34	1.52	1.59
34	BA	1075	U	O3'-P	-7.34	1.52	1.61
34	BA	1833	G	C6-N1	-7.34	1.34	1.39
35	BB	111	C	C2'-C1'	-7.34	1.45	1.53
35	BB	648	G	C2-N2	-7.34	1.27	1.34
35	BB	793	A	C3'-C2'	-7.34	1.44	1.52
36	BC	73	U	C2'-C1'	-7.34	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	97	U	C4'-O4'	-7.34	1.36	1.45
85	AA	162	A	O3'-P	-7.34	1.52	1.61
85	AA	164	G	N9-C8	-7.34	1.32	1.37
85	AA	720	A	P-O5'	-7.34	1.52	1.59
85	AA	1221	G	N7-C5	-7.34	1.34	1.39
85	AA	2236	U	C3'-C2'	-7.34	1.44	1.52
34	BA	257	G	C3'-C2'	-7.34	1.44	1.52
34	BA	1816	G	N9-C8	-7.34	1.32	1.37
35	BB	1306	G	C1'-N9	-7.34	1.36	1.46
35	BB	1336	G	P-O5'	-7.34	1.52	1.59
36	BC	43	A	P-O5'	-7.34	1.52	1.59
85	AA	939	A	N9-C8	-7.34	1.31	1.37
34	BA	246	G	C6-N1	-7.34	1.34	1.39
34	BA	854	A	O3'-P	-7.34	1.52	1.61
34	BA	1226	G	C1'-N9	-7.34	1.36	1.46
34	BA	1443	U	N1-C2	-7.34	1.31	1.38
34	BA	1720	U	C4'-C3'	-7.34	1.45	1.53
35	BB	1071	G	P-O5'	-7.34	1.52	1.59
35	BB	1096	G	C3'-C2'	-7.34	1.44	1.52
35	BB	1356	G	P-O5'	-7.34	1.52	1.59
37	BD	38	U	O3'-P	-7.34	1.52	1.61
37	BD	79	G	C1'-N9	-7.34	1.36	1.46
85	AA	75	U	O3'-P	-7.34	1.52	1.61
85	AA	422	G	N1-C2	-7.34	1.31	1.37
85	AA	925	G	C6-N1	-7.34	1.34	1.39
34	BA	392	A	O3'-P	-7.33	1.52	1.61
34	BA	755	G	P-O5'	-7.33	1.52	1.59
34	BA	948	C	C2'-C1'	-7.33	1.45	1.53
85	AA	386	G	C3'-C2'	-7.33	1.44	1.52
85	AA	741	G	N1-C2	-7.33	1.31	1.37
35	BB	651	G	N9-C4	-7.33	1.32	1.38
35	BB	665	A	C6-N6	-7.33	1.28	1.33
85	AA	18	C	C2-N3	-7.33	1.29	1.35
85	AA	332	A	O3'-P	-7.33	1.52	1.61
85	AA	335	G	P-O5'	-7.33	1.52	1.59
85	AA	1563	U	O3'-P	-7.33	1.52	1.61
85	AA	1955	U	C2'-C1'	-7.33	1.45	1.53
34	BA	37	A	P-O5'	-7.33	1.52	1.59
34	BA	252	A	C1'-N9	-7.33	1.36	1.46
34	BA	749	G	C2'-C1'	-7.33	1.45	1.53
34	BA	889	U	C1'-N1	-7.33	1.36	1.46
34	BA	1063	G	C6-N1	-7.33	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1255	G	N7-C5	-7.33	1.34	1.39
34	BA	1430	C	P-O5'	-7.33	1.52	1.59
34	BA	1481	U	C4'-C3'	-7.33	1.45	1.53
35	BB	1102	U	C1'-N1	-7.33	1.36	1.46
35	BB	1214	U	P-O5'	-7.33	1.52	1.59
34	BA	37	A	O3'-P	-7.33	1.52	1.61
35	BB	1037	A	C1'-N9	-7.33	1.36	1.46
35	BB	1398	A	C5-C4	-7.33	1.33	1.38
85	AA	2095	U	P-O5'	-7.33	1.52	1.59
34	BA	743	A	C8-N7	7.33	1.36	1.31
34	BA	1305	A	C2'-C1'	-7.33	1.45	1.53
35	BB	485	U	O3'-P	-7.33	1.52	1.61
40	BG	81	G	C2'-C1'	-7.33	1.45	1.53
40	BG	125	C	C2'-C1'	-7.33	1.45	1.53
85	AA	702	G	C5-C4	-7.33	1.33	1.38
35	BB	787	A	C2'-C1'	-7.33	1.45	1.53
85	AA	701	C	C2'-C1'	-7.33	1.45	1.53
85	AA	775	C	N1-C6	-7.33	1.32	1.37
34	BA	497	U	C2-N3	-7.33	1.32	1.37
34	BA	856	G	C6-N1	-7.33	1.34	1.39
34	BA	914	G	N7-C5	-7.33	1.34	1.39
34	BA	986	G	C5-C4	-7.33	1.33	1.38
34	BA	992	A	C5-C4	-7.33	1.33	1.38
34	BA	1288	U	C3'-C2'	-7.33	1.44	1.52
35	BB	126	C	C4-C5	-7.33	1.37	1.43
35	BB	1301	U	O3'-P	-7.33	1.52	1.61
36	BC	104	A	C4'-C3'	-7.33	1.45	1.53
38	BE	160	C	O3'-P	-7.33	1.52	1.61
85	AA	413	G	C5'-C4'	-7.33	1.42	1.51
85	AA	1290	G	C4'-C3'	-7.33	1.45	1.53
85	AA	2221	A	N3-C4	-7.33	1.30	1.34
35	BB	592	G	C6-N1	-7.32	1.34	1.39
35	BB	983	C	O3'-P	-7.32	1.52	1.61
36	BC	41	A	C3'-C2'	-7.32	1.44	1.52
36	BC	88	A	N7-C5	-7.32	1.34	1.39
40	BG	108	G	C6-N1	-7.32	1.34	1.39
85	AA	40	A	N7-C5	-7.32	1.34	1.39
85	AA	503	A	N9-C4	-7.32	1.33	1.37
85	AA	1784	G	P-O5'	-7.32	1.52	1.59
34	BA	17	A	C4'-C3'	-7.32	1.45	1.53
34	BA	1611	A	P-O5'	-7.32	1.52	1.59
35	BB	25	A	C5'-C4'	-7.32	1.42	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	134	G	C2-N2	-7.32	1.27	1.34
85	AA	899	A	C5'-C4'	7.32	1.60	1.51
85	AA	1372	C	C2'-C1'	-7.32	1.45	1.53
34	BA	460	G	C3'-C2'	-7.32	1.44	1.52
34	BA	1520	A	C3'-C2'	-7.32	1.44	1.52
35	BB	35	G	C2-N2	-7.32	1.27	1.34
35	BB	998	G	O3'-P	-7.32	1.52	1.61
38	BE	178	G	N9-C4	-7.32	1.32	1.38
40	BG	38	A	C1'-N9	-7.32	1.36	1.46
85	AA	1708	A	C1'-N9	-7.32	1.36	1.46
85	AA	1857	G	P-O5'	-7.32	1.52	1.59
85	AA	2240	G	O3'-P	-7.32	1.52	1.61
35	BB	877	A	N9-C4	7.32	1.42	1.37
85	AA	816	A	N9-C4	7.32	1.42	1.37
85	AA	1197	U	C2'-C1'	-7.32	1.45	1.53
34	BA	1430	C	C1'-N1	-7.32	1.36	1.46
34	BA	1539	A	O3'-P	-7.32	1.52	1.61
35	BB	662	G	C6-N1	-7.32	1.34	1.39
35	BB	870	C	P-O5'	-7.32	1.52	1.59
35	BB	969	C	C2'-C1'	-7.32	1.45	1.53
35	BB	1384	A	O3'-P	-7.32	1.52	1.61
37	BD	105	G	C1'-N9	-7.32	1.36	1.46
85	AA	44	C	O3'-P	-7.32	1.52	1.61
85	AA	567	G	C2'-C1'	-7.32	1.45	1.53
85	AA	1129	A	N7-C5	-7.32	1.34	1.39
85	AA	1795	C	C2'-C1'	-7.32	1.45	1.53
85	AA	2197	A	C3'-C2'	-7.32	1.44	1.52
34	BA	70	C	C2'-C1'	-7.32	1.45	1.53
34	BA	235	C	P-O5'	-7.32	1.52	1.59
34	BA	722	A	N7-C5	-7.32	1.34	1.39
34	BA	1093	G	P-O5'	-7.32	1.52	1.59
34	BA	1408	C	C3'-C2'	-7.32	1.44	1.52
40	BG	100	G	C2'-C1'	-7.32	1.45	1.53
40	BG	148	C	O3'-P	-7.32	1.52	1.61
85	AA	2	A	N7-C5	-7.32	1.34	1.39
85	AA	1520	A	P-O5'	-7.32	1.52	1.59
35	BB	1067	G	N3-C4	-7.31	1.30	1.35
36	BC	129	C	C2'-C1'	-7.31	1.45	1.53
40	BG	4	A	N7-C5	-7.31	1.34	1.39
85	AA	1547	G	C2'-C1'	-7.31	1.45	1.53
34	BA	785	G	C6-N1	-7.31	1.34	1.39
34	BA	1552	C	O3'-P	-7.31	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	376	A	C5-C4	-7.31	1.33	1.38
35	BB	450	A	C4'-C3'	-7.31	1.45	1.53
35	BB	1232	A	O3'-P	-7.31	1.52	1.61
35	BB	1237	C	C1'-N1	-7.31	1.36	1.46
36	BC	41	A	O3'-P	-7.31	1.52	1.61
34	BA	382	G	N9-C4	7.31	1.43	1.38
35	BB	101	U	N3-C4	-7.31	1.31	1.38
35	BB	621	C	C3'-C2'	-7.31	1.44	1.52
36	BC	166	G	N7-C5	-7.31	1.34	1.39
37	BD	103	C	O3'-P	-7.31	1.52	1.61
40	BG	93	U	P-O5'	-7.31	1.52	1.59
85	AA	1667	C	C3'-C2'	-7.31	1.44	1.52
85	AA	2124	G	C1'-N9	-7.31	1.36	1.46
34	BA	541	C	C2'-C1'	-7.31	1.45	1.53
34	BA	1793	G	N3-C4	-7.31	1.30	1.35
41	BH	124	C	C4'-C3'	-7.31	1.45	1.53
85	AA	1468	G	C4'-C3'	7.31	1.61	1.53
34	BA	835	U	O3'-P	-7.31	1.52	1.61
34	BA	1516	G	C2'-C1'	-7.31	1.45	1.53
35	BB	381	C	P-O5'	-7.31	1.52	1.59
35	BB	1200	A	C3'-C2'	-7.31	1.44	1.52
35	BB	1242	C	O3'-P	-7.31	1.52	1.61
40	BG	129	G	O3'-P	-7.31	1.52	1.61
85	AA	312	G	C6-N1	-7.31	1.34	1.39
85	AA	2190	U	N3-C4	-7.31	1.31	1.38
34	BA	277	A	O3'-P	-7.31	1.52	1.61
34	BA	1156	U	C1'-N1	-7.31	1.36	1.46
34	BA	1547	G	O3'-P	-7.31	1.52	1.61
35	BB	1056	A	P-O5'	-7.31	1.52	1.59
40	BG	4	A	C2'-C1'	-7.31	1.45	1.53
85	AA	30	G	N9-C4	-7.31	1.32	1.38
34	BA	55	G	C2'-C1'	-7.30	1.45	1.53
34	BA	330	A	C5-C4	-7.30	1.33	1.38
34	BA	1552	C	C2-N3	-7.30	1.29	1.35
34	BA	1686	G	C5-C4	-7.30	1.33	1.38
35	BB	75	A	P-O5'	-7.30	1.52	1.59
35	BB	136	A	C4'-C3'	7.30	1.61	1.53
35	BB	431	U	O4'-C1'	-7.30	1.32	1.41
85	AA	575	G	O3'-P	-7.30	1.52	1.61
85	AA	1483	A	O3'-P	-7.30	1.52	1.61
34	BA	116	G	C4'-O4'	-7.30	1.36	1.45
34	BA	1559	C	C1'-N1	-7.30	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	119	G	C6-N1	-7.30	1.34	1.39
35	BB	1089	A	O3'-P	-7.30	1.52	1.61
35	BB	1186	A	P-O5'	-7.30	1.52	1.59
85	AA	1098	C	C2'-C1'	-7.30	1.45	1.53
34	BA	25	C	P-O5'	-7.30	1.52	1.59
34	BA	36	A	O3'-P	-7.30	1.52	1.61
34	BA	123	C	C2'-C1'	-7.30	1.45	1.53
34	BA	1601	C	O3'-P	-7.30	1.52	1.61
35	BB	443	A	O3'-P	-7.30	1.52	1.61
35	BB	587	A	C8-N7	-7.30	1.26	1.31
35	BB	617	C	C3'-C2'	-7.30	1.44	1.52
85	AA	96	C	C2'-C1'	-7.30	1.45	1.53
34	BA	373	G	N9-C4	-7.30	1.32	1.38
34	BA	937	G	O3'-P	-7.30	1.52	1.61
34	BA	1025	A	N7-C5	-7.30	1.34	1.39
34	BA	1219	G	O3'-P	-7.30	1.52	1.61
34	BA	1621	U	P-O5'	-7.30	1.52	1.59
34	BA	1654	G	N9-C4	-7.30	1.32	1.38
35	BB	80	C	N3-C4	-7.30	1.28	1.33
35	BB	1126	A	P-O5'	-7.30	1.52	1.59
36	BC	143	C	C2-N3	-7.30	1.29	1.35
38	BE	35	A	O3'-P	-7.30	1.52	1.61
38	BE	97	G	N9-C4	-7.30	1.32	1.38
85	AA	424	A	C6-N6	-7.30	1.28	1.33
85	AA	659	A	C3'-C2'	-7.30	1.44	1.52
85	AA	1117	G	O4'-C1'	-7.30	1.32	1.41
34	BA	401	A	C2'-C1'	-7.30	1.45	1.53
35	BB	664	A	N9-C4	-7.30	1.33	1.37
38	BE	178	G	C1'-N9	-7.30	1.36	1.46
40	BG	152	G	O3'-P	-7.30	1.52	1.61
85	AA	688	C	O3'-P	-7.30	1.52	1.61
85	AA	918	U	P-O5'	-7.30	1.52	1.59
85	AA	1987	G	O3'-P	-7.30	1.52	1.61
34	BA	1158	A	O3'-P	-7.30	1.52	1.61
35	BB	839	G	C2-N2	-7.30	1.27	1.34
85	AA	521	A	C1'-N9	-7.30	1.36	1.46
85	AA	652	U	O3'-P	-7.30	1.52	1.61
85	AA	869	A	N3-C4	-7.30	1.30	1.34
85	AA	881	C	P-O5'	-7.30	1.52	1.59
85	AA	1269	A	C5-C4	-7.30	1.33	1.38
85	AA	1524	A	C3'-C2'	-7.30	1.44	1.52
85	AA	2014	G	N9-C4	-7.30	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	79	C	C2'-C1'	-7.29	1.45	1.53
34	BA	1456	C	C2'-C1'	-7.29	1.45	1.53
85	AA	1655	G	N9-C8	-7.29	1.32	1.37
34	BA	234	A	C3'-C2'	-7.29	1.44	1.52
34	BA	297	A	N7-C5	-7.29	1.34	1.39
34	BA	982	A	C5-C4	-7.29	1.33	1.38
34	BA	1136	A	P-O5'	-7.29	1.52	1.59
34	BA	1238	C	O3'-P	-7.29	1.52	1.61
35	BB	1214	U	C3'-C2'	-7.29	1.44	1.52
35	BB	1258	G	N9-C4	-7.29	1.32	1.38
38	BE	130	G	C2'-C1'	-7.29	1.45	1.53
34	BA	605	G	C2-N3	7.29	1.38	1.32
34	BA	702	G	C2-N2	-7.29	1.27	1.34
34	BA	1676	A	C3'-C2'	-7.29	1.44	1.52
34	BA	1707	C	C2-N3	-7.29	1.29	1.35
34	BA	1806	A	C5-C4	-7.29	1.33	1.38
34	BA	1831	A	P-O5'	-7.29	1.52	1.59
40	BG	126	G	C1'-N9	-7.29	1.36	1.46
85	AA	1263	G	N3-C4	-7.29	1.30	1.35
34	BA	395	G	N9-C4	-7.29	1.32	1.38
34	BA	784	C	O3'-P	-7.29	1.52	1.61
34	BA	1642	A	C1'-N9	-7.29	1.36	1.46
35	BB	692	G	O3'-P	-7.29	1.52	1.61
36	BC	66	G	C1'-N9	-7.29	1.36	1.46
85	AA	482	C	O3'-P	-7.29	1.52	1.61
86	AB	8	U	O3'-P	-7.29	1.52	1.61
35	BB	408	U	P-O5'	-7.29	1.52	1.59
85	AA	1891	U	P-O5'	-7.29	1.52	1.59
34	BA	1031	U	O3'-P	-7.29	1.52	1.61
34	BA	1840	C	C2-N3	-7.29	1.29	1.35
85	AA	1485	G	N9-C4	-7.29	1.32	1.38
85	AA	1594	G	C2'-C1'	-7.29	1.45	1.53
34	BA	198	U	C1'-N1	-7.29	1.36	1.46
34	BA	376	U	C1'-N1	-7.29	1.36	1.46
34	BA	400	A	C6-N1	-7.29	1.30	1.35
34	BA	581	U	C3'-O3'	7.29	1.52	1.42
34	BA	811	C	C2'-C1'	-7.29	1.45	1.53
34	BA	1473	A	P-O5'	-7.29	1.52	1.59
35	BB	768	A	P-O5'	-7.29	1.52	1.59
35	BB	1100	C	C4'-C3'	-7.29	1.45	1.53
85	AA	423	G	C2'-C1'	-7.29	1.45	1.53
85	AA	2053	A	C2'-C1'	-7.29	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	487	A	N3-C4	-7.28	1.30	1.34
34	BA	987	C	C3'-C2'	-7.28	1.44	1.52
34	BA	1249	G	C1'-N9	-7.28	1.36	1.46
34	BA	1615	A	C1'-N9	-7.28	1.36	1.46
35	BB	482	A	N7-C5	-7.28	1.34	1.39
35	BB	695	U	N3-C4	-7.28	1.31	1.38
38	BE	163	A	C8-N7	-7.28	1.26	1.31
40	BG	135	C	C2'-C1'	-7.28	1.45	1.53
40	BG	155	A	O3'-P	-7.28	1.52	1.61
40	BG	182	G	C8-N7	-7.28	1.26	1.30
85	AA	247	G	C2'-C1'	-7.28	1.45	1.53
85	AA	1117	G	P-O5'	-7.28	1.52	1.59
34	BA	3	G	C5'-C4'	-7.28	1.42	1.51
36	BC	57	C	O3'-P	-7.28	1.52	1.61
85	AA	168	A	N9-C4	7.28	1.42	1.37
85	AA	1127	G	C2'-C1'	-7.28	1.45	1.53
34	BA	169	C	C2'-C1'	-7.28	1.45	1.53
34	BA	572	G	C1'-N9	7.28	1.59	1.48
34	BA	1337	A	P-O5'	-7.28	1.52	1.59
35	BB	27	C	C2-N3	-7.28	1.29	1.35
35	BB	637	G	N7-C5	-7.28	1.34	1.39
35	BB	658	G	C1'-N9	-7.28	1.36	1.46
35	BB	1247	C	P-O5'	-7.28	1.52	1.59
38	BE	42	C	O3'-P	-7.28	1.52	1.61
85	AA	754	C	C4'-C3'	-7.28	1.45	1.53
85	AA	1209	U	N3-C4	-7.28	1.31	1.38
34	BA	911	G	C5-C4	-7.28	1.33	1.38
35	BB	980	G	P-O5'	-7.28	1.52	1.59
37	BD	75	G	C2'-C1'	-7.28	1.45	1.53
40	BG	1	G	N1-C2	-7.28	1.31	1.37
40	BG	142	A	C5-C4	-7.28	1.33	1.38
34	BA	36	A	N7-C5	-7.28	1.34	1.39
34	BA	91	C	P-O5'	-7.28	1.52	1.59
34	BA	355	U	O3'-P	-7.28	1.52	1.61
34	BA	1648	G	N7-C5	-7.28	1.34	1.39
35	BB	79	U	C2-N3	-7.28	1.32	1.37
35	BB	1004	A	C3'-C2'	-7.28	1.44	1.52
36	BC	149	A	N9-C4	7.28	1.42	1.37
37	BD	54	A	P-O5'	-7.28	1.52	1.59
40	BG	106	G	C1'-N9	-7.28	1.36	1.46
85	AA	286	C	C3'-C2'	-7.28	1.44	1.52
85	AA	428	G	C5-C4	-7.28	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	459	C	O3'-P	-7.28	1.52	1.61
85	AA	805	A	O3'-P	-7.28	1.52	1.61
85	AA	1515	A	C4'-C3'	-7.28	1.45	1.53
85	AA	1657	C	P-O5'	-7.28	1.52	1.59
85	AA	2167	A	C2'-C1'	-7.28	1.45	1.53
34	BA	443	U	O3'-P	-7.28	1.52	1.61
34	BA	852	C	N3-C4	-7.28	1.28	1.33
34	BA	946	A	C4'-O4'	-7.28	1.36	1.45
35	BB	961	G	P-O5'	-7.28	1.52	1.59
35	BB	1372	G	N1-C2	-7.28	1.31	1.37
38	BE	8	G	O3'-P	-7.28	1.52	1.61
85	AA	151	A	O3'-P	-7.28	1.52	1.61
85	AA	2139	G	C4'-C3'	-7.28	1.45	1.53
85	AA	2184	A	C2'-C1'	-7.28	1.45	1.53
34	BA	1287	G	O3'-P	-7.27	1.52	1.61
35	BB	560	C	O3'-P	-7.27	1.52	1.61
35	BB	837	A	C3'-C2'	-7.27	1.44	1.52
85	AA	258	G	C5-C4	-7.27	1.33	1.38
34	BA	135	G	C2'-C1'	-7.27	1.45	1.53
34	BA	344	G	C5-C4	-7.27	1.33	1.38
34	BA	496	G	C6-N1	-7.27	1.34	1.39
34	BA	1006	G	N1-C2	-7.27	1.31	1.37
34	BA	1089	U	P-O5'	-7.27	1.52	1.59
34	BA	1443	U	C4'-O4'	-7.27	1.36	1.45
35	BB	498	G	P-O5'	-7.27	1.52	1.59
35	BB	675	U	N3-C4	-7.27	1.31	1.38
35	BB	1219	A	N9-C4	-7.27	1.33	1.37
35	BB	1482	A	N9-C4	-7.27	1.33	1.37
37	BD	13	A	P-O5'	-7.27	1.52	1.59
38	BE	200	A	C1'-N9	-7.27	1.36	1.46
39	BF	69	A	C1'-N9	-7.27	1.36	1.46
85	AA	1998	A	C2'-C1'	-7.27	1.45	1.53
34	BA	49	A	C8-N7	-7.27	1.26	1.31
34	BA	698	U	C2'-C1'	-7.27	1.45	1.53
35	BB	116	G	C6-N1	-7.27	1.34	1.39
38	BE	182	U	P-O5'	-7.27	1.52	1.59
41	BH	106	G	C1'-N9	-7.27	1.36	1.46
85	AA	107	A	C2'-C1'	-7.27	1.45	1.53
85	AA	1496	U	C3'-C2'	-7.27	1.44	1.52
85	AA	1551	G	N7-C5	-7.27	1.34	1.39
86	AB	61	C	P-O5'	-7.27	1.52	1.59
34	BA	75	U	N3-C4	-7.27	1.31	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	529	A	P-O5'	-7.27	1.52	1.59
34	BA	1483	U	C3'-C2'	-7.27	1.44	1.52
35	BB	841	U	N3-C4	-7.27	1.31	1.38
35	BB	872	A	C3'-C2'	-7.27	1.44	1.52
35	BB	1021	C	C1'-N1	-7.27	1.36	1.46
36	BC	160	C	C2-N3	-7.27	1.29	1.35
40	BG	65	C	C3'-C2'	-7.27	1.44	1.52
40	BG	92	U	C1'-N1	-7.27	1.36	1.46
86	AB	49	C	O3'-P	-7.27	1.52	1.61
34	BA	709	C	O3'-P	-7.27	1.52	1.61
34	BA	1575	U	C3'-C2'	-7.27	1.44	1.52
35	BB	397	C	C2-N3	-7.27	1.29	1.35
35	BB	1341	U	C5'-C4'	-7.27	1.42	1.51
35	BB	1395	G	C2'-C1'	-7.27	1.45	1.53
37	BD	74	A	C1'-N9	-7.27	1.36	1.46
38	BE	165	U	P-O5'	-7.27	1.52	1.59
40	BG	148	C	C5'-C4'	-7.27	1.42	1.51
85	AA	307	G	N7-C5	-7.27	1.34	1.39
85	AA	1229	G	C6-N1	-7.27	1.34	1.39
85	AA	1883	C	P-O5'	-7.27	1.52	1.59
35	BB	495	A	C1'-N9	-7.27	1.36	1.46
35	BB	833	G	C2'-C1'	-7.27	1.45	1.53
85	AA	1548	A	C2'-C1'	-7.27	1.45	1.53
34	BA	255	G	P-O5'	-7.26	1.52	1.59
34	BA	1032	A	O3'-P	-7.26	1.52	1.61
34	BA	1056	C	O3'-P	-7.26	1.52	1.61
35	BB	1182	A	C3'-C2'	-7.26	1.44	1.52
35	BB	1359	G	C1'-N9	-7.26	1.36	1.46
40	BG	105	A	C3'-C2'	-7.26	1.44	1.52
41	BH	21	G	C4'-C3'	-7.26	1.45	1.53
34	BA	700	G	C1'-N9	-7.26	1.36	1.46
38	BE	171	U	P-O5'	-7.26	1.52	1.59
38	BE	196	C	C2-N3	-7.26	1.29	1.35
34	BA	575	U	C5'-C4'	7.26	1.60	1.51
35	BB	37	C	O3'-P	-7.26	1.52	1.61
35	BB	831	C	C2-N3	-7.26	1.29	1.35
35	BB	1155	U	C3'-C2'	-7.26	1.44	1.52
35	BB	1382	U	C3'-C2'	-7.26	1.44	1.52
37	BD	87	G	N9-C4	-7.26	1.32	1.38
38	BE	32	U	C4'-O4'	-7.26	1.36	1.45
40	BG	61	A	C1'-N9	-7.26	1.36	1.46
40	BG	118	U	O3'-P	-7.26	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	118	U	C3'-C2'	-7.26	1.44	1.52
85	AA	310	U	P-O5'	-7.26	1.52	1.59
85	AA	1207	C	P-O5'	-7.26	1.52	1.59
85	AA	1281	G	C5-C4	-7.26	1.33	1.38
34	BA	1413	G	N7-C5	-7.26	1.34	1.39
34	BA	1590	G	N1-C2	-7.26	1.31	1.37
35	BB	1237	C	O3'-P	-7.26	1.52	1.61
35	BB	1524	G	C2'-C1'	-7.26	1.45	1.53
36	BC	18	G	C2-N2	-7.26	1.27	1.34
38	BE	198	A	C2'-C1'	-7.26	1.45	1.53
85	AA	102	A	C2'-C1'	-7.26	1.45	1.53
34	BA	830	U	C2'-C1'	-7.26	1.45	1.53
35	BB	808	U	N3-C4	-7.26	1.31	1.38
85	AA	1116	G	C2'-C1'	-7.26	1.45	1.53
34	BA	1334	G	C2'-C1'	-7.26	1.45	1.53
35	BB	598	C	C2'-C1'	-7.26	1.45	1.53
35	BB	648	G	C2'-C1'	-7.26	1.45	1.53
35	BB	801	G	P-O5'	-7.26	1.52	1.59
35	BB	1109	A	C2'-C1'	-7.26	1.45	1.53
35	BB	1194	A	N7-C5	-7.26	1.34	1.39
85	AA	105	A	C1'-N9	-7.26	1.36	1.46
85	AA	329	G	C2'-C1'	-7.26	1.45	1.53
85	AA	1216	A	C5'-C4'	-7.26	1.42	1.51
35	BB	485	U	C3'-C2'	-7.25	1.44	1.52
35	BB	543	G	C6-N1	-7.25	1.34	1.39
35	BB	1023	G	C2'-C1'	-7.25	1.45	1.53
85	AA	488	G	N9-C8	-7.25	1.32	1.37
85	AA	1137	C	O3'-P	-7.25	1.52	1.61
85	AA	2181	G	O4'-C1'	-7.25	1.32	1.41
34	BA	1840	C	C4'-C3'	-7.25	1.45	1.53
35	BB	669	A	N7-C5	-7.25	1.34	1.39
35	BB	1096	G	C5-C4	-7.25	1.33	1.38
37	BD	60	C	C2-N3	-7.25	1.29	1.35
85	AA	736	U	C3'-C2'	-7.25	1.44	1.52
85	AA	2192	A	C5-C4	-7.25	1.33	1.38
34	BA	363	G	O3'-P	-7.25	1.52	1.61
35	BB	675	U	C3'-C2'	-7.25	1.44	1.52
40	BG	31	G	C1'-N9	-7.25	1.36	1.46
40	BG	109	C	C2'-C1'	-7.25	1.45	1.53
85	AA	104	C	C2-N3	-7.25	1.29	1.35
85	AA	1267	A	C2'-C1'	-7.25	1.45	1.53
85	AA	2121	G	C6-N1	-7.25	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	183	G	C2'-C1'	-7.25	1.45	1.53
35	BB	808	U	P-O5'	-7.25	1.52	1.59
35	BB	1018	U	C3'-C2'	-7.25	1.44	1.52
35	BB	1317	U	P-O5'	-7.25	1.52	1.59
34	BA	134	U	C2'-C1'	-7.25	1.45	1.53
34	BA	308	C	C3'-C2'	-7.25	1.44	1.52
34	BA	394	A	O3'-P	-7.25	1.52	1.61
35	BB	816	U	C4'-C3'	-7.25	1.45	1.53
36	BC	119	G	C1'-N9	-7.25	1.36	1.46
36	BC	140	U	C4-C5	-7.25	1.37	1.43
38	BE	108	U	P-O5'	-7.25	1.52	1.59
34	BA	371	U	C3'-C2'	-7.25	1.44	1.52
34	BA	1454	G	C5-C6	-7.25	1.35	1.42
35	BB	845	C	C2'-C1'	-7.25	1.45	1.53
38	BE	28	C	C3'-C2'	-7.25	1.44	1.52
38	BE	36	U	C2-N3	-7.25	1.32	1.37
85	AA	28	A	N9-C4	-7.25	1.33	1.37
85	AA	197	C	C5'-C4'	7.25	1.60	1.51
85	AA	885	A	C2'-C1'	-7.25	1.45	1.53
85	AA	2072	G	O3'-P	-7.25	1.52	1.61
35	BB	806	U	C2-N3	-7.25	1.32	1.37
36	BC	6	G	C5-C4	-7.25	1.33	1.38
41	BH	31	A	C2'-C1'	-7.25	1.45	1.53
85	AA	518	A	C5'-C4'	7.25	1.60	1.51
85	AA	1125	G	C5-C4	-7.25	1.33	1.38
85	AA	1203	G	O3'-P	-7.25	1.52	1.61
85	AA	1487	G	C5-C4	-7.25	1.33	1.38
85	AA	1687	U	P-O5'	-7.25	1.52	1.59
34	BA	15	G	C5-C6	-7.24	1.35	1.42
34	BA	305	C	O3'-P	-7.24	1.52	1.61
35	BB	90	G	N9-C4	-7.24	1.32	1.38
35	BB	1024	G	N9-C4	7.24	1.43	1.38
36	BC	37	U	C1'-N1	-7.24	1.36	1.46
38	BE	129	G	N9-C4	-7.24	1.32	1.38
38	BE	149	A	N7-C5	-7.24	1.34	1.39
40	BG	118	U	C2'-C1'	-7.24	1.45	1.53
85	AA	819	G	C1'-N9	-7.24	1.36	1.46
86	AB	48	C	O3'-P	-7.24	1.52	1.61
34	BA	1210	A	O3'-P	-7.24	1.52	1.61
35	BB	432	C	C4'-O4'	-7.24	1.36	1.45
35	BB	1440	A	C5-C4	-7.24	1.33	1.38
37	BD	76	U	C2'-C1'	-7.24	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	128	U	C2'-C1'	-7.24	1.45	1.53
85	AA	307	G	N9-C4	-7.24	1.32	1.38
85	AA	381	A	O3'-P	-7.24	1.52	1.61
85	AA	660	G	C2'-C1'	-7.24	1.45	1.53
34	BA	269	G	C2-N3	-7.24	1.26	1.32
34	BA	1566	G	N7-C5	-7.24	1.34	1.39
35	BB	399	A	N7-C5	-7.24	1.34	1.39
36	BC	52	A	C2'-C1'	-7.24	1.45	1.53
36	BC	63	G	C1'-N9	-7.24	1.36	1.46
41	BH	21	G	N9-C4	-7.24	1.32	1.38
85	AA	752	C	O3'-P	-7.24	1.52	1.61
85	AA	886	A	C4'-C3'	-7.24	1.45	1.53
85	AA	892	C	P-O5'	-7.24	1.52	1.59
85	AA	1222	A	O3'-P	-7.24	1.52	1.61
85	AA	1484	G	C5-C4	-7.24	1.33	1.38
85	AA	1841	G	O3'-P	-7.24	1.52	1.61
34	BA	805	A	N3-C4	-7.24	1.30	1.34
34	BA	854	A	P-O5'	-7.24	1.52	1.59
34	BA	997	U	N3-C4	-7.24	1.31	1.38
35	BB	1102	U	C2'-C1'	-7.24	1.45	1.53
85	AA	256	A	P-O5'	-7.24	1.52	1.59
34	BA	1202	G	C5-C4	-7.24	1.33	1.38
40	BG	36	G	C6-N1	-7.24	1.34	1.39
85	AA	1638	C	O3'-P	-7.24	1.52	1.61
34	BA	3	G	O3'-P	-7.24	1.52	1.61
34	BA	351	A	C4'-C3'	-7.24	1.45	1.53
34	BA	1099	U	C4'-C3'	-7.24	1.45	1.53
34	BA	1105	A	P-O5'	-7.24	1.52	1.59
34	BA	1281	U	N3-C4	-7.24	1.31	1.38
34	BA	1722	U	C2'-C1'	-7.24	1.45	1.53
35	BB	68	G	N3-C4	-7.24	1.30	1.35
85	AA	710	A	N3-C4	-7.24	1.30	1.34
85	AA	2193	A	C4'-C3'	-7.24	1.45	1.53
34	BA	254	U	N3-C4	-7.23	1.31	1.38
34	BA	796	G	C8-N7	-7.23	1.26	1.30
35	BB	868	C	P-O5'	-7.23	1.52	1.59
35	BB	1111	C	P-O5'	-7.23	1.52	1.59
35	BB	1388	A	C1'-N9	-7.23	1.36	1.46
40	BG	45	G	C1'-N9	-7.23	1.36	1.46
34	BA	463	A	C1'-N9	-7.23	1.36	1.46
35	BB	558	U	C2'-C1'	-7.23	1.45	1.53
35	BB	1286	G	N9-C4	-7.23	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	132	C	O3'-P	-7.23	1.52	1.61
85	AA	480	U	P-O5'	-7.23	1.52	1.59
85	AA	1491	G	O3'-P	-7.23	1.52	1.61
34	BA	312	U	C2-N3	-7.23	1.32	1.37
34	BA	450	G	C4'-C3'	-7.23	1.45	1.53
34	BA	953	G	C6-N1	-7.23	1.34	1.39
35	BB	1355	C	C4'-C3'	-7.23	1.45	1.53
36	BC	129	C	O3'-P	-7.23	1.52	1.61
38	BE	122	G	O3'-P	-7.23	1.52	1.61
85	AA	1156	A	C2'-C1'	-7.23	1.45	1.53
85	AA	1180	C	C4'-C3'	-7.23	1.45	1.53
40	BG	100	G	P-O5'	-7.23	1.52	1.59
41	BH	113	G	C5-C4	-7.23	1.33	1.38
85	AA	882	C	C2'-C1'	-7.23	1.45	1.53
85	AA	2150	G	P-O5'	-7.23	1.52	1.59
35	BB	877	A	O3'-P	-7.23	1.52	1.61
35	BB	1226	G	C5-C4	-7.23	1.33	1.38
85	AA	165	C	C4'-C3'	-7.23	1.45	1.53
85	AA	1168	C	O3'-P	-7.23	1.52	1.61
34	BA	376	U	C3'-C2'	-7.23	1.44	1.52
34	BA	763	U	N1-C6	-7.23	1.31	1.38
34	BA	1149	C	C4-N4	-7.23	1.27	1.33
34	BA	1320	A	C3'-C2'	-7.23	1.44	1.52
34	BA	1385	U	P-O5'	-7.23	1.52	1.59
34	BA	659	U	O3'-P	-7.22	1.52	1.61
34	BA	1181	G	C3'-C2'	-7.22	1.44	1.52
35	BB	553	U	O3'-P	-7.22	1.52	1.61
35	BB	674	C	P-O5'	-7.22	1.52	1.59
35	BB	1004	A	O3'-P	-7.22	1.52	1.61
35	BB	1054	G	P-O5'	-7.22	1.52	1.59
37	BD	96	C	C3'-C2'	-7.22	1.44	1.52
85	AA	35	U	C3'-C2'	-7.22	1.44	1.52
85	AA	2051	G	C2'-C1'	-7.22	1.45	1.53
34	BA	264	A	O3'-P	-7.22	1.52	1.61
34	BA	1225	A	N3-C4	-7.22	1.30	1.34
34	BA	1271	C	C3'-C2'	-7.22	1.44	1.52
34	BA	1646	U	C2'-C1'	-7.22	1.45	1.53
35	BB	518	G	O3'-P	-7.22	1.52	1.61
35	BB	698	C	P-O5'	-7.22	1.52	1.59
35	BB	1506	C	C5'-C4'	7.22	1.60	1.51
36	BC	80	A	N9-C4	-7.22	1.33	1.37
85	AA	95	U	O3'-P	-7.22	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	379	U	O3'-P	-7.22	1.52	1.61
85	AA	2073	U	C2-N3	-7.22	1.32	1.37
34	BA	18	G	C6-N1	7.22	1.44	1.39
34	BA	735	A	C1'-N9	-7.22	1.36	1.46
35	BB	121	A	O3'-P	-7.22	1.52	1.61
35	BB	1093	C	C2'-C1'	-7.22	1.45	1.53
35	BB	1259	A	N3-C4	-7.22	1.30	1.34
65	Bf	376	ARG	CD-NE	7.22	1.58	1.46
85	AA	365	G	C6-N1	-7.22	1.34	1.39
85	AA	2174	G	C6-N1	-7.22	1.34	1.39
34	BA	1412	G	N7-C5	-7.22	1.34	1.39
35	BB	831	C	P-O5'	-7.22	1.52	1.59
35	BB	1354	C	C5'-C4'	-7.22	1.42	1.51
35	BB	1462	G	C4'-C3'	7.22	1.61	1.53
41	BH	51	C	C2'-C1'	-7.22	1.45	1.53
85	AA	490	A	C4'-C3'	-7.22	1.45	1.53
85	AA	2035	C	O3'-P	-7.22	1.52	1.61
85	AA	2067	A	C2'-C1'	-7.22	1.45	1.53
85	AA	2112	G	P-O5'	-7.22	1.52	1.59
85	AA	2218	G	N9-C8	-7.22	1.32	1.37
34	BA	421	G	C3'-C2'	-7.22	1.44	1.52
35	BB	385	C	O3'-P	-7.22	1.52	1.61
35	BB	388	C	O3'-P	-7.22	1.52	1.61
41	BH	44	A	C5-C4	-7.22	1.33	1.38
85	AA	1371	C	P-O5'	-7.22	1.52	1.59
85	AA	1712	A	C2'-C1'	-7.22	1.45	1.53
85	AA	2083	G	C4'-C3'	-7.22	1.45	1.53
34	BA	239	C	N3-C4	7.22	1.39	1.33
34	BA	1203	G	C6-N1	-7.22	1.34	1.39
35	BB	34	G	N7-C5	-7.22	1.34	1.39
35	BB	66	G	C2'-C1'	-7.22	1.45	1.53
37	BD	66	G	C1'-N9	-7.22	1.36	1.46
38	BE	32	U	C2-N3	-7.22	1.32	1.37
39	BF	37	C	C2'-C1'	-7.22	1.45	1.53
40	BG	104	A	N7-C5	-7.22	1.34	1.39
34	BA	39	C	O3'-P	-7.21	1.52	1.61
34	BA	527	C	C2'-C1'	-7.21	1.45	1.53
34	BA	1529	G	C2-N2	-7.21	1.27	1.34
34	BA	1477	C	O3'-P	-7.21	1.52	1.61
34	BA	1828	A	N7-C5	-7.21	1.34	1.39
35	BB	1286	G	C1'-N9	-7.21	1.36	1.46
35	BB	1294	C	C5'-C4'	-7.21	1.42	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	104	A	C6-N1	-7.21	1.30	1.35
85	AA	1872	G	C2'-C1'	-7.21	1.45	1.53
34	BA	128	C	C2-N3	-7.21	1.29	1.35
34	BA	403	A	O3'-P	-7.21	1.52	1.61
34	BA	891	C	C2-N3	-7.21	1.29	1.35
34	BA	1561	C	C1'-N1	-7.21	1.36	1.46
34	BA	1683	C	N1-C6	-7.21	1.32	1.37
35	BB	80	C	C3'-C2'	-7.21	1.44	1.52
35	BB	487	A	O3'-P	-7.21	1.52	1.61
36	BC	65	G	C2'-C1'	-7.21	1.45	1.53
36	BC	78	G	C5-C4	-7.21	1.33	1.38
40	BG	95	U	N1-C6	-7.21	1.31	1.38
85	AA	1467	U	P-O5'	-7.21	1.52	1.59
85	AA	1470	A	C2'-C1'	-7.21	1.45	1.53
85	AA	2127	G	O3'-P	-7.21	1.52	1.61
34	BA	1562	G	C1'-N9	-7.21	1.36	1.46
35	BB	508	U	C3'-C2'	-7.21	1.44	1.52
35	BB	622	G	C6-N1	-7.21	1.34	1.39
35	BB	626	C	O3'-P	-7.21	1.52	1.61
35	BB	1048	A	C4'-C3'	-7.21	1.45	1.53
35	BB	1194	A	C1'-N9	-7.21	1.36	1.46
85	AA	1269	A	P-O5'	-7.21	1.52	1.59
34	BA	1172	C	O3'-P	-7.21	1.52	1.61
34	BA	1435	A	P-O5'	-7.21	1.52	1.59
34	BA	1467	U	C2-N3	-7.21	1.32	1.37
34	BA	1697	U	O4'-C1'	-7.21	1.32	1.41
85	AA	931	G	C1'-N9	-7.21	1.36	1.46
85	AA	2145	G	N9-C8	-7.21	1.32	1.37
34	BA	116	G	P-O5'	-7.21	1.52	1.59
34	BA	373	G	C2-N2	-7.21	1.27	1.34
34	BA	548	G	P-O5'	-7.21	1.52	1.59
34	BA	718	U	N1-C2	-7.21	1.32	1.38
34	BA	1346	U	P-O5'	-7.21	1.52	1.59
34	BA	1827	C	P-O5'	-7.21	1.52	1.59
35	BB	679	G	C1'-N9	-7.21	1.36	1.46
35	BB	692	G	P-O5'	-7.21	1.52	1.59
37	BD	49	A	O3'-P	-7.21	1.52	1.61
85	AA	1159	C	P-O5'	-7.21	1.52	1.59
85	AA	1680	U	C2-N3	-7.21	1.32	1.37
85	AA	2075	C	C1'-N1	-7.21	1.36	1.46
34	BA	671	C	P-O5'	-7.21	1.52	1.59
34	BA	1233	U	O3'-P	-7.21	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1433	U	C3'-C2'	-7.21	1.44	1.52
34	BA	1514	A	O4'-C1'	-7.21	1.32	1.41
40	BG	142	A	C3'-C2'	-7.21	1.44	1.52
41	BH	125	U	C2'-C1'	-7.21	1.45	1.53
85	AA	119	G	C1'-N9	-7.21	1.36	1.46
85	AA	2123	U	C2'-C1'	-7.21	1.45	1.53
34	BA	1830	A	C2'-C1'	-7.20	1.45	1.53
35	BB	15	C	O3'-P	-7.20	1.52	1.61
35	BB	458	U	C2-N3	-7.20	1.32	1.37
35	BB	628	A	C1'-N9	-7.20	1.36	1.46
85	AA	155	U	C2'-C1'	-7.20	1.45	1.53
85	AA	449	G	C1'-N9	-7.20	1.36	1.46
35	BB	73	G	O4'-C1'	-7.20	1.32	1.41
35	BB	1252	G	C1'-N9	-7.20	1.36	1.46
40	BG	91	U	C3'-C2'	-7.20	1.44	1.52
34	BA	1231	C	C2'-C1'	-7.20	1.45	1.53
35	BB	377	A	C4'-C3'	-7.20	1.45	1.53
35	BB	1050	A	C5-C4	-7.20	1.33	1.38
35	BB	1186	A	C1'-N9	-7.20	1.36	1.46
35	BB	1314	G	P-O5'	-7.20	1.52	1.59
40	BG	113	G	O3'-P	-7.20	1.52	1.61
85	AA	506	G	O3'-P	-7.20	1.52	1.61
85	AA	1456	A	C2'-C1'	-7.20	1.45	1.53
85	AA	1848	G	N9-C4	-7.20	1.32	1.38
34	BA	83	G	N9-C4	-7.20	1.32	1.38
34	BA	275	C	P-O5'	-7.20	1.52	1.59
34	BA	333	A	C2'-C1'	-7.20	1.45	1.53
34	BA	1323	G	C2-N2	-7.20	1.27	1.34
35	BB	1171	U	C2-N3	-7.20	1.32	1.37
37	BD	71	G	O3'-P	-7.20	1.52	1.61
37	BD	105	G	C5-C4	-7.20	1.33	1.38
40	BG	71	C	O3'-P	-7.20	1.52	1.61
85	AA	719	C	C2'-C1'	-7.20	1.45	1.53
34	BA	376	U	P-O5'	-7.20	1.52	1.59
35	BB	1033	U	O3'-P	-7.20	1.52	1.61
36	BC	116	C	C3'-C2'	-7.20	1.44	1.52
85	AA	959	C	C4'-C3'	-7.20	1.45	1.53
85	AA	1000	U	O3'-P	-7.20	1.52	1.61
85	AA	1470	A	C1'-N9	-7.20	1.36	1.46
34	BA	875	G	C2-N2	-7.20	1.27	1.34
34	BA	982	A	P-O5'	-7.20	1.52	1.59
34	BA	1417	C	C2-N3	-7.20	1.29	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	625	A	C2'-C1'	-7.20	1.45	1.53
35	BB	1429	A	P-O5'	-7.20	1.52	1.59
35	BB	1437	U	O3'-P	-7.20	1.52	1.61
41	BH	6	U	O3'-P	-7.20	1.52	1.61
85	AA	93	G	N7-C5	-7.20	1.34	1.39
85	AA	413	G	N1-C2	-7.20	1.31	1.37
85	AA	613	G	N9-C4	-7.20	1.32	1.38
85	AA	931	G	P-O5'	-7.20	1.52	1.59
85	AA	1227	A	N3-C4	-7.20	1.30	1.34
85	AA	1509	A	N7-C5	-7.20	1.34	1.39
85	AA	1893	G	N9-C4	-7.20	1.32	1.38
34	BA	724	A	C2'-C1'	-7.19	1.45	1.53
34	BA	1272	U	C2'-C1'	-7.19	1.45	1.53
35	BB	1241	U	P-O5'	-7.19	1.52	1.59
35	BB	1246	C	C2-N3	-7.19	1.29	1.35
35	BB	1302	C	P-O5'	-7.19	1.52	1.59
35	BB	1368	A	C5-C4	-7.19	1.33	1.38
85	AA	1268	C	P-O5'	-7.19	1.52	1.59
34	BA	591	G	N9-C4	-7.19	1.32	1.38
34	BA	816	G	C6-N1	-7.19	1.34	1.39
34	BA	1235	C	P-O5'	-7.19	1.52	1.59
34	BA	1419	A	O3'-P	-7.19	1.52	1.61
34	BA	1696	G	O3'-P	-7.19	1.52	1.61
35	BB	642	G	C3'-C2'	-7.19	1.44	1.52
35	BB	794	G	C3'-C2'	-7.19	1.44	1.52
35	BB	874	G	P-O5'	-7.19	1.52	1.59
40	BG	2	U	O3'-P	-7.19	1.52	1.61
40	BG	172	C	C2'-C1'	-7.19	1.45	1.53
85	AA	1126	G	N1-C2	-7.19	1.31	1.37
34	BA	390	A	C1'-N9	-7.19	1.36	1.46
35	BB	673	C	O3'-P	-7.19	1.52	1.61
35	BB	1006	C	P-O5'	-7.19	1.52	1.59
35	BB	1166	A	O3'-P	-7.19	1.52	1.61
40	BG	118	U	C1'-N1	-7.19	1.36	1.46
41	BH	17	A	N9-C4	-7.19	1.33	1.37
85	AA	20	G	C2-N2	-7.19	1.27	1.34
85	AA	31	C	C3'-C2'	-7.19	1.44	1.52
85	AA	529	G	N7-C5	-7.19	1.34	1.39
85	AA	867	G	N9-C8	-7.19	1.32	1.37
34	BA	336	A	N9-C4	-7.19	1.33	1.37
35	BB	1383	C	O3'-P	-7.19	1.52	1.61
35	BB	1470	G	C2'-C1'	-7.19	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	152	G	N1-C2	-7.19	1.31	1.37
34	BA	1299	G	P-O5'	-7.19	1.52	1.59
34	BA	1412	G	C1'-N9	-7.19	1.36	1.46
35	BB	63	A	N7-C5	-7.19	1.34	1.39
35	BB	1075	A	N3-C4	-7.19	1.30	1.34
35	BB	1407	U	P-O5'	-7.19	1.52	1.59
38	BE	194	A	C5-C6	-7.19	1.34	1.41
85	AA	409	C	C3'-C2'	-7.19	1.44	1.52
85	AA	1199	C	C5'-C4'	-7.19	1.42	1.51
34	BA	196	A	C4'-C3'	-7.19	1.45	1.53
34	BA	1180	A	O3'-P	-7.19	1.52	1.61
34	BA	1421	A	C1'-N9	-7.19	1.36	1.46
34	BA	1682	A	C4'-C3'	-7.19	1.45	1.53
40	BG	101	G	C6-N1	-7.19	1.34	1.39
40	BG	149	U	C2'-C1'	-7.19	1.45	1.53
34	BA	125	G	C1'-N9	-7.18	1.36	1.46
34	BA	1061	A	P-O5'	-7.18	1.52	1.59
34	BA	1157	A	C2'-C1'	-7.18	1.45	1.53
35	BB	576	A	C5-C4	-7.18	1.33	1.38
35	BB	678	U	C4'-O4'	-7.18	1.36	1.45
40	BG	33	G	N9-C8	-7.18	1.32	1.37
40	BG	40	G	C2'-C1'	-7.18	1.45	1.53
85	AA	654	A	N9-C4	-7.18	1.33	1.37
34	BA	543	A	C4'-C3'	-7.18	1.45	1.53
34	BA	1725	U	O3'-P	-7.18	1.52	1.61
35	BB	1172	U	P-O5'	-7.18	1.52	1.59
35	BB	1248	A	C2'-C1'	-7.18	1.45	1.53
41	BH	33	G	C3'-C2'	-7.18	1.44	1.52
85	AA	684	G	C5-C4	-7.18	1.33	1.38
85	AA	971	U	C2'-C1'	-7.18	1.45	1.53
85	AA	2139	G	C8-N7	-7.18	1.26	1.30
35	BB	1019	C	C4'-C3'	-7.18	1.45	1.53
35	BB	1064	U	O3'-P	-7.18	1.52	1.61
85	AA	210	G	N9-C4	-7.18	1.32	1.38
34	BA	101	G	C2'-C1'	-7.18	1.45	1.53
34	BA	1213	A	C2'-C1'	-7.18	1.45	1.53
34	BA	1776	G	O3'-P	-7.18	1.52	1.61
35	BB	567	G	O3'-P	-7.18	1.52	1.61
35	BB	1003	G	P-O5'	-7.18	1.52	1.59
38	BE	50	G	C6-N1	-7.18	1.34	1.39
40	BG	19	C	P-O5'	-7.18	1.52	1.59
85	AA	339	A	N9-C4	-7.18	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	817	G	N7-C5	-7.18	1.34	1.39
34	BA	1796	A	C5-C4	-7.18	1.33	1.38
34	BA	525	A	C2'-C1'	-7.18	1.45	1.53
34	BA	1205	A	N9-C4	-7.18	1.33	1.37
34	BA	1239	G	N1-C2	-7.18	1.32	1.37
38	BE	124	G	C2-N2	-7.18	1.27	1.34
40	BG	130	G	C2'-C1'	-7.18	1.45	1.53
41	BH	24	U	P-O5'	-7.18	1.52	1.59
85	AA	97	A	P-O5'	-7.18	1.52	1.59
85	AA	329	G	N9-C4	-7.18	1.32	1.38
85	AA	525	C	O3'-P	-7.18	1.52	1.61
85	AA	918	U	C4'-C3'	-7.18	1.45	1.53
85	AA	1460	G	N9-C8	-7.18	1.32	1.37
34	BA	1163	G	N7-C5	-7.17	1.34	1.39
34	BA	1403	G	N9-C4	-7.17	1.32	1.38
34	BA	1527	G	N9-C4	-7.17	1.32	1.38
35	BB	663	G	C6-N1	-7.17	1.34	1.39
35	BB	1026	G	N7-C5	-7.17	1.34	1.39
37	BD	38	U	P-O5'	-7.17	1.52	1.59
85	AA	49	C	O3'-P	-7.17	1.52	1.61
85	AA	755	G	N9-C4	7.17	1.43	1.38
34	BA	4	A	C3'-C2'	-7.17	1.44	1.52
34	BA	1118	C	C2'-C1'	-7.17	1.45	1.53
85	AA	2133	A	C8-N7	-7.17	1.26	1.31
34	BA	273	G	O3'-P	-7.17	1.52	1.61
34	BA	1485	U	N3-C4	-7.17	1.31	1.38
34	BA	1596	C	P-O5'	-7.17	1.52	1.59
34	BA	1650	G	P-O5'	-7.17	1.52	1.59
35	BB	128	C	P-O5'	-7.17	1.52	1.59
35	BB	372	U	O3'-P	-7.17	1.52	1.61
35	BB	558	U	P-O5'	-7.17	1.52	1.59
38	BE	203	C	C4-C5	-7.17	1.37	1.43
85	AA	108	C	O3'-P	-7.17	1.52	1.61
85	AA	1200	A	C1'-N9	-7.17	1.36	1.46
85	AA	1527	G	O3'-P	-7.17	1.52	1.61
85	AA	1593	C	O3'-P	-7.17	1.52	1.61
85	AA	2229	G	O3'-P	-7.17	1.52	1.61
34	BA	268	U	C1'-N1	-7.17	1.36	1.46
35	BB	1196	A	C2'-C1'	-7.17	1.45	1.53
37	BD	88	U	O3'-P	-7.17	1.52	1.61
34	BA	1010	C	C2-N3	-7.17	1.30	1.35
34	BA	1074	C	C3'-C2'	-7.17	1.44	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	87	G	N9-C4	-7.17	1.32	1.38
35	BB	466	A	O3'-P	-7.17	1.52	1.61
37	BD	62	A	P-O5'	-7.17	1.52	1.59
37	BD	107	G	C2-N2	-7.17	1.27	1.34
85	AA	210	G	C4'-C3'	-7.17	1.45	1.53
34	BA	2	A	N7-C5	-7.17	1.34	1.39
34	BA	934	G	N9-C4	-7.17	1.32	1.38
34	BA	1264	U	O3'-P	-7.17	1.52	1.61
35	BB	1082	A	N7-C5	-7.17	1.34	1.39
35	BB	1262	A	N7-C5	-7.17	1.34	1.39
36	BC	12	A	C1'-N9	-7.17	1.36	1.46
85	AA	925	G	C2'-C1'	-7.17	1.45	1.53
85	AA	1265	C	C2'-C1'	-7.17	1.45	1.53
34	BA	728	A	N7-C5	-7.17	1.34	1.39
34	BA	1161	G	C4'-C3'	7.17	1.61	1.53
34	BA	1652	G	C5-C4	-7.17	1.33	1.38
35	BB	1378	U	N3-C4	-7.17	1.32	1.38
85	AA	178	U	N1-C2	-7.17	1.32	1.38
85	AA	204	U	O3'-P	-7.17	1.52	1.61
85	AA	638	G	P-O5'	-7.17	1.52	1.59
34	BA	321	G	P-O5'	-7.16	1.52	1.59
34	BA	347	A	O3'-P	-7.16	1.52	1.61
34	BA	606	G	C3'-O3'	-7.16	1.32	1.42
34	BA	625	U	P-O5'	-7.16	1.52	1.59
34	BA	989	C	O3'-P	-7.16	1.52	1.61
34	BA	1026	C	C2-N3	-7.16	1.30	1.35
34	BA	1686	G	C4'-C3'	-7.16	1.45	1.53
34	BA	1740	U	P-O5'	-7.16	1.52	1.59
35	BB	137	A	C6-N6	-7.16	1.28	1.33
35	BB	591	A	O3'-P	-7.16	1.52	1.61
35	BB	1243	A	C2'-C1'	-7.16	1.45	1.53
38	BE	4	A	C2'-C1'	-7.16	1.45	1.53
85	AA	1472	G	N9-C4	-7.16	1.32	1.38
85	AA	1502	A	C3'-C2'	-7.16	1.44	1.52
85	AA	2027	U	P-O5'	-7.16	1.52	1.59
34	BA	218	G	C1'-N9	-7.16	1.36	1.46
34	BA	429	G	O3'-P	-7.16	1.52	1.61
34	BA	1701	U	C2'-C1'	-7.16	1.45	1.53
34	BA	786	U	O3'-P	-7.16	1.52	1.61
34	BA	1103	G	N1-C2	-7.16	1.32	1.37
34	BA	1251	A	P-O5'	-7.16	1.52	1.59
34	BA	1593	U	C2-N3	-7.16	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1272	G	N7-C5	-7.16	1.34	1.39
39	BF	23	G	N7-C5	-7.16	1.34	1.39
40	BG	80	G	C6-N1	-7.16	1.34	1.39
41	BH	29	G	C5-C6	-7.16	1.35	1.42
85	AA	384	C	C1'-N1	-7.16	1.36	1.46
85	AA	1680	U	P-O5'	-7.16	1.52	1.59
85	AA	1708	A	O3'-P	-7.16	1.52	1.61
34	BA	452	A	C5-C4	-7.16	1.33	1.38
35	BB	4	C	C3'-C2'	7.16	1.60	1.52
35	BB	1299	G	C3'-C2'	-7.16	1.44	1.52
35	BB	1305	A	O3'-P	-7.16	1.52	1.61
38	BE	13	A	N7-C5	-7.16	1.34	1.39
40	BG	35	G	O3'-P	-7.16	1.52	1.61
41	BH	39	G	C5-C4	-7.16	1.33	1.38
41	BH	114	G	C3'-C2'	-7.16	1.44	1.52
85	AA	644	A	N7-C5	-7.16	1.34	1.39
85	AA	1929	G	O3'-P	-7.16	1.52	1.61
34	BA	29	U	C3'-C2'	-7.16	1.44	1.52
34	BA	1845	G	C4'-C3'	-7.16	1.45	1.53
35	BB	94	A	N3-C4	-7.16	1.30	1.34
40	BG	162	A	C5-C4	-7.16	1.33	1.38
85	AA	1460	G	N3-C4	-7.16	1.30	1.35
34	BA	875	G	N9-C8	-7.16	1.32	1.37
34	BA	986	G	C3'-C2'	-7.16	1.44	1.52
34	BA	1493	U	C3'-O3'	7.16	1.52	1.42
34	BA	1806	A	N9-C8	-7.16	1.32	1.37
35	BB	1262	A	C2'-C1'	-7.16	1.45	1.53
41	BH	116	A	O3'-P	-7.16	1.52	1.61
85	AA	88	G	P-O5'	-7.16	1.52	1.59
85	AA	177	A	C3'-C2'	-7.16	1.44	1.52
85	AA	505	U	O4'-C1'	-7.16	1.32	1.41
85	AA	924	A	C2'-C1'	-7.16	1.45	1.53
85	AA	2229	G	N7-C5	-7.16	1.34	1.39
34	BA	735	A	N3-C4	-7.15	1.30	1.34
34	BA	1513	G	N9-C4	7.15	1.43	1.38
35	BB	49	A	N3-C4	-7.15	1.30	1.34
35	BB	976	U	C3'-C2'	-7.15	1.44	1.52
35	BB	1110	G	N7-C5	-7.15	1.34	1.39
35	BB	1294	C	O4'-C1'	-7.15	1.32	1.41
35	BB	1319	U	C5'-C4'	7.15	1.59	1.51
36	BC	91	G	C2'-C1'	-7.15	1.45	1.53
85	AA	302	C	C4'-C3'	7.15	1.61	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	975	G	C3'-C2'	-7.15	1.44	1.52
34	BA	105	U	C2-N3	-7.15	1.32	1.37
34	BA	565	U	C2-N3	-7.15	1.32	1.37
34	BA	579	U	C5'-C4'	-7.15	1.42	1.51
35	BB	379	U	C2-N3	-7.15	1.32	1.37
85	AA	7	G	C2'-C1'	-7.15	1.45	1.53
85	AA	1291	A	P-O5'	-7.15	1.52	1.59
34	BA	1679	C	P-O5'	-7.15	1.52	1.59
35	BB	121	A	C1'-N9	-7.15	1.36	1.46
35	BB	661	G	N9-C4	-7.15	1.32	1.38
35	BB	1446	C	C2-N3	-7.15	1.30	1.35
38	BE	2	G	C1'-N9	-7.15	1.36	1.46
38	BE	2	G	C6-N1	-7.15	1.34	1.39
40	BG	129	G	C6-N1	-7.15	1.34	1.39
41	BH	121	A	N7-C5	-7.15	1.34	1.39
85	AA	627	A	C1'-N9	-7.15	1.36	1.46
85	AA	1723	U	O3'-P	-7.15	1.52	1.61
34	BA	51	C	N3-C4	-7.15	1.28	1.33
35	BB	363	A	C2'-C1'	-7.15	1.45	1.53
35	BB	1137	G	N9-C4	-7.15	1.32	1.38
34	BA	46	C	O3'-P	-7.15	1.52	1.61
34	BA	103	G	C3'-C2'	-7.15	1.44	1.52
34	BA	337	C	C2-N3	-7.15	1.30	1.35
35	BB	119	G	N1-C2	-7.15	1.32	1.37
35	BB	1081	U	N3-C4	-7.15	1.32	1.38
35	BB	1321	G	P-O5'	-7.15	1.52	1.59
38	BE	62	C	C3'-C2'	-7.15	1.44	1.52
40	BG	65	C	C2-N3	-7.15	1.30	1.35
85	AA	4	C	C3'-C2'	-7.15	1.44	1.52
85	AA	249	C	C3'-C2'	-7.15	1.44	1.52
85	AA	309	G	N1-C2	-7.15	1.32	1.37
85	AA	1345	C	P-O5'	-7.15	1.52	1.59
85	AA	1466	U	C2-N3	-7.15	1.32	1.37
34	BA	803	U	O3'-P	-7.15	1.52	1.61
34	BA	1194	G	N9-C4	-7.15	1.32	1.38
35	BB	432	C	O3'-P	-7.15	1.52	1.61
38	BE	38	C	C2'-C1'	-7.15	1.45	1.53
34	BA	909	G	O4'-C1'	-7.14	1.32	1.41
34	BA	1089	U	O3'-P	-7.14	1.52	1.61
35	BB	409	U	O3'-P	-7.14	1.52	1.61
35	BB	981	A	N3-C4	-7.14	1.30	1.34
35	BB	1229	A	P-O5'	-7.14	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1515	C	C3'-C2'	-7.14	1.44	1.52
85	AA	63	G	C2-N2	-7.14	1.27	1.34
34	BA	445	C	O3'-P	-7.14	1.52	1.61
34	BA	664	C	O3'-P	-7.14	1.52	1.61
34	BA	687	G	C1'-N9	-7.14	1.36	1.46
34	BA	703	U	C3'-C2'	-7.14	1.44	1.52
35	BB	1127	A	C1'-N9	-7.14	1.36	1.46
34	BA	20	A	C5-C4	-7.14	1.33	1.38
35	BB	26	C	P-O5'	-7.14	1.52	1.59
36	BC	63	G	N7-C5	-7.14	1.34	1.39
39	BF	56	C	C1'-N1	7.14	1.59	1.48
85	AA	548	G	C3'-C2'	-7.14	1.44	1.52
85	AA	747	U	C2'-C1'	-7.14	1.45	1.53
85	AA	909	C	O3'-P	-7.14	1.52	1.61
85	AA	2191	C	C2'-C1'	-7.14	1.45	1.53
34	BA	150	C	O3'-P	-7.14	1.52	1.61
34	BA	435	U	O3'-P	-7.14	1.52	1.61
34	BA	774	A	C4'-C3'	-7.14	1.45	1.53
34	BA	1796	A	N9-C8	-7.14	1.32	1.37
41	BH	132	C	C2'-C1'	-7.14	1.45	1.53
85	AA	10	G	N1-C2	-7.14	1.32	1.37
85	AA	414	C	C3'-C2'	-7.14	1.44	1.52
85	AA	644	A	C8-N7	-7.14	1.26	1.31
85	AA	1554	C	O3'-P	-7.14	1.52	1.61
34	BA	83	G	P-O5'	-7.14	1.52	1.59
34	BA	130	U	C2-N3	-7.14	1.32	1.37
34	BA	200	C	C2'-C1'	-7.14	1.45	1.53
35	BB	838	G	P-O5'	-7.14	1.52	1.59
35	BB	1159	U	C4'-C3'	-7.14	1.45	1.53
36	BC	164	G	C3'-C2'	-7.14	1.44	1.52
40	BG	1	G	O3'-P	-7.14	1.52	1.61
85	AA	850	U	P-O5'	-7.14	1.52	1.59
85	AA	1832	G	C3'-C2'	-7.14	1.44	1.52
34	BA	82	A	O4'-C1'	-7.14	1.32	1.41
34	BA	421	G	C1'-N9	-7.14	1.36	1.46
34	BA	504	A	N9-C8	-7.14	1.32	1.37
34	BA	701	G	C2'-C1'	-7.14	1.45	1.53
34	BA	1592	U	C3'-C2'	-7.14	1.45	1.52
35	BB	62	C	C2'-C1'	-7.14	1.45	1.53
35	BB	1479	C	C3'-C2'	-7.14	1.44	1.52
36	BC	48	A	C1'-N9	-7.14	1.36	1.46
38	BE	23	G	N1-C2	-7.14	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	58	G	N7-C5	-7.14	1.34	1.39
85	AA	1458	G	C2'-C1'	-7.14	1.45	1.53
85	AA	2071	U	O3'-P	-7.14	1.52	1.61
34	BA	48	C	P-O5'	-7.13	1.52	1.59
34	BA	563	A	O3'-P	-7.13	1.52	1.61
34	BA	1242	A	P-O5'	-7.13	1.52	1.59
34	BA	1472	G	C1'-N9	-7.13	1.36	1.46
34	BA	1530	G	C6-N1	-7.13	1.34	1.39
34	BA	1721	U	N1-C2	-7.13	1.32	1.38
35	BB	102	G	N1-C2	-7.13	1.32	1.37
35	BB	993	A	C1'-N9	-7.13	1.36	1.46
35	BB	1127	A	P-O5'	-7.13	1.52	1.59
85	AA	335	G	N7-C5	-7.13	1.34	1.39
85	AA	383	C	N3-C4	-7.13	1.28	1.33
85	AA	820	G	O3'-P	-7.13	1.52	1.61
85	AA	1474	U	C2'-C1'	-7.13	1.45	1.53
34	BA	380	A	C1'-N9	-7.13	1.36	1.46
34	BA	417	A	C4'-O4'	-7.13	1.36	1.45
34	BA	1041	U	C1'-N1	-7.13	1.36	1.46
34	BA	1439	C	O3'-P	-7.13	1.52	1.61
35	BB	833	G	O3'-P	-7.13	1.52	1.61
36	BC	152	C	C4'-C3'	-7.13	1.45	1.53
38	BE	129	G	C8-N7	-7.13	1.26	1.30
40	BG	51	U	C2-N3	-7.13	1.32	1.37
85	AA	1469	G	C5-C4	-7.13	1.33	1.38
34	BA	156	U	O3'-P	-7.13	1.52	1.61
34	BA	244	A	O3'-P	-7.13	1.52	1.61
34	BA	329	G	C4'-O4'	-7.13	1.36	1.45
34	BA	1400	A	N9-C4	-7.13	1.33	1.37
34	BA	1416	C	C5'-C4'	7.13	1.59	1.51
39	BF	46	G	N1-C2	-7.13	1.32	1.37
85	AA	165	C	C2-N3	-7.13	1.30	1.35
85	AA	291	G	P-O5'	-7.13	1.52	1.59
85	AA	965	G	C3'-C2'	-7.13	1.45	1.52
85	AA	1294	U	O3'-P	-7.13	1.52	1.61
85	AA	1680	U	N3-C4	-7.13	1.32	1.38
34	BA	1070	G	C5-C6	-7.13	1.35	1.42
34	BA	1110	A	C3'-C2'	-7.13	1.45	1.52
35	BB	607	G	O4'-C1'	-7.13	1.32	1.41
35	BB	1014	U	C3'-C2'	-7.13	1.45	1.52
35	BB	1153	G	C5-C4	-7.13	1.33	1.38
36	BC	166	G	O3'-P	-7.13	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1486	G	N9-C4	-7.13	1.32	1.38
85	AA	1857	G	C2'-C1'	-7.13	1.45	1.53
34	BA	1033	G	O3'-P	-7.13	1.52	1.61
34	BA	1783	C	O3'-P	-7.13	1.52	1.61
35	BB	110	U	C2'-C1'	-7.13	1.45	1.53
35	BB	132	G	C8-N7	-7.13	1.26	1.30
36	BC	96	A	C5-C4	-7.13	1.33	1.38
36	BC	156	A	C8-N7	-7.13	1.26	1.31
40	BG	72	G	P-O5'	-7.13	1.52	1.59
40	BG	168	A	O3'-P	-7.13	1.52	1.61
85	AA	938	A	N9-C4	-7.13	1.33	1.37
34	BA	451	A	P-O5'	-7.13	1.52	1.59
35	BB	829	C	C2-N3	-7.13	1.30	1.35
85	AA	424	A	C5-C6	-7.13	1.34	1.41
85	AA	751	C	P-O5'	-7.13	1.52	1.59
85	AA	1847	U	O3'-P	-7.13	1.52	1.61
34	BA	89	G	N9-C8	-7.12	1.32	1.37
35	BB	428	G	C6-N1	-7.12	1.34	1.39
36	BC	78	G	C3'-C2'	-7.12	1.45	1.52
34	BA	516	U	C2-N3	-7.12	1.32	1.37
35	BB	658	G	C2'-C1'	-7.12	1.45	1.53
35	BB	1075	A	C1'-N9	-7.12	1.36	1.46
35	BB	1380	G	C1'-N9	-7.12	1.36	1.46
85	AA	129	U	C2-N3	-7.12	1.32	1.37
85	AA	803	C	P-O5'	-7.12	1.52	1.59
34	BA	430	A	O3'-P	-7.12	1.52	1.61
34	BA	617	G	C4'-O4'	-7.12	1.36	1.45
34	BA	930	A	C2'-C1'	-7.12	1.45	1.53
34	BA	1834	A	P-O5'	-7.12	1.52	1.59
35	BB	392	G	N1-C2	-7.12	1.32	1.37
35	BB	782	A	C3'-C2'	-7.12	1.45	1.52
35	BB	1335	G	P-O5'	-7.12	1.52	1.59
35	BB	1513	U	N3-C4	-7.12	1.32	1.38
35	BB	1548	C	C2'-C1'	-7.12	1.45	1.53
37	BD	91	U	C4'-C3'	-7.12	1.45	1.53
38	BE	124	G	C3'-C2'	-7.12	1.45	1.52
85	AA	67	C	O3'-P	-7.12	1.52	1.61
85	AA	685	U	P-O5'	-7.12	1.52	1.59
34	BA	202	A	N9-C4	-7.12	1.33	1.37
34	BA	1093	G	O3'-P	-7.12	1.52	1.61
40	BG	44	G	C2'-C1'	-7.12	1.45	1.53
85	AA	112	A	C4'-C3'	-7.12	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1462	A	P-O5'	-7.12	1.52	1.59
85	AA	2222	G	O3'-P	-7.12	1.52	1.61
34	BA	104	A	C3'-C2'	-7.12	1.45	1.52
34	BA	1010	C	P-O5'	-7.12	1.52	1.59
34	BA	1277	G	C3'-C2'	-7.12	1.45	1.52
34	BA	1323	G	C2'-C1'	-7.12	1.45	1.53
34	BA	1451	A	C5-C4	-7.12	1.33	1.38
35	BB	1164	U	C3'-C2'	-7.12	1.45	1.52
35	BB	1297	G	P-O5'	-7.12	1.52	1.59
35	BB	1431	G	C6-N1	-7.12	1.34	1.39
85	AA	767	A	C2'-C1'	-7.12	1.45	1.53
85	AA	1527	G	C2-N2	-7.12	1.27	1.34
85	AA	1709	U	P-O5'	-7.12	1.52	1.59
35	BB	736	G	C2'-C1'	-7.12	1.45	1.53
35	BB	1168	G	O4'-C1'	-7.12	1.32	1.41
85	AA	2075	C	C2-N3	-7.12	1.30	1.35
34	BA	866	C	N3-C4	7.12	1.39	1.33
34	BA	1835	A	C4'-C3'	-7.12	1.45	1.53
41	BH	45	G	O3'-P	-7.12	1.52	1.61
41	BH	123	G	C2-N2	-7.12	1.27	1.34
85	AA	404	A	C2'-C1'	-7.12	1.45	1.53
85	AA	771	A	P-O5'	-7.12	1.52	1.59
34	BA	124	G	C3'-C2'	-7.11	1.45	1.52
34	BA	720	A	C1'-N9	-7.11	1.36	1.46
34	BA	1573	C	P-O5'	-7.11	1.52	1.59
35	BB	477	U	O3'-P	-7.11	1.52	1.61
35	BB	1445	A	O4'-C1'	-7.11	1.32	1.41
40	BG	80	G	N7-C5	-7.11	1.34	1.39
85	AA	247	G	N9-C4	-7.11	1.32	1.38
34	BA	755	G	N3-C4	-7.11	1.30	1.35
35	BB	72	G	C2-N2	-7.11	1.27	1.34
35	BB	1313	C	C2-N3	-7.11	1.30	1.35
85	AA	262	G	C6-N1	-7.11	1.34	1.39
85	AA	439	U	O3'-P	-7.11	1.52	1.61
85	AA	2110	U	C3'-C2'	-7.11	1.45	1.52
34	BA	1406	U	C4'-C3'	-7.11	1.45	1.53
35	BB	27	C	C3'-C2'	-7.11	1.45	1.52
35	BB	124	G	C6-N1	-7.11	1.34	1.39
35	BB	1316	U	C2-N3	-7.11	1.32	1.37
37	BD	48	G	N9-C8	-7.11	1.32	1.37
40	BG	43	U	C2-N3	-7.11	1.32	1.37
85	AA	417	U	N1-C2	-7.11	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1134	G	P-O5'	-7.11	1.52	1.59
85	AA	1203	G	C3'-C2'	-7.11	1.45	1.52
85	AA	1209	U	C4'-C3'	-7.11	1.45	1.53
85	AA	2213	A	O3'-P	-7.11	1.52	1.61
34	BA	61	G	O4'-C1'	-7.11	1.32	1.41
34	BA	339	G	O3'-P	-7.11	1.52	1.61
35	BB	842	G	C2'-C1'	-7.11	1.45	1.53
40	BG	6	A	N7-C5	-7.11	1.34	1.39
34	BA	594	G	N7-C5	-7.11	1.34	1.39
34	BA	1409	A	C3'-C2'	-7.11	1.45	1.52
35	BB	619	A	C2'-C1'	-7.11	1.45	1.53
35	BB	1208	G	N7-C5	-7.11	1.34	1.39
36	BC	92	C	O3'-P	-7.11	1.52	1.61
85	AA	349	C	C2'-C1'	-7.11	1.45	1.53
85	AA	1497	U	O3'-P	-7.11	1.52	1.61
85	AA	1538	C	P-O5'	-7.11	1.52	1.59
85	AA	1710	C	O3'-P	-7.11	1.52	1.61
85	AA	2006	G	O3'-P	-7.11	1.52	1.61
34	BA	212	A	C5-C4	-7.11	1.33	1.38
34	BA	333	A	C5-C4	-7.11	1.33	1.38
34	BA	1196	C	C4'-O4'	-7.11	1.36	1.45
34	BA	1650	G	N9-C4	-7.11	1.32	1.38
34	BA	1699	A	O4'-C1'	7.11	1.50	1.41
38	BE	198	A	C5-C4	-7.11	1.33	1.38
85	AA	1976	G	C2-N2	-7.11	1.27	1.34
34	BA	459	U	C2'-C1'	-7.10	1.45	1.53
34	BA	584	A	O4'-C1'	-7.10	1.32	1.41
34	BA	1036	G	C6-N1	-7.10	1.34	1.39
34	BA	1046	G	O3'-P	-7.10	1.52	1.61
34	BA	1153	C	C2'-C1'	-7.10	1.45	1.53
85	AA	351	C	C2'-C1'	-7.10	1.45	1.53
85	AA	617	C	O3'-P	-7.10	1.52	1.61
85	AA	675	A	C2'-C1'	-7.10	1.45	1.53
34	BA	88	C	C3'-C2'	-7.10	1.45	1.52
34	BA	1054	U	O3'-P	-7.10	1.52	1.61
34	BA	1309	U	O3'-P	-7.10	1.52	1.61
35	BB	1369	A	C5-C4	-7.10	1.33	1.38
41	BH	119	U	C5'-C4'	-7.10	1.42	1.51
85	AA	29	U	C2-N3	-7.10	1.32	1.37
85	AA	1301	C	O3'-P	-7.10	1.52	1.61
85	AA	1305	A	P-O5'	-7.10	1.52	1.59
85	AA	2060	G	N9-C8	-7.10	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2085	C	C1'-N1	-7.10	1.36	1.46
34	BA	125	G	C2'-C1'	-7.10	1.45	1.53
34	BA	126	G	P-O5'	-7.10	1.52	1.59
34	BA	416	A	N3-C4	-7.10	1.30	1.34
34	BA	1253	G	N7-C5	-7.10	1.34	1.39
34	BA	1466	U	O3'-P	-7.10	1.52	1.61
35	BB	1283	C	O3'-P	-7.10	1.52	1.61
41	BH	43	G	P-O5'	-7.10	1.52	1.59
85	AA	2055	G	C2'-C1'	-7.10	1.45	1.53
34	BA	45	A	N3-C4	-7.10	1.30	1.34
34	BA	339	G	C6-N1	-7.10	1.34	1.39
34	BA	752	A	P-O5'	-7.10	1.52	1.59
34	BA	1270	G	C2'-C1'	-7.10	1.45	1.53
34	BA	1504	A	C2'-C1'	-7.10	1.45	1.53
35	BB	812	G	C2'-C1'	-7.10	1.45	1.53
35	BB	861	C	C2'-C1'	-7.10	1.45	1.53
36	BC	156	A	P-O5'	-7.10	1.52	1.59
38	BE	154	A	P-O5'	-7.10	1.52	1.59
40	BG	82	U	P-O5'	-7.10	1.52	1.59
40	BG	176	G	C2'-C1'	-7.10	1.45	1.53
41	BH	5	G	O3'-P	-7.10	1.52	1.61
85	AA	597	A	C8-N7	-7.10	1.26	1.31
85	AA	668	A	C1'-N9	-7.10	1.36	1.46
85	AA	1134	G	C4'-O4'	-7.10	1.36	1.45
85	AA	1897	A	P-O5'	-7.10	1.52	1.59
34	BA	77	C	P-O5'	-7.10	1.52	1.59
35	BB	403	U	C3'-C2'	-7.10	1.45	1.52
35	BB	449	C	C4-N4	-7.10	1.27	1.33
35	BB	1093	C	O3'-P	-7.10	1.52	1.61
85	AA	313	A	C3'-C2'	-7.10	1.45	1.52
85	AA	1848	G	P-O5'	-7.10	1.52	1.59
34	BA	315	U	O3'-P	-7.10	1.52	1.61
34	BA	440	A	C3'-C2'	-7.09	1.45	1.52
34	BA	721	A	N3-C4	-7.09	1.30	1.34
34	BA	1025	A	N3-C4	-7.09	1.30	1.34
34	BA	1599	A	N3-C4	-7.09	1.30	1.34
34	BA	1743	U	O3'-P	-7.09	1.52	1.61
35	BB	500	C	C2'-C1'	-7.09	1.45	1.53
35	BB	643	G	O3'-P	-7.09	1.52	1.61
35	BB	650	A	C5-C4	-7.09	1.33	1.38
36	BC	60	U	P-O5'	-7.09	1.52	1.59
85	AA	80	G	N9-C4	-7.09	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	436	G	C5-C4	-7.09	1.33	1.38
85	AA	486	G	C2'-C1'	-7.09	1.45	1.53
85	AA	783	C	P-O5'	-7.09	1.52	1.59
34	BA	115	U	O3'-P	-7.09	1.52	1.61
34	BA	898	G	P-O5'	-7.09	1.52	1.59
34	BA	1027	C	C2'-C1'	-7.09	1.45	1.53
35	BB	33	A	N9-C8	-7.09	1.32	1.37
35	BB	538	A	N9-C4	-7.09	1.33	1.37
85	AA	891	G	C5'-C4'	7.09	1.59	1.51
85	AA	1235	G	C1'-N9	-7.09	1.36	1.46
85	AA	1470	A	O3'-P	-7.09	1.52	1.61
85	AA	1679	U	C2'-C1'	-7.09	1.45	1.53
34	BA	519	G	N7-C5	-7.09	1.34	1.39
34	BA	1201	G	C4'-C3'	-7.09	1.45	1.53
34	BA	1285	G	O3'-P	-7.09	1.52	1.61
35	BB	73	G	N3-C4	-7.09	1.30	1.35
35	BB	125	G	P-O5'	-7.09	1.52	1.59
35	BB	536	U	P-O5'	-7.09	1.52	1.59
35	BB	542	A	C5-C4	-7.09	1.33	1.38
35	BB	674	C	C1'-N1	-7.09	1.36	1.46
41	BH	111	U	C3'-C2'	-7.09	1.45	1.52
85	AA	33	U	P-O5'	-7.09	1.52	1.59
85	AA	1485	G	C5-C4	-7.09	1.33	1.38
34	BA	354	G	O3'-P	-7.09	1.52	1.61
35	BB	114	A	N7-C5	-7.09	1.34	1.39
35	BB	1075	A	N7-C5	-7.09	1.34	1.39
35	BB	1321	G	C2'-C1'	-7.09	1.45	1.53
36	BC	39	G	O4'-C1'	-7.09	1.32	1.41
37	BD	14	C	O3'-P	-7.09	1.52	1.61
41	BH	24	U	C1'-N1	-7.09	1.36	1.46
41	BH	38	G	N7-C5	-7.09	1.34	1.39
85	AA	446	C	P-O5'	-7.09	1.52	1.59
85	AA	658	C	O3'-P	-7.09	1.52	1.61
85	AA	750	A	O4'-C1'	-7.09	1.32	1.41
85	AA	1677	A	P-O5'	-7.09	1.52	1.59
35	BB	804	U	C4'-O4'	-7.09	1.36	1.45
35	BB	1095	G	C6-N1	-7.09	1.34	1.39
35	BB	1302	C	C2'-C1'	-7.09	1.45	1.53
35	BB	1320	U	O3'-P	-7.09	1.52	1.61
85	AA	1672	G	O3'-P	-7.09	1.52	1.61
85	AA	1962	U	C3'-C2'	-7.09	1.45	1.52
34	BA	41	U	O3'-P	-7.09	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	693	G	O3'-P	-7.09	1.52	1.61
34	BA	760	G	C1'-N9	-7.09	1.36	1.46
34	BA	968	G	C5-C4	-7.09	1.33	1.38
34	BA	1150	A	C2'-C1'	-7.09	1.45	1.53
34	BA	1280	A	O3'-P	-7.09	1.52	1.61
34	BA	1806	A	O3'-P	-7.09	1.52	1.61
35	BB	1221	G	C2'-C1'	-7.09	1.45	1.53
37	BD	75	G	C5-C4	-7.09	1.33	1.38
41	BH	4	U	C2'-C1'	-7.09	1.45	1.53
41	BH	35	G	C3'-C2'	-7.09	1.45	1.52
85	AA	1807	A	O3'-P	-7.09	1.52	1.61
34	BA	1431	G	O3'-P	-7.08	1.52	1.61
36	BC	93	C	C4'-C3'	-7.08	1.45	1.53
85	AA	421	G	P-O5'	-7.08	1.52	1.59
85	AA	1700	C	C2-N3	-7.08	1.30	1.35
34	BA	1618	A	C1'-N9	-7.08	1.36	1.46
34	BA	1807	G	C2'-C1'	-7.08	1.45	1.53
35	BB	7	C	C1'-N1	-7.08	1.36	1.46
35	BB	444	U	N3-C4	-7.08	1.32	1.38
35	BB	997	G	O3'-P	-7.08	1.52	1.61
35	BB	1478	G	C2'-C1'	-7.08	1.45	1.53
37	BD	12	U	C3'-C2'	-7.08	1.45	1.52
85	AA	1209	U	C2'-C1'	-7.08	1.45	1.53
34	BA	1163	G	N1-C2	-7.08	1.32	1.37
34	BA	1177	C	C3'-C2'	-7.08	1.45	1.52
34	BA	1282	G	N3-C4	-7.08	1.30	1.35
34	BA	1288	U	P-O5'	-7.08	1.52	1.59
34	BA	1516	G	N1-C2	-7.08	1.32	1.37
37	BD	77	A	C2'-C1'	-7.08	1.45	1.53
85	AA	1024	G	O3'-P	-7.08	1.52	1.61
35	BB	1067	G	N9-C4	-7.08	1.32	1.38
35	BB	1321	G	C1'-N9	-7.08	1.36	1.46
34	BA	979	G	C1'-N9	-7.08	1.36	1.46
34	BA	1284	G	N7-C5	-7.08	1.35	1.39
34	BA	1295	U	P-O5'	-7.08	1.52	1.59
35	BB	988	G	N9-C4	-7.08	1.32	1.38
85	AA	1153	G	N9-C8	-7.08	1.32	1.37
85	AA	1674	G	O3'-P	-7.08	1.52	1.61
85	AA	2237	G	O3'-P	-7.08	1.52	1.61
34	BA	1156	U	N3-C4	-7.08	1.32	1.38
34	BA	1285	G	N1-C2	-7.08	1.32	1.37
34	BA	1416	C	C2'-C1'	-7.08	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	43	U	C3'-C2'	-7.08	1.45	1.52
37	BD	80	G	C1'-N9	-7.08	1.36	1.46
41	BH	22	A	C1'-N9	-7.08	1.36	1.46
85	AA	11	A	O3'-P	-7.08	1.52	1.61
85	AA	41	G	C3'-C2'	-7.08	1.45	1.52
85	AA	495	G	C5'-C4'	-7.08	1.42	1.51
85	AA	1826	U	O3'-P	-7.08	1.52	1.61
34	BA	480	G	C5-C6	-7.08	1.35	1.42
34	BA	973	U	C4'-O4'	-7.08	1.36	1.45
35	BB	802	G	N9-C4	7.08	1.43	1.38
35	BB	1426	G	C2'-C1'	-7.08	1.45	1.53
38	BE	105	A	C8-N7	-7.08	1.26	1.31
85	AA	439	U	C2'-C1'	-7.08	1.45	1.53
85	AA	493	A	O3'-P	-7.08	1.52	1.61
85	AA	1266	C	C4'-C3'	-7.08	1.45	1.53
85	AA	2053	A	N7-C5	-7.08	1.35	1.39
85	AA	2171	A	P-O5'	-7.08	1.52	1.59
85	AA	2202	G	C2-N2	-7.08	1.27	1.34
34	BA	123	C	C4'-C3'	-7.07	1.45	1.53
34	BA	784	C	C2-N3	-7.07	1.30	1.35
34	BA	856	G	N9-C8	-7.07	1.32	1.37
34	BA	983	A	C1'-N9	-7.07	1.36	1.46
34	BA	992	A	O3'-P	-7.07	1.52	1.61
34	BA	1225	A	C6-N1	-7.07	1.30	1.35
34	BA	1669	C	C2'-C1'	-7.07	1.45	1.53
35	BB	1147	G	C2'-C1'	-7.07	1.45	1.53
35	BB	1343	C	C2'-C1'	-7.07	1.45	1.53
35	BB	1435	G	N3-C4	-7.07	1.30	1.35
35	BB	1490	G	O3'-P	-7.07	1.52	1.61
37	BD	3	G	C3'-C2'	-7.07	1.45	1.52
85	AA	475	A	C5-C4	-7.07	1.33	1.38
85	AA	563	U	P-O5'	-7.07	1.52	1.59
85	AA	1133	C	C3'-C2'	-7.07	1.45	1.52
85	AA	1486	G	C1'-N9	-7.07	1.36	1.46
85	AA	2128	G	C1'-N9	-7.07	1.36	1.46
34	BA	12	G	N3-C4	-7.07	1.30	1.35
34	BA	157	U	P-O5'	-7.07	1.52	1.59
34	BA	739	A	C4'-O4'	-7.07	1.36	1.45
34	BA	927	A	C3'-C2'	-7.07	1.45	1.52
34	BA	1567	G	C5-C4	-7.07	1.33	1.38
34	BA	1722	U	C1'-N1	-7.07	1.36	1.46
35	BB	597	C	C3'-C2'	-7.07	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	36	C	C2-N3	-7.07	1.30	1.35
85	AA	891	G	N7-C5	-7.07	1.35	1.39
34	BA	68	A	N7-C5	-7.07	1.35	1.39
34	BA	568	G	O3'-P	-7.07	1.52	1.61
34	BA	795	G	N7-C5	-7.07	1.35	1.39
34	BA	857	C	O3'-P	-7.07	1.52	1.61
34	BA	1289	C	O4'-C1'	-7.07	1.32	1.41
37	BD	56	G	C3'-C2'	-7.07	1.45	1.52
40	BG	26	G	C5-C4	-7.07	1.33	1.38
85	AA	714	U	C2-N3	-7.07	1.32	1.37
34	BA	362	G	C6-N1	-7.07	1.34	1.39
35	BB	612	A	C3'-C2'	-7.07	1.45	1.52
37	BD	81	C	C2-N3	-7.07	1.30	1.35
85	AA	448	G	O3'-P	-7.07	1.52	1.61
34	BA	336	A	C5-C4	-7.07	1.33	1.38
34	BA	769	U	P-O5'	-7.07	1.52	1.59
34	BA	1637	G	C6-N1	-7.07	1.34	1.39
34	BA	1714	A	O3'-P	-7.07	1.52	1.61
39	BF	69	A	P-O5'	-7.07	1.52	1.59
85	AA	388	G	O3'-P	-7.07	1.52	1.61
85	AA	426	C	P-O5'	-7.07	1.52	1.59
85	AA	673	A	P-O5'	-7.07	1.52	1.59
85	AA	944	C	C4-C5	-7.07	1.37	1.43
85	AA	1136	A	N9-C8	-7.07	1.32	1.37
85	AA	1816	C	O3'-P	-7.07	1.52	1.61
34	BA	572	G	C8-N7	7.07	1.35	1.30
34	BA	1122	G	N7-C5	-7.07	1.35	1.39
34	BA	1613	G	N1-C2	-7.07	1.32	1.37
35	BB	435	A	O3'-P	-7.07	1.52	1.61
35	BB	518	G	P-O5'	-7.07	1.52	1.59
35	BB	818	U	O4'-C1'	-7.07	1.32	1.41
35	BB	1176	G	C2-N2	-7.07	1.27	1.34
35	BB	1402	U	O3'-P	-7.07	1.52	1.61
37	BD	56	G	N9-C4	-7.07	1.32	1.38
40	BG	166	C	C3'-C2'	-7.07	1.45	1.52
85	AA	244	G	N9-C8	-7.07	1.32	1.37
34	BA	1652	G	C4'-C3'	-7.06	1.45	1.53
35	BB	124	G	N7-C5	-7.06	1.35	1.39
35	BB	677	U	O3'-P	-7.06	1.52	1.61
36	BC	19	A	O3'-P	-7.06	1.52	1.61
41	BH	43	G	O3'-P	-7.06	1.52	1.61
34	BA	48	C	C1'-N1	-7.06	1.36	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	49	A	N3-C4	-7.06	1.30	1.34
34	BA	826	C	O3'-P	-7.06	1.52	1.61
35	BB	1171	U	P-O5'	-7.06	1.52	1.59
36	BC	107	C	C3'-C2'	-7.06	1.45	1.52
36	BC	144	C	C1'-N1	-7.06	1.36	1.46
38	BE	52	U	C2'-C1'	-7.06	1.45	1.53
85	AA	422	G	N7-C5	-7.06	1.35	1.39
85	AA	680	U	C2-N3	-7.06	1.32	1.37
34	BA	147	U	O3'-P	-7.06	1.52	1.61
34	BA	335	C	C3'-C2'	-7.06	1.45	1.52
34	BA	977	G	C2'-C1'	-7.06	1.45	1.53
34	BA	1055	U	C3'-C2'	-7.06	1.45	1.52
34	BA	1477	C	C1'-N1	-7.06	1.36	1.46
85	AA	242	G	C2'-C1'	-7.06	1.45	1.53
85	AA	573	U	C2'-C1'	-7.06	1.45	1.53
34	BA	79	C	C4'-C3'	-7.06	1.45	1.53
34	BA	899	G	N3-C4	-7.06	1.30	1.35
34	BA	1278	A	P-O5'	-7.06	1.52	1.59
34	BA	1473	A	C3'-C2'	-7.06	1.45	1.52
40	BG	111	C	C3'-C2'	-7.06	1.45	1.52
85	AA	11	A	C3'-C2'	-7.06	1.45	1.52
85	AA	600	C	C2'-C1'	-7.06	1.45	1.53
85	AA	1846	G	C6-N1	-7.06	1.34	1.39
34	BA	1794	A	C5'-C4'	7.06	1.59	1.51
35	BB	1019	C	C4-N4	-7.06	1.27	1.33
35	BB	1488	G	C1'-N9	-7.06	1.36	1.46
36	BC	72	A	C4'-C3'	-7.06	1.45	1.53
85	AA	421	G	O3'-P	-7.06	1.52	1.61
85	AA	690	G	C2-N2	-7.06	1.27	1.34
85	AA	1139	G	C6-N1	-7.06	1.34	1.39
35	BB	665	A	N3-C4	-7.06	1.30	1.34
38	BE	114	G	C5-C4	-7.06	1.33	1.38
85	AA	87	C	O3'-P	-7.06	1.52	1.61
34	BA	43	U	C3'-C2'	-7.05	1.45	1.52
34	BA	1049	G	C8-N7	-7.05	1.26	1.30
34	BA	1258	G	C4'-C3'	-7.05	1.45	1.53
35	BB	439	G	C2-N2	-7.05	1.27	1.34
35	BB	1335	G	C5-C4	-7.05	1.33	1.38
35	BB	1428	C	C2-N3	-7.05	1.30	1.35
36	BC	15	G	N9-C8	-7.05	1.32	1.37
38	BE	144	A	O3'-P	-7.05	1.52	1.61
38	BE	161	G	C4'-C3'	-7.05	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	773	G	P-O5'	-7.05	1.52	1.59
85	AA	1809	G	C2'-C1'	-7.05	1.45	1.53
85	AA	2199	G	C2-N2	-7.05	1.27	1.34
34	BA	1062	G	N9-C4	-7.05	1.32	1.38
34	BA	1178	U	C4'-C3'	-7.05	1.45	1.53
35	BB	645	C	P-O5'	-7.05	1.52	1.59
34	BA	121	A	C1'-N9	-7.05	1.36	1.46
34	BA	611	A	C8-N7	-7.05	1.26	1.31
34	BA	1315	C	C4'-C3'	-7.05	1.45	1.53
34	BA	1327	G	C1'-N9	-7.05	1.36	1.46
35	BB	1104	A	O3'-P	-7.05	1.52	1.61
35	BB	1153	G	C4'-C3'	-7.05	1.45	1.53
35	BB	1200	A	P-O5'	-7.05	1.52	1.59
36	BC	122	A	P-O5'	-7.05	1.52	1.59
37	BD	3	G	P-O5'	-7.05	1.52	1.59
38	BE	93	U	C2-N3	-7.05	1.32	1.37
40	BG	66	C	C3'-C2'	-7.05	1.45	1.52
85	AA	462	A	O3'-P	-7.05	1.52	1.61
85	AA	1517	G	C3'-C2'	-7.05	1.45	1.52
85	AA	1654	G	O3'-P	-7.05	1.52	1.61
85	AA	1662	U	O3'-P	-7.05	1.52	1.61
34	BA	632	U	P-O5'	-7.05	1.52	1.59
35	BB	52	G	O3'-P	-7.05	1.52	1.61
35	BB	614	U	C2'-C1'	-7.05	1.45	1.53
35	BB	1227	G	P-O5'	-7.05	1.52	1.59
37	BD	56	G	P-O5'	-7.05	1.52	1.59
85	AA	1112	G	C5-C6	-7.05	1.35	1.42
85	AA	1547	G	P-O5'	-7.05	1.52	1.59
34	BA	456	G	C5-C4	-7.05	1.33	1.38
34	BA	613	A	C1'-N9	-7.05	1.36	1.46
34	BA	1115	A	O3'-P	-7.05	1.52	1.61
35	BB	648	G	C5-C6	-7.05	1.35	1.42
35	BB	787	A	C5-C4	-7.05	1.33	1.38
85	AA	153	C	C2'-C1'	-7.05	1.45	1.53
34	BA	792	A	O3'-P	-7.05	1.52	1.61
34	BA	1711	G	C5'-C4'	-7.05	1.42	1.51
35	BB	58	G	C6-N1	-7.05	1.34	1.39
38	BE	48	G	N1-C2	-7.05	1.32	1.37
38	BE	175	U	P-O5'	-7.05	1.52	1.59
85	AA	1673	A	O3'-P	-7.05	1.52	1.61
34	BA	672	G	N3-C4	-7.04	1.30	1.35
34	BA	1455	C	C2'-C1'	-7.04	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	459	U	C4'-C3'	-7.04	1.45	1.53
35	BB	1027	U	C4'-C3'	-7.04	1.45	1.53
35	BB	1159	U	O3'-P	-7.04	1.52	1.61
85	AA	1680	U	O3'-P	-7.04	1.52	1.61
34	BA	74	A	C4'-C3'	-7.04	1.45	1.53
34	BA	96	G	C8-N7	-7.04	1.26	1.30
34	BA	379	C	O3'-P	-7.04	1.52	1.61
34	BA	504	A	C3'-C2'	-7.04	1.45	1.52
34	BA	705	C	C2-N3	-7.04	1.30	1.35
35	BB	423	G	N1-C2	-7.04	1.32	1.37
35	BB	1026	G	C4'-C3'	7.04	1.60	1.53
35	BB	1120	A	C1'-N9	-7.04	1.36	1.46
35	BB	1254	G	N7-C5	-7.04	1.35	1.39
35	BB	1335	G	C2-N2	-7.04	1.27	1.34
35	BB	1375	G	C3'-C2'	-7.04	1.45	1.52
35	BB	1406	C	C2-N3	-7.04	1.30	1.35
85	AA	334	A	C3'-C2'	-7.04	1.45	1.52
34	BA	140	C	P-O5'	-7.04	1.52	1.59
34	BA	306	G	C2'-C1'	-7.04	1.45	1.53
34	BA	1500	G	C2'-C1'	-7.04	1.45	1.53
35	BB	745	C	O3'-P	-7.04	1.52	1.61
37	BD	58	G	O3'-P	-7.04	1.52	1.61
40	BG	48	U	N3-C4	-7.04	1.32	1.38
85	AA	765	U	O3'-P	-7.04	1.52	1.61
85	AA	1465	C	O3'-P	-7.04	1.52	1.61
85	AA	2046	G	O3'-P	-7.04	1.52	1.61
35	BB	1184	C	C2-N3	-7.04	1.30	1.35
40	BG	46	G	C2-N2	-7.04	1.27	1.34
34	BA	126	G	C1'-N9	-7.04	1.36	1.46
34	BA	1422	A	O3'-P	-7.04	1.52	1.61
35	BB	1186	A	O3'-P	-7.04	1.52	1.61
38	BE	109	C	O3'-P	-7.04	1.52	1.61
39	BF	61	A	O3'-P	-7.04	1.52	1.61
85	AA	1652	A	N9-C4	-7.04	1.33	1.37
85	AA	2082	C	N1-C6	-7.04	1.32	1.37
34	BA	411	C	O3'-P	-7.04	1.52	1.61
34	BA	1543	A	C3'-C2'	-7.04	1.45	1.52
35	BB	607	G	C3'-C2'	-7.04	1.45	1.52
35	BB	1180	G	N9-C4	-7.04	1.32	1.38
34	BA	352	G	C4'-C3'	-7.04	1.45	1.53
34	BA	566	G	C6-N1	-7.04	1.34	1.39
34	BA	924	U	C2'-C1'	-7.04	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1252	G	N1-C2	-7.04	1.32	1.37
34	BA	1738	G	C2'-C1'	-7.04	1.45	1.53
35	BB	481	A	C2'-C1'	-7.04	1.45	1.53
35	BB	542	A	N3-C4	-7.04	1.30	1.34
35	BB	1052	G	N7-C5	-7.04	1.35	1.39
85	AA	456	A	C3'-C2'	-7.04	1.45	1.52
85	AA	512	U	O3'-P	-7.04	1.52	1.61
85	AA	1669	G	O3'-P	-7.04	1.52	1.61
34	BA	19	G	C4'-C3'	-7.03	1.45	1.53
34	BA	493	G	C3'-C2'	-7.03	1.45	1.52
35	BB	993	A	C8-N7	-7.03	1.26	1.31
35	BB	1398	A	C8-N7	-7.03	1.26	1.31
36	BC	70	C	C1'-N1	-7.03	1.37	1.46
85	AA	11	A	C5'-C4'	-7.03	1.43	1.51
85	AA	157	G	C2-N2	-7.03	1.27	1.34
85	AA	755	G	O3'-P	-7.03	1.52	1.61
85	AA	1482	C	C2'-C1'	-7.03	1.45	1.53
34	BA	78	U	P-O5'	-7.03	1.52	1.59
34	BA	377	G	C2-N2	-7.03	1.27	1.34
40	BG	82	U	C2-N3	-7.03	1.32	1.37
40	BG	163	G	O3'-P	-7.03	1.52	1.61
85	AA	1270	C	N1-C6	-7.03	1.32	1.37
85	AA	1513	U	C2'-C1'	-7.03	1.45	1.53
85	AA	1527	G	N3-C4	-7.03	1.30	1.35
85	AA	2188	C	C4'-C3'	-7.03	1.45	1.53
34	BA	46	C	C4'-C3'	-7.03	1.45	1.53
34	BA	414	A	C1'-N9	-7.03	1.37	1.46
35	BB	29	C	P-O5'	-7.03	1.52	1.59
35	BB	59	U	C4'-C3'	-7.03	1.45	1.53
35	BB	574	G	C6-N1	-7.03	1.34	1.39
35	BB	837	A	C4'-O4'	-7.03	1.36	1.45
35	BB	1281	G	O3'-P	-7.03	1.52	1.61
36	BC	100	U	C3'-C2'	-7.03	1.45	1.52
37	BD	34	C	O3'-P	-7.03	1.52	1.61
41	BH	20	A	C3'-C2'	-7.03	1.45	1.52
41	BH	121	A	C2'-C1'	-7.03	1.45	1.53
85	AA	869	A	O3'-P	-7.03	1.52	1.61
85	AA	2218	G	N9-C4	-7.03	1.32	1.38
34	BA	1247	G	C5-C4	-7.03	1.33	1.38
35	BB	448	G	C2-N2	-7.03	1.27	1.34
41	BH	15	A	C5-C4	-7.03	1.33	1.38
41	BH	20	A	C2'-C1'	-7.03	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1693	C	P-O5'	-7.03	1.52	1.59
34	BA	1067	G	C1'-N9	-7.03	1.37	1.46
34	BA	1301	G	C1'-N9	-7.03	1.37	1.46
34	BA	1328	U	C2-N3	-7.03	1.32	1.37
34	BA	1490	U	N1-C2	-7.03	1.32	1.38
35	BB	1013	U	C2'-C1'	-7.03	1.45	1.53
35	BB	1072	C	C2'-C1'	-7.03	1.45	1.53
39	BF	45	G	C2'-C1'	-7.03	1.45	1.53
40	BG	136	G	C1'-N9	-7.03	1.37	1.46
34	BA	196	A	C6-N6	-7.03	1.28	1.33
34	BA	318	U	O3'-P	-7.03	1.52	1.61
34	BA	1469	G	N7-C5	-7.03	1.35	1.39
35	BB	689	C	C2'-C1'	-7.03	1.45	1.53
35	BB	1036	G	C2-N2	-7.03	1.27	1.34
35	BB	1408	G	N9-C8	-7.03	1.32	1.37
38	BE	32	U	C1'-N1	-7.03	1.37	1.46
40	BG	130	G	N9-C4	-7.03	1.32	1.38
85	AA	170	C	C3'-C2'	-7.03	1.45	1.52
85	AA	982	G	C2-N3	-7.03	1.27	1.32
85	AA	1562	U	C4'-C3'	-7.03	1.45	1.53
85	AA	1823	G	P-O5'	-7.03	1.52	1.59
35	BB	1446	C	O3'-P	-7.02	1.52	1.61
85	AA	123	A	C2'-C1'	-7.02	1.45	1.53
34	BA	40	A	N7-C5	-7.02	1.35	1.39
35	BB	792	G	O3'-P	-7.02	1.52	1.61
36	BC	49	G	N1-C2	-7.02	1.32	1.37
38	BE	115	U	C2'-C1'	-7.02	1.45	1.53
38	BE	185	G	N9-C4	-7.02	1.32	1.38
85	AA	188	G	P-O5'	-7.02	1.52	1.59
85	AA	380	C	C2'-C1'	-7.02	1.45	1.53
34	BA	166	G	C3'-C2'	-7.02	1.45	1.52
34	BA	1520	A	C4'-C3'	-7.02	1.45	1.53
35	BB	57	G	C3'-C2'	-7.02	1.45	1.52
35	BB	505	G	N9-C8	-7.02	1.32	1.37
35	BB	1166	A	N9-C4	-7.02	1.33	1.37
34	BA	131	A	N9-C4	-7.02	1.33	1.37
34	BA	183	G	N7-C5	-7.02	1.35	1.39
34	BA	408	U	C2'-C1'	-7.02	1.45	1.53
34	BA	676	G	C1'-N9	-7.02	1.37	1.46
35	BB	438	G	C1'-N9	-7.02	1.37	1.46
35	BB	501	G	C2'-C1'	-7.02	1.45	1.53
35	BB	1395	G	N1-C2	-7.02	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	15	G	P-O5'	-7.02	1.52	1.59
36	BC	111	C	C4-C5	-7.02	1.37	1.43
85	AA	305	A	C4'-C3'	7.02	1.60	1.53
85	AA	334	A	O3'-P	-7.02	1.52	1.61
85	AA	1517	G	C5'-C4'	-7.02	1.43	1.51
34	BA	43	U	C1'-N1	-7.02	1.37	1.46
34	BA	67	A	C1'-N9	-7.02	1.37	1.46
34	BA	216	C	C3'-C2'	-7.02	1.45	1.52
34	BA	301	U	C1'-N1	-7.02	1.37	1.46
35	BB	580	A	C5'-C4'	7.02	1.59	1.51
35	BB	1027	U	C2'-C1'	-7.02	1.45	1.53
36	BC	6	G	N3-C4	-7.02	1.30	1.35
36	BC	51	A	C5-C6	-7.02	1.34	1.41
40	BG	53	C	O3'-P	-7.02	1.52	1.61
40	BG	109	C	C2-N3	-7.02	1.30	1.35
85	AA	149	A	C2'-C1'	-7.02	1.45	1.53
85	AA	335	G	C3'-C2'	-7.02	1.45	1.52
85	AA	964	C	C5'-C4'	7.02	1.59	1.51
86	AB	70	G	C2'-C1'	-7.02	1.45	1.53
34	BA	289	A	O4'-C1'	-7.02	1.32	1.41
34	BA	1015	G	C2'-C1'	-7.02	1.45	1.53
34	BA	1807	G	N1-C2	-7.02	1.32	1.37
35	BB	139	G	C2'-C1'	-7.02	1.45	1.53
36	BC	5	U	O3'-P	-7.02	1.52	1.61
85	AA	247	G	C5-C4	-7.02	1.33	1.38
85	AA	320	U	O3'-P	-7.02	1.52	1.61
85	AA	1094	G	P-O5'	-7.02	1.52	1.59
85	AA	1497	U	C3'-C2'	-7.02	1.45	1.52
85	AA	2228	G	C1'-N9	-7.02	1.37	1.46
34	BA	319	C	P-O5'	-7.01	1.52	1.59
34	BA	339	G	C5-C4	-7.01	1.33	1.38
34	BA	1547	G	C1'-N9	-7.01	1.37	1.46
35	BB	35	G	C1'-N9	-7.01	1.37	1.46
85	AA	19	A	C2'-C1'	-7.01	1.45	1.53
85	AA	1482	C	C4'-C3'	-7.01	1.45	1.53
85	AA	1812	C	O3'-P	-7.01	1.52	1.61
85	AA	1916	A	P-O5'	-7.01	1.52	1.59
85	AA	2229	G	N1-C2	-7.01	1.32	1.37
34	BA	1704	G	C1'-N9	-7.01	1.37	1.46
40	BG	94	G	O3'-P	-7.01	1.52	1.61
34	BA	1093	G	N1-C2	-7.01	1.32	1.37
34	BA	1094	U	C3'-C2'	-7.01	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	785	G	C2-N2	-7.01	1.27	1.34
38	BE	5	A	P-O5'	-7.01	1.52	1.59
40	BG	37	G	N7-C5	-7.01	1.35	1.39
40	BG	159	A	C5-C4	-7.01	1.33	1.38
85	AA	1552	U	P-O5'	-7.01	1.52	1.59
34	BA	313	C	P-O5'	-7.01	1.52	1.59
34	BA	326	A	P-O5'	-7.01	1.52	1.59
35	BB	100	A	P-O5'	-7.01	1.52	1.59
35	BB	1029	U	O3'-P	-7.01	1.52	1.61
35	BB	1255	U	P-O5'	-7.01	1.52	1.59
35	BB	1278	A	P-O5'	-7.01	1.52	1.59
36	BC	28	C	O3'-P	-7.01	1.52	1.61
37	BD	113	G	C3'-C2'	-7.01	1.45	1.52
41	BH	23	G	C4'-C3'	-7.01	1.45	1.53
36	BC	97	U	C3'-C2'	-7.01	1.45	1.52
85	AA	323	U	C2-N3	-7.01	1.32	1.37
85	AA	1537	A	C2'-C1'	-7.01	1.45	1.53
85	AA	1570	A	P-O5'	-7.01	1.52	1.59
85	AA	2138	G	N3-C4	-7.01	1.30	1.35
34	BA	912	G	N9-C8	-7.01	1.32	1.37
34	BA	1270	G	C1'-N9	-7.01	1.37	1.46
34	BA	1801	G	C5-C4	-7.01	1.33	1.38
34	BA	1802	C	C1'-N1	-7.01	1.37	1.46
41	BH	106	G	C5-C4	-7.01	1.33	1.38
85	AA	2024	U	O3'-P	-7.01	1.52	1.61
34	BA	106	U	C4'-C3'	-7.00	1.45	1.53
34	BA	1512	C	O3'-P	-7.00	1.52	1.61
36	BC	133	C	C5'-C4'	7.00	1.59	1.51
85	AA	1286	C	O3'-P	-7.00	1.52	1.61
34	BA	429	G	P-O5'	-7.00	1.52	1.59
34	BA	717	U	O3'-P	-7.00	1.52	1.61
34	BA	1333	G	C2'-C1'	-7.00	1.45	1.53
35	BB	458	U	C3'-C2'	-7.00	1.45	1.52
35	BB	484	G	C4'-O4'	-7.00	1.36	1.45
35	BB	1351	G	C5-C4	-7.00	1.33	1.38
35	BB	1404	A	C1'-N9	-7.00	1.37	1.46
40	BG	46	G	N3-C4	-7.00	1.30	1.35
40	BG	90	G	C5-C4	-7.00	1.33	1.38
85	AA	1217	U	C2-N3	-7.00	1.32	1.37
85	AA	2198	G	N7-C5	-7.00	1.35	1.39
34	BA	1167	A	N9-C4	-7.00	1.33	1.37
35	BB	1294	C	C1'-N1	-7.00	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	41	A	C5-C4	-7.00	1.33	1.38
85	AA	1169	A	C2'-C1'	-7.00	1.45	1.53
85	AA	1472	G	C5-C4	-7.00	1.33	1.38
34	BA	1462	U	C2-N3	-7.00	1.32	1.37
35	BB	58	G	N1-C2	-7.00	1.32	1.37
37	BD	29	C	C3'-C2'	-7.00	1.45	1.52
37	BD	118	C	O3'-P	-7.00	1.52	1.61
85	AA	445	U	P-O5'	-7.00	1.52	1.59
85	AA	1187	G	N9-C4	-7.00	1.32	1.38
85	AA	1450	U	O3'-P	-7.00	1.52	1.61
85	AA	1471	G	C4'-C3'	-7.00	1.45	1.53
34	BA	194	G	N9-C4	-7.00	1.32	1.38
34	BA	367	G	C6-N1	-7.00	1.34	1.39
34	BA	380	A	C2'-C1'	-7.00	1.45	1.53
34	BA	432	A	C1'-N9	-7.00	1.37	1.46
34	BA	531	C	C4'-C3'	-7.00	1.45	1.53
35	BB	676	G	C5-C4	-7.00	1.33	1.38
40	BG	4	A	C3'-C2'	-7.00	1.45	1.52
40	BG	180	C	P-O5'	-7.00	1.52	1.59
85	AA	1485	G	N7-C5	-7.00	1.35	1.39
34	BA	277	A	C3'-C2'	-7.00	1.45	1.52
34	BA	342	U	O3'-P	-7.00	1.52	1.61
34	BA	362	G	P-O5'	-7.00	1.52	1.59
34	BA	487	A	P-O5'	-7.00	1.52	1.59
34	BA	1104	C	C1'-N1	-7.00	1.37	1.46
34	BA	1242	A	O3'-P	-7.00	1.52	1.61
34	BA	1330	G	C3'-C2'	-7.00	1.45	1.52
34	BA	1614	G	N7-C5	-7.00	1.35	1.39
35	BB	665	A	O3'-P	-7.00	1.52	1.61
37	BD	33	U	P-O5'	-7.00	1.52	1.59
85	AA	29	U	C2'-C1'	-7.00	1.45	1.53
85	AA	1988	A	N9-C4	-7.00	1.33	1.37
85	AA	2125	A	C2'-C1'	-7.00	1.45	1.53
34	BA	1219	G	C6-N1	-7.00	1.34	1.39
35	BB	1427	A	N9-C8	-7.00	1.32	1.37
36	BC	44	A	N9-C4	-7.00	1.33	1.37
37	BD	47	U	N3-C4	-7.00	1.32	1.38
85	AA	479	C	C2'-C1'	-7.00	1.45	1.53
85	AA	608	A	C2'-C1'	-7.00	1.45	1.53
34	BA	1270	G	P-O5'	-6.99	1.52	1.59
34	BA	1654	G	C2-N2	-6.99	1.27	1.34
34	BA	1832	A	C2'-C1'	-6.99	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	491	A	C5-C4	-6.99	1.33	1.38
35	BB	582	G	N3-C4	-6.99	1.30	1.35
85	AA	17	C	C2'-C1'	-6.99	1.45	1.53
85	AA	424	A	N3-C4	-6.99	1.30	1.34
85	AA	1420	U	C5'-C4'	6.99	1.59	1.51
34	BA	354	G	P-O5'	-6.99	1.52	1.59
34	BA	1529	G	O3'-P	-6.99	1.52	1.61
35	BB	830	G	N1-C2	-6.99	1.32	1.37
40	BG	177	U	C3'-C2'	-6.99	1.45	1.52
34	BA	126	G	C6-N1	-6.99	1.34	1.39
34	BA	564	C	C2'-C1'	-6.99	1.45	1.53
34	BA	1243	A	C5-C4	-6.99	1.33	1.38
34	BA	1467	U	C2'-C1'	-6.99	1.45	1.53
34	BA	1499	A	O3'-P	-6.99	1.52	1.61
35	BB	115	A	C4'-O4'	-6.99	1.36	1.45
35	BB	1142	C	C2-N3	-6.99	1.30	1.35
41	BH	119	U	C4'-O4'	-6.99	1.36	1.45
85	AA	688	C	C2-N3	-6.99	1.30	1.35
85	AA	1490	A	C2'-C1'	-6.99	1.45	1.53
85	AA	1579	A	C2'-C1'	-6.99	1.45	1.53
85	AA	1660	U	O3'-P	-6.99	1.52	1.61
85	AA	2068	A	C5'-C4'	6.99	1.59	1.51
34	BA	1063	G	N1-C2	-6.99	1.32	1.37
34	BA	1203	G	N1-C2	-6.99	1.32	1.37
85	AA	3	U	O3'-P	-6.99	1.52	1.61
85	AA	160	A	N9-C4	-6.99	1.33	1.37
85	AA	749	C	P-O5'	-6.99	1.52	1.59
85	AA	812	C	P-O5'	-6.99	1.52	1.59
85	AA	1611	A	O3'-P	-6.99	1.52	1.61
34	BA	68	A	C4'-C3'	-6.99	1.45	1.53
34	BA	937	G	N1-C2	-6.99	1.32	1.37
35	BB	1488	G	N7-C5	-6.99	1.35	1.39
37	BD	85	C	C4'-C3'	-6.99	1.45	1.53
85	AA	65	A	O3'-P	-6.99	1.52	1.61
85	AA	496	C	C2'-C1'	-6.99	1.45	1.53
85	AA	1464	G	C1'-N9	-6.99	1.37	1.46
85	AA	1968	A	N9-C4	-6.99	1.33	1.37
34	BA	61	G	C3'-C2'	-6.99	1.45	1.52
34	BA	386	A	C6-N1	-6.99	1.30	1.35
34	BA	1541	G	C1'-N9	-6.99	1.37	1.46
34	BA	1720	U	C1'-N1	-6.99	1.37	1.46
35	BB	593	A	P-O5'	-6.99	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1236	A	C5-C4	-6.99	1.33	1.38
36	BC	120	G	P-O5'	-6.99	1.52	1.59
38	BE	97	G	C2'-C1'	-6.99	1.45	1.53
40	BG	4	A	N1-C2	-6.99	1.28	1.34
85	AA	1599	G	P-O5'	-6.99	1.52	1.59
34	BA	1067	G	C5-C4	-6.98	1.33	1.38
34	BA	205	G	N7-C5	-6.98	1.35	1.39
34	BA	207	A	C2'-C1'	-6.98	1.45	1.53
34	BA	761	U	C5'-C4'	-6.98	1.43	1.51
35	BB	1224	C	O3'-P	-6.98	1.52	1.61
35	BB	1247	C	C2'-C1'	-6.98	1.45	1.53
35	BB	1257	A	N9-C4	-6.98	1.33	1.37
38	BE	58	U	O3'-P	-6.98	1.52	1.61
39	BF	22	U	N1-C6	-6.98	1.31	1.38
41	BH	21	G	N7-C5	-6.98	1.35	1.39
85	AA	94	C	O3'-P	-6.98	1.52	1.61
85	AA	1530	U	P-O5'	-6.98	1.52	1.59
34	BA	150	C	P-O5'	-6.98	1.52	1.59
34	BA	1826	C	C4-N4	-6.98	1.27	1.33
35	BB	42	A	O3'-P	-6.98	1.52	1.61
35	BB	505	G	N1-C2	-6.98	1.32	1.37
35	BB	1176	G	C2'-C1'	-6.98	1.45	1.53
37	BD	46	G	C2'-C1'	-6.98	1.45	1.53
37	BD	72	U	C2-N3	-6.98	1.32	1.37
38	BE	33	C	C4'-C3'	-6.98	1.45	1.53
41	BH	19	G	C2'-C1'	-6.98	1.45	1.53
85	AA	667	A	C5-C4	-6.98	1.33	1.38
85	AA	1824	G	P-O5'	-6.98	1.52	1.59
34	BA	602	G	N7-C5	-6.98	1.35	1.39
34	BA	607	C	C4-N4	-6.98	1.27	1.33
35	BB	447	C	P-O5'	-6.98	1.52	1.59
35	BB	1359	G	C8-N7	-6.98	1.26	1.30
85	AA	414	C	O3'-P	-6.98	1.52	1.61
85	AA	2239	A	C2'-C1'	-6.98	1.45	1.53
34	BA	14	G	C4'-O4'	-6.98	1.36	1.45
34	BA	924	U	C2-N3	-6.98	1.32	1.37
34	BA	1472	G	C3'-C2'	-6.98	1.45	1.52
35	BB	669	A	C2'-C1'	-6.98	1.45	1.53
35	BB	704	G	N1-C2	-6.98	1.32	1.37
35	BB	1356	G	N1-C2	-6.98	1.32	1.37
37	BD	105	G	P-O5'	-6.98	1.52	1.59
38	BE	105	A	C4'-C3'	-6.98	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	140	C	P-O5'	-6.98	1.52	1.59
34	BA	68	A	O3'-P	-6.98	1.52	1.61
34	BA	428	C	C3'-C2'	-6.98	1.45	1.52
34	BA	1556	A	C4'-O4'	-6.98	1.36	1.45
35	BB	1136	G	C1'-N9	-6.98	1.37	1.46
85	AA	182	C	O3'-P	-6.98	1.52	1.61
85	AA	1604	A	O3'-P	-6.98	1.52	1.61
34	BA	627	U	C2-N3	-6.97	1.32	1.37
34	BA	913	U	C2'-C1'	-6.97	1.45	1.53
34	BA	1331	G	O3'-P	-6.97	1.52	1.61
35	BB	433	C	C3'-C2'	-6.97	1.45	1.52
35	BB	690	C	C1'-N1	-6.97	1.37	1.46
35	BB	1299	G	C4'-C3'	-6.97	1.45	1.53
35	BB	1376	G	C5'-C4'	-6.97	1.43	1.51
39	BF	66	C	P-O5'	-6.97	1.52	1.59
85	AA	492	C	O3'-P	-6.97	1.52	1.61
85	AA	682	C	P-O5'	-6.97	1.52	1.59
85	AA	1200	A	C2'-C1'	-6.97	1.45	1.53
34	BA	383	G	C2-N3	-6.97	1.27	1.32
34	BA	437	G	C8-N7	-6.97	1.26	1.30
34	BA	970	U	O3'-P	-6.97	1.52	1.61
35	BB	499	A	O3'-P	-6.97	1.52	1.61
36	BC	21	U	O3'-P	-6.97	1.52	1.61
85	AA	470	C	N1-C6	-6.97	1.32	1.37
85	AA	476	C	C2-N3	-6.97	1.30	1.35
85	AA	2137	A	N7-C5	-6.97	1.35	1.39
34	BA	629	G	N9-C4	-6.97	1.32	1.38
85	AA	196	U	P-O5'	-6.97	1.52	1.59
85	AA	319	U	P-O5'	-6.97	1.52	1.59
85	AA	1507	G	N3-C4	-6.97	1.30	1.35
34	BA	16	C	O3'-P	-6.97	1.52	1.61
34	BA	1408	C	N1-C6	-6.97	1.32	1.37
35	BB	368	C	C2-N3	-6.97	1.30	1.35
35	BB	660	G	N9-C4	-6.97	1.32	1.38
35	BB	1067	G	C6-N1	-6.97	1.34	1.39
35	BB	1250	A	C3'-C2'	-6.97	1.45	1.52
36	BC	61	A	O3'-P	-6.97	1.52	1.61
38	BE	107	U	P-O5'	-6.97	1.52	1.59
38	BE	128	G	C3'-C2'	-6.97	1.45	1.52
85	AA	1497	U	C2'-C1'	-6.97	1.45	1.53
34	BA	1658	G	O3'-P	-6.97	1.52	1.61
34	BA	1816	G	C5-C6	-6.97	1.35	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	474	G	C2'-C1'	-6.97	1.45	1.53
35	BB	591	A	C1'-N9	-6.97	1.37	1.46
35	BB	1382	U	P-O5'	-6.97	1.52	1.59
40	BG	132	U	C1'-N1	-6.97	1.37	1.46
34	BA	143	A	C2'-C1'	-6.97	1.45	1.53
34	BA	894	G	N1-C2	-6.97	1.32	1.37
34	BA	1611	A	C5-C4	-6.97	1.33	1.38
35	BB	411	A	O3'-P	-6.97	1.52	1.61
35	BB	1281	G	P-O5'	-6.97	1.52	1.59
36	BC	24	G	C4'-C3'	-6.97	1.45	1.53
39	BF	50	C	N1-C6	-6.97	1.32	1.37
40	BG	1	G	C3'-C2'	-6.97	1.45	1.52
40	BG	55	A	C2'-C1'	-6.97	1.45	1.53
85	AA	778	C	P-O5'	-6.97	1.52	1.59
85	AA	881	C	N3-C4	-6.97	1.29	1.33
85	AA	1705	G	N9-C4	-6.97	1.32	1.38
2	A1	100	TYR	CB-CG	-6.96	1.41	1.51
34	BA	346	A	N7-C5	-6.96	1.35	1.39
34	BA	459	U	P-O5'	-6.96	1.52	1.59
34	BA	1262	A	C5'-C4'	-6.96	1.43	1.51
34	BA	1706	A	N7-C5	-6.96	1.35	1.39
36	BC	164	G	C2'-C1'	-6.96	1.45	1.53
40	BG	156	G	C4'-C3'	-6.96	1.45	1.53
85	AA	344	U	C5'-C4'	6.96	1.59	1.51
85	AA	387	U	P-O5'	-6.96	1.52	1.59
85	AA	435	A	O3'-P	-6.96	1.52	1.61
85	AA	513	G	C2'-C1'	-6.96	1.45	1.53
85	AA	553	G	O3'-P	-6.96	1.52	1.61
34	BA	108	A	N3-C4	-6.96	1.30	1.34
34	BA	1427	U	P-O5'	-6.96	1.52	1.59
34	BA	1477	C	C5'-C4'	-6.96	1.43	1.51
37	BD	112	U	O3'-P	-6.96	1.52	1.61
85	AA	517	A	C3'-C2'	-6.96	1.45	1.52
85	AA	1527	G	C6-N1	-6.96	1.34	1.39
85	AA	2082	C	O3'-P	-6.96	1.52	1.61
34	BA	176	G	O3'-P	-6.96	1.52	1.61
34	BA	451	A	N9-C4	-6.96	1.33	1.37
34	BA	616	G	N9-C4	-6.96	1.32	1.38
34	BA	1101	A	C1'-N9	-6.96	1.37	1.46
34	BA	1221	A	C2'-C1'	-6.96	1.45	1.53
35	BB	617	C	N3-C4	-6.96	1.29	1.33
35	BB	901	U	C4'-C3'	-6.96	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1071	G	C3'-C2'	-6.96	1.45	1.52
35	BB	1234	G	O3'-P	-6.96	1.52	1.61
35	BB	1438	U	P-O5'	-6.96	1.52	1.59
85	AA	1129	A	O3'-P	-6.96	1.52	1.61
85	AA	1517	G	C2-N2	-6.96	1.27	1.34
85	AA	1991	C	C4'-O4'	-6.96	1.36	1.45
85	AA	2116	U	O3'-P	-6.96	1.52	1.61
34	BA	32	A	C2'-C1'	-6.96	1.45	1.53
34	BA	843	G	N9-C4	6.96	1.43	1.38
34	BA	1572	G	C6-N1	-6.96	1.34	1.39
35	BB	35	G	N9-C8	-6.96	1.32	1.37
35	BB	641	C	C2'-C1'	-6.96	1.45	1.53
35	BB	680	A	P-O5'	-6.96	1.52	1.59
36	BC	59	A	C5-C4	-6.96	1.33	1.38
40	BG	125	C	C3'-C2'	-6.96	1.45	1.52
85	AA	12	U	C2-N3	-6.96	1.32	1.37
85	AA	1228	A	O3'-P	-6.96	1.52	1.61
34	BA	5	C	C2'-C1'	-6.96	1.45	1.53
34	BA	293	A	C5-C4	-6.96	1.33	1.38
34	BA	1507	C	C1'-N1	-6.96	1.37	1.46
34	BA	1697	U	C4-C5	-6.96	1.37	1.43
34	BA	1843	G	P-O5'	-6.96	1.52	1.59
35	BB	617	C	O3'-P	-6.96	1.52	1.61
35	BB	654	C	C2-N3	-6.96	1.30	1.35
35	BB	1272	G	N1-C2	-6.96	1.32	1.37
35	BB	1404	A	N9-C8	-6.96	1.32	1.37
37	BD	8	A	O3'-P	-6.96	1.52	1.61
37	BD	94	C	O3'-P	-6.96	1.52	1.61
40	BG	126	G	C5-C4	-6.96	1.33	1.38
85	AA	80	G	O3'-P	-6.96	1.52	1.61
85	AA	322	A	O3'-P	-6.96	1.52	1.61
85	AA	927	A	C2'-C1'	-6.96	1.45	1.53
85	AA	1495	G	C4'-O4'	-6.96	1.36	1.45
34	BA	818	G	C2'-C1'	-6.96	1.45	1.53
34	BA	927	A	O3'-P	-6.96	1.52	1.61
35	BB	73	G	C5-C4	-6.96	1.33	1.38
35	BB	394	A	C5-C4	-6.96	1.33	1.38
35	BB	1196	A	P-O5'	-6.96	1.52	1.59
36	BC	147	G	N1-C2	-6.96	1.32	1.37
41	BH	17	A	C4'-C3'	-6.96	1.45	1.53
85	AA	962	U	O3'-P	-6.96	1.52	1.61
85	AA	1144	G	O3'-P	-6.96	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	91	C	C3'-C2'	-6.96	1.45	1.52
34	BA	140	C	O3'-P	-6.96	1.52	1.61
34	BA	704	G	P-O5'	-6.96	1.52	1.59
38	BE	32	U	C4'-C3'	-6.96	1.45	1.53
38	BE	198	A	C1'-N9	-6.96	1.37	1.46
85	AA	665	A	P-O5'	-6.96	1.52	1.59
85	AA	1735	U	O3'-P	-6.96	1.52	1.61
85	AA	2222	G	C6-N1	-6.96	1.34	1.39
34	BA	464	U	O3'-P	-6.95	1.52	1.61
34	BA	502	U	C2-N3	-6.95	1.32	1.37
34	BA	569	C	C3'-C2'	-6.95	1.45	1.52
34	BA	1305	A	P-O5'	-6.95	1.52	1.59
34	BA	1695	G	C2'-C1'	-6.95	1.45	1.53
35	BB	587	A	C5-C4	-6.95	1.33	1.38
40	BG	158	A	O3'-P	-6.95	1.52	1.61
41	BH	133	U	C2-N3	-6.95	1.32	1.37
85	AA	246	C	C2-N3	-6.95	1.30	1.35
85	AA	687	G	O3'-P	-6.95	1.52	1.61
85	AA	1221	G	N3-C4	-6.95	1.30	1.35
85	AA	1829	C	C3'-C2'	-6.95	1.45	1.52
34	BA	585	G	C4'-C3'	-6.95	1.45	1.53
34	BA	1031	U	C2-N3	-6.95	1.32	1.37
35	BB	562	A	C5-C4	-6.95	1.33	1.38
35	BB	705	C	C2-N3	-6.95	1.30	1.35
35	BB	1089	A	C5-C4	-6.95	1.33	1.38
85	AA	159	G	N9-C4	-6.95	1.32	1.38
85	AA	994	A	C4'-C3'	-6.95	1.45	1.53
85	AA	1531	G	C5-C4	-6.95	1.33	1.38
34	BA	456	G	N9-C4	-6.95	1.32	1.38
34	BA	1059	U	C4'-O4'	-6.95	1.36	1.45
35	BB	1224	C	C4'-C3'	-6.95	1.45	1.53
37	BD	79	G	C2-N2	-6.95	1.27	1.34
40	BG	79	U	C3'-C2'	-6.95	1.45	1.52
85	AA	440	U	C2-N3	-6.95	1.32	1.37
85	AA	907	G	N9-C8	-6.95	1.32	1.37
85	AA	962	U	P-O5'	-6.95	1.52	1.59
85	AA	1235	G	N7-C5	-6.95	1.35	1.39
34	BA	447	U	O3'-P	-6.95	1.52	1.61
34	BA	492	G	N9-C8	-6.95	1.32	1.37
34	BA	752	A	C1'-N9	-6.95	1.37	1.46
34	BA	901	C	P-O5'	-6.95	1.52	1.59
35	BB	777	C	O3'-P	-6.95	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1160	U	C2-N3	-6.95	1.32	1.37
35	BB	1185	G	C4'-C3'	-6.95	1.45	1.53
36	BC	36	G	C1'-N9	-6.95	1.37	1.46
36	BC	40	A	O3'-P	-6.95	1.52	1.61
38	BE	18	U	C3'-C2'	-6.95	1.45	1.52
85	AA	1221	G	C6-N1	-6.95	1.34	1.39
85	AA	2039	G	C4'-O4'	-6.95	1.36	1.45
34	BA	1437	G	C5-C4	-6.95	1.33	1.38
35	BB	1182	A	O3'-P	-6.95	1.52	1.61
35	BB	1340	U	O3'-P	-6.95	1.52	1.61
38	BE	131	C	C3'-C2'	-6.95	1.45	1.52
85	AA	428	G	C2-N2	-6.95	1.27	1.34
85	AA	662	U	C4'-C3'	-6.95	1.45	1.53
34	BA	441	A	O3'-P	-6.95	1.52	1.61
34	BA	1735	G	C8-N7	-6.95	1.26	1.30
35	BB	788	U	C3'-C2'	-6.95	1.45	1.52
35	BB	1150	A	P-O5'	-6.95	1.52	1.59
38	BE	198	A	P-O5'	-6.95	1.52	1.59
39	BF	30	C	O3'-P	-6.95	1.52	1.61
41	BH	99	G	P-O5'	-6.95	1.52	1.59
41	BH	104	U	C4'-C3'	-6.95	1.45	1.53
85	AA	456	A	N9-C8	-6.95	1.32	1.37
85	AA	1526	G	C1'-N9	-6.95	1.37	1.46
85	AA	1676	G	N7-C5	-6.95	1.35	1.39
85	AA	1692	U	O3'-P	-6.95	1.52	1.61
34	BA	1428	G	C5-C4	-6.94	1.33	1.38
35	BB	73	G	N9-C4	-6.94	1.32	1.38
35	BB	638	G	C5-C4	-6.94	1.33	1.38
36	BC	151	G	C5-C4	-6.94	1.33	1.38
37	BD	45	U	P-O5'	-6.94	1.52	1.59
85	AA	1672	G	C2'-C1'	-6.94	1.45	1.53
86	AB	48	C	C2-N3	-6.94	1.30	1.35
34	BA	1241	U	O3'-P	-6.94	1.52	1.61
35	BB	40	C	C1'-N1	-6.94	1.37	1.46
35	BB	1176	G	C1'-N9	-6.94	1.37	1.46
38	BE	49	A	C3'-C2'	-6.94	1.45	1.52
85	AA	157	G	C5-C4	-6.94	1.33	1.38
85	AA	637	U	P-O5'	-6.94	1.52	1.59
85	AA	2184	A	C1'-N9	-6.94	1.37	1.46
35	BB	498	G	C2'-C1'	-6.94	1.45	1.53
35	BB	658	G	N1-C2	-6.94	1.32	1.37
35	BB	1545	U	P-O5'	-6.94	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2239	A	O3'-P	-6.94	1.52	1.61
34	BA	794	G	C6-N1	-6.94	1.34	1.39
35	BB	390	G	C2'-C1'	-6.94	1.45	1.53
85	AA	404	A	C4'-O4'	-6.94	1.36	1.45
25	AR	48	GLY	CA-C	-6.94	1.40	1.51
34	BA	34	U	C3'-C2'	-6.94	1.45	1.52
34	BA	469	C	C4'-C3'	-6.94	1.45	1.53
34	BA	619	U	N3-C4	-6.94	1.32	1.38
34	BA	1032	A	N7-C5	-6.94	1.35	1.39
34	BA	1039	G	C2'-C1'	-6.94	1.45	1.53
34	BA	1214	U	C3'-C2'	-6.94	1.45	1.52
34	BA	1379	G	C2-N3	-6.94	1.27	1.32
35	BB	122	U	N3-C4	-6.94	1.32	1.38
36	BC	139	A	C3'-C2'	-6.94	1.45	1.52
37	BD	7	G	C1'-N9	-6.94	1.37	1.46
85	AA	653	A	O3'-P	-6.94	1.52	1.61
85	AA	794	A	C1'-N9	-6.94	1.37	1.46
85	AA	1109	G	C6-N1	-6.94	1.34	1.39
85	AA	2034	G	N7-C5	-6.94	1.35	1.39
35	BB	1179	C	C3'-C2'	-6.94	1.45	1.52
85	AA	654	A	P-O5'	-6.94	1.52	1.59
85	AA	1502	A	P-O5'	-6.94	1.52	1.59
34	BA	58	A	C5-C4	-6.93	1.33	1.38
34	BA	413	A	N9-C8	-6.93	1.32	1.37
34	BA	498	A	P-O5'	-6.93	1.52	1.59
34	BA	755	G	C5-C4	-6.93	1.33	1.38
34	BA	925	G	C2-N2	-6.93	1.27	1.34
34	BA	1190	A	C4'-C3'	-6.93	1.45	1.53
35	BB	558	U	C3'-C2'	-6.93	1.45	1.52
35	BB	1442	C	C2'-C1'	-6.93	1.45	1.53
37	BD	55	A	O3'-P	-6.93	1.52	1.61
41	BH	44	A	P-O5'	-6.93	1.52	1.59
85	AA	493	A	C3'-C2'	-6.93	1.45	1.52
85	AA	1262	A	C3'-C2'	-6.93	1.45	1.52
85	AA	1519	A	O3'-P	-6.93	1.52	1.61
85	AA	1728	G	O3'-P	-6.93	1.52	1.61
34	BA	757	G	C4'-C3'	-6.93	1.45	1.53
34	BA	1040	G	N7-C5	-6.93	1.35	1.39
34	BA	1076	U	N3-C4	-6.93	1.32	1.38
34	BA	1148	U	O3'-P	-6.93	1.52	1.61
34	BA	1285	G	C5-C4	-6.93	1.33	1.38
35	BB	1070	G	N9-C4	-6.93	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1277	A	C2'-C1'	-6.93	1.45	1.53
35	BB	1351	G	C1'-N9	-6.93	1.37	1.46
85	AA	364	C	O3'-P	-6.93	1.52	1.61
85	AA	1148	G	C2-N2	-6.93	1.27	1.34
85	AA	1269	A	C2'-C1'	-6.93	1.45	1.53
34	BA	77	C	C2-N3	-6.93	1.30	1.35
34	BA	238	C	N1-C2	-6.93	1.33	1.40
41	BH	47	G	C5-C4	-6.93	1.33	1.38
85	AA	937	G	C3'-C2'	-6.93	1.45	1.52
34	BA	420	A	C8-N7	-6.93	1.26	1.31
34	BA	623	U	O3'-P	-6.93	1.52	1.61
34	BA	726	G	N7-C5	-6.93	1.35	1.39
35	BB	423	G	C2-N2	-6.93	1.27	1.34
35	BB	556	U	P-O5'	-6.93	1.52	1.59
35	BB	750	G	C2'-C1'	-6.93	1.45	1.53
40	BG	5	G	C2'-C1'	-6.93	1.45	1.53
40	BG	115	C	O3'-P	-6.93	1.52	1.61
85	AA	115	U	C2-N3	-6.93	1.32	1.37
85	AA	1373	U	O3'-P	-6.93	1.52	1.61
85	AA	1919	G	O3'-P	-6.93	1.52	1.61
85	AA	2127	G	N9-C8	-6.93	1.32	1.37
85	AA	2240	G	C3'-C2'	-6.93	1.45	1.52
34	BA	258	C	C3'-C2'	-6.93	1.45	1.52
34	BA	1188	U	O3'-P	-6.93	1.52	1.61
34	BA	1341	A	N7-C5	-6.93	1.35	1.39
36	BC	56	G	C2'-C1'	-6.93	1.45	1.53
34	BA	350	C	C3'-C2'	-6.93	1.45	1.52
34	BA	394	A	P-O5'	-6.93	1.52	1.59
34	BA	417	A	C2'-C1'	-6.93	1.45	1.53
34	BA	693	G	C4'-C3'	-6.93	1.45	1.53
34	BA	1431	G	C5-C4	-6.93	1.33	1.38
34	BA	1527	G	C3'-C2'	-6.93	1.45	1.52
34	BA	1561	C	C4'-O4'	-6.93	1.36	1.45
35	BB	814	A	C2'-C1'	-6.93	1.45	1.53
35	BB	1107	C	C3'-C2'	-6.93	1.45	1.52
36	BC	52	A	N7-C5	-6.93	1.35	1.39
38	BE	4	A	N7-C5	-6.93	1.35	1.39
85	AA	600	C	C4'-C3'	6.93	1.60	1.53
85	AA	910	G	C2-N3	-6.93	1.27	1.32
85	AA	1928	A	O3'-P	-6.93	1.52	1.61
34	BA	144	C	C3'-C2'	-6.92	1.45	1.52
34	BA	195	G	N9-C8	-6.92	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	975	A	P-O5'	-6.92	1.52	1.59
34	BA	1005	C	O3'-P	-6.92	1.52	1.61
34	BA	1009	G	C6-N1	-6.92	1.34	1.39
34	BA	1041	U	C2-N3	-6.92	1.32	1.37
34	BA	1646	U	C3'-C2'	-6.92	1.45	1.52
36	BC	112	G	N3-C4	-6.92	1.30	1.35
37	BD	75	G	N7-C5	-6.92	1.35	1.39
85	AA	410	A	O3'-P	-6.92	1.52	1.61
85	AA	461	G	C8-N7	-6.92	1.26	1.30
85	AA	1451	U	C4-C5	-6.92	1.37	1.43
34	BA	718	U	O3'-P	-6.92	1.52	1.61
40	BG	109	C	C1'-N1	-6.92	1.37	1.46
85	AA	1704	C	O3'-P	-6.92	1.52	1.61
34	BA	3	G	O4'-C1'	-6.92	1.32	1.41
34	BA	26	C	O3'-P	-6.92	1.52	1.61
34	BA	112	C	C4'-O4'	-6.92	1.36	1.45
34	BA	1078	U	P-O5'	-6.92	1.52	1.59
34	BA	1321	A	C5-C4	-6.92	1.33	1.38
35	BB	878	G	C2'-C1'	-6.92	1.45	1.53
35	BB	957	A	P-O5'	-6.92	1.52	1.59
35	BB	1168	G	C4'-C3'	-6.92	1.45	1.53
35	BB	1188	A	O4'-C1'	-6.92	1.32	1.41
35	BB	1313	C	C4'-C3'	-6.92	1.45	1.53
35	BB	1510	G	C1'-N9	-6.92	1.37	1.46
36	BC	165	U	O3'-P	-6.92	1.52	1.61
37	BD	33	U	C4'-C3'	-6.92	1.45	1.53
40	BG	74	G	N7-C5	-6.92	1.35	1.39
85	AA	499	G	N9-C4	-6.92	1.32	1.38
85	AA	700	U	P-O5'	-6.92	1.52	1.59
85	AA	969	U	C5'-C4'	6.92	1.59	1.51
85	AA	1489	G	P-O5'	-6.92	1.52	1.59
85	AA	2198	G	P-O5'	-6.92	1.52	1.59
34	BA	475	A	C2'-C1'	-6.92	1.45	1.53
85	AA	989	U	C2'-C1'	-6.92	1.45	1.53
34	BA	9	A	P-O5'	-6.92	1.52	1.59
34	BA	1409	A	C5-C4	-6.92	1.33	1.38
34	BA	1491	U	N3-C4	-6.92	1.32	1.38
35	BB	808	U	C2-N3	-6.92	1.32	1.37
35	BB	1153	G	C1'-N9	-6.92	1.37	1.46
35	BB	1298	C	C4-N4	-6.92	1.27	1.33
37	BD	9	C	O3'-P	-6.92	1.52	1.61
37	BD	82	G	C4'-O4'	-6.92	1.36	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2146	G	C6-N1	-6.92	1.34	1.39
34	BA	541	C	C3'-C2'	-6.92	1.45	1.52
34	BA	706	C	C4'-C3'	-6.92	1.45	1.53
34	BA	913	U	P-O5'	-6.92	1.52	1.59
35	BB	511	A	O3'-P	-6.92	1.52	1.61
35	BB	653	G	N9-C4	-6.92	1.32	1.38
35	BB	1143	A	P-O5'	-6.92	1.52	1.59
35	BB	1264	U	O3'-P	-6.92	1.52	1.61
37	BD	25	G	O3'-P	-6.92	1.52	1.61
85	AA	661	C	C2'-C1'	-6.92	1.45	1.53
85	AA	903	G	O3'-P	-6.92	1.52	1.61
34	BA	987	C	O3'-P	-6.92	1.52	1.61
34	BA	1658	G	N9-C8	-6.92	1.33	1.37
34	BA	1701	U	C1'-N1	-6.92	1.37	1.46
35	BB	823	G	C2'-C1'	-6.92	1.45	1.53
35	BB	967	G	C3'-C2'	-6.92	1.45	1.52
35	BB	1293	C	C2-N3	-6.92	1.30	1.35
35	BB	1295	A	C2'-C1'	-6.92	1.45	1.53
85	AA	131	C	O3'-P	-6.92	1.52	1.61
85	AA	389	A	C2'-C1'	-6.92	1.45	1.53
85	AA	673	A	O3'-P	-6.92	1.52	1.61
34	BA	922	C	C3'-C2'	-6.91	1.45	1.52
34	BA	1040	G	N9-C8	-6.91	1.33	1.37
34	BA	1158	A	C5-C4	-6.91	1.33	1.38
34	BA	1300	G	C1'-N9	-6.91	1.37	1.46
34	BA	1341	A	C1'-N9	-6.91	1.37	1.46
34	BA	1555	G	C5-C4	-6.91	1.33	1.38
35	BB	102	G	C5-C4	-6.91	1.33	1.38
35	BB	386	G	O3'-P	-6.91	1.52	1.61
35	BB	1066	G	N7-C5	-6.91	1.35	1.39
35	BB	1227	G	C3'-C2'	-6.91	1.45	1.52
39	BF	47	C	C2-N3	-6.91	1.30	1.35
85	AA	149	A	N9-C4	-6.91	1.33	1.37
85	AA	422	G	O3'-P	-6.91	1.52	1.61
34	BA	65	A	N9-C4	-6.91	1.33	1.37
34	BA	1178	U	C2'-C1'	-6.91	1.45	1.53
85	AA	395	G	N1-C2	-6.91	1.32	1.37
85	AA	1703	A	O3'-P	-6.91	1.52	1.61
34	BA	161	U	P-O5'	-6.91	1.52	1.59
34	BA	745	A	O3'-P	-6.91	1.52	1.61
34	BA	876	C	C1'-N1	-6.91	1.37	1.46
35	BB	391	G	C2'-C1'	-6.91	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1295	A	P-O5'	-6.91	1.52	1.59
35	BB	1335	G	O3'-P	-6.91	1.52	1.61
85	AA	90	A	N7-C5	-6.91	1.35	1.39
85	AA	455	G	O4'-C1'	-6.91	1.32	1.41
85	AA	900	G	O3'-P	-6.91	1.52	1.61
85	AA	1665	G	C3'-C2'	-6.91	1.45	1.52
85	AA	1677	A	C2'-C1'	-6.91	1.45	1.53
34	BA	60	A	C2'-C1'	-6.91	1.45	1.53
34	BA	497	U	N3-C4	-6.91	1.32	1.38
34	BA	828	A	N9-C4	-6.91	1.33	1.37
34	BA	1428	G	C2'-C1'	-6.91	1.45	1.53
35	BB	393	A	C1'-N9	-6.91	1.37	1.46
35	BB	615	A	C8-N7	-6.91	1.26	1.31
35	BB	1250	A	C1'-N9	-6.91	1.37	1.46
36	BC	142	C	O3'-P	-6.91	1.52	1.61
40	BG	175	G	C5'-C4'	6.91	1.59	1.51
34	BA	957	A	C2'-C1'	-6.91	1.45	1.53
37	BD	99	G	N3-C4	-6.91	1.30	1.35
34	BA	495	A	C5-C4	-6.91	1.33	1.38
34	BA	822	U	C2'-C1'	-6.91	1.45	1.53
34	BA	958	G	C2-N2	-6.91	1.27	1.34
34	BA	1055	U	O3'-P	-6.91	1.52	1.61
34	BA	1517	U	O3'-P	-6.91	1.52	1.61
34	BA	1816	G	C8-N7	-6.91	1.26	1.30
85	AA	329	G	N9-C8	-6.91	1.33	1.37
85	AA	1458	G	C3'-C2'	-6.91	1.45	1.52
85	AA	1668	G	C2-N3	-6.91	1.27	1.32
34	BA	93	A	C8-N7	-6.90	1.26	1.31
34	BA	484	A	C1'-N9	-6.90	1.37	1.46
37	BD	106	G	C5-C4	-6.90	1.33	1.38
85	AA	5	U	O3'-P	-6.90	1.52	1.61
85	AA	861	G	P-O5'	-6.90	1.52	1.59
34	BA	3	G	C2'-C1'	-6.90	1.45	1.53
34	BA	789	U	C1'-N1	-6.90	1.37	1.46
34	BA	1162	U	C2-N3	-6.90	1.32	1.37
35	BB	1207	C	C4'-C3'	-6.90	1.45	1.53
35	BB	1251	G	N1-C2	-6.90	1.32	1.37
35	BB	1426	G	C2-N2	-6.90	1.27	1.34
85	AA	1238	U	P-O5'	-6.90	1.52	1.59
85	AA	2111	C	P-O5'	-6.90	1.52	1.59
34	BA	233	U	O3'-P	-6.90	1.52	1.61
34	BA	754	G	C4'-C3'	-6.90	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1642	A	C3'-C2'	-6.90	1.45	1.52
35	BB	455	G	C6-N1	-6.90	1.34	1.39
37	BD	30	A	C2'-C1'	-6.90	1.45	1.53
38	BE	201	A	O3'-P	-6.90	1.52	1.61
40	BG	97	G	C4'-C3'	-6.90	1.45	1.53
85	AA	2198	G	C6-N1	-6.90	1.34	1.39
34	BA	290	G	C2'-C1'	-6.90	1.45	1.53
34	BA	1837	U	C2'-C1'	-6.90	1.45	1.53
35	BB	1485	G	N7-C5	-6.90	1.35	1.39
37	BD	50	A	N9-C4	-6.90	1.33	1.37
85	AA	97	A	C8-N7	-6.90	1.26	1.31
85	AA	171	U	P-O5'	-6.90	1.52	1.59
85	AA	442	G	N9-C4	-6.90	1.32	1.38
34	BA	76	U	C3'-C2'	-6.90	1.45	1.52
34	BA	87	G	C3'-C2'	-6.90	1.45	1.52
34	BA	1028	A	C1'-N9	-6.90	1.37	1.46
34	BA	1514	A	P-O5'	-6.90	1.52	1.59
38	BE	10	G	N3-C4	-6.90	1.30	1.35
67	Bh	134	PHE	CB-CG	-6.90	1.39	1.51
85	AA	338	G	C5'-C4'	-6.90	1.43	1.51
34	BA	24	C	C2'-C1'	-6.90	1.45	1.53
34	BA	1407	C	O3'-P	-6.90	1.52	1.61
35	BB	1309	A	O3'-P	-6.90	1.52	1.61
85	AA	485	A	C4'-C3'	-6.90	1.45	1.53
34	BA	478	G	C8-N7	-6.89	1.26	1.30
35	BB	803	U	C2'-C1'	-6.89	1.45	1.53
36	BC	127	C	O3'-P	-6.89	1.52	1.61
40	BG	58	G	N9-C4	-6.89	1.32	1.38
34	BA	223	U	C3'-C2'	-6.89	1.45	1.52
34	BA	936	A	C1'-N9	-6.89	1.37	1.46
34	BA	1176	C	C2-N3	-6.89	1.30	1.35
34	BA	1655	G	C5-C4	-6.89	1.33	1.38
34	BA	1701	U	O3'-P	-6.89	1.52	1.61
36	BC	155	C	C2-N3	-6.89	1.30	1.35
37	BD	111	U	O3'-P	-6.89	1.52	1.61
85	AA	583	U	P-O5'	-6.89	1.52	1.59
85	AA	619	A	C2'-C1'	-6.89	1.45	1.53
85	AA	1228	A	N3-C4	-6.89	1.30	1.34
34	BA	55	G	N7-C5	-6.89	1.35	1.39
34	BA	466	G	O3'-P	-6.89	1.52	1.61
34	BA	925	G	P-O5'	-6.89	1.52	1.59
34	BA	1167	A	C1'-N9	-6.89	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	368	C	C1'-N1	-6.89	1.37	1.46
35	BB	456	A	C2'-C1'	-6.89	1.45	1.53
36	BC	54	G	C6-N1	-6.89	1.34	1.39
85	AA	2140	U	C2-N3	-6.89	1.32	1.37
34	BA	395	G	O4'-C1'	-6.89	1.32	1.41
34	BA	1115	A	N3-C4	-6.89	1.30	1.34
34	BA	1666	U	C3'-C2'	-6.89	1.45	1.52
35	BB	19	C	O3'-P	-6.89	1.52	1.61
35	BB	956	G	P-O5'	-6.89	1.52	1.59
35	BB	1167	C	C2'-C1'	6.89	1.60	1.53
35	BB	1304	U	O3'-P	-6.89	1.52	1.61
39	BF	19	A	C5'-C4'	6.89	1.59	1.51
85	AA	1152	U	O3'-P	-6.89	1.52	1.61
85	AA	1645	G	C3'-C2'	-6.89	1.45	1.52
34	BA	146	G	C6-N1	-6.89	1.34	1.39
35	BB	563	A	C8-N7	-6.89	1.26	1.31
35	BB	870	C	C4-C5	-6.89	1.37	1.43
41	BH	37	U	C4'-O4'	-6.89	1.36	1.45
34	BA	169	C	O3'-P	-6.89	1.52	1.61
34	BA	234	A	N7-C5	-6.89	1.35	1.39
34	BA	505	U	O3'-P	-6.89	1.52	1.61
34	BA	705	C	C1'-N1	-6.89	1.37	1.46
34	BA	1527	G	C2-N2	-6.89	1.27	1.34
34	BA	1544	G	N7-C5	-6.89	1.35	1.39
35	BB	1052	G	C2'-C1'	-6.89	1.45	1.53
35	BB	1069	C	C3'-C2'	-6.89	1.45	1.52
35	BB	1335	G	C2'-C1'	-6.89	1.45	1.53
35	BB	1406	C	C4-N4	-6.89	1.27	1.33
41	BH	133	U	C5'-C4'	-6.89	1.43	1.51
85	AA	186	U	C3'-C2'	-6.89	1.45	1.52
85	AA	416	U	C1'-N1	-6.89	1.37	1.46
85	AA	423	G	C5-C4	-6.89	1.33	1.38
85	AA	514	U	C1'-N1	-6.89	1.37	1.46
85	AA	1485	G	C2'-C1'	-6.89	1.45	1.53
34	BA	151	A	C4'-O4'	6.88	1.54	1.45
35	BB	67	A	O3'-P	-6.88	1.52	1.61
35	BB	1079	G	C3'-C2'	-6.88	1.45	1.52
35	BB	1079	G	C5-C4	-6.88	1.33	1.38
35	BB	1120	A	O3'-P	-6.88	1.52	1.61
35	BB	1356	G	C1'-N9	-6.88	1.37	1.46
35	BB	1414	A	N7-C5	-6.88	1.35	1.39
85	AA	1097	G	P-O5'	-6.88	1.52	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	781	U	C4'-C3'	-6.88	1.45	1.53
34	BA	857	C	O4'-C1'	-6.88	1.32	1.41
34	BA	1593	U	C3'-C2'	-6.88	1.45	1.52
40	BG	130	G	C3'-C2'	-6.88	1.45	1.52
85	AA	1207	C	O3'-P	-6.88	1.52	1.61
85	AA	1895	C	C4'-C3'	-6.88	1.45	1.53
34	BA	595	U	C4'-C3'	6.88	1.60	1.53
34	BA	1190	A	O3'-P	-6.88	1.52	1.61
34	BA	1329	U	C3'-C2'	-6.88	1.45	1.52
35	BB	28	G	C2-N2	-6.88	1.27	1.34
35	BB	395	U	C2'-C1'	-6.88	1.45	1.53
35	BB	505	G	C5-C4	-6.88	1.33	1.38
35	BB	658	G	O3'-P	-6.88	1.52	1.61
35	BB	809	U	N3-C4	-6.88	1.32	1.38
35	BB	869	G	C2-N2	-6.88	1.27	1.34
35	BB	1415	G	O3'-P	-6.88	1.52	1.61
37	BD	31	U	O3'-P	-6.88	1.52	1.61
40	BG	158	A	C2'-C1'	-6.88	1.45	1.53
85	AA	1016	G	N9-C4	-6.88	1.32	1.38
85	AA	2135	A	O3'-P	-6.88	1.52	1.61
34	BA	493	G	C6-N1	-6.88	1.34	1.39
34	BA	747	G	N9-C8	-6.88	1.33	1.37
34	BA	1194	G	O3'-P	-6.88	1.52	1.61
34	BA	1682	A	N7-C5	-6.88	1.35	1.39
35	BB	1158	C	O3'-P	-6.88	1.52	1.61
37	BD	71	G	N1-C2	-6.88	1.32	1.37
40	BG	76	C	P-O5'	-6.88	1.52	1.59
34	BA	15	G	N7-C5	-6.88	1.35	1.39
34	BA	118	C	C2'-C1'	-6.88	1.45	1.53
34	BA	826	C	C3'-C2'	-6.88	1.45	1.52
34	BA	896	U	C4'-C3'	-6.88	1.45	1.53
34	BA	1193	A	C3'-C2'	-6.88	1.45	1.52
34	BA	1210	A	C2'-C1'	-6.88	1.45	1.53
34	BA	1431	G	N9-C4	-6.88	1.32	1.38
34	BA	1661	U	C4'-O4'	-6.88	1.36	1.45
35	BB	703	U	C2-N3	-6.88	1.32	1.37
35	BB	898	U	P-O5'	-6.88	1.52	1.59
39	BF	33	C	C2'-C1'	-6.88	1.45	1.53
39	BF	52	A	N7-C5	-6.88	1.35	1.39
85	AA	285	C	C2'-C1'	-6.88	1.45	1.53
85	AA	1353	U	C5'-C4'	6.88	1.59	1.51
85	AA	1464	G	O4'-C1'	-6.88	1.32	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	885	A	N9-C4	-6.88	1.33	1.37
34	BA	953	G	C5-C4	-6.88	1.33	1.38
35	BB	604	C	C2'-C1'	-6.88	1.45	1.53
35	BB	666	A	C5-C4	-6.88	1.33	1.38
36	BC	57	C	P-O5'	-6.88	1.52	1.59
40	BG	163	G	C8-N7	-6.88	1.26	1.30
85	AA	124	A	P-O5'	-6.88	1.52	1.59
85	AA	438	G	C5-C4	-6.88	1.33	1.38
85	AA	1207	C	C5'-C4'	-6.88	1.43	1.51
85	AA	983	A	N7-C5	-6.88	1.35	1.39
34	BA	355	U	C3'-C2'	-6.87	1.45	1.52
34	BA	371	U	C1'-N1	-6.87	1.37	1.46
34	BA	941	G	P-O5'	-6.87	1.52	1.59
34	BA	1465	C	C2-N3	-6.87	1.30	1.35
34	BA	1557	G	P-O5'	-6.87	1.52	1.59
35	BB	444	U	C1'-N1	-6.87	1.37	1.46
85	AA	139	G	P-O5'	-6.87	1.52	1.59
85	AA	851	G	C8-N7	-6.87	1.26	1.30
85	AA	2017	U	O3'-P	-6.87	1.52	1.61
34	BA	44	U	C4'-C3'	-6.87	1.45	1.53
34	BA	955	G	O3'-P	-6.87	1.52	1.61
35	BB	531	U	C2-N3	-6.87	1.32	1.37
35	BB	550	G	O3'-P	-6.87	1.52	1.61
35	BB	1427	A	C6-N6	-6.87	1.28	1.33
38	BE	10	G	C1'-N9	6.87	1.59	1.48
85	AA	651	G	O3'-P	-6.87	1.52	1.61
85	AA	1067	G	O3'-P	-6.87	1.52	1.61
85	AA	1529	A	O3'-P	-6.87	1.52	1.61
85	AA	1658	G	P-O5'	-6.87	1.52	1.59
34	BA	590	U	C2-N3	-6.87	1.32	1.37
34	BA	1665	G	C2-N2	-6.87	1.27	1.34
35	BB	385	C	C3'-C2'	-6.87	1.45	1.52
35	BB	588	A	P-O5'	-6.87	1.52	1.59
35	BB	795	A	P-O5'	-6.87	1.52	1.59
35	BB	1496	C	C3'-C2'	-6.87	1.45	1.52
40	BG	29	U	C3'-C2'	-6.87	1.45	1.52
85	AA	794	A	C8-N7	-6.87	1.26	1.31
85	AA	1140	G	N9-C4	-6.87	1.32	1.38
34	BA	1004	U	C2-N3	-6.87	1.32	1.37
35	BB	72	G	C2'-C1'	-6.87	1.45	1.53
35	BB	784	C	C2-N3	-6.87	1.30	1.35
35	BB	1204	C	C3'-C2'	-6.87	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1494	G	O3'-P	-6.87	1.52	1.61
85	AA	1257	A	O3'-P	-6.87	1.52	1.61
34	BA	34	U	C2'-C1'	-6.87	1.45	1.53
34	BA	50	G	C6-N1	-6.87	1.34	1.39
34	BA	955	G	C3'-C2'	-6.87	1.45	1.52
34	BA	1070	G	C3'-C2'	-6.87	1.45	1.52
35	BB	602	G	N9-C8	-6.87	1.33	1.37
36	BC	164	G	C4'-O4'	-6.87	1.36	1.45
85	AA	862	U	C2-N3	-6.87	1.32	1.37
85	AA	1143	C	C4'-C3'	-6.87	1.45	1.53
85	AA	1621	U	C2'-C1'	-6.87	1.45	1.53
34	BA	235	C	O3'-P	-6.87	1.52	1.61
34	BA	602	G	N3-C4	-6.87	1.30	1.35
34	BA	1399	A	O3'-P	-6.87	1.52	1.61
35	BB	2	C	C2'-C1'	-6.87	1.45	1.53
35	BB	659	C	C4'-O4'	-6.87	1.36	1.45
40	BG	7	U	O3'-P	-6.87	1.52	1.61
40	BG	59	G	C2'-C1'	-6.87	1.45	1.53
40	BG	129	G	C3'-C2'	-6.87	1.45	1.52
41	BH	123	G	C3'-C2'	-6.87	1.45	1.52
85	AA	451	G	O3'-P	-6.87	1.52	1.61
85	AA	889	G	N3-C4	-6.87	1.30	1.35
85	AA	1282	A	C1'-N9	-6.87	1.37	1.46
85	AA	1283	C	C4'-O4'	-6.87	1.36	1.45
85	AA	2176	U	C1'-N1	-6.87	1.37	1.46
34	BA	274	C	C2-N3	-6.86	1.30	1.35
34	BA	418	G	N9-C4	-6.86	1.32	1.38
34	BA	859	G	C5-C4	-6.86	1.33	1.38
34	BA	1515	U	P-O5'	-6.86	1.52	1.59
34	BA	1833	G	N7-C5	-6.86	1.35	1.39
35	BB	554	C	P-O5'	-6.86	1.52	1.59
35	BB	1434	G	C1'-N9	-6.86	1.37	1.46
35	BB	1514	G	O3'-P	-6.86	1.52	1.61
38	BE	130	G	C2-N2	-6.86	1.27	1.34
40	BG	128	U	C2'-C1'	-6.86	1.45	1.53
41	BH	54	U	C2'-C1'	-6.86	1.45	1.53
85	AA	116	G	C1'-N9	-6.86	1.37	1.46
85	AA	689	U	C3'-C2'	-6.86	1.45	1.52
34	BA	1103	G	C3'-C2'	-6.86	1.45	1.52
34	BA	1406	U	O3'-P	-6.86	1.52	1.61
35	BB	1253	U	C2-N3	-6.86	1.32	1.37
35	BB	1324	C	O3'-P	-6.86	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	94	C	C2'-C1'	-6.86	1.45	1.53
39	BF	42	G	P-O5'	-6.86	1.52	1.59
40	BG	115	C	C1'-N1	-6.86	1.37	1.46
85	AA	30	G	C3'-C2'	-6.86	1.45	1.52
34	BA	23	A	C6-N1	-6.86	1.30	1.35
34	BA	384	U	O3'-P	-6.86	1.52	1.61
34	BA	596	G	C4'-C3'	6.86	1.60	1.53
34	BA	1644	A	P-O5'	-6.86	1.52	1.59
35	BB	1139	A	C1'-N9	-6.86	1.37	1.46
37	BD	62	A	C1'-N9	-6.86	1.37	1.46
39	BF	50	C	O3'-P	-6.86	1.52	1.61
85	AA	385	A	C4'-O4'	-6.86	1.36	1.45
85	AA	750	A	P-O5'	-6.86	1.52	1.59
34	BA	949	C	P-O5'	-6.86	1.52	1.59
35	BB	683	U	P-O5'	-6.86	1.52	1.59
35	BB	1091	C	P-O5'	-6.86	1.52	1.59
38	BE	8	G	C1'-N9	-6.86	1.37	1.46
34	BA	291	C	C5'-C4'	-6.86	1.43	1.51
34	BA	599	U	C4'-C3'	-6.86	1.45	1.53
34	BA	1291	A	P-O5'	-6.86	1.52	1.59
34	BA	1795	A	C3'-C2'	-6.86	1.45	1.52
35	BB	579	A	P-O5'	-6.86	1.52	1.59
37	BD	37	G	C2'-C1'	-6.86	1.45	1.53
39	BF	43	U	P-O5'	-6.86	1.52	1.59
85	AA	186	U	C2'-C1'	-6.86	1.45	1.53
85	AA	709	A	C5-C4	-6.86	1.33	1.38
85	AA	1262	A	P-O5'	-6.86	1.52	1.59
85	AA	1577	G	N7-C5	-6.86	1.35	1.39
85	AA	1648	G	C2'-C1'	-6.86	1.45	1.53
85	AA	2139	G	C1'-N9	-6.86	1.37	1.46
34	BA	315	U	N3-C4	-6.86	1.32	1.38
34	BA	1724	G	C5'-C4'	6.86	1.59	1.51
35	BB	1141	A	C3'-C2'	-6.86	1.45	1.52
35	BB	1508	G	C1'-N9	-6.86	1.37	1.46
85	AA	15	U	C1'-N1	-6.86	1.37	1.46
85	AA	313	A	C8-N7	-6.86	1.26	1.31
85	AA	470	C	C2'-C1'	-6.86	1.45	1.53
34	BA	1124	U	P-O5'	-6.85	1.52	1.59
40	BG	126	G	C6-N1	-6.85	1.34	1.39
85	AA	408	C	C2'-C1'	-6.85	1.45	1.53
85	AA	1155	A	C5-C4	-6.85	1.33	1.38
85	AA	1751	G	O3'-P	-6.85	1.52	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	81	C	N1-C6	-6.85	1.33	1.37
34	BA	146	G	C2-N2	-6.85	1.27	1.34
34	BA	371	U	C2'-C1'	-6.85	1.45	1.53
34	BA	1203	G	C4'-C3'	-6.85	1.45	1.53
34	BA	1544	G	O3'-P	-6.85	1.52	1.61
34	BA	1642	A	C5'-C4'	-6.85	1.43	1.51
34	BA	1654	G	C6-N1	-6.85	1.34	1.39
34	BA	1836	A	C3'-C2'	-6.85	1.45	1.52
35	BB	80	C	C2-N3	-6.85	1.30	1.35
35	BB	566	A	N7-C5	-6.85	1.35	1.39
38	BE	63	C	C2-N3	-6.85	1.30	1.35
40	BG	7	U	P-O5'	-6.85	1.52	1.59
72	Bm	40	LEU	C-N	-6.85	1.21	1.34
85	AA	533	C	C3'-C2'	-6.85	1.45	1.52
85	AA	641	A	C4'-C3'	-6.85	1.45	1.53
85	AA	687	G	C6-N1	-6.85	1.34	1.39
85	AA	1251	G	O3'-P	-6.85	1.52	1.61
86	AB	31	A	P-O5'	-6.85	1.52	1.59
34	BA	69	C	P-O5'	-6.85	1.52	1.59
34	BA	943	G	C3'-C2'	-6.85	1.45	1.52
34	BA	1242	A	C2'-C1'	-6.85	1.45	1.53
35	BB	1491	G	P-O5'	-6.85	1.52	1.59
41	BH	26	C	C4-C5	-6.85	1.37	1.43
85	AA	1354	A	N9-C4	6.85	1.42	1.37
85	AA	2069	A	C1'-N9	-6.85	1.37	1.46
85	AA	2109	G	C2'-C1'	-6.85	1.45	1.53
34	BA	31	A	C2'-C1'	-6.85	1.45	1.53
34	BA	51	C	C2'-C1'	-6.85	1.45	1.53
34	BA	351	A	C5-C4	-6.85	1.33	1.38
34	BA	431	A	N7-C5	-6.85	1.35	1.39
34	BA	817	U	O3'-P	-6.85	1.52	1.61
35	BB	1057	G	C1'-N9	-6.85	1.37	1.46
35	BB	1117	G	C2'-C1'	-6.85	1.45	1.53
35	BB	1244	U	C2'-C1'	-6.85	1.45	1.53
85	AA	310	U	C2-N3	-6.85	1.32	1.37
85	AA	407	G	C5-C4	-6.85	1.33	1.38
85	AA	605	A	C1'-N9	-6.85	1.37	1.46
34	BA	32	A	O3'-P	-6.85	1.52	1.61
34	BA	1484	A	C3'-C2'	-6.85	1.45	1.52
34	BA	1514	A	C1'-N9	-6.85	1.37	1.46
37	BD	79	G	N3-C4	-6.85	1.30	1.35
38	BE	144	A	C2'-C1'	-6.85	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	243	A	N9-C4	-6.85	1.33	1.37
85	AA	1509	A	C3'-C2'	-6.85	1.45	1.52
85	AA	1655	G	C3'-C2'	-6.85	1.45	1.52
85	AA	1918	U	P-O5'	-6.85	1.52	1.59
34	BA	126	G	O3'-P	-6.85	1.52	1.61
34	BA	1154	U	O3'-P	-6.85	1.52	1.61
34	BA	1202	G	N1-C2	-6.85	1.32	1.37
34	BA	1226	G	C2'-C1'	-6.85	1.45	1.53
34	BA	1605	G	C2'-C1'	-6.85	1.45	1.53
35	BB	631	G	C2'-C1'	-6.85	1.45	1.53
35	BB	1078	U	C5'-C4'	-6.85	1.43	1.51
35	BB	1131	C	C4'-O4'	-6.85	1.36	1.45
85	AA	157	G	O3'-P	-6.85	1.52	1.61
34	BA	147	U	N1-C2	-6.84	1.32	1.38
34	BA	472	G	C4'-C3'	-6.84	1.45	1.53
35	BB	108	G	N3-C4	-6.84	1.30	1.35
35	BB	468	U	O3'-P	-6.84	1.52	1.61
35	BB	643	G	C2'-C1'	-6.84	1.45	1.53
35	BB	1372	G	C6-N1	-6.84	1.34	1.39
40	BG	160	C	O3'-P	-6.84	1.52	1.61
85	AA	642	G	O3'-P	-6.84	1.52	1.61
85	AA	1991	C	P-O5'	-6.84	1.52	1.59
85	AA	2110	U	C2'-C1'	-6.84	1.45	1.53
34	BA	908	G	C2'-C1'	-6.84	1.45	1.53
35	BB	129	U	C1'-N1	-6.84	1.37	1.46
35	BB	430	A	C5-C4	-6.84	1.33	1.38
35	BB	1336	G	C2'-C1'	-6.84	1.45	1.53
41	BH	39	G	N9-C8	-6.84	1.33	1.37
34	BA	617	G	C3'-C2'	-6.84	1.45	1.52
34	BA	802	G	C2'-C1'	-6.84	1.45	1.53
34	BA	899	G	C2-N3	-6.84	1.27	1.32
34	BA	1270	G	C2-N2	-6.84	1.27	1.34
34	BA	1612	C	C2-N3	-6.84	1.30	1.35
35	BB	100	A	C5-C4	-6.84	1.33	1.38
35	BB	1031	G	C4'-C3'	-6.84	1.45	1.53
35	BB	1408	G	O3'-P	-6.84	1.52	1.61
35	BB	1497	C	C2'-C1'	-6.84	1.45	1.53
36	BC	153	C	C4-C5	-6.84	1.37	1.43
41	BH	35	G	C8-N7	-6.84	1.26	1.30
85	AA	807	A	C2'-C1'	-6.84	1.45	1.53
34	BA	147	U	C1'-N1	-6.84	1.37	1.46
34	BA	825	G	C2'-C1'	-6.84	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1071	G	C4'-C3'	-6.84	1.45	1.53
34	BA	1160	U	C4'-C3'	-6.84	1.45	1.53
34	BA	1181	G	P-O5'	-6.84	1.52	1.59
35	BB	44	C	C2-N3	-6.84	1.30	1.35
37	BD	22	A	O3'-P	-6.84	1.52	1.61
85	AA	557	G	N9-C8	-6.84	1.33	1.37
85	AA	708	G	C2-N2	-6.84	1.27	1.34
34	BA	714	G	C4'-C3'	-6.84	1.45	1.53
34	BA	904	G	N9-C4	-6.84	1.32	1.38
34	BA	1460	U	O3'-P	-6.84	1.52	1.61
36	BC	101	U	C3'-C2'	-6.84	1.45	1.52
40	BG	172	C	C4'-C3'	-6.84	1.45	1.53
85	AA	352	G	O3'-P	-6.84	1.52	1.61
34	BA	410	G	P-O5'	-6.84	1.52	1.59
34	BA	444	A	O3'-P	-6.84	1.52	1.61
34	BA	490	A	O3'-P	-6.84	1.52	1.61
34	BA	860	G	C2-N2	-6.84	1.27	1.34
35	BB	1201	G	C2'-C1'	-6.84	1.45	1.53
38	BE	96	G	O3'-P	-6.84	1.52	1.61
40	BG	76	C	C4'-O4'	-6.84	1.36	1.45
41	BH	125	U	C3'-C2'	-6.84	1.45	1.52
85	AA	9	U	N3-C4	-6.84	1.32	1.38
85	AA	136	U	P-O5'	-6.84	1.52	1.59
85	AA	535	G	C2'-C1'	-6.84	1.45	1.53
85	AA	636	G	C2'-C1'	-6.84	1.45	1.53
85	AA	816	A	C5'-C4'	6.84	1.59	1.51
85	AA	1511	C	O3'-P	-6.84	1.52	1.61
85	AA	1611	A	N9-C4	-6.84	1.33	1.37
34	BA	890	G	O3'-P	-6.83	1.52	1.61
34	BA	941	G	O3'-P	-6.83	1.52	1.61
35	BB	117	A	C4'-C3'	-6.83	1.45	1.53
35	BB	579	A	N7-C5	-6.83	1.35	1.39
35	BB	600	C	C2'-C1'	-6.83	1.45	1.53
35	BB	607	G	C2-N3	-6.83	1.27	1.32
35	BB	1318	U	O3'-P	-6.83	1.52	1.61
85	AA	277	G	P-O5'	-6.83	1.52	1.59
34	BA	1001	G	N9-C4	-6.83	1.32	1.38
34	BA	1213	A	C3'-C2'	-6.83	1.45	1.52
34	BA	1415	C	O3'-P	-6.83	1.52	1.61
34	BA	1795	A	O3'-P	-6.83	1.52	1.61
35	BB	1022	C	C1'-N1	-6.83	1.37	1.46
41	BH	128	G	C2'-C1'	-6.83	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	128	G	C5-C4	-6.83	1.33	1.38
85	AA	120	C	O3'-P	-6.83	1.52	1.61
85	AA	280	U	O3'-P	-6.83	1.52	1.61
85	AA	678	A	C3'-C2'	-6.83	1.45	1.52
85	AA	959	C	P-O5'	-6.83	1.52	1.59
85	AA	1136	A	N9-C4	-6.83	1.33	1.37
85	AA	1499	G	C3'-C2'	-6.83	1.45	1.52
34	BA	423	G	C3'-C2'	-6.83	1.45	1.52
34	BA	765	U	C4'-O4'	6.83	1.54	1.45
34	BA	1033	G	C2-N3	-6.83	1.27	1.32
34	BA	1587	C	C3'-C2'	-6.83	1.45	1.52
34	BA	1822	U	C2'-C1'	-6.83	1.45	1.53
35	BB	1059	U	C2'-C1'	-6.83	1.45	1.53
37	BD	56	G	O3'-P	-6.83	1.52	1.61
40	BG	21	C	O3'-P	-6.83	1.52	1.61
40	BG	90	G	O3'-P	-6.83	1.52	1.61
41	BH	5	G	P-O5'	-6.83	1.52	1.59
85	AA	789	A	O3'-P	-6.83	1.52	1.61
85	AA	837	C	O3'-P	-6.83	1.52	1.61
85	AA	978	U	O3'-P	-6.83	1.52	1.61
35	BB	432	C	C3'-C2'	-6.83	1.45	1.52
35	BB	839	G	C6-N1	-6.83	1.34	1.39
35	BB	1403	G	C4'-C3'	-6.83	1.45	1.53
38	BE	27	A	C5'-C4'	-6.83	1.43	1.51
34	BA	775	C	C2-N3	-6.83	1.30	1.35
34	BA	1244	G	P-O5'	-6.83	1.52	1.59
35	BB	85	A	C5-C4	-6.83	1.33	1.38
35	BB	439	G	N7-C5	-6.83	1.35	1.39
35	BB	1260	A	C2'-C1'	-6.83	1.45	1.53
37	BD	112	U	N3-C4	-6.83	1.32	1.38
38	BE	109	C	C2'-C1'	-6.83	1.45	1.53
85	AA	557	G	C1'-N9	-6.83	1.37	1.46
35	BB	118	A	C2'-C1'	-6.83	1.45	1.53
35	BB	1022	C	P-O5'	-6.83	1.52	1.59
35	BB	1440	A	C1'-N9	-6.83	1.37	1.46
38	BE	66	A	C8-N7	-6.83	1.26	1.31
38	BE	199	A	C1'-N9	-6.83	1.37	1.46
85	AA	367	A	C3'-C2'	-6.83	1.45	1.52
85	AA	444	U	O3'-P	-6.83	1.52	1.61
85	AA	1186	C	C4-N4	-6.83	1.27	1.33
34	BA	326	A	C4'-O4'	-6.83	1.36	1.45
34	BA	1516	G	C2-N2	-6.83	1.27	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1522	G	N7-C5	-6.83	1.35	1.39
34	BA	1532	G	C1'-N9	-6.83	1.37	1.46
35	BB	624	A	N9-C4	-6.83	1.33	1.37
35	BB	804	U	N3-C4	-6.83	1.32	1.38
35	BB	1126	A	N3-C4	-6.83	1.30	1.34
85	AA	156	G	C2'-C1'	-6.83	1.45	1.53
85	AA	258	G	O3'-P	-6.83	1.52	1.61
85	AA	579	U	C4'-C3'	-6.83	1.45	1.53
85	AA	642	G	N1-C2	-6.83	1.32	1.37
34	BA	677	U	N1-C2	-6.82	1.32	1.38
34	BA	686	U	C3'-C2'	-6.82	1.45	1.52
34	BA	1033	G	C5-C6	-6.82	1.35	1.42
34	BA	1696	G	C2-N2	-6.82	1.27	1.34
35	BB	1116	U	C3'-C2'	-6.82	1.45	1.52
35	BB	1380	G	N9-C4	-6.82	1.32	1.38
36	BC	11	G	C2-N2	-6.82	1.27	1.34
39	BF	44	C	O3'-P	-6.82	1.52	1.61
39	BF	71	G	C5-C4	-6.82	1.33	1.38
85	AA	244	G	N7-C5	-6.82	1.35	1.39
85	AA	407	G	C1'-N9	-6.82	1.37	1.46
85	AA	896	C	C4'-C3'	-6.82	1.45	1.53
85	AA	993	G	C2'-C1'	-6.82	1.45	1.53
85	AA	1839	G	C2'-C1'	-6.82	1.45	1.53
85	AA	1927	G	N9-C4	-6.82	1.32	1.38
85	AA	1932	C	C4'-C3'	6.82	1.60	1.53
34	BA	48	C	C2'-C1'	-6.82	1.45	1.53
34	BA	706	C	C4'-O4'	-6.82	1.36	1.45
34	BA	1532	G	O3'-P	-6.82	1.52	1.61
35	BB	1077	C	O3'-P	-6.82	1.52	1.61
35	BB	1407	U	C4'-O4'	-6.82	1.36	1.45
35	BB	1540	U	C3'-C2'	-6.82	1.45	1.52
34	BA	96	G	C3'-C2'	-6.82	1.45	1.52
34	BA	607	C	P-O5'	-6.82	1.52	1.59
34	BA	888	G	N1-C2	-6.82	1.32	1.37
34	BA	1681	U	N3-C4	-6.82	1.32	1.38
35	BB	1485	G	O3'-P	-6.82	1.52	1.61
38	BE	166	G	O3'-P	-6.82	1.52	1.61
85	AA	385	A	C3'-C2'	-6.82	1.45	1.52
34	BA	740	A	C5-C6	6.82	1.47	1.41
34	BA	1039	G	C5-C4	-6.82	1.33	1.38
34	BA	1613	G	C5-C6	-6.82	1.35	1.42
34	BA	1719	G	C2-N2	-6.82	1.27	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	651	G	C6-N1	-6.82	1.34	1.39
85	AA	428	G	C1'-N9	-6.82	1.37	1.46
34	BA	442	G	O3'-P	-6.82	1.52	1.61
34	BA	909	G	P-O5'	-6.82	1.52	1.59
34	BA	1470	G	C1'-N9	-6.82	1.37	1.46
35	BB	1443	C	C4'-C3'	-6.82	1.45	1.53
40	BG	57	A	C6-N1	6.82	1.40	1.35
85	AA	1266	C	C2'-C1'	-6.82	1.45	1.53
34	BA	793	A	C8-N7	-6.82	1.26	1.31
34	BA	1461	A	N7-C5	-6.82	1.35	1.39
34	BA	1477	C	C2'-C1'	-6.82	1.45	1.53
35	BB	439	G	P-O5'	-6.82	1.52	1.59
35	BB	1207	C	C2'-C1'	-6.82	1.45	1.53
38	BE	194	A	C1'-N9	-6.82	1.37	1.46
40	BG	5	G	C6-N1	-6.82	1.34	1.39
40	BG	68	U	C2'-C1'	-6.82	1.45	1.53
85	AA	366	A	C4'-C3'	-6.82	1.45	1.53
85	AA	419	A	C8-N7	-6.82	1.26	1.31
85	AA	848	C	P-O5'	-6.82	1.52	1.59
34	BA	57	A	C8-N7	-6.81	1.26	1.31
34	BA	1410	C	C2'-C1'	-6.81	1.45	1.53
35	BB	431	U	C3'-C2'	-6.81	1.45	1.52
35	BB	486	G	C6-N1	-6.81	1.34	1.39
35	BB	1334	C	N1-C6	-6.81	1.33	1.37
40	BG	15	G	N3-C4	-6.81	1.30	1.35
41	BH	68	G	C2'-C1'	-6.81	1.45	1.53
85	AA	1904	C	N1-C2	-6.81	1.33	1.40
34	BA	395	G	O3'-P	-6.81	1.52	1.61
34	BA	796	G	C5-C6	-6.81	1.35	1.42
35	BB	468	U	C2-N3	-6.81	1.32	1.37
35	BB	784	C	O3'-P	-6.81	1.52	1.61
38	BE	40	C	C4'-C3'	-6.81	1.45	1.53
40	BG	156	G	O3'-P	-6.81	1.52	1.61
85	AA	400	G	O3'-P	-6.81	1.52	1.61
85	AA	743	C	C3'-C2'	-6.81	1.45	1.52
85	AA	745	C	N1-C6	-6.81	1.33	1.37
85	AA	1268	C	C4'-C3'	-6.81	1.45	1.53
85	AA	1464	G	C2'-C1'	-6.81	1.45	1.53
85	AA	1927	G	C2-N2	-6.81	1.27	1.34
85	AA	2122	A	P-O5'	-6.81	1.52	1.59
34	BA	409	A	N3-C4	-6.81	1.30	1.34
35	BB	535	U	C3'-C2'	-6.81	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	4	U	C2'-C1'	-6.81	1.45	1.53
37	BD	41	G	C2'-C1'	-6.81	1.45	1.53
39	BF	12	U	C5'-C4'	-6.81	1.43	1.51
34	BA	911	G	C2'-C1'	-6.81	1.45	1.53
34	BA	1399	A	P-O5'	-6.81	1.52	1.59
35	BB	130	G	C6-N1	-6.81	1.34	1.39
35	BB	496	C	P-O5'	-6.81	1.52	1.59
35	BB	813	C	C3'-C2'	-6.81	1.45	1.52
35	BB	1137	G	N1-C2	-6.81	1.32	1.37
34	BA	202	A	C2'-C1'	-6.81	1.45	1.53
34	BA	253	U	C4'-C3'	-6.81	1.45	1.53
34	BA	706	C	O3'-P	-6.81	1.52	1.61
34	BA	796	G	O3'-P	-6.81	1.52	1.61
35	BB	54	U	N1-C2	-6.81	1.32	1.38
35	BB	1427	A	C2'-C1'	-6.81	1.45	1.53
38	BE	193	A	C8-N7	-6.81	1.26	1.31
39	BF	45	G	P-O5'	-6.81	1.52	1.59
40	BG	6	A	C5-C4	-6.81	1.33	1.38
85	AA	93	G	C2-N2	-6.81	1.27	1.34
34	BA	740	A	C6-N1	6.81	1.40	1.35
34	BA	1280	A	C5-C4	-6.81	1.33	1.38
34	BA	1578	A	C3'-C2'	-6.81	1.45	1.52
35	BB	450	A	N7-C5	-6.81	1.35	1.39
35	BB	661	G	C3'-C2'	-6.81	1.45	1.52
36	BC	45	C	C3'-C2'	-6.81	1.45	1.52
36	BC	147	G	C2-N2	-6.81	1.27	1.34
85	AA	1524	A	N7-C5	-6.81	1.35	1.39
34	BA	1224	A	C1'-N9	-6.80	1.37	1.46
35	BB	702	G	N7-C5	-6.80	1.35	1.39
85	AA	413	G	C3'-C2'	-6.80	1.45	1.52
85	AA	791	C	O3'-P	-6.80	1.52	1.61
85	AA	1508	A	O3'-P	-6.80	1.52	1.61
85	AA	1550	C	O3'-P	-6.80	1.52	1.61
34	BA	1551	G	C6-N1	-6.80	1.34	1.39
34	BA	1652	G	O3'-P	-6.80	1.52	1.61
35	BB	1239	A	C2'-C1'	-6.80	1.45	1.53
35	BB	1284	U	C2'-C1'	-6.80	1.45	1.53
85	AA	2087	C	C3'-C2'	-6.80	1.45	1.52
34	BA	7	U	O3'-P	-6.80	1.52	1.61
34	BA	469	C	P-O5'	-6.80	1.52	1.59
34	BA	596	G	C6-N1	-6.80	1.34	1.39
34	BA	1544	G	C6-N1	-6.80	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1830	A	N7-C5	-6.80	1.35	1.39
35	BB	634	A	C3'-C2'	-6.80	1.45	1.52
35	BB	1181	A	C2'-C1'	-6.80	1.45	1.53
37	BD	99	G	N9-C8	-6.80	1.33	1.37
41	BH	1	U	C2'-C1'	-6.80	1.45	1.53
41	BH	128	G	C8-N7	-6.80	1.26	1.30
85	AA	1868	G	N9-C4	-6.80	1.32	1.38
85	AA	1978	G	C5-C6	-6.80	1.35	1.42
85	AA	2145	G	C6-N1	-6.80	1.34	1.39
34	BA	417	A	C5-C4	-6.80	1.33	1.38
34	BA	810	A	C2'-C1'	-6.80	1.45	1.53
34	BA	1023	G	C5-C6	-6.80	1.35	1.42
34	BA	1170	A	C1'-N9	-6.80	1.37	1.46
34	BA	1470	G	C2'-C1'	-6.80	1.45	1.53
34	BA	1804	A	C2'-C1'	-6.80	1.45	1.53
35	BB	1404	A	C5-C6	-6.80	1.34	1.41
39	BF	48	G	C5-C4	-6.80	1.33	1.38
85	AA	189	G	C6-N1	-6.80	1.34	1.39
85	AA	1198	U	C2-N3	-6.80	1.32	1.37
34	BA	418	G	N3-C4	-6.80	1.30	1.35
34	BA	1221	A	C3'-C2'	-6.80	1.45	1.52
34	BA	1641	G	C1'-N9	-6.80	1.37	1.46
34	BA	1816	G	C6-N1	-6.80	1.34	1.39
35	BB	787	A	C4'-O4'	-6.80	1.36	1.45
35	BB	1342	C	C3'-C2'	-6.80	1.45	1.52
38	BE	39	U	O3'-P	-6.80	1.52	1.61
41	BH	19	G	N1-C2	-6.80	1.32	1.37
85	AA	1510	A	P-O5'	-6.80	1.52	1.59
34	BA	220	U	C2-N3	-6.80	1.32	1.37
34	BA	541	C	O3'-P	-6.80	1.52	1.61
34	BA	1522	G	O3'-P	-6.80	1.52	1.61
35	BB	621	C	O3'-P	-6.80	1.52	1.61
36	BC	107	C	O3'-P	-6.80	1.52	1.61
85	AA	1200	A	P-O5'	-6.80	1.52	1.59
85	AA	1471	G	C8-N7	-6.80	1.26	1.30
36	BC	68	A	N3-C4	-6.79	1.30	1.34
85	AA	1706	A	P-O5'	-6.79	1.52	1.59
85	AA	1734	A	C4'-C3'	6.79	1.60	1.53
35	BB	1159	U	O4'-C1'	-6.79	1.32	1.41
36	BC	19	A	C5-C4	-6.79	1.33	1.38
36	BC	147	G	N7-C5	-6.79	1.35	1.39
37	BD	70	C	C3'-C2'	-6.79	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	60	C	N3-C4	-6.79	1.29	1.33
39	BF	18	U	P-O5'	-6.79	1.52	1.59
85	AA	167	A	P-O5'	-6.79	1.52	1.59
85	AA	521	A	O3'-P	-6.79	1.52	1.61
85	AA	698	G	N7-C5	-6.79	1.35	1.39
85	AA	907	G	O3'-P	-6.79	1.52	1.61
85	AA	1200	A	N7-C5	-6.79	1.35	1.39
85	AA	1479	U	C2'-C1'	-6.79	1.45	1.53
85	AA	1513	U	P-O5'	-6.79	1.52	1.59
85	AA	1706	A	N9-C4	-6.79	1.33	1.37
34	BA	195	G	N7-C5	-6.79	1.35	1.39
34	BA	815	C	C3'-C2'	-6.79	1.45	1.52
34	BA	943	G	N9-C4	-6.79	1.32	1.38
34	BA	991	U	N3-C4	-6.79	1.32	1.38
34	BA	1024	A	N7-C5	-6.79	1.35	1.39
34	BA	1278	A	N7-C5	-6.79	1.35	1.39
35	BB	1153	G	C6-N1	-6.79	1.34	1.39
35	BB	1406	C	C4'-C3'	-6.79	1.45	1.53
38	BE	24	G	O3'-P	-6.79	1.52	1.61
57	BX	82	PHE	CA-CB	6.79	1.68	1.53
85	AA	83	U	O3'-P	-6.79	1.53	1.61
85	AA	280	U	C2-N3	-6.79	1.32	1.37
85	AA	534	A	N1-C2	-6.79	1.28	1.34
85	AA	676	U	O3'-P	-6.79	1.53	1.61
85	AA	1509	A	C6-N1	-6.79	1.30	1.35
35	BB	399	A	C3'-C2'	-6.79	1.45	1.52
35	BB	1055	G	P-O5'	-6.79	1.52	1.59
40	BG	10	U	C2-N3	-6.79	1.32	1.37
85	AA	670	C	C2'-C1'	-6.79	1.45	1.53
85	AA	1294	U	P-O5'	-6.79	1.52	1.59
85	AA	2076	C	C3'-C2'	-6.79	1.45	1.52
34	BA	104	A	C8-N7	-6.79	1.26	1.31
34	BA	313	C	C2-N3	-6.79	1.30	1.35
34	BA	600	G	C2-N2	-6.79	1.27	1.34
34	BA	1271	C	C2'-C1'	-6.79	1.45	1.53
34	BA	1526	C	C3'-C2'	-6.79	1.45	1.52
85	AA	424	A	C6-N1	-6.79	1.30	1.35
85	AA	466	A	O3'-P	-6.79	1.53	1.61
85	AA	630	A	N7-C5	-6.79	1.35	1.39
85	AA	1133	C	C4'-C3'	-6.79	1.45	1.53
85	AA	1295	G	C3'-C2'	-6.79	1.45	1.52
34	BA	333	A	C5-C6	-6.79	1.34	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	373	G	C3'-C2'	-6.79	1.45	1.52
34	BA	523	A	O4'-C1'	-6.79	1.32	1.41
35	BB	1293	C	C3'-C2'	-6.79	1.45	1.52
85	AA	629	A	C1'-N9	-6.79	1.37	1.46
85	AA	2128	G	N1-C2	-6.79	1.32	1.37
34	BA	14	G	C5-C4	-6.79	1.33	1.38
34	BA	48	C	C3'-C2'	-6.79	1.45	1.52
34	BA	1482	A	C5-C4	-6.79	1.33	1.38
34	BA	1560	U	C1'-N1	-6.79	1.37	1.46
34	BA	1665	G	C6-N1	-6.79	1.34	1.39
35	BB	397	C	N3-C4	-6.79	1.29	1.33
35	BB	419	G	C2'-C1'	-6.79	1.45	1.53
35	BB	1103	A	P-O5'	-6.79	1.52	1.59
85	AA	384	C	C2-N3	-6.79	1.30	1.35
85	AA	444	U	N1-C2	-6.79	1.32	1.38
85	AA	766	G	C2'-C1'	-6.79	1.45	1.53
85	AA	807	A	O3'-P	-6.79	1.53	1.61
85	AA	840	A	P-O5'	-6.79	1.52	1.59
85	AA	1101	C	C2'-C1'	-6.79	1.45	1.53
85	AA	2076	C	P-O5'	-6.79	1.52	1.59
85	AA	2189	U	O3'-P	-6.79	1.53	1.61
34	BA	648	C	P-O5'	-6.78	1.52	1.59
34	BA	941	G	C3'-C2'	-6.78	1.45	1.52
34	BA	1613	G	C1'-N9	-6.78	1.37	1.46
35	BB	374	A	N9-C8	-6.78	1.32	1.37
35	BB	1084	A	P-O5'	-6.78	1.52	1.59
35	BB	1243	A	C5-C4	-6.78	1.34	1.38
35	BB	1517	G	C5'-C4'	6.78	1.59	1.51
38	BE	123	A	C5-C6	-6.78	1.34	1.41
40	BG	19	C	C2-N3	-6.78	1.30	1.35
40	BG	71	C	C4-N4	-6.78	1.27	1.33
85	AA	750	A	C5-C4	-6.78	1.34	1.38
85	AA	1295	G	C2'-C1'	-6.78	1.45	1.53
85	AA	2022	A	O3'-P	-6.78	1.53	1.61
85	AA	2082	C	C4'-O4'	-6.78	1.36	1.45
34	BA	1544	G	N9-C8	-6.78	1.33	1.37
34	BA	1804	A	N9-C8	-6.78	1.32	1.37
35	BB	1405	G	C2-N2	-6.78	1.27	1.34
36	BC	75	G	N7-C5	-6.78	1.35	1.39
37	BD	91	U	O3'-P	-6.78	1.53	1.61
41	BH	20	A	C1'-N9	-6.78	1.37	1.46
85	AA	453	G	N9-C4	6.78	1.43	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	142	A	N7-C5	-6.78	1.35	1.39
34	BA	1334	G	C6-N1	-6.78	1.34	1.39
34	BA	1682	A	N3-C4	-6.78	1.30	1.34
35	BB	677	U	N1-C2	-6.78	1.32	1.38
35	BB	736	G	P-O5'	-6.78	1.52	1.59
36	BC	75	G	N1-C2	-6.78	1.32	1.37
38	BE	12	A	C2'-C1'	-6.78	1.45	1.53
85	AA	119	G	C2-N3	-6.78	1.27	1.32
85	AA	365	G	N9-C4	-6.78	1.32	1.38
85	AA	368	C	C2-N3	-6.78	1.30	1.35
85	AA	507	C	C3'-C2'	-6.78	1.45	1.52
85	AA	1654	G	C2'-C1'	-6.78	1.45	1.53
85	AA	1863	A	P-O5'	-6.78	1.52	1.59
35	BB	1461	C	O3'-P	-6.78	1.53	1.61
85	AA	21	U	C4'-C3'	-6.78	1.45	1.53
85	AA	720	A	C2'-C1'	-6.78	1.45	1.53
85	AA	1450	U	C4-O4	-6.78	1.18	1.23
85	AA	2165	C	P-O5'	-6.78	1.52	1.59
34	BA	275	C	C2'-C1'	-6.78	1.45	1.53
34	BA	341	U	C1'-N1	-6.78	1.37	1.46
34	BA	398	G	N3-C4	-6.78	1.30	1.35
34	BA	410	G	O3'-P	-6.78	1.53	1.61
34	BA	1225	A	C3'-C2'	-6.78	1.45	1.52
35	BB	113	C	C2'-C1'	-6.78	1.45	1.53
35	BB	118	A	N7-C5	-6.78	1.35	1.39
35	BB	1271	A	N7-C5	-6.78	1.35	1.39
39	BF	11	C	O5'-C5'	6.78	1.55	1.44
41	BH	31	A	C5-C4	-6.78	1.34	1.38
85	AA	255	A	C2'-C1'	-6.78	1.45	1.53
34	BA	12	G	C2'-C1'	-6.78	1.45	1.53
34	BA	182	U	C2-N3	-6.78	1.33	1.37
34	BA	1067	G	N9-C8	-6.78	1.33	1.37
34	BA	1592	U	C4'-C3'	-6.78	1.45	1.53
35	BB	29	C	C1'-N1	-6.78	1.37	1.46
35	BB	1445	A	C3'-C2'	-6.78	1.45	1.52
38	BE	94	U	C1'-N1	-6.78	1.37	1.46
40	BG	151	A	C1'-N9	-6.78	1.37	1.46
85	AA	1365	U	O3'-P	-6.78	1.53	1.61
85	AA	1869	U	P-O5'	-6.78	1.52	1.59
34	BA	932	G	C5-C6	-6.77	1.35	1.42
34	BA	944	G	C5-C4	-6.77	1.33	1.38
34	BA	1695	G	C4'-C3'	-6.77	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	695	U	O3'-P	-6.77	1.53	1.61
35	BB	839	G	N3-C4	-6.77	1.30	1.35
35	BB	1488	G	O3'-P	-6.77	1.53	1.61
39	BF	27	G	N9-C4	-6.77	1.32	1.38
41	BH	123	G	N9-C4	-6.77	1.32	1.38
85	AA	139	G	O3'-P	-6.77	1.53	1.61
85	AA	160	A	O3'-P	-6.77	1.53	1.61
85	AA	251	A	O3'-P	-6.77	1.53	1.61
85	AA	1453	U	C2-N3	-6.77	1.33	1.37
34	BA	199	U	C2-N3	-6.77	1.33	1.37
34	BA	771	A	C5'-C4'	6.77	1.59	1.51
35	BB	53	C	C2-N3	-6.77	1.30	1.35
35	BB	1358	A	C2'-C1'	-6.77	1.46	1.53
40	BG	182	G	C5-C6	-6.77	1.35	1.42
85	AA	119	G	O3'-P	-6.77	1.53	1.61
85	AA	472	A	N9-C4	-6.77	1.33	1.37
85	AA	543	A	C3'-C2'	-6.77	1.45	1.52
85	AA	2191	C	C3'-C2'	-6.77	1.45	1.52
34	BA	619	U	C2-N3	-6.77	1.33	1.37
34	BA	690	G	P-O5'	-6.77	1.52	1.59
34	BA	725	C	O3'-P	-6.77	1.53	1.61
35	BB	1056	A	O3'-P	-6.77	1.53	1.61
36	BC	41	A	C5-C6	-6.77	1.34	1.41
38	BE	148	C	C5'-C4'	-6.77	1.43	1.51
41	BH	47	G	C1'-N9	-6.77	1.37	1.46
85	AA	686	U	C5'-C4'	-6.77	1.43	1.51
85	AA	1789	C	O3'-P	-6.77	1.53	1.61
34	BA	303	C	P-O5'	-6.77	1.52	1.59
34	BA	523	A	C4'-O4'	-6.77	1.36	1.45
34	BA	530	A	P-O5'	-6.77	1.52	1.59
34	BA	602	G	N9-C4	-6.77	1.32	1.38
34	BA	698	U	N1-C2	-6.77	1.32	1.38
34	BA	712	C	C2'-C1'	-6.77	1.46	1.53
34	BA	768	G	C1'-N9	-6.77	1.37	1.46
34	BA	1596	C	C2'-C1'	-6.77	1.46	1.53
34	BA	1647	G	N7-C5	-6.77	1.35	1.39
36	BC	129	C	C3'-C2'	-6.77	1.45	1.52
41	BH	21	G	C5-C4	-6.77	1.33	1.38
85	AA	701	C	C3'-C2'	-6.77	1.45	1.52
85	AA	748	C	O3'-P	-6.77	1.53	1.61
85	AA	1122	U	C3'-C2'	-6.77	1.45	1.52
34	BA	255	G	C5-C4	-6.77	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	890	G	C5-C4	-6.77	1.33	1.38
34	BA	900	A	N9-C8	-6.77	1.32	1.37
34	BA	1208	U	P-O5'	-6.77	1.52	1.59
34	BA	1284	G	P-O5'	-6.77	1.52	1.59
34	BA	1469	G	N1-C2	-6.77	1.32	1.37
35	BB	68	G	C5-C4	-6.77	1.33	1.38
35	BB	637	G	C2-N2	-6.77	1.27	1.34
35	BB	1310	C	C3'-C2'	-6.77	1.45	1.52
35	BB	1436	U	O3'-P	-6.77	1.53	1.61
35	BB	1445	A	C8-N7	-6.77	1.26	1.31
37	BD	87	G	O3'-P	-6.77	1.53	1.61
38	BE	4	A	N9-C4	-6.77	1.33	1.37
39	BF	18	U	N1-C6	-6.77	1.31	1.38
40	BG	50	G	C2'-C1'	-6.77	1.46	1.53
85	AA	505	U	C5'-C4'	-6.77	1.43	1.51
85	AA	648	G	P-O5'	-6.77	1.52	1.59
85	AA	895	C	P-O5'	-6.77	1.52	1.59
34	BA	313	C	C1'-N1	-6.77	1.37	1.46
34	BA	1833	G	N1-C2	-6.77	1.32	1.37
35	BB	780	U	P-O5'	-6.77	1.52	1.59
40	BG	137	G	C6-N1	-6.77	1.34	1.39
85	AA	2244	G	O3'-P	-6.77	1.53	1.61
34	BA	109	A	C2'-C1'	-6.76	1.46	1.53
34	BA	697	A	P-O5'	-6.76	1.52	1.59
34	BA	1516	G	O3'-P	-6.76	1.53	1.61
34	BA	1640	G	P-O5'	-6.76	1.52	1.59
35	BB	75	A	C2'-C1'	-6.76	1.46	1.53
35	BB	689	C	P-O5'	-6.76	1.52	1.59
35	BB	843	G	C3'-C2'	-6.76	1.45	1.52
38	BE	124	G	C6-N1	-6.76	1.34	1.39
40	BG	113	G	P-O5'	-6.76	1.52	1.59
85	AA	404	A	P-O5'	-6.76	1.52	1.59
34	BA	1074	C	P-O5'	-6.76	1.52	1.59
34	BA	1147	C	C4'-C3'	-6.76	1.45	1.53
34	BA	1561	C	O4'-C1'	-6.76	1.32	1.41
34	BA	1565	U	C2'-C1'	-6.76	1.46	1.53
35	BB	1048	A	C5-C4	-6.76	1.34	1.38
35	BB	1457	A	C4'-O4'	-6.76	1.36	1.45
85	AA	859	G	C2'-C1'	-6.76	1.46	1.53
85	AA	1241	A	C2'-C1'	-6.76	1.46	1.53
34	BA	236	A	C2'-C1'	-6.76	1.46	1.53
34	BA	736	G	C1'-N9	-6.76	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1573	C	C2'-C1'	-6.76	1.46	1.53
34	BA	1687	A	C3'-C2'	-6.76	1.45	1.52
34	BA	1728	G	C6-N1	-6.76	1.34	1.39
35	BB	268	G	C2'-C1'	-6.76	1.46	1.53
35	BB	428	G	O3'-P	-6.76	1.53	1.61
35	BB	811	C	C3'-C2'	-6.76	1.45	1.52
35	BB	1174	C	O3'-P	-6.76	1.53	1.61
37	BD	48	G	O4'-C1'	-6.76	1.32	1.41
38	BE	69	C	P-O5'	-6.76	1.52	1.59
38	BE	127	G	C5-C4	-6.76	1.33	1.38
34	BA	1086	A	N7-C5	-6.76	1.35	1.39
35	BB	96	A	C2'-C1'	-6.76	1.46	1.53
35	BB	127	U	O3'-P	-6.76	1.53	1.61
35	BB	996	G	C5-C4	-6.76	1.33	1.38
35	BB	1138	A	C3'-C2'	-6.76	1.45	1.52
35	BB	1157	G	C3'-C2'	-6.76	1.45	1.52
36	BC	156	A	C2'-C1'	-6.76	1.46	1.53
85	AA	24	U	C2-N3	-6.76	1.33	1.37
85	AA	189	G	C5-C4	-6.76	1.33	1.38
85	AA	1460	G	P-O5'	-6.76	1.52	1.59
85	AA	1655	G	C1'-N9	-6.76	1.37	1.46
85	AA	1708	A	P-O5'	-6.76	1.52	1.59
34	BA	903	C	O3'-P	-6.76	1.53	1.61
34	BA	1736	A	C8-N7	-6.76	1.26	1.31
35	BB	540	G	O3'-P	-6.76	1.53	1.61
38	BE	62	C	O3'-P	-6.76	1.53	1.61
85	AA	866	U	C4'-C3'	-6.76	1.45	1.53
35	BB	387	G	C8-N7	-6.76	1.26	1.30
35	BB	577	U	O3'-P	-6.76	1.53	1.61
35	BB	578	G	C2'-C1'	-6.76	1.46	1.53
40	BG	90	G	N1-C2	-6.76	1.32	1.37
41	BH	13	C	C3'-C2'	-6.76	1.45	1.52
85	AA	2045	U	C3'-C2'	-6.76	1.45	1.52
34	BA	858	C	O3'-P	-6.75	1.53	1.61
34	BA	1163	G	C1'-N9	-6.75	1.37	1.46
35	BB	544	C	P-O5'	-6.75	1.52	1.59
39	BF	7	G	C2'-C1'	-6.75	1.46	1.53
85	AA	532	G	N7-C5	-6.75	1.35	1.39
85	AA	667	A	C1'-N9	-6.75	1.37	1.46
85	AA	772	C	O3'-P	-6.75	1.53	1.61
85	AA	2193	A	C8-N7	-6.75	1.26	1.31
34	BA	231	U	C5'-C4'	6.75	1.59	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	395	G	C5-C4	-6.75	1.33	1.38
34	BA	1691	G	N7-C5	-6.75	1.35	1.39
34	BA	1728	G	C2-N2	-6.75	1.27	1.34
35	BB	770	G	C2'-C1'	-6.75	1.46	1.53
35	BB	1434	G	C2'-C1'	-6.75	1.46	1.53
36	BC	19	A	P-O5'	-6.75	1.52	1.59
36	BC	34	U	O3'-P	-6.75	1.53	1.61
85	AA	20	G	C5-C4	-6.75	1.33	1.38
85	AA	116	G	C3'-C2'	-6.75	1.45	1.52
85	AA	430	G	C1'-N9	-6.75	1.37	1.46
85	AA	631	G	C6-N1	-6.75	1.34	1.39
34	BA	146	G	O3'-P	-6.75	1.53	1.61
34	BA	406	G	N9-C8	-6.75	1.33	1.37
34	BA	468	A	C4'-C3'	6.75	1.60	1.53
34	BA	691	A	C1'-N9	-6.75	1.37	1.46
34	BA	1266	A	C4'-O4'	-6.75	1.36	1.45
34	BA	1396	A	N7-C5	-6.75	1.35	1.39
34	BA	1632	G	C6-N1	-6.75	1.34	1.39
35	BB	1375	G	C2'-C1'	-6.75	1.46	1.53
37	BD	95	G	C3'-C2'	-6.75	1.45	1.52
38	BE	195	G	N9-C8	-6.75	1.33	1.37
85	AA	59	C	O3'-P	-6.75	1.53	1.61
85	AA	492	C	C2-N3	-6.75	1.30	1.35
85	AA	918	U	C2-N3	-6.75	1.33	1.37
85	AA	926	C	C1'-N1	-6.75	1.37	1.46
85	AA	1209	U	C4-C5	-6.75	1.37	1.43
85	AA	1502	A	N9-C8	-6.75	1.32	1.37
34	BA	11	U	C2-N3	-6.75	1.33	1.37
34	BA	598	G	N9-C4	-6.75	1.32	1.38
34	BA	631	G	C2-N3	-6.75	1.27	1.32
34	BA	760	G	O3'-P	-6.75	1.53	1.61
34	BA	1657	A	C5-C4	-6.75	1.34	1.38
34	BA	1827	C	O3'-P	-6.75	1.53	1.61
35	BB	1070	G	O3'-P	-6.75	1.53	1.61
85	AA	324	U	C3'-C2'	-6.75	1.45	1.52
85	AA	543	A	C5-C4	-6.75	1.34	1.38
85	AA	1462	A	C1'-N9	-6.75	1.37	1.46
34	BA	1577	U	C3'-C2'	-6.75	1.45	1.52
35	BB	112	G	O3'-P	-6.75	1.53	1.61
36	BC	82	C	P-O5'	-6.75	1.53	1.59
85	AA	99	U	C3'-C2'	-6.75	1.45	1.52
85	AA	1144	G	N3-C4	-6.75	1.30	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1514	A	P-O5'	-6.75	1.53	1.59
85	AA	1570	A	O3'-P	-6.75	1.53	1.61
85	AA	1701	G	C3'-C2'	-6.75	1.45	1.52
85	AA	2133	A	C1'-N9	-6.75	1.37	1.46
34	BA	397	A	N9-C4	-6.75	1.33	1.37
34	BA	964	U	N1-C2	-6.75	1.32	1.38
35	BB	490	G	C5-C4	-6.75	1.33	1.38
35	BB	1421	C	N3-C4	-6.75	1.29	1.33
39	BF	42	G	C2'-C1'	-6.75	1.46	1.53
41	BH	128	G	N9-C8	-6.75	1.33	1.37
34	BA	225	A	O3'-P	-6.75	1.53	1.61
34	BA	543	A	C3'-C2'	-6.75	1.45	1.52
34	BA	942	G	C3'-C2'	-6.75	1.45	1.52
34	BA	1073	G	C2-N2	-6.75	1.27	1.34
35	BB	1022	C	C2'-C1'	-6.75	1.46	1.53
40	BG	110	U	C2-N3	-6.75	1.33	1.37
85	AA	309	G	C3'-C2'	-6.75	1.45	1.52
85	AA	876	U	C2'-C1'	-6.75	1.46	1.53
34	BA	1203	G	C5-C4	-6.74	1.33	1.38
34	BA	1668	C	C3'-C2'	-6.74	1.45	1.52
34	BA	1684	A	O3'-P	-6.74	1.53	1.61
35	BB	469	G	C1'-N9	-6.74	1.37	1.46
35	BB	558	U	C2-N3	-6.74	1.33	1.37
35	BB	687	C	C3'-C2'	-6.74	1.45	1.52
35	BB	1158	C	P-O5'	-6.74	1.53	1.59
35	BB	1322	A	P-O5'	-6.74	1.53	1.59
36	BC	146	U	P-O5'	-6.74	1.53	1.59
85	AA	1903	G	C2'-C1'	-6.74	1.46	1.53
35	BB	692	G	O4'-C1'	-6.74	1.32	1.41
36	BC	119	G	C6-N1	-6.74	1.34	1.39
85	AA	575	G	P-O5'	-6.74	1.53	1.59
85	AA	720	A	C3'-C2'	-6.74	1.45	1.52
85	AA	1120	G	C3'-C2'	-6.74	1.45	1.52
34	BA	427	G	C5-C4	-6.74	1.33	1.38
34	BA	821	G	C4'-C3'	6.74	1.60	1.53
34	BA	990	G	C6-N1	-6.74	1.34	1.39
34	BA	1279	U	C3'-C2'	-6.74	1.45	1.52
34	BA	1309	U	C5'-C4'	6.74	1.59	1.51
35	BB	454	U	O3'-P	-6.74	1.53	1.61
35	BB	1084	A	C5-C4	-6.74	1.34	1.38
85	AA	1126	G	O3'-P	-6.74	1.53	1.61
85	AA	1221	G	C2-N2	-6.74	1.27	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1491	G	N3-C4	-6.74	1.30	1.35
34	BA	49	A	C5-C4	-6.74	1.34	1.38
34	BA	206	C	P-O5'	6.74	1.66	1.59
35	BB	430	A	O3'-P	-6.74	1.53	1.61
35	BB	559	U	C2-N3	-6.74	1.33	1.37
35	BB	1201	G	O3'-P	6.74	1.69	1.61
36	BC	97	U	C2-N3	-6.74	1.33	1.37
85	AA	980	U	P-O5'	-6.74	1.53	1.59
34	BA	23	A	C6-N6	-6.74	1.28	1.33
35	BB	648	G	P-O5'	-6.74	1.53	1.59
35	BB	1519	U	C3'-C2'	-6.74	1.45	1.52
38	BE	181	U	P-O5'	-6.74	1.53	1.59
85	AA	26	A	O3'-P	-6.74	1.53	1.61
85	AA	979	U	P-O5'	-6.74	1.53	1.59
34	BA	479	U	C3'-C2'	-6.74	1.45	1.52
34	BA	1473	A	C1'-N9	-6.74	1.37	1.46
35	BB	508	U	C2'-C1'	-6.74	1.46	1.53
35	BB	556	U	C2'-C1'	-6.74	1.46	1.53
35	BB	1239	A	O3'-P	-6.74	1.53	1.61
35	BB	1263	A	C5-C4	-6.74	1.34	1.38
35	BB	1410	G	C2'-C1'	-6.74	1.46	1.53
38	BE	30	C	O3'-P	-6.74	1.53	1.61
85	AA	989	U	C3'-C2'	-6.74	1.45	1.52
85	AA	1125	G	C6-N1	-6.74	1.34	1.39
34	BA	248	G	N9-C4	-6.73	1.32	1.38
35	BB	1356	G	N7-C5	-6.73	1.35	1.39
85	AA	2036	A	N9-C8	-6.73	1.32	1.37
34	BA	86	A	C5-C4	-6.73	1.34	1.38
34	BA	94	G	C4'-C3'	-6.73	1.45	1.53
34	BA	146	G	N3-C4	-6.73	1.30	1.35
34	BA	187	G	O3'-P	-6.73	1.53	1.61
34	BA	491	U	N1-C2	-6.73	1.32	1.38
34	BA	916	A	N9-C4	-6.73	1.33	1.37
35	BB	486	G	P-O5'	-6.73	1.53	1.59
35	BB	1110	G	P-O5'	-6.73	1.53	1.59
35	BB	1113	C	O3'-P	-6.73	1.53	1.61
35	BB	1332	G	C6-N1	-6.73	1.34	1.39
85	AA	1896	G	O3'-P	-6.73	1.53	1.61
34	BA	15	G	O3'-P	-6.73	1.53	1.61
34	BA	247	U	P-O5'	-6.73	1.53	1.59
34	BA	339	G	C2-N2	-6.73	1.27	1.34
34	BA	908	G	C6-N1	-6.73	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	919	A	C6-N1	-6.73	1.30	1.35
34	BA	1165	A	C3'-C2'	-6.73	1.45	1.52
34	BA	1847	G	C8-N7	-6.73	1.26	1.30
35	BB	691	A	C2'-C1'	-6.73	1.46	1.53
35	BB	1068	G	O3'-P	-6.73	1.53	1.61
40	BG	138	C	C5'-C4'	-6.73	1.43	1.51
85	AA	1175	A	C4'-C3'	-6.73	1.45	1.53
85	AA	1291	A	C2'-C1'	-6.73	1.46	1.53
85	AA	1527	G	N9-C4	-6.73	1.32	1.38
34	BA	1650	G	C3'-C2'	-6.73	1.45	1.52
35	BB	1230	A	O4'-C1'	-6.73	1.32	1.41
35	BB	1410	G	O3'-P	-6.73	1.53	1.61
34	BA	166	G	C2'-C1'	-6.73	1.46	1.53
34	BA	325	A	N7-C5	-6.73	1.35	1.39
34	BA	798	G	N3-C4	-6.73	1.30	1.35
34	BA	1702	G	O3'-P	-6.73	1.53	1.61
35	BB	612	A	C2'-C1'	-6.73	1.46	1.53
35	BB	1082	A	N9-C4	-6.73	1.33	1.37
35	BB	1435	G	N7-C5	-6.73	1.35	1.39
36	BC	3	C	C4'-C3'	-6.73	1.45	1.53
37	BD	41	G	O4'-C1'	-6.73	1.32	1.41
40	BG	34	A	C5-C4	-6.73	1.34	1.38
40	BG	76	C	C5'-C4'	-6.73	1.43	1.51
85	AA	385	A	N3-C4	-6.73	1.30	1.34
85	AA	385	A	N7-C5	-6.73	1.35	1.39
85	AA	903	G	P-O5'	-6.73	1.53	1.59
85	AA	1214	C	P-O5'	-6.73	1.53	1.59
85	AA	1476	C	P-O5'	-6.73	1.53	1.59
85	AA	1508	A	N7-C5	-6.73	1.35	1.39
85	AA	2060	G	C5-C4	-6.73	1.33	1.38
34	BA	109	A	O3'-P	-6.73	1.53	1.61
34	BA	600	G	N9-C4	-6.73	1.32	1.38
34	BA	1316	G	O3'-P	-6.73	1.53	1.61
85	AA	867	G	N7-C5	-6.73	1.35	1.39
85	AA	1115	G	C2-N2	-6.73	1.27	1.34
34	BA	126	G	C8-N7	-6.72	1.26	1.30
34	BA	877	U	N3-C4	-6.72	1.32	1.38
34	BA	1310	C	N3-C4	-6.72	1.29	1.33
35	BB	1359	G	C3'-C2'	-6.72	1.45	1.52
35	BB	1451	C	O3'-P	-6.72	1.53	1.61
38	BE	154	A	C4'-C3'	-6.72	1.45	1.53
85	AA	254	G	O3'-P	-6.72	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	584	G	C5-C6	-6.72	1.35	1.42
85	AA	1578	G	C6-N1	-6.72	1.34	1.39
34	BA	349	G	N9-C8	-6.72	1.33	1.37
34	BA	653	U	O3'-P	-6.72	1.53	1.61
34	BA	1735	G	C2'-C1'	-6.72	1.46	1.53
35	BB	366	G	P-O5'	-6.72	1.53	1.59
35	BB	544	C	O3'-P	-6.72	1.53	1.61
35	BB	803	U	O3'-P	-6.72	1.53	1.61
35	BB	1448	U	C2-N3	-6.72	1.33	1.37
36	BC	132	U	C2'-C1'	-6.72	1.46	1.53
85	AA	597	A	C4'-C3'	-6.72	1.45	1.53
85	AA	1480	C	P-O5'	-6.72	1.53	1.59
85	AA	1665	G	O3'-P	-6.72	1.53	1.61
34	BA	382	G	O3'-P	-6.72	1.53	1.61
34	BA	464	U	P-O5'	-6.72	1.53	1.59
34	BA	1535	G	C2'-C1'	-6.72	1.46	1.53
35	BB	1215	U	C2-N3	-6.72	1.33	1.37
35	BB	1399	A	O3'-P	-6.72	1.53	1.61
85	AA	386	G	C2'-C1'	-6.72	1.46	1.53
85	AA	2128	G	C2-N2	-6.72	1.27	1.34
34	BA	180	G	O3'-P	-6.72	1.53	1.61
34	BA	191	G	C6-N1	-6.72	1.34	1.39
34	BA	341	U	C2'-C1'	-6.72	1.46	1.53
34	BA	753	G	C2-N2	-6.72	1.27	1.34
34	BA	1164	C	C2-N3	-6.72	1.30	1.35
34	BA	1165	A	N7-C5	-6.72	1.35	1.39
34	BA	1533	G	N1-C2	-6.72	1.32	1.37
35	BB	418	G	O3'-P	-6.72	1.53	1.61
35	BB	470	C	C1'-N1	-6.72	1.37	1.46
35	BB	984	U	O3'-P	-6.72	1.53	1.61
35	BB	1048	A	C1'-N9	-6.72	1.37	1.46
35	BB	1449	G	C5-C4	-6.72	1.33	1.38
85	AA	159	G	C2-N3	-6.72	1.27	1.32
85	AA	1553	G	C2'-C1'	-6.72	1.46	1.53
34	BA	1262	A	O4'-C1'	-6.72	1.32	1.41
35	BB	494	C	C2-N3	-6.72	1.30	1.35
35	BB	964	G	C5-C4	-6.72	1.33	1.38
85	AA	432	A	N7-C5	-6.72	1.35	1.39
85	AA	2198	G	C2-N2	-6.72	1.27	1.34
34	BA	411	C	P-O5'	-6.72	1.53	1.59
34	BA	752	A	O3'-P	-6.72	1.53	1.61
34	BA	1497	A	P-O5'	-6.72	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	841	U	C2-N3	-6.72	1.33	1.37
35	BB	1302	C	O3'-P	-6.72	1.53	1.61
37	BD	35	C	C3'-C2'	-6.72	1.45	1.52
85	AA	180	A	P-O5'	-6.72	1.53	1.59
85	AA	668	A	C2'-C1'	-6.72	1.46	1.53
34	BA	683	C	C4'-C3'	6.71	1.60	1.53
34	BA	1494	G	C5-C4	-6.71	1.33	1.38
35	BB	362	A	O3'-P	-6.71	1.53	1.61
35	BB	552	C	C3'-C2'	-6.71	1.45	1.52
35	BB	1194	A	C4'-C3'	-6.71	1.45	1.53
35	BB	1394	A	C3'-C2'	-6.71	1.45	1.52
35	BB	1467	A	C2'-C1'	-6.71	1.46	1.53
39	BF	29	U	C2'-C1'	-6.71	1.46	1.53
34	BA	435	U	C2-N3	-6.71	1.33	1.37
34	BA	1073	G	N3-C4	-6.71	1.30	1.35
34	BA	1173	C	C3'-C2'	-6.71	1.45	1.52
34	BA	1222	C	C4'-O4'	-6.71	1.36	1.45
34	BA	1644	A	O3'-P	-6.71	1.53	1.61
35	BB	108	G	N9-C8	-6.71	1.33	1.37
35	BB	1002	G	C2'-C1'	-6.71	1.46	1.53
36	BC	30	U	N3-C4	-6.71	1.32	1.38
85	AA	84	C	O3'-P	-6.71	1.53	1.61
85	AA	1479	U	O3'-P	-6.71	1.53	1.61
85	AA	1688	U	P-O5'	-6.71	1.53	1.59
34	BA	207	A	C5-C4	-6.71	1.34	1.38
34	BA	326	A	N3-C4	-6.71	1.30	1.34
34	BA	330	A	N3-C4	-6.71	1.30	1.34
34	BA	417	A	O4'-C1'	-6.71	1.32	1.41
34	BA	1226	G	C3'-C2'	-6.71	1.45	1.52
35	BB	596	C	C3'-C2'	-6.71	1.45	1.52
35	BB	681	G	O3'-P	-6.71	1.53	1.61
35	BB	707	G	C1'-N9	-6.71	1.37	1.46
35	BB	815	G	C5-C4	-6.71	1.33	1.38
35	BB	1365	G	C1'-N9	-6.71	1.37	1.46
35	BB	1375	G	C2-N2	-6.71	1.27	1.34
85	AA	74	U	C2'-C1'	-6.71	1.46	1.53
85	AA	722	G	C8-N7	-6.71	1.26	1.30
85	AA	1299	A	C3'-C2'	-6.71	1.45	1.52
34	BA	801	U	O3'-P	-6.71	1.53	1.61
34	BA	1086	A	O3'-P	-6.71	1.53	1.61
34	BA	1580	U	P-O5'	-6.71	1.53	1.59
35	BB	119	G	N9-C8	-6.71	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	6	A	N9-C4	-6.71	1.33	1.37
85	AA	173	A	N7-C5	-6.71	1.35	1.39
34	BA	374	U	P-O5'	-6.71	1.53	1.59
34	BA	572	G	C3'-O3'	6.71	1.51	1.42
34	BA	1487	U	N3-C4	-6.71	1.32	1.38
35	BB	337	U	P-O5'	-6.71	1.53	1.59
37	BD	71	G	C2-N2	-6.71	1.27	1.34
41	BH	104	U	C3'-C2'	-6.71	1.45	1.52
85	AA	1302	A	N9-C4	6.71	1.41	1.37
85	AA	1685	G	O3'-P	-6.71	1.53	1.61
85	AA	1714	G	N9-C4	-6.71	1.32	1.38
85	AA	1799	C	C4-C5	-6.71	1.37	1.43
34	BA	454	G	C1'-N9	-6.71	1.37	1.46
34	BA	1414	C	C2'-C1'	-6.71	1.46	1.53
34	BA	1488	C	O3'-P	-6.71	1.53	1.61
34	BA	1570	C	C3'-C2'	-6.71	1.45	1.52
38	BE	104	G	N9-C4	6.71	1.43	1.38
38	BE	110	U	O3'-P	-6.71	1.53	1.61
40	BG	81	G	C5-C4	-6.71	1.33	1.38
41	BH	35	G	N1-C2	-6.71	1.32	1.37
34	BA	240	C	O3'-P	-6.71	1.53	1.61
34	BA	529	A	C5-C6	-6.71	1.35	1.41
34	BA	1568	A	O3'-P	-6.71	1.53	1.61
35	BB	428	G	N3-C4	-6.71	1.30	1.35
35	BB	629	C	C2-N3	-6.71	1.30	1.35
35	BB	747	A	P-O5'	-6.71	1.53	1.59
40	BG	32	U	N1-C2	-6.71	1.32	1.38
85	AA	116	G	C2'-C1'	-6.71	1.46	1.53
85	AA	2172	A	N9-C4	-6.71	1.33	1.37
34	BA	407	A	N3-C4	-6.70	1.30	1.34
34	BA	988	U	O3'-P	-6.70	1.53	1.61
34	BA	1072	U	P-O5'	-6.70	1.53	1.59
34	BA	1246	G	C5-C4	-6.70	1.33	1.38
35	BB	133	G	C5-C4	-6.70	1.33	1.38
35	BB	487	A	C5-C4	-6.70	1.34	1.38
35	BB	1381	U	C2-N3	-6.70	1.33	1.37
35	BB	1488	G	C2'-C1'	-6.70	1.46	1.53
85	AA	367	A	O4'-C1'	-6.70	1.32	1.41
85	AA	677	U	P-O5'	-6.70	1.53	1.59
85	AA	1263	G	N1-C2	-6.70	1.32	1.37
85	AA	2067	A	C4'-C3'	-6.70	1.45	1.53
34	BA	1823	A	O3'-P	-6.70	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	393	A	C3'-C2'	-6.70	1.45	1.52
37	BD	91	U	C2'-C1'	-6.70	1.46	1.53
39	BF	47	C	C3'-C2'	-6.70	1.45	1.52
85	AA	69	C	C1'-N1	-6.70	1.37	1.46
34	BA	223	U	P-O5'	-6.70	1.53	1.59
34	BA	426	A	C4'-C3'	-6.70	1.45	1.53
34	BA	1280	A	C3'-C2'	-6.70	1.45	1.52
34	BA	1844	U	C2'-C1'	-6.70	1.46	1.53
35	BB	155	G	P-O5'	-6.70	1.53	1.59
35	BB	658	G	P-O5'	-6.70	1.53	1.59
35	BB	702	G	O3'-P	-6.70	1.53	1.61
85	AA	312	G	O3'-P	-6.70	1.53	1.61
85	AA	442	G	C1'-N9	-6.70	1.37	1.46
85	AA	443	A	C1'-N9	-6.70	1.37	1.46
85	AA	642	G	C2-N2	-6.70	1.27	1.34
85	AA	1887	G	C2'-C1'	-6.70	1.46	1.53
34	BA	515	U	C2'-C1'	-6.70	1.46	1.53
34	BA	704	G	O3'-P	-6.70	1.53	1.61
35	BB	538	A	N7-C5	-6.70	1.35	1.39
35	BB	542	A	P-O5'	-6.70	1.53	1.59
35	BB	833	G	N9-C8	-6.70	1.33	1.37
35	BB	1128	U	P-O5'	-6.70	1.53	1.59
85	AA	1891	U	O3'-P	-6.70	1.53	1.61
85	AA	2173	A	C3'-C2'	-6.70	1.45	1.52
34	BA	1090	A	C2'-C1'	-6.70	1.46	1.53
34	BA	1257	U	C4'-C3'	-6.70	1.45	1.53
34	BA	1650	G	C2'-C1'	-6.70	1.46	1.53
35	BB	1457	A	O3'-P	-6.70	1.53	1.61
85	AA	316	C	N3-C4	-6.70	1.29	1.33
34	BA	463	A	C2'-C1'	-6.70	1.46	1.53
34	BA	591	G	C1'-N9	-6.70	1.37	1.46
34	BA	819	G	P-O5'	-6.70	1.53	1.59
34	BA	984	U	P-O5'	-6.70	1.53	1.59
34	BA	1685	C	C1'-N1	-6.70	1.37	1.46
34	BA	1799	G	C6-N1	-6.70	1.34	1.39
35	BB	383	U	O3'-P	-6.70	1.53	1.61
35	BB	454	U	C3'-C2'	-6.70	1.45	1.52
35	BB	501	G	C3'-C2'	-6.70	1.45	1.52
35	BB	1015	U	C3'-C2'	-6.70	1.45	1.52
85	AA	318	A	C1'-N9	-6.70	1.37	1.46
34	BA	260	A	C3'-C2'	-6.69	1.45	1.52
36	BC	61	A	P-O5'	-6.69	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	47	U	C1'-N1	-6.69	1.37	1.46
85	AA	2014	G	N3-C4	-6.69	1.30	1.35
34	BA	1039	G	N9-C8	-6.69	1.33	1.37
34	BA	1162	U	N3-C4	-6.69	1.32	1.38
35	BB	1108	G	N1-C2	-6.69	1.32	1.37
35	BB	1359	G	O3'-P	-6.69	1.53	1.61
40	BG	50	G	N9-C8	-6.69	1.33	1.37
85	AA	596	A	O3'-P	-6.69	1.53	1.61
85	AA	2188	C	C4-N4	-6.69	1.27	1.33
85	AA	2217	A	N9-C4	-6.69	1.33	1.37
34	BA	253	U	C3'-C2'	-6.69	1.45	1.52
34	BA	710	A	O3'-P	-6.69	1.53	1.61
34	BA	1583	A	O3'-P	-6.69	1.53	1.61
34	BA	1601	C	C2-N3	-6.69	1.30	1.35
35	BB	1420	U	C4'-C3'	-6.69	1.45	1.53
36	BC	15	G	C1'-N9	-6.69	1.37	1.46
40	BG	55	A	C5-C4	-6.69	1.34	1.38
34	BA	1672	C	N1-C6	-6.69	1.33	1.37
34	BA	1682	A	C1'-N9	-6.69	1.37	1.46
35	BB	441	G	C5-C4	-6.69	1.33	1.38
35	BB	1229	A	C1'-N9	-6.69	1.37	1.46
85	AA	1657	C	C2-N3	-6.69	1.30	1.35
34	BA	1058	C	C3'-C2'	-6.69	1.45	1.52
34	BA	1433	U	C4'-O4'	-6.69	1.36	1.45
35	BB	728	A	C1'-N9	-6.69	1.37	1.46
36	BC	18	G	N7-C5	-6.69	1.35	1.39
38	BE	73	A	C2'-C1'	-6.69	1.46	1.53
85	AA	309	G	O3'-P	-6.69	1.53	1.61
85	AA	1201	A	P-O5'	-6.69	1.53	1.59
85	AA	1263	G	C1'-N9	-6.69	1.37	1.46
85	AA	1670	U	C3'-C2'	-6.69	1.45	1.52
34	BA	334	G	P-O5'	-6.69	1.53	1.59
34	BA	1591	G	C6-N1	-6.69	1.34	1.39
34	BA	1685	C	C4'-C3'	-6.69	1.45	1.53
36	BC	9	G	N3-C4	-6.69	1.30	1.35
36	BC	27	U	N1-C2	-6.69	1.32	1.38
85	AA	435	A	C5-C4	-6.69	1.34	1.38
85	AA	1266	C	O3'-P	-6.69	1.53	1.61
34	BA	291	C	C2-N3	-6.68	1.30	1.35
34	BA	431	A	C4'-C3'	-6.68	1.45	1.53
34	BA	487	A	N1-C2	-6.68	1.28	1.34
34	BA	1183	U	C5'-C4'	6.68	1.59	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	380	G	C2-N3	-6.68	1.27	1.32
35	BB	580	A	C2'-C1'	-6.68	1.46	1.53
35	BB	1330	A	O3'-P	-6.68	1.53	1.61
85	AA	405	C	O3'-P	-6.68	1.53	1.61
85	AA	792	A	C6-N6	-6.68	1.28	1.33
85	AA	1460	G	O3'-P	-6.68	1.53	1.61
85	AA	1476	C	C2'-C1'	-6.68	1.46	1.53
85	AA	1629	C	O3'-P	-6.68	1.53	1.61
34	BA	105	U	C3'-C2'	-6.68	1.45	1.52
34	BA	855	C	C2'-C1'	-6.68	1.46	1.53
34	BA	1150	A	C1'-N9	-6.68	1.37	1.46
34	BA	1178	U	C3'-C2'	-6.68	1.45	1.52
35	BB	37	C	C2'-C1'	-6.68	1.46	1.53
35	BB	469	G	P-O5'	-6.68	1.53	1.59
35	BB	828	G	C3'-C2'	-6.68	1.45	1.52
35	BB	1123	A	C1'-N9	-6.68	1.37	1.46
35	BB	1408	G	N9-C4	-6.68	1.32	1.38
40	BG	62	C	C2'-C1'	-6.68	1.46	1.53
40	BG	98	A	C3'-C2'	-6.68	1.45	1.52
40	BG	135	C	N1-C6	-6.68	1.33	1.37
85	AA	102	A	C1'-N9	-6.68	1.37	1.46
85	AA	2136	C	C2-N3	-6.68	1.30	1.35
34	BA	536	C	C2'-C1'	-6.68	1.46	1.53
35	BB	3	C	C4-C5	-6.68	1.37	1.43
35	BB	109	U	C2-N3	-6.68	1.33	1.37
38	BE	50	G	C4'-C3'	-6.68	1.45	1.53
85	AA	76	G	O3'-P	-6.68	1.53	1.61
34	BA	18	G	C5-C4	-6.68	1.33	1.38
34	BA	278	U	O3'-P	-6.68	1.53	1.61
34	BA	1305	A	N7-C5	-6.68	1.35	1.39
34	BA	1431	G	P-O5'	-6.68	1.53	1.59
35	BB	44	C	C4'-C3'	-6.68	1.45	1.53
35	BB	70	A	C5'-C4'	6.68	1.59	1.51
35	BB	680	A	C8-N7	-6.68	1.26	1.31
35	BB	1345	A	N9-C4	-6.68	1.33	1.37
38	BE	26	G	P-O5'	-6.68	1.53	1.59
40	BG	129	G	C2'-C1'	-6.68	1.46	1.53
85	AA	20	G	N3-C4	-6.68	1.30	1.35
85	AA	2205	A	O3'-P	-6.68	1.53	1.61
34	BA	751	A	C2'-C1'	-6.68	1.46	1.53
34	BA	784	C	C3'-C2'	-6.68	1.45	1.52
34	BA	1058	C	P-O5'	-6.68	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	126	C	C2-N3	-6.68	1.30	1.35
35	BB	1113	C	C2'-C1'	-6.68	1.46	1.53
85	AA	176	C	C2'-C1'	-6.68	1.46	1.53
85	AA	1546	G	N9-C4	-6.68	1.32	1.38
85	AA	2139	G	C5-C4	-6.68	1.33	1.38
34	BA	257	G	P-O5'	-6.68	1.53	1.59
34	BA	299	C	C2'-C1'	-6.68	1.46	1.53
34	BA	480	G	C2'-C1'	-6.68	1.46	1.53
34	BA	588	C	C1'-N1	-6.68	1.37	1.46
34	BA	932	G	C2'-C1'	-6.68	1.46	1.53
34	BA	1066	A	C2'-C1'	-6.68	1.46	1.53
34	BA	1446	G	C6-N1	-6.68	1.34	1.39
35	BB	30	A	N9-C4	-6.68	1.33	1.37
35	BB	62	C	O3'-P	-6.68	1.53	1.61
35	BB	575	C	C2'-C1'	-6.68	1.46	1.53
35	BB	1063	C	C1'-N1	-6.68	1.37	1.46
35	BB	1175	A	N3-C4	-6.68	1.30	1.34
36	BC	128	U	C2'-C1'	-6.68	1.46	1.53
40	BG	74	G	N1-C2	-6.68	1.32	1.37
40	BG	167	C	C2'-C1'	-6.68	1.46	1.53
85	AA	532	G	C2-N2	-6.68	1.27	1.34
85	AA	770	C	P-O5'	-6.68	1.53	1.59
85	AA	1118	U	P-O5'	-6.68	1.53	1.59
85	AA	1578	G	N7-C5	-6.68	1.35	1.39
85	AA	2132	A	P-O5'	-6.68	1.53	1.59
34	BA	516	U	C4'-O4'	-6.67	1.36	1.45
35	BB	1102	U	N1-C6	-6.67	1.31	1.38
36	BC	154	A	P-O5'	-6.67	1.53	1.59
37	BD	96	C	C4-N4	-6.67	1.27	1.33
40	BG	75	C	C1'-N1	-6.67	1.37	1.46
40	BG	79	U	P-O5'	-6.67	1.53	1.59
85	AA	333	A	N9-C4	-6.67	1.33	1.37
85	AA	1704	C	C2'-C1'	-6.67	1.46	1.53
85	AA	2170	G	C2'-C1'	-6.67	1.46	1.53
34	BA	1515	U	O3'-P	-6.67	1.53	1.61
35	BB	595	U	C2'-C1'	-6.67	1.46	1.53
35	BB	1220	A	C3'-C2'	-6.67	1.45	1.52
85	AA	56	U	C2'-C1'	-6.67	1.46	1.53
85	AA	483	G	C4'-C3'	-6.67	1.45	1.53
85	AA	537	G	C2'-C1'	-6.67	1.46	1.53
34	BA	389	U	C2'-C1'	-6.67	1.46	1.53
34	BA	578	C	O3'-P	-6.67	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	822	U	P-O5'	-6.67	1.53	1.59
34	BA	892	C	O4'-C1'	-6.67	1.32	1.41
34	BA	1530	G	N9-C8	-6.67	1.33	1.37
35	BB	515	C	P-O5'	-6.67	1.53	1.59
41	BH	66	G	C2'-C1'	-6.67	1.46	1.53
85	AA	506	G	C6-N1	-6.67	1.34	1.39
85	AA	936	C	C4-N4	-6.67	1.27	1.33
85	AA	1813	C	C2'-C1'	-6.67	1.46	1.53
85	AA	2172	A	N3-C4	-6.67	1.30	1.34
34	BA	1006	G	C1'-N9	-6.67	1.37	1.46
35	BB	606	C	C3'-C2'	-6.67	1.45	1.52
85	AA	421	G	N1-C2	-6.67	1.32	1.37
85	AA	2007	G	O3'-P	-6.67	1.53	1.61
34	BA	124	G	O3'-P	-6.67	1.53	1.61
34	BA	839	U	C2-N3	-6.67	1.33	1.37
34	BA	1005	C	C4'-C3'	-6.67	1.45	1.53
34	BA	1171	C	C2-N3	-6.67	1.30	1.35
35	BB	92	C	P-O5'	-6.67	1.53	1.59
35	BB	135	C	N1-C6	-6.67	1.33	1.37
35	BB	433	C	O3'-P	-6.67	1.53	1.61
35	BB	481	A	C3'-C2'	-6.67	1.45	1.52
35	BB	1144	A	C5-C4	-6.67	1.34	1.38
37	BD	16	U	C2'-C1'	-6.67	1.46	1.53
41	BH	3	U	C3'-C2'	-6.67	1.45	1.52
85	AA	458	C	C2-N3	-6.67	1.30	1.35
85	AA	671	G	C5'-C4'	-6.67	1.43	1.51
85	AA	1177	G	C6-N1	-6.67	1.34	1.39
85	AA	1284	A	C1'-N9	-6.67	1.37	1.46
34	BA	38	G	C2-N2	-6.67	1.27	1.34
34	BA	104	A	N3-C4	-6.67	1.30	1.34
34	BA	705	C	O3'-P	-6.67	1.53	1.61
34	BA	943	G	C2-N2	-6.67	1.27	1.34
35	BB	1165	A	P-O5'	-6.67	1.53	1.59
85	AA	161	A	C1'-N9	-6.67	1.37	1.46
85	AA	439	U	P-O5'	-6.67	1.53	1.59
34	BA	1273	U	C4'-C3'	-6.67	1.45	1.53
35	BB	659	C	O3'-P	-6.67	1.53	1.61
85	AA	2102	A	C1'-N9	-6.67	1.37	1.46
34	BA	857	C	C1'-N1	-6.66	1.37	1.46
34	BA	1538	G	N7-C5	-6.66	1.35	1.39
34	BA	1795	A	C5-C4	-6.66	1.34	1.38
35	BB	65	A	C5-C4	-6.66	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	815	G	P-O5'	-6.66	1.53	1.59
35	BB	981	A	N7-C5	-6.66	1.35	1.39
35	BB	1205	A	O3'-P	-6.66	1.53	1.61
40	BG	118	U	O4'-C1'	-6.66	1.32	1.41
41	BH	38	G	N1-C2	-6.66	1.32	1.37
85	AA	1491	G	C2-N3	-6.66	1.27	1.32
34	BA	195	G	C2-N2	-6.66	1.27	1.34
34	BA	304	G	C2'-C1'	-6.66	1.46	1.53
34	BA	500	C	C3'-O3'	-6.66	1.32	1.42
34	BA	1075	U	C2'-C1'	-6.66	1.46	1.53
35	BB	51	U	C2-N3	-6.66	1.33	1.37
35	BB	1369	A	C2'-C1'	-6.66	1.46	1.53
36	BC	18	G	C4'-C3'	-6.66	1.45	1.53
38	BE	61	A	N9-C4	-6.66	1.33	1.37
40	BG	72	G	C2'-C1'	-6.66	1.46	1.53
85	AA	147	G	C2'-C1'	-6.66	1.46	1.53
85	AA	820	G	C5'-C4'	-6.66	1.43	1.51
85	AA	1297	G	O3'-P	-6.66	1.53	1.61
85	AA	2057	G	N9-C4	-6.66	1.32	1.38
34	BA	249	A	C5'-C4'	6.66	1.59	1.51
34	BA	542	A	O3'-P	-6.66	1.53	1.61
34	BA	1105	A	C1'-N9	-6.66	1.37	1.46
34	BA	1258	G	N9-C8	-6.66	1.33	1.37
40	BG	100	G	N9-C4	-6.66	1.32	1.38
40	BG	118	U	N1-C6	-6.66	1.31	1.38
41	BH	8	C	O3'-P	-6.66	1.53	1.61
85	AA	527	A	O3'-P	-6.66	1.53	1.61
34	BA	199	U	O3'-P	-6.66	1.53	1.61
34	BA	259	C	C3'-C2'	-6.66	1.45	1.52
34	BA	1161	G	O3'-P	-6.66	1.53	1.61
34	BA	1469	G	C6-N1	-6.66	1.34	1.39
35	BB	116	G	C2-N2	-6.66	1.27	1.34
35	BB	484	G	C2-N2	-6.66	1.27	1.34
35	BB	493	U	C3'-C2'	-6.66	1.45	1.52
35	BB	828	G	O3'-P	-6.66	1.53	1.61
35	BB	1074	U	C2'-C1'	-6.66	1.46	1.53
35	BB	1458	U	P-O5'	-6.66	1.53	1.59
38	BE	113	C	C2'-C1'	-6.66	1.46	1.53
83	Bx	48	GLY	N-CA	6.66	1.56	1.46
85	AA	810	C	C2'-C1'	-6.66	1.46	1.53
85	AA	855	G	C4'-C3'	-6.66	1.45	1.53
85	AA	876	U	O3'-P	-6.66	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2139	G	C3'-C2'	-6.66	1.45	1.52
34	BA	381	A	C3'-C2'	-6.66	1.45	1.52
41	BH	45	G	C2'-C1'	-6.66	1.46	1.53
34	BA	713	C	P-O5'	-6.66	1.53	1.59
34	BA	802	G	O3'-P	-6.66	1.53	1.61
34	BA	885	A	C1'-N9	-6.66	1.37	1.46
34	BA	1052	G	C2'-C1'	-6.66	1.46	1.53
34	BA	1063	G	P-O5'	-6.66	1.53	1.59
34	BA	1254	C	C2'-C1'	-6.66	1.46	1.53
34	BA	1815	G	N1-C2	-6.66	1.32	1.37
35	BB	555	G	N7-C5	-6.66	1.35	1.39
35	BB	1350	A	P-O5'	-6.66	1.53	1.59
35	BB	1405	G	C3'-C2'	-6.66	1.45	1.52
36	BC	26	U	C4'-C3'	-6.66	1.45	1.53
36	BC	158	U	P-O5'	-6.66	1.53	1.59
37	BD	78	C	C2'-C1'	-6.66	1.46	1.53
85	AA	560	C	C2'-C1'	-6.66	1.46	1.53
85	AA	741	G	P-O5'	-6.66	1.53	1.59
85	AA	769	C	C3'-C2'	-6.66	1.45	1.52
85	AA	1038	U	C5'-C4'	6.66	1.59	1.51
34	BA	447	U	C2-N3	-6.65	1.33	1.37
34	BA	1809	G	C5-C4	-6.65	1.33	1.38
40	BG	18	U	O3'-P	-6.65	1.53	1.61
85	AA	903	G	C2'-C1'	-6.65	1.46	1.53
34	BA	358	A	O3'-P	-6.65	1.53	1.61
34	BA	513	U	C5'-C4'	-6.65	1.43	1.51
35	BB	101	U	C5'-C4'	-6.65	1.43	1.51
35	BB	405	U	C2-N3	-6.65	1.33	1.37
35	BB	526	A	N3-C4	-6.65	1.30	1.34
35	BB	1033	U	C1'-N1	-6.65	1.37	1.46
35	BB	1063	C	C4'-O4'	-6.65	1.36	1.45
35	BB	1205	A	C4'-C3'	-6.65	1.45	1.53
35	BB	1529	G	C2'-C1'	-6.65	1.46	1.53
41	BH	45	G	C6-N1	-6.65	1.34	1.39
85	AA	597	A	C2'-C1'	-6.65	1.46	1.53
85	AA	988	C	O3'-P	-6.65	1.53	1.61
85	AA	1515	A	O4'-C1'	-6.65	1.33	1.41
85	AA	2210	C	C2'-C1'	-6.65	1.46	1.53
34	BA	67	A	C4'-C3'	-6.65	1.45	1.53
34	BA	245	U	C2-N3	-6.65	1.33	1.37
34	BA	894	G	C8-N7	-6.65	1.26	1.30
34	BA	971	G	C5-C4	-6.65	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1641	G	C4'-C3'	-6.65	1.45	1.53
35	BB	521	U	O3'-P	-6.65	1.53	1.61
85	AA	1542	A	O4'-C1'	-6.65	1.33	1.41
85	AA	1888	U	P-O5'	-6.65	1.53	1.59
85	AA	2139	G	C5-C6	-6.65	1.35	1.42
34	BA	1048	C	C2-N3	-6.65	1.30	1.35
34	BA	1156	U	O3'-P	-6.65	1.53	1.61
35	BB	563	A	C1'-N9	-6.65	1.37	1.46
35	BB	1209	A	N3-C4	-6.65	1.30	1.34
85	AA	558	U	C2'-C1'	-6.65	1.46	1.53
85	AA	1188	A	C4'-C3'	-6.65	1.45	1.53
34	BA	3	G	C2-N3	-6.65	1.27	1.32
34	BA	280	A	N9-C4	-6.65	1.33	1.37
34	BA	932	G	O3'-P	-6.65	1.53	1.61
34	BA	965	A	C4'-C3'	-6.65	1.45	1.53
34	BA	1248	A	N3-C4	-6.65	1.30	1.34
34	BA	1797	A	C5-C4	-6.65	1.34	1.38
35	BB	398	A	O3'-P	-6.65	1.53	1.61
35	BB	1025	A	O5'-C5'	6.65	1.55	1.44
36	BC	136	G	C6-N1	-6.65	1.34	1.39
38	BE	50	G	N1-C2	-6.65	1.32	1.37
85	AA	337	C	C2'-C1'	-6.65	1.46	1.53
85	AA	369	A	C5-C4	-6.65	1.34	1.38
85	AA	667	A	O3'-P	-6.65	1.53	1.61
85	AA	1145	U	O3'-P	-6.65	1.53	1.61
85	AA	2152	C	O3'-P	-6.65	1.53	1.61
85	AA	2193	A	N7-C5	-6.65	1.35	1.39
30	AW	48	TYR	CB-CG	-6.65	1.41	1.51
34	BA	138	C	C2-N3	-6.65	1.30	1.35
34	BA	774	A	N7-C5	-6.65	1.35	1.39
85	AA	363	A	P-O5'	-6.65	1.53	1.59
34	BA	483	A	N9-C8	-6.64	1.32	1.37
34	BA	518	C	C1'-N1	-6.64	1.37	1.46
34	BA	891	C	C2'-C1'	-6.64	1.46	1.53
35	BB	108	G	C5-C4	-6.64	1.33	1.38
35	BB	1209	A	C5'-C4'	6.64	1.59	1.51
35	BB	1291	G	C5-C4	-6.64	1.33	1.38
36	BC	92	C	C2'-C1'	-6.64	1.46	1.53
85	AA	790	A	C4'-O4'	-6.64	1.36	1.45
34	BA	289	A	C1'-N9	-6.64	1.37	1.46
34	BA	487	A	C6-N1	-6.64	1.30	1.35
34	BA	1175	G	N1-C2	-6.64	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	66	G	C6-N1	-6.64	1.34	1.39
35	BB	598	C	C2-N3	-6.64	1.30	1.35
35	BB	1214	U	C2-N3	-6.64	1.33	1.37
35	BB	1387	C	O3'-P	-6.64	1.53	1.61
36	BC	65	G	C3'-C2'	-6.64	1.45	1.52
40	BG	168	A	O4'-C1'	-6.64	1.33	1.41
85	AA	40	A	O3'-P	-6.64	1.53	1.61
85	AA	321	C	O3'-P	-6.64	1.53	1.61
85	AA	397	G	C1'-N9	-6.64	1.37	1.46
85	AA	491	G	C2-N2	-6.64	1.27	1.34
85	AA	1173	A	N9-C4	-6.64	1.33	1.37
34	BA	934	G	O4'-C1'	-6.64	1.33	1.41
34	BA	1252	G	C5-C4	-6.64	1.33	1.38
34	BA	1616	A	C2'-C1'	-6.64	1.46	1.53
35	BB	520	G	C5-C6	-6.64	1.35	1.42
35	BB	1002	G	C2-N2	-6.64	1.27	1.34
38	BE	98	C	O3'-P	-6.64	1.53	1.61
85	AA	95	U	C2'-C1'	-6.64	1.46	1.53
85	AA	608	A	P-O5'	-6.64	1.53	1.59
85	AA	2039	G	C1'-N9	-6.64	1.37	1.46
34	BA	1495	A	N7-C5	-6.64	1.35	1.39
35	BB	1375	G	C1'-N9	-6.64	1.37	1.46
36	BC	12	A	C6-N6	-6.64	1.28	1.33
36	BC	77	A	P-O5'	-6.64	1.53	1.59
37	BD	72	U	C1'-N1	-6.64	1.37	1.46
38	BE	193	A	N3-C4	-6.64	1.30	1.34
40	BG	40	G	N7-C5	-6.64	1.35	1.39
85	AA	40	A	C2'-C1'	-6.64	1.46	1.53
85	AA	481	A	P-O5'	-6.64	1.53	1.59
85	AA	668	A	N3-C4	-6.64	1.30	1.34
85	AA	685	U	O3'-P	-6.64	1.53	1.61
85	AA	1170	C	C2'-C1'	-6.64	1.46	1.53
38	BE	46	G	C1'-N9	-6.64	1.37	1.46
34	BA	29	U	P-O5'	-6.64	1.53	1.59
34	BA	61	G	N7-C5	-6.64	1.35	1.39
34	BA	125	G	C3'-C2'	-6.64	1.45	1.52
34	BA	1310	C	C2'-C1'	-6.64	1.46	1.53
34	BA	1566	G	N9-C4	-6.64	1.32	1.38
35	BB	102	G	C1'-N9	-6.64	1.37	1.46
35	BB	604	C	C3'-C2'	-6.64	1.45	1.52
35	BB	1338	U	C3'-C2'	-6.64	1.45	1.52
40	BG	9	G	C5-C4	-6.64	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	466	A	N9-C8	-6.64	1.32	1.37
85	AA	557	G	O3'-P	-6.64	1.53	1.61
85	AA	804	A	C2'-C1'	-6.64	1.46	1.53
85	AA	1268	C	C2-N3	-6.64	1.30	1.35
85	AA	1811	C	C5'-C4'	6.64	1.59	1.51
85	AA	2028	G	N7-C5	-6.64	1.35	1.39
34	BA	791	A	P-O5'	-6.63	1.53	1.59
34	BA	1172	C	P-O5'	-6.63	1.53	1.59
34	BA	1597	G	P-O5'	-6.63	1.53	1.59
34	BA	1598	U	C2-N3	-6.63	1.33	1.37
35	BB	9	G	N9-C4	-6.63	1.32	1.38
35	BB	375	G	C2-N2	-6.63	1.27	1.34
35	BB	628	A	C5-C4	-6.63	1.34	1.38
35	BB	649	A	C5-C4	-6.63	1.34	1.38
39	BF	13	U	N1-C2	-6.63	1.32	1.38
85	AA	109	G	P-O5'	-6.63	1.53	1.59
34	BA	124	G	P-O5'	-6.63	1.53	1.59
35	BB	1086	G	N7-C5	-6.63	1.35	1.39
85	AA	2003	C	C2'-C1'	-6.63	1.46	1.53
85	AA	2071	U	C2-N3	-6.63	1.33	1.37
34	BA	23	A	O3'-P	-6.63	1.53	1.61
34	BA	85	C	O3'-P	-6.63	1.53	1.61
34	BA	266	G	O3'-P	-6.63	1.53	1.61
35	BB	631	G	N7-C5	-6.63	1.35	1.39
35	BB	999	G	C3'-C2'	-6.63	1.45	1.52
36	BC	64	U	P-O5'	-6.63	1.53	1.59
36	BC	103	A	O3'-P	-6.63	1.53	1.61
38	BE	114	G	N9-C8	-6.63	1.33	1.37
39	BF	31	U	P-O5'	-6.63	1.53	1.59
41	BH	120	C	C2-N3	-6.63	1.30	1.35
85	AA	987	C	C2-N3	-6.63	1.30	1.35
34	BA	859	G	C2'-C1'	-6.63	1.46	1.53
34	BA	1088	G	P-O5'	-6.63	1.53	1.59
35	BB	459	U	O3'-P	-6.63	1.53	1.61
35	BB	1298	C	C4'-C3'	-6.63	1.45	1.53
38	BE	12	A	O3'-P	-6.63	1.53	1.61
85	AA	471	U	O4'-C1'	-6.63	1.33	1.41
85	AA	927	A	O3'-P	-6.63	1.53	1.61
85	AA	1482	C	N3-C4	-6.63	1.29	1.33
34	BA	837	U	C2'-C1'	-6.63	1.46	1.53
34	BA	1278	A	N9-C4	-6.63	1.33	1.37
35	BB	502	C	C1'-N1	-6.63	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	846	A	N9-C4	-6.63	1.33	1.37
35	BB	999	G	C5-C4	-6.63	1.33	1.38
35	BB	1292	G	C5'-C4'	-6.63	1.43	1.51
35	BB	1296	A	C2'-C1'	-6.63	1.46	1.53
38	BE	130	G	C1'-N9	-6.63	1.37	1.46
40	BG	116	G	C6-N1	-6.63	1.34	1.39
85	AA	395	G	C5'-C4'	-6.63	1.43	1.51
85	AA	661	C	C3'-C2'	-6.63	1.45	1.52
85	AA	1219	A	O3'-P	-6.63	1.53	1.61
85	AA	1693	C	C2'-C1'	-6.63	1.46	1.53
34	BA	577	U	O3'-P	-6.63	1.53	1.61
34	BA	1332	U	C3'-C2'	-6.63	1.45	1.52
35	BB	35	G	C6-N1	-6.63	1.34	1.39
35	BB	1034	U	C1'-N1	-6.63	1.37	1.46
35	BB	1036	G	C1'-N9	-6.63	1.37	1.46
36	BC	169	G	P-O5'	-6.63	1.53	1.59
37	BD	6	C	P-O5'	-6.63	1.53	1.59
38	BE	176	G	N3-C4	-6.63	1.30	1.35
39	BF	51	C	C2-N3	-6.63	1.30	1.35
85	AA	102	A	N7-C5	-6.63	1.35	1.39
85	AA	1136	A	N7-C5	-6.63	1.35	1.39
85	AA	1510	A	N7-C5	-6.63	1.35	1.39
34	BA	723	C	C2'-C1'	-6.62	1.46	1.53
35	BB	58	G	C1'-N9	-6.62	1.37	1.46
35	BB	994	A	C3'-C2'	6.62	1.60	1.52
35	BB	1497	C	P-O5'	-6.62	1.53	1.59
38	BE	141	A	C8-N7	-6.62	1.26	1.31
34	BA	189	G	P-O5'	-6.62	1.53	1.59
34	BA	266	G	C2-N2	-6.62	1.27	1.34
34	BA	1321	A	C1'-N9	-6.62	1.37	1.46
34	BA	1419	A	C2'-C1'	-6.62	1.46	1.53
35	BB	97	U	C2'-C1'	-6.62	1.46	1.53
35	BB	592	G	P-O5'	-6.62	1.53	1.59
35	BB	1534	U	C3'-C2'	-6.62	1.45	1.52
38	BE	96	G	N7-C5	-6.62	1.35	1.39
38	BE	147	G	N9-C4	-6.62	1.32	1.38
85	AA	688	C	P-O5'	-6.62	1.53	1.59
34	BA	361	C	O3'-P	-6.62	1.53	1.61
34	BA	409	A	C3'-C2'	-6.62	1.45	1.52
34	BA	1598	U	C4'-C3'	-6.62	1.45	1.53
35	BB	1062	G	C5-C4	-6.62	1.33	1.38
35	BB	1444	U	C3'-C2'	-6.62	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1471	A	N7-C5	-6.62	1.35	1.39
36	BC	16	A	C3'-C2'	-6.62	1.45	1.52
36	BC	120	G	C1'-N9	-6.62	1.37	1.46
37	BD	49	A	N9-C4	-6.62	1.33	1.37
85	AA	1105	G	N1-C2	-6.62	1.32	1.37
85	AA	1910	A	P-O5'	-6.62	1.53	1.59
35	BB	18	A	C2'-C1'	-6.62	1.46	1.53
35	BB	664	A	O4'-C1'	-6.62	1.33	1.41
35	BB	1153	G	P-O5'	-6.62	1.53	1.59
40	BG	98	A	N9-C8	-6.62	1.32	1.37
85	AA	1450	U	C2'-C1'	-6.62	1.46	1.53
34	BA	688	G	C3'-C2'	-6.62	1.45	1.52
34	BA	987	C	C2-N3	-6.62	1.30	1.35
34	BA	1006	G	O3'-P	-6.62	1.53	1.61
36	BC	46	G	O3'-P	-6.62	1.53	1.61
85	AA	352	G	C3'-C2'	-6.62	1.45	1.52
34	BA	145	U	C2-N3	-6.62	1.33	1.37
35	BB	25	A	C2'-C1'	-6.62	1.46	1.53
35	BB	33	A	C2'-C1'	-6.62	1.46	1.53
34	BA	76	U	N3-C4	-6.62	1.32	1.38
34	BA	1530	G	C1'-N9	-6.62	1.37	1.46
35	BB	39	C	C2'-C1'	-6.62	1.46	1.53
35	BB	1200	A	O3'-P	-6.62	1.53	1.61
36	BC	148	C	N1-C6	-6.62	1.33	1.37
37	BD	85	C	C2'-C1'	-6.62	1.46	1.53
40	BG	115	C	N1-C6	-6.62	1.33	1.37
41	BH	34	G	C8-N7	-6.62	1.26	1.30
85	AA	643	C	P-O5'	-6.62	1.53	1.59
34	BA	891	C	O3'-P	-6.61	1.53	1.61
34	BA	1504	A	C3'-C2'	-6.61	1.45	1.52
35	BB	630	A	N9-C4	-6.61	1.33	1.37
35	BB	1052	G	P-O5'	-6.61	1.53	1.59
35	BB	1486	C	C2'-C1'	-6.61	1.46	1.53
38	BE	162	U	N1-C6	-6.61	1.31	1.38
40	BG	32	U	P-O5'	-6.61	1.53	1.59
40	BG	88	G	C1'-N9	-6.61	1.37	1.46
85	AA	30	G	O3'-P	-6.61	1.53	1.61
85	AA	112	A	C1'-N9	-6.61	1.37	1.46
85	AA	461	G	N9-C4	-6.61	1.32	1.38
85	AA	471	U	C3'-C2'	-6.61	1.45	1.52
85	AA	1299	A	O3'-P	-6.61	1.53	1.61
85	AA	1864	G	O3'-P	-6.61	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
86	AB	13	C	C2'-C1'	-6.61	1.46	1.53
34	BA	662	U	C2'-C1'	-6.61	1.46	1.53
34	BA	908	G	N9-C4	-6.61	1.32	1.38
34	BA	1151	A	O3'-P	-6.61	1.53	1.61
34	BA	1436	A	C5-C4	-6.61	1.34	1.38
35	BB	595	U	O3'-P	-6.61	1.53	1.61
85	AA	1868	G	P-O5'	-6.61	1.53	1.59
34	BA	290	G	C6-N1	-6.61	1.34	1.39
34	BA	679	U	O4'-C1'	-6.61	1.33	1.41
34	BA	1450	G	C4'-O4'	-6.61	1.36	1.45
35	BB	412	A	N9-C8	-6.61	1.32	1.37
35	BB	638	G	C3'-C2'	-6.61	1.45	1.52
35	BB	690	C	C2-N3	-6.61	1.30	1.35
39	BF	71	G	N9-C8	-6.61	1.33	1.37
40	BG	28	A	C1'-N9	-6.61	1.37	1.46
85	AA	728	U	P-O5'	-6.61	1.53	1.59
85	AA	1224	C	C2-N3	-6.61	1.30	1.35
34	BA	315	U	P-O5'	-6.61	1.53	1.59
34	BA	358	A	C2'-C1'	-6.61	1.46	1.53
34	BA	1472	G	O3'-P	-6.61	1.53	1.61
35	BB	359	A	O3'-P	-6.61	1.53	1.61
35	BB	622	G	O3'-P	-6.61	1.53	1.61
85	AA	679	A	C5-C4	-6.61	1.34	1.38
34	BA	17	A	C1'-N9	-6.61	1.37	1.46
34	BA	334	G	N9-C8	-6.61	1.33	1.37
34	BA	545	U	C2'-C1'	-6.61	1.46	1.53
34	BA	892	C	N1-C6	-6.61	1.33	1.37
34	BA	1675	C	C2-N3	-6.61	1.30	1.35
35	BB	969	C	P-O5'	-6.61	1.53	1.59
35	BB	1050	A	N9-C4	-6.61	1.33	1.37
35	BB	1389	C	O3'-P	-6.61	1.53	1.61
38	BE	133	C	C2'-C1'	-6.61	1.46	1.53
41	BH	59	G	P-O5'	-6.61	1.53	1.59
85	AA	535	G	O3'-P	-6.61	1.53	1.61
85	AA	821	U	P-O5'	-6.61	1.53	1.59
85	AA	2082	C	N3-C4	-6.61	1.29	1.33
34	BA	828	A	P-O5'	-6.61	1.53	1.59
34	BA	1601	C	C2'-C1'	-6.61	1.46	1.53
34	BA	1667	G	C5-C4	-6.61	1.33	1.38
34	BA	1785	G	N9-C4	-6.61	1.32	1.38
35	BB	375	G	N9-C8	-6.61	1.33	1.37
35	BB	582	G	C2'-C1'	-6.61	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	857	G	N7-C5	-6.61	1.35	1.39
35	BB	1504	U	C5'-C4'	6.61	1.59	1.51
40	BG	96	C	C1'-N1	-6.61	1.37	1.46
41	BH	104	U	P-O5'	-6.61	1.53	1.59
85	AA	875	C	C2'-C1'	-6.61	1.46	1.53
85	AA	1167	G	O3'-P	-6.61	1.53	1.61
85	AA	1796	C	P-O5'	-6.61	1.53	1.59
86	AB	70	G	N7-C5	-6.61	1.35	1.39
35	BB	54	U	C2-N3	-6.60	1.33	1.37
34	BA	211	C	O3'-P	-6.60	1.53	1.61
34	BA	388	A	N7-C5	-6.60	1.35	1.39
34	BA	426	A	C5-C4	-6.60	1.34	1.38
34	BA	813	C	O4'-C1'	-6.60	1.33	1.41
34	BA	870	C	N1-C6	6.60	1.41	1.37
34	BA	1033	G	C2'-C1'	-6.60	1.46	1.53
34	BA	1145	U	O3'-P	-6.60	1.53	1.61
35	BB	32	C	P-O5'	-6.60	1.53	1.59
35	BB	81	A	C3'-C2'	-6.60	1.45	1.52
35	BB	1362	G	C5-C4	-6.60	1.33	1.38
35	BB	1396	G	C1'-N9	-6.60	1.37	1.46
36	BC	121	G	C3'-C2'	-6.60	1.45	1.52
85	AA	104	C	N1-C6	-6.60	1.33	1.37
85	AA	546	U	O3'-P	-6.60	1.53	1.61
85	AA	880	A	O3'-P	-6.60	1.53	1.61
85	AA	1549	G	C3'-C2'	-6.60	1.45	1.52
34	BA	690	G	C5-C4	-6.60	1.33	1.38
35	BB	133	G	P-O5'	-6.60	1.53	1.59
85	AA	487	G	N1-C2	-6.60	1.32	1.37
85	AA	687	G	C1'-N9	-6.60	1.37	1.46
85	AA	2201	A	O3'-P	-6.60	1.53	1.61
34	BA	307	C	C3'-C2'	-6.60	1.45	1.52
34	BA	968	G	O3'-P	-6.60	1.53	1.61
34	BA	1286	C	C3'-C2'	-6.60	1.45	1.52
34	BA	1312	A	C3'-C2'	-6.60	1.45	1.52
34	BA	1562	G	N1-C2	-6.60	1.32	1.37
35	BB	402	G	C6-N1	-6.60	1.34	1.39
35	BB	1039	A	C4'-C3'	-6.60	1.45	1.53
36	BC	8	C	O3'-P	-6.60	1.53	1.61
36	BC	106	G	C2-N2	-6.60	1.27	1.34
38	BE	49	A	P-O5'	-6.60	1.53	1.59
85	AA	881	C	C2'-C1'	-6.60	1.46	1.53
85	AA	2141	G	C8-N7	-6.60	1.26	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	402	G	O3'-P	-6.60	1.53	1.61
34	BA	1229	G	C2-N2	-6.60	1.27	1.34
34	BA	1793	G	O3'-P	-6.60	1.53	1.61
35	BB	1202	G	O3'-P	-6.60	1.53	1.61
35	BB	1213	U	C2'-C1'	-6.60	1.46	1.53
35	BB	1421	C	O3'-P	-6.60	1.53	1.61
35	BB	1515	C	O3'-P	-6.60	1.53	1.61
36	BC	34	U	O4'-C1'	-6.60	1.33	1.41
39	BF	72	A	N7-C5	-6.60	1.35	1.39
85	AA	57	G	N1-C2	-6.60	1.32	1.37
85	AA	393	C	C5'-C4'	-6.60	1.43	1.51
85	AA	715	G	N9-C4	6.60	1.43	1.38
85	AA	965	G	C2-N2	-6.60	1.27	1.34
34	BA	27	G	C5-C4	-6.60	1.33	1.38
34	BA	1697	U	P-O5'	-6.60	1.53	1.59
35	BB	1137	G	C2-N3	-6.60	1.27	1.32
85	AA	1219	A	C5-C6	-6.60	1.35	1.41
85	AA	1349	A	P-O5'	-6.60	1.53	1.59
85	AA	1503	G	C5-C4	-6.60	1.33	1.38
85	AA	2141	G	N7-C5	-6.60	1.35	1.39
85	AA	2146	G	O4'-C1'	-6.60	1.33	1.41
34	BA	719	G	C2-N2	-6.59	1.27	1.34
34	BA	902	C	C2'-C1'	-6.59	1.46	1.53
34	BA	1472	G	C2-N2	-6.59	1.27	1.34
34	BA	1474	G	N9-C4	-6.59	1.32	1.38
34	BA	1818	A	N7-C5	-6.59	1.35	1.39
35	BB	1508	G	C6-N1	-6.59	1.34	1.39
37	BD	47	U	C5'-C4'	6.59	1.59	1.51
85	AA	118	C	C1'-N1	-6.59	1.37	1.46
85	AA	156	G	C5-C6	-6.59	1.35	1.42
85	AA	311	U	C2-N3	-6.59	1.33	1.37
85	AA	900	G	P-O5'	-6.59	1.53	1.59
85	AA	1186	C	C2-N3	-6.59	1.30	1.35
85	AA	1206	A	C2'-C1'	-6.59	1.46	1.53
85	AA	1292	A	C1'-N9	-6.59	1.37	1.46
34	BA	1335	A	C4'-C3'	-6.59	1.45	1.53
34	BA	1712	U	C1'-N1	-6.59	1.37	1.46
36	BC	145	G	P-O5'	-6.59	1.53	1.59
85	AA	427	G	C6-N1	-6.59	1.34	1.39
85	AA	1674	G	P-O5'	-6.59	1.53	1.59
34	BA	980	C	O3'-P	-6.59	1.53	1.61
34	BA	1000	G	C3'-C2'	-6.59	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1047	U	C2-N3	-6.59	1.33	1.37
34	BA	1427	U	C4'-C3'	-6.59	1.45	1.53
35	BB	57	G	C6-N1	-6.59	1.34	1.39
35	BB	415	A	C5-C4	-6.59	1.34	1.38
35	BB	1004	A	C2'-C1'	-6.59	1.46	1.53
85	AA	327	G	C6-N1	-6.59	1.34	1.39
34	BA	377	G	N1-C2	-6.59	1.32	1.37
34	BA	436	U	N3-C4	-6.59	1.32	1.38
34	BA	1156	U	C2'-C1'	-6.59	1.46	1.53
34	BA	1440	C	P-O5'	-6.59	1.53	1.59
34	BA	1543	A	P-O5'	-6.59	1.53	1.59
34	BA	1550	G	O3'-P	-6.59	1.53	1.61
37	BD	20	C	O3'-P	-6.59	1.53	1.61
40	BG	58	G	C6-N1	-6.59	1.34	1.39
85	AA	309	G	C2-N2	-6.59	1.27	1.34
85	AA	747	U	C3'-C2'	-6.59	1.45	1.52
85	AA	897	A	O3'-P	-6.59	1.53	1.61
85	AA	965	G	C6-N1	-6.59	1.34	1.39
85	AA	1203	G	N9-C4	-6.59	1.32	1.38
85	AA	1518	A	C1'-N9	-6.59	1.37	1.46
85	AA	1578	G	C2'-C1'	-6.59	1.46	1.53
85	AA	2114	U	C3'-C2'	-6.59	1.45	1.52
34	BA	1576	C	C3'-C2'	-6.59	1.45	1.52
34	BA	1614	G	O3'-P	-6.59	1.53	1.61
35	BB	81	A	C2'-C1'	-6.59	1.46	1.53
35	BB	370	A	C1'-N9	-6.59	1.37	1.46
35	BB	630	A	N7-C5	-6.59	1.35	1.39
36	BC	64	U	C2-N3	-6.59	1.33	1.37
37	BD	79	G	C3'-C2'	-6.59	1.45	1.52
41	BH	109	G	C4'-O4'	-6.59	1.36	1.45
34	BA	723	C	N1-C2	-6.59	1.33	1.40
34	BA	1013	A	C2'-C1'	-6.59	1.46	1.53
34	BA	1574	C	C2-N3	-6.59	1.30	1.35
35	BB	1205	A	C8-N7	-6.59	1.26	1.31
35	BB	1364	C	O3'-P	-6.59	1.53	1.61
35	BB	1430	G	C6-N1	-6.59	1.34	1.39
35	BB	1446	C	P-O5'	-6.59	1.53	1.59
38	BE	177	U	C3'-C2'	-6.59	1.45	1.52
85	AA	2127	G	N7-C5	-6.59	1.35	1.39
34	BA	356	C	C4'-C3'	-6.58	1.46	1.53
34	BA	751	A	N7-C5	-6.58	1.35	1.39
34	BA	1591	G	C3'-C2'	-6.58	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	454	U	C2-N3	-6.58	1.33	1.37
35	BB	999	G	C6-N1	-6.58	1.34	1.39
85	AA	88	G	C5-C4	-6.58	1.33	1.38
85	AA	2147	A	C8-N7	-6.58	1.26	1.31
34	BA	57	A	C3'-C2'	-6.58	1.45	1.52
34	BA	592	G	P-O5'	-6.58	1.53	1.59
34	BA	759	A	C4'-C3'	-6.58	1.46	1.53
34	BA	806	U	P-O5'	-6.58	1.53	1.59
34	BA	1001	G	C2-N2	-6.58	1.27	1.34
34	BA	1007	G	N3-C4	-6.58	1.30	1.35
34	BA	1224	A	O3'-P	-6.58	1.53	1.61
34	BA	1273	U	O3'-P	-6.58	1.53	1.61
35	BB	98	A	P-O5'	-6.58	1.53	1.59
35	BB	547	A	N9-C4	-6.58	1.33	1.37
35	BB	623	A	P-O5'	-6.58	1.53	1.59
36	BC	31	A	O4'-C1'	-6.58	1.33	1.41
40	BG	33	G	C2-N3	-6.58	1.27	1.32
41	BH	17	A	C2'-C1'	-6.58	1.46	1.53
85	AA	889	G	C2-N2	-6.58	1.27	1.34
85	AA	1140	G	N1-C2	-6.58	1.32	1.37
85	AA	1697	C	O3'-P	-6.58	1.53	1.61
34	BA	1704	G	O3'-P	-6.58	1.53	1.61
35	BB	366	G	O3'-P	-6.58	1.53	1.61
40	BG	58	G	C2'-C1'	-6.58	1.46	1.53
40	BG	141	A	C1'-N9	-6.58	1.37	1.46
85	AA	485	A	C5-C4	-6.58	1.34	1.38
85	AA	1470	A	N9-C8	-6.58	1.32	1.37
34	BA	53	G	N1-C2	-6.58	1.32	1.37
34	BA	65	A	N3-C4	-6.58	1.30	1.34
34	BA	1251	A	O3'-P	-6.58	1.53	1.61
35	BB	964	G	C3'-C2'	-6.58	1.45	1.52
38	BE	94	U	C4'-O4'	-6.58	1.36	1.45
38	BE	128	G	C1'-N9	-6.58	1.37	1.46
38	BE	131	C	N1-C6	-6.58	1.33	1.37
85	AA	571	G	C2'-C1'	-6.58	1.46	1.53
85	AA	1525	C	O3'-P	-6.58	1.53	1.61
34	BA	725	C	P-O5'	-6.58	1.53	1.59
34	BA	832	C	P-O5'	-6.58	1.53	1.59
34	BA	1243	A	N7-C5	-6.58	1.35	1.39
34	BA	1642	A	O4'-C1'	-6.58	1.33	1.41
34	BA	1804	A	O3'-P	-6.58	1.53	1.61
35	BB	356	C	P-O5'	-6.58	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	668	A	P-O5'	-6.58	1.53	1.59
35	BB	971	A	C8-N7	-6.58	1.26	1.31
35	BB	1070	G	C3'-C2'	-6.58	1.45	1.52
35	BB	1118	G	O3'-P	-6.58	1.53	1.61
85	AA	921	C	O3'-P	-6.58	1.53	1.61
85	AA	1589	G	C6-N1	-6.58	1.34	1.39
34	BA	811	C	O3'-P	-6.58	1.53	1.61
35	BB	438	G	C2'-C1'	-6.58	1.46	1.53
35	BB	1081	U	P-O5'	-6.58	1.53	1.59
34	BA	76	U	C2'-C1'	-6.58	1.46	1.53
34	BA	267	G	N9-C8	-6.58	1.33	1.37
34	BA	277	A	C5-C4	-6.58	1.34	1.38
34	BA	521	C	C4'-C3'	-6.58	1.46	1.53
34	BA	803	U	C2-N3	-6.58	1.33	1.37
34	BA	1287	G	C2-N2	-6.58	1.27	1.34
34	BA	1804	A	N3-C4	-6.58	1.30	1.34
35	BB	509	A	P-O5'	-6.58	1.53	1.59
35	BB	567	G	P-O5'	-6.58	1.53	1.59
35	BB	1086	G	C6-N1	-6.58	1.34	1.39
35	BB	1147	G	C1'-N9	-6.58	1.37	1.46
35	BB	1292	G	C2-N2	-6.58	1.27	1.34
35	BB	1399	A	C5-C4	-6.58	1.34	1.38
38	BE	8	G	C3'-C2'	-6.58	1.45	1.52
38	BE	59	U	C3'-C2'	-6.58	1.45	1.52
38	BE	139	U	N1-C6	-6.58	1.32	1.38
40	BG	179	C	C4'-C3'	-6.58	1.46	1.53
85	AA	756	G	N7-C5	-6.58	1.35	1.39
85	AA	802	A	C8-N7	-6.58	1.26	1.31
85	AA	1176	C	C2'-C1'	-6.58	1.46	1.53
34	BA	514	U	C4'-C3'	-6.57	1.46	1.53
34	BA	532	C	C4'-C3'	-6.57	1.46	1.53
34	BA	1593	U	P-O5'	-6.57	1.53	1.59
35	BB	1360	A	O3'-P	-6.57	1.53	1.61
35	BB	1457	A	O4'-C1'	-6.57	1.33	1.41
36	BC	86	U	O3'-P	-6.57	1.53	1.61
38	BE	186	C	C1'-N1	-6.57	1.37	1.46
40	BG	87	G	O3'-P	-6.57	1.53	1.61
40	BG	91	U	C2'-C1'	-6.57	1.46	1.53
85	AA	96	C	C3'-C2'	-6.57	1.45	1.52
85	AA	211	C	O4'-C1'	-6.57	1.33	1.41
85	AA	533	C	C4-C5	-6.57	1.37	1.43
85	AA	930	G	O3'-P	-6.57	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	977	U	C3'-C2'	-6.57	1.45	1.52
85	AA	1509	A	C8-N7	-6.57	1.26	1.31
34	BA	87	G	C2'-C1'	-6.57	1.46	1.53
34	BA	191	G	N1-C2	-6.57	1.32	1.37
34	BA	1071	G	P-O5'	-6.57	1.53	1.59
35	BB	101	U	C3'-C2'	-6.57	1.45	1.52
39	BF	48	G	N7-C5	-6.57	1.35	1.39
85	AA	366	A	C3'-C2'	-6.57	1.45	1.52
85	AA	700	U	C3'-C2'	-6.57	1.45	1.52
85	AA	1468	G	P-O5'	-6.57	1.53	1.59
34	BA	700	G	C2-N2	-6.57	1.27	1.34
35	BB	88	U	C3'-C2'	-6.57	1.45	1.52
35	BB	1537	C	C3'-C2'	-6.57	1.45	1.52
77	Br	140	GLY	CA-C	-6.57	1.41	1.51
85	AA	1810	C	C4'-C3'	-6.57	1.46	1.53
85	AA	2048	C	C2-N3	-6.57	1.30	1.35
34	BA	455	A	P-O5'	-6.57	1.53	1.59
40	BG	146	C	O3'-P	-6.57	1.53	1.61
85	AA	965	G	N7-C5	-6.57	1.35	1.39
34	BA	166	G	N9-C8	-6.57	1.33	1.37
34	BA	658	C	C2-N3	-6.57	1.30	1.35
34	BA	776	U	N3-C4	-6.57	1.32	1.38
34	BA	979	G	N3-C4	-6.57	1.30	1.35
35	BB	448	G	N7-C5	-6.57	1.35	1.39
35	BB	508	U	P-O5'	-6.57	1.53	1.59
35	BB	637	G	N9-C4	-6.57	1.32	1.38
35	BB	640	A	N7-C5	-6.57	1.35	1.39
35	BB	794	G	O3'-P	-6.57	1.53	1.61
35	BB	832	C	N1-C6	-6.57	1.33	1.37
35	BB	1086	G	C5-C4	-6.57	1.33	1.38
35	BB	1313	C	C3'-C2'	-6.57	1.45	1.52
35	BB	1364	C	C2-N3	-6.57	1.30	1.35
38	BE	46	G	C5-C4	-6.57	1.33	1.38
38	BE	114	G	N9-C4	-6.57	1.32	1.38
40	BG	110	U	O3'-P	-6.57	1.53	1.61
85	AA	18	C	C3'-C2'	-6.57	1.45	1.52
85	AA	313	A	C2'-C1'	-6.57	1.46	1.53
85	AA	710	A	P-O5'	-6.57	1.53	1.59
85	AA	731	U	C3'-C2'	-6.57	1.45	1.52
85	AA	2004	U	C4'-O4'	-6.57	1.37	1.45
85	AA	2075	C	O4'-C1'	-6.57	1.33	1.41
85	AA	2192	A	O3'-P	-6.57	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1001	G	C2'-C1'	-6.57	1.46	1.53
34	BA	1019	C	C2'-C1'	-6.57	1.46	1.53
35	BB	514	G	C3'-C2'	-6.57	1.45	1.52
35	BB	1204	C	C2'-C1'	-6.57	1.46	1.53
35	BB	1306	G	C2'-C1'	-6.57	1.46	1.53
40	BG	75	C	C3'-C2'	-6.57	1.45	1.52
85	AA	313	A	N9-C8	-6.57	1.32	1.37
85	AA	467	U	C2-N3	-6.57	1.33	1.37
85	AA	1482	C	P-O5'	-6.57	1.53	1.59
85	AA	2150	G	C2'-C1'	-6.57	1.46	1.53
85	AA	2195	A	C3'-C2'	-6.57	1.45	1.52
34	BA	798	G	O3'-P	-6.56	1.53	1.61
34	BA	1152	A	C1'-N9	-6.56	1.37	1.46
34	BA	1314	A	P-O5'	-6.56	1.53	1.59
34	BA	1516	G	C6-N1	-6.56	1.34	1.39
35	BB	487	A	P-O5'	-6.56	1.53	1.59
35	BB	644	A	N7-C5	-6.56	1.35	1.39
35	BB	810	G	C6-N1	-6.56	1.34	1.39
36	BC	41	A	C8-N7	-6.56	1.26	1.31
36	BC	151	G	C3'-C2'	-6.56	1.45	1.52
85	AA	176	C	C2-N3	-6.56	1.30	1.35
85	AA	1578	G	N9-C4	-6.56	1.32	1.38
34	BA	65	A	C5-C4	-6.56	1.34	1.38
34	BA	356	C	P-O5'	-6.56	1.53	1.59
34	BA	908	G	O3'-P	-6.56	1.53	1.61
35	BB	1349	U	N3-C4	-6.56	1.32	1.38
36	BC	58	G	C1'-N9	-6.56	1.37	1.46
41	BH	52	G	O3'-P	-6.56	1.53	1.61
44	BK	181	TYR	CB-CG	-6.56	1.41	1.51
85	AA	2055	G	C1'-N9	-6.56	1.37	1.46
85	AA	2197	A	N7-C5	-6.56	1.35	1.39
34	BA	1838	U	C2'-C1'	-6.56	1.46	1.53
35	BB	560	C	C1'-N1	-6.56	1.37	1.46
35	BB	1221	G	O3'-P	-6.56	1.53	1.61
35	BB	1452	U	O3'-P	-6.56	1.53	1.61
36	BC	75	G	C2-N2	-6.56	1.27	1.34
38	BE	33	C	O3'-P	-6.56	1.53	1.61
85	AA	737	G	O3'-P	-6.56	1.53	1.61
34	BA	298	G	O3'-P	-6.56	1.53	1.61
34	BA	1836	A	C2'-C1'	-6.56	1.46	1.53
35	BB	1408	G	P-O5'	-6.56	1.53	1.59
37	BD	46	G	C4'-C3'	-6.56	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1117	G	O3'-P	-6.56	1.53	1.61
85	AA	1281	G	N9-C8	-6.56	1.33	1.37
85	AA	1586	C	P-O5'	-6.56	1.53	1.59
85	AA	2180	C	C3'-C2'	-6.56	1.45	1.52
34	BA	417	A	P-O5'	-6.56	1.53	1.59
34	BA	539	C	O3'-P	-6.56	1.53	1.61
34	BA	661	C	O3'-P	-6.56	1.53	1.61
34	BA	1421	A	O3'-P	-6.56	1.53	1.61
34	BA	1687	A	N9-C4	-6.56	1.33	1.37
35	BB	646	U	O3'-P	-6.56	1.53	1.61
38	BE	163	A	C2'-C1'	-6.56	1.46	1.53
39	BF	49	C	C2'-C1'	-6.56	1.46	1.53
85	AA	353	G	C1'-N9	-6.56	1.37	1.46
85	AA	1509	A	C1'-N9	-6.56	1.37	1.46
85	AA	1734	A	C2'-C1'	-6.56	1.46	1.53
85	AA	2178	A	C1'-N9	-6.56	1.37	1.46
35	BB	1284	U	C3'-C2'	-6.56	1.45	1.52
41	BH	29	G	N3-C4	-6.56	1.30	1.35
41	BH	47	G	C2-N2	-6.56	1.27	1.34
85	AA	1519	A	C5-C4	-6.56	1.34	1.38
34	BA	698	U	P-O5'	-6.55	1.53	1.59
34	BA	1152	A	C4'-C3'	-6.55	1.46	1.53
34	BA	1409	A	N7-C5	-6.55	1.35	1.39
34	BA	1791	C	P-O5'	-6.55	1.53	1.59
35	BB	549	U	C3'-C2'	-6.55	1.45	1.52
41	BH	63	G	C8-N7	-6.55	1.27	1.30
85	AA	469	G	C5-C6	-6.55	1.35	1.42
85	AA	540	A	C3'-C2'	-6.55	1.45	1.52
85	AA	1599	G	O3'-P	-6.55	1.53	1.61
34	BA	981	A	N7-C5	-6.55	1.35	1.39
34	BA	1801	G	C3'-C2'	-6.55	1.45	1.52
37	BD	47	U	C2'-C1'	-6.55	1.46	1.53
39	BF	38	C	O3'-P	-6.55	1.53	1.61
40	BG	99	A	P-O5'	-6.55	1.53	1.59
85	AA	1532	G	C1'-N9	-6.55	1.37	1.46
34	BA	626	G	P-O5'	-6.55	1.53	1.59
34	BA	969	A	C1'-N9	-6.55	1.37	1.46
34	BA	1011	G	N3-C4	-6.55	1.30	1.35
34	BA	1230	G	C6-N1	-6.55	1.34	1.39
34	BA	1401	C	C2'-C1'	-6.55	1.46	1.53
34	BA	1431	G	N3-C4	-6.55	1.30	1.35
34	BA	1531	G	C6-N1	-6.55	1.34	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1599	A	C1'-N9	-6.55	1.37	1.46
34	BA	1635	A	O3'-P	-6.55	1.53	1.61
35	BB	583	G	O3'-P	-6.55	1.53	1.61
35	BB	1180	G	C2-N2	-6.55	1.27	1.34
39	BF	65	U	O4'-C1'	-6.55	1.33	1.41
41	BH	115	A	C5-C4	-6.55	1.34	1.38
85	AA	753	U	C5'-C4'	6.55	1.59	1.51
34	BA	456	G	C2'-C1'	-6.55	1.46	1.53
34	BA	890	G	C4'-O4'	-6.55	1.37	1.45
34	BA	1034	U	P-O5'	-6.55	1.53	1.59
34	BA	1050	A	C2'-C1'	-6.55	1.46	1.53
34	BA	1093	G	C2'-C1'	-6.55	1.46	1.53
34	BA	1523	U	C3'-C2'	-6.55	1.45	1.52
34	BA	1808	A	N3-C4	-6.55	1.30	1.34
35	BB	615	A	N7-C5	-6.55	1.35	1.39
35	BB	838	G	N9-C4	6.55	1.43	1.38
35	BB	1181	A	P-O5'	-6.55	1.53	1.59
85	AA	175	A	O3'-P	-6.55	1.53	1.61
85	AA	432	A	O3'-P	-6.55	1.53	1.61
85	AA	859	G	C5-C4	-6.55	1.33	1.38
85	AA	1167	G	N3-C4	-6.55	1.30	1.35
85	AA	1178	A	O3'-P	-6.55	1.53	1.61
34	BA	817	U	P-O5'	-6.55	1.53	1.59
34	BA	1059	U	C4'-C3'	-6.55	1.46	1.53
34	BA	1550	G	N1-C2	-6.55	1.32	1.37
34	BA	1655	G	C1'-N9	-6.55	1.37	1.46
34	BA	1711	G	O3'-P	-6.55	1.53	1.61
35	BB	495	A	C4'-C3'	-6.55	1.46	1.53
34	BA	80	U	O4'-C1'	-6.55	1.33	1.41
34	BA	96	G	C1'-N9	-6.55	1.37	1.46
34	BA	120	A	C5'-C4'	6.55	1.59	1.51
34	BA	597	C	C4-C5	-6.55	1.37	1.43
34	BA	1089	U	C4'-C3'	-6.55	1.46	1.53
35	BB	401	U	O3'-P	-6.55	1.53	1.61
35	BB	511	A	C2'-C1'	-6.55	1.46	1.53
35	BB	684	U	C3'-C2'	-6.55	1.45	1.52
35	BB	790	A	N9-C8	-6.55	1.32	1.37
35	BB	978	C	C4-N4	-6.55	1.28	1.33
35	BB	1088	C	P-O5'	-6.55	1.53	1.59
35	BB	1407	U	C2-N3	-6.55	1.33	1.37
40	BG	88	G	C4'-C3'	-6.55	1.46	1.53
85	AA	27	U	P-O5'	-6.55	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1215	A	C3'-C2'	-6.55	1.45	1.52
85	AA	2184	A	C5-C4	-6.55	1.34	1.38
85	AA	931	G	N7-C5	-6.54	1.35	1.39
34	BA	1253	G	P-O5'	-6.54	1.53	1.59
34	BA	1643	U	O3'-P	-6.54	1.53	1.61
35	BB	816	U	O4'-C1'	-6.54	1.33	1.41
35	BB	1252	G	C8-N7	-6.54	1.27	1.30
35	BB	1349	U	C2-N3	-6.54	1.33	1.37
35	BB	1387	C	N3-C4	-6.54	1.29	1.33
36	BC	75	G	O3'-P	-6.54	1.53	1.61
85	AA	513	G	C5-C4	-6.54	1.33	1.38
85	AA	631	G	C4'-O4'	-6.54	1.37	1.45
85	AA	721	C	O3'-P	-6.54	1.53	1.61
85	AA	920	A	C5'-C4'	6.54	1.59	1.51
34	BA	182	U	C3'-C2'	-6.54	1.45	1.52
34	BA	207	A	O3'-P	-6.54	1.53	1.61
34	BA	403	A	C2'-C1'	-6.54	1.46	1.53
34	BA	496	G	O3'-P	-6.54	1.53	1.61
34	BA	1103	G	C2'-C1'	-6.54	1.46	1.53
34	BA	1585	A	P-O5'	-6.54	1.53	1.59
35	BB	559	U	O3'-P	-6.54	1.53	1.61
35	BB	809	U	C2-N3	-6.54	1.33	1.37
35	BB	1190	U	N3-C4	-6.54	1.32	1.38
35	BB	1278	A	C5-C4	-6.54	1.34	1.38
36	BC	23	G	C4'-C3'	6.54	1.60	1.53
38	BE	7	U	P-O5'	-6.54	1.53	1.59
38	BE	96	G	C2'-C1'	-6.54	1.46	1.53
40	BG	168	A	C6-N1	-6.54	1.30	1.35
85	AA	39	A	C2'-C1'	-6.54	1.46	1.53
85	AA	640	C	C4'-C3'	-6.54	1.46	1.53
85	AA	1540	A	O3'-P	-6.54	1.53	1.61
35	BB	1309	A	C2'-C1'	-6.54	1.46	1.53
35	BB	1447	U	P-O5'	-6.54	1.53	1.59
85	AA	698	G	C2'-C1'	-6.54	1.46	1.53
85	AA	1218	C	O3'-P	-6.54	1.53	1.61
85	AA	1686	G	C2-N2	-6.54	1.28	1.34
34	BA	202	A	O3'-P	-6.54	1.53	1.61
34	BA	935	A	P-O5'	-6.54	1.53	1.59
34	BA	1591	G	N1-C2	-6.54	1.32	1.37
35	BB	996	G	P-O5'	-6.54	1.53	1.59
37	BD	82	G	C2-N2	-6.54	1.28	1.34
85	AA	372	U	C1'-N1	-6.54	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1933	G	C5'-C4'	6.54	1.59	1.51
35	BB	868	C	C3'-C2'	-6.54	1.45	1.52
35	BB	1047	C	C1'-N1	-6.54	1.37	1.46
85	AA	496	C	O3'-P	-6.54	1.53	1.61
34	BA	348	U	C2-N3	-6.54	1.33	1.37
34	BA	585	G	N7-C5	-6.54	1.35	1.39
34	BA	704	G	C3'-C2'	-6.54	1.45	1.52
34	BA	1461	A	C3'-C2'	-6.54	1.45	1.52
34	BA	1510	C	C4'-C3'	-6.54	1.46	1.53
34	BA	1839	G	N7-C5	-6.54	1.35	1.39
35	BB	843	G	C2'-C1'	-6.54	1.46	1.53
35	BB	1098	G	N9-C8	-6.54	1.33	1.37
35	BB	1111	C	O3'-P	-6.54	1.53	1.61
35	BB	1172	U	C2'-C1'	-6.54	1.46	1.53
35	BB	1380	G	C6-N1	-6.54	1.34	1.39
35	BB	1439	U	P-O5'	-6.54	1.53	1.59
35	BB	1477	C	C2'-C1'	-6.54	1.46	1.53
36	BC	48	A	P-O5'	-6.54	1.53	1.59
37	BD	57	C	C2'-C1'	-6.54	1.46	1.53
37	BD	74	A	C3'-C2'	-6.54	1.45	1.52
85	AA	324	U	C2-N3	-6.54	1.33	1.37
85	AA	932	A	N7-C5	-6.54	1.35	1.39
85	AA	2110	U	C1'-N1	-6.54	1.37	1.46
34	BA	56	G	N7-C5	-6.53	1.35	1.39
34	BA	290	G	C2-N2	-6.53	1.28	1.34
34	BA	351	A	O3'-P	-6.53	1.53	1.61
34	BA	1173	C	C2'-C1'	-6.53	1.46	1.53
34	BA	1597	G	C2-N2	-6.53	1.28	1.34
34	BA	1739	G	C2-N2	-6.53	1.28	1.34
35	BB	550	G	C2-N2	-6.53	1.28	1.34
35	BB	626	C	C3'-C2'	-6.53	1.45	1.52
35	BB	696	G	C2'-C1'	-6.53	1.46	1.53
35	BB	1335	G	C1'-N9	-6.53	1.37	1.46
35	BB	1488	G	P-O5'	-6.53	1.53	1.59
37	BD	103	C	C3'-C2'	-6.53	1.45	1.52
38	BE	56	U	O4'-C1'	-6.53	1.33	1.41
85	AA	1023	U	P-O5'	-6.53	1.53	1.59
85	AA	1363	U	O3'-P	-6.53	1.53	1.61
85	AA	1488	G	C1'-N9	-6.53	1.37	1.46
34	BA	585	G	C2'-C1'	-6.53	1.46	1.53
34	BA	823	G	C2-N2	-6.53	1.28	1.34
35	BB	1411	U	O3'-P	-6.53	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	200	A	P-O5'	-6.53	1.53	1.59
40	BG	11	G	N9-C4	-6.53	1.32	1.38
40	BG	101	G	C1'-N9	-6.53	1.37	1.46
85	AA	526	G	O3'-P	-6.53	1.53	1.61
85	AA	1107	A	C5'-C4'	-6.53	1.43	1.51
85	AA	1644	G	N9-C4	6.53	1.43	1.38
34	BA	166	G	N9-C4	-6.53	1.32	1.38
34	BA	664	C	C3'-C2'	-6.53	1.45	1.52
34	BA	766	A	P-O5'	-6.53	1.53	1.59
34	BA	1260	G	C3'-C2'	-6.53	1.45	1.52
34	BA	1494	G	C5-C6	-6.53	1.35	1.42
35	BB	1372	G	C1'-N9	-6.53	1.37	1.46
36	BC	42	G	O4'-C1'	-6.53	1.33	1.41
38	BE	70	C	O3'-P	-6.53	1.53	1.61
38	BE	186	C	C2'-C1'	-6.53	1.46	1.53
85	AA	435	A	N7-C5	-6.53	1.35	1.39
85	AA	1123	C	O3'-P	-6.53	1.53	1.61
85	AA	2196	G	N3-C4	-6.53	1.30	1.35
85	AA	2196	G	O4'-C1'	-6.53	1.33	1.41
34	BA	462	C	O3'-P	-6.53	1.53	1.61
34	BA	1641	G	C2-N2	-6.53	1.28	1.34
35	BB	506	G	C6-N1	-6.53	1.34	1.39
35	BB	520	G	C6-N1	-6.53	1.34	1.39
35	BB	812	G	C2-N2	-6.53	1.28	1.34
36	BC	41	A	N7-C5	-6.53	1.35	1.39
85	AA	11	A	C2'-C1'	-6.53	1.46	1.53
85	AA	2116	U	P-O5'	-6.53	1.53	1.59
34	BA	183	G	C3'-C2'	-6.53	1.45	1.52
34	BA	215	C	C4-C5	-6.53	1.37	1.43
34	BA	239	C	C5-C6	-6.53	1.29	1.34
34	BA	354	G	C6-N1	-6.53	1.34	1.39
34	BA	748	C	O3'-P	-6.53	1.53	1.61
34	BA	1196	C	N3-C4	-6.53	1.29	1.33
35	BB	132	G	O3'-P	-6.53	1.53	1.61
35	BB	785	G	C6-N1	-6.53	1.34	1.39
35	BB	1135	U	C2-N3	-6.53	1.33	1.37
35	BB	1193	G	C6-N1	-6.53	1.34	1.39
41	BH	28	U	C2'-C1'	-6.53	1.46	1.53
85	AA	271	A	N9-C4	-6.53	1.33	1.37
85	AA	526	G	C2'-C1'	-6.53	1.46	1.53
85	AA	1700	C	C3'-C2'	-6.53	1.45	1.52
34	BA	89	G	C4'-O4'	-6.53	1.37	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1519	G	N7-C5	-6.53	1.35	1.39
34	BA	1722	U	N3-C4	-6.53	1.32	1.38
34	BA	1739	G	N7-C5	-6.53	1.35	1.39
35	BB	484	G	N7-C5	-6.53	1.35	1.39
35	BB	1018	U	O3'-P	-6.53	1.53	1.61
35	BB	1172	U	C4'-C3'	-6.53	1.46	1.53
35	BB	1316	U	C3'-C2'	-6.53	1.45	1.52
35	BB	1545	U	C2-N3	-6.53	1.33	1.37
85	AA	316	C	C3'-C2'	-6.53	1.45	1.52
85	AA	671	G	P-O5'	-6.53	1.53	1.59
85	AA	894	A	C5'-C4'	6.53	1.59	1.51
85	AA	1180	C	C4'-O4'	-6.53	1.37	1.45
85	AA	1624	U	P-O5'	-6.53	1.53	1.59
85	AA	1830	U	C2'-C1'	-6.53	1.46	1.53
85	AA	1887	G	C1'-N9	-6.53	1.37	1.46
85	AA	2098	A	C2'-C1'	-6.53	1.46	1.53
34	BA	1005	C	C2'-C1'	-6.52	1.46	1.53
34	BA	1485	U	C3'-C2'	-6.52	1.45	1.52
35	BB	627	G	C5-C4	-6.52	1.33	1.38
35	BB	1170	U	C3'-C2'	-6.52	1.45	1.52
40	BG	39	A	C2'-C1'	-6.52	1.46	1.53
85	AA	90	A	N3-C4	-6.52	1.30	1.34
85	AA	495	G	P-O5'	-6.52	1.53	1.59
85	AA	889	G	N9-C4	-6.52	1.32	1.38
85	AA	1252	A	C1'-N9	-6.52	1.37	1.46
34	BA	908	G	C5-C4	-6.52	1.33	1.38
34	BA	1021	U	C2-N3	-6.52	1.33	1.37
34	BA	1073	G	C8-N7	-6.52	1.27	1.30
34	BA	1091	U	O4'-C1'	-6.52	1.33	1.41
34	BA	1101	A	N9-C4	-6.52	1.33	1.37
34	BA	1258	G	O3'-P	-6.52	1.53	1.61
34	BA	1709	A	O4'-C1'	-6.52	1.33	1.41
35	BB	36	U	N1-C2	-6.52	1.32	1.38
35	BB	550	G	N3-C4	-6.52	1.30	1.35
35	BB	1143	A	N9-C4	-6.52	1.33	1.37
38	BE	120	C	C4-N4	-6.52	1.28	1.33
34	BA	31	A	N3-C4	-6.52	1.30	1.34
34	BA	196	A	C2'-C1'	-6.52	1.46	1.53
35	BB	1068	G	N7-C5	-6.52	1.35	1.39
37	BD	2	G	C1'-N9	-6.52	1.37	1.46
38	BE	140	G	C4'-C3'	-6.52	1.46	1.53
85	AA	203	C	P-O5'	-6.52	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	695	A	P-O5'	-6.52	1.53	1.59
34	BA	208	A	P-O5'	-6.52	1.53	1.59
34	BA	582	U	P-O5'	-6.52	1.53	1.59
34	BA	969	A	P-O5'	-6.52	1.53	1.59
35	BB	13	A	C4'-C3'	-6.52	1.46	1.53
35	BB	429	C	C3'-C2'	-6.52	1.45	1.52
35	BB	1115	G	C3'-C2'	-6.52	1.45	1.52
35	BB	1176	G	N7-C5	-6.52	1.35	1.39
35	BB	1204	C	P-O5'	-6.52	1.53	1.59
38	BE	69	C	N1-C6	-6.52	1.33	1.37
40	BG	23	C	C3'-O3'	-6.52	1.33	1.42
85	AA	431	G	C3'-C2'	-6.52	1.45	1.52
85	AA	1495	G	C2'-C1'	-6.52	1.46	1.53
34	BA	726	G	N9-C4	-6.52	1.32	1.38
34	BA	733	G	N3-C4	-6.52	1.30	1.35
34	BA	1077	G	C5-C4	-6.52	1.33	1.38
34	BA	1148	U	O4'-C1'	-6.52	1.33	1.41
34	BA	1414	C	C3'-C2'	-6.52	1.45	1.52
34	BA	1721	U	C3'-C2'	6.52	1.60	1.52
35	BB	66	G	C1'-N9	-6.52	1.37	1.46
35	BB	117	A	C2'-C1'	-6.52	1.46	1.53
35	BB	1106	G	N7-C5	-6.52	1.35	1.39
35	BB	1307	C	C4'-C3'	-6.52	1.46	1.53
38	BE	171	U	N1-C2	-6.52	1.32	1.38
40	BG	63	U	P-O5'	-6.52	1.53	1.59
40	BG	174	G	C5-C4	-6.52	1.33	1.38
85	AA	15	U	C2'-C1'	-6.52	1.46	1.53
85	AA	530	A	O3'-P	-6.52	1.53	1.61
85	AA	972	G	C2'-C1'	-6.52	1.46	1.53
34	BA	1092	U	N3-C4	-6.52	1.32	1.38
35	BB	135	C	C2-N3	-6.52	1.30	1.35
35	BB	1037	A	O3'-P	-6.52	1.53	1.61
35	BB	1071	G	N7-C5	-6.52	1.35	1.39
36	BC	147	G	C3'-C2'	-6.52	1.45	1.52
37	BD	77	A	O3'-P	-6.52	1.53	1.61
85	AA	1722	G	N9-C4	-6.52	1.32	1.38
85	AA	2054	G	C1'-N9	-6.52	1.37	1.46
85	AA	2222	G	C1'-N9	-6.52	1.37	1.46
34	BA	453	A	C1'-N9	-6.51	1.37	1.46
34	BA	581	U	C3'-C2'	6.51	1.60	1.52
34	BA	1413	G	N9-C8	-6.51	1.33	1.37
34	BA	1688	G	O4'-C1'	-6.51	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	37	G	C2-N2	-6.51	1.28	1.34
85	AA	1117	G	C1'-N9	-6.51	1.37	1.46
85	AA	2218	G	C5-C4	-6.51	1.33	1.38
34	BA	244	A	N9-C4	-6.51	1.33	1.37
34	BA	307	C	O3'-P	-6.51	1.53	1.61
34	BA	1074	C	C2'-C1'	-6.51	1.46	1.53
34	BA	1696	G	C2'-C1'	-6.51	1.46	1.53
35	BB	599	U	O3'-P	-6.51	1.53	1.61
35	BB	1274	G	O3'-P	-6.51	1.53	1.61
85	AA	207	G	C6-N1	-6.51	1.34	1.39
85	AA	669	G	C2'-C1'	-6.51	1.46	1.53
85	AA	1698	A	N7-C5	-6.51	1.35	1.39
34	BA	589	A	C2'-C1'	-6.51	1.46	1.53
34	BA	628	U	C2-N3	-6.51	1.33	1.37
34	BA	804	G	C8-N7	-6.51	1.27	1.30
34	BA	940	C	C2'-C1'	-6.51	1.46	1.53
34	BA	1411	C	C3'-C2'	-6.51	1.45	1.52
34	BA	1590	G	N7-C5	-6.51	1.35	1.39
35	BB	426	A	N7-C5	-6.51	1.35	1.39
35	BB	582	G	C6-N1	-6.51	1.34	1.39
35	BB	1134	G	N7-C5	-6.51	1.35	1.39
35	BB	1498	G	C2-N3	-6.51	1.27	1.32
39	BF	13	U	C2'-C1'	-6.51	1.46	1.53
39	BF	29	U	P-O5'	-6.51	1.53	1.59
40	BG	59	G	N7-C5	-6.51	1.35	1.39
67	Bh	161	GLY	CA-C	-6.51	1.41	1.51
85	AA	2135	A	C3'-C2'	-6.51	1.45	1.52
34	BA	100	A	N3-C4	-6.51	1.30	1.34
34	BA	522	C	O3'-P	-6.51	1.53	1.61
34	BA	850	C	C2'-C1'	-6.51	1.46	1.53
35	BB	1016	C	O3'-P	-6.51	1.53	1.61
35	BB	1220	A	N7-C5	-6.51	1.35	1.39
35	BB	1302	C	C3'-C2'	-6.51	1.45	1.52
85	AA	2060	G	C1'-N9	-6.51	1.37	1.46
34	BA	575	U	C2-N3	-6.51	1.33	1.37
34	BA	1557	G	C2-N2	-6.51	1.28	1.34
35	BB	375	G	N1-C2	-6.51	1.32	1.37
35	BB	681	G	C6-N1	-6.51	1.34	1.39
85	AA	33	U	C2-N3	-6.51	1.33	1.37
85	AA	884	A	C5'-C4'	6.51	1.59	1.51
85	AA	1240	A	C8-N7	-6.51	1.26	1.31
85	AA	1499	G	C5-C4	-6.51	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1671	G	C2-N2	-6.51	1.28	1.34
34	BA	5	C	P-O5'	-6.51	1.53	1.59
34	BA	102	G	C4'-C3'	-6.51	1.46	1.53
34	BA	745	A	C2'-C1'	-6.51	1.46	1.53
34	BA	999	G	C6-N1	-6.51	1.34	1.39
34	BA	1588	U	N3-C4	-6.51	1.32	1.38
35	BB	1016	C	P-O5'	-6.51	1.53	1.59
35	BB	1168	G	C2-N2	-6.51	1.28	1.34
85	AA	48	G	N7-C5	-6.51	1.35	1.39
85	AA	357	C	N1-C6	-6.51	1.33	1.37
85	AA	674	U	C2-N3	-6.51	1.33	1.37
85	AA	982	G	O3'-P	-6.51	1.53	1.61
85	AA	1140	G	C1'-N9	-6.51	1.37	1.46
85	AA	1260	G	N9-C4	-6.51	1.32	1.38
85	AA	2169	C	C2'-C1'	-6.51	1.46	1.53
34	BA	1104	C	C2'-C1'	-6.50	1.46	1.53
36	BC	113	G	N1-C2	-6.50	1.32	1.37
38	BE	22	A	P-O5'	-6.50	1.53	1.59
85	AA	982	G	C2-N2	-6.50	1.28	1.34
85	AA	1443	U	C2-N3	-6.50	1.33	1.37
34	BA	515	U	C2-N3	-6.50	1.33	1.37
34	BA	1016	A	N3-C4	-6.50	1.30	1.34
34	BA	1271	C	C1'-N1	-6.50	1.37	1.46
35	BB	1157	G	C1'-N9	-6.50	1.37	1.46
35	BB	1375	G	N1-C2	-6.50	1.32	1.37
36	BC	6	G	C2-N2	-6.50	1.28	1.34
36	BC	119	G	N1-C2	-6.50	1.32	1.37
38	BE	194	A	C2'-C1'	-6.50	1.46	1.53
85	AA	922	A	C2'-C1'	-6.50	1.46	1.53
85	AA	1652	A	C2'-C1'	-6.50	1.46	1.53
34	BA	502	U	C4'-C3'	-6.50	1.46	1.53
34	BA	991	U	O3'-P	-6.50	1.53	1.61
34	BA	1292	A	O3'-P	-6.50	1.53	1.61
34	BA	1776	G	C2'-C1'	-6.50	1.46	1.53
35	BB	22	A	C2'-C1'	-6.50	1.46	1.53
35	BB	520	G	N9-C4	-6.50	1.32	1.38
35	BB	867	C	O3'-P	-6.50	1.53	1.61
35	BB	1446	C	C4-N4	-6.50	1.28	1.33
36	BC	96	A	C2'-C1'	-6.50	1.46	1.53
36	BC	114	C	C4'-C3'	-6.50	1.46	1.53
38	BE	116	U	C3'-C2'	-6.50	1.45	1.52
40	BG	81	G	C1'-N9	-6.50	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	124	C	P-O5'	-6.50	1.53	1.59
85	AA	1452	C	P-O5'	-6.50	1.53	1.59
85	AA	1633	A	O3'-P	-6.50	1.53	1.61
85	AA	2008	G	O4'-C1'	-6.50	1.33	1.41
34	BA	260	A	C8-N7	-6.50	1.27	1.31
34	BA	801	U	C2-N3	-6.50	1.33	1.37
34	BA	809	U	C1'-N1	-6.50	1.37	1.46
35	BB	364	U	C4-C5	-6.50	1.37	1.43
35	BB	708	C	C3'-C2'	-6.50	1.45	1.52
85	AA	1282	A	O4'-C1'	-6.50	1.33	1.41
34	BA	513	U	C2'-C1'	-6.50	1.46	1.53
34	BA	1176	C	N3-C4	-6.50	1.29	1.33
34	BA	1289	C	C2'-C1'	-6.50	1.46	1.53
34	BA	1806	A	C2'-C1'	-6.50	1.46	1.53
35	BB	1411	U	P-O5'	-6.50	1.53	1.59
35	BB	1541	G	C2'-C1'	-6.50	1.46	1.53
85	AA	505	U	C2-N3	-6.50	1.33	1.37
85	AA	971	U	O3'-P	-6.50	1.53	1.61
85	AA	1764	C	P-O5'	-6.50	1.53	1.59
85	AA	2044	A	N9-C4	-6.50	1.33	1.37
85	AA	2062	U	C4'-C3'	-6.50	1.46	1.53
34	BA	24	C	C1'-N1	-6.50	1.37	1.46
34	BA	186	G	C2'-C1'	-6.50	1.46	1.53
35	BB	444	U	C2-N3	-6.50	1.33	1.37
35	BB	567	G	C5-C4	-6.50	1.33	1.38
37	BD	27	A	C2'-C1'	-6.50	1.46	1.53
85	AA	1953	G	P-O5'	-6.50	1.53	1.59
85	AA	2152	C	P-O5'	-6.50	1.53	1.59
34	BA	1474	G	C5-C6	-6.50	1.35	1.42
85	AA	924	A	N9-C4	-6.50	1.33	1.37
85	AA	937	G	O3'-P	-6.50	1.53	1.61
34	BA	915	A	N7-C5	-6.49	1.35	1.39
34	BA	1724	G	C4'-C3'	6.49	1.60	1.53
34	BA	1801	G	O4'-C1'	-6.49	1.33	1.41
35	BB	614	U	C2-N3	-6.49	1.33	1.37
35	BB	675	U	C2-N3	-6.49	1.33	1.37
35	BB	1362	G	C2'-C1'	-6.49	1.46	1.53
36	BC	98	C	C2-N3	-6.49	1.30	1.35
38	BE	60	C	C2'-C1'	-6.49	1.46	1.53
38	BE	167	U	P-O5'	-6.49	1.53	1.59
85	AA	515	C	O4'-C1'	-6.49	1.33	1.41
85	AA	613	G	O3'-P	-6.49	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	852	C	O3'-P	-6.49	1.53	1.61
85	AA	903	G	C3'-C2'	-6.49	1.45	1.52
85	AA	2187	G	C6-N1	-6.49	1.35	1.39
34	BA	561	U	O3'-P	-6.49	1.53	1.61
34	BA	786	U	N1-C2	-6.49	1.32	1.38
34	BA	1556	A	N3-C4	-6.49	1.30	1.34
34	BA	1802	C	C2-N3	-6.49	1.30	1.35
35	BB	1267	C	C2'-C1'	-6.49	1.46	1.53
36	BC	140	U	C2'-C1'	-6.49	1.46	1.53
40	BG	33	G	N3-C4	-6.49	1.30	1.35
85	AA	1231	G	P-O5'	-6.49	1.53	1.59
34	BA	674	G	C4'-C3'	-6.49	1.46	1.53
34	BA	709	C	C2'-C1'	-6.49	1.46	1.53
34	BA	1475	G	C3'-O3'	-6.49	1.33	1.42
38	BE	187	G	C2'-C1'	-6.49	1.46	1.53
40	BG	61	A	C3'-C2'	-6.49	1.45	1.52
85	AA	77	C	C2'-C1'	-6.49	1.46	1.53
85	AA	110	U	O3'-P	-6.49	1.53	1.61
85	AA	305	A	C5'-C4'	6.49	1.59	1.51
85	AA	2109	G	C2-N2	-6.49	1.28	1.34
34	BA	224	G	C2'-C1'	-6.49	1.46	1.53
34	BA	1225	A	C6-N6	-6.49	1.28	1.33
34	BA	1635	A	N9-C4	6.49	1.41	1.37
35	BB	962	U	C2-N3	-6.49	1.33	1.37
36	BC	101	U	C1'-N1	-6.49	1.37	1.46
36	BC	162	C	C2-N3	-6.49	1.30	1.35
38	BE	106	C	C2'-C1'	-6.49	1.46	1.53
85	AA	106	G	N1-C2	-6.49	1.32	1.37
85	AA	1149	A	N9-C4	-6.49	1.33	1.37
34	BA	52	G	C4'-C3'	-6.49	1.46	1.53
41	BH	133	U	O4'-C1'	-6.49	1.33	1.41
85	AA	411	U	C1'-N1	-6.49	1.37	1.46
34	BA	45	A	O4'-C1'	-6.49	1.33	1.41
34	BA	937	G	C6-N1	-6.49	1.35	1.39
39	BF	38	C	C2'-C1'	-6.49	1.46	1.53
85	AA	404	A	N9-C4	-6.49	1.33	1.37
85	AA	757	A	N9-C4	-6.49	1.33	1.37
85	AA	1558	U	C3'-C2'	-6.49	1.45	1.52
34	BA	1080	U	C5'-C4'	6.48	1.59	1.51
34	BA	1476	G	O3'-P	-6.48	1.53	1.61
35	BB	728	A	O3'-P	-6.48	1.53	1.61
85	AA	2178	A	C3'-C2'	-6.48	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	427	G	N7-C5	-6.48	1.35	1.39
34	BA	484	A	O4'-C1'	-6.48	1.33	1.41
34	BA	686	U	O3'-P	-6.48	1.53	1.61
34	BA	1140	A	P-O5'	-6.48	1.53	1.59
34	BA	1800	G	C5-C4	-6.48	1.33	1.38
37	BD	66	G	C2'-C1'	-6.48	1.46	1.53
39	BF	24	G	N7-C5	-6.48	1.35	1.39
40	BG	161	C	C2-N3	-6.48	1.30	1.35
85	AA	55	A	C3'-C2'	-6.48	1.45	1.52
85	AA	469	G	C2'-C1'	-6.48	1.46	1.53
85	AA	929	G	C4'-C3'	-6.48	1.46	1.53
85	AA	1790	G	O3'-P	-6.48	1.53	1.61
85	AA	2083	G	C6-N1	-6.48	1.35	1.39
85	AA	2092	A	C5-C4	-6.48	1.34	1.38
85	AA	2131	C	O3'-P	-6.48	1.53	1.61
34	BA	362	G	N1-C2	-6.48	1.32	1.37
34	BA	660	C	O3'-P	-6.48	1.53	1.61
34	BA	1098	G	N9-C8	-6.48	1.33	1.37
34	BA	1550	G	N9-C4	-6.48	1.32	1.38
35	BB	1065	G	C2'-C1'	-6.48	1.46	1.53
35	BB	1314	G	C6-N1	-6.48	1.35	1.39
35	BB	1480	G	N9-C4	-6.48	1.32	1.38
36	BC	87	C	C2'-C1'	-6.48	1.46	1.53
38	BE	49	A	C6-N1	-6.48	1.31	1.35
85	AA	258	G	C1'-N9	-6.48	1.37	1.46
34	BA	1288	U	C4'-C3'	-6.48	1.46	1.53
35	BB	717	A	N9-C4	6.48	1.41	1.37
35	BB	1124	G	C6-N1	-6.48	1.35	1.39
85	AA	1001	G	C2'-C1'	-6.48	1.46	1.53
85	AA	1712	A	C1'-N9	-6.48	1.37	1.46
34	BA	1512	C	C1'-N1	-6.48	1.37	1.46
34	BA	1517	U	C4'-C3'	-6.48	1.46	1.53
35	BB	28	G	C6-N1	-6.48	1.35	1.39
35	BB	130	G	C5-C4	-6.48	1.33	1.38
35	BB	394	A	C1'-N9	-6.48	1.37	1.46
35	BB	682	U	C3'-C2'	-6.48	1.45	1.52
38	BE	202	C	P-O5'	-6.48	1.53	1.59
85	AA	154	U	C2-N3	-6.48	1.33	1.37
85	AA	616	A	C4'-C3'	-6.48	1.46	1.53
85	AA	1487	G	C2-N3	-6.48	1.27	1.32
34	BA	1739	G	N3-C4	-6.48	1.30	1.35
35	BB	624	A	C1'-N9	-6.48	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	151	C	O3'-P	-6.48	1.53	1.61
85	AA	766	G	C3'-C2'	-6.48	1.45	1.52
34	BA	112	C	C2'-C1'	-6.47	1.46	1.53
34	BA	192	G	C3'-C2'	-6.47	1.45	1.52
34	BA	344	G	P-O5'	-6.47	1.53	1.59
34	BA	1345	U	P-O5'	-6.47	1.53	1.59
34	BA	1634	A	O3'-P	-6.47	1.53	1.61
34	BA	1839	G	C6-N1	-6.47	1.35	1.39
35	BB	91	G	N3-C4	-6.47	1.30	1.35
35	BB	555	G	O3'-P	-6.47	1.53	1.61
35	BB	1254	G	N9-C8	-6.47	1.33	1.37
37	BD	92	G	C5-C4	-6.47	1.33	1.38
38	BE	142	A	O3'-P	-6.47	1.53	1.61
85	AA	634	U	O3'-P	-6.47	1.53	1.61
85	AA	1463	A	C5-C4	-6.47	1.34	1.38
85	AA	2082	C	C2'-C1'	-6.47	1.46	1.53
34	BA	269	G	C6-N1	-6.47	1.35	1.39
34	BA	329	G	C1'-N9	-6.47	1.37	1.46
34	BA	783	U	C4'-C3'	-6.47	1.46	1.53
34	BA	1246	G	C2'-C1'	-6.47	1.46	1.53
35	BB	451	A	N3-C4	-6.47	1.30	1.34
36	BC	96	A	C3'-C2'	-6.47	1.45	1.52
39	BF	8	C	O3'-P	-6.47	1.53	1.61
40	BG	84	U	O3'-P	-6.47	1.53	1.61
40	BG	146	C	P-O5'	-6.47	1.53	1.59
85	AA	430	G	O4'-C1'	-6.47	1.33	1.41
85	AA	552	C	O3'-P	-6.47	1.53	1.61
85	AA	1133	C	C2'-C1'	-6.47	1.46	1.53
85	AA	2228	G	C2-N2	-6.47	1.28	1.34
34	BA	668	G	C2'-C1'	-6.47	1.46	1.53
34	BA	748	C	C1'-N1	-6.47	1.37	1.46
34	BA	1039	G	N9-C4	-6.47	1.32	1.38
34	BA	1519	G	N1-C2	-6.47	1.32	1.37
34	BA	1555	G	C2-N3	-6.47	1.27	1.32
38	BE	199	A	C5-C4	-6.47	1.34	1.38
85	AA	824	C	O3'-P	-6.47	1.53	1.61
85	AA	1874	G	O3'-P	-6.47	1.53	1.61
34	BA	379	C	P-O5'	-6.47	1.53	1.59
34	BA	829	U	C2'-C1'	-6.47	1.46	1.53
35	BB	373	C	C2'-C1'	-6.47	1.46	1.53
35	BB	623	A	N9-C4	-6.47	1.33	1.37
35	BB	957	A	O3'-P	-6.47	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1191	G	N9-C4	-6.47	1.32	1.38
35	BB	1544	A	C5-C4	-6.47	1.34	1.38
36	BC	59	A	N9-C8	-6.47	1.32	1.37
36	BC	86	U	P-O5'	-6.47	1.53	1.59
38	BE	149	A	O3'-P	-6.47	1.53	1.61
41	BH	18	C	O3'-P	-6.47	1.53	1.61
85	AA	491	G	P-O5'	-6.47	1.53	1.59
35	BB	1435	G	C1'-N9	-6.47	1.37	1.46
40	BG	95	U	C4'-C3'	-6.47	1.46	1.53
40	BG	95	U	N3-C4	-6.47	1.32	1.38
34	BA	30	A	N9-C8	-6.47	1.32	1.37
34	BA	952	G	O3'-P	-6.47	1.53	1.61
34	BA	1052	G	C1'-N9	-6.47	1.37	1.46
34	BA	1104	C	P-O5'	-6.47	1.53	1.59
34	BA	1154	U	C4'-C3'	-6.47	1.46	1.53
35	BB	2	C	O3'-P	-6.47	1.53	1.61
35	BB	1298	C	P-O5'	-6.47	1.53	1.59
35	BB	1458	U	C2'-C1'	-6.47	1.46	1.53
38	BE	120	C	C2'-C1'	-6.47	1.46	1.53
40	BG	122	G	N9-C4	-6.47	1.32	1.38
85	AA	1190	G	C3'-C2'	-6.47	1.45	1.52
85	AA	2127	G	C3'-C2'	-6.47	1.45	1.52
34	BA	1023	G	N9-C4	-6.46	1.32	1.38
34	BA	1051	A	C2'-C1'	-6.46	1.46	1.53
34	BA	1236	U	O3'-P	-6.46	1.53	1.61
34	BA	1600	G	C5-C4	-6.46	1.33	1.38
34	BA	1604	A	C8-N7	-6.46	1.27	1.31
34	BA	1661	U	O4'-C1'	-6.46	1.33	1.41
35	BB	412	A	C2'-C1'	-6.46	1.46	1.53
35	BB	722	U	P-O5'	-6.46	1.53	1.59
35	BB	1404	A	C5-C4	-6.46	1.34	1.38
35	BB	1432	U	N3-C4	-6.46	1.32	1.38
85	AA	1576	G	P-O5'	-6.46	1.53	1.59
34	BA	47	U	O3'-P	-6.46	1.53	1.61
34	BA	1090	A	N9-C4	-6.46	1.33	1.37
35	BB	583	G	C6-N1	-6.46	1.35	1.39
35	BB	693	U	C2'-C1'	-6.46	1.46	1.53
38	BE	180	G	N9-C8	-6.46	1.33	1.37
85	AA	444	U	P-O5'	-6.46	1.53	1.59
85	AA	1898	C	O3'-P	-6.46	1.53	1.61
34	BA	57	A	C5-C6	-6.46	1.35	1.41
34	BA	1002	U	C2-N3	-6.46	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1566	G	C5-C6	-6.46	1.35	1.42
35	BB	429	C	O3'-P	-6.46	1.53	1.61
35	BB	1513	U	P-O5'	-6.46	1.53	1.59
36	BC	136	G	C3'-C2'	-6.46	1.45	1.52
38	BE	17	U	C2'-C1'	-6.46	1.46	1.53
38	BE	144	A	C1'-N9	-6.46	1.37	1.46
39	BF	71	G	C1'-N9	-6.46	1.37	1.46
40	BG	16	G	C4'-C3'	-6.46	1.46	1.53
85	AA	463	G	C5-C4	-6.46	1.33	1.38
37	BD	79	G	C5'-C4'	6.46	1.59	1.51
85	AA	1535	C	P-O5'	-6.46	1.53	1.59
34	BA	300	C	N1-C6	-6.46	1.33	1.37
34	BA	438	A	N7-C5	-6.46	1.35	1.39
35	BB	514	G	C6-N1	-6.46	1.35	1.39
35	BB	620	G	C4'-C3'	-6.46	1.46	1.53
35	BB	680	A	C5-C4	-6.46	1.34	1.38
35	BB	1057	G	C2'-C1'	-6.46	1.46	1.53
35	BB	1130	U	C3'-C2'	-6.46	1.45	1.52
36	BC	43	A	C1'-N9	-6.46	1.37	1.46
41	BH	44	A	O3'-P	-6.46	1.53	1.61
85	AA	420	C	N1-C6	-6.46	1.33	1.37
85	AA	805	A	N9-C4	6.46	1.41	1.37
85	AA	1868	G	C6-N1	-6.46	1.35	1.39
85	AA	2211	G	C3'-C2'	-6.46	1.45	1.52
86	AB	7	A	C2'-C1'	-6.46	1.46	1.53
34	BA	110	C	P-O5'	-6.46	1.53	1.59
34	BA	448	U	N3-C4	-6.46	1.32	1.38
34	BA	891	C	C1'-N1	-6.46	1.37	1.46
34	BA	1224	A	N7-C5	-6.46	1.35	1.39
34	BA	1572	G	C2'-C1'	-6.46	1.46	1.53
35	BB	363	A	C4'-C3'	-6.46	1.46	1.53
35	BB	1140	C	O3'-P	-6.46	1.53	1.61
35	BB	1207	C	C4'-O4'	-6.46	1.37	1.45
35	BB	1273	G	N9-C4	-6.46	1.32	1.38
37	BD	88	U	P-O5'	-6.46	1.53	1.59
38	BE	63	C	P-O5'	-6.46	1.53	1.59
38	BE	114	G	O3'-P	-6.46	1.53	1.61
85	AA	1106	A	O3'-P	-6.46	1.53	1.61
34	BA	378	C	C3'-C2'	-6.46	1.45	1.52
34	BA	722	A	P-O5'	-6.46	1.53	1.59
34	BA	749	G	C5-C4	-6.46	1.33	1.38
34	BA	1016	A	C5-C4	-6.46	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1428	G	O3'-P	-6.46	1.53	1.61
34	BA	1461	A	C5-C4	-6.46	1.34	1.38
35	BB	868	C	C2-N3	-6.46	1.30	1.35
85	AA	51	A	C2'-C1'	-6.46	1.46	1.53
85	AA	441	C	C2-N3	-6.46	1.30	1.35
85	AA	1488	G	C2'-C1'	-6.46	1.46	1.53
34	BA	499	C	C3'-C2'	-6.45	1.45	1.52
34	BA	617	G	C4'-C3'	-6.45	1.46	1.53
34	BA	743	A	O3'-P	-6.45	1.53	1.61
34	BA	1486	U	N1-C6	-6.45	1.32	1.38
34	BA	1653	G	C5-C4	-6.45	1.33	1.38
35	BB	125	G	N1-C2	-6.45	1.32	1.37
35	BB	1109	A	C1'-N9	-6.45	1.37	1.46
35	BB	1299	G	C6-N1	-6.45	1.35	1.39
35	BB	1478	G	C4'-C3'	-6.45	1.46	1.53
37	BD	2	G	C2'-C1'	-6.45	1.46	1.53
37	BD	3	G	N1-C2	-6.45	1.32	1.37
41	BH	17	A	C5'-C4'	-6.45	1.43	1.51
85	AA	487	G	C6-N1	-6.45	1.35	1.39
85	AA	1112	G	C2-N2	-6.45	1.28	1.34
34	BA	508	C	C2'-C1'	-6.45	1.46	1.53
34	BA	1572	G	P-O5'	-6.45	1.53	1.59
34	BA	1638	U	O3'-P	-6.45	1.53	1.61
35	BB	121	A	C6-N6	-6.45	1.28	1.33
35	BB	1032	U	C2'-C1'	-6.45	1.46	1.53
36	BC	154	A	C1'-N9	-6.45	1.37	1.46
34	BA	234	A	C8-N7	-6.45	1.27	1.31
34	BA	369	A	N9-C4	-6.45	1.33	1.37
34	BA	1322	A	N7-C5	-6.45	1.35	1.39
34	BA	1428	G	C1'-N9	-6.45	1.37	1.46
34	BA	1804	A	C8-N7	-6.45	1.27	1.31
35	BB	124	G	C2-N2	-6.45	1.28	1.34
35	BB	137	A	C5-C4	-6.45	1.34	1.38
35	BB	494	C	C1'-N1	-6.45	1.37	1.46
35	BB	681	G	P-O5'	-6.45	1.53	1.59
35	BB	1187	G	N3-C4	-6.45	1.30	1.35
37	BD	98	G	C5-C4	-6.45	1.33	1.38
40	BG	165	C	C2'-C1'	-6.45	1.46	1.53
85	AA	2138	G	C5-C4	-6.45	1.33	1.38
34	BA	51	C	C4-N4	-6.45	1.28	1.33
34	BA	102	G	O3'-P	-6.45	1.53	1.61
34	BA	1465	C	C1'-N1	-6.45	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1570	C	C2'-C1'	-6.45	1.46	1.53
35	BB	307	A	N9-C4	-6.45	1.33	1.37
35	BB	430	A	C3'-C2'	-6.45	1.45	1.52
35	BB	875	G	C2'-C1'	-6.45	1.46	1.53
35	BB	1230	A	N9-C4	-6.45	1.33	1.37
35	BB	1405	G	C2'-C1'	-6.45	1.46	1.53
37	BD	61	C	C3'-C2'	-6.45	1.45	1.52
38	BE	178	G	C5-C4	-6.45	1.33	1.38
40	BG	52	A	N7-C5	-6.45	1.35	1.39
85	AA	56	U	O3'-P	-6.45	1.53	1.61
85	AA	392	G	C3'-C2'	-6.45	1.45	1.52
85	AA	409	C	C2-N3	-6.45	1.30	1.35
85	AA	452	A	C5-C4	-6.45	1.34	1.38
85	AA	1299	A	C1'-N9	-6.45	1.37	1.46
85	AA	1645	G	C2'-C1'	-6.45	1.46	1.53
85	AA	2105	G	C2'-C1'	-6.45	1.46	1.53
34	BA	334	G	N1-C2	-6.45	1.32	1.37
40	BG	66	C	C2-N3	-6.45	1.30	1.35
85	AA	410	A	N3-C4	-6.45	1.30	1.34
85	AA	2008	G	C2'-C1'	-6.45	1.46	1.53
34	BA	544	U	C4-C5	-6.45	1.37	1.43
34	BA	1207	A	O3'-P	-6.45	1.53	1.61
35	BB	578	G	C3'-C2'	-6.45	1.45	1.52
35	BB	679	G	C5-C4	-6.45	1.33	1.38
35	BB	1212	C	C3'-C2'	-6.45	1.45	1.52
36	BC	116	C	C1'-N1	-6.45	1.37	1.46
38	BE	69	C	C2'-C1'	-6.45	1.46	1.53
85	AA	725	G	C2'-C1'	-6.45	1.46	1.53
85	AA	1175	A	C2'-C1'	-6.45	1.46	1.53
34	BA	81	C	C1'-N1	-6.44	1.37	1.46
34	BA	257	G	C5'-C4'	-6.44	1.43	1.51
34	BA	1279	U	C1'-N1	-6.44	1.37	1.46
35	BB	830	G	C5-C4	-6.44	1.33	1.38
39	BF	66	C	O3'-P	-6.44	1.53	1.61
85	AA	171	U	O3'-P	-6.44	1.53	1.61
34	BA	42	A	C3'-C2'	-6.44	1.45	1.52
34	BA	525	A	C5-C4	-6.44	1.34	1.38
34	BA	696	A	C4'-C3'	-6.44	1.46	1.53
34	BA	1237	U	O3'-P	-6.44	1.53	1.61
34	BA	1275	G	C1'-N9	-6.44	1.37	1.46
35	BB	77	A	C3'-C2'	-6.44	1.45	1.52
35	BB	360	C	O3'-P	-6.44	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	368	C	C3'-C2'	-6.44	1.45	1.52
35	BB	441	G	N7-C5	-6.44	1.35	1.39
35	BB	596	C	O3'-P	-6.44	1.53	1.61
35	BB	804	U	C2-N3	-6.44	1.33	1.37
35	BB	1268	C	C3'-C2'	-6.44	1.45	1.52
35	BB	1383	C	P-O5'	-6.44	1.53	1.59
37	BD	48	G	C2'-C1'	-6.44	1.46	1.53
38	BE	124	G	N9-C8	-6.44	1.33	1.37
41	BH	115	A	N9-C4	-6.44	1.33	1.37
85	AA	128	U	C4'-C3'	-6.44	1.46	1.53
85	AA	396	U	C4'-C3'	-6.44	1.46	1.53
85	AA	404	A	C4'-C3'	-6.44	1.46	1.53
85	AA	866	U	P-O5'	-6.44	1.53	1.59
85	AA	1004	G	N7-C5	-6.44	1.35	1.39
85	AA	1108	U	P-O5'	-6.44	1.53	1.59
85	AA	2237	G	C6-N1	-6.44	1.35	1.39
34	BA	357	A	C1'-N9	-6.44	1.37	1.46
34	BA	471	U	C4'-O4'	-6.44	1.37	1.45
34	BA	718	U	C2-N3	-6.44	1.33	1.37
34	BA	1418	G	C1'-N9	-6.44	1.37	1.46
35	BB	612	A	P-O5'	-6.44	1.53	1.59
35	BB	1118	G	C1'-N9	-6.44	1.37	1.46
37	BD	70	C	O3'-P	-6.44	1.53	1.61
40	BG	87	G	C2'-C1'	-6.44	1.46	1.53
40	BG	89	A	C4'-C3'	-6.44	1.46	1.53
85	AA	1144	G	C2-N2	-6.44	1.28	1.34
85	AA	2056	C	O3'-P	-6.44	1.53	1.61
85	AA	2123	U	O3'-P	-6.44	1.53	1.61
34	BA	38	G	N3-C4	-6.44	1.30	1.35
34	BA	495	A	C1'-N9	-6.44	1.37	1.46
34	BA	881	C	N1-C6	-6.44	1.33	1.37
35	BB	602	G	N9-C4	-6.44	1.32	1.38
35	BB	1416	A	C5-C4	-6.44	1.34	1.38
36	BC	83	A	O3'-P	-6.44	1.53	1.61
85	AA	846	U	P-O5'	-6.44	1.53	1.59
85	AA	1240	A	N7-C5	-6.44	1.35	1.39
85	AA	2113	U	C3'-C2'	-6.44	1.45	1.52
34	BA	196	A	N9-C8	-6.44	1.32	1.37
34	BA	280	A	C5-C6	-6.44	1.35	1.41
34	BA	1601	C	C4'-C3'	-6.44	1.46	1.53
35	BB	71	A	C2'-C1'	-6.44	1.46	1.53
35	BB	998	G	C1'-N9	-6.44	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1100	C	C4-N4	-6.44	1.28	1.33
36	BC	21	U	C1'-N1	-6.44	1.37	1.46
40	BG	114	A	O3'-P	-6.44	1.53	1.61
85	AA	722	G	N9-C4	-6.44	1.32	1.38
85	AA	2012	G	O3'-P	-6.44	1.53	1.61
34	BA	308	C	C2'-C1'	-6.44	1.46	1.53
34	BA	1191	C	C2-N3	-6.44	1.30	1.35
34	BA	1463	U	C3'-C2'	-6.44	1.45	1.52
34	BA	1844	U	O3'-P	-6.44	1.53	1.61
35	BB	408	U	O3'-P	-6.44	1.53	1.61
38	BE	197	A	N9-C8	-6.44	1.32	1.37
39	BF	27	G	C1'-N9	-6.44	1.37	1.46
40	BG	181	C	C4'-C3'	-6.44	1.46	1.53
85	AA	23	G	C3'-C2'	-6.44	1.45	1.52
85	AA	24	U	N1-C6	-6.44	1.32	1.38
85	AA	662	U	N1-C2	-6.44	1.32	1.38
34	BA	729	C	O3'-P	-6.43	1.53	1.61
36	BC	78	G	C6-N1	-6.43	1.35	1.39
37	BD	44	U	C2-N3	-6.43	1.33	1.37
41	BH	115	A	C1'-N9	-6.43	1.37	1.46
85	AA	711	C	C2-N3	-6.43	1.30	1.35
85	AA	781	G	C2'-C1'	-6.43	1.46	1.53
85	AA	1719	C	O3'-P	-6.43	1.53	1.61
34	BA	138	C	C4'-C3'	-6.43	1.46	1.53
34	BA	423	G	P-O5'	-6.43	1.53	1.59
34	BA	682	A	N7-C5	-6.43	1.35	1.39
34	BA	1285	G	C2-N2	-6.43	1.28	1.34
35	BB	369	A	O3'-P	-6.43	1.53	1.61
36	BC	71	A	O3'-P	-6.43	1.53	1.61
85	AA	331	G	C6-N1	-6.43	1.35	1.39
41	BH	113	G	C5-C6	-6.43	1.35	1.42
34	BA	7	U	N1-C2	6.43	1.44	1.38
34	BA	536	C	P-O5'	-6.43	1.53	1.59
34	BA	1436	A	N9-C4	-6.43	1.33	1.37
34	BA	1579	G	N1-C2	-6.43	1.32	1.37
35	BB	1248	A	O3'-P	-6.43	1.53	1.61
85	AA	110	U	C1'-N1	-6.43	1.37	1.46
85	AA	145	C	C4'-C3'	-6.43	1.46	1.53
85	AA	247	G	P-O5'	-6.43	1.53	1.59
85	AA	503	A	P-O5'	-6.43	1.53	1.59
85	AA	613	G	C4'-C3'	-6.43	1.46	1.53
85	AA	818	C	C2-N3	-6.43	1.30	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1111	A	C5-C4	-6.43	1.34	1.38
34	BA	797	A	N7-C5	-6.43	1.35	1.39
35	BB	579	A	C5-C6	-6.43	1.35	1.41
85	AA	307	G	O3'-P	6.43	1.68	1.61
34	BA	102	G	C2-N2	-6.43	1.28	1.34
34	BA	423	G	C1'-N9	-6.43	1.37	1.46
34	BA	1064	A	C3'-C2'	-6.43	1.45	1.52
34	BA	1294	C	P-O5'	-6.43	1.53	1.59
34	BA	1634	A	C5'-C4'	6.43	1.59	1.51
34	BA	1664	C	C2'-C1'	-6.43	1.46	1.53
35	BB	18	A	N7-C5	-6.43	1.35	1.39
35	BB	46	U	O3'-P	-6.43	1.53	1.61
35	BB	1078	U	C3'-C2'	-6.43	1.45	1.52
38	BE	45	G	C2-N2	-6.43	1.28	1.34
40	BG	106	G	C2'-C1'	-6.43	1.46	1.53
85	AA	89	C	O3'-P	-6.43	1.53	1.61
85	AA	361	U	O3'-P	-6.43	1.53	1.61
85	AA	621	U	C3'-C2'	-6.43	1.45	1.52
85	AA	670	C	C2-N3	-6.43	1.30	1.35
85	AA	1369	U	P-O5'	-6.43	1.53	1.59
85	AA	1951	U	O3'-P	-6.43	1.53	1.61
85	AA	2131	C	C2'-C1'	-6.43	1.46	1.53
85	AA	2183	U	C4'-C3'	-6.43	1.46	1.53
34	BA	380	A	P-O5'	-6.42	1.53	1.59
34	BA	1098	G	C1'-N9	-6.42	1.37	1.46
35	BB	87	G	C2-N3	-6.42	1.27	1.32
35	BB	1141	A	C4'-C3'	-6.42	1.46	1.53
35	BB	1497	C	C3'-C2'	-6.42	1.45	1.52
41	BH	16	A	C2'-C1'	-6.42	1.46	1.53
85	AA	1184	A	O3'-P	-6.42	1.53	1.61
85	AA	1902	C	C2'-C1'	-6.42	1.46	1.53
34	BA	651	U	P-O5'	-6.42	1.53	1.59
34	BA	947	A	P-O5'	-6.42	1.53	1.59
34	BA	1032	A	C2'-C1'	-6.42	1.46	1.53
34	BA	1074	C	C2-N3	-6.42	1.30	1.35
34	BA	1653	G	C3'-C2'	-6.42	1.45	1.52
34	BA	1667	G	O3'-P	-6.42	1.53	1.61
34	BA	1825	U	N1-C2	-6.42	1.32	1.38
35	BB	365	U	C2-N3	-6.42	1.33	1.37
35	BB	380	G	N3-C4	-6.42	1.30	1.35
35	BB	640	A	C2'-C1'	-6.42	1.46	1.53
35	BB	1143	A	C2'-C1'	-6.42	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1159	U	P-O5'	-6.42	1.53	1.59
35	BB	1202	G	N3-C4	6.42	1.40	1.35
40	BG	71	C	C5-C6	-6.42	1.29	1.34
34	BA	769	U	C3'-C2'	6.42	1.60	1.52
34	BA	1473	A	C5-C4	-6.42	1.34	1.38
34	BA	1501	U	C1'-N1	-6.42	1.37	1.46
34	BA	1647	G	N9-C8	-6.42	1.33	1.37
34	BA	1813	C	C2-N3	-6.42	1.30	1.35
35	BB	620	G	C5-C4	-6.42	1.33	1.38
35	BB	1185	G	C3'-C2'	-6.42	1.45	1.52
35	BB	1345	A	P-O5'	-6.42	1.53	1.59
40	BG	13	A	O3'-P	-6.42	1.53	1.61
40	BG	150	A	C1'-N9	-6.42	1.37	1.46
41	BH	53	C	O3'-P	-6.42	1.53	1.61
85	AA	140	C	O3'-P	-6.42	1.53	1.61
85	AA	776	C	P-O5'	-6.42	1.53	1.59
85	AA	906	U	O3'-P	-6.42	1.53	1.61
85	AA	1228	A	N9-C8	-6.42	1.32	1.37
85	AA	2174	G	C5-C4	-6.42	1.33	1.38
37	BD	28	C	C2'-C1'	-6.42	1.46	1.53
38	BE	19	G	C2-N2	-6.42	1.28	1.34
85	AA	374	C	O3'-P	-6.42	1.53	1.61
85	AA	2011	C	O3'-P	-6.42	1.53	1.61
34	BA	38	G	P-O5'	-6.42	1.53	1.59
34	BA	189	G	N7-C5	-6.42	1.35	1.39
34	BA	806	U	C2'-C1'	-6.42	1.46	1.53
34	BA	1344	G	C2'-C1'	-6.42	1.46	1.53
34	BA	1628	A	P-O5'	-6.42	1.53	1.59
34	BA	1639	U	N3-C4	-6.42	1.32	1.38
34	BA	1707	C	N1-C6	-6.42	1.33	1.37
35	BB	48	G	P-O5'	-6.42	1.53	1.59
35	BB	465	C	C2-N3	-6.42	1.30	1.35
35	BB	856	U	C2'-C1'	-6.42	1.46	1.53
35	BB	1168	G	C4'-O4'	-6.42	1.37	1.45
35	BB	1371	G	C3'-C2'	-6.42	1.45	1.52
35	BB	1393	C	P-O5'	-6.42	1.53	1.59
35	BB	1394	A	C1'-N9	-6.42	1.37	1.46
37	BD	81	C	C5'-C4'	-6.42	1.43	1.51
39	BF	56	C	C2'-C1'	6.42	1.60	1.53
41	BH	31	A	C4'-O4'	-6.42	1.37	1.45
85	AA	113	U	C2'-C1'	-6.42	1.46	1.53
85	AA	160	A	N3-C4	-6.42	1.30	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	536	C	O3'-P	-6.42	1.53	1.61
85	AA	1130	G	C2-N2	-6.42	1.28	1.34
85	AA	1763	G	P-O5'	-6.42	1.53	1.59
34	BA	12	G	N7-C5	-6.42	1.35	1.39
34	BA	927	A	C5-C4	-6.42	1.34	1.38
34	BA	1062	G	C2-N2	-6.42	1.28	1.34
34	BA	1246	G	C6-N1	-6.42	1.35	1.39
35	BB	411	A	C8-N7	-6.42	1.27	1.31
35	BB	579	A	C2'-C1'	-6.42	1.46	1.53
35	BB	1505	U	P-O5'	-6.42	1.53	1.59
40	BG	148	C	N1-C2	-6.42	1.33	1.40
85	AA	117	C	O3'-P	-6.42	1.53	1.61
85	AA	395	G	C6-N1	-6.42	1.35	1.39
85	AA	757	A	C2'-C1'	-6.42	1.46	1.53
85	AA	1122	U	C2'-C1'	-6.42	1.46	1.53
85	AA	1244	A	C5-C4	-6.42	1.34	1.38
85	AA	1456	A	C8-N7	-6.42	1.27	1.31
34	BA	425	G	N7-C5	-6.42	1.35	1.39
34	BA	1437	G	N1-C2	-6.42	1.32	1.37
34	BA	43	U	C2-N3	-6.41	1.33	1.37
34	BA	386	A	O3'-P	-6.41	1.53	1.61
34	BA	1409	A	O3'-P	-6.41	1.53	1.61
34	BA	1656	A	O3'-P	-6.41	1.53	1.61
34	BA	1706	A	C3'-C2'	-6.41	1.45	1.52
35	BB	852	G	C6-N1	-6.41	1.35	1.39
38	BE	43	A	C6-N1	-6.41	1.31	1.35
34	BA	33	C	C4'-C3'	-6.41	1.46	1.53
34	BA	1819	U	C2'-C1'	-6.41	1.46	1.53
34	BA	1837	U	N3-C4	-6.41	1.32	1.38
35	BB	104	G	C1'-N9	-6.41	1.37	1.46
35	BB	310	U	P-O5'	-6.41	1.53	1.59
35	BB	435	A	C8-N7	-6.41	1.27	1.31
35	BB	505	G	N7-C5	-6.41	1.35	1.39
35	BB	587	A	C3'-C2'	-6.41	1.45	1.52
39	BF	53	G	C2'-C1'	-6.41	1.46	1.53
85	AA	648	G	C5-C4	-6.41	1.33	1.38
85	AA	1153	G	C5-C6	-6.41	1.35	1.42
85	AA	1451	U	P-O5'	-6.41	1.53	1.59
85	AA	1554	C	C2-N3	-6.41	1.30	1.35
34	BA	611	A	C4'-C3'	-6.41	1.46	1.53
34	BA	714	G	O3'-P	-6.41	1.53	1.61
34	BA	881	C	C2-N3	-6.41	1.30	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	974	G	O3'-P	-6.41	1.53	1.61
34	BA	1014	A	C6-N1	-6.41	1.31	1.35
34	BA	1051	A	N9-C4	-6.41	1.34	1.37
34	BA	1114	G	N1-C2	-6.41	1.32	1.37
35	BB	528	G	N9-C4	-6.41	1.32	1.38
35	BB	1336	G	C5-C4	-6.41	1.33	1.38
35	BB	1485	G	N1-C2	-6.41	1.32	1.37
35	BB	1507	U	C4'-C3'	-6.41	1.46	1.53
36	BC	102	G	C1'-N9	-6.41	1.37	1.46
38	BE	161	G	C2'-C1'	-6.41	1.46	1.53
40	BG	117	C	O3'-P	-6.41	1.53	1.61
85	AA	6	G	O3'-P	-6.41	1.53	1.61
85	AA	157	G	C6-N1	-6.41	1.35	1.39
85	AA	943	U	C4-C5	-6.41	1.37	1.43
34	BA	151	A	C5'-C4'	6.41	1.59	1.51
34	BA	309	U	P-O5'	-6.41	1.53	1.59
34	BA	672	G	N7-C5	-6.41	1.35	1.39
34	BA	709	C	C3'-C2'	-6.41	1.45	1.52
34	BA	1033	G	C2-N2	-6.41	1.28	1.34
34	BA	1091	U	C3'-C2'	-6.41	1.45	1.52
34	BA	1486	U	C3'-C2'	-6.41	1.45	1.52
34	BA	1784	G	C2-N2	-6.41	1.28	1.34
35	BB	697	G	C1'-N9	-6.41	1.37	1.46
35	BB	1048	A	O3'-P	-6.41	1.53	1.61
35	BB	1483	A	C8-N7	-6.41	1.27	1.31
39	BF	28	C	C3'-C2'	-6.41	1.45	1.52
85	AA	665	A	C1'-N9	-6.41	1.37	1.46
85	AA	1199	C	N1-C6	-6.41	1.33	1.37
85	AA	1262	A	N7-C5	-6.41	1.35	1.39
34	BA	457	A	O3'-P	-6.41	1.53	1.61
34	BA	515	U	C1'-N1	-6.41	1.37	1.46
34	BA	1505	G	N3-C4	-6.41	1.30	1.35
34	BA	1641	G	C2-N3	-6.41	1.27	1.32
37	BD	37	G	O3'-P	-6.41	1.53	1.61
37	BD	74	A	N7-C5	-6.41	1.35	1.39
40	BG	116	G	P-O5'	-6.41	1.53	1.59
41	BH	21	G	N1-C2	-6.41	1.32	1.37
85	AA	104	C	C1'-N1	-6.41	1.37	1.46
85	AA	1238	U	C4'-C3'	-6.41	1.46	1.53
34	BA	910	U	C1'-N1	-6.41	1.37	1.46
34	BA	1016	A	N7-C5	-6.41	1.35	1.39
34	BA	1178	U	N3-C4	-6.41	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1497	A	O4'-C1'	-6.41	1.33	1.41
34	BA	1610	A	C4'-O4'	-6.41	1.37	1.45
35	BB	59	U	O4'-C1'	-6.41	1.33	1.41
35	BB	376	A	N3-C4	-6.41	1.31	1.34
40	BG	91	U	O3'-P	-6.41	1.53	1.61
41	BH	21	G	C2-N2	-6.41	1.28	1.34
41	BH	41	A	O3'-P	-6.41	1.53	1.61
85	AA	1215	A	C2'-C1'	-6.41	1.46	1.53
85	AA	1347	C	P-O5'	-6.41	1.53	1.59
34	BA	93	A	C3'-C2'	-6.40	1.45	1.52
34	BA	957	A	N3-C4	-6.40	1.31	1.34
35	BB	1163	U	O3'-P	-6.40	1.53	1.61
38	BE	104	G	C5'-C4'	6.40	1.59	1.51
85	AA	1493	A	C4'-O4'	-6.40	1.37	1.45
34	BA	253	U	N3-C4	-6.40	1.32	1.38
34	BA	588	C	C4'-O4'	-6.40	1.37	1.45
34	BA	1011	G	C8-N7	-6.40	1.27	1.30
34	BA	1201	G	N7-C5	-6.40	1.35	1.39
34	BA	1322	A	C4'-C3'	-6.40	1.46	1.53
34	BA	1584	G	N7-C5	-6.40	1.35	1.39
34	BA	1660	A	O3'-P	-6.40	1.53	1.61
35	BB	1161	G	N7-C5	-6.40	1.35	1.39
35	BB	1297	G	C4'-C3'	-6.40	1.46	1.53
37	BD	78	C	N3-C4	-6.40	1.29	1.33
85	AA	460	U	C3'-C2'	-6.40	1.45	1.52
85	AA	1502	A	C4'-C3'	-6.40	1.46	1.53
85	AA	1690	A	C1'-N9	-6.40	1.37	1.46
85	AA	2021	A	N7-C5	-6.40	1.35	1.39
34	BA	214	A	O4'-C1'	-6.40	1.33	1.41
34	BA	957	A	N9-C4	-6.40	1.34	1.37
34	BA	1295	U	O4'-C1'	-6.40	1.33	1.41
34	BA	1686	G	C6-N1	-6.40	1.35	1.39
34	BA	1712	U	P-O5'	-6.40	1.53	1.59
35	BB	417	A	N7-C5	-6.40	1.35	1.39
35	BB	710	A	C2'-C1'	-6.40	1.46	1.53
37	BD	9	C	C1'-N1	-6.40	1.37	1.46
39	BF	46	G	C2'-C1'	-6.40	1.46	1.53
40	BG	16	G	C5-C4	-6.40	1.33	1.38
40	BG	66	C	N1-C6	-6.40	1.33	1.37
40	BG	157	A	C8-N7	-6.40	1.27	1.31
40	BG	160	C	P-O5'	-6.40	1.53	1.59
85	AA	498	C	C3'-C2'	-6.40	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	835	C	C2'-C1'	-6.40	1.46	1.53
85	AA	1174	G	C2-N2	-6.40	1.28	1.34
85	AA	1519	A	N3-C4	-6.40	1.31	1.34
34	BA	990	G	C2-N2	-6.40	1.28	1.34
35	BB	670	G	C1'-N9	-6.40	1.37	1.46
35	BB	1280	U	N3-C4	-6.40	1.32	1.38
35	BB	1297	G	C6-N1	-6.40	1.35	1.39
38	BE	13	A	C3'-C2'	-6.40	1.45	1.52
38	BE	75	C	N1-C6	-6.40	1.33	1.37
85	AA	397	G	C6-N1	-6.40	1.35	1.39
85	AA	464	A	C5-C6	-6.40	1.35	1.41
85	AA	495	G	C2'-C1'	-6.40	1.46	1.53
85	AA	2196	G	C2'-C1'	-6.40	1.46	1.53
34	BA	1496	G	N7-C5	-6.40	1.35	1.39
35	BB	5	A	C3'-C2'	-6.40	1.45	1.52
35	BB	124	G	N3-C4	-6.40	1.30	1.35
35	BB	1123	A	C2'-C1'	-6.40	1.46	1.53
41	BH	38	G	N9-C8	-6.40	1.33	1.37
85	AA	709	A	C1'-N9	-6.40	1.37	1.46
85	AA	2114	U	P-O5'	-6.40	1.53	1.59
34	BA	128	C	C4'-C3'	-6.40	1.46	1.53
34	BA	572	G	C2'-C1'	6.40	1.60	1.53
34	BA	1050	A	C3'-C2'	-6.40	1.45	1.52
34	BA	1474	G	C1'-N9	-6.40	1.37	1.46
34	BA	1523	U	N3-C4	-6.40	1.32	1.38
41	BH	58	C	O3'-P	-6.40	1.53	1.61
85	AA	66	U	O3'-P	-6.40	1.53	1.61
85	AA	880	A	C6-N1	-6.40	1.31	1.35
34	BA	982	A	N3-C4	-6.39	1.31	1.34
34	BA	1312	A	N9-C4	-6.39	1.34	1.37
34	BA	1600	G	C2-N2	-6.39	1.28	1.34
35	BB	1197	G	N9-C4	-6.39	1.32	1.38
36	BC	59	A	C1'-N9	-6.39	1.37	1.46
37	BD	87	G	C3'-C2'	-6.39	1.45	1.52
85	AA	533	C	O3'-P	-6.39	1.53	1.61
85	AA	1509	A	C5-C6	-6.39	1.35	1.41
34	BA	306	G	C1'-N9	-6.39	1.37	1.46
34	BA	1564	A	C1'-N9	-6.39	1.38	1.46
34	BA	1800	G	C2-N2	-6.39	1.28	1.34
35	BB	1047	C	P-O5'	-6.39	1.53	1.59
35	BB	1434	G	C5-C4	-6.39	1.33	1.38
40	BG	73	U	C1'-N1	-6.39	1.37	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	11	A	N7-C5	-6.39	1.35	1.39
85	AA	88	G	N1-C2	-6.39	1.32	1.37
85	AA	105	A	C2'-C1'	-6.39	1.46	1.53
85	AA	155	U	P-O5'	-6.39	1.53	1.59
85	AA	1185	G	C2'-C1'	-6.39	1.46	1.53
85	AA	1272	G	N9-C4	-6.39	1.32	1.38
85	AA	1444	U	O3'-P	-6.39	1.53	1.61
85	AA	1925	A	O4'-C1'	-6.39	1.33	1.41
34	BA	88	C	C2'-C1'	-6.39	1.46	1.53
34	BA	185	A	O4'-C1'	-6.39	1.33	1.41
34	BA	742	C	O3'-P	-6.39	1.53	1.61
34	BA	946	A	C2'-C1'	-6.39	1.46	1.53
85	AA	268	A	C3'-C2'	-6.39	1.45	1.52
85	AA	388	G	P-O5'	-6.39	1.53	1.59
85	AA	776	C	O3'-P	-6.39	1.53	1.61
34	BA	475	A	N7-C5	-6.39	1.35	1.39
34	BA	531	C	C3'-C2'	-6.39	1.45	1.52
34	BA	1403	G	C2-N2	-6.39	1.28	1.34
34	BA	1552	C	C1'-N1	-6.39	1.38	1.46
34	BA	1684	A	C2'-C1'	-6.39	1.46	1.53
34	BA	1691	G	C1'-N9	-6.39	1.38	1.46
35	BB	1156	U	O3'-P	-6.39	1.53	1.61
35	BB	1288	G	C1'-N9	-6.39	1.38	1.46
37	BD	30	A	C5-C4	-6.39	1.34	1.38
38	BE	3	G	C6-N1	-6.39	1.35	1.39
40	BG	112	C	C2-N3	-6.39	1.30	1.35
85	AA	185	A	C3'-C2'	-6.39	1.45	1.52
85	AA	423	G	C3'-C2'	-6.39	1.45	1.52
85	AA	550	G	N9-C4	-6.39	1.32	1.38
85	AA	665	A	N9-C4	-6.39	1.34	1.37
85	AA	1198	U	C4'-C3'	-6.39	1.46	1.53
85	AA	1484	G	C1'-N9	-6.39	1.38	1.46
85	AA	1496	U	O3'-P	-6.39	1.53	1.61
86	AB	26	A	P-O5'	-6.39	1.53	1.59
35	BB	1516	C	C2-N3	-6.39	1.30	1.35
37	BD	115	A	N9-C4	-6.39	1.34	1.37
85	AA	706	U	C2'-C1'	-6.39	1.46	1.53
85	AA	1519	A	C2'-C1'	-6.39	1.46	1.53
85	AA	1752	C	O3'-P	-6.39	1.53	1.61
34	BA	51	C	C4'-C3'	-6.39	1.46	1.53
34	BA	275	C	C1'-N1	-6.39	1.38	1.46
34	BA	747	G	C6-N1	-6.39	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	907	A	C2'-C1'	-6.39	1.46	1.53
34	BA	1500	G	O3'-P	-6.39	1.53	1.61
34	BA	1503	U	P-O5'	-6.39	1.53	1.59
35	BB	374	A	P-O5'	-6.39	1.53	1.59
35	BB	1371	G	C5-C4	-6.39	1.33	1.38
35	BB	1538	G	C5-C4	-6.39	1.33	1.38
38	BE	161	G	C4'-O4'	-6.39	1.37	1.45
40	BG	17	A	C4'-O4'	-6.39	1.37	1.45
40	BG	42	A	N9-C4	-6.39	1.34	1.37
86	AB	51	U	O3'-P	-6.39	1.53	1.61
34	BA	1632	G	C2-N2	-6.38	1.28	1.34
34	BA	1667	G	C4'-C3'	-6.38	1.46	1.53
35	BB	1430	G	N1-C2	-6.38	1.32	1.37
41	BH	107	A	O3'-P	-6.38	1.53	1.61
85	AA	23	G	C4'-C3'	6.38	1.60	1.53
85	AA	367	A	O3'-P	-6.38	1.53	1.61
85	AA	1002	G	N7-C5	-6.38	1.35	1.39
85	AA	1999	C	C2'-C1'	-6.38	1.46	1.53
34	BA	244	A	C5-C4	-6.38	1.34	1.38
34	BA	629	G	N7-C5	-6.38	1.35	1.39
34	BA	958	G	C6-N1	-6.38	1.35	1.39
35	BB	104	G	O3'-P	-6.38	1.53	1.61
35	BB	1253	U	C1'-N1	-6.38	1.38	1.46
35	BB	1461	C	P-O5'	-6.38	1.53	1.59
40	BG	77	U	N1-C2	-6.38	1.32	1.38
40	BG	82	U	O4'-C1'	-6.38	1.33	1.41
85	AA	125	A	N9-C4	-6.38	1.34	1.37
85	AA	662	U	C2'-C1'	-6.38	1.46	1.53
85	AA	2123	U	C5'-C4'	-6.38	1.43	1.51
34	BA	933	U	C2'-C1'	-6.38	1.46	1.53
34	BA	1286	C	P-O5'	-6.38	1.53	1.59
34	BA	1556	A	N9-C8	-6.38	1.32	1.37
35	BB	389	G	C4'-C3'	-6.38	1.46	1.53
38	BE	195	G	C1'-N9	-6.38	1.38	1.46
85	AA	544	A	P-O5'	-6.38	1.53	1.59
85	AA	1128	G	O3'-P	-6.38	1.53	1.61
85	AA	1480	C	C2'-C1'	-6.38	1.46	1.53
85	AA	1505	G	C2'-C1'	-6.38	1.46	1.53
34	BA	1048	C	P-O5'	-6.38	1.53	1.59
34	BA	1101	A	O3'-P	-6.38	1.53	1.61
35	BB	370	A	N9-C4	-6.38	1.34	1.37
35	BB	1421	C	C4-N4	-6.38	1.28	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	38	C	C3'-C2'	-6.38	1.45	1.52
85	AA	247	G	C5-C6	-6.38	1.35	1.42
85	AA	435	A	C2'-C1'	-6.38	1.46	1.53
85	AA	1227	A	O3'-P	-6.38	1.53	1.61
34	BA	67	A	N7-C5	-6.38	1.35	1.39
34	BA	989	C	P-O5'	-6.38	1.53	1.59
35	BB	62	C	C2-N3	-6.38	1.30	1.35
35	BB	1246	C	C5'-C4'	-6.38	1.43	1.51
35	BB	1483	A	N3-C4	-6.38	1.31	1.34
37	BD	36	C	C4'-C3'	-6.38	1.46	1.53
38	BE	8	G	N9-C4	6.38	1.43	1.38
41	BH	10	U	O3'-P	-6.38	1.53	1.61
85	AA	1148	G	C1'-N9	-6.38	1.38	1.46
34	BA	10	G	N9-C8	-6.38	1.33	1.37
34	BA	1505	G	N9-C4	-6.38	1.32	1.38
34	BA	1610	A	C2'-C1'	-6.38	1.46	1.53
35	BB	32	C	C2'-C1'	-6.38	1.46	1.53
35	BB	92	C	O3'-P	-6.38	1.53	1.61
35	BB	1309	A	N7-C5	-6.38	1.35	1.39
35	BB	1522	G	C2-N2	-6.38	1.28	1.34
39	BF	46	G	N9-C4	-6.38	1.32	1.38
40	BG	94	G	C3'-C2'	-6.38	1.45	1.52
85	AA	166	C	C2-N3	-6.38	1.30	1.35
85	AA	180	A	N3-C4	-6.38	1.31	1.34
85	AA	429	G	C1'-N9	-6.38	1.38	1.46
85	AA	725	G	C6-N1	-6.38	1.35	1.39
85	AA	978	U	C3'-C2'	-6.38	1.45	1.52
85	AA	1525	C	C3'-C2'	-6.38	1.45	1.52
85	AA	1905	A	P-O5'	-6.38	1.53	1.59
34	BA	593	G	C3'-C2'	6.38	1.59	1.52
34	BA	1138	C	C2'-C1'	-6.38	1.46	1.53
34	BA	1524	G	C5-C4	-6.38	1.33	1.38
35	BB	596	C	C2-N3	-6.38	1.30	1.35
35	BB	1172	U	C1'-N1	-6.38	1.38	1.46
35	BB	1343	C	C1'-N1	-6.38	1.38	1.46
39	BF	4	A	N9-C8	-6.38	1.32	1.37
62	Bc	13	ARG	CD-NE	6.38	1.57	1.46
85	AA	103	U	N1-C2	-6.38	1.32	1.38
34	BA	686	U	C2-N3	-6.37	1.33	1.37
38	BE	29	C	P-O5'	-6.37	1.53	1.59
38	BE	155	C	O3'-P	-6.37	1.53	1.61
39	BF	40	U	O3'-P	-6.37	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1191	G	C5-C4	-6.37	1.33	1.38
85	AA	1295	G	N1-C2	-6.37	1.32	1.37
34	BA	237	A	C2'-C1'	-6.37	1.46	1.53
34	BA	1410	C	C2-N3	-6.37	1.30	1.35
34	BA	1708	A	N9-C8	-6.37	1.32	1.37
35	BB	435	A	P-O5'	-6.37	1.53	1.59
35	BB	486	G	N7-C5	-6.37	1.35	1.39
35	BB	565	U	C2'-C1'	-6.37	1.46	1.53
35	BB	686	A	C3'-C2'	-6.37	1.45	1.52
35	BB	1144	A	C4'-C3'	-6.37	1.46	1.53
35	BB	1308	G	C1'-N9	-6.37	1.38	1.46
38	BE	149	A	N3-C4	-6.37	1.31	1.34
40	BG	75	C	C2-N3	-6.37	1.30	1.35
40	BG	125	C	O3'-P	-6.37	1.53	1.61
41	BH	127	A	C3'-C2'	-6.37	1.45	1.52
85	AA	480	U	O4'-C1'	-6.37	1.33	1.41
85	AA	2190	U	C2'-C1'	-6.37	1.46	1.53
86	AB	71	G	C4'-C3'	-6.37	1.46	1.53
34	BA	45	A	C1'-N9	-6.37	1.38	1.46
34	BA	349	G	N3-C4	-6.37	1.30	1.35
34	BA	364	C	C2'-C1'	-6.37	1.46	1.53
34	BA	1400	A	P-O5'	-6.37	1.53	1.59
35	BB	1233	U	N1-C2	-6.37	1.32	1.38
85	AA	176	C	C1'-N1	-6.37	1.38	1.46
85	AA	1173	A	O3'-P	-6.37	1.53	1.61
34	BA	589	A	N7-C5	-6.37	1.35	1.39
34	BA	836	U	P-O5'	-6.37	1.53	1.59
34	BA	966	G	C3'-C2'	-6.37	1.45	1.52
34	BA	1028	A	C3'-C2'	-6.37	1.45	1.52
34	BA	1545	C	O4'-C1'	-6.37	1.33	1.41
35	BB	9	G	C2'-C1'	-6.37	1.46	1.53
35	BB	94	A	C5-C4	-6.37	1.34	1.38
35	BB	114	A	O3'-P	-6.37	1.53	1.61
35	BB	1094	A	O3'-P	-6.37	1.53	1.61
37	BD	97	U	C2'-C1'	-6.37	1.46	1.53
40	BG	89	A	C3'-C2'	-6.37	1.45	1.52
40	BG	121	C	P-O5'	-6.37	1.53	1.59
85	AA	1510	A	C2'-C1'	-6.37	1.46	1.53
85	AA	2186	U	N1-C2	-6.37	1.32	1.38
85	AA	2211	G	N9-C4	-6.37	1.32	1.38
34	BA	840	U	C3'-C2'	-6.37	1.45	1.52
34	BA	1033	G	N1-C2	-6.37	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1208	U	C2'-C1'	-6.37	1.46	1.53
34	BA	1670	A	C6-N1	-6.37	1.31	1.35
34	BA	1820	G	C1'-N9	-6.37	1.38	1.46
35	BB	90	G	C3'-C2'	-6.37	1.45	1.52
34	BA	35	U	C2'-C1'	-6.37	1.46	1.53
34	BA	441	A	P-O5'	-6.37	1.53	1.59
34	BA	1289	C	C4'-O4'	-6.37	1.37	1.45
34	BA	1735	G	N9-C4	-6.37	1.32	1.38
35	BB	1155	U	O3'-P	-6.37	1.53	1.61
40	BG	4	A	N3-C4	-6.37	1.31	1.34
85	AA	86	G	O3'-P	-6.37	1.53	1.61
85	AA	1521	U	C2'-C1'	-6.37	1.46	1.53
85	AA	2084	U	C2'-C1'	-6.37	1.46	1.53
34	BA	70	C	P-O5'	-6.36	1.53	1.59
34	BA	773	A	O3'-P	-6.36	1.53	1.61
34	BA	1161	G	N1-C2	-6.36	1.32	1.37
34	BA	1428	G	N1-C2	-6.36	1.32	1.37
40	BG	169	A	C5-C4	-6.36	1.34	1.38
41	BH	75	G	N7-C5	6.36	1.43	1.39
85	AA	42	G	C2'-C1'	-6.36	1.46	1.53
85	AA	117	C	C2'-C1'	-6.36	1.46	1.53
85	AA	157	G	C4'-C3'	-6.36	1.46	1.53
85	AA	400	G	N3-C4	-6.36	1.30	1.35
85	AA	453	G	N1-C2	-6.36	1.32	1.37
85	AA	880	A	N9-C4	-6.36	1.34	1.37
85	AA	1156	A	O3'-P	-6.36	1.53	1.61
85	AA	1271	U	N1-C2	-6.36	1.32	1.38
85	AA	1463	A	C8-N7	-6.36	1.27	1.31
86	AB	67	C	N3-C4	-6.36	1.29	1.33
34	BA	1109	G	C5-C4	-6.36	1.33	1.38
34	BA	1276	G	C5-C4	-6.36	1.33	1.38
34	BA	1456	C	C3'-C2'	-6.36	1.45	1.52
34	BA	1503	U	C1'-N1	-6.36	1.38	1.46
85	AA	2225	G	C2'-C1'	-6.36	1.46	1.53
34	BA	651	U	C3'-C2'	-6.36	1.45	1.52
35	BB	1225	A	N9-C4	-6.36	1.34	1.37
35	BB	1236	A	P-O5'	-6.36	1.53	1.59
35	BB	1293	C	C4'-C3'	-6.36	1.46	1.53
35	BB	1508	G	O3'-P	-6.36	1.53	1.61
40	BG	50	G	C4'-C3'	-6.36	1.46	1.53
85	AA	358	U	C3'-C2'	-6.36	1.45	1.52
85	AA	710	A	C5'-C4'	-6.36	1.43	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1112	G	C6-N1	-6.36	1.35	1.39
34	BA	197	A	P-O5'	-6.36	1.53	1.59
34	BA	416	A	P-O5'	-6.36	1.53	1.59
34	BA	808	U	C4'-C3'	-6.36	1.46	1.53
34	BA	859	G	N3-C4	-6.36	1.30	1.35
34	BA	1195	G	O3'-P	-6.36	1.53	1.61
34	BA	1707	C	N3-C4	-6.36	1.29	1.33
35	BB	728	A	C2'-C1'	-6.36	1.46	1.53
35	BB	951	U	P-O5'	-6.36	1.53	1.59
35	BB	1305	A	N3-C4	-6.36	1.31	1.34
35	BB	1418	C	N3-C4	-6.36	1.29	1.33
37	BD	4	U	C4'-C3'	-6.36	1.46	1.53
85	AA	188	G	C3'-C2'	-6.36	1.45	1.52
34	BA	733	G	O3'-P	-6.36	1.53	1.61
34	BA	825	G	O3'-P	-6.36	1.53	1.61
34	BA	973	U	C5'-C4'	-6.36	1.43	1.51
34	BA	1255	G	C1'-N9	-6.36	1.38	1.46
35	BB	1331	U	C3'-C2'	-6.36	1.45	1.52
35	BB	1446	C	C3'-C2'	-6.36	1.45	1.52
35	BB	1519	U	O3'-P	-6.36	1.53	1.61
35	BB	1540	U	C2-N3	-6.36	1.33	1.37
38	BE	203	C	O4'-C1'	-6.36	1.33	1.41
40	BG	26	G	C2'-C1'	-6.36	1.46	1.53
85	AA	732	G	O3'-P	-6.36	1.53	1.61
85	AA	1485	G	O4'-C1'	-6.36	1.33	1.41
85	AA	2217	A	O3'-P	-6.36	1.53	1.61
34	BA	606	G	C2-N3	6.36	1.37	1.32
36	BC	11	G	C2'-C1'	-6.36	1.46	1.53
37	BD	96	C	N3-C4	-6.36	1.29	1.33
40	BG	12	A	N7-C5	-6.36	1.35	1.39
41	BH	40	C	P-O5'	-6.36	1.53	1.59
85	AA	390	U	N1-C2	-6.36	1.32	1.38
85	AA	637	U	C3'-C2'	-6.36	1.45	1.52
85	AA	654	A	C2'-C1'	-6.36	1.46	1.53
85	AA	1891	U	N3-C4	-6.36	1.32	1.38
34	BA	718	U	C1'-N1	-6.35	1.38	1.46
34	BA	1519	G	N9-C4	-6.35	1.32	1.38
34	BA	1545	C	C2-N3	-6.35	1.30	1.35
35	BB	541	U	P-O5'	-6.35	1.53	1.59
35	BB	1304	U	P-O5'	-6.35	1.53	1.59
37	BD	116	C	C3'-C2'	-6.35	1.45	1.52
85	AA	78	A	P-O5'	-6.35	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1272	G	C6-N1	-6.35	1.35	1.39
85	AA	2138	G	C2-N2	-6.35	1.28	1.34
34	BA	428	C	C2'-C1'	-6.35	1.46	1.53
34	BA	472	G	O3'-P	-6.35	1.53	1.61
34	BA	624	G	C2'-C1'	-6.35	1.46	1.53
34	BA	928	C	C2-N3	-6.35	1.30	1.35
34	BA	1728	G	C1'-N9	-6.35	1.38	1.46
35	BB	372	U	N3-C4	-6.35	1.32	1.38
35	BB	1092	G	C5'-C4'	6.35	1.58	1.51
35	BB	1460	G	C1'-N9	-6.35	1.38	1.46
38	BE	111	C	C2'-C1'	-6.35	1.46	1.53
85	AA	148	G	C6-N1	-6.35	1.35	1.39
85	AA	198	U	O3'-P	-6.35	1.53	1.61
85	AA	486	G	C1'-N9	-6.35	1.38	1.46
85	AA	842	G	O3'-P	-6.35	1.53	1.61
85	AA	1180	C	C2'-C1'	-6.35	1.46	1.53
85	AA	1366	A	C2'-C1'	-6.35	1.46	1.53
85	AA	2238	C	C2'-C1'	-6.35	1.46	1.53
34	BA	359	G	C5'-C4'	6.35	1.58	1.51
34	BA	542	A	C2'-C1'	-6.35	1.46	1.53
34	BA	843	G	C3'-C2'	-6.35	1.45	1.52
37	BD	68	C	C2-N3	-6.35	1.30	1.35
85	AA	69	C	C3'-C2'	-6.35	1.45	1.52
85	AA	773	G	C5-C4	-6.35	1.33	1.38
34	BA	1559	C	C4'-C3'	-6.35	1.46	1.53
35	BB	575	C	N1-C6	-6.35	1.33	1.37
35	BB	1354	C	C2-N3	-6.35	1.30	1.35
41	BH	32	U	C4'-C3'	-6.35	1.46	1.53
41	BH	121	A	N3-C4	-6.35	1.31	1.34
85	AA	333	A	C1'-N9	-6.35	1.38	1.46
85	AA	569	A	P-O5'	-6.35	1.53	1.59
85	AA	991	G	N3-C4	-6.35	1.31	1.35
34	BA	1108	U	C2'-C1'	-6.35	1.46	1.53
34	BA	1445	U	O3'-P	-6.35	1.53	1.61
35	BB	542	A	C1'-N9	-6.35	1.38	1.46
35	BB	1265	U	O3'-P	-6.35	1.53	1.61
57	BX	87	TYR	CA-CB	6.35	1.68	1.53
85	AA	766	G	C6-N1	-6.35	1.35	1.39
85	AA	867	G	C3'-O3'	-6.35	1.33	1.42
85	AA	912	C	N1-C6	-6.35	1.33	1.37
85	AA	1263	G	N9-C8	-6.35	1.33	1.37
86	AB	15	G	C5'-C4'	6.35	1.58	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	767	U	O3'-P	-6.35	1.53	1.61
35	BB	790	A	N9-C4	-6.35	1.34	1.37
38	BE	33	C	C2'-C1'	-6.35	1.46	1.53
40	BG	10	U	C3'-C2'	-6.35	1.45	1.52
85	AA	1490	A	P-O5'	-6.35	1.53	1.59
85	AA	2039	G	C5-C4	-6.35	1.33	1.38
85	AA	2083	G	P-O5'	-6.35	1.53	1.59
34	BA	255	G	N9-C4	-6.34	1.32	1.38
34	BA	309	U	O3'-P	-6.34	1.53	1.61
34	BA	499	C	P-O5'	-6.34	1.53	1.59
34	BA	518	C	C4'-C3'	-6.34	1.46	1.53
34	BA	628	U	C1'-N1	-6.34	1.38	1.46
34	BA	1582	C	C1'-N1	-6.34	1.38	1.46
34	BA	1642	A	C4'-O4'	-6.34	1.37	1.45
35	BB	555	G	C1'-N9	-6.34	1.38	1.46
35	BB	786	A	C5-C4	-6.34	1.34	1.38
35	BB	1080	U	C1'-N1	-6.34	1.38	1.46
35	BB	1145	G	C5-C6	-6.34	1.36	1.42
35	BB	1272	G	N9-C4	-6.34	1.32	1.38
35	BB	1534	U	C2-N3	-6.34	1.33	1.37
38	BE	145	A	C2'-C1'	-6.34	1.46	1.53
40	BG	19	C	O3'-P	-6.34	1.53	1.61
85	AA	1198	U	P-O5'	-6.34	1.53	1.59
85	AA	1292	A	C8-N7	-6.34	1.27	1.31
85	AA	1297	G	C2'-C1'	-6.34	1.46	1.53
85	AA	2086	C	C2'-C1'	-6.34	1.46	1.53
34	BA	840	U	P-O5'	-6.34	1.53	1.59
34	BA	955	G	C2'-C1'	-6.34	1.46	1.53
34	BA	1088	G	O3'-P	-6.34	1.53	1.61
34	BA	1269	C	C2'-C1'	-6.34	1.46	1.53
35	BB	501	G	N1-C2	-6.34	1.32	1.37
35	BB	1336	G	N1-C2	-6.34	1.32	1.37
35	BB	1492	C	O4'-C1'	-6.34	1.33	1.41
36	BC	117	A	C1'-N9	-6.34	1.38	1.46
85	AA	2186	U	C2'-C1'	-6.34	1.46	1.53
34	BA	456	G	O3'-P	-6.34	1.53	1.61
34	BA	977	G	N9-C4	-6.34	1.32	1.38
34	BA	1487	U	O4'-C1'	-6.34	1.33	1.41
34	BA	1521	C	C2-N3	-6.34	1.30	1.35
34	BA	1582	C	C2'-C1'	-6.34	1.46	1.53
41	BH	41	A	N7-C5	-6.34	1.35	1.39
85	AA	178	U	O3'-P	-6.34	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	879	G	C5-C4	-6.34	1.33	1.38
85	AA	1710	C	P-O5'	-6.34	1.53	1.59
34	BA	1045	C	C4-N4	-6.34	1.28	1.33
34	BA	1601	C	C4'-O4'	-6.34	1.37	1.45
35	BB	38	C	C3'-C2'	-6.34	1.45	1.52
35	BB	439	G	N3-C4	-6.34	1.31	1.35
37	BD	93	G	N1-C2	-6.34	1.32	1.37
39	BF	2	G	N9-C8	-6.34	1.33	1.37
39	BF	53	G	P-O5'	-6.34	1.53	1.59
40	BG	166	C	C2'-C1'	-6.34	1.46	1.53
85	AA	541	A	C3'-C2'	-6.34	1.45	1.52
85	AA	1486	G	C4'-C3'	-6.34	1.46	1.53
85	AA	2070	C	O3'-P	-6.34	1.53	1.61
85	AA	2220	U	C4'-C3'	-6.34	1.46	1.53
85	AA	2222	G	N7-C5	-6.34	1.35	1.39
35	BB	608	A	C3'-C2'	-6.34	1.45	1.52
35	BB	1025	A	C3'-C2'	6.34	1.59	1.52
85	AA	79	G	C4'-C3'	-6.34	1.46	1.53
85	AA	1485	G	C5-C6	-6.34	1.36	1.42
85	AA	1508	A	N9-C4	-6.34	1.34	1.37
34	BA	115	U	C4'-O4'	-6.34	1.37	1.45
34	BA	455	A	C3'-C2'	-6.34	1.45	1.52
34	BA	690	G	N9-C4	-6.34	1.32	1.38
34	BA	1222	C	P-O5'	-6.34	1.53	1.59
34	BA	1711	G	C8-N7	-6.34	1.27	1.30
38	BE	112	G	C6-N1	-6.34	1.35	1.39
85	AA	462	A	N3-C4	-6.34	1.31	1.34
85	AA	1977	G	C3'-C2'	-6.34	1.45	1.52
35	BB	95	A	O3'-P	-6.33	1.53	1.61
85	AA	1289	U	C2-N3	-6.33	1.33	1.37
85	AA	1477	A	P-O5'	-6.33	1.53	1.59
34	BA	684	G	P-O5'	6.33	1.66	1.59
34	BA	957	A	C4'-C3'	-6.33	1.46	1.53
35	BB	681	G	C2'-C1'	-6.33	1.46	1.53
35	BB	827	U	C3'-C2'	-6.33	1.45	1.52
35	BB	1251	G	O4'-C1'	-6.33	1.33	1.41
35	BB	1424	G	N3-C4	-6.33	1.31	1.35
38	BE	31	A	N3-C4	-6.33	1.31	1.34
85	AA	923	A	C3'-C2'	-6.33	1.45	1.52
85	AA	1673	A	C5-C4	-6.33	1.34	1.38
34	BA	216	C	N1-C2	-6.33	1.33	1.40
34	BA	746	C	C2-N3	-6.33	1.30	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1253	G	O3'-P	-6.33	1.53	1.61
35	BB	72	G	C1'-N9	-6.33	1.38	1.46
35	BB	1426	G	N9-C4	6.33	1.43	1.38
38	BE	59	U	C1'-N1	-6.33	1.38	1.46
85	AA	104	C	P-O5'	-6.33	1.53	1.59
85	AA	450	A	C5'-C4'	-6.33	1.43	1.51
85	AA	499	G	C2-N2	-6.33	1.28	1.34
85	AA	629	A	P-O5'	-6.33	1.53	1.59
85	AA	720	A	N7-C5	-6.33	1.35	1.39
85	AA	2030	U	C2'-C1'	-6.33	1.46	1.53
85	AA	2225	G	C5-C4	-6.33	1.33	1.38
34	BA	139	U	N3-C4	-6.33	1.32	1.38
34	BA	747	G	C3'-C2'	-6.33	1.45	1.52
35	BB	804	U	C2'-C1'	-6.33	1.46	1.53
38	BE	202	C	C2'-C1'	-6.33	1.46	1.53
40	BG	80	G	C2'-C1'	-6.33	1.46	1.53
34	BA	129	U	O3'-P	-6.33	1.53	1.61
34	BA	354	G	C3'-C2'	-6.33	1.45	1.52
34	BA	545	U	O3'-P	-6.33	1.53	1.61
34	BA	915	A	O4'-C1'	-6.33	1.33	1.41
34	BA	974	G	C6-N1	-6.33	1.35	1.39
34	BA	1246	G	N9-C4	-6.33	1.32	1.38
34	BA	1690	U	N3-C4	-6.33	1.32	1.38
35	BB	1020	U	C2-N3	-6.33	1.33	1.37
35	BB	1297	G	C2'-C1'	-6.33	1.46	1.53
35	BB	1370	G	C4'-C3'	-6.33	1.46	1.53
36	BC	21	U	P-O5'	-6.33	1.53	1.59
38	BE	24	G	P-O5'	-6.33	1.53	1.59
38	BE	163	A	P-O5'	-6.33	1.53	1.59
39	BF	54	U	C5'-C4'	6.33	1.58	1.51
85	AA	103	U	P-O5'	-6.33	1.53	1.59
85	AA	284	C	P-O5'	-6.33	1.53	1.59
85	AA	579	U	O3'-P	-6.33	1.53	1.61
85	AA	1465	C	O4'-C1'	-6.33	1.33	1.41
34	BA	671	C	O3'-P	-6.33	1.53	1.61
34	BA	1460	U	C2'-C1'	-6.33	1.46	1.53
35	BB	1491	G	C2'-C1'	-6.33	1.46	1.53
85	AA	1872	G	N7-C5	-6.33	1.35	1.39
34	BA	148	G	O3'-P	-6.33	1.53	1.61
34	BA	351	A	C2'-C1'	-6.33	1.46	1.53
34	BA	612	U	P-O5'	-6.33	1.53	1.59
34	BA	742	C	O4'-C1'	-6.33	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	862	C	O3'-P	6.33	1.68	1.61
34	BA	925	G	N9-C4	-6.33	1.32	1.38
34	BA	1203	G	C2-N2	-6.33	1.28	1.34
34	BA	1347	G	P-O5'	-6.33	1.53	1.59
34	BA	1450	G	P-O5'	-6.33	1.53	1.59
34	BA	1820	G	C2-N2	-6.33	1.28	1.34
35	BB	16	G	C6-N1	-6.33	1.35	1.39
35	BB	498	G	C5-C4	-6.33	1.33	1.38
35	BB	1459	U	C2'-C1'	-6.33	1.46	1.53
36	BC	49	G	C2'-C1'	-6.33	1.46	1.53
40	BG	31	G	N7-C5	-6.33	1.35	1.39
41	BH	16	A	O3'-P	-6.33	1.53	1.61
85	AA	655	U	N1-C2	-6.33	1.32	1.38
85	AA	1473	U	O3'-P	-6.33	1.53	1.61
85	AA	1501	A	C2'-C1'	-6.33	1.46	1.53
85	AA	1793	A	O3'-P	-6.33	1.53	1.61
85	AA	2142	A	C8-N7	-6.33	1.27	1.31
85	AA	2231	G	O3'-P	-6.33	1.53	1.61
34	BA	58	A	N7-C5	-6.32	1.35	1.39
34	BA	484	A	C4'-O4'	-6.32	1.37	1.45
34	BA	720	A	N9-C8	-6.32	1.32	1.37
34	BA	726	G	C2'-C1'	-6.32	1.46	1.53
34	BA	1671	A	N3-C4	-6.32	1.31	1.34
35	BB	52	G	C5-C4	-6.32	1.33	1.38
35	BB	807	U	O3'-P	-6.32	1.53	1.61
35	BB	892	U	P-O5'	-6.32	1.53	1.59
35	BB	1027	U	C2-N3	-6.32	1.33	1.37
37	BD	42	A	O3'-P	-6.32	1.53	1.61
40	BG	64	C	C3'-C2'	-6.32	1.45	1.52
85	AA	57	G	C5'-C4'	-6.32	1.43	1.51
85	AA	1668	G	C2'-C1'	-6.32	1.46	1.53
34	BA	344	G	N7-C5	-6.32	1.35	1.39
34	BA	827	A	P-O5'	-6.32	1.53	1.59
34	BA	1015	G	C6-N1	-6.32	1.35	1.39
34	BA	1706	A	C6-N1	-6.32	1.31	1.35
35	BB	1195	A	C8-N7	-6.32	1.27	1.31
40	BG	1	G	C1'-N9	-6.32	1.38	1.46
86	AB	9	A	N9-C4	-6.32	1.34	1.37
34	BA	83	G	C2'-C1'	-6.32	1.46	1.53
34	BA	875	G	O4'-C1'	-6.32	1.33	1.41
34	BA	1255	G	N9-C8	-6.32	1.33	1.37
34	BA	1267	A	O4'-C1'	-6.32	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1724	G	C4'-O4'	-6.32	1.37	1.45
35	BB	664	A	N7-C5	-6.32	1.35	1.39
35	BB	1376	G	C5-C4	-6.32	1.33	1.38
35	BB	1449	G	C1'-N9	-6.32	1.38	1.46
36	BC	151	G	N9-C4	-6.32	1.32	1.38
38	BE	12	A	C5-C4	-6.32	1.34	1.38
38	BE	58	U	C2'-C1'	-6.32	1.46	1.53
40	BG	27	C	C2'-C1'	-6.32	1.46	1.53
85	AA	270	A	N9-C4	-6.32	1.34	1.37
85	AA	482	C	P-O5'	-6.32	1.53	1.59
85	AA	674	U	N3-C4	-6.32	1.32	1.38
85	AA	790	A	N7-C5	-6.32	1.35	1.39
85	AA	1520	A	C2'-C1'	-6.32	1.46	1.53
36	BC	49	G	P-O5'	-6.32	1.53	1.59
36	BC	102	G	N3-C4	-6.32	1.31	1.35
36	BC	152	C	C2'-C1'	-6.32	1.46	1.53
85	AA	273	C	C3'-C2'	-6.32	1.45	1.52
85	AA	1163	G	N9-C8	-6.32	1.33	1.37
85	AA	1221	G	C8-N7	-6.32	1.27	1.30
85	AA	1953	G	C2'-C1'	-6.32	1.46	1.53
85	AA	2155	U	O3'-P	-6.32	1.53	1.61
34	BA	193	C	C1'-N1	-6.32	1.38	1.46
34	BA	675	C	N1-C6	-6.32	1.33	1.37
34	BA	692	U	O4'-C1'	-6.32	1.33	1.41
34	BA	703	U	C2'-C1'	-6.32	1.46	1.53
34	BA	787	A	N3-C4	-6.32	1.31	1.34
34	BA	1531	G	C3'-C2'	-6.32	1.45	1.52
35	BB	1253	U	C2'-C1'	-6.32	1.46	1.53
35	BB	1262	A	N9-C4	-6.32	1.34	1.37
35	BB	1467	A	C4'-C3'	6.32	1.60	1.53
38	BE	94	U	C4'-C3'	-6.32	1.46	1.53
85	AA	194	U	O3'-P	-6.32	1.53	1.61
85	AA	442	G	C5'-C4'	-6.32	1.43	1.51
85	AA	1881	C	N1-C6	-6.32	1.33	1.37
34	BA	412	G	C2-N2	-6.32	1.28	1.34
34	BA	513	U	C4-O4	-6.32	1.18	1.23
34	BA	776	U	C2-N3	-6.32	1.33	1.37
34	BA	1039	G	C6-N1	-6.32	1.35	1.39
34	BA	1210	A	P-O5'	-6.32	1.53	1.59
34	BA	1401	C	O3'-P	-6.32	1.53	1.61
34	BA	1630	A	C4'-C3'	6.32	1.60	1.53
35	BB	26	C	C2'-C1'	-6.32	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	614	U	N3-C4	-6.32	1.32	1.38
35	BB	622	G	N9-C8	-6.32	1.33	1.37
35	BB	1425	A	P-O5'	-6.32	1.53	1.59
85	AA	1651	C	P-O5'	-6.32	1.53	1.59
85	AA	1797	U	C2'-C1'	-6.32	1.46	1.53
85	AA	2186	U	C1'-N1	-6.32	1.38	1.46
34	BA	748	C	C2-N3	-6.31	1.30	1.35
35	BB	538	A	C1'-N9	-6.31	1.38	1.46
35	BB	661	G	C2-N2	-6.31	1.28	1.34
35	BB	1352	C	C3'-C2'	-6.31	1.45	1.52
35	BB	1439	U	C2'-C1'	-6.31	1.46	1.53
35	BB	1466	A	N7-C5	-6.31	1.35	1.39
36	BC	62	A	O3'-P	-6.31	1.53	1.61
37	BD	23	A	C1'-N9	-6.31	1.38	1.46
34	BA	536	C	C3'-C2'	-6.31	1.45	1.52
34	BA	683	C	P-O5'	-6.31	1.53	1.59
34	BA	1035	A	N9-C4	-6.31	1.34	1.37
34	BA	1081	U	O3'-P	-6.31	1.53	1.61
34	BA	1644	A	C5-C4	-6.31	1.34	1.38
34	BA	1707	C	C2'-C1'	-6.31	1.46	1.53
34	BA	1708	A	O3'-P	-6.31	1.53	1.61
35	BB	402	G	C2'-C1'	-6.31	1.46	1.53
35	BB	472	C	C2-N3	-6.31	1.30	1.35
35	BB	878	G	C5'-C4'	6.31	1.58	1.51
35	BB	967	G	N1-C2	-6.31	1.32	1.37
35	BB	1057	G	N3-C4	-6.31	1.31	1.35
36	BC	14	G	O3'-P	-6.31	1.53	1.61
40	BG	94	G	N1-C2	-6.31	1.32	1.37
85	AA	97	A	N9-C8	-6.31	1.32	1.37
85	AA	289	G	C4'-O4'	6.31	1.53	1.45
85	AA	957	A	P-O5'	-6.31	1.53	1.59
85	AA	2093	U	P-O5'	-6.31	1.53	1.59
86	AB	2	C	C2'-C1'	-6.31	1.46	1.53
34	BA	1017	C	C1'-N1	-6.31	1.38	1.46
35	BB	16	G	N9-C4	-6.31	1.32	1.38
41	BH	19	G	C4'-C3'	-6.31	1.46	1.53
85	AA	2147	A	C5-C6	-6.31	1.35	1.41
34	BA	118	C	O4'-C1'	-6.31	1.33	1.41
34	BA	678	C	C4-N4	-6.31	1.28	1.33
34	BA	681	G	C8-N7	-6.31	1.27	1.30
34	BA	1546	C	C1'-N1	-6.31	1.38	1.46
35	BB	452	A	C5-C4	-6.31	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1161	G	C6-N1	-6.31	1.35	1.39
36	BC	108	A	C1'-N9	-6.31	1.38	1.46
40	BG	86	U	C4'-C3'	-6.31	1.46	1.53
40	BG	116	G	N3-C4	-6.31	1.31	1.35
85	AA	707	U	N3-C4	-6.31	1.32	1.38
85	AA	1282	A	C2'-C1'	-6.31	1.46	1.53
85	AA	1560	A	O3'-P	-6.31	1.53	1.61
85	AA	1654	G	C1'-N9	-6.31	1.38	1.46
85	AA	2122	A	C2'-C1'	-6.31	1.46	1.53
85	AA	2208	G	C3'-C2'	-6.31	1.45	1.52
34	BA	480	G	N7-C5	-6.31	1.35	1.39
34	BA	800	G	C3'-C2'	-6.31	1.45	1.52
34	BA	950	C	C2'-C1'	-6.31	1.46	1.53
34	BA	1062	G	N1-C2	-6.31	1.32	1.37
34	BA	1803	A	P-O5'	-6.31	1.53	1.59
36	BC	35	C	C2-N3	-6.31	1.30	1.35
40	BG	113	G	C5-C4	-6.31	1.33	1.38
85	AA	447	C	P-O5'	-6.31	1.53	1.59
85	AA	2176	U	C4'-O4'	-6.31	1.37	1.45
85	AA	2192	A	C1'-N9	-6.31	1.38	1.46
34	BA	952	G	C6-N1	-6.31	1.35	1.39
35	BB	996	G	N1-C2	-6.31	1.32	1.37
41	BH	21	G	C1'-N9	-6.31	1.38	1.46
34	BA	788	C	N1-C6	-6.30	1.33	1.37
34	BA	1073	G	C3'-C2'	-6.30	1.45	1.52
34	BA	1333	G	C2-N2	-6.30	1.28	1.34
34	BA	1412	G	C4'-C3'	-6.30	1.46	1.53
36	BC	59	A	P-O5'	-6.30	1.53	1.59
40	BG	71	C	C2'-C1'	-6.30	1.46	1.53
85	AA	381	A	O4'-C1'	-6.30	1.33	1.41
85	AA	506	G	C2'-C1'	-6.30	1.46	1.53
34	BA	246	G	C1'-N9	-6.30	1.38	1.46
34	BA	991	U	P-O5'	-6.30	1.53	1.59
34	BA	1451	A	P-O5'	-6.30	1.53	1.59
35	BB	1033	U	C3'-C2'	-6.30	1.45	1.52
35	BB	1421	C	C3'-C2'	-6.30	1.45	1.52
85	AA	626	G	C3'-C2'	-6.30	1.45	1.52
34	BA	429	G	N9-C4	-6.30	1.32	1.38
34	BA	574	U	P-O5'	-6.30	1.53	1.59
35	BB	44	C	O3'-P	-6.30	1.53	1.61
35	BB	126	C	N3-C4	-6.30	1.29	1.33
35	BB	542	A	N7-C5	-6.30	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	795	A	N7-C5	-6.30	1.35	1.39
35	BB	887	G	N9-C4	6.30	1.43	1.38
35	BB	1405	G	P-O5'	-6.30	1.53	1.59
39	BF	57	C	C1'-N1	-6.30	1.38	1.46
41	BH	13	C	P-O5'	-6.30	1.53	1.59
41	BH	21	G	O3'-P	-6.30	1.53	1.61
85	AA	369	A	C2'-C1'	-6.30	1.46	1.53
85	AA	1149	A	P-O5'	-6.30	1.53	1.59
85	AA	1459	C	O3'-P	-6.30	1.53	1.61
85	AA	2061	C	C2'-C1'	-6.30	1.46	1.53
86	AB	69	G	C2-N2	-6.30	1.28	1.34
34	BA	35	U	C2-N3	-6.30	1.33	1.37
34	BA	612	U	O3'-P	-6.30	1.53	1.61
34	BA	770	G	O3'-P	-6.30	1.53	1.61
34	BA	1082	U	O4'-C1'	-6.30	1.33	1.41
34	BA	1544	G	N3-C4	-6.30	1.31	1.35
35	BB	436	G	C8-N7	-6.30	1.27	1.30
36	BC	56	G	N7-C5	-6.30	1.35	1.39
36	BC	161	U	N3-C4	-6.30	1.32	1.38
37	BD	25	G	C3'-C2'	-6.30	1.45	1.52
37	BD	49	A	C6-N1	-6.30	1.31	1.35
38	BE	25	U	N1-C6	-6.30	1.32	1.38
40	BG	44	G	O3'-P	-6.30	1.53	1.61
40	BG	108	G	C3'-C2'	-6.30	1.45	1.52
85	AA	449	G	C5-C4	-6.30	1.33	1.38
85	AA	1526	G	C2-N2	-6.30	1.28	1.34
85	AA	2223	C	C2-N3	-6.30	1.30	1.35
35	BB	520	G	C1'-N9	-6.30	1.38	1.46
35	BB	607	G	N1-C2	-6.30	1.32	1.37
36	BC	76	C	N1-C6	-6.30	1.33	1.37
34	BA	762	A	O3'-P	-6.30	1.53	1.61
34	BA	843	G	C2'-C1'	-6.30	1.46	1.53
35	BB	265	C	P-O5'	-6.30	1.53	1.59
35	BB	841	U	C3'-C2'	-6.30	1.45	1.52
35	BB	1538	G	C2'-C1'	-6.30	1.46	1.53
36	BC	121	G	C5-C6	-6.30	1.36	1.42
85	AA	2	A	O3'-P	-6.30	1.53	1.61
85	AA	402	G	P-O5'	-6.30	1.53	1.59
85	AA	1296	G	C2'-C1'	-6.30	1.46	1.53
34	BA	1164	C	C4'-C3'	-6.29	1.46	1.53
34	BA	1466	U	N3-C4	-6.29	1.32	1.38
34	BA	1556	A	O4'-C1'	-6.29	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1706	A	C1'-N9	-6.29	1.38	1.46
85	AA	1129	A	C1'-N9	-6.29	1.38	1.46
85	AA	1501	A	C1'-N9	-6.29	1.38	1.46
34	BA	15	G	N1-C2	-6.29	1.32	1.37
34	BA	160	G	N9-C8	-6.29	1.33	1.37
34	BA	205	G	C5-C6	-6.29	1.36	1.42
34	BA	222	C	C2-N3	-6.29	1.30	1.35
34	BA	596	G	N1-C2	-6.29	1.32	1.37
34	BA	998	U	O3'-P	-6.29	1.53	1.61
34	BA	1477	C	N1-C6	-6.29	1.33	1.37
34	BA	1685	C	C4'-O4'	-6.29	1.37	1.45
35	BB	1048	A	N7-C5	-6.29	1.35	1.39
36	BC	88	A	C8-N7	-6.29	1.27	1.31
85	AA	107	A	N7-C5	-6.29	1.35	1.39
85	AA	1528	A	C1'-N9	-6.29	1.38	1.46
85	AA	1697	C	N1-C6	-6.29	1.33	1.37
85	AA	2012	G	N7-C5	-6.29	1.35	1.39
85	AA	2187	G	C5-C4	-6.29	1.33	1.38
34	BA	109	A	P-O5'	-6.29	1.53	1.59
34	BA	761	U	C4'-C3'	-6.29	1.46	1.53
34	BA	809	U	N3-C4	-6.29	1.32	1.38
34	BA	852	C	C2-N3	-6.29	1.30	1.35
34	BA	935	A	C2'-C1'	-6.29	1.46	1.53
34	BA	1273	U	P-O5'	-6.29	1.53	1.59
34	BA	1640	G	C5-C4	-6.29	1.33	1.38
35	BB	621	C	N1-C6	-6.29	1.33	1.37
35	BB	837	A	N3-C4	-6.29	1.31	1.34
35	BB	879	G	N7-C5	-6.29	1.35	1.39
35	BB	1110	G	C2'-C1'	-6.29	1.46	1.53
39	BF	41	U	C2'-C1'	-6.29	1.46	1.53
85	AA	386	G	C8-N7	-6.29	1.27	1.30
85	AA	523	U	O3'-P	-6.29	1.53	1.61
85	AA	1206	A	C5-C4	-6.29	1.34	1.38
34	BA	919	A	C2'-C1'	-6.29	1.46	1.53
34	BA	1052	G	C5-C6	-6.29	1.36	1.42
34	BA	1190	A	C1'-N9	-6.29	1.38	1.46
35	BB	601	U	C4'-C3'	-6.29	1.46	1.53
41	BH	106	G	N7-C5	-6.29	1.35	1.39
85	AA	585	G	N9-C4	-6.29	1.32	1.38
85	AA	867	G	C2'-C1'	-6.29	1.46	1.53
85	AA	2042	G	C1'-N9	-6.29	1.38	1.46
34	BA	1842	U	C3'-C2'	-6.29	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	55	C	O3'-P	-6.29	1.53	1.61
35	BB	1277	A	C6-N1	-6.29	1.31	1.35
35	BB	1396	G	C5-C4	-6.29	1.33	1.38
36	BC	113	G	C2-N2	-6.29	1.28	1.34
36	BC	163	A	C5-C4	-6.29	1.34	1.38
39	BF	57	C	O3'-P	-6.29	1.53	1.61
85	AA	18	C	O3'-P	-6.29	1.53	1.61
85	AA	79	G	C1'-N9	-6.29	1.38	1.46
85	AA	432	A	C3'-C2'	-6.29	1.45	1.52
85	AA	995	G	C2-N2	-6.29	1.28	1.34
34	BA	572	G	C5-C6	6.29	1.48	1.42
34	BA	764	G	N1-C2	-6.29	1.32	1.37
34	BA	1276	G	C2'-C1'	-6.29	1.46	1.53
35	BB	1201	G	O4'-C1'	-6.29	1.33	1.41
41	BH	106	G	C2'-C1'	-6.29	1.46	1.53
34	BA	456	G	C1'-N9	-6.29	1.38	1.46
34	BA	625	U	C2'-C1'	-6.29	1.46	1.53
34	BA	1096	C	C2'-C1'	-6.29	1.46	1.53
34	BA	1502	G	O3'-P	-6.29	1.53	1.61
34	BA	1668	C	C2'-C1'	-6.29	1.46	1.53
35	BB	775	U	C2'-C1'	-6.29	1.46	1.53
35	BB	1288	G	N9-C4	-6.29	1.32	1.38
35	BB	1359	G	N9-C8	-6.29	1.33	1.37
35	BB	1504	U	O3'-P	-6.29	1.53	1.61
85	AA	426	C	C3'-C2'	-6.29	1.45	1.52
85	AA	611	G	C5-C4	-6.29	1.33	1.38
85	AA	1460	G	O4'-C1'	-6.29	1.33	1.41
34	BA	83	G	N9-C8	-6.28	1.33	1.37
34	BA	107	C	O3'-P	-6.28	1.53	1.61
34	BA	364	C	C3'-C2'	-6.28	1.45	1.52
34	BA	971	G	N7-C5	-6.28	1.35	1.39
35	BB	104	G	C6-N1	-6.28	1.35	1.39
35	BB	459	U	P-O5'	-6.28	1.53	1.59
35	BB	810	G	N9-C8	-6.28	1.33	1.37
37	BD	91	U	N3-C4	-6.28	1.32	1.38
40	BG	162	A	O3'-P	-6.28	1.53	1.61
85	AA	164	G	C5'-C4'	-6.28	1.43	1.51
85	AA	543	A	N3-C4	-6.28	1.31	1.34
85	AA	626	G	C5'-C4'	-6.28	1.43	1.51
85	AA	1733	G	C4'-C3'	6.28	1.60	1.53
34	BA	341	U	C4'-O4'	-6.28	1.37	1.45
34	BA	1141	C	P-O5'	-6.28	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	28	C	C4'-O4'	-6.28	1.37	1.45
39	BF	4	A	N7-C5	-6.28	1.35	1.39
83	Bx	200	GLY	CA-C	-6.28	1.41	1.51
85	AA	779	G	C2'-C1'	-6.28	1.46	1.53
34	BA	14	G	C1'-N9	-6.28	1.38	1.46
34	BA	102	G	N7-C5	-6.28	1.35	1.39
34	BA	240	C	C3'-C2'	-6.28	1.45	1.52
34	BA	875	G	C3'-C2'	-6.28	1.45	1.52
34	BA	1003	A	N9-C8	-6.28	1.32	1.37
34	BA	1024	A	O3'-P	-6.28	1.53	1.61
34	BA	1226	G	N9-C8	-6.28	1.33	1.37
35	BB	528	G	O3'-P	-6.28	1.53	1.61
35	BB	902	C	O3'-P	-6.28	1.53	1.61
38	BE	143	A	O3'-P	-6.28	1.53	1.61
39	BF	35	C	C4-C5	-6.28	1.38	1.43
85	AA	17	C	O3'-P	-6.28	1.53	1.61
85	AA	93	G	C6-N1	-6.28	1.35	1.39
85	AA	1002	G	P-O5'	-6.28	1.53	1.59
85	AA	2141	G	N9-C8	-6.28	1.33	1.37
34	BA	366	G	C4'-C3'	-6.28	1.46	1.53
34	BA	890	G	C6-N1	-6.28	1.35	1.39
34	BA	1113	A	C3'-C2'	-6.28	1.45	1.52
34	BA	1458	A	O3'-P	-6.28	1.53	1.61
38	BE	165	U	C4'-C3'	-6.28	1.46	1.53
85	AA	116	G	O3'-P	-6.28	1.53	1.61
85	AA	1699	A	C2'-C1'	-6.28	1.46	1.53
34	BA	912	G	C6-N1	-6.28	1.35	1.39
34	BA	1062	G	C2'-C1'	-6.28	1.46	1.53
35	BB	404	A	C5-C4	-6.28	1.34	1.38
35	BB	1108	G	C5-C4	-6.28	1.33	1.38
38	BE	67	A	C3'-C2'	-6.28	1.45	1.52
38	BE	176	G	C6-N1	-6.28	1.35	1.39
85	AA	1035	C	C2'-C1'	-6.28	1.46	1.53
34	BA	151	A	C3'-C2'	6.28	1.59	1.52
34	BA	736	G	C8-N7	-6.28	1.27	1.30
34	BA	1279	U	O3'-P	-6.28	1.53	1.61
34	BA	1293	A	N3-C4	-6.28	1.31	1.34
34	BA	1312	A	C1'-N9	-6.28	1.38	1.46
34	BA	1560	U	C2'-C1'	-6.28	1.46	1.53
34	BA	1672	C	C2'-C1'	-6.28	1.46	1.53
34	BA	1700	C	C2-N3	-6.28	1.30	1.35
35	BB	96	A	C5-C6	-6.28	1.35	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	654	C	N1-C6	-6.28	1.33	1.37
35	BB	1102	U	C3'-C2'	-6.28	1.45	1.52
35	BB	1546	C	C2'-C1'	-6.28	1.46	1.53
41	BH	46	C	C2-N3	-6.28	1.30	1.35
85	AA	369	A	C3'-C2'	-6.28	1.45	1.52
85	AA	923	A	C3'-O3'	-6.28	1.33	1.42
36	BC	138	C	C5'-C4'	-6.27	1.43	1.51
39	BF	57	C	C5'-C4'	-6.27	1.43	1.51
85	AA	1289	U	C3'-C2'	-6.27	1.45	1.52
85	AA	2146	G	C4'-C3'	-6.27	1.46	1.53
34	BA	194	G	C4'-C3'	-6.27	1.46	1.53
34	BA	245	U	C3'-C2'	-6.27	1.45	1.52
34	BA	288	U	N3-C4	-6.27	1.32	1.38
34	BA	699	G	O3'-P	-6.27	1.53	1.61
34	BA	1292	A	C5-C4	-6.27	1.34	1.38
34	BA	1740	U	C2-N3	-6.27	1.33	1.37
35	BB	1115	G	C6-N1	-6.27	1.35	1.39
36	BC	52	A	N3-C4	-6.27	1.31	1.34
36	BC	121	G	N7-C5	-6.27	1.35	1.39
85	AA	126	U	C2'-C1'	-6.27	1.46	1.53
85	AA	672	U	C4'-C3'	-6.27	1.46	1.53
85	AA	927	A	C1'-N9	-6.27	1.38	1.46
4	A3	49	TYR	CB-CG	-6.27	1.42	1.51
34	BA	1283	U	C4'-C3'	-6.27	1.46	1.53
35	BB	1308	G	C2-N2	-6.27	1.28	1.34
35	BB	1459	U	N3-C4	-6.27	1.32	1.38
85	AA	696	G	P-O5'	-6.27	1.53	1.59
86	AB	24	G	C2-N2	-6.27	1.28	1.34
34	BA	401	A	O3'-P	-6.27	1.53	1.61
34	BA	490	A	C2'-C1'	-6.27	1.46	1.53
34	BA	979	G	C5-C6	-6.27	1.36	1.42
34	BA	1305	A	C6-N1	-6.27	1.31	1.35
35	BB	1507	U	C2'-C1'	-6.27	1.46	1.53
36	BC	112	G	C2'-C1'	-6.27	1.46	1.53
37	BD	117	U	O3'-P	-6.27	1.53	1.61
38	BE	176	G	C3'-C2'	-6.27	1.45	1.52
85	AA	21	U	P-O5'	-6.27	1.53	1.59
85	AA	57	G	C4'-C3'	-6.27	1.46	1.53
85	AA	407	G	N1-C2	-6.27	1.32	1.37
85	AA	580	C	C2'-C1'	-6.27	1.46	1.53
85	AA	2130	G	N1-C2	-6.27	1.32	1.37
85	AA	2242	U	P-O5'	-6.27	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	162	G	C5-C6	-6.27	1.36	1.42
34	BA	524	G	N9-C4	-6.27	1.32	1.38
34	BA	702	G	O3'-P	-6.27	1.53	1.61
34	BA	786	U	C4-C5	-6.27	1.38	1.43
34	BA	1155	U	C2'-C1'	-6.27	1.46	1.53
34	BA	1333	G	C5-C4	-6.27	1.33	1.38
34	BA	1579	G	C2'-C1'	-6.27	1.46	1.53
34	BA	1604	A	C1'-N9	-6.27	1.38	1.46
35	BB	12	G	C3'-C2'	-6.27	1.45	1.52
35	BB	421	U	P-O5'	-6.27	1.53	1.59
35	BB	1243	A	C3'-C2'	-6.27	1.45	1.52
35	BB	1469	A	N9-C4	-6.27	1.34	1.37
36	BC	50	C	C2-N3	-6.27	1.30	1.35
36	BC	92	C	C4'-C3'	-6.27	1.46	1.53
41	BH	23	G	C2-N2	-6.27	1.28	1.34
50	BQ	100	LYS	C-N	-6.27	1.22	1.34
85	AA	337	C	P-O5'	-6.27	1.53	1.59
85	AA	495	G	N7-C5	-6.27	1.35	1.39
85	AA	635	G	O3'-P	-6.27	1.53	1.61
85	AA	1179	A	N7-C5	-6.27	1.35	1.39
85	AA	1258	U	O3'-P	-6.27	1.53	1.61
85	AA	1449	C	C2'-C1'	-6.27	1.46	1.53
34	BA	461	A	C5-C4	-6.27	1.34	1.38
36	BC	163	A	C2'-C1'	-6.27	1.46	1.53
85	AA	472	A	N7-C5	-6.27	1.35	1.39
85	AA	867	G	C1'-N9	-6.27	1.38	1.46
34	BA	82	A	C4'-O4'	-6.26	1.37	1.45
34	BA	92	G	N1-C2	-6.26	1.32	1.37
34	BA	387	A	C1'-N9	-6.26	1.38	1.46
34	BA	1165	A	C2'-C1'	-6.26	1.46	1.53
34	BA	1320	A	O3'-P	-6.26	1.53	1.61
34	BA	1557	G	C5-C6	-6.26	1.36	1.42
34	BA	1563	G	C4'-C3'	-6.26	1.46	1.53
34	BA	1793	G	C2-N3	-6.26	1.27	1.32
35	BB	33	A	N3-C4	-6.26	1.31	1.34
35	BB	1075	A	O3'-P	-6.26	1.53	1.61
36	BC	128	U	C2-N3	-6.26	1.33	1.37
40	BG	88	G	C3'-C2'	-6.26	1.45	1.52
85	AA	80	G	C2-N2	-6.26	1.28	1.34
85	AA	241	U	C4'-C3'	-6.26	1.46	1.53
85	AA	333	A	C8-N7	-6.26	1.27	1.31
85	AA	418	G	C1'-N9	-6.26	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	431	G	N9-C4	-6.26	1.32	1.38
85	AA	666	A	P-O5'	-6.26	1.53	1.59
85	AA	758	C	C2-N3	-6.26	1.30	1.35
85	AA	1185	G	O3'-P	-6.26	1.53	1.61
34	BA	373	G	O3'-P	-6.26	1.53	1.61
34	BA	427	G	C3'-C2'	-6.26	1.45	1.52
34	BA	714	G	P-O5'	-6.26	1.53	1.59
34	BA	1546	C	C3'-C2'	-6.26	1.45	1.52
35	BB	503	G	O3'-P	-6.26	1.53	1.61
85	AA	352	G	C5-C4	-6.26	1.33	1.38
85	AA	443	A	C4'-C3'	-6.26	1.46	1.53
85	AA	595	A	O3'-P	-6.26	1.53	1.61
34	BA	122	U	C2-N3	-6.26	1.33	1.37
34	BA	432	A	P-O5'	-6.26	1.53	1.59
34	BA	849	G	O3'-P	-6.26	1.53	1.61
34	BA	865	C	N3-C4	6.26	1.38	1.33
34	BA	943	G	C2'-C1'	-6.26	1.46	1.53
35	BB	546	A	P-O5'	-6.26	1.53	1.59
35	BB	596	C	C4'-C3'	-6.26	1.46	1.53
35	BB	830	G	N9-C4	-6.26	1.32	1.38
35	BB	1284	U	N3-C4	-6.26	1.32	1.38
37	BD	77	A	N9-C8	-6.26	1.32	1.37
39	BF	58	U	C1'-N1	-6.26	1.38	1.46
85	AA	65	A	P-O5'	-6.26	1.53	1.59
85	AA	192	G	O3'-P	-6.26	1.53	1.61
85	AA	504	U	C3'-C2'	-6.26	1.45	1.52
85	AA	706	U	N3-C4	-6.26	1.32	1.38
85	AA	941	C	P-O5'	-6.26	1.53	1.59
85	AA	1251	G	C8-N7	-6.26	1.27	1.30
85	AA	1674	G	C5-C4	-6.26	1.33	1.38
85	AA	2201	A	N9-C8	-6.26	1.32	1.37
34	BA	367	G	C5-C6	-6.26	1.36	1.42
34	BA	794	G	C4'-C3'	-6.26	1.46	1.53
34	BA	986	G	C2'-C1'	-6.26	1.46	1.53
34	BA	1213	A	C8-N7	-6.26	1.27	1.31
35	BB	1391	G	N9-C4	-6.26	1.32	1.38
39	BF	30	C	P-O5'	-6.26	1.53	1.59
82	Bw	137	PRO	CA-C	-6.26	1.40	1.52
85	AA	395	G	C4'-O4'	-6.26	1.37	1.45
85	AA	2130	G	C5-C6	-6.26	1.36	1.42
34	BA	1688	G	C5-C4	-6.26	1.33	1.38
35	BB	702	G	C3'-C2'	-6.26	1.45	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	782	A	C4'-C3'	-6.26	1.46	1.53
35	BB	1233	U	P-O5'	-6.26	1.53	1.59
85	AA	759	G	C2'-C1'	-6.26	1.46	1.53
34	BA	263	G	O3'-P	-6.26	1.53	1.61
34	BA	991	U	C2'-C1'	-6.26	1.46	1.53
34	BA	1250	C	C3'-C2'	-6.26	1.45	1.52
34	BA	1514	A	C5-C4	-6.26	1.34	1.38
34	BA	1604	A	C2'-C1'	-6.26	1.46	1.53
34	BA	1637	G	C2'-C1'	-6.26	1.46	1.53
35	BB	1205	A	N9-C8	-6.26	1.32	1.37
39	BF	5	U	N3-C4	-6.26	1.32	1.38
85	AA	156	G	N1-C2	-6.26	1.32	1.37
85	AA	883	A	C4'-C3'	-6.26	1.46	1.53
85	AA	1508	A	C5-C4	-6.26	1.34	1.38
34	BA	1175	G	C2-N2	-6.25	1.28	1.34
34	BA	1416	C	C1'-N1	-6.25	1.38	1.46
34	BA	1498	A	P-O5'	-6.25	1.53	1.59
34	BA	1514	A	O3'-P	-6.25	1.53	1.61
35	BB	22	A	P-O5'	-6.25	1.53	1.59
35	BB	82	G	C3'-C2'	-6.25	1.45	1.52
35	BB	785	G	O3'-P	-6.25	1.53	1.61
40	BG	72	G	C1'-N9	-6.25	1.38	1.46
41	BH	13	C	C4'-C3'	-6.25	1.46	1.53
85	AA	9	U	P-O5'	-6.25	1.53	1.59
34	BA	15	G	C5-C4	-6.25	1.33	1.38
34	BA	855	C	C3'-C2'	-6.25	1.45	1.52
34	BA	966	G	N9-C8	-6.25	1.33	1.37
34	BA	1704	G	N9-C4	-6.25	1.32	1.38
34	BA	1800	G	C2-N3	-6.25	1.27	1.32
35	BB	1084	A	C1'-N9	-6.25	1.38	1.46
35	BB	1544	A	P-O5'	-6.25	1.53	1.59
36	BC	66	G	C5-C4	-6.25	1.33	1.38
40	BG	56	G	C4'-C3'	-6.25	1.46	1.53
40	BG	141	A	C5-C4	-6.25	1.34	1.38
85	AA	30	G	C1'-N9	-6.25	1.38	1.46
85	AA	2114	U	O3'-P	-6.25	1.53	1.61
85	AA	2204	A	P-O5'	-6.25	1.53	1.59
34	BA	793	A	N9-C8	-6.25	1.32	1.37
34	BA	1343	A	P-O5'	-6.25	1.53	1.59
34	BA	1711	G	C4'-C3'	-6.25	1.46	1.53
35	BB	367	C	P-O5'	-6.25	1.53	1.59
35	BB	564	U	O3'-P	-6.25	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	80	G	C2-N2	-6.25	1.28	1.34
39	BF	54	U	C4-C5	-6.25	1.38	1.43
41	BH	21	G	C6-N1	-6.25	1.35	1.39
85	AA	718	C	C2-N3	-6.25	1.30	1.35
85	AA	875	C	C2-N3	6.25	1.40	1.35
85	AA	1151	G	C2'-C1'	-6.25	1.46	1.53
85	AA	1223	A	C5-C4	-6.25	1.34	1.38
85	AA	1866	A	O3'-P	-6.25	1.53	1.61
34	BA	783	U	C2'-C1'	-6.25	1.46	1.53
34	BA	1062	G	C5-C4	-6.25	1.33	1.38
34	BA	1211	G	C4'-O4'	-6.25	1.37	1.45
40	BG	99	A	N7-C5	-6.25	1.35	1.39
85	AA	999	A	N9-C4	6.25	1.41	1.37
85	AA	1707	G	P-O5'	-6.25	1.53	1.59
85	AA	2156	C	P-O5'	-6.25	1.53	1.59
34	BA	173	U	C2-N3	-6.25	1.33	1.37
34	BA	196	A	N7-C5	-6.25	1.35	1.39
34	BA	521	C	C2'-C1'	-6.25	1.46	1.53
34	BA	794	G	N9-C8	-6.25	1.33	1.37
34	BA	1812	C	C2'-C1'	-6.25	1.46	1.53
35	BB	461	U	C4-C5	-6.25	1.38	1.43
35	BB	1117	G	P-O5'	-6.25	1.53	1.59
37	BD	101	A	C2'-C1'	-6.25	1.46	1.53
38	BE	90	G	P-O5'	-6.25	1.53	1.59
38	BE	197	A	O3'-P	-6.25	1.53	1.61
85	AA	9	U	C2'-C1'	-6.25	1.46	1.53
85	AA	24	U	C3'-C2'	-6.25	1.45	1.52
85	AA	107	A	N3-C4	-6.25	1.31	1.34
85	AA	109	G	C2'-C1'	-6.25	1.46	1.53
85	AA	395	G	N7-C5	-6.25	1.35	1.39
85	AA	442	G	C2-N2	-6.25	1.28	1.34
34	BA	406	G	N7-C5	-6.25	1.35	1.39
34	BA	416	A	O3'-P	-6.25	1.53	1.61
34	BA	496	G	C1'-N9	-6.25	1.38	1.46
34	BA	959	G	O4'-C1'	-6.25	1.33	1.41
35	BB	27	C	C3'-O3'	-6.25	1.33	1.42
35	BB	333	C	P-O5'	-6.25	1.53	1.59
35	BB	963	G	C3'-C2'	-6.25	1.45	1.52
35	BB	1125	A	C2'-C1'	-6.25	1.46	1.53
36	BC	62	A	C1'-N9	-6.25	1.38	1.46
85	AA	634	U	P-O5'	-6.25	1.53	1.59
34	BA	1210	A	N3-C4	-6.25	1.31	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1645	C	C2-N3	-6.25	1.30	1.35
35	BB	1046	C	C4'-C3'	-6.25	1.46	1.53
38	BE	20	C	N1-C6	-6.25	1.33	1.37
38	BE	91	G	C5-C4	-6.25	1.33	1.38
85	AA	1665	G	N7-C5	-6.25	1.35	1.39
34	BA	254	U	C2'-C1'	-6.24	1.46	1.53
34	BA	294	C	O3'-P	-6.24	1.53	1.61
34	BA	752	A	O4'-C1'	-6.24	1.33	1.41
34	BA	1240	G	C3'-C2'	-6.24	1.45	1.52
35	BB	377	A	N9-C4	-6.24	1.34	1.37
35	BB	533	U	C2'-C1'	-6.24	1.46	1.53
35	BB	551	C	C3'-C2'	-6.24	1.45	1.52
35	BB	629	C	O3'-P	-6.24	1.53	1.61
35	BB	635	A	C2'-C1'	-6.24	1.46	1.53
35	BB	677	U	C4'-C3'	-6.24	1.46	1.53
36	BC	112	G	N9-C4	-6.24	1.32	1.38
38	BE	50	G	N7-C5	-6.24	1.35	1.39
85	AA	860	C	N1-C2	6.24	1.46	1.40
85	AA	1221	G	C5-C4	-6.24	1.33	1.38
85	AA	1576	G	C2'-C1'	-6.24	1.46	1.53
85	AA	2016	A	C1'-N9	-6.24	1.38	1.46
85	AA	2114	U	C2'-C1'	-6.24	1.46	1.53
34	BA	221	G	C2-N2	-6.24	1.28	1.34
34	BA	1161	G	C5'-C4'	6.24	1.58	1.51
34	BA	1226	G	C6-N1	-6.24	1.35	1.39
34	BA	1708	A	P-O5'	-6.24	1.53	1.59
35	BB	490	G	C2-N2	-6.24	1.28	1.34
35	BB	976	U	N1-C6	-6.24	1.32	1.38
35	BB	1286	G	C3'-C2'	-6.24	1.45	1.52
37	BD	3	G	O4'-C1'	-6.24	1.33	1.41
34	BA	741	A	C4'-C3'	-6.24	1.46	1.53
34	BA	779	U	C2-N3	6.24	1.42	1.37
34	BA	1221	A	N9-C4	-6.24	1.34	1.37
34	BA	1427	U	C2'-C1'	-6.24	1.46	1.53
34	BA	1572	G	C1'-N9	-6.24	1.38	1.46
35	BB	659	C	C2-N3	-6.24	1.30	1.35
35	BB	1496	C	C2'-C1'	-6.24	1.46	1.53
35	BB	1502	U	O3'-P	-6.24	1.53	1.61
36	BC	102	G	N9-C8	-6.24	1.33	1.37
37	BD	40	C	O3'-P	-6.24	1.53	1.61
38	BE	91	G	O3'-P	-6.24	1.53	1.61
40	BG	63	U	C4'-C3'	-6.24	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	625	G	N7-C5	-6.24	1.35	1.39
34	BA	54	A	C2'-C1'	-6.24	1.46	1.53
34	BA	162	G	C1'-N9	-6.24	1.38	1.46
34	BA	210	G	C4'-C3'	6.24	1.60	1.53
34	BA	535	G	C2-N3	6.24	1.37	1.32
34	BA	1232	C	O3'-P	-6.24	1.53	1.61
34	BA	1720	U	O4'-C1'	-6.24	1.33	1.41
35	BB	77	A	O4'-C1'	-6.24	1.33	1.41
35	BB	1206	G	C1'-N9	-6.24	1.38	1.46
36	BC	104	A	C1'-N9	-6.24	1.38	1.46
38	BE	138	U	C2-N3	-6.24	1.33	1.37
41	BH	113	G	C2-N3	-6.24	1.27	1.32
85	AA	410	A	P-O5'	-6.24	1.53	1.59
85	AA	1109	G	C3'-C2'	-6.24	1.45	1.52
85	AA	1116	G	C1'-N9	-6.24	1.38	1.46
85	AA	1698	A	C1'-N9	-6.24	1.38	1.46
34	BA	334	G	C1'-N9	-6.24	1.38	1.46
34	BA	1419	A	N9-C4	-6.24	1.34	1.37
36	BC	118	U	C2-N3	-6.24	1.33	1.37
39	BF	55	A	P-O5'	-6.24	1.53	1.59
85	AA	101	C	O3'-P	-6.24	1.53	1.61
85	AA	107	A	P-O5'	-6.24	1.53	1.59
85	AA	340	G	C2'-C1'	-6.24	1.46	1.53
85	AA	1130	G	C1'-N9	-6.24	1.38	1.46
34	BA	27	G	P-O5'	-6.24	1.53	1.59
34	BA	39	C	N1-C6	-6.24	1.33	1.37
34	BA	525	A	N7-C5	-6.24	1.35	1.39
34	BA	629	G	N9-C8	-6.24	1.33	1.37
34	BA	1155	U	N1-C2	-6.24	1.32	1.38
35	BB	376	A	P-O5'	-6.24	1.53	1.59
35	BB	970	C	C3'-C2'	-6.24	1.45	1.52
35	BB	1394	A	C2'-C1'	-6.24	1.46	1.53
36	BC	40	A	N3-C4	-6.24	1.31	1.34
85	AA	129	U	O3'-P	-6.24	1.53	1.61
85	AA	276	C	C1'-N1	-6.24	1.38	1.46
85	AA	747	U	O3'-P	-6.24	1.53	1.61
85	AA	1670	U	N3-C4	-6.24	1.32	1.38
34	BA	439	A	C2'-C1'	-6.23	1.46	1.53
34	BA	458	G	C5'-C4'	-6.23	1.43	1.51
34	BA	887	U	O3'-P	-6.23	1.53	1.61
34	BA	890	G	C3'-C2'	-6.23	1.45	1.52
35	BB	86	A	C2'-C1'	-6.23	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1074	U	C2-N3	-6.23	1.33	1.37
85	AA	717	G	N1-C2	-6.23	1.32	1.37
34	BA	915	A	C1'-N9	-6.23	1.38	1.46
34	BA	1046	G	C5-C4	-6.23	1.33	1.38
34	BA	1135	U	C2-N3	-6.23	1.33	1.37
34	BA	1671	A	N7-C5	-6.23	1.35	1.39
35	BB	1067	G	O3'-P	-6.23	1.53	1.61
35	BB	1272	G	C5-C4	-6.23	1.33	1.38
40	BG	102	G	C2-N2	-6.23	1.28	1.34
85	AA	28	A	C5-C4	-6.23	1.34	1.38
85	AA	1703	A	C3'-C2'	-6.23	1.45	1.52
85	AA	1889	U	O3'-P	-6.23	1.53	1.61
85	AA	1948	A	C2'-C1'	-6.23	1.46	1.53
34	BA	729	C	C2-N3	-6.23	1.30	1.35
35	BB	608	A	C5-C4	-6.23	1.34	1.38
35	BB	1372	G	C5-C4	-6.23	1.33	1.38
36	BC	2	A	C5-C4	-6.23	1.34	1.38
39	BF	14	C	C3'-C2'	-6.23	1.45	1.52
40	BG	121	C	O3'-P	-6.23	1.53	1.61
41	BH	43	G	C2-N2	-6.23	1.28	1.34
85	AA	965	G	N9-C4	-6.23	1.32	1.38
85	AA	1293	U	C2-N3	-6.23	1.33	1.37
85	AA	1495	G	C5'-C4'	-6.23	1.43	1.51
85	AA	2192	A	C3'-C2'	-6.23	1.45	1.52
35	BB	1035	C	C3'-C2'	-6.23	1.45	1.52
35	BB	1487	G	N1-C2	-6.23	1.32	1.37
34	BA	9	A	C3'-C2'	-6.23	1.45	1.52
34	BA	373	G	P-O5'	-6.23	1.53	1.59
34	BA	484	A	C5-C4	-6.23	1.34	1.38
34	BA	744	G	C6-N1	-6.23	1.35	1.39
34	BA	946	A	C4'-C3'	-6.23	1.46	1.53
34	BA	1820	G	C2'-C1'	-6.23	1.46	1.53
35	BB	390	G	N1-C2	-6.23	1.32	1.37
35	BB	815	G	O3'-P	-6.23	1.53	1.61
35	BB	1102	U	O4'-C1'	-6.23	1.33	1.41
35	BB	1220	A	C1'-N9	-6.23	1.38	1.46
35	BB	1273	G	O3'-P	-6.23	1.53	1.61
38	BE	34	C	N1-C6	-6.23	1.33	1.37
40	BG	145	C	P-O5'	-6.23	1.53	1.59
85	AA	1221	G	C4'-C3'	-6.23	1.46	1.53
85	AA	1734	A	C1'-N9	-6.23	1.38	1.46
85	AA	2172	A	C2'-C1'	-6.23	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	523	A	P-O5'	-6.23	1.53	1.59
34	BA	864	G	C2-N3	-6.23	1.27	1.32
35	BB	101	U	C4'-C3'	-6.23	1.46	1.53
35	BB	1265	U	C2-N3	-6.23	1.33	1.37
35	BB	1314	G	C4'-C3'	-6.23	1.46	1.53
34	BA	43	U	O4'-C1'	-6.22	1.33	1.41
34	BA	220	U	C2'-C1'	-6.22	1.46	1.53
34	BA	726	G	C2-N2	-6.22	1.28	1.34
34	BA	1093	G	C5-C4	-6.22	1.33	1.38
34	BA	1098	G	C6-N1	-6.22	1.35	1.39
35	BB	570	A	C2'-C1'	-6.22	1.46	1.53
35	BB	622	G	C5-C4	-6.22	1.33	1.38
35	BB	634	A	C1'-N9	-6.22	1.38	1.46
35	BB	874	G	C3'-C2'	-6.22	1.46	1.52
85	AA	350	U	C2'-C1'	-6.22	1.46	1.53
85	AA	886	A	C3'-C2'	-6.22	1.46	1.52
85	AA	1923	A	O3'-P	-6.22	1.53	1.61
34	BA	1287	G	C5-C4	-6.22	1.33	1.38
34	BA	1468	U	C2-N3	-6.22	1.33	1.37
34	BA	1706	A	C6-N6	-6.22	1.28	1.33
35	BB	493	U	C2-N3	-6.22	1.33	1.37
35	BB	1196	A	N7-C5	-6.22	1.35	1.39
36	BC	61	A	N7-C5	-6.22	1.35	1.39
38	BE	197	A	N7-C5	-6.22	1.35	1.39
41	BH	15	A	C4'-C3'	-6.22	1.46	1.53
41	BH	55	C	C2'-C1'	-6.22	1.46	1.53
67	Bh	142	GLY	CA-C	-6.22	1.41	1.51
85	AA	397	G	N1-C2	-6.22	1.32	1.37
85	AA	664	C	P-O5'	-6.22	1.53	1.59
85	AA	924	A	C3'-C2'	-6.22	1.46	1.52
85	AA	1190	G	C1'-N9	-6.22	1.38	1.46
85	AA	1275	A	C6-N1	-6.22	1.31	1.35
85	AA	1477	A	C2'-C1'	-6.22	1.46	1.53
85	AA	1634	U	C2-N3	-6.22	1.33	1.37
85	AA	1669	G	C1'-N9	-6.22	1.38	1.46
85	AA	2141	G	C4'-O4'	-6.22	1.37	1.45
34	BA	399	G	N7-C5	-6.22	1.35	1.39
34	BA	1653	G	P-O5'	-6.22	1.53	1.59
40	BG	174	G	C5-C6	-6.22	1.36	1.42
85	AA	910	G	N9-C4	-6.22	1.32	1.38
85	AA	1679	U	C3'-C2'	-6.22	1.46	1.52
34	BA	18	G	N9-C8	-6.22	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	362	G	O3'-P	-6.22	1.53	1.61
34	BA	456	G	C3'-C2'	-6.22	1.46	1.52
34	BA	1069	U	O3'-P	-6.22	1.53	1.61
34	BA	1073	G	C2'-C1'	-6.22	1.46	1.53
85	AA	1	G	N3-C4	-6.22	1.31	1.35
85	AA	585	G	C2-N2	-6.22	1.28	1.34
34	BA	198	U	N1-C2	-6.22	1.32	1.38
34	BA	1195	G	O4'-C1'	-6.22	1.33	1.41
34	BA	1545	C	O3'-P	-6.22	1.53	1.61
35	BB	850	U	O4'-C1'	-6.22	1.33	1.41
34	BA	267	G	C3'-C2'	-6.22	1.46	1.52
34	BA	431	A	C1'-N9	-6.22	1.38	1.46
35	BB	450	A	C2'-C1'	-6.22	1.46	1.53
35	BB	734	A	O3'-P	-6.22	1.53	1.61
35	BB	1144	A	C6-N6	-6.22	1.28	1.33
35	BB	1153	G	N1-C2	-6.22	1.32	1.37
37	BD	104	C	C3'-C2'	-6.22	1.46	1.52
85	AA	419	A	C3'-C2'	-6.22	1.46	1.52
85	AA	555	C	O3'-P	-6.22	1.53	1.61
85	AA	2032	G	O3'-P	-6.22	1.53	1.61
34	BA	108	A	C8-N7	-6.21	1.27	1.31
34	BA	785	G	C2'-C1'	-6.21	1.46	1.53
35	BB	1133	C	C4'-C3'	-6.21	1.46	1.53
35	BB	1152	U	C2-N3	-6.21	1.33	1.37
35	BB	1414	A	N9-C8	-6.21	1.32	1.37
36	BC	55	U	N3-C4	-6.21	1.32	1.38
36	BC	104	A	N9-C4	-6.21	1.34	1.37
40	BG	173	C	N1-C6	-6.21	1.33	1.37
85	AA	1272	G	N3-C4	-6.21	1.31	1.35
34	BA	774	A	C5-C4	-6.21	1.34	1.38
34	BA	1140	A	O3'-P	-6.21	1.53	1.61
35	BB	112	G	N3-C4	-6.21	1.31	1.35
35	BB	813	C	C2-N3	-6.21	1.30	1.35
35	BB	1137	G	C2-N2	-6.21	1.28	1.34
85	AA	1668	G	N9-C4	-6.21	1.32	1.38
85	AA	1799	C	C2'-C1'	-6.21	1.46	1.53
34	BA	191	G	C1'-N9	-6.21	1.38	1.46
34	BA	239	C	P-O5'	-6.21	1.53	1.59
34	BA	1109	G	N1-C2	-6.21	1.32	1.37
34	BA	1197	U	C2-N3	-6.21	1.33	1.37
34	BA	1299	G	C3'-C2'	-6.21	1.46	1.52
34	BA	1515	U	C2'-C1'	-6.21	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	483	C	C1'-N1	-6.21	1.38	1.46
35	BB	628	A	N7-C5	-6.21	1.35	1.39
35	BB	1453	G	N9-C4	-6.21	1.32	1.38
38	BE	7	U	C2-N3	-6.21	1.33	1.37
38	BE	163	A	O3'-P	-6.21	1.53	1.61
40	BG	8	U	O3'-P	-6.21	1.53	1.61
85	AA	254	G	C2'-C1'	-6.21	1.46	1.53
85	AA	1170	C	O3'-P	-6.21	1.53	1.61
85	AA	1674	G	C1'-N9	-6.21	1.38	1.46
85	AA	2138	G	N9-C8	-6.21	1.33	1.37
34	BA	584	A	O3'-P	-6.21	1.53	1.61
36	BC	163	A	C1'-N9	-6.21	1.38	1.46
37	BD	61	C	C2-N3	-6.21	1.30	1.35
38	BE	35	A	N9-C4	-6.21	1.34	1.37
34	BA	273	G	P-O5'	-6.21	1.53	1.59
34	BA	324	C	O4'-C1'	-6.21	1.33	1.41
34	BA	566	G	C1'-N9	-6.21	1.38	1.46
34	BA	727	G	C1'-N9	-6.21	1.38	1.46
34	BA	1143	U	N3-C4	-6.21	1.32	1.38
35	BB	792	G	C2'-C1'	-6.21	1.46	1.53
35	BB	1104	A	C4'-O4'	-6.21	1.37	1.45
36	BC	90	U	P-O5'	-6.21	1.53	1.59
85	AA	24	U	C1'-N1	-6.21	1.38	1.46
85	AA	253	C	O3'-P	-6.21	1.53	1.61
34	BA	406	G	N3-C4	-6.21	1.31	1.35
35	BB	443	A	C5-C4	-6.21	1.34	1.38
35	BB	702	G	C2-N2	-6.21	1.28	1.34
35	BB	1134	G	C5-C6	-6.21	1.36	1.42
35	BB	1369	A	N3-C4	-6.21	1.31	1.34
35	BB	1445	A	N3-C4	-6.21	1.31	1.34
35	BB	1480	G	O3'-P	-6.21	1.53	1.61
38	BE	65	U	C4'-C3'	-6.21	1.46	1.53
39	BF	36	G	C3'-C2'	-6.21	1.46	1.52
85	AA	1102	C	P-O5'	-6.21	1.53	1.59
34	BA	418	G	N7-C5	-6.21	1.35	1.39
34	BA	897	U	C3'-C2'	-6.21	1.46	1.52
34	BA	1054	U	O4'-C1'	-6.21	1.33	1.41
34	BA	1546	C	C2-N3	-6.21	1.30	1.35
34	BA	1557	G	C3'-C2'	-6.21	1.46	1.52
35	BB	642	G	N7-C5	-6.21	1.35	1.39
23	AP	50	GLY	CA-C	-6.20	1.42	1.51
34	BA	127	U	O4'-C1'	-6.20	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	395	G	C4'-O4'	-6.20	1.37	1.45
34	BA	1035	A	O3'-P	-6.20	1.53	1.61
34	BA	1282	G	C5-C4	-6.20	1.34	1.38
35	BB	102	G	C6-N1	-6.20	1.35	1.39
35	BB	789	G	C2'-C1'	-6.20	1.46	1.53
35	BB	829	C	C3'-C2'	-6.20	1.46	1.52
35	BB	1200	A	N9-C4	6.20	1.41	1.37
39	BF	9	C	C2-N3	-6.20	1.30	1.35
40	BG	169	A	N9-C8	-6.20	1.32	1.37
85	AA	863	C	C2'-C1'	-6.20	1.46	1.53
85	AA	1036	A	O3'-P	-6.20	1.53	1.61
85	AA	1656	C	P-O5'	-6.20	1.53	1.59
85	AA	1823	G	C2'-C1'	-6.20	1.46	1.53
85	AA	1915	C	O3'-P	-6.20	1.53	1.61
85	AA	2105	G	C2-N2	-6.20	1.28	1.34
85	AA	2200	A	C2'-C1'	-6.20	1.46	1.53
85	AA	2203	C	C2-N3	-6.20	1.30	1.35
86	AB	14	A	C4'-C3'	6.20	1.59	1.53
34	BA	329	G	C6-N1	-6.20	1.35	1.39
34	BA	1256	A	P-O5'	-6.20	1.53	1.59
35	BB	43	G	C6-N1	-6.20	1.35	1.39
35	BB	652	G	P-O5'	-6.20	1.53	1.59
35	BB	1036	G	N9-C4	-6.20	1.32	1.38
35	BB	1063	C	C4'-C3'	-6.20	1.46	1.53
37	BD	69	U	O3'-P	-6.20	1.53	1.61
85	AA	1290	G	C2'-C1'	-6.20	1.46	1.53
85	AA	1583	U	O3'-P	-6.20	1.53	1.61
34	BA	422	C	C4'-C3'	-6.20	1.46	1.53
34	BA	1302	C	O3'-P	-6.20	1.53	1.61
35	BB	132	G	N9-C8	-6.20	1.33	1.37
35	BB	1070	G	C2-N2	-6.20	1.28	1.34
36	BC	57	C	C2-N3	-6.20	1.30	1.35
40	BG	105	A	C2'-C1'	-6.20	1.46	1.53
85	AA	377	U	N3-C4	-6.20	1.32	1.38
85	AA	637	U	C2-N3	-6.20	1.33	1.37
85	AA	1692	U	N3-C4	-6.20	1.32	1.38
85	AA	2226	U	O3'-P	-6.20	1.53	1.61
34	BA	41	U	C2'-C1'	-6.20	1.46	1.53
34	BA	98	A	P-O5'	-6.20	1.53	1.59
34	BA	358	A	C5-C4	-6.20	1.34	1.38
34	BA	383	G	C2-N2	-6.20	1.28	1.34
34	BA	418	G	C5-C4	-6.20	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	456	G	N7-C5	-6.20	1.35	1.39
34	BA	674	G	C2'-C1'	-6.20	1.46	1.53
34	BA	751	A	C8-N7	-6.20	1.27	1.31
34	BA	1428	G	N7-C5	-6.20	1.35	1.39
36	BC	119	G	C2'-C1'	-6.20	1.46	1.53
36	BC	133	C	C2'-C1'	-6.20	1.46	1.53
85	AA	10	G	C6-N1	-6.20	1.35	1.39
85	AA	451	G	P-O5'	-6.20	1.53	1.59
85	AA	537	G	O3'-P	-6.20	1.53	1.61
85	AA	877	G	C6-N1	-6.20	1.35	1.39
85	AA	1211	C	C1'-N1	-6.20	1.38	1.46
85	AA	1494	C	O3'-P	-6.20	1.53	1.61
85	AA	2220	U	P-O5'	-6.20	1.53	1.59
40	BG	179	C	P-O5'	-6.20	1.53	1.59
85	AA	1218	C	N1-C6	-6.20	1.33	1.37
85	AA	1462	A	N7-C5	-6.20	1.35	1.39
34	BA	151	A	N9-C4	6.20	1.41	1.37
34	BA	453	A	C3'-C2'	-6.20	1.46	1.52
35	BB	107	A	C3'-C2'	-6.20	1.46	1.52
35	BB	1347	C	O3'-P	-6.20	1.53	1.61
36	BC	21	U	C2-N3	-6.20	1.33	1.37
85	AA	492	C	C2'-C1'	-6.20	1.46	1.53
85	AA	506	G	C1'-N9	-6.20	1.38	1.46
85	AA	709	A	C4'-C3'	-6.20	1.46	1.53
85	AA	1110	A	C5-C4	-6.20	1.34	1.38
85	AA	1164	A	C3'-C2'	-6.20	1.46	1.52
85	AA	1203	G	P-O5'	-6.20	1.53	1.59
85	AA	2227	A	C1'-N9	-6.20	1.38	1.46
34	BA	510	U	O3'-P	-6.19	1.53	1.61
34	BA	764	G	C6-N1	-6.19	1.35	1.39
35	BB	114	A	C2'-C1'	-6.19	1.46	1.53
40	BG	21	C	C5'-C4'	6.19	1.58	1.51
85	AA	1476	C	C1'-N1	-6.19	1.38	1.46
85	AA	1492	U	C2'-C1'	-6.19	1.46	1.53
85	AA	2095	U	C2-N3	-6.19	1.33	1.37
34	BA	580	U	N1-C2	-6.19	1.32	1.38
34	BA	650	C	O3'-P	-6.19	1.53	1.61
34	BA	960	C	O3'-P	-6.19	1.53	1.61
34	BA	1261	G	C3'-C2'	-6.19	1.46	1.52
35	BB	75	A	C1'-N9	-6.19	1.38	1.46
35	BB	670	G	N9-C4	-6.19	1.32	1.38
40	BG	74	G	O3'-P	-6.19	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	123	C	C2-N3	-6.19	1.30	1.35
85	AA	1451	U	C2-N3	-6.19	1.33	1.37
85	AA	1500	C	P-O5'	-6.19	1.53	1.59
85	AA	1707	G	C2'-C1'	-6.19	1.46	1.53
34	BA	1147	C	C2'-C1'	-6.19	1.46	1.53
35	BB	402	G	C2-N2	-6.19	1.28	1.34
35	BB	546	A	O4'-C1'	-6.19	1.33	1.41
35	BB	1136	G	N7-C5	-6.19	1.35	1.39
35	BB	1213	U	O3'-P	-6.19	1.53	1.61
35	BB	1289	G	O3'-P	-6.19	1.53	1.61
36	BC	34	U	C4'-O4'	-6.19	1.37	1.45
38	BE	129	G	N1-C2	-6.19	1.32	1.37
40	BG	156	G	P-O5'	-6.19	1.53	1.59
85	AA	256	A	N9-C4	-6.19	1.34	1.37
85	AA	411	U	N3-C4	-6.19	1.32	1.38
85	AA	1659	C	O3'-P	-6.19	1.53	1.61
85	AA	1987	G	P-O5'	-6.19	1.53	1.59
35	BB	445	G	C2'-C1'	-6.19	1.46	1.53
35	BB	470	C	C3'-C2'	-6.19	1.46	1.52
35	BB	806	U	C4-O4	-6.19	1.18	1.23
36	BC	90	U	C3'-C2'	-6.19	1.46	1.52
36	BC	109	A	C1'-N9	-6.19	1.38	1.46
40	BG	74	G	C5-C4	-6.19	1.34	1.38
41	BH	36	C	C4'-O4'	-6.19	1.37	1.45
85	AA	405	C	P-O5'	-6.19	1.53	1.59
85	AA	1147	A	N7-C5	-6.19	1.35	1.39
34	BA	89	G	O4'-C1'	-6.19	1.33	1.41
34	BA	1409	A	C8-N7	-6.19	1.27	1.31
36	BC	152	C	C4'-O4'	-6.19	1.37	1.45
37	BD	13	A	O3'-P	-6.19	1.53	1.61
38	BE	32	U	N1-C2	-6.19	1.32	1.38
40	BG	35	G	C3'-C2'	-6.19	1.46	1.52
40	BG	73	U	N3-C4	-6.19	1.32	1.38
40	BG	112	C	C4'-C3'	-6.19	1.46	1.53
40	BG	132	U	C2'-C1'	-6.19	1.46	1.53
85	AA	181	A	P-O5'	-6.19	1.53	1.59
85	AA	775	C	C2-N3	-6.19	1.30	1.35
34	BA	32	A	P-O5'	-6.19	1.53	1.59
34	BA	419	U	O3'-P	-6.19	1.53	1.61
34	BA	689	C	O3'-P	-6.19	1.53	1.61
34	BA	1209	A	O3'-P	-6.19	1.53	1.61
34	BA	1590	G	C2-N2	-6.19	1.28	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1735	G	C5'-C4'	-6.19	1.44	1.51
34	BA	1842	U	O3'-P	-6.19	1.53	1.61
35	BB	73	G	C6-N1	-6.19	1.35	1.39
35	BB	421	U	C2'-C1'	-6.19	1.46	1.53
35	BB	575	C	O3'-P	-6.19	1.53	1.61
85	AA	42	G	C2-N2	-6.19	1.28	1.34
85	AA	392	G	C2-N2	-6.19	1.28	1.34
85	AA	464	A	C8-N7	-6.19	1.27	1.31
85	AA	2141	G	C5-C6	-6.19	1.36	1.42
34	BA	271	C	C5'-C4'	6.18	1.58	1.51
34	BA	697	A	C2'-C1'	-6.18	1.46	1.53
34	BA	730	C	C3'-C2'	-6.18	1.46	1.52
35	BB	479	U	C2-N3	-6.18	1.33	1.37
35	BB	504	C	N1-C2	-6.18	1.33	1.40
35	BB	544	C	C2-N3	-6.18	1.30	1.35
35	BB	788	U	C1'-N1	-6.18	1.38	1.46
35	BB	1159	U	C3'-C2'	-6.18	1.46	1.52
38	BE	34	C	C3'-C2'	-6.18	1.46	1.52
34	BA	72	U	C2-N3	-6.18	1.33	1.37
34	BA	352	G	C2'-C1'	-6.18	1.46	1.53
34	BA	482	C	C5'-C4'	-6.18	1.44	1.51
34	BA	484	A	C3'-C2'	-6.18	1.46	1.52
34	BA	749	G	N9-C4	-6.18	1.33	1.38
34	BA	1147	C	C3'-C2'	-6.18	1.46	1.52
34	BA	1230	G	C5-C4	-6.18	1.34	1.38
34	BA	1452	U	C4'-C3'	-6.18	1.46	1.53
34	BA	1671	A	C3'-C2'	-6.18	1.46	1.52
35	BB	456	A	C4'-C3'	-6.18	1.46	1.53
35	BB	1035	C	O3'-P	-6.18	1.53	1.61
40	BG	21	C	C4'-O4'	6.18	1.53	1.45
85	AA	15	U	C2-N3	-6.18	1.33	1.37
85	AA	325	C	O3'-P	-6.18	1.53	1.61
85	AA	943	U	N1-C6	-6.18	1.32	1.38
85	AA	1540	A	N7-C5	-6.18	1.35	1.39
35	BB	1296	A	C3'-C2'	-6.18	1.46	1.52
85	AA	1114	A	C1'-N9	-6.18	1.38	1.46
34	BA	53	G	N9-C4	-6.18	1.33	1.38
34	BA	467	A	C5-C4	-6.18	1.34	1.38
34	BA	1226	G	C4'-C3'	-6.18	1.46	1.53
34	BA	1652	G	N7-C5	-6.18	1.35	1.39
35	BB	501	G	N9-C8	-6.18	1.33	1.37
36	BC	59	A	C2'-C1'	-6.18	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	66	G	C2-N2	-6.18	1.28	1.34
37	BD	63	C	P-O5'	-6.18	1.53	1.59
38	BE	35	A	C4'-C3'	-6.18	1.46	1.53
85	AA	992	G	C3'-C2'	-6.18	1.46	1.52
85	AA	1298	G	N7-C5	-6.18	1.35	1.39
35	BB	670	G	C2'-C1'	-6.18	1.46	1.53
35	BB	877	A	C3'-C2'	-6.18	1.46	1.52
35	BB	1174	C	C2-N3	-6.18	1.30	1.35
85	AA	2054	G	N9-C8	-6.18	1.33	1.37
85	AA	2132	A	C1'-N9	-6.18	1.38	1.46
34	BA	124	G	C1'-N9	-6.18	1.38	1.46
34	BA	237	A	P-O5'	-6.18	1.53	1.59
34	BA	247	U	C3'-O3'	6.18	1.50	1.42
34	BA	324	C	C4'-O4'	-6.18	1.37	1.45
34	BA	621	G	C3'-C2'	-6.18	1.46	1.52
34	BA	719	G	C6-N1	-6.18	1.35	1.39
34	BA	885	A	C2'-C1'	-6.18	1.46	1.53
34	BA	1564	A	C5-C4	-6.18	1.34	1.38
34	BA	1679	C	N3-C4	-6.18	1.29	1.33
35	BB	507	G	C2'-C1'	-6.18	1.46	1.53
35	BB	617	C	C4-N4	-6.18	1.28	1.33
35	BB	949	G	O3'-P	-6.18	1.53	1.61
41	BH	68	G	P-O5'	-6.18	1.53	1.59
85	AA	501	A	N9-C4	-6.18	1.34	1.37
85	AA	526	G	N7-C5	-6.18	1.35	1.39
85	AA	900	G	N9-C4	-6.18	1.33	1.38
85	AA	1247	A	C5'-C4'	-6.18	1.44	1.51
85	AA	1283	C	C1'-N1	-6.18	1.38	1.46
85	AA	2005	U	O3'-P	-6.18	1.53	1.61
85	AA	2112	G	C3'-C2'	-6.18	1.46	1.52
34	BA	83	G	C8-N7	-6.17	1.27	1.30
34	BA	371	U	O3'-P	-6.17	1.53	1.61
34	BA	1668	C	C4'-C3'	-6.17	1.46	1.53
35	BB	122	U	C2'-C1'	-6.17	1.46	1.53
35	BB	788	U	O3'-P	-6.17	1.53	1.61
35	BB	1235	A	N7-C5	-6.17	1.35	1.39
35	BB	1267	C	C1'-N1	-6.17	1.38	1.46
36	BC	26	U	O4'-C1'	-6.17	1.33	1.41
37	BD	110	G	N9-C8	-6.17	1.33	1.37
38	BE	124	G	C4'-C3'	-6.17	1.46	1.53
40	BG	83	U	P-O5'	-6.17	1.53	1.59
40	BG	124	A	C1'-N9	-6.17	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	142	A	N3-C4	-6.17	1.31	1.34
85	AA	855	G	O3'-P	-6.17	1.53	1.61
85	AA	2142	A	O3'-P	-6.17	1.53	1.61
34	BA	586	G	C2'-C1'	-6.17	1.46	1.53
34	BA	1048	C	C2'-C1'	-6.17	1.46	1.53
35	BB	415	A	C8-N7	-6.17	1.27	1.31
35	BB	660	G	N7-C5	-6.17	1.35	1.39
35	BB	795	A	C5-C4	-6.17	1.34	1.38
35	BB	1111	C	C2-N3	-6.17	1.30	1.35
35	BB	1145	G	C2'-C1'	-6.17	1.46	1.53
37	BD	80	G	C2'-C1'	-6.17	1.46	1.53
85	AA	104	C	C3'-C2'	-6.17	1.46	1.52
34	BA	352	G	C6-N1	-6.17	1.35	1.39
34	BA	1197	U	O4'-C1'	-6.17	1.33	1.41
34	BA	1431	G	N7-C5	-6.17	1.35	1.39
35	BB	1132	A	C3'-C2'	-6.17	1.46	1.52
35	BB	1260	A	C5-C4	-6.17	1.34	1.38
35	BB	1470	G	C5-C6	-6.17	1.36	1.42
40	BG	52	A	C2'-C1'	-6.17	1.46	1.53
85	AA	518	A	O3'-P	-6.17	1.53	1.61
85	AA	759	G	C1'-N9	-6.17	1.38	1.46
85	AA	1143	C	C4'-O4'	-6.17	1.37	1.45
85	AA	1265	C	C3'-C2'	-6.17	1.46	1.52
85	AA	1295	G	N7-C5	-6.17	1.35	1.39
85	AA	1598	A	P-O5'	-6.17	1.53	1.59
34	BA	1105	A	C3'-O3'	6.17	1.50	1.42
34	BA	1195	G	N9-C8	-6.17	1.33	1.37
34	BA	1505	G	C2-N2	-6.17	1.28	1.34
34	BA	1529	G	N3-C4	-6.17	1.31	1.35
34	BA	1648	G	C2-N2	-6.17	1.28	1.34
35	BB	52	G	N9-C8	-6.17	1.33	1.37
85	AA	2178	A	N9-C4	-6.17	1.34	1.37
34	BA	50	G	C2-N2	-6.17	1.28	1.34
34	BA	241	U	C3'-C2'	-6.17	1.46	1.52
34	BA	1489	U	N1-C2	-6.17	1.32	1.38
35	BB	1459	U	O3'-P	-6.17	1.53	1.61
37	BD	87	G	N9-C8	-6.17	1.33	1.37
38	BE	34	C	C4'-C3'	-6.17	1.46	1.53
38	BE	45	G	O3'-P	-6.17	1.53	1.61
41	BH	43	G	C4'-O4'	-6.17	1.37	1.45
85	AA	256	A	O3'-P	-6.17	1.53	1.61
85	AA	286	C	C2'-C1'	-6.17	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2007	G	C2-N3	-6.17	1.27	1.32
85	AA	2137	A	O3'-P	-6.17	1.53	1.61
34	BA	1441	C	N3-C4	-6.17	1.29	1.33
38	BE	174	U	O3'-P	-6.17	1.53	1.61
85	AA	258	G	C2'-C1'	-6.17	1.46	1.53
85	AA	369	A	C1'-N9	-6.17	1.38	1.46
85	AA	717	G	C6-N1	-6.17	1.35	1.39
85	AA	2062	U	P-O5'	-6.17	1.53	1.59
86	AB	45	U	O3'-P	-6.17	1.53	1.61
34	BA	29	U	N3-C4	-6.17	1.32	1.38
34	BA	32	A	C3'-C2'	-6.17	1.46	1.52
34	BA	687	G	N1-C2	-6.17	1.32	1.37
34	BA	1564	A	C8-N7	-6.17	1.27	1.31
40	BG	154	C	P-O5'	-6.17	1.53	1.59
85	AA	449	G	C3'-C2'	-6.17	1.46	1.52
85	AA	1687	U	O3'-P	-6.17	1.53	1.61
34	BA	368	U	O3'-P	-6.16	1.53	1.61
34	BA	1254	C	C4'-C3'	-6.16	1.46	1.53
34	BA	1533	G	C5-C4	-6.16	1.34	1.38
35	BB	53	C	O3'-P	-6.16	1.53	1.61
35	BB	1089	A	C2'-C1'	-6.16	1.46	1.53
35	BB	1452	U	C2-N3	-6.16	1.33	1.37
36	BC	164	G	C1'-N9	-6.16	1.38	1.46
40	BG	119	A	O4'-C1'	-6.16	1.33	1.41
85	AA	279	C	P-O5'	-6.16	1.53	1.59
85	AA	316	C	O3'-P	-6.16	1.53	1.61
85	AA	718	C	O3'-P	-6.16	1.53	1.61
85	AA	1465	C	C2-N3	-6.16	1.30	1.35
85	AA	2152	C	C2'-C1'	-6.16	1.46	1.53
34	BA	141	G	N7-C5	-6.16	1.35	1.39
34	BA	602	G	C2-N3	-6.16	1.27	1.32
34	BA	909	G	C4'-O4'	-6.16	1.37	1.45
34	BA	931	G	N1-C2	-6.16	1.32	1.37
34	BA	934	G	C2'-C1'	-6.16	1.46	1.53
34	BA	1530	G	N9-C4	-6.16	1.33	1.38
35	BB	576	A	P-O5'	-6.16	1.53	1.59
35	BB	985	A	O3'-P	-6.16	1.53	1.61
39	BF	58	U	C2-N3	-6.16	1.33	1.37
34	BA	79	C	C1'-N1	-6.16	1.38	1.46
34	BA	248	G	C4'-C3'	6.16	1.59	1.53
34	BA	494	A	C5-C4	-6.16	1.34	1.38
34	BA	1218	G	C2'-C1'	-6.16	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1403	G	P-O5'	-6.16	1.53	1.59
35	BB	1173	C	O3'-P	-6.16	1.53	1.61
35	BB	1188	A	O3'-P	-6.16	1.53	1.61
35	BB	1355	C	P-O5'	-6.16	1.53	1.59
36	BC	136	G	N1-C2	-6.16	1.32	1.37
38	BE	30	C	C4-N4	-6.16	1.28	1.33
85	AA	516	G	C5-C4	-6.16	1.34	1.38
85	AA	1134	G	O3'-P	-6.16	1.53	1.61
34	BA	396	U	N1-C2	-6.16	1.33	1.38
34	BA	1241	U	N1-C2	-6.16	1.33	1.38
35	BB	518	G	C1'-N9	-6.16	1.38	1.46
35	BB	1026	G	C3'-O3'	6.16	1.50	1.42
35	BB	1032	U	N1-C2	-6.16	1.33	1.38
37	BD	71	G	C5-C4	-6.16	1.34	1.38
37	BD	87	G	C2-N2	-6.16	1.28	1.34
40	BG	51	U	C4'-O4'	-6.16	1.37	1.45
40	BG	102	G	N7-C5	-6.16	1.35	1.39
85	AA	696	G	C5-C4	-6.16	1.34	1.38
85	AA	815	G	C2-N3	6.16	1.37	1.32
85	AA	984	A	O3'-P	-6.16	1.53	1.61
85	AA	1558	U	O3'-P	-6.16	1.53	1.61
34	BA	447	U	C5'-C4'	-6.16	1.44	1.51
34	BA	1654	G	N1-C2	-6.16	1.32	1.37
34	BA	1703	A	C5-C4	-6.16	1.34	1.38
35	BB	135	C	C2'-C1'	-6.16	1.46	1.53
35	BB	1060	U	C4'-C3'	-6.16	1.46	1.53
40	BG	162	A	N7-C5	-6.16	1.35	1.39
85	AA	46	U	C2-N3	-6.16	1.33	1.37
85	AA	2231	G	C5'-C4'	6.16	1.58	1.51
34	BA	12	G	C5-C4	-6.16	1.34	1.38
34	BA	31	A	C5-C4	-6.16	1.34	1.38
34	BA	270	U	O3'-P	-6.16	1.53	1.61
34	BA	500	C	C2-N3	-6.16	1.30	1.35
34	BA	628	U	C2'-C1'	-6.16	1.46	1.53
34	BA	744	G	N1-C2	-6.16	1.32	1.37
34	BA	902	C	N1-C6	-6.16	1.33	1.37
34	BA	976	C	C3'-C2'	-6.16	1.46	1.52
35	BB	548	A	P-O5'	-6.16	1.53	1.59
35	BB	1052	G	C5-C4	-6.16	1.34	1.38
35	BB	1498	G	C2-N2	-6.16	1.28	1.34
36	BC	73	U	O3'-P	-6.16	1.53	1.61
41	BH	35	G	N9-C8	-6.16	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	11	A	C4'-C3'	-6.16	1.46	1.53
85	AA	484	G	C4'-C3'	-6.16	1.46	1.53
85	AA	1030	U	O3'-P	-6.16	1.53	1.61
85	AA	1141	U	P-O5'	-6.16	1.53	1.59
85	AA	1201	A	C2'-C1'	-6.16	1.46	1.53
85	AA	1279	A	N7-C5	-6.16	1.35	1.39
85	AA	1520	A	C5-C4	-6.16	1.34	1.38
85	AA	2150	G	C2-N2	-6.16	1.28	1.34
34	BA	1253	G	N3-C4	-6.15	1.31	1.35
38	BE	66	A	C3'-C2'	-6.15	1.46	1.52
40	BG	151	A	C3'-C2'	-6.15	1.46	1.52
85	AA	718	C	C3'-C2'	-6.15	1.46	1.52
85	AA	1515	A	N9-C4	-6.15	1.34	1.37
85	AA	1660	U	C2'-C1'	-6.15	1.46	1.53
34	BA	32	A	N7-C5	-6.15	1.35	1.39
34	BA	478	G	O3'-P	-6.15	1.53	1.61
34	BA	1240	G	C8-N7	-6.15	1.27	1.30
35	BB	100	A	C4'-C3'	-6.15	1.46	1.53
35	BB	541	U	C2'-C1'	-6.15	1.46	1.53
35	BB	1098	G	C2'-C1'	-6.15	1.46	1.53
35	BB	1372	G	N9-C4	-6.15	1.33	1.38
36	BC	133	C	O3'-P	-6.15	1.53	1.61
38	BE	21	C	C2-N3	-6.15	1.30	1.35
38	BE	126	G	C4'-C3'	6.15	1.59	1.53
41	BH	44	A	C5-C6	-6.15	1.35	1.41
85	AA	802	A	C5'-C4'	6.15	1.58	1.51
85	AA	1268	C	C2'-C1'	-6.15	1.46	1.53
85	AA	1518	A	C3'-C2'	-6.15	1.46	1.52
85	AA	2171	A	O3'-P	-6.15	1.53	1.61
34	BA	94	G	C2-N2	-6.15	1.28	1.34
34	BA	824	C	C2-N3	-6.15	1.30	1.35
34	BA	1368	G	P-O5'	-6.15	1.53	1.59
35	BB	1307	C	C2-N3	-6.15	1.30	1.35
36	BC	152	C	C1'-N1	-6.15	1.38	1.46
37	BD	92	G	N9-C8	-6.15	1.33	1.37
40	BG	93	U	C3'-C2'	-6.15	1.46	1.52
40	BG	125	C	P-O5'	-6.15	1.53	1.59
41	BH	64	U	C2-N3	-6.15	1.33	1.37
85	AA	256	A	C1'-N9	-6.15	1.38	1.46
85	AA	353	G	N9-C4	-6.15	1.33	1.38
85	AA	1902	C	P-O5'	-6.15	1.53	1.59
34	BA	1558	C	C2-N3	-6.15	1.30	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1084	A	N3-C4	-6.15	1.31	1.34
34	BA	293	A	N7-C5	-6.15	1.35	1.39
34	BA	624	G	O3'-P	-6.15	1.53	1.61
34	BA	949	C	C4'-C3'	-6.15	1.46	1.53
34	BA	1210	A	C5-C4	-6.15	1.34	1.38
34	BA	1333	G	C6-N1	-6.15	1.35	1.39
35	BB	572	G	O3'-P	-6.15	1.53	1.61
35	BB	1288	G	C3'-C2'	-6.15	1.46	1.52
40	BG	5	G	C4'-C3'	-6.15	1.46	1.53
40	BG	76	C	O4'-C1'	-6.15	1.33	1.41
85	AA	314	C	C4-C5	-6.15	1.38	1.43
85	AA	421	G	C2-N2	-6.15	1.28	1.34
85	AA	573	U	O3'-P	-6.15	1.53	1.61
85	AA	2179	C	C2-N3	-6.15	1.30	1.35
35	BB	75	A	C3'-C2'	-6.15	1.46	1.52
35	BB	87	G	C2'-C1'	-6.15	1.46	1.53
85	AA	452	A	N9-C4	-6.15	1.34	1.37
34	BA	340	U	C4'-C3'	-6.14	1.46	1.53
34	BA	860	G	N3-C4	-6.14	1.31	1.35
34	BA	1704	G	N9-C8	-6.14	1.33	1.37
35	BB	19	C	C3'-C2'	-6.14	1.46	1.52
35	BB	816	U	C2'-C1'	-6.14	1.46	1.53
35	BB	1108	G	C1'-N9	-6.14	1.38	1.46
35	BB	1147	G	N9-C4	-6.14	1.33	1.38
38	BE	175	U	C2-N3	-6.14	1.33	1.37
41	BH	43	G	C5-C4	-6.14	1.34	1.38
85	AA	430	G	O3'-P	-6.14	1.53	1.61
85	AA	672	U	C2-N3	-6.14	1.33	1.37
34	BA	184	C	O3'-P	-6.14	1.53	1.61
34	BA	203	U	C4'-C3'	-6.14	1.46	1.53
34	BA	302	A	P-O5'	-6.14	1.53	1.59
34	BA	655	U	P-O5'	-6.14	1.53	1.59
34	BA	982	A	N7-C5	-6.14	1.35	1.39
34	BA	1657	A	O3'-P	-6.14	1.53	1.61
34	BA	1693	U	C3'-C2'	-6.14	1.46	1.52
35	BB	126	C	C4-N4	-6.14	1.28	1.33
35	BB	412	A	P-O5'	-6.14	1.53	1.59
35	BB	436	G	C5-C6	-6.14	1.36	1.42
35	BB	553	U	C2-N3	-6.14	1.33	1.37
35	BB	1090	A	P-O5'	-6.14	1.53	1.59
85	AA	658	C	C3'-C2'	-6.14	1.46	1.52
85	AA	1129	A	C3'-C2'	-6.14	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1487	G	N1-C2	-6.14	1.32	1.37
34	BA	1722	U	O3'-P	-6.14	1.53	1.61
35	BB	495	A	C3'-C2'	-6.14	1.46	1.52
37	BD	26	C	C3'-C2'	-6.14	1.46	1.52
40	BG	115	C	C4'-C3'	-6.14	1.46	1.53
85	AA	46	U	P-O5'	-6.14	1.53	1.59
85	AA	141	A	C2'-C1'	-6.14	1.46	1.53
85	AA	821	U	O3'-P	-6.14	1.53	1.61
85	AA	997	U	C4'-C3'	-6.14	1.46	1.53
34	BA	319	C	C1'-N1	-6.14	1.38	1.46
34	BA	616	G	N7-C5	-6.14	1.35	1.39
34	BA	1265	G	C3'-C2'	-6.14	1.46	1.52
35	BB	365	U	P-O5'	-6.14	1.53	1.59
35	BB	1111	C	C3'-C2'	-6.14	1.46	1.52
40	BG	76	C	C2'-C1'	-6.14	1.46	1.53
41	BH	33	G	N7-C5	-6.14	1.35	1.39
64	Be	163	ARG	CD-NE	6.14	1.56	1.46
85	AA	681	G	C2'-C1'	-6.14	1.46	1.53
85	AA	1118	U	N3-C4	-6.14	1.32	1.38
85	AA	1820	G	O3'-P	-6.14	1.53	1.61
85	AA	1923	A	N3-C4	-6.14	1.31	1.34
85	AA	2117	U	C2'-C1'	-6.14	1.46	1.53
34	BA	956	G	C6-N1	-6.14	1.35	1.39
34	BA	1249	G	C2'-C1'	-6.14	1.46	1.53
35	BB	1441	C	P-O5'	-6.14	1.53	1.59
85	AA	9	U	C1'-N1	-6.14	1.38	1.46
85	AA	1808	G	N7-C5	-6.14	1.35	1.39
86	AB	62	C	P-O5'	-6.14	1.53	1.59
34	BA	237	A	N7-C5	-6.14	1.35	1.39
34	BA	681	G	O4'-C1'	-6.14	1.33	1.41
34	BA	690	G	C4'-O4'	-6.14	1.37	1.45
34	BA	1232	C	C2-N3	-6.14	1.30	1.35
35	BB	1023	G	N7-C5	-6.14	1.35	1.39
36	BC	63	G	O3'-P	-6.14	1.53	1.61
85	AA	628	C	P-O5'	-6.14	1.53	1.59
85	AA	1167	G	C2'-C1'	-6.14	1.46	1.53
34	BA	229	C	O3'-P	-6.13	1.53	1.61
34	BA	363	G	C5-C4	-6.13	1.34	1.38
34	BA	909	G	C6-N1	-6.13	1.35	1.39
34	BA	913	U	C2-N3	-6.13	1.33	1.37
34	BA	1663	U	C3'-C2'	-6.13	1.46	1.52
34	BA	1798	G	O4'-C1'	-6.13	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	462	G	C1'-N9	-6.13	1.38	1.46
85	AA	149	A	C1'-N9	-6.13	1.38	1.46
85	AA	741	G	C2-N2	-6.13	1.28	1.34
85	AA	811	A	C3'-C2'	-6.13	1.46	1.52
85	AA	868	A	N9-C4	-6.13	1.34	1.37
85	AA	1541	G	O4'-C1'	-6.13	1.33	1.41
85	AA	2138	G	P-O5'	-6.13	1.53	1.59
34	BA	217	C	O3'-P	-6.13	1.53	1.61
34	BA	407	A	C2'-C1'	-6.13	1.46	1.53
34	BA	741	A	N9-C8	-6.13	1.32	1.37
34	BA	1011	G	N7-C5	-6.13	1.35	1.39
35	BB	1235	A	C3'-C2'	-6.13	1.46	1.52
37	BD	1	G	N3-C4	-6.13	1.31	1.35
38	BE	126	G	C3'-O3'	6.13	1.50	1.42
85	AA	1834	U	P-O5'	-6.13	1.53	1.59
34	BA	89	G	O3'-P	-6.13	1.53	1.61
34	BA	111	U	O4'-C1'	-6.13	1.33	1.41
34	BA	143	A	C1'-N9	-6.13	1.38	1.46
34	BA	1036	G	N9-C8	-6.13	1.33	1.37
34	BA	1228	G	O3'-P	-6.13	1.53	1.61
35	BB	57	G	O3'-P	-6.13	1.53	1.61
35	BB	1139	A	C4'-C3'	-6.13	1.46	1.53
39	BF	4	A	N3-C4	-6.13	1.31	1.34
39	BF	23	G	C1'-N9	-6.13	1.38	1.46
34	BA	321	G	C1'-N9	-6.13	1.38	1.46
34	BA	1072	U	O3'-P	-6.13	1.53	1.61
34	BA	1506	C	C5'-C4'	6.13	1.58	1.51
35	BB	1484	A	C6-N6	-6.13	1.29	1.33
36	BC	146	U	C5'-C4'	-6.13	1.44	1.51
37	BD	56	G	N3-C4	-6.13	1.31	1.35
40	BG	55	A	O3'-P	-6.13	1.53	1.61
40	BG	101	G	O3'-P	-6.13	1.53	1.61
41	BH	29	G	C6-O6	-6.13	1.18	1.24
85	AA	1703	A	C5'-C4'	-6.13	1.44	1.51
34	BA	166	G	C5-C4	-6.13	1.34	1.38
34	BA	404	C	C2'-C1'	-6.13	1.46	1.53
34	BA	490	A	C5-C4	-6.13	1.34	1.38
34	BA	904	G	O3'-P	-6.13	1.53	1.61
35	BB	95	A	N9-C8	-6.13	1.32	1.37
35	BB	756	C	C2'-C1'	-6.13	1.46	1.53
35	BB	1184	C	C2'-C1'	-6.13	1.46	1.53
36	BC	122	A	N9-C4	-6.13	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	139	U	O4'-C1'	-6.13	1.33	1.41
38	BE	165	U	O3'-P	-6.13	1.53	1.61
40	BG	88	G	C2-N2	-6.13	1.28	1.34
85	AA	1595	G	P-O5'	-6.13	1.53	1.59
86	AB	28	G	P-O5'	-6.13	1.53	1.59
34	BA	381	A	C1'-N9	-6.13	1.38	1.46
34	BA	502	U	C5'-C4'	-6.13	1.44	1.51
34	BA	1296	U	C2-N3	-6.13	1.33	1.37
34	BA	1703	A	O3'-P	-6.13	1.53	1.61
34	BA	1798	G	C2'-C1'	-6.13	1.46	1.53
35	BB	655	U	C3'-C2'	-6.13	1.46	1.52
35	BB	984	U	P-O5'	-6.13	1.53	1.59
35	BB	1392	A	N9-C4	-6.13	1.34	1.37
39	BF	68	C	O3'-P	-6.13	1.53	1.61
40	BG	30	C	C4'-C3'	-6.13	1.46	1.53
40	BG	135	C	C1'-N1	-6.13	1.38	1.46
41	BH	5	G	C2'-C1'	-6.13	1.46	1.53
41	BH	65	G	O3'-P	-6.13	1.53	1.61
85	AA	453	G	O3'-P	-6.13	1.53	1.61
85	AA	687	G	C5-C6	-6.13	1.36	1.42
85	AA	849	A	C2'-C1'	-6.13	1.46	1.53
85	AA	1163	G	C6-N1	-6.13	1.35	1.39
85	AA	2130	G	C2'-C1'	-6.13	1.46	1.53
85	AA	2135	A	C2'-C1'	-6.13	1.46	1.53
34	BA	1004	U	O3'-P	-6.12	1.53	1.61
34	BA	1664	C	C3'-C2'	-6.12	1.46	1.52
35	BB	587	A	N7-C5	-6.12	1.35	1.39
35	BB	1005	A	O3'-P	-6.12	1.53	1.61
35	BB	1355	C	C3'-C2'	-6.12	1.46	1.52
40	BG	77	U	C4'-C3'	-6.12	1.46	1.53
41	BH	21	G	N9-C8	-6.12	1.33	1.37
85	AA	79	G	C2'-C1'	-6.12	1.46	1.53
85	AA	1095	C	C2'-C1'	-6.12	1.46	1.53
34	BA	795	G	O3'-P	-6.12	1.53	1.61
34	BA	1094	U	N3-C4	-6.12	1.32	1.38
36	BC	36	G	C5-C6	-6.12	1.36	1.42
40	BG	74	G	C3'-C2'	-6.12	1.46	1.52
85	AA	317	A	C2'-C1'	-6.12	1.46	1.53
85	AA	442	G	C2-N3	-6.12	1.27	1.32
85	AA	567	G	P-O5'	-6.12	1.53	1.59
85	AA	983	A	C8-N7	-6.12	1.27	1.31
85	AA	1154	A	C5'-C4'	6.12	1.58	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1355	U	C4'-C3'	6.12	1.59	1.53
85	AA	1839	G	N9-C4	-6.12	1.33	1.38
85	AA	1852	U	C2-N3	-6.12	1.33	1.37
85	AA	2176	U	O4'-C1'	-6.12	1.33	1.41
86	AB	26	A	C2'-C1'	-6.12	1.46	1.53
34	BA	63	A	C2'-C1'	-6.12	1.46	1.53
34	BA	131	A	C2'-C1'	-6.12	1.46	1.53
34	BA	446	U	P-O5'	-6.12	1.53	1.59
34	BA	732	A	C3'-C2'	-6.12	1.46	1.52
35	BB	28	G	C8-N7	-6.12	1.27	1.30
35	BB	113	C	O4'-C1'	-6.12	1.33	1.41
35	BB	384	A	N7-C5	-6.12	1.35	1.39
35	BB	395	U	O3'-P	-6.12	1.53	1.61
35	BB	766	G	O3'-P	-6.12	1.53	1.61
35	BB	1105	G	N3-C4	-6.12	1.31	1.35
35	BB	1328	C	P-O5'	-6.12	1.53	1.59
35	BB	1424	G	C5-C4	-6.12	1.34	1.38
40	BG	81	G	N7-C5	-6.12	1.35	1.39
40	BG	91	U	C2-N3	-6.12	1.33	1.37
85	AA	155	U	N3-C4	-6.12	1.32	1.38
85	AA	1265	C	C2-N3	-6.12	1.30	1.35
34	BA	1040	G	C1'-N9	-6.12	1.38	1.46
34	BA	1044	A	C1'-N9	-6.12	1.38	1.46
35	BB	415	A	O3'-P	-6.12	1.53	1.61
35	BB	1371	G	N7-C5	-6.12	1.35	1.39
41	BH	22	A	P-O5'	-6.12	1.53	1.59
85	AA	1548	A	N9-C4	-6.12	1.34	1.37
34	BA	11	U	N1-C2	-6.12	1.33	1.38
34	BA	324	C	C2'-C1'	-6.12	1.46	1.53
34	BA	373	G	C6-N1	-6.12	1.35	1.39
34	BA	698	U	C3'-C2'	-6.12	1.46	1.52
34	BA	818	G	C5-C4	-6.12	1.34	1.38
34	BA	971	G	C2'-C1'	-6.12	1.46	1.53
34	BA	1044	A	C3'-C2'	-6.12	1.46	1.52
34	BA	1462	U	C5'-C4'	-6.12	1.44	1.51
34	BA	1688	G	C1'-N9	-6.12	1.38	1.46
35	BB	431	U	C2-N3	-6.12	1.33	1.37
35	BB	511	A	N7-C5	-6.12	1.35	1.39
35	BB	975	G	N7-C5	-6.12	1.35	1.39
36	BC	69	U	C2'-C1'	-6.12	1.46	1.53
36	BC	113	G	C3'-C2'	-6.12	1.46	1.52
38	BE	172	U	C3'-C2'	-6.12	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	178	G	N1-C2	-6.12	1.32	1.37
85	AA	554	A	C2'-C1'	-6.12	1.46	1.53
85	AA	1305	A	N7-C5	-6.12	1.35	1.39
85	AA	1698	A	O3'-P	-6.12	1.53	1.61
34	BA	1817	G	N9-C4	-6.12	1.33	1.38
35	BB	51	U	O3'-P	-6.12	1.53	1.61
85	AA	77	C	N1-C6	6.12	1.40	1.37
85	AA	638	G	O3'-P	-6.12	1.53	1.61
85	AA	1157	U	C2-N3	-6.12	1.33	1.37
85	AA	1992	A	P-O5'	-6.12	1.53	1.59
34	BA	115	U	C1'-N1	-6.12	1.38	1.46
34	BA	316	G	C5-C4	-6.12	1.34	1.38
34	BA	722	A	O3'-P	-6.12	1.53	1.61
34	BA	882	G	C5-C4	-6.12	1.34	1.38
34	BA	969	A	C8-N7	-6.12	1.27	1.31
34	BA	1158	A	N3-C4	-6.12	1.31	1.34
34	BA	1527	G	N1-C2	-6.12	1.32	1.37
38	BE	26	G	C2'-C1'	-6.12	1.46	1.53
40	BG	123	C	C2'-C1'	-6.12	1.46	1.53
85	AA	400	G	C1'-N9	-6.12	1.38	1.46
85	AA	1545	U	C2-N3	-6.12	1.33	1.37
34	BA	98	A	N9-C8	-6.11	1.32	1.37
34	BA	900	A	N3-C4	-6.11	1.31	1.34
34	BA	1530	G	C5-C4	-6.11	1.34	1.38
35	BB	519	A	O3'-P	-6.11	1.53	1.61
35	BB	700	C	C2-N3	-6.11	1.30	1.35
35	BB	1090	A	O3'-P	-6.11	1.53	1.61
35	BB	1092	G	C2'-C1'	-6.11	1.46	1.53
35	BB	1249	G	C6-N1	-6.11	1.35	1.39
35	BB	1404	A	O3'-P	-6.11	1.53	1.61
38	BE	14	C	N3-C4	-6.11	1.29	1.33
40	BG	148	C	C4'-C3'	-6.11	1.46	1.53
41	BH	36	C	C4'-C3'	-6.11	1.46	1.53
85	AA	115	U	C3'-C2'	-6.11	1.46	1.52
85	AA	1177	G	C2-N2	-6.11	1.28	1.34
85	AA	1887	G	O3'-P	-6.11	1.53	1.61
85	AA	1910	A	N3-C4	-6.11	1.31	1.34
34	BA	943	G	C2-N3	-6.11	1.27	1.32
34	BA	1498	A	C1'-N9	-6.11	1.38	1.46
34	BA	1522	G	N1-C2	-6.11	1.32	1.37
35	BB	471	U	C2-N3	-6.11	1.33	1.37
35	BB	1272	G	C6-N1	-6.11	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	239	G	P-O5'	-6.11	1.53	1.59
34	BA	112	C	C4'-C3'	-6.11	1.46	1.53
34	BA	215	C	O3'-P	-6.11	1.53	1.61
34	BA	321	G	N9-C4	-6.11	1.33	1.38
34	BA	1232	C	N1-C6	-6.11	1.33	1.37
35	BB	23	U	C2'-C1'	-6.11	1.46	1.53
35	BB	616	U	C4'-C3'	-6.11	1.46	1.53
35	BB	684	U	N3-C4	-6.11	1.32	1.38
35	BB	1358	A	O3'-P	-6.11	1.53	1.61
37	BD	49	A	P-O5'	-6.11	1.53	1.59
40	BG	28	A	C4'-C3'	-6.11	1.46	1.53
85	AA	309	G	C2'-C1'	-6.11	1.46	1.53
85	AA	1197	U	C4'-C3'	-6.11	1.46	1.53
85	AA	2210	C	C3'-C2'	-6.11	1.46	1.52
85	AA	2226	U	C5'-C4'	6.11	1.58	1.51
34	BA	267	G	P-O5'	-6.11	1.53	1.59
34	BA	1470	G	C5-C4	-6.11	1.34	1.38
35	BB	643	G	C4'-C3'	-6.11	1.46	1.53
85	AA	2141	G	P-O5'	-6.11	1.53	1.59
34	BA	753	G	C4'-C3'	-6.11	1.46	1.53
34	BA	1273	U	N1-C2	-6.11	1.33	1.38
34	BA	1278	A	C5-C4	-6.11	1.34	1.38
34	BA	1639	U	C4'-O4'	-6.11	1.37	1.45
34	BA	1683	C	O3'-P	-6.11	1.53	1.61
34	BA	1688	G	O3'-P	-6.11	1.53	1.61
35	BB	665	A	C5-C4	-6.11	1.34	1.38
35	BB	1250	A	N9-C4	-6.11	1.34	1.37
35	BB	1333	U	O3'-P	-6.11	1.53	1.61
37	BD	6	C	C4'-C3'	-6.11	1.46	1.53
40	BG	15	G	C3'-C2'	-6.11	1.46	1.52
40	BG	178	G	C6-N1	-6.11	1.35	1.39
41	BH	27	A	C4'-C3'	6.11	1.59	1.53
85	AA	120	C	C2-N3	-6.11	1.30	1.35
85	AA	181	A	N9-C4	-6.11	1.34	1.37
85	AA	1506	U	O4'-C1'	-6.11	1.33	1.41
85	AA	1991	C	O3'-P	-6.11	1.53	1.61
85	AA	2194	U	C1'-N1	-6.11	1.38	1.46
34	BA	676	G	C2-N2	-6.11	1.28	1.34
34	BA	957	A	C1'-N9	-6.11	1.38	1.46
34	BA	1031	U	C2'-C1'	-6.11	1.46	1.53
34	BA	1405	A	O4'-C1'	-6.11	1.33	1.41
34	BA	1451	A	N9-C8	-6.11	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1641	G	N1-C2	-6.11	1.32	1.37
34	BA	1806	A	P-O5'	-6.11	1.53	1.59
35	BB	688	U	C3'-C2'	-6.11	1.46	1.52
35	BB	773	G	C1'-N9	-6.11	1.38	1.46
36	BC	167	U	N1-C2	-6.11	1.33	1.38
37	BD	109	U	O3'-P	-6.11	1.53	1.61
41	BH	27	A	C3'-O3'	6.11	1.50	1.42
85	AA	285	C	C5'-C4'	6.11	1.58	1.51
85	AA	377	U	C3'-C2'	-6.11	1.46	1.52
85	AA	1980	A	O3'-P	-6.11	1.53	1.61
35	BB	436	G	O3'-P	-6.10	1.53	1.61
35	BB	515	C	C3'-C2'	-6.10	1.46	1.52
36	BC	101	U	C4'-C3'	-6.10	1.46	1.53
38	BE	40	C	C4'-O4'	-6.10	1.37	1.45
85	AA	2017	U	C2'-C1'	-6.10	1.46	1.53
34	BA	67	A	N3-C4	-6.10	1.31	1.34
34	BA	88	C	C4'-C3'	-6.10	1.46	1.53
34	BA	681	G	C4'-O4'	-6.10	1.37	1.45
34	BA	794	G	N1-C2	-6.10	1.32	1.37
34	BA	886	G	C3'-C2'	-6.10	1.46	1.52
34	BA	1017	C	N1-C6	-6.10	1.33	1.37
34	BA	1214	U	C2-N3	-6.10	1.33	1.37
34	BA	1691	G	C5-C6	-6.10	1.36	1.42
35	BB	13	A	N9-C4	-6.10	1.34	1.37
35	BB	361	A	P-O5'	-6.10	1.53	1.59
35	BB	608	A	C2'-C1'	-6.10	1.46	1.53
35	BB	1176	G	C6-N1	-6.10	1.35	1.39
35	BB	1181	A	C5-C4	-6.10	1.34	1.38
35	BB	1286	G	C5-C4	-6.10	1.34	1.38
40	BG	154	C	C4'-C3'	-6.10	1.46	1.53
47	BN	34	PRO	CA-C	-6.10	1.40	1.52
85	AA	474	C	C5'-C4'	-6.10	1.44	1.51
85	AA	1212	C	O3'-P	-6.10	1.53	1.61
85	AA	1450	U	N3-C4	-6.10	1.32	1.38
85	AA	1712	A	P-O5'	-6.10	1.53	1.59
85	AA	1986	G	P-O5'	-6.10	1.53	1.59
34	BA	1634	A	C8-N7	-6.10	1.27	1.31
35	BB	1240	A	C4'-C3'	-6.10	1.46	1.53
41	BH	131	A	C2'-C1'	-6.10	1.46	1.53
34	BA	500	C	C1'-N1	-6.10	1.38	1.46
34	BA	943	G	C5-C4	-6.10	1.34	1.38
34	BA	1064	A	C1'-N9	-6.10	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1252	G	P-O5'	-6.10	1.53	1.59
34	BA	1433	U	O4'-C1'	-6.10	1.33	1.41
34	BA	1478	G	N7-C5	-6.10	1.35	1.39
34	BA	1693	U	C2'-C1'	-6.10	1.46	1.53
35	BB	8	U	O4'-C1'	-6.10	1.33	1.41
35	BB	43	G	N7-C5	-6.10	1.35	1.39
35	BB	261	C	O3'-P	-6.10	1.53	1.61
35	BB	699	U	C2-N3	-6.10	1.33	1.37
35	BB	882	U	O3'-P	-6.10	1.53	1.61
69	Bj	67	LEU	N-CA	-6.10	1.34	1.46
85	AA	1213	U	O4'-C1'	-6.10	1.33	1.41
34	BA	136	A	C5-C4	-6.10	1.34	1.38
34	BA	430	A	C1'-N9	-6.10	1.38	1.46
34	BA	603	U	C5'-C4'	-6.10	1.44	1.51
34	BA	720	A	C2'-C1'	-6.10	1.46	1.53
34	BA	730	C	N1-C6	-6.10	1.33	1.37
34	BA	1248	A	N9-C4	-6.10	1.34	1.37
34	BA	1531	G	C2-N2	-6.10	1.28	1.34
35	BB	12	G	C2'-C1'	-6.10	1.46	1.53
35	BB	164	U	P-O5'	-6.10	1.53	1.59
35	BB	1311	G	N1-C2	-6.10	1.32	1.37
35	BB	1460	G	N9-C4	-6.10	1.33	1.38
36	BC	47	C	O3'-P	-6.10	1.53	1.61
38	BE	49	A	C5-C4	-6.10	1.34	1.38
39	BF	34	C	O3'-P	-6.10	1.53	1.61
40	BG	12	A	O3'-P	-6.10	1.53	1.61
41	BH	36	C	C1'-N1	-6.10	1.38	1.46
85	AA	2091	C	C3'-C2'	-6.10	1.46	1.52
34	BA	102	G	N1-C2	-6.10	1.32	1.37
34	BA	432	A	C2'-C1'	-6.10	1.46	1.53
34	BA	1466	U	P-O5'	-6.10	1.53	1.59
85	AA	661	C	N1-C6	-6.10	1.33	1.37
85	AA	1708	A	N7-C5	-6.10	1.35	1.39
34	BA	305	C	C2'-C1'	-6.09	1.46	1.53
34	BA	339	G	N1-C2	-6.09	1.32	1.37
34	BA	732	A	P-O5'	-6.09	1.53	1.59
34	BA	868	C	C2'-C1'	-6.09	1.46	1.53
34	BA	911	G	C2-N2	-6.09	1.28	1.34
34	BA	974	G	C4'-C3'	-6.09	1.46	1.53
34	BA	1746	G	P-O5'	-6.09	1.53	1.59
35	BB	456	A	N3-C4	-6.09	1.31	1.34
35	BB	482	A	C8-N7	-6.09	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	792	G	C2-N3	-6.09	1.27	1.32
37	BD	2	G	P-O5'	-6.09	1.53	1.59
38	BE	47	U	N3-C4	-6.09	1.32	1.38
38	BE	92	C	O3'-P	-6.09	1.53	1.61
85	AA	915	G	N9-C4	-6.09	1.33	1.38
85	AA	1153	G	N9-C4	-6.09	1.33	1.38
85	AA	1857	G	O3'-P	-6.09	1.53	1.61
34	BA	42	A	C5-C4	-6.09	1.34	1.38
35	BB	122	U	C3'-C2'	-6.09	1.46	1.52
35	BB	416	U	C2-N3	-6.09	1.33	1.37
35	BB	1543	C	C2-N3	-6.09	1.30	1.35
85	AA	1107	A	C6-N1	-6.09	1.31	1.35
85	AA	1245	U	P-O5'	-6.09	1.53	1.59
34	BA	179	U	C4'-C3'	6.09	1.59	1.53
34	BA	253	U	O3'-P	-6.09	1.53	1.61
34	BA	1258	G	N3-C4	-6.09	1.31	1.35
34	BA	1736	A	C2'-C1'	-6.09	1.46	1.53
35	BB	43	G	C4'-C3'	-6.09	1.46	1.53
35	BB	1070	G	C1'-N9	-6.09	1.38	1.46
35	BB	1172	U	C3'-C2'	-6.09	1.46	1.52
35	BB	1374	U	O3'-P	-6.09	1.53	1.61
40	BG	65	C	C1'-N1	-6.09	1.38	1.46
40	BG	130	G	P-O5'	-6.09	1.53	1.59
85	AA	427	G	C2'-C1'	-6.09	1.46	1.53
34	BA	26	C	C1'-N1	-6.09	1.38	1.46
34	BA	127	U	C2-N3	-6.09	1.33	1.37
34	BA	582	U	O3'-P	-6.09	1.53	1.61
34	BA	774	A	C3'-C2'	-6.09	1.46	1.52
34	BA	784	C	C4-N4	-6.09	1.28	1.33
34	BA	854	A	C4'-C3'	-6.09	1.46	1.53
34	BA	1729	G	C3'-O3'	6.09	1.50	1.42
35	BB	630	A	C5-C4	-6.09	1.34	1.38
35	BB	709	G	O3'-P	-6.09	1.53	1.61
35	BB	1030	U	O3'-P	-6.09	1.53	1.61
35	BB	1417	C	C2'-C1'	-6.09	1.46	1.53
38	BE	50	G	C3'-C2'	-6.09	1.46	1.52
85	AA	8	U	C2'-C1'	-6.09	1.46	1.53
85	AA	537	G	C3'-C2'	-6.09	1.46	1.52
85	AA	553	G	C3'-C2'	-6.09	1.46	1.52
85	AA	889	G	C8-N7	-6.09	1.27	1.30
85	AA	916	A	N9-C4	-6.09	1.34	1.37
85	AA	1543	C	C4'-C3'	6.09	1.59	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	76	U	C4'-C3'	-6.09	1.46	1.53
34	BA	86	A	O3'-P	-6.09	1.53	1.61
36	BC	72	A	O3'-P	-6.09	1.53	1.61
34	BA	182	U	N3-C4	-6.09	1.32	1.38
34	BA	696	A	N9-C4	-6.09	1.34	1.37
34	BA	950	C	C3'-C2'	-6.09	1.46	1.52
34	BA	1161	G	C3'-C2'	-6.09	1.46	1.52
34	BA	1776	G	P-O5'	-6.09	1.53	1.59
35	BB	812	G	O3'-P	-6.09	1.53	1.61
35	BB	841	U	C2'-C1'	-6.09	1.46	1.53
35	BB	960	C	O3'-P	-6.09	1.53	1.61
35	BB	1470	G	C2-N2	-6.09	1.28	1.34
36	BC	14	G	C1'-N9	-6.09	1.38	1.46
37	BD	36	C	C3'-C2'	-6.09	1.46	1.52
38	BE	123	A	C3'-C2'	-6.09	1.46	1.52
38	BE	130	G	C6-N1	-6.09	1.35	1.39
85	AA	113	U	N1-C2	-6.09	1.33	1.38
85	AA	1217	U	O4'-C1'	-6.09	1.33	1.41
34	BA	492	G	C2'-C1'	-6.08	1.46	1.53
85	AA	365	G	C3'-C2'	-6.08	1.46	1.52
85	AA	943	U	P-O5'	-6.08	1.53	1.59
85	AA	1558	U	N3-C4	-6.08	1.32	1.38
34	BA	781	U	O3'-P	-6.08	1.53	1.61
34	BA	947	A	C2'-C1'	-6.08	1.46	1.53
34	BA	1190	A	C3'-C2'	-6.08	1.46	1.52
34	BA	1799	G	C3'-O3'	6.08	1.50	1.42
35	BB	35	G	N9-C4	-6.08	1.33	1.38
35	BB	143	G	P-O5'	-6.08	1.53	1.59
35	BB	1135	U	N1-C2	-6.08	1.33	1.38
40	BG	37	G	C2-N2	-6.08	1.28	1.34
85	AA	406	U	O3'-P	-6.08	1.53	1.61
85	AA	1561	A	C2'-C1'	-6.08	1.46	1.53
85	AA	2090	C	C3'-C2'	-6.08	1.46	1.52
34	BA	257	G	C2-N2	-6.08	1.28	1.34
34	BA	325	A	C8-N7	-6.08	1.27	1.31
34	BA	1175	G	N7-C5	-6.08	1.35	1.39
34	BA	1246	G	C3'-C2'	-6.08	1.46	1.52
34	BA	1432	C	C4-N4	-6.08	1.28	1.33
34	BA	1589	U	O3'-P	-6.08	1.53	1.61
34	BA	1649	A	C5-C4	-6.08	1.34	1.38
34	BA	1740	U	C2'-C1'	-6.08	1.46	1.53
35	BB	548	A	O3'-P	-6.08	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	727	U	P-O5'	-6.08	1.53	1.59
35	BB	1193	G	C1'-N9	-6.08	1.38	1.46
36	BC	71	A	C2'-C1'	-6.08	1.46	1.53
85	AA	157	G	P-O5'	-6.08	1.53	1.59
85	AA	159	G	P-O5'	-6.08	1.53	1.59
85	AA	363	A	C4'-C3'	-6.08	1.46	1.53
85	AA	585	G	N9-C8	-6.08	1.33	1.37
85	AA	2227	A	C2'-C1'	-6.08	1.46	1.53
34	BA	674	G	N9-C8	-6.08	1.33	1.37
34	BA	1410	C	C4'-C3'	-6.08	1.46	1.53
34	BA	1524	G	C2-N2	-6.08	1.28	1.34
35	BB	481	A	C6-N1	-6.08	1.31	1.35
35	BB	680	A	O3'-P	-6.08	1.53	1.61
35	BB	691	A	C5-C4	-6.08	1.34	1.38
34	BA	57	A	C4'-C3'	-6.08	1.46	1.53
34	BA	357	A	P-O5'	-6.08	1.53	1.59
34	BA	891	C	N1-C6	-6.08	1.33	1.37
34	BA	1119	A	O3'-P	-6.08	1.53	1.61
34	BA	1568	A	C1'-N9	-6.08	1.38	1.46
35	BB	806	U	C2'-C1'	-6.08	1.46	1.53
35	BB	806	U	N3-C4	-6.08	1.32	1.38
37	BD	95	G	O4'-C1'	-6.08	1.33	1.41
38	BE	9	C	O3'-P	-6.08	1.53	1.61
85	AA	312	G	C2'-C1'	-6.08	1.46	1.53
85	AA	1432	C	O3'-P	-6.08	1.53	1.61
85	AA	1855	U	C3'-C2'	-6.08	1.46	1.52
85	AA	1957	C	C2'-C1'	-6.08	1.46	1.53
85	AA	2105	G	C1'-N9	-6.08	1.38	1.46
85	AA	2237	G	C2-N2	-6.08	1.28	1.34
34	BA	385	U	C3'-C2'	-6.08	1.46	1.52
34	BA	869	C	N3-C4	6.08	1.38	1.33
35	BB	1159	U	N3-C4	-6.08	1.32	1.38
37	BD	77	A	C4'-C3'	-6.08	1.46	1.53
85	AA	1665	G	C2'-C1'	-6.08	1.46	1.53
34	BA	246	G	C4'-O4'	-6.08	1.37	1.45
34	BA	493	G	C4'-C3'	-6.08	1.46	1.53
34	BA	616	G	C5'-C4'	-6.08	1.44	1.51
34	BA	1312	A	N7-C5	-6.08	1.35	1.39
34	BA	1421	A	P-O5'	-6.08	1.53	1.59
34	BA	1451	A	N9-C4	-6.08	1.34	1.37
35	BB	505	G	C2'-C1'	-6.08	1.46	1.53
35	BB	702	G	C5-C4	-6.08	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	831	C	C4'-C3'	-6.08	1.46	1.53
35	BB	1410	G	C3'-C2'	-6.08	1.46	1.52
35	BB	1410	G	N3-C4	-6.08	1.31	1.35
37	BD	104	C	O3'-P	-6.08	1.53	1.61
39	BF	35	C	O3'-P	-6.08	1.53	1.61
85	AA	2096	G	N9-C8	-6.08	1.33	1.37
85	AA	2148	C	C4-N4	-6.08	1.28	1.33
34	BA	1291	A	C1'-N9	-6.07	1.38	1.46
34	BA	1574	C	P-O5'	-6.07	1.53	1.59
35	BB	1149	A	C2'-C1'	-6.07	1.46	1.53
36	BC	15	G	O3'-P	-6.07	1.53	1.61
37	BD	5	A	C2'-C1'	-6.07	1.46	1.53
85	AA	566	U	C2'-C1'	-6.07	1.46	1.53
85	AA	1105	G	O3'-P	-6.07	1.53	1.61
85	AA	2117	U	C3'-C2'	-6.07	1.46	1.52
85	AA	2188	C	C2-N3	-6.07	1.30	1.35
35	BB	133	G	C2'-C1'	-6.07	1.46	1.53
38	BE	36	U	N1-C2	6.07	1.44	1.38
40	BG	140	G	N9-C4	-6.07	1.33	1.38
41	BH	120	C	C2'-C1'	-6.07	1.46	1.53
85	AA	936	C	O4'-C1'	-6.07	1.33	1.41
35	BB	1079	G	C2-N2	-6.07	1.28	1.34
35	BB	1249	G	P-O5'	-6.07	1.53	1.59
36	BC	96	A	N9-C8	-6.07	1.32	1.37
38	BE	20	C	P-O5'	6.07	1.65	1.59
40	BG	47	G	O3'-P	-6.07	1.53	1.61
85	AA	459	C	C3'-C2'	-6.07	1.46	1.52
85	AA	669	G	O3'-P	-6.07	1.53	1.61
85	AA	893	G	N9-C4	-6.07	1.33	1.38
85	AA	1262	A	N3-C4	-6.07	1.31	1.34
85	AA	1813	C	P-O5'	-6.07	1.53	1.59
85	AA	1963	G	C4'-O4'	-6.07	1.37	1.45
34	BA	164	C	C2'-C1'	-6.07	1.46	1.53
35	BB	115	A	O3'-P	-6.07	1.53	1.61
35	BB	399	A	C5-C4	-6.07	1.34	1.38
85	AA	1892	G	C2'-C1'	-6.07	1.46	1.53
85	AA	2026	U	N1-C2	-6.07	1.33	1.38
34	BA	279	U	C2'-C1'	-6.07	1.46	1.53
34	BA	847	U	C5'-C4'	6.07	1.58	1.51
34	BA	1087	A	C2'-C1'	-6.07	1.46	1.53
34	BA	1278	A	C1'-N9	-6.07	1.38	1.46
34	BA	1521	C	P-O5'	-6.07	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	837	A	O4'-C1'	-6.07	1.33	1.41
35	BB	1136	G	C2-N2	-6.07	1.28	1.34
36	BC	122	A	C5-C4	-6.07	1.34	1.38
38	BE	137	A	C5-C4	-6.07	1.34	1.38
38	BE	173	G	P-O5'	-6.07	1.53	1.59
39	BF	46	G	C6-N1	-6.07	1.35	1.39
40	BG	114	A	C2'-C1'	-6.07	1.46	1.53
41	BH	33	G	N9-C8	-6.07	1.33	1.37
85	AA	398	U	C3'-C2'	-6.07	1.46	1.52
85	AA	1495	G	C5-C4	-6.07	1.34	1.38
85	AA	1504	A	N9-C8	-6.07	1.32	1.37
85	AA	1697	C	C2'-C1'	-6.07	1.46	1.53
85	AA	2062	U	C5'-C4'	-6.07	1.44	1.51
85	AA	2069	A	O3'-P	-6.07	1.53	1.61
85	AA	2094	U	C2-N3	-6.07	1.33	1.37
85	AA	2201	A	C5-C4	-6.07	1.34	1.38
86	AB	5	G	C3'-C2'	-6.07	1.46	1.52
34	BA	277	A	N9-C4	-6.07	1.34	1.37
34	BA	1333	G	N9-C8	-6.07	1.33	1.37
35	BB	131	A	N1-C2	-6.07	1.28	1.34
35	BB	1015	U	O3'-P	-6.07	1.53	1.61
35	BB	1091	C	O3'-P	-6.07	1.53	1.61
38	BE	170	U	C4'-O4'	-6.07	1.37	1.45
38	BE	203	C	O3'-P	-6.07	1.53	1.61
85	AA	897	A	N7-C5	-6.07	1.35	1.39
85	AA	1877	G	C4'-C3'	-6.07	1.46	1.53
34	BA	489	A	C2'-C1'	-6.06	1.46	1.53
34	BA	1653	G	N1-C2	-6.06	1.32	1.37
37	BD	94	C	C4'-C3'	-6.06	1.46	1.53
37	BD	97	U	C3'-C2'	-6.06	1.46	1.52
85	AA	1257	A	P-O5'	-6.06	1.53	1.59
34	BA	373	G	C2-N3	-6.06	1.27	1.32
34	BA	674	G	O3'-P	-6.06	1.53	1.61
35	BB	1109	A	C5-C4	-6.06	1.34	1.38
35	BB	1462	G	C5'-C4'	6.06	1.58	1.51
35	BB	1512	C	N1-C6	6.06	1.40	1.37
35	BB	1523	U	O3'-P	-6.06	1.53	1.61
85	AA	422	G	C1'-N9	-6.06	1.38	1.46
85	AA	924	A	O3'-P	-6.06	1.53	1.61
85	AA	2098	A	N7-C5	-6.06	1.35	1.39
85	AA	2111	C	O3'-P	-6.06	1.53	1.61
34	BA	438	A	C2'-C1'	-6.06	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1028	A	C5'-C4'	-6.06	1.44	1.51
35	BB	448	G	N3-C4	-6.06	1.31	1.35
35	BB	1404	A	C8-N7	-6.06	1.27	1.31
37	BD	54	A	O3'-P	-6.06	1.53	1.61
45	BL	112	GLY	CA-C	-6.06	1.42	1.51
85	AA	37	U	O3'-P	-6.06	1.53	1.61
34	BA	89	G	C3'-C2'	-6.06	1.46	1.52
34	BA	277	A	C2'-C1'	-6.06	1.46	1.53
34	BA	438	A	C4'-C3'	6.06	1.59	1.53
34	BA	1312	A	C4'-O4'	-6.06	1.37	1.45
34	BA	1403	G	O3'-P	-6.06	1.53	1.61
35	BB	585	U	O3'-P	-6.06	1.53	1.61
35	BB	1076	U	C2-N3	-6.06	1.33	1.37
35	BB	1079	G	C6-N1	-6.06	1.35	1.39
38	BE	128	G	N1-C2	-6.06	1.32	1.37
85	AA	19	A	C1'-N9	-6.06	1.38	1.46
85	AA	356	U	N1-C2	-6.06	1.33	1.38
85	AA	732	G	C4'-C3'	-6.06	1.46	1.53
85	AA	1228	A	C8-N7	-6.06	1.27	1.31
34	BA	236	A	C4'-C3'	-6.06	1.46	1.53
34	BA	655	U	O4'-C1'	-6.06	1.33	1.41
34	BA	1057	C	C4'-C3'	-6.06	1.46	1.53
34	BA	1323	G	C1'-N9	-6.06	1.38	1.46
35	BB	582	G	O3'-P	-6.06	1.53	1.61
35	BB	666	A	O4'-C1'	-6.06	1.33	1.41
35	BB	1330	A	C2'-C1'	-6.06	1.46	1.53
36	BC	68	A	C3'-C2'	-6.06	1.46	1.52
38	BE	114	G	P-O5'	-6.06	1.53	1.59
40	BG	34	A	O3'-P	-6.06	1.53	1.61
59	BZ	80	ASP	N-CA	-6.06	1.34	1.46
85	AA	206	U	P-O5'	-6.06	1.53	1.59
85	AA	504	U	C4'-O4'	-6.06	1.37	1.45
85	AA	1253	G	C1'-N9	-6.06	1.38	1.46
85	AA	1858	G	O3'-P	-6.06	1.53	1.61
34	BA	517	A	N9-C8	-6.06	1.32	1.37
34	BA	524	G	N7-C5	-6.06	1.35	1.39
34	BA	557	U	P-OP2	-6.06	1.38	1.49
34	BA	627	U	O3'-P	-6.06	1.53	1.61
34	BA	1522	G	C8-N7	-6.06	1.27	1.30
34	BA	1731	A	C8-N7	-6.06	1.27	1.31
35	BB	563	A	O3'-P	-6.06	1.53	1.61
35	BB	1258	G	C5-C4	-6.06	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1462	G	N1-C2	-6.06	1.32	1.37
34	BA	409	A	N7-C5	-6.05	1.35	1.39
34	BA	563	A	C4'-C3'	-6.05	1.46	1.53
34	BA	1390	C	P-O5'	-6.05	1.53	1.59
34	BA	1547	G	C5-C4	-6.05	1.34	1.38
35	BB	42	A	C1'-N9	-6.05	1.38	1.46
35	BB	688	U	N3-C4	-6.05	1.33	1.38
35	BB	832	C	O3'-P	-6.05	1.53	1.61
35	BB	973	G	N9-C4	6.05	1.42	1.38
35	BB	1311	G	C5-C4	-6.05	1.34	1.38
35	BB	1439	U	C2-N3	-6.05	1.33	1.37
38	BE	124	G	C5-C4	-6.05	1.34	1.38
39	BF	27	G	P-O5'	-6.05	1.53	1.59
85	AA	867	G	C2-N2	-6.05	1.28	1.34
85	AA	982	G	N3-C4	-6.05	1.31	1.35
85	AA	1112	G	C8-N7	-6.05	1.27	1.30
85	AA	1498	C	C3'-C2'	-6.05	1.46	1.52
85	AA	1505	G	C5-C4	-6.05	1.34	1.38
85	AA	1645	G	O3'-P	-6.05	1.53	1.61
85	AA	2105	G	C6-N1	-6.05	1.35	1.39
34	BA	790	G	C5-C6	-6.05	1.36	1.42
34	BA	901	C	C4'-O4'	-6.05	1.37	1.45
34	BA	1292	A	N3-C4	-6.05	1.31	1.34
34	BA	1584	G	P-O5'	-6.05	1.53	1.59
66	Bg	61	PRO	CA-C	-6.05	1.40	1.52
85	AA	681	G	C5-C4	-6.05	1.34	1.38
85	AA	1283	C	N3-C4	-6.05	1.29	1.33
34	BA	401	A	C5-C4	-6.05	1.34	1.38
34	BA	406	G	C1'-N9	-6.05	1.38	1.46
35	BB	72	G	N3-C4	-6.05	1.31	1.35
35	BB	1350	A	C2'-C1'	-6.05	1.46	1.53
35	BB	1414	A	C6-N1	-6.05	1.31	1.35
36	BC	144	C	C4'-O4'	-6.05	1.37	1.45
36	BC	162	C	N3-C4	-6.05	1.29	1.33
38	BE	178	G	C2-N2	-6.05	1.28	1.34
85	AA	513	G	N7-C5	-6.05	1.35	1.39
85	AA	742	U	N3-C4	-6.05	1.33	1.38
85	AA	2047	U	P-O5'	-6.05	1.53	1.59
34	BA	565	U	C2'-C1'	-6.05	1.46	1.53
34	BA	580	U	N3-C4	-6.05	1.33	1.38
34	BA	750	C	P-O5'	-6.05	1.53	1.59
34	BA	1034	U	C2'-C1'	-6.05	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1816	G	O3'-P	-6.05	1.53	1.61
35	BB	889	U	O3'-P	-6.05	1.53	1.61
36	BC	109	A	P-O5'	-6.05	1.53	1.59
37	BD	78	C	C4'-C3'	-6.05	1.46	1.53
69	Bj	45	THR	C-N	-6.05	1.22	1.34
85	AA	252	G	C5-C6	-6.05	1.36	1.42
85	AA	833	U	O3'-P	-6.05	1.53	1.61
85	AA	1255	C	C3'-C2'	-6.05	1.46	1.52
34	BA	414	A	C4'-O4'	-6.05	1.37	1.45
34	BA	1255	G	C3'-C2'	-6.05	1.46	1.52
35	BB	1161	G	N3-C4	-6.05	1.31	1.35
85	AA	2175	U	C3'-C2'	-6.05	1.46	1.52
34	BA	1827	C	C4-N4	-6.05	1.28	1.33
35	BB	488	G	N9-C4	-6.05	1.33	1.38
35	BB	753	A	P-O5'	-6.05	1.53	1.59
40	BG	32	U	O3'-P	-6.05	1.53	1.61
40	BG	131	U	C2'-C1'	-6.05	1.46	1.53
85	AA	133	G	C3'-C2'	-6.05	1.46	1.52
85	AA	1177	G	C2'-C1'	-6.05	1.46	1.53
85	AA	2191	C	N1-C2	-6.05	1.34	1.40
34	BA	432	A	C3'-C2'	-6.04	1.46	1.52
34	BA	1631	U	C2'-C1'	-6.04	1.46	1.53
38	BE	14	C	O3'-P	-6.04	1.53	1.61
85	AA	1367	C	C4'-C3'	-6.04	1.46	1.53
34	BA	140	C	C2'-C1'	-6.04	1.46	1.53
34	BA	401	A	C6-N6	-6.04	1.29	1.33
34	BA	454	G	C3'-C2'	-6.04	1.46	1.52
34	BA	536	C	O3'-P	-6.04	1.53	1.61
34	BA	772	G	C5'-C4'	6.04	1.58	1.51
34	BA	1203	G	C5-C6	-6.04	1.36	1.42
34	BA	1670	A	C4'-C3'	-6.04	1.46	1.53
35	BB	261	C	P-O5'	-6.04	1.53	1.59
35	BB	661	G	C4'-C3'	-6.04	1.46	1.53
35	BB	1286	G	C6-N1	-6.04	1.35	1.39
38	BE	51	C	O3'-P	-6.04	1.53	1.61
40	BG	108	G	O3'-P	-6.04	1.53	1.61
41	BH	43	G	N3-C4	-6.04	1.31	1.35
41	BH	133	U	N3-C4	-6.04	1.33	1.38
85	AA	421	G	C2'-C1'	-6.04	1.46	1.53
85	AA	678	A	C5-C4	-6.04	1.34	1.38
85	AA	776	C	C2-N3	-6.04	1.30	1.35
85	AA	795	C	N1-C6	-6.04	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1155	A	O3'-P	-6.04	1.53	1.61
85	AA	2009	A	O3'-P	-6.04	1.53	1.61
34	BA	327	G	C6-N1	-6.04	1.35	1.39
34	BA	375	C	N1-C6	-6.04	1.33	1.37
34	BA	413	A	C4'-C3'	-6.04	1.46	1.53
34	BA	782	C	C4'-C3'	-6.04	1.46	1.53
35	BB	403	U	P-O5'	-6.04	1.53	1.59
35	BB	1035	C	C5'-C4'	-6.04	1.44	1.51
35	BB	1191	G	C2-N2	-6.04	1.28	1.34
38	BE	3	G	N3-C4	-6.04	1.31	1.35
40	BG	134	U	C2'-C1'	-6.04	1.46	1.53
41	BH	10	U	C2-N3	-6.04	1.33	1.37
85	AA	27	U	C4'-C3'	6.04	1.59	1.53
85	AA	335	G	C2-N2	-6.04	1.28	1.34
85	AA	355	G	P-O5'	-6.04	1.53	1.59
85	AA	497	G	C2'-C1'	-6.04	1.46	1.53
85	AA	1142	G	C1'-N9	-6.04	1.38	1.46
85	AA	1558	U	C2'-C1'	-6.04	1.46	1.53
35	BB	802	G	C2-N2	-6.04	1.28	1.34
35	BB	971	A	C3'-C2'	-6.04	1.46	1.52
37	BD	72	U	C3'-C2'	-6.04	1.46	1.52
38	BE	191	U	C2-N3	-6.04	1.33	1.37
40	BG	8	U	N1-C6	-6.04	1.32	1.38
85	AA	706	U	C3'-C2'	-6.04	1.46	1.52
85	AA	1701	G	P-O5'	-6.04	1.53	1.59
85	AA	1794	U	C1'-N1	-6.04	1.38	1.46
85	AA	1844	A	C1'-N9	-6.04	1.38	1.46
34	BA	593	G	C6-N1	-6.04	1.35	1.39
34	BA	866	C	N1-C6	6.04	1.40	1.37
34	BA	1063	G	O3'-P	-6.04	1.53	1.61
34	BA	1210	A	C8-N7	-6.04	1.27	1.31
35	BB	869	G	N9-C4	-6.04	1.33	1.38
35	BB	1280	U	P-O5'	-6.04	1.53	1.59
36	BC	1	A	O3'-P	-6.04	1.53	1.61
36	BC	161	U	O3'-P	-6.04	1.53	1.61
38	BE	26	G	C3'-C2'	-6.04	1.46	1.52
40	BG	25	G	C5-C4	-6.04	1.34	1.38
85	AA	113	U	C2-N3	-6.04	1.33	1.37
85	AA	464	A	C5-C4	-6.04	1.34	1.38
85	AA	635	G	C2-N2	-6.04	1.28	1.34
85	AA	1228	A	N7-C5	-6.04	1.35	1.39
85	AA	1854	U	O3'-P	-6.04	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1024	A	C2'-C1'	-6.04	1.46	1.53
34	BA	1209	A	P-O5'	-6.04	1.53	1.59
35	BB	1133	C	C3'-C2'	-6.04	1.46	1.52
35	BB	1254	G	C1'-N9	-6.04	1.38	1.46
38	BE	12	A	N7-C5	-6.04	1.35	1.39
85	AA	2104	C	P-O5'	-6.04	1.53	1.59
34	BA	93	A	C5-C4	-6.04	1.34	1.38
34	BA	100	A	P-O5'	-6.04	1.53	1.59
34	BA	820	C	C2-N3	-6.04	1.30	1.35
34	BA	857	C	C5'-C4'	-6.04	1.44	1.51
34	BA	1067	G	C2'-C1'	-6.04	1.46	1.53
34	BA	1556	A	C8-N7	-6.04	1.27	1.31
35	BB	59	U	N1-C2	-6.04	1.33	1.38
35	BB	125	G	C2-N2	-6.04	1.28	1.34
35	BB	1086	G	C8-N7	-6.04	1.27	1.30
35	BB	1094	A	C2'-C1'	-6.04	1.46	1.53
38	BE	11	A	N7-C5	-6.04	1.35	1.39
38	BE	136	G	C1'-N9	-6.04	1.38	1.46
39	BF	36	G	C2'-C1'	-6.04	1.46	1.53
85	AA	95	U	C3'-C2'	-6.04	1.46	1.52
85	AA	1269	A	C4'-C3'	-6.04	1.46	1.53
34	BA	372	U	O4'-C1'	-6.03	1.33	1.41
34	BA	769	U	C2'-C1'	-6.03	1.46	1.53
34	BA	791	A	N7-C5	-6.03	1.35	1.39
34	BA	1834	A	C6-N1	-6.03	1.31	1.35
35	BB	685	G	O3'-P	-6.03	1.53	1.61
36	BC	109	A	O3'-P	-6.03	1.53	1.61
85	AA	141	A	O3'-P	-6.03	1.53	1.61
85	AA	717	G	C3'-C2'	-6.03	1.46	1.52
85	AA	1209	U	C3'-C2'	-6.03	1.46	1.52
85	AA	1436	A	C5'-C4'	6.03	1.58	1.51
85	AA	1578	G	O3'-P	-6.03	1.53	1.61
85	AA	1713	A	C2'-C1'	-6.03	1.46	1.53
34	BA	606	G	C6-N1	-6.03	1.35	1.39
34	BA	953	G	C2-N2	-6.03	1.28	1.34
35	BB	134	G	C2-N2	-6.03	1.28	1.34
35	BB	1500	U	O3'-P	-6.03	1.53	1.61
36	BC	6	G	N7-C5	-6.03	1.35	1.39
85	AA	423	G	C6-N1	-6.03	1.35	1.39
85	AA	1463	A	N7-C5	-6.03	1.35	1.39
34	BA	1496	G	C5'-C4'	-6.03	1.44	1.51
34	BA	1837	U	C3'-C2'	-6.03	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	19	C	P-O5'	-6.03	1.53	1.59
35	BB	102	G	C3'-C2'	-6.03	1.46	1.52
35	BB	409	U	C3'-C2'	-6.03	1.46	1.52
35	BB	422	U	O3'-P	-6.03	1.53	1.61
35	BB	651	G	C2-N2	-6.03	1.28	1.34
35	BB	1136	G	C2-N3	-6.03	1.27	1.32
35	BB	1336	G	N7-C5	-6.03	1.35	1.39
39	BF	40	U	C3'-C2'	6.03	1.59	1.52
40	BG	121	C	C4-N4	-6.03	1.28	1.33
85	AA	112	A	O4'-C1'	-6.03	1.33	1.41
85	AA	395	G	C4'-C3'	-6.03	1.46	1.53
85	AA	615	A	C2'-C1'	-6.03	1.46	1.53
85	AA	730	G	C2-N2	-6.03	1.28	1.34
85	AA	794	A	C3'-C2'	-6.03	1.46	1.52
85	AA	1000	U	C4'-O4'	-6.03	1.37	1.45
85	AA	2149	C	O3'-P	-6.03	1.53	1.61
34	BA	12	G	N1-C2	-6.03	1.32	1.37
34	BA	233	U	N3-C4	-6.03	1.33	1.38
34	BA	1110	A	N9-C8	-6.03	1.32	1.37
35	BB	36	U	C2-N3	-6.03	1.33	1.37
85	AA	1356	U	P-O5'	-6.03	1.53	1.59
85	AA	1600	G	C2'-C1'	-6.03	1.46	1.53
85	AA	2025	A	N9-C4	-6.03	1.34	1.37
34	BA	60	A	C3'-C2'	-6.03	1.46	1.52
34	BA	177	G	O3'-P	-6.03	1.53	1.61
34	BA	731	A	N7-C5	-6.03	1.35	1.39
34	BA	1696	G	C1'-N9	-6.03	1.38	1.46
35	BB	28	G	N1-C2	-6.03	1.32	1.37
35	BB	545	C	C4-N4	-6.03	1.28	1.33
35	BB	1262	A	O3'-P	-6.03	1.53	1.61
40	BG	54	G	N9-C4	-6.03	1.33	1.38
85	AA	14	C	O3'-P	-6.03	1.53	1.61
85	AA	499	G	P-O5'	-6.03	1.53	1.59
85	AA	634	U	C3'-C2'	-6.03	1.46	1.52
85	AA	1736	U	C3'-O3'	6.03	1.50	1.42
85	AA	1825	A	N7-C5	-6.03	1.35	1.39
34	BA	335	C	C2-N3	-6.03	1.30	1.35
34	BA	582	U	C1'-N1	-6.03	1.38	1.46
34	BA	904	G	P-O5'	-6.03	1.53	1.59
34	BA	985	C	C4'-C3'	-6.03	1.46	1.53
34	BA	1609	U	N3-C4	-6.03	1.33	1.38
34	BA	1703	A	N3-C4	-6.03	1.31	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1722	U	C2-N3	-6.03	1.33	1.37
35	BB	845	C	P-O5'	-6.03	1.53	1.59
35	BB	1455	A	C5'-C4'	6.03	1.58	1.51
36	BC	122	A	C1'-N9	-6.03	1.38	1.46
38	BE	12	A	N3-C4	-6.03	1.31	1.34
85	AA	542	G	C2-N2	-6.03	1.28	1.34
85	AA	763	U	N1-C2	-6.03	1.33	1.38
85	AA	1004	G	C3'-C2'	-6.03	1.46	1.52
85	AA	1130	G	O3'-P	-6.03	1.53	1.61
85	AA	1658	G	O3'-P	-6.03	1.53	1.61
34	BA	990	G	P-O5'	-6.02	1.53	1.59
35	BB	488	G	C4'-O4'	-6.02	1.37	1.45
40	BG	114	A	P-O5'	-6.02	1.53	1.59
85	AA	54	C	C2'-C1'	-6.02	1.46	1.53
85	AA	317	A	C1'-N9	-6.02	1.38	1.46
85	AA	455	G	N3-C4	-6.02	1.31	1.35
85	AA	2062	U	C2-N3	-6.02	1.33	1.37
34	BA	114	U	C4'-C3'	6.02	1.59	1.53
34	BA	344	G	C1'-N9	-6.02	1.38	1.46
34	BA	364	C	C2-N3	-6.02	1.30	1.35
34	BA	409	A	C4'-C3'	-6.02	1.46	1.53
34	BA	916	A	O3'-P	-6.02	1.53	1.61
34	BA	1577	U	N3-C4	-6.02	1.33	1.38
35	BB	7	C	N3-C4	-6.02	1.29	1.33
35	BB	612	A	C5-C4	-6.02	1.34	1.38
35	BB	1213	U	C4'-O4'	-6.02	1.37	1.45
36	BC	106	G	C3'-C2'	-6.02	1.46	1.52
73	Bn	39	TYR	C-N	-6.02	1.22	1.34
85	AA	165	C	C1'-N1	-6.02	1.38	1.46
85	AA	709	A	C6-N6	-6.02	1.29	1.33
85	AA	739	C	O3'-P	-6.02	1.53	1.61
85	AA	1825	A	O3'-P	-6.02	1.53	1.61
34	BA	165	C	C2'-C1'	-6.02	1.46	1.53
34	BA	617	G	C1'-N9	-6.02	1.38	1.46
34	BA	688	G	C2-N2	-6.02	1.28	1.34
34	BA	1011	G	C5-C6	-6.02	1.36	1.42
34	BA	1531	G	N1-C2	-6.02	1.32	1.37
35	BB	546	A	O3'-P	-6.02	1.53	1.61
35	BB	702	G	C2'-C1'	-6.02	1.46	1.53
39	BF	29	U	C3'-C2'	-6.02	1.46	1.52
41	BH	121	A	C4'-C3'	-6.02	1.46	1.53
85	AA	286	C	O3'-P	-6.02	1.53	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1482	C	C3'-C2'	-6.02	1.46	1.52
85	AA	1598	A	C2'-C1'	-6.02	1.46	1.53
34	BA	301	U	C4'-C3'	-6.02	1.46	1.53
34	BA	458	G	C3'-C2'	-6.02	1.46	1.52
34	BA	542	A	N3-C4	-6.02	1.31	1.34
34	BA	633	G	C2'-C1'	-6.02	1.46	1.53
34	BA	858	C	C2-N3	-6.02	1.30	1.35
34	BA	992	A	P-O5'	-6.02	1.53	1.59
34	BA	1109	G	C6-N1	-6.02	1.35	1.39
34	BA	1660	A	N3-C4	-6.02	1.31	1.34
34	BA	1719	G	O3'-P	-6.02	1.53	1.61
35	BB	583	G	C2-N2	-6.02	1.28	1.34
38	BE	129	G	N3-C4	-6.02	1.31	1.35
85	AA	293	A	C3'-C2'	-6.02	1.46	1.52
85	AA	427	G	O3'-P	-6.02	1.53	1.61
85	AA	1242	A	N9-C4	-6.02	1.34	1.37
85	AA	1984	A	C2'-C1'	-6.02	1.46	1.53
34	BA	312	U	C3'-C2'	-6.02	1.46	1.52
34	BA	1249	G	P-O5'	-6.02	1.53	1.59
34	BA	1527	G	C5-C4	-6.02	1.34	1.38
35	BB	134	G	N9-C4	-6.02	1.33	1.38
36	BC	155	C	C4'-C3'	-6.02	1.46	1.53
37	BD	48	G	C4'-O4'	-6.02	1.37	1.45
38	BE	36	U	O3'-P	-6.02	1.53	1.61
41	BH	135	U	N3-C4	-6.02	1.33	1.38
85	AA	87	C	P-O5'	-6.02	1.53	1.59
85	AA	537	G	N9-C4	-6.02	1.33	1.38
85	AA	1060	U	O3'-P	-6.02	1.53	1.61
85	AA	1106	A	N3-C4	-6.02	1.31	1.34
85	AA	2142	A	C3'-C2'	-6.02	1.46	1.52
35	BB	116	G	O3'-P	-6.02	1.53	1.61
35	BB	663	G	C4'-C3'	-6.02	1.46	1.53
37	BD	93	G	C5-C4	-6.02	1.34	1.38
38	BE	19	G	C2-N3	-6.02	1.27	1.32
85	AA	76	G	C4'-C3'	-6.02	1.46	1.53
85	AA	640	C	C3'-C2'	-6.02	1.46	1.52
85	AA	896	C	O3'-P	-6.02	1.53	1.61
85	AA	1967	A	P-O5'	-6.02	1.53	1.59
85	AA	1978	G	C2'-C1'	-6.02	1.46	1.53
34	BA	446	U	C4'-C3'	-6.01	1.46	1.53
34	BA	625	U	C2-N3	-6.01	1.33	1.37
34	BA	747	G	C1'-N9	-6.01	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1321	A	C3'-C2'	-6.01	1.46	1.52
35	BB	370	A	C4'-C3'	-6.01	1.46	1.53
35	BB	704	G	O3'-P	-6.01	1.53	1.61
35	BB	777	C	C4-C5	-6.01	1.38	1.43
35	BB	1164	U	C2-N3	-6.01	1.33	1.37
35	BB	1380	G	N3-C4	-6.01	1.31	1.35
35	BB	1468	A	C3'-C2'	-6.01	1.46	1.52
35	BB	1513	U	C2-N3	-6.01	1.33	1.37
36	BC	9	G	C2-N2	-6.01	1.28	1.34
37	BD	60	C	C3'-C2'	-6.01	1.46	1.52
38	BE	50	G	N9-C8	-6.01	1.33	1.37
38	BE	85	G	O3'-P	-6.01	1.53	1.61
40	BG	62	C	O3'-P	-6.01	1.53	1.61
41	BH	106	G	P-O5'	-6.01	1.53	1.59
85	AA	362	G	N9-C4	-6.01	1.33	1.38
85	AA	423	G	C2-N2	-6.01	1.28	1.34
34	BA	1533	G	N7-C5	-6.01	1.35	1.39
35	BB	29	C	C4'-C3'	6.01	1.59	1.53
38	BE	138	U	P-O5'	-6.01	1.53	1.59
85	AA	450	A	C2'-C1'	-6.01	1.46	1.53
85	AA	869	A	N1-C2	-6.01	1.28	1.34
85	AA	1611	A	C4'-C3'	-6.01	1.46	1.53
85	AA	2008	G	C5-C4	-6.01	1.34	1.38
34	BA	106	U	N1-C2	-6.01	1.33	1.38
34	BA	329	G	C5-C4	-6.01	1.34	1.38
34	BA	1338	G	C2-N2	-6.01	1.28	1.34
34	BA	1482	A	O3'-P	-6.01	1.53	1.61
34	BA	1671	A	C6-N1	-6.01	1.31	1.35
35	BB	1102	U	N1-C2	-6.01	1.33	1.38
35	BB	1264	U	C3'-C2'	-6.01	1.46	1.52
85	AA	1883	C	N1-C6	-6.01	1.33	1.37
34	BA	734	G	O3'-P	-6.01	1.53	1.61
34	BA	1344	G	O3'-P	-6.01	1.53	1.61
34	BA	1833	G	C2-N2	-6.01	1.28	1.34
35	BB	22	A	C1'-N9	-6.01	1.38	1.46
35	BB	660	G	C2'-C1'	-6.01	1.46	1.53
35	BB	700	C	C2'-C1'	-6.01	1.46	1.53
38	BE	161	G	N7-C5	-6.01	1.35	1.39
40	BG	58	G	O4'-C1'	-6.01	1.33	1.41
40	BG	153	C	C2-N3	-6.01	1.30	1.35
85	AA	100	A	C5-C4	-6.01	1.34	1.38
85	AA	995	G	C2-N3	-6.01	1.27	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1270	C	C2'-C1'	-6.01	1.46	1.53
85	AA	2097	U	O3'-P	-6.01	1.53	1.61
34	BA	27	G	C2-N2	-6.01	1.28	1.34
34	BA	1276	G	N7-C5	-6.01	1.35	1.39
35	BB	1497	C	C2-N3	-6.01	1.30	1.35
36	BC	66	G	C6-N1	-6.01	1.35	1.39
38	BE	10	G	C3'-C2'	6.01	1.59	1.52
85	AA	18	C	C2'-C1'	-6.01	1.46	1.53
86	AB	4	C	O3'-P	-6.01	1.53	1.61
34	BA	65	A	C5'-C4'	6.01	1.58	1.51
34	BA	191	G	C4'-C3'	-6.01	1.46	1.53
34	BA	1723	U	C2-N3	-6.01	1.33	1.37
35	BB	517	G	C6-N1	-6.01	1.35	1.39
35	BB	619	A	P-O5'	-6.01	1.53	1.59
35	BB	1230	A	O3'-P	-6.01	1.53	1.61
85	AA	1150	G	N9-C8	-6.01	1.33	1.37
34	BA	116	G	C1'-N9	-6.00	1.38	1.46
34	BA	766	A	C6-N1	-6.00	1.31	1.35
34	BA	1500	G	C2-N3	6.00	1.37	1.32
35	BB	986	C	O3'-P	-6.00	1.53	1.61
35	BB	1480	G	P-O5'	-6.00	1.53	1.59
85	AA	662	U	C2-N3	-6.00	1.33	1.37
85	AA	703	U	C2'-C1'	-6.00	1.46	1.53
85	AA	1127	G	O4'-C1'	-6.00	1.33	1.41
85	AA	2100	A	P-O5'	-6.00	1.53	1.59
85	AA	2104	C	O3'-P	-6.00	1.53	1.61
34	BA	436	U	C2-N3	-6.00	1.33	1.37
34	BA	599	U	N3-C4	-6.00	1.33	1.38
34	BA	685	C	C2'-C1'	-6.00	1.46	1.53
34	BA	910	U	P-O5'	-6.00	1.53	1.59
34	BA	962	U	C3'-C2'	-6.00	1.46	1.52
34	BA	1174	A	C3'-C2'	-6.00	1.46	1.52
34	BA	1506	C	O3'-P	-6.00	1.53	1.61
34	BA	1550	G	O4'-C1'	-6.00	1.33	1.41
34	BA	1604	A	P-O5'	-6.00	1.53	1.59
35	BB	593	A	N7-C5	-6.00	1.35	1.39
35	BB	1456	G	N7-C5	-6.00	1.35	1.39
35	BB	1511	U	P-O5'	-6.00	1.53	1.59
36	BC	2	A	N9-C8	-6.00	1.32	1.37
40	BG	5	G	C2-N2	-6.00	1.28	1.34
85	AA	408	C	N3-C4	-6.00	1.29	1.33
85	AA	994	A	N9-C4	-6.00	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1665	G	P-O5'	-6.00	1.53	1.59
85	AA	2171	A	C3'-C2'	-6.00	1.46	1.52
34	BA	395	G	C3'-C2'	-6.00	1.46	1.52
34	BA	695	A	C1'-N9	-6.00	1.38	1.46
34	BA	718	U	C3'-C2'	-6.00	1.46	1.52
34	BA	1035	A	P-O5'	-6.00	1.53	1.59
34	BA	1233	U	C2-N3	-6.00	1.33	1.37
34	BA	1412	G	C2'-C1'	-6.00	1.46	1.53
34	BA	1438	C	C2-N3	-6.00	1.30	1.35
35	BB	408	U	C4'-O4'	-6.00	1.37	1.45
35	BB	1239	A	C1'-N9	-6.00	1.38	1.46
40	BG	119	A	O3'-P	-6.00	1.53	1.61
85	AA	396	U	C4'-O4'	-6.00	1.37	1.45
85	AA	666	A	O3'-P	-6.00	1.53	1.61
85	AA	1251	G	N7-C5	-6.00	1.35	1.39
85	AA	1262	A	C1'-N9	-6.00	1.38	1.46
85	AA	1472	G	O3'-P	-6.00	1.53	1.61
85	AA	1482	C	C2-N3	-6.00	1.30	1.35
34	BA	87	G	C6-N1	-6.00	1.35	1.39
34	BA	196	A	C8-N7	-6.00	1.27	1.31
34	BA	1532	G	C3'-C2'	-6.00	1.46	1.52
85	AA	446	C	C2-N3	-6.00	1.30	1.35
85	AA	989	U	O5'-C5'	-6.00	1.33	1.42
85	AA	1434	U	O3'-P	-6.00	1.53	1.61
34	BA	176	G	C3'-C2'	-6.00	1.46	1.52
34	BA	544	U	C3'-C2'	-6.00	1.46	1.52
34	BA	697	A	C3'-C2'	-6.00	1.46	1.52
34	BA	787	A	C2'-C1'	-6.00	1.46	1.53
35	BB	117	A	P-O5'	-6.00	1.53	1.59
35	BB	692	G	N1-C2	-6.00	1.32	1.37
35	BB	1308	G	O3'-P	-6.00	1.53	1.61
35	BB	1494	G	C3'-C2'	-6.00	1.46	1.52
35	BB	1510	G	C2'-C1'	-6.00	1.46	1.53
36	BC	54	G	N7-C5	-6.00	1.35	1.39
36	BC	153	C	C2-N3	-6.00	1.30	1.35
85	AA	296	A	C3'-C2'	-6.00	1.46	1.52
85	AA	1506	U	C4'-O4'	-6.00	1.37	1.45
34	BA	243	C	C2-N3	-6.00	1.30	1.35
34	BA	572	G	C3'-C2'	6.00	1.59	1.52
34	BA	972	C	C2'-C1'	-6.00	1.46	1.53
34	BA	1000	G	C5-C4	-6.00	1.34	1.38
34	BA	1163	G	C2-N2	-6.00	1.28	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1532	G	C5-C4	-6.00	1.34	1.38
35	BB	21	C	C3'-C2'	-6.00	1.46	1.52
35	BB	775	U	C2-N3	-6.00	1.33	1.37
35	BB	896	C	P-O5'	-6.00	1.53	1.59
35	BB	1015	U	N3-C4	-6.00	1.33	1.38
35	BB	1207	C	C2-N3	-6.00	1.30	1.35
35	BB	1319	U	O3'-P	-6.00	1.53	1.61
39	BF	23	G	N9-C8	-6.00	1.33	1.37
40	BG	67	A	C4'-C3'	-6.00	1.46	1.52
40	BG	136	G	N9-C4	-6.00	1.33	1.38
85	AA	258	G	P-O5'	-6.00	1.53	1.59
85	AA	636	G	N7-C5	-6.00	1.35	1.39
85	AA	645	C	C2'-C1'	-6.00	1.46	1.53
85	AA	1925	A	C4'-C3'	-6.00	1.46	1.52
34	BA	590	U	O3'-P	-6.00	1.53	1.61
35	BB	626	C	C2'-C1'	-6.00	1.46	1.53
35	BB	1099	U	C2'-C1'	-6.00	1.46	1.53
36	BC	157	U	C2'-C1'	-6.00	1.46	1.53
37	BD	110	G	C8-N7	-6.00	1.27	1.30
40	BG	75	C	N1-C6	-6.00	1.33	1.37
85	AA	539	A	C8-N7	-6.00	1.27	1.31
85	AA	1107	A	N9-C4	6.00	1.41	1.37
85	AA	2204	A	C5-C4	-6.00	1.34	1.38
34	BA	73	G	C4'-O4'	-5.99	1.37	1.45
34	BA	85	C	P-O5'	-5.99	1.53	1.59
34	BA	127	U	O3'-P	-5.99	1.53	1.61
34	BA	492	G	O3'-P	-5.99	1.53	1.61
34	BA	755	G	C1'-N9	-5.99	1.38	1.46
34	BA	1299	G	C2-N3	-5.99	1.27	1.32
34	BA	1530	G	N3-C4	-5.99	1.31	1.35
35	BB	622	G	N3-C4	-5.99	1.31	1.35
35	BB	690	C	C4'-C3'	-5.99	1.46	1.52
36	BC	69	U	C2-N3	-5.99	1.33	1.37
38	BE	193	A	P-O5'	-5.99	1.53	1.59
40	BG	31	G	N3-C4	-5.99	1.31	1.35
40	BG	35	G	N7-C5	-5.99	1.35	1.39
41	BH	115	A	N7-C5	-5.99	1.35	1.39
85	AA	5	U	N3-C4	-5.99	1.33	1.38
85	AA	1166	C	C3'-C2'	-5.99	1.46	1.52
34	BA	10	G	O4'-C1'	-5.99	1.33	1.41
34	BA	41	U	C4'-C3'	-5.99	1.46	1.52
34	BA	694	G	N1-C2	-5.99	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	731	A	O3'-P	-5.99	1.53	1.61
34	BA	1466	U	N1-C2	-5.99	1.33	1.38
35	BB	631	G	C5-C6	-5.99	1.36	1.42
35	BB	1045	G	N3-C4	-5.99	1.31	1.35
35	BB	1220	A	C5-C6	-5.99	1.35	1.41
41	BH	48	G	P-O5'	-5.99	1.53	1.59
85	AA	717	G	O3'-P	-5.99	1.53	1.61
34	BA	62	A	C1'-N9	-5.99	1.38	1.46
34	BA	147	U	C4'-C3'	-5.99	1.46	1.52
35	BB	988	G	N9-C8	-5.99	1.33	1.37
35	BB	1049	G	C3'-C2'	-5.99	1.46	1.52
40	BG	158	A	N3-C4	-5.99	1.31	1.34
85	AA	162	A	C4'-C3'	-5.99	1.46	1.52
85	AA	540	A	N9-C4	-5.99	1.34	1.37
85	AA	650	G	N9-C4	-5.99	1.33	1.38
85	AA	914	U	O3'-P	-5.99	1.53	1.61
34	BA	64	A	C4'-O4'	-5.99	1.37	1.45
34	BA	280	A	O3'-P	-5.99	1.53	1.61
34	BA	736	G	O4'-C1'	-5.99	1.33	1.41
34	BA	807	U	O4'-C1'	-5.99	1.33	1.41
34	BA	1199	U	P-O5'	-5.99	1.53	1.59
34	BA	1592	U	C2'-C1'	-5.99	1.46	1.53
34	BA	1609	U	C1'-N1	-5.99	1.38	1.46
34	BA	1665	G	N7-C5	-5.99	1.35	1.39
35	BB	490	G	C5-C6	-5.99	1.36	1.42
35	BB	557	C	C2'-C1'	-5.99	1.46	1.53
35	BB	814	A	O3'-P	-5.99	1.53	1.61
35	BB	1353	G	C3'-C2'	-5.99	1.46	1.52
36	BC	46	G	C6-N1	-5.99	1.35	1.39
36	BC	71	A	C3'-C2'	-5.99	1.46	1.52
40	BG	165	C	C2-N3	-5.99	1.30	1.35
41	BH	7	C	C2'-C1'	-5.99	1.46	1.53
85	AA	393	C	P-O5'	-5.99	1.53	1.59
85	AA	418	G	C2'-C1'	-5.99	1.46	1.53
34	BA	425	G	C2'-C1'	-5.99	1.46	1.53
34	BA	1719	G	C5-C4	-5.99	1.34	1.38
35	BB	97	U	P-O5'	-5.99	1.53	1.59
35	BB	1154	C	P-O5'	-5.99	1.53	1.59
36	BC	123	G	C2-N3	-5.99	1.27	1.32
40	BG	3	G	N9-C4	-5.99	1.33	1.38
34	BA	378	C	C4-N4	-5.99	1.28	1.33
34	BA	790	G	C8-N7	-5.99	1.27	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1634	A	C6-N1	-5.99	1.31	1.35
34	BA	1668	C	C4'-O4'	-5.99	1.37	1.45
35	BB	522	A	C4'-O4'	-5.99	1.37	1.45
35	BB	1197	G	C2'-C1'	-5.99	1.46	1.53
37	BD	99	G	C2'-C1'	-5.99	1.46	1.53
38	BE	72	C	C2'-C1'	-5.99	1.46	1.53
85	AA	109	G	N9-C4	5.99	1.42	1.38
85	AA	2205	A	C2'-C1'	-5.99	1.46	1.53
34	BA	25	C	O3'-P	-5.98	1.53	1.61
34	BA	1295	U	C3'-C2'	-5.98	1.46	1.52
35	BB	658	G	C5-C4	-5.98	1.34	1.38
39	BF	11	C	P-O5'	5.98	1.65	1.59
41	BH	72	G	C3'-C2'	-5.98	1.46	1.52
85	AA	275	A	C2'-C1'	-5.98	1.46	1.53
85	AA	315	U	P-O5'	-5.98	1.53	1.59
85	AA	869	A	C1'-N9	-5.98	1.38	1.46
85	AA	1852	U	C4'-C3'	5.98	1.59	1.53
85	AA	2196	G	C1'-N9	-5.98	1.38	1.46
86	AB	68	C	N1-C6	-5.98	1.33	1.37
34	BA	233	U	P-O5'	-5.98	1.53	1.59
34	BA	373	G	N1-C2	-5.98	1.32	1.37
34	BA	568	G	C2'-C1'	-5.98	1.46	1.53
34	BA	615	A	N3-C4	-5.98	1.31	1.34
34	BA	699	G	C2'-C1'	-5.98	1.46	1.53
34	BA	789	U	C2-N3	-5.98	1.33	1.37
34	BA	1075	U	C2-N3	-5.98	1.33	1.37
34	BA	1289	C	C2-N3	-5.98	1.30	1.35
34	BA	1544	G	C5-C4	-5.98	1.34	1.38
34	BA	1585	A	O4'-C1'	-5.98	1.33	1.41
35	BB	94	A	C2'-C1'	-5.98	1.46	1.53
35	BB	826	G	N7-C5	-5.98	1.35	1.39
35	BB	1354	C	C3'-C2'	-5.98	1.46	1.52
35	BB	1534	U	O3'-P	-5.98	1.53	1.61
40	BG	116	G	N7-C5	-5.98	1.35	1.39
85	AA	1277	C	O4'-C1'	-5.98	1.33	1.41
85	AA	1437	G	C1'-N9	-5.98	1.38	1.46
85	AA	2204	A	C3'-C2'	-5.98	1.46	1.52
34	BA	371	U	C4'-O4'	-5.98	1.37	1.45
34	BA	710	A	C1'-N9	-5.98	1.38	1.46
34	BA	1022	C	C1'-N1	-5.98	1.38	1.46
34	BA	1507	C	C2-N3	-5.98	1.30	1.35
35	BB	1145	G	N3-C4	-5.98	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	118	U	O3'-P	-5.98	1.53	1.61
37	BD	28	C	O3'-P	-5.98	1.53	1.61
38	BE	192	A	C6-N1	-5.98	1.31	1.35
40	BG	115	C	P-O5'	-5.98	1.53	1.59
85	AA	885	A	N9-C4	5.98	1.41	1.37
85	AA	1580	A	N9-C4	-5.98	1.34	1.37
85	AA	1753	A	C3'-O3'	5.98	1.50	1.42
85	AA	2058	C	C2-N3	-5.98	1.30	1.35
34	BA	44	U	C4'-O4'	-5.98	1.37	1.45
34	BA	1431	G	N9-C8	-5.98	1.33	1.37
35	BB	505	G	N9-C4	5.98	1.42	1.38
35	BB	1172	U	C4'-O4'	-5.98	1.37	1.45
35	BB	1274	G	N3-C4	-5.98	1.31	1.35
38	BE	147	G	N9-C8	-5.98	1.33	1.37
38	BE	158	U	C2-N3	-5.98	1.33	1.37
41	BH	4	U	N1-C6	-5.98	1.32	1.38
71	Bl	39	TYR	CB-CG	-5.98	1.42	1.51
85	AA	379	U	C1'-N1	-5.98	1.38	1.46
85	AA	413	G	C5-C4	-5.98	1.34	1.38
85	AA	1669	G	C5-C4	-5.98	1.34	1.38
34	BA	228	A	P-O5'	-5.98	1.53	1.59
34	BA	559	C	N1-C6	-5.98	1.33	1.37
34	BA	783	U	N3-C4	-5.98	1.33	1.38
34	BA	909	G	N1-C2	-5.98	1.32	1.37
34	BA	1036	G	C5-C6	-5.98	1.36	1.42
34	BA	1333	G	N9-C4	-5.98	1.33	1.38
34	BA	1839	G	C1'-N9	-5.98	1.38	1.46
35	BB	791	A	N7-C5	-5.98	1.35	1.39
35	BB	1202	G	C4'-C3'	-5.98	1.46	1.52
35	BB	1323	U	O3'-P	-5.98	1.53	1.61
35	BB	1427	A	C1'-N9	-5.98	1.38	1.46
35	BB	1479	C	C2'-C1'	-5.98	1.46	1.53
37	BD	7	G	C2'-C1'	-5.98	1.46	1.53
40	BG	34	A	C2'-C1'	-5.98	1.46	1.53
41	BH	15	A	C2'-C1'	-5.98	1.46	1.53
85	AA	48	G	C2'-C1'	-5.98	1.46	1.53
85	AA	760	U	O3'-P	-5.98	1.53	1.61
85	AA	2208	G	C5'-C4'	-5.98	1.44	1.51
34	BA	1508	C	N3-C4	-5.98	1.29	1.33
34	BA	1552	C	N1-C6	-5.98	1.33	1.37
35	BB	424	U	O3'-P	-5.98	1.53	1.61
34	BA	28	C	C3'-C2'	-5.97	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	57	A	C1'-N9	-5.97	1.38	1.46
34	BA	118	C	C2-N3	-5.97	1.30	1.35
34	BA	1000	G	C4'-C3'	-5.97	1.46	1.52
34	BA	1196	C	P-O5'	-5.97	1.53	1.59
35	BB	408	U	C4'-C3'	-5.97	1.46	1.52
35	BB	1057	G	C5-C4	-5.97	1.34	1.38
35	BB	1120	A	C3'-C2'	-5.97	1.46	1.52
35	BB	1177	U	C2-N3	-5.97	1.33	1.37
35	BB	1183	U	P-O5'	-5.97	1.53	1.59
35	BB	1258	G	C6-N1	-5.97	1.35	1.39
36	BC	16	A	C4'-C3'	-5.97	1.46	1.52
37	BD	111	U	N3-C4	-5.97	1.33	1.38
40	BG	46	G	C1'-N9	-5.97	1.38	1.46
85	AA	311	U	C3'-C2'	-5.97	1.46	1.52
85	AA	461	G	O3'-P	-5.97	1.53	1.61
85	AA	976	G	O3'-P	-5.97	1.53	1.61
85	AA	976	G	N9-C4	-5.97	1.33	1.38
85	AA	1458	G	O3'-P	-5.97	1.53	1.61
85	AA	1598	A	C4'-C3'	-5.97	1.46	1.52
34	BA	234	A	C5'-C4'	-5.97	1.44	1.51
34	BA	332	U	O3'-P	-5.97	1.53	1.61
34	BA	1516	G	N9-C4	-5.97	1.33	1.38
34	BA	1588	U	C3'-C2'	-5.97	1.46	1.52
34	BA	1596	C	O3'-P	-5.97	1.53	1.61
35	BB	386	G	N7-C5	-5.97	1.35	1.39
35	BB	464	C	C2'-C1'	-5.97	1.46	1.53
35	BB	483	C	C2'-C1'	-5.97	1.46	1.53
35	BB	1525	G	P-O5'	-5.97	1.53	1.59
36	BC	67	U	O4'-C1'	-5.97	1.33	1.41
36	BC	86	U	N1-C2	5.97	1.44	1.38
37	BD	25	G	N7-C5	-5.97	1.35	1.39
38	BE	136	G	P-O5'	-5.97	1.53	1.59
38	BE	140	G	N1-C2	-5.97	1.32	1.37
38	BE	170	U	O3'-P	-5.97	1.53	1.61
40	BG	60	A	O3'-P	-5.97	1.53	1.61
40	BG	171	A	O3'-P	-5.97	1.53	1.61
85	AA	130	G	C5'-C4'	-5.97	1.44	1.51
85	AA	312	G	C8-N7	-5.97	1.27	1.30
85	AA	1325	C	O3'-P	-5.97	1.53	1.61
85	AA	1527	G	C2'-C1'	-5.97	1.46	1.53
34	BA	1112	U	O3'-P	-5.97	1.53	1.61
34	BA	1134	A	C2'-C1'	-5.97	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1486	U	C2-N3	-5.97	1.33	1.37
34	BA	1644	A	C8-N7	-5.97	1.27	1.31
35	BB	424	U	P-O5'	-5.97	1.53	1.59
35	BB	1335	G	N1-C2	-5.97	1.32	1.37
85	AA	147	G	P-O5'	-5.97	1.53	1.59
85	AA	1707	G	O3'-P	-5.97	1.53	1.61
34	BA	257	G	C8-N7	-5.97	1.27	1.30
34	BA	448	U	O3'-P	-5.97	1.53	1.61
34	BA	540	G	C5-C4	-5.97	1.34	1.38
34	BA	953	G	N9-C4	-5.97	1.33	1.38
34	BA	1203	G	C1'-N9	-5.97	1.38	1.46
34	BA	1435	A	C5-C4	-5.97	1.34	1.38
34	BA	1827	C	C3'-C2'	-5.97	1.46	1.52
35	BB	121	A	N9-C8	-5.97	1.32	1.37
35	BB	514	G	N1-C2	-5.97	1.32	1.37
35	BB	712	U	O3'-P	-5.97	1.53	1.61
36	BC	71	A	N9-C4	-5.97	1.34	1.37
37	BD	84	U	N3-C4	-5.97	1.33	1.38
85	AA	714	U	N3-C4	-5.97	1.33	1.38
85	AA	744	C	C4-C5	-5.97	1.38	1.43
85	AA	807	A	C4'-C3'	-5.97	1.46	1.52
85	AA	1129	A	C2'-C1'	-5.97	1.46	1.53
85	AA	1282	A	P-O5'	-5.97	1.53	1.59
85	AA	1364	U	C2'-C1'	-5.97	1.46	1.53
85	AA	1555	G	C6-N1	-5.97	1.35	1.39
85	AA	1952	C	C2'-C1'	-5.97	1.46	1.53
34	BA	1197	U	N3-C4	-5.97	1.33	1.38
34	BA	1201	G	C2'-C1'	-5.97	1.46	1.53
35	BB	64	U	N1-C2	-5.97	1.33	1.38
35	BB	1147	G	C5'-C4'	-5.97	1.44	1.51
38	BE	192	A	C8-N7	-5.97	1.27	1.31
85	AA	1927	G	C2-N3	-5.97	1.27	1.32
34	BA	344	G	C8-N7	-5.97	1.27	1.30
34	BA	810	A	O3'-P	-5.97	1.53	1.61
34	BA	1122	G	P-O5'	-5.97	1.53	1.59
34	BA	1266	A	N3-C4	-5.97	1.31	1.34
35	BB	522	A	P-O5'	-5.97	1.53	1.59
35	BB	1298	C	N3-C4	-5.97	1.29	1.33
34	BA	527	C	C3'-C2'	-5.96	1.46	1.52
34	BA	1062	G	C5-C6	-5.96	1.36	1.42
34	BA	1260	G	C2-N2	-5.96	1.28	1.34
34	BA	1643	U	P-O5'	-5.96	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1682	A	C2'-C1'	-5.96	1.46	1.53
34	BA	1687	A	C4'-O4'	-5.96	1.37	1.45
35	BB	65	A	C2'-C1'	-5.96	1.46	1.53
35	BB	93	A	C5-C6	-5.96	1.35	1.41
35	BB	571	C	C1'-N1	-5.96	1.38	1.46
35	BB	628	A	C6-N1	-5.96	1.31	1.35
36	BC	18	G	P-O5'	-5.96	1.53	1.59
38	BE	17	U	C1'-N1	-5.96	1.38	1.46
38	BE	96	G	C6-N1	-5.96	1.35	1.39
40	BG	93	U	N3-C4	-5.96	1.33	1.38
85	AA	65	A	N9-C8	-5.96	1.32	1.37
85	AA	74	U	O3'-P	-5.96	1.53	1.61
85	AA	359	A	C3'-C2'	-5.96	1.46	1.52
85	AA	483	G	N9-C8	-5.96	1.33	1.37
85	AA	519	A	C5-C4	-5.96	1.34	1.38
85	AA	931	G	C2-N2	-5.96	1.28	1.34
85	AA	1560	A	P-O5'	-5.96	1.53	1.59
85	AA	1933	G	C4'-C3'	5.96	1.59	1.53
85	AA	2039	G	C2'-C1'	-5.96	1.46	1.53
35	BB	71	A	C3'-C2'	-5.96	1.46	1.52
35	BB	87	G	N7-C5	-5.96	1.35	1.39
35	BB	833	G	C2-N3	-5.96	1.27	1.32
40	BG	19	C	C3'-C2'	-5.96	1.46	1.52
41	BH	105	U	C4'-C3'	-5.96	1.46	1.52
85	AA	660	G	O3'-P	-5.96	1.53	1.61
34	BA	684	G	C4'-C3'	5.96	1.59	1.53
34	BA	731	A	N9-C4	-5.96	1.34	1.37
34	BA	877	U	C2'-C1'	-5.96	1.46	1.53
34	BA	1363	A	N9-C4	-5.96	1.34	1.37
34	BA	1540	C	P-O5'	-5.96	1.53	1.59
34	BA	1668	C	O4'-C1'	-5.96	1.33	1.41
35	BB	637	G	P-O5'	-5.96	1.53	1.59
35	BB	1228	A	C2'-C1'	-5.96	1.46	1.53
35	BB	1413	U	N3-C4	-5.96	1.33	1.38
85	AA	252	G	N9-C4	-5.96	1.33	1.38
85	AA	574	U	P-O5'	-5.96	1.53	1.59
85	AA	684	G	N1-C2	-5.96	1.32	1.37
85	AA	778	C	C3'-C2'	-5.96	1.46	1.52
85	AA	811	A	C5-C4	-5.96	1.34	1.38
85	AA	1992	A	C6-N1	-5.96	1.31	1.35
34	BA	622	G	P-O5'	-5.96	1.53	1.59
34	BA	629	G	C6-N1	-5.96	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1838	U	C3'-C2'	-5.96	1.46	1.52
35	BB	607	G	C6-N1	-5.96	1.35	1.39
35	BB	772	U	O3'-P	-5.96	1.53	1.61
37	BD	27	A	C3'-C2'	-5.96	1.46	1.52
85	AA	185	A	C8-N7	-5.96	1.27	1.31
85	AA	375	C	C2-N3	-5.96	1.30	1.35
85	AA	963	U	C3'-C2'	-5.96	1.46	1.52
85	AA	1108	U	O3'-P	-5.96	1.53	1.61
85	AA	2132	A	C5-C4	-5.96	1.34	1.38
34	BA	103	G	N9-C8	-5.96	1.33	1.37
34	BA	603	U	C4-C5	-5.96	1.38	1.43
34	BA	1072	U	C2'-C1'	-5.96	1.46	1.53
34	BA	1148	U	C2-N3	-5.96	1.33	1.37
34	BA	1452	U	C2-N3	-5.96	1.33	1.37
34	BA	1519	G	C5-C4	-5.96	1.34	1.38
35	BB	74	U	C2-N3	-5.96	1.33	1.37
35	BB	1092	G	C3'-C2'	-5.96	1.46	1.52
36	BC	145	G	C1'-N9	-5.96	1.38	1.46
85	AA	159	G	C5-C4	-5.96	1.34	1.38
85	AA	868	A	C3'-C2'	-5.96	1.46	1.52
34	BA	1252	G	N9-C8	-5.96	1.33	1.37
35	BB	118	A	C5-C4	-5.96	1.34	1.38
35	BB	567	G	C2-N2	-5.96	1.28	1.34
35	BB	1061	G	N3-C4	-5.96	1.31	1.35
36	BC	116	C	N1-C6	-5.96	1.33	1.37
38	BE	64	A	C5-C4	-5.96	1.34	1.38
40	BG	148	C	O4'-C1'	-5.96	1.33	1.41
41	BH	68	G	N7-C5	-5.96	1.35	1.39
85	AA	1510	A	C3'-C2'	-5.96	1.46	1.52
85	AA	1622	G	O3'-P	-5.96	1.54	1.61
85	AA	1813	C	O3'-P	-5.96	1.54	1.61
85	AA	1894	G	O4'-C1'	-5.96	1.33	1.41
85	AA	1960	C	C2'-C1'	-5.96	1.46	1.53
85	AA	2086	C	C4'-C3'	-5.96	1.46	1.52
35	BB	648	G	C4'-O4'	-5.96	1.37	1.45
35	BB	1407	U	C2'-C1'	-5.96	1.46	1.53
40	BG	107	U	C1'-N1	-5.96	1.38	1.46
41	BH	131	A	N3-C4	-5.96	1.31	1.34
85	AA	431	G	C2-N3	-5.96	1.27	1.32
85	AA	497	G	C2-N2	-5.96	1.28	1.34
85	AA	1618	G	N9-C4	-5.96	1.33	1.38
34	BA	185	A	N9-C4	-5.95	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	201	A	N9-C4	-5.95	1.34	1.37
34	BA	532	C	C3'-C2'	-5.95	1.46	1.52
34	BA	557	U	C2'-O2'	5.95	1.49	1.41
34	BA	804	G	C5'-C4'	-5.95	1.44	1.51
34	BA	1195	G	N9-C4	-5.95	1.33	1.38
35	BB	1396	G	P-O5'	-5.95	1.53	1.59
35	BB	1437	U	C2'-C1'	-5.95	1.46	1.53
37	BD	5	A	C5-C4	-5.95	1.34	1.38
38	BE	129	G	C5-C6	-5.95	1.36	1.42
38	BE	141	A	C5-C4	-5.95	1.34	1.38
85	AA	333	A	N7-C5	-5.95	1.35	1.39
85	AA	693	A	C6-N1	-5.95	1.31	1.35
85	AA	966	G	O3'-P	-5.95	1.54	1.61
34	BA	146	G	P-O5'	-5.95	1.53	1.59
35	BB	442	U	C4'-O4'	-5.95	1.37	1.45
35	BB	1045	G	C2-N2	-5.95	1.28	1.34
35	BB	1336	G	C2-N2	-5.95	1.28	1.34
38	BE	121	G	P-O5'	-5.95	1.53	1.59
85	AA	412	G	C6-N1	-5.95	1.35	1.39
35	BB	968	C	P-O5'	-5.95	1.53	1.59
35	BB	1144	A	P-O5'	-5.95	1.53	1.59
36	BC	79	A	C1'-N9	-5.95	1.38	1.46
38	BE	46	G	N9-C4	-5.95	1.33	1.38
85	AA	400	G	C2-N3	-5.95	1.27	1.32
85	AA	993	G	N7-C5	-5.95	1.35	1.39
85	AA	1243	G	C5-C4	-5.95	1.34	1.38
34	BA	955	G	C5-C4	-5.95	1.34	1.38
34	BA	1094	U	C1'-N1	-5.95	1.38	1.46
34	BA	1310	C	C4-N4	-5.95	1.28	1.33
35	BB	1212	C	O3'-P	-5.95	1.54	1.61
37	BD	99	G	C5-C6	-5.95	1.36	1.42
85	AA	557	G	C3'-C2'	-5.95	1.46	1.52
85	AA	824	C	P-O5'	-5.95	1.53	1.59
85	AA	1142	G	C3'-C2'	-5.95	1.46	1.52
34	BA	445	C	P-O5'	-5.95	1.53	1.59
35	BB	78	C	P-O5'	-5.95	1.53	1.59
35	BB	580	A	C4'-C3'	5.95	1.59	1.53
36	BC	90	U	C2'-C1'	-5.95	1.46	1.53
85	AA	167	A	C3'-C2'	-5.95	1.46	1.52
85	AA	373	G	C5-C4	-5.95	1.34	1.38
85	AA	1499	G	N9-C8	-5.95	1.33	1.37
85	AA	1619	A	N9-C4	-5.95	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	20	A	C5-C6	-5.95	1.35	1.41
34	BA	609	G	C5-C4	-5.95	1.34	1.38
34	BA	1414	C	O3'-P	-5.95	1.54	1.61
34	BA	1474	G	C5-C4	-5.95	1.34	1.38
35	BB	498	G	O3'-P	-5.95	1.54	1.61
35	BB	678	U	C4'-C3'	-5.95	1.46	1.52
35	BB	802	G	O3'-P	-5.95	1.54	1.61
35	BB	1271	A	C2'-C1'	-5.95	1.46	1.53
36	BC	154	A	C5-C4	-5.95	1.34	1.38
37	BD	44	U	N3-C4	-5.95	1.33	1.38
38	BE	184	G	O3'-P	-5.95	1.54	1.61
85	AA	991	G	C2-N2	-5.95	1.28	1.34
85	AA	1158	U	O3'-P	-5.95	1.54	1.61
34	BA	1640	G	C2-N2	-5.94	1.28	1.34
34	BA	1656	A	C8-N7	-5.94	1.27	1.31
35	BB	1045	G	C3'-C2'	-5.94	1.46	1.52
85	AA	1125	G	N9-C8	-5.94	1.33	1.37
85	AA	1864	G	N3-C4	-5.94	1.31	1.35
34	BA	78	U	C2-N3	-5.94	1.33	1.37
34	BA	106	U	P-O5'	-5.94	1.53	1.59
34	BA	301	U	C3'-C2'	-5.94	1.46	1.52
34	BA	339	G	N9-C4	-5.94	1.33	1.38
34	BA	679	U	C1'-N1	-5.94	1.38	1.46
35	BB	372	U	C3'-C2'	-5.94	1.46	1.52
35	BB	874	G	C2'-C1'	-5.94	1.46	1.53
35	BB	1058	U	N3-C4	-5.94	1.33	1.38
35	BB	1488	G	C5-C4	-5.94	1.34	1.38
41	BH	130	G	P-O5'	-5.94	1.53	1.59
85	AA	873	U	C3'-O3'	5.94	1.50	1.42
85	AA	1448	A	C4'-C3'	-5.94	1.46	1.52
85	AA	1884	A	O3'-P	-5.94	1.54	1.61
34	BA	56	G	N9-C8	-5.94	1.33	1.37
34	BA	440	A	C2'-C1'	-5.94	1.46	1.53
34	BA	737	U	C3'-C2'	-5.94	1.46	1.52
34	BA	961	C	C2-N3	-5.94	1.30	1.35
34	BA	990	G	N1-C2	-5.94	1.32	1.37
34	BA	996	U	N3-C4	-5.94	1.33	1.38
34	BA	1327	G	N9-C8	-5.94	1.33	1.37
34	BA	1768	G	C2'-C1'	-5.94	1.46	1.53
35	BB	450	A	C3'-C2'	-5.94	1.46	1.52
35	BB	496	C	C4'-C3'	-5.94	1.46	1.52
35	BB	1101	C	P-O5'	-5.94	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1469	A	C2'-C1'	-5.94	1.46	1.53
36	BC	36	G	N3-C4	-5.94	1.31	1.35
38	BE	68	U	N3-C4	-5.94	1.33	1.38
85	AA	352	G	N7-C5	-5.94	1.35	1.39
85	AA	1162	A	O3'-P	-5.94	1.54	1.61
85	AA	1227	A	C5-C4	-5.94	1.34	1.38
85	AA	1471	G	C6-O6	-5.94	1.18	1.24
34	BA	114	U	C4'-O4'	-5.94	1.37	1.45
38	BE	130	G	O3'-P	-5.94	1.54	1.61
40	BG	169	A	C8-N7	-5.94	1.27	1.31
85	AA	1369	U	O3'-P	-5.94	1.54	1.61
34	BA	40	A	N9-C4	-5.94	1.34	1.37
34	BA	306	G	O3'-P	-5.94	1.54	1.61
34	BA	741	A	O4'-C1'	-5.94	1.33	1.41
34	BA	786	U	N1-C6	-5.94	1.32	1.38
34	BA	907	A	P-O5'	-5.94	1.53	1.59
34	BA	1098	G	N9-C4	-5.94	1.33	1.38
34	BA	1100	A	O3'-P	-5.94	1.54	1.61
34	BA	1274	A	N3-C4	-5.94	1.31	1.34
35	BB	627	G	C2-N2	-5.94	1.28	1.34
35	BB	1038	G	C5-C4	-5.94	1.34	1.38
35	BB	1384	A	P-O5'	-5.94	1.53	1.59
35	BB	1444	U	C2'-C1'	-5.94	1.46	1.53
36	BC	56	G	C1'-N9	-5.94	1.38	1.46
38	BE	46	G	O4'-C1'	-5.94	1.33	1.41
38	BE	53	U	C2'-C1'	-5.94	1.46	1.53
38	BE	166	G	C3'-C2'	-5.94	1.46	1.52
85	AA	34	G	C6-N1	-5.94	1.35	1.39
85	AA	526	G	C5'-C4'	-5.94	1.44	1.51
85	AA	693	A	C3'-C2'	5.94	1.59	1.52
85	AA	929	G	N7-C5	-5.94	1.35	1.39
85	AA	1523	G	C5-C4	-5.94	1.34	1.38
85	AA	1648	G	O3'-P	-5.94	1.54	1.61
34	BA	905	A	O3'-P	-5.94	1.54	1.61
34	BA	1550	G	C4'-C3'	-5.94	1.46	1.52
35	BB	1161	G	C2-N3	-5.94	1.28	1.32
36	BC	25	C	C4-C5	-5.94	1.38	1.43
85	AA	390	U	C1'-N1	-5.94	1.38	1.46
85	AA	1202	G	C6-N1	-5.94	1.35	1.39
34	BA	433	G	C5-C4	-5.93	1.34	1.38
34	BA	579	U	N1-C2	-5.93	1.33	1.38
34	BA	585	G	C5-C6	-5.93	1.36	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	754	G	C2-N2	-5.93	1.28	1.34
35	BB	103	C	O3'-P	-5.93	1.54	1.61
35	BB	714	U	N3-C4	-5.93	1.33	1.38
35	BB	785	G	N9-C8	-5.93	1.33	1.37
35	BB	812	G	C6-N1	-5.93	1.35	1.39
35	BB	891	U	O3'-P	-5.93	1.54	1.61
35	BB	1050	A	N7-C5	-5.93	1.35	1.39
35	BB	1430	G	N7-C5	-5.93	1.35	1.39
38	BE	107	U	C5'-C4'	5.93	1.58	1.51
85	AA	1859	C	O3'-P	-5.93	1.54	1.61
34	BA	236	A	C4'-O4'	-5.93	1.37	1.45
34	BA	244	A	C2'-C1'	-5.93	1.46	1.53
34	BA	790	G	C6-N1	-5.93	1.35	1.39
34	BA	1472	G	C5-C4	-5.93	1.34	1.38
34	BA	1714	A	C1'-N9	-5.93	1.38	1.46
35	BB	314	A	O3'-P	-5.93	1.54	1.61
35	BB	449	C	N3-C4	-5.93	1.29	1.33
35	BB	1082	A	C5-C4	-5.93	1.34	1.38
35	BB	1426	G	C3'-C2'	-5.93	1.46	1.52
40	BG	138	C	C3'-C2'	-5.93	1.46	1.52
40	BG	152	G	C2'-C1'	-5.93	1.46	1.53
85	AA	378	A	C6-N6	-5.93	1.29	1.33
85	AA	1130	G	N7-C5	-5.93	1.35	1.39
85	AA	1220	A	C4'-C3'	-5.93	1.46	1.52
85	AA	1496	U	O4'-C1'	-5.93	1.33	1.41
85	AA	1544	G	C3'-C2'	-5.93	1.46	1.52
85	AA	1869	U	C2'-C1'	-5.93	1.46	1.53
85	AA	2217	A	C1'-N9	-5.93	1.38	1.46
34	BA	40	A	N3-C4	-5.93	1.31	1.34
34	BA	419	U	N1-C2	-5.93	1.33	1.38
34	BA	493	G	C2-N2	-5.93	1.28	1.34
34	BA	1023	G	C6-N1	-5.93	1.35	1.39
35	BB	439	G	C5-C4	-5.93	1.34	1.38
35	BB	1037	A	N9-C4	-5.93	1.34	1.37
35	BB	1155	U	N3-C4	-5.93	1.33	1.38
38	BE	13	A	C1'-N9	-5.93	1.38	1.46
39	BF	67	A	O3'-P	-5.93	1.54	1.61
85	AA	277	G	C1'-N9	-5.93	1.38	1.46
85	AA	2016	A	C2'-C1'	-5.93	1.46	1.53
34	BA	442	G	N9-C4	-5.93	1.33	1.38
34	BA	956	G	O3'-P	-5.93	1.54	1.61
34	BA	1017	C	C2'-C1'	-5.93	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	497	C	C3'-C2'	-5.93	1.46	1.52
36	BC	154	A	C8-N7	-5.93	1.27	1.31
38	BE	20	C	C4'-C3'	-5.93	1.46	1.52
85	AA	877	G	C5-C4	-5.93	1.34	1.38
34	BA	1542	A	C5-C4	-5.93	1.34	1.38
35	BB	673	C	N1-C6	-5.93	1.33	1.37
35	BB	1220	A	C5-C4	-5.93	1.34	1.38
40	BG	80	G	C1'-N9	-5.93	1.38	1.46
85	AA	90	A	O3'-P	-5.93	1.54	1.61
85	AA	1006	C	C3'-C2'	-5.93	1.46	1.52
86	AB	19	G	C2-N2	-5.93	1.28	1.34
34	BA	148	G	N9-C4	-5.93	1.33	1.38
34	BA	331	G	N7-C5	-5.93	1.35	1.39
34	BA	471	U	C2'-C1'	-5.93	1.46	1.53
34	BA	496	G	C2-N2	-5.93	1.28	1.34
34	BA	626	G	C6-N1	-5.93	1.35	1.39
34	BA	633	G	C3'-C2'	-5.93	1.46	1.52
34	BA	675	C	C4-C5	-5.93	1.38	1.43
35	BB	775	U	O3'-P	-5.93	1.54	1.61
35	BB	1192	C	C4'-C3'	-5.93	1.46	1.52
38	BE	149	A	N9-C8	-5.93	1.33	1.37
40	BG	48	U	C2'-C1'	-5.93	1.46	1.53
85	AA	747	U	C2-N3	-5.93	1.33	1.37
85	AA	1630	U	O3'-P	-5.93	1.54	1.61
85	AA	2129	U	C4'-C3'	-5.93	1.46	1.52
34	BA	187	G	C6-N1	-5.92	1.35	1.39
34	BA	353	U	P-O5'	-5.92	1.53	1.59
34	BA	426	A	N9-C8	-5.92	1.33	1.37
34	BA	745	A	C4'-C3'	-5.92	1.46	1.52
34	BA	757	G	C3'-C2'	-5.92	1.46	1.52
34	BA	992	A	C5-C6	-5.92	1.35	1.41
34	BA	1251	A	C5-C4	-5.92	1.34	1.38
34	BA	1257	U	N1-C2	-5.92	1.33	1.38
34	BA	1542	A	C2'-C1'	-5.92	1.46	1.53
35	BB	79	U	C3'-C2'	-5.92	1.46	1.52
35	BB	789	G	N9-C4	-5.92	1.33	1.38
35	BB	812	G	C1'-N9	-5.92	1.38	1.46
35	BB	1357	C	C2-N3	-5.92	1.31	1.35
37	BD	93	G	C2-N2	-5.92	1.28	1.34
40	BG	25	G	N9-C4	-5.92	1.33	1.38
85	AA	69	C	P-O5'	-5.92	1.53	1.59
85	AA	379	U	C2-N3	-5.92	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1169	A	P-O5'	-5.92	1.53	1.59
85	AA	1695	G	N3-C4	-5.92	1.31	1.35
34	BA	184	C	C5'-C4'	-5.92	1.44	1.51
35	BB	377	A	C1'-N9	-5.92	1.38	1.46
34	BA	140	C	C4'-O4'	-5.92	1.37	1.45
34	BA	678	C	N1-C6	-5.92	1.33	1.37
34	BA	1324	G	N7-C5	-5.92	1.35	1.39
34	BA	1529	G	N7-C5	-5.92	1.35	1.39
34	BA	1654	G	P-O5'	-5.92	1.53	1.59
35	BB	372	U	C4'-C3'	-5.92	1.46	1.52
35	BB	793	A	O3'-P	-5.92	1.54	1.61
35	BB	806	U	C3'-C2'	-5.92	1.46	1.52
35	BB	1442	C	O3'-P	-5.92	1.54	1.61
85	AA	32	U	N1-C2	-5.92	1.33	1.38
85	AA	562	C	P-O5'	-5.92	1.53	1.59
85	AA	631	G	C1'-N9	-5.92	1.38	1.46
34	BA	1097	G	C6-N1	-5.92	1.35	1.39
34	BA	162	G	N3-C4	-5.92	1.31	1.35
34	BA	939	C	N3-C4	-5.92	1.29	1.33
34	BA	1087	A	C1'-N9	-5.92	1.38	1.46
34	BA	1520	A	C1'-N9	-5.92	1.38	1.46
34	BA	1627	U	C5'-C4'	5.92	1.58	1.51
34	BA	1655	G	C2-N2	-5.92	1.28	1.34
35	BB	480	C	O3'-P	-5.92	1.54	1.61
35	BB	618	U	C1'-N1	-5.92	1.38	1.46
35	BB	1052	G	C5-C6	-5.92	1.36	1.42
35	BB	1199	A	C4'-O4'	-5.92	1.37	1.45
35	BB	1238	A	P-O5'	-5.92	1.53	1.59
37	BD	94	C	C3'-C2'	-5.92	1.46	1.52
41	BH	41	A	C8-N7	-5.92	1.27	1.31
83	Bx	47	PHE	C-N	5.92	1.43	1.33
85	AA	194	U	C3'-C2'	-5.92	1.46	1.52
85	AA	739	C	C3'-C2'	-5.92	1.46	1.52
85	AA	1533	C	O3'-P	-5.92	1.54	1.61
85	AA	1541	G	C5-C4	-5.92	1.34	1.38
85	AA	2007	G	C6-N1	-5.92	1.35	1.39
34	BA	208	A	N9-C4	5.92	1.41	1.37
34	BA	936	A	C5-C4	-5.92	1.34	1.38
34	BA	1114	G	N7-C5	-5.92	1.35	1.39
34	BA	1780	U	C2'-C1'	-5.92	1.46	1.53
35	BB	35	G	C3'-C2'	-5.92	1.46	1.52
35	BB	596	C	C1'-N1	-5.92	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	664	A	C4'-O4'	-5.92	1.37	1.45
35	BB	1526	C	P-O5'	-5.92	1.53	1.59
37	BD	118	C	P-O5'	-5.92	1.53	1.59
39	BF	17	U	C2-N3	-5.92	1.33	1.37
41	BH	130	G	N7-C5	-5.92	1.35	1.39
85	AA	548	G	O3'-P	-5.92	1.54	1.61
85	AA	782	G	C2'-C1'	-5.92	1.46	1.53
34	BA	421	G	C2'-C1'	-5.92	1.46	1.53
34	BA	755	G	C2-N3	-5.92	1.28	1.32
35	BB	1099	U	N3-C4	-5.92	1.33	1.38
35	BB	1371	G	P-O5'	-5.92	1.53	1.59
38	BE	146	U	C3'-C2'	-5.92	1.46	1.52
40	BG	31	G	C5-C6	-5.92	1.36	1.42
41	BH	19	G	C5-C4	-5.92	1.34	1.38
85	AA	432	A	O4'-C1'	-5.92	1.33	1.41
34	BA	14	G	C2-N2	-5.91	1.28	1.34
34	BA	244	A	C1'-N9	-5.91	1.38	1.46
34	BA	1221	A	N1-C2	-5.91	1.29	1.34
34	BA	1226	G	N7-C5	-5.91	1.35	1.39
34	BA	1342	C	O3'-P	-5.91	1.54	1.61
34	BA	1836	A	C5-C4	-5.91	1.34	1.38
35	BB	545	C	C4-C5	-5.91	1.38	1.43
35	BB	1151	A	N3-C4	-5.91	1.31	1.34
35	BB	1235	A	C1'-N9	-5.91	1.38	1.46
36	BC	4	G	N3-C4	-5.91	1.31	1.35
85	AA	276	C	O3'-P	-5.91	1.54	1.61
85	AA	361	U	C2-N3	-5.91	1.33	1.37
85	AA	489	C	C4-N4	-5.91	1.28	1.33
85	AA	767	A	C5'-C4'	5.91	1.58	1.51
85	AA	2145	G	P-O5'	-5.91	1.53	1.59
34	BA	691	A	O3'-P	-5.91	1.54	1.61
35	BB	618	U	O3'-P	-5.91	1.54	1.61
35	BB	1146	C	C4'-C3'	-5.91	1.46	1.52
35	BB	1159	U	C1'-N1	-5.91	1.38	1.46
35	BB	1258	G	N3-C4	-5.91	1.31	1.35
35	BB	1271	A	C6-N1	-5.91	1.31	1.35
40	BG	84	U	C2'-C1'	-5.91	1.46	1.53
85	AA	992	G	C5-C4	-5.91	1.34	1.38
85	AA	1129	A	N9-C4	-5.91	1.34	1.37
85	AA	1913	G	O3'-P	-5.91	1.54	1.61
85	AA	2071	U	C3'-C2'	-5.91	1.46	1.52
34	BA	112	C	C1'-N1	-5.91	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	181	G	C1'-N9	-5.91	1.38	1.46
34	BA	590	U	C3'-C2'	-5.91	1.46	1.52
34	BA	814	C	C1'-N1	-5.91	1.38	1.46
34	BA	1224	A	C5-C4	-5.91	1.34	1.38
35	BB	417	A	C2'-C1'	-5.91	1.46	1.53
35	BB	1415	G	N9-C4	-5.91	1.33	1.38
35	BB	1424	G	C4'-C3'	-5.91	1.46	1.52
38	BE	53	U	C4-C5	-5.91	1.38	1.43
38	BE	115	U	C2-N3	-5.91	1.33	1.37
40	BG	22	G	C3'-O3'	-5.91	1.33	1.42
85	AA	355	G	C1'-N9	-5.91	1.38	1.46
85	AA	679	A	N3-C4	-5.91	1.31	1.34
85	AA	774	C	C3'-C2'	-5.91	1.46	1.52
85	AA	802	A	O3'-P	-5.91	1.54	1.61
85	AA	1298	G	C6-N1	-5.91	1.35	1.39
85	AA	1528	A	N9-C8	-5.91	1.33	1.37
34	BA	444	A	C5-C4	-5.91	1.34	1.38
35	BB	473	U	O3'-P	-5.91	1.54	1.61
35	BB	833	G	C2-N2	-5.91	1.28	1.34
35	BB	1394	A	C5-C4	-5.91	1.34	1.38
37	BD	21	G	C6-N1	-5.91	1.35	1.39
39	BF	48	G	C8-N7	-5.91	1.27	1.30
40	BG	76	C	N3-C4	5.91	1.38	1.33
85	AA	477	U	C2'-C1'	-5.91	1.46	1.53
85	AA	622	G	N9-C8	-5.91	1.33	1.37
85	AA	945	A	C4'-C3'	-5.91	1.46	1.52
85	AA	1258	U	C2'-C1'	-5.91	1.46	1.53
85	AA	1844	A	C2'-C1'	-5.91	1.46	1.53
34	BA	1332	U	C1'-N1	-5.91	1.38	1.46
34	BA	1437	G	C2-N2	-5.91	1.28	1.34
35	BB	100	A	C3'-C2'	-5.91	1.46	1.52
35	BB	510	A	C6-N1	-5.91	1.31	1.35
35	BB	616	U	C3'-C2'	-5.91	1.46	1.52
85	AA	742	U	O3'-P	-5.91	1.54	1.61
85	AA	764	U	O3'-P	-5.91	1.54	1.61
85	AA	2130	G	C2-N2	-5.91	1.28	1.34
34	BA	121	A	C4'-O4'	-5.91	1.37	1.45
34	BA	473	A	C2'-C1'	-5.91	1.46	1.53
34	BA	479	U	C2-N3	-5.91	1.33	1.37
34	BA	540	G	O3'-P	-5.91	1.54	1.61
34	BA	707	C	C3'-C2'	-5.91	1.46	1.52
34	BA	900	A	O4'-C1'	-5.91	1.33	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1048	C	O3'-P	-5.91	1.54	1.61
34	BA	1548	A	C4'-C3'	-5.91	1.46	1.52
35	BB	426	A	C5-C4	-5.91	1.34	1.38
35	BB	518	G	C3'-C2'	-5.91	1.46	1.52
35	BB	691	A	O3'-P	-5.91	1.54	1.61
35	BB	853	U	P-O5'	-5.91	1.53	1.59
35	BB	1407	U	C4'-C3'	-5.91	1.46	1.52
37	BD	98	G	C1'-N9	-5.91	1.38	1.46
39	BF	62	U	C1'-N1	-5.91	1.38	1.46
85	AA	146	U	C2-N3	-5.91	1.33	1.37
85	AA	704	A	P-O5'	-5.91	1.53	1.59
85	AA	1511	C	N1-C6	-5.91	1.33	1.37
85	AA	1986	G	O3'-P	-5.91	1.54	1.61
34	BA	1193	A	C1'-N9	-5.90	1.38	1.46
34	BA	1418	G	C5-C6	-5.90	1.36	1.42
35	BB	790	A	N7-C5	-5.90	1.35	1.39
85	AA	168	A	C3'-C2'	-5.90	1.46	1.52
85	AA	424	A	C5-C4	-5.90	1.34	1.38
85	AA	696	G	N9-C4	-5.90	1.33	1.38
85	AA	2035	C	C4'-C3'	-5.90	1.46	1.52
34	BA	129	U	C2'-C1'	-5.90	1.46	1.53
34	BA	683	C	C3'-O3'	5.90	1.50	1.42
34	BA	1149	C	C3'-C2'	-5.90	1.46	1.52
34	BA	1190	A	N9-C4	-5.90	1.34	1.37
35	BB	488	G	N9-C8	-5.90	1.33	1.37
35	BB	1124	G	C5-C6	-5.90	1.36	1.42
38	BE	194	A	C6-N1	-5.90	1.31	1.35
40	BG	174	G	C4'-C3'	-5.90	1.46	1.52
85	AA	378	A	C5-C4	-5.90	1.34	1.38
85	AA	515	C	C4'-C3'	-5.90	1.46	1.52
85	AA	879	G	C4'-C3'	-5.90	1.46	1.52
34	BA	293	A	N9-C8	-5.90	1.33	1.37
34	BA	655	U	C3'-C2'	-5.90	1.46	1.52
34	BA	934	G	N3-C4	-5.90	1.31	1.35
34	BA	1287	G	N3-C4	-5.90	1.31	1.35
34	BA	1405	A	C4'-C3'	-5.90	1.46	1.52
34	BA	1452	U	O3'-P	-5.90	1.54	1.61
35	BB	1227	G	C4'-C3'	-5.90	1.46	1.52
35	BB	1353	G	C6-N1	-5.90	1.35	1.39
38	BE	140	G	N9-C8	-5.90	1.33	1.37
85	AA	55	A	C5-C4	-5.90	1.34	1.38
85	AA	103	U	N3-C4	-5.90	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	174	U	P-O5'	-5.90	1.53	1.59
85	AA	686	U	C2-N3	-5.90	1.33	1.37
85	AA	730	G	C2'-C1'	-5.90	1.46	1.53
85	AA	1000	U	C1'-N1	-5.90	1.38	1.46
85	AA	1671	G	C2'-C1'	-5.90	1.46	1.53
85	AA	1931	C	C5'-C4'	5.90	1.58	1.51
34	BA	187	G	N1-C2	-5.90	1.33	1.37
34	BA	848	U	O3'-P	-5.90	1.54	1.61
34	BA	888	G	C3'-C2'	-5.90	1.46	1.52
34	BA	1069	U	C2-N3	-5.90	1.33	1.37
34	BA	1412	G	P-O5'	-5.90	1.53	1.59
35	BB	29	C	O3'-P	-5.90	1.54	1.61
35	BB	693	U	N1-C2	-5.90	1.33	1.38
35	BB	1197	G	P-O5'	-5.90	1.53	1.59
40	BG	106	G	N3-C4	-5.90	1.31	1.35
85	AA	572	G	C2-N3	-5.90	1.28	1.32
85	AA	2200	A	C4'-C3'	-5.90	1.46	1.52
86	AB	7	A	O3'-P	-5.90	1.54	1.61
34	BA	42	A	C1'-N9	-5.90	1.38	1.46
35	BB	1003	G	O3'-P	-5.90	1.54	1.61
35	BB	1186	A	N9-C4	-5.90	1.34	1.37
35	BB	1281	G	N3-C4	-5.90	1.31	1.35
35	BB	1401	G	C5-C4	-5.90	1.34	1.38
36	BC	10	C	C1'-N1	-5.90	1.38	1.46
38	BE	160	C	C4'-C3'	-5.90	1.46	1.52
39	BF	8	C	P-O5'	-5.90	1.53	1.59
85	AA	113	U	O3'-P	-5.90	1.54	1.61
85	AA	403	G	O3'-P	-5.90	1.54	1.61
85	AA	1856	G	C1'-N9	-5.90	1.38	1.46
34	BA	419	U	C3'-C2'	-5.90	1.46	1.52
34	BA	484	A	N7-C5	-5.90	1.35	1.39
34	BA	487	A	C6-N6	-5.90	1.29	1.33
34	BA	755	G	O4'-C1'	-5.90	1.33	1.41
34	BA	849	G	C1'-N9	-5.90	1.38	1.46
34	BA	1126	U	P-O5'	-5.90	1.53	1.59
34	BA	1252	G	C3'-C2'	-5.90	1.46	1.52
34	BA	1590	G	C4'-C3'	-5.90	1.46	1.52
35	BB	895	U	N1-C2	5.90	1.43	1.38
85	AA	57	G	C2-N2	-5.90	1.28	1.34
85	AA	475	A	O4'-C1'	-5.90	1.33	1.41
85	AA	764	U	N3-C4	-5.90	1.33	1.38
85	AA	917	A	O3'-P	-5.90	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2000	C	O3'-P	-5.90	1.54	1.61
85	AA	2123	U	C4'-O4'	-5.90	1.37	1.45
85	AA	2130	G	C8-N7	-5.90	1.27	1.30
25	AR	51	ASP	N-CA	-5.89	1.34	1.46
34	BA	255	G	C2-N2	-5.89	1.28	1.34
34	BA	392	A	P-O5'	-5.89	1.53	1.59
34	BA	487	A	O3'-P	-5.89	1.54	1.61
34	BA	1049	G	C2-N2	-5.89	1.28	1.34
34	BA	1415	C	C2'-C1'	-5.89	1.46	1.53
35	BB	602	G	N7-C5	-5.89	1.35	1.39
35	BB	1133	C	N3-C4	-5.89	1.29	1.33
36	BC	147	G	C5-C6	-5.89	1.36	1.42
38	BE	11	A	N9-C4	-5.89	1.34	1.37
49	BP	93	ARG	N-CA	-5.89	1.34	1.46
85	AA	430	G	C6-N1	-5.89	1.35	1.39
85	AA	784	C	C3'-C2'	-5.89	1.46	1.52
85	AA	1099	U	N3-C4	-5.89	1.33	1.38
85	AA	1248	U	C2-N3	-5.89	1.33	1.37
85	AA	1306	U	O3'-P	-5.89	1.54	1.61
85	AA	1456	A	P-O5'	-5.89	1.53	1.59
85	AA	2157	G	C2-N2	-5.89	1.28	1.34
34	BA	851	C	C3'-C2'	-5.89	1.46	1.52
34	BA	854	A	C2'-C1'	-5.89	1.46	1.53
34	BA	1262	A	O3'-P	-5.89	1.54	1.61
34	BA	1516	G	N9-C8	-5.89	1.33	1.37
34	BA	1544	G	C1'-N9	-5.89	1.38	1.46
35	BB	615	A	C5-C4	-5.89	1.34	1.38
35	BB	1187	G	C2'-C1'	-5.89	1.46	1.53
35	BB	1537	C	O3'-P	-5.89	1.54	1.61
38	BE	41	C	C3'-C2'	-5.89	1.46	1.52
38	BE	59	U	N3-C4	-5.89	1.33	1.38
38	BE	160	C	C4-N4	-5.89	1.28	1.33
39	BF	10	A	P-O5'	-5.89	1.53	1.59
85	AA	708	G	C5-C4	-5.89	1.34	1.38
85	AA	918	U	N3-C4	-5.89	1.33	1.38
34	BA	420	A	C2'-C1'	-5.89	1.46	1.53
34	BA	490	A	N9-C8	-5.89	1.33	1.37
34	BA	1424	G	N3-C4	-5.89	1.31	1.35
34	BA	1803	A	C8-N7	-5.89	1.27	1.31
35	BB	123	U	C2-N3	-5.89	1.33	1.37
35	BB	818	U	O3'-P	-5.89	1.54	1.61
85	AA	248	U	C2'-C1'	-5.89	1.46	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	588	G	N9-C4	-5.89	1.33	1.38
85	AA	621	U	C2-N3	-5.89	1.33	1.37
85	AA	2127	G	C8-N7	-5.89	1.27	1.30
34	BA	112	C	N1-C6	-5.89	1.33	1.37
34	BA	331	G	C2'-C1'	-5.89	1.46	1.53
34	BA	699	G	C3'-C2'	-5.89	1.46	1.52
34	BA	1505	G	C2'-C1'	-5.89	1.46	1.53
34	BA	1590	G	N9-C4	-5.89	1.33	1.38
35	BB	104	G	P-O5'	-5.89	1.53	1.59
35	BB	465	C	C1'-N1	-5.89	1.38	1.46
35	BB	693	U	C2-N3	-5.89	1.33	1.37
35	BB	1197	G	O3'-P	-5.89	1.54	1.61
35	BB	1226	G	C2-N3	-5.89	1.28	1.32
35	BB	1259	A	O3'-P	-5.89	1.54	1.61
35	BB	1477	C	O3'-P	-5.89	1.54	1.61
35	BB	1511	U	O4'-C1'	-5.89	1.33	1.41
38	BE	210	G	P-O5'	-5.89	1.53	1.59
85	AA	324	U	O3'-P	-5.89	1.54	1.61
85	AA	879	G	N7-C5	-5.89	1.35	1.39
85	AA	1274	A	N3-C4	-5.89	1.31	1.34
85	AA	1698	A	C5-C4	-5.89	1.34	1.38
85	AA	1912	U	O3'-P	-5.89	1.54	1.61
34	BA	272	A	N3-C4	-5.89	1.31	1.34
34	BA	878	G	O3'-P	-5.89	1.54	1.61
34	BA	1245	C	C1'-N1	-5.89	1.38	1.46
34	BA	1741	G	C6-N1	-5.89	1.35	1.39
35	BB	1386	C	C2-N3	-5.89	1.31	1.35
35	BB	1430	G	N9-C4	-5.89	1.33	1.38
38	BE	91	G	C3'-C2'	-5.89	1.46	1.52
40	BG	156	G	C1'-N9	-5.89	1.38	1.46
85	AA	665	A	O3'-P	-5.89	1.54	1.61
85	AA	863	C	C2-N3	-5.89	1.31	1.35
34	BA	220	U	N3-C4	-5.89	1.33	1.38
34	BA	429	G	N9-C8	-5.89	1.33	1.37
34	BA	682	A	P-O5'	-5.89	1.53	1.59
34	BA	711	C	C2'-C1'	-5.89	1.46	1.53
34	BA	1331	G	C5-C4	-5.89	1.34	1.38
35	BB	269	A	C2'-C1'	-5.89	1.46	1.53
35	BB	1039	A	O3'-P	-5.89	1.54	1.61
36	BC	73	U	P-O5'	-5.89	1.53	1.59
36	BC	128	U	N3-C4	-5.89	1.33	1.38
38	BE	5	A	C5-C4	-5.89	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
39	BF	44	C	C4'-C3'	-5.89	1.46	1.52
40	BG	137	G	C3'-C2'	-5.89	1.46	1.52
85	AA	111	A	N7-C5	-5.89	1.35	1.39
85	AA	133	G	C2-N2	-5.89	1.28	1.34
85	AA	404	A	C1'-N9	-5.89	1.38	1.46
85	AA	747	U	P-O5'	-5.89	1.53	1.59
85	AA	1136	A	C8-N7	-5.89	1.27	1.31
85	AA	1734	A	N9-C4	5.89	1.41	1.37
85	AA	1978	G	C2-N2	-5.89	1.28	1.34
85	AA	2028	G	O3'-P	-5.89	1.54	1.61
34	BA	432	A	C8-N7	-5.88	1.27	1.31
34	BA	573	U	C2'-C1'	-5.88	1.46	1.53
34	BA	684	G	C6-N1	-5.88	1.35	1.39
34	BA	1457	C	O3'-P	-5.88	1.54	1.61
35	BB	695	U	C2-N3	-5.88	1.33	1.37
35	BB	1094	A	N9-C4	-5.88	1.34	1.37
35	BB	1268	C	O3'-P	-5.88	1.54	1.61
85	AA	159	G	N9-C8	-5.88	1.33	1.37
85	AA	258	G	N9-C4	-5.88	1.33	1.38
85	AA	318	A	C4'-O4'	-5.88	1.38	1.45
85	AA	678	A	N7-C5	-5.88	1.35	1.39
34	BA	713	C	C2-N3	-5.88	1.31	1.35
34	BA	896	U	O3'-P	-5.88	1.54	1.61
35	BB	1169	A	O3'-P	-5.88	1.54	1.61
85	AA	693	A	N3-C4	-5.88	1.31	1.34
85	AA	2031	C	C4'-C3'	-5.88	1.46	1.52
85	AA	2178	A	O3'-P	-5.88	1.54	1.61
34	BA	168	U	C3'-C2'	-5.88	1.46	1.52
34	BA	187	G	C1'-N9	-5.88	1.38	1.46
34	BA	759	A	C8-N7	-5.88	1.27	1.31
34	BA	860	G	N9-C8	-5.88	1.33	1.37
34	BA	1200	U	C2'-C1'	-5.88	1.46	1.53
35	BB	60	A	C5-C4	-5.88	1.34	1.38
35	BB	657	A	C6-N1	-5.88	1.31	1.35
35	BB	697	G	N1-C2	-5.88	1.33	1.37
35	BB	1086	G	N1-C2	-5.88	1.33	1.37
36	BC	18	G	C3'-C2'	-5.88	1.46	1.52
36	BC	86	U	C4'-C3'	5.88	1.59	1.53
39	BF	38	C	C2-N3	-5.88	1.31	1.35
39	BF	65	U	C4'-O4'	-5.88	1.38	1.45
85	AA	111	A	C6-N1	-5.88	1.31	1.35
85	AA	940	G	C8-N7	-5.88	1.27	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	608	G	P-O5'	5.88	1.65	1.59
34	BA	1288	U	O3'-P	-5.88	1.54	1.61
34	BA	1403	G	C1'-N9	-5.88	1.38	1.46
35	BB	679	G	N1-C2	-5.88	1.33	1.37
34	BA	242	U	C3'-C2'	-5.88	1.46	1.52
34	BA	841	G	C6-N1	-5.88	1.35	1.39
34	BA	1030	C	P-O5'	-5.88	1.53	1.59
34	BA	1285	G	N7-C5	-5.88	1.35	1.39
34	BA	1347	G	C2-N2	-5.88	1.28	1.34
34	BA	1605	G	C4'-C3'	-5.88	1.46	1.52
34	BA	1816	G	N9-C4	-5.88	1.33	1.38
35	BB	1299	G	C2-N2	-5.88	1.28	1.34
36	BC	11	G	N1-C2	-5.88	1.33	1.37
37	BD	99	G	C2-N2	-5.88	1.28	1.34
38	BE	199	A	O3'-P	-5.88	1.54	1.61
40	BG	131	U	C1'-N1	-5.88	1.38	1.46
41	BH	134	U	C4'-C3'	-5.88	1.46	1.52
85	AA	336	C	C2'-C1'	-5.88	1.46	1.53
85	AA	656	U	O3'-P	-5.88	1.54	1.61
85	AA	1257	A	C3'-C2'	-5.88	1.46	1.52
85	AA	2126	U	C4'-O4'	-5.88	1.38	1.45
34	BA	745	A	C6-N1	-5.88	1.31	1.35
34	BA	1345	U	O3'-P	-5.88	1.54	1.61
35	BB	468	U	C3'-C2'	-5.88	1.46	1.52
35	BB	613	C	C3'-C2'	-5.88	1.46	1.52
38	BE	15	A	C3'-C2'	-5.88	1.46	1.52
40	BG	115	C	C2-N3	-5.88	1.31	1.35
41	BH	13	C	N1-C2	-5.88	1.34	1.40
85	AA	395	G	C5-C6	-5.88	1.36	1.42
85	AA	816	A	O3'-P	-5.88	1.54	1.61
85	AA	1268	C	C3'-C2'	-5.88	1.46	1.52
85	AA	1938	G	C2'-C1'	-5.88	1.46	1.53
85	AA	2145	G	N7-C5	-5.88	1.35	1.39
34	BA	280	A	C8-N7	-5.88	1.27	1.31
34	BA	1025	A	N9-C8	-5.88	1.33	1.37
35	BB	527	U	O3'-P	-5.88	1.54	1.61
35	BB	1459	U	C2-N3	-5.88	1.33	1.37
40	BG	168	A	C5-C4	-5.88	1.34	1.38
41	BH	45	G	C8-N7	-5.88	1.27	1.30
85	AA	358	U	O3'-P	-5.88	1.54	1.61
85	AA	463	G	N9-C8	-5.88	1.33	1.37
34	BA	351	A	C6-N6	-5.87	1.29	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1311	G	C5-C4	-5.87	1.34	1.38
34	BA	1379	G	N3-C4	-5.87	1.31	1.35
34	BA	1716	A	N3-C4	-5.87	1.31	1.34
35	BB	135	C	C1'-N1	-5.87	1.38	1.46
35	BB	544	C	C3'-C2'	-5.87	1.46	1.52
35	BB	1250	A	C5-C4	-5.87	1.34	1.38
37	BD	54	A	C8-N7	-5.87	1.27	1.31
38	BE	148	C	C3'-C2'	-5.87	1.46	1.52
85	AA	464	A	O3'-P	-5.87	1.54	1.61
85	AA	607	U	C2'-C1'	-5.87	1.46	1.53
85	AA	669	G	C6-N1	-5.87	1.35	1.39
85	AA	1921	G	C3'-C2'	-5.87	1.46	1.52
85	AA	2230	U	C3'-O3'	5.87	1.50	1.42
34	BA	588	C	O3'-P	-5.87	1.54	1.61
34	BA	1555	G	C4'-C3'	-5.87	1.46	1.52
35	BB	41	A	P-O5'	-5.87	1.53	1.59
35	BB	44	C	C2'-C1'	-5.87	1.46	1.53
35	BB	94	A	N9-C8	-5.87	1.33	1.37
35	BB	127	U	N1-C6	-5.87	1.32	1.38
35	BB	822	G	C2-N3	5.87	1.37	1.32
35	BB	1353	G	O3'-P	-5.87	1.54	1.61
40	BG	14	G	C2-N2	-5.87	1.28	1.34
85	AA	72	C	C2'-C1'	-5.87	1.46	1.53
85	AA	586	G	O3'-P	-5.87	1.54	1.61
85	AA	705	G	C2'-C1'	-5.87	1.46	1.53
85	AA	1162	A	C1'-N9	-5.87	1.38	1.46
85	AA	1480	C	C1'-N1	-5.87	1.38	1.46
34	BA	410	G	N9-C4	-5.87	1.33	1.38
34	BA	884	G	O3'-P	-5.87	1.54	1.61
34	BA	1160	U	N1-C2	-5.87	1.33	1.38
34	BA	1272	U	C4'-C3'	-5.87	1.46	1.52
34	BA	1490	U	C1'-N1	-5.87	1.38	1.46
34	BA	1567	G	C6-N1	-5.87	1.35	1.39
35	BB	87	G	N3-C4	-5.87	1.31	1.35
35	BB	386	G	C2'-C1'	-5.87	1.46	1.53
38	BE	28	C	C4'-C3'	-5.87	1.46	1.52
85	AA	400	G	C4'-O4'	-5.87	1.38	1.45
85	AA	442	G	C8-N7	-5.87	1.27	1.30
85	AA	1447	U	C2-N3	-5.87	1.33	1.37
34	BA	3	G	N7-C5	-5.87	1.35	1.39
34	BA	27	G	N7-C5	-5.87	1.35	1.39
34	BA	177	G	N9-C4	-5.87	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	242	U	C2'-C1'	-5.87	1.46	1.53
34	BA	348	U	N3-C4	-5.87	1.33	1.38
34	BA	686	U	C2'-C1'	-5.87	1.46	1.53
34	BA	1093	G	N7-C5	-5.87	1.35	1.39
35	BB	6	A	O3'-P	-5.87	1.54	1.61
35	BB	435	A	C2'-C1'	-5.87	1.46	1.53
35	BB	443	A	P-O5'	-5.87	1.53	1.59
35	BB	659	C	O4'-C1'	-5.87	1.34	1.41
35	BB	801	G	C2'-C1'	-5.87	1.46	1.53
35	BB	1169	A	C2'-C1'	-5.87	1.46	1.53
35	BB	1422	G	C2-N3	-5.87	1.28	1.32
35	BB	1445	A	C6-N1	-5.87	1.31	1.35
41	BH	37	U	N1-C2	-5.87	1.33	1.38
85	AA	12	U	C4'-C3'	-5.87	1.46	1.52
85	AA	242	G	N9-C4	-5.87	1.33	1.38
85	AA	1120	G	O3'-P	-5.87	1.54	1.61
34	BA	295	G	N9-C8	-5.87	1.33	1.37
34	BA	1781	A	N7-C5	-5.87	1.35	1.39
35	BB	1180	G	C3'-C2'	-5.87	1.46	1.52
35	BB	1396	G	C6-N1	-5.87	1.35	1.39
40	BG	161	C	O4'-C1'	-5.87	1.34	1.41
41	BH	104	U	O4'-C1'	-5.87	1.34	1.41
85	AA	375	C	O3'-P	-5.87	1.54	1.61
85	AA	1885	A	N7-C5	-5.87	1.35	1.39
34	BA	168	U	C2'-C1'	-5.87	1.46	1.53
34	BA	421	G	C6-N1	-5.87	1.35	1.39
34	BA	764	G	C4'-C3'	5.87	1.59	1.53
34	BA	961	C	C4'-C3'	-5.87	1.46	1.52
34	BA	1807	G	C5-C6	-5.87	1.36	1.42
35	BB	651	G	O3'-P	-5.87	1.54	1.61
35	BB	1263	A	C1'-N9	-5.87	1.38	1.46
35	BB	1359	G	C5-C6	-5.87	1.36	1.42
35	BB	1381	U	P-O5'	-5.87	1.53	1.59
36	BC	91	G	C5-C4	-5.87	1.34	1.38
36	BC	154	A	N9-C4	-5.87	1.34	1.37
37	BD	9	C	P-O5'	-5.87	1.53	1.59
40	BG	175	G	N7-C5	-5.87	1.35	1.39
41	BH	20	A	C5'-C4'	-5.87	1.44	1.51
48	BO	208	MET	C-N	-5.87	1.23	1.34
85	AA	105	A	N7-C5	-5.87	1.35	1.39
85	AA	430	G	N9-C4	-5.87	1.33	1.38
85	AA	571	G	C5'-C4'	5.87	1.58	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	582	A	C3'-C2'	-5.87	1.46	1.52
85	AA	1274	A	P-O5'	-5.87	1.53	1.59
85	AA	1914	U	P-O5'	-5.87	1.53	1.59
85	AA	2020	C	C5'-C4'	-5.87	1.44	1.51
85	AA	2195	A	P-O5'	-5.87	1.53	1.59
34	BA	401	A	N9-C8	-5.86	1.33	1.37
34	BA	526	C	C1'-N1	-5.86	1.38	1.46
34	BA	614	A	N9-C4	-5.86	1.34	1.37
34	BA	621	G	C6-N1	-5.86	1.35	1.39
34	BA	722	A	C8-N7	-5.86	1.27	1.31
34	BA	1195	G	C5-C4	-5.86	1.34	1.38
34	BA	1638	U	N3-C4	-5.86	1.33	1.38
34	BA	1638	U	P-O5'	-5.86	1.53	1.59
35	BB	24	C	P-O5'	-5.86	1.53	1.59
35	BB	56	U	C3'-C2'	-5.86	1.46	1.52
35	BB	1087	A	N3-C4	-5.86	1.31	1.34
35	BB	1119	G	N9-C4	-5.86	1.33	1.38
35	BB	1299	G	N7-C5	-5.86	1.35	1.39
38	BE	47	U	C3'-C2'	-5.86	1.46	1.52
38	BE	98	C	C3'-C2'	-5.86	1.46	1.52
40	BG	27	C	C2-N3	-5.86	1.31	1.35
85	AA	339	A	C3'-C2'	-5.86	1.46	1.52
85	AA	794	A	P-O5'	-5.86	1.53	1.59
85	AA	1247	A	C1'-N9	-5.86	1.38	1.46
34	BA	81	C	C2-N3	-5.86	1.31	1.35
34	BA	906	A	O4'-C1'	-5.86	1.34	1.41
34	BA	1276	G	C3'-C2'	-5.86	1.46	1.52
34	BA	1440	C	N1-C6	-5.86	1.33	1.37
35	BB	1367	U	N3-C4	-5.86	1.33	1.38
40	BG	88	G	C5-C4	-5.86	1.34	1.38
85	AA	462	A	P-O5'	-5.86	1.53	1.59
85	AA	2067	A	P-O5'	-5.86	1.53	1.59
34	BA	816	G	N1-C2	-5.86	1.33	1.37
34	BA	1466	U	C4'-C3'	-5.86	1.46	1.52
35	BB	546	A	C2'-C1'	-5.86	1.47	1.53
35	BB	833	G	P-O5'	-5.86	1.53	1.59
35	BB	1165	A	C3'-C2'	-5.86	1.46	1.52
35	BB	1467	A	C8-N7	-5.86	1.27	1.31
38	BE	88	G	N7-C5	-5.86	1.35	1.39
38	BE	188	C	C2'-C1'	-5.86	1.47	1.53
85	AA	447	C	N3-C4	-5.86	1.29	1.33
85	AA	823	C	C3'-C2'	-5.86	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	67	A	N9-C8	-5.86	1.33	1.37
34	BA	209	A	C5-C4	-5.86	1.34	1.38
34	BA	795	G	C2'-C1'	-5.86	1.47	1.53
37	BD	65	G	N9-C8	-5.86	1.33	1.37
34	BA	362	G	C2'-C1'	-5.86	1.47	1.53
34	BA	574	U	C1'-N1	5.86	1.57	1.48
34	BA	585	G	C8-N7	-5.86	1.27	1.30
34	BA	751	A	C5-C4	-5.86	1.34	1.38
34	BA	975	A	C1'-N9	-5.86	1.38	1.46
34	BA	1475	G	C6-N1	-5.86	1.35	1.39
34	BA	1590	G	C2'-C1'	-5.86	1.47	1.53
35	BB	425	G	C5-C4	-5.86	1.34	1.38
35	BB	462	G	O3'-P	-5.86	1.54	1.61
35	BB	731	U	O3'-P	-5.86	1.54	1.61
35	BB	1450	G	C5-C4	-5.86	1.34	1.38
35	BB	1494	G	C6-N1	-5.86	1.35	1.39
36	BC	4	G	P-O5'	-5.86	1.53	1.59
36	BC	30	U	C4'-O4'	-5.86	1.38	1.45
38	BE	105	A	N7-C5	-5.86	1.35	1.39
39	BF	51	C	C4'-C3'	5.86	1.59	1.53
41	BH	63	G	N9-C4	5.86	1.42	1.38
85	AA	9	U	C3'-C2'	-5.86	1.46	1.52
85	AA	322	A	C5-C4	-5.86	1.34	1.38
85	AA	1978	G	C5-C4	-5.86	1.34	1.38
34	BA	148	G	N9-C8	-5.86	1.33	1.37
34	BA	389	U	N3-C4	-5.86	1.33	1.38
34	BA	540	G	C3'-C2'	-5.86	1.46	1.52
34	BA	874	G	C2-N3	5.86	1.37	1.32
34	BA	1311	G	N1-C2	-5.86	1.33	1.37
34	BA	1532	G	C2-N2	-5.86	1.28	1.34
34	BA	1653	G	C2-N2	-5.86	1.28	1.34
34	BA	1678	U	C2'-C1'	-5.86	1.47	1.53
35	BB	998	G	C3'-C2'	-5.86	1.46	1.52
35	BB	1281	G	C1'-N9	-5.86	1.38	1.46
35	BB	1347	C	C4'-C3'	-5.86	1.46	1.52
35	BB	1363	A	C1'-N9	-5.86	1.38	1.46
35	BB	1424	G	C2'-C1'	-5.86	1.47	1.53
41	BH	10	U	P-O5'	-5.86	1.53	1.59
85	AA	282	C	P-O5'	-5.86	1.53	1.59
85	AA	522	A	O3'-P	-5.86	1.54	1.61
85	AA	588	G	O4'-C1'	-5.86	1.34	1.41
85	AA	775	C	C2'-C1'	-5.86	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	817	G	C6-N1	-5.86	1.35	1.39
85	AA	1541	G	C5'-C4'	-5.86	1.44	1.51
34	BA	95	C	O4'-C1'	-5.85	1.34	1.41
34	BA	570	G	P-O5'	-5.85	1.53	1.59
34	BA	988	U	C3'-C2'	-5.85	1.46	1.52
35	BB	467	G	C1'-N9	-5.85	1.38	1.46
37	BD	73	U	C1'-N1	-5.85	1.38	1.46
41	BH	35	G	C2'-C1'	-5.85	1.47	1.53
85	AA	1833	C	O3'-P	-5.85	1.54	1.61
86	AB	63	G	N9-C4	-5.85	1.33	1.38
34	BA	349	G	C2-N2	-5.85	1.28	1.34
34	BA	491	U	C1'-N1	-5.85	1.38	1.46
34	BA	496	G	C3'-C2'	-5.85	1.46	1.52
34	BA	754	G	N3-C4	-5.85	1.31	1.35
34	BA	859	G	O3'-P	-5.85	1.54	1.61
34	BA	906	A	C6-N1	-5.85	1.31	1.35
34	BA	1131	G	C2'-C1'	-5.85	1.47	1.53
34	BA	1435	A	C4'-O4'	-5.85	1.38	1.45
34	BA	1454	G	C2-N3	-5.85	1.28	1.32
34	BA	1539	A	C2'-C1'	-5.85	1.47	1.53
34	BA	1572	G	C2-N2	-5.85	1.28	1.34
35	BB	58	G	N9-C8	-5.85	1.33	1.37
35	BB	130	G	N1-C2	-5.85	1.33	1.37
35	BB	658	G	N9-C8	-5.85	1.33	1.37
35	BB	1103	A	C6-N1	-5.85	1.31	1.35
35	BB	1425	A	C1'-N9	-5.85	1.38	1.46
36	BC	163	A	N7-C5	-5.85	1.35	1.39
37	BD	16	U	C1'-N1	-5.85	1.38	1.46
38	BE	89	G	P-O5'	-5.85	1.53	1.59
38	BE	111	C	N1-C2	-5.85	1.34	1.40
39	BF	3	A	C2'-C1'	-5.85	1.47	1.53
85	AA	631	G	P-O5'	-5.85	1.53	1.59
85	AA	696	G	C2-N2	-5.85	1.28	1.34
85	AA	1221	G	N1-C2	-5.85	1.33	1.37
34	BA	327	G	C4'-C3'	-5.85	1.46	1.52
34	BA	1652	G	C2-N2	-5.85	1.28	1.34
34	BA	1833	G	P-O5'	-5.85	1.53	1.59
35	BB	634	A	C5'-C4'	-5.85	1.44	1.51
36	BC	90	U	C2-N3	-5.85	1.33	1.37
85	AA	31	C	C1'-N1	-5.85	1.38	1.46
85	AA	1509	A	N9-C4	-5.85	1.34	1.37
34	BA	900	A	P-O5'	-5.85	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1033	G	N7-C5	-5.85	1.35	1.39
34	BA	1496	G	C6-N1	-5.85	1.35	1.39
35	BB	361	A	N9-C4	-5.85	1.34	1.37
35	BB	644	A	C1'-N9	-5.85	1.38	1.46
35	BB	1081	U	C3'-C2'	-5.85	1.46	1.52
35	BB	1335	G	C6-N1	-5.85	1.35	1.39
35	BB	1478	G	P-O5'	-5.85	1.53	1.59
36	BC	109	A	N7-C5	-5.85	1.35	1.39
37	BD	107	G	C3'-C2'	-5.85	1.46	1.52
40	BG	71	C	C3'-C2'	-5.85	1.46	1.52
85	AA	102	A	C5-C4	-5.85	1.34	1.38
85	AA	388	G	N9-C4	-5.85	1.33	1.38
85	AA	403	G	C5-C4	-5.85	1.34	1.38
85	AA	740	A	P-O5'	-5.85	1.53	1.59
85	AA	757	A	C3'-C2'	-5.85	1.46	1.52
85	AA	960	G	N3-C4	-5.85	1.31	1.35
85	AA	992	G	C4'-C3'	-5.85	1.46	1.52
85	AA	1237	A	C4'-C3'	-5.85	1.46	1.52
85	AA	1786	G	C2'-C1'	-5.85	1.47	1.53
85	AA	1825	A	P-O5'	-5.85	1.53	1.59
85	AA	2103	C	C2'-C1'	-5.85	1.47	1.53
85	AA	2201	A	C3'-C2'	-5.85	1.46	1.52
34	BA	346	A	N3-C4	-5.85	1.31	1.34
34	BA	661	C	C2-N3	-5.85	1.31	1.35
34	BA	832	C	O3'-P	-5.85	1.54	1.61
34	BA	1220	C	C4-N4	-5.85	1.28	1.33
34	BA	1648	G	C3'-C2'	-5.85	1.46	1.52
35	BB	654	C	C2'-C1'	-5.85	1.47	1.53
35	BB	1328	C	C5'-C4'	5.85	1.58	1.51
35	BB	1469	A	C5-C4	-5.85	1.34	1.38
36	BC	155	C	C3'-C2'	-5.85	1.46	1.52
36	BC	167	U	P-O5'	-5.85	1.53	1.59
37	BD	88	U	C2'-C1'	-5.85	1.47	1.53
39	BF	25	G	C5-C4	-5.85	1.34	1.38
40	BG	97	G	C2-N2	-5.85	1.28	1.34
85	AA	363	A	C4'-O4'	-5.85	1.38	1.45
85	AA	1784	G	O3'-P	-5.85	1.54	1.61
34	BA	346	A	P-O5'	-5.85	1.53	1.59
34	BA	969	A	O4'-C1'	-5.85	1.34	1.41
35	BB	126	C	C2'-C1'	-5.85	1.47	1.53
35	BB	317	C	P-O5'	-5.85	1.53	1.59
35	BB	1547	U	C2'-C1'	-5.85	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	506	G	C5-C6	-5.85	1.36	1.42
85	AA	526	G	C3'-C2'	-5.85	1.46	1.52
85	AA	1170	C	C2-N3	-5.85	1.31	1.35
34	BA	444	A	C1'-N9	-5.84	1.38	1.46
34	BA	830	U	P-O5'	-5.84	1.53	1.59
34	BA	911	G	O3'-P	-5.84	1.54	1.61
34	BA	918	U	O3'-P	-5.84	1.54	1.61
35	BB	38	C	C2-N3	-5.84	1.31	1.35
35	BB	502	C	C3'-C2'	-5.84	1.46	1.52
35	BB	537	A	O3'-P	-5.84	1.54	1.61
35	BB	700	C	N1-C6	-5.84	1.33	1.37
35	BB	828	G	C6-N1	-5.84	1.35	1.39
35	BB	1406	C	N3-C4	-5.84	1.29	1.33
36	BC	105	C	O3'-P	-5.84	1.54	1.61
38	BE	89	G	O3'-P	-5.84	1.54	1.61
41	BH	121	A	C6-N1	-5.84	1.31	1.35
59	BZ	40	TYR	CB-CG	-5.84	1.42	1.51
85	AA	78	A	O3'-P	-5.84	1.54	1.61
85	AA	272	C	P-O5'	-5.84	1.53	1.59
85	AA	502	A	O3'-P	-5.84	1.54	1.61
85	AA	768	C	C3'-C2'	-5.84	1.46	1.52
85	AA	789	A	P-O5'	5.84	1.65	1.59
85	AA	1850	G	O3'-P	-5.84	1.54	1.61
85	AA	1896	G	P-O5'	-5.84	1.53	1.59
85	AA	2119	C	N1-C2	5.84	1.46	1.40
34	BA	1698	C	C2'-C1'	-5.84	1.47	1.53
35	BB	545	C	P-O5'	-5.84	1.53	1.59
35	BB	568	A	P-O5'	-5.84	1.53	1.59
36	BC	94	C	C3'-C2'	-5.84	1.46	1.52
36	BC	108	A	O3'-P	-5.84	1.54	1.61
41	BH	4	U	N1-C2	-5.84	1.33	1.38
85	AA	1350	A	O3'-P	-5.84	1.54	1.61
85	AA	1907	U	P-O5'	-5.84	1.53	1.59
34	BA	339	G	N3-C4	-5.84	1.31	1.35
34	BA	1509	U	O3'-P	-5.84	1.54	1.61
34	BA	1714	A	C2'-C1'	-5.84	1.47	1.53
35	BB	704	G	C5'-C4'	-5.84	1.44	1.51
35	BB	806	U	C1'-N1	-5.84	1.38	1.46
35	BB	1484	A	C2'-C1'	-5.84	1.47	1.53
39	BF	25	G	N9-C4	-5.84	1.33	1.38
85	AA	146	U	C2'-C1'	-5.84	1.47	1.53
85	AA	590	U	P-O5'	-5.84	1.53	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1220	A	C6-N1	-5.84	1.31	1.35
34	BA	12	G	C8-N7	-5.84	1.27	1.30
34	BA	1723	U	C1'-N1	5.84	1.57	1.48
35	BB	1507	U	C2-N3	-5.84	1.33	1.37
85	AA	420	C	C2'-C1'	-5.84	1.47	1.53
85	AA	910	G	C3'-C2'	-5.84	1.46	1.52
34	BA	506	U	C4'-C3'	-5.84	1.46	1.52
34	BA	907	A	C3'-C2'	-5.84	1.46	1.52
34	BA	1381	A	C2'-C1'	-5.84	1.47	1.53
35	BB	25	A	C5-C4	-5.84	1.34	1.38
35	BB	780	U	C2-N3	-5.84	1.33	1.37
35	BB	1222	A	C2'-C1'	-5.84	1.47	1.53
85	AA	10	G	P-O5'	-5.84	1.53	1.59
85	AA	484	G	C1'-N9	-5.84	1.38	1.46
85	AA	695	A	C2'-C1'	-5.84	1.47	1.53
34	BA	340	U	N3-C4	-5.84	1.33	1.38
34	BA	454	G	C6-N1	-5.84	1.35	1.39
34	BA	472	G	N3-C4	-5.84	1.31	1.35
34	BA	853	A	O4'-C1'	-5.84	1.34	1.41
34	BA	1510	C	N3-C4	-5.84	1.29	1.33
35	BB	486	G	C1'-N9	-5.84	1.38	1.46
35	BB	1156	U	C3'-C2'	-5.84	1.46	1.52
35	BB	1226	G	P-O5'	-5.84	1.53	1.59
37	BD	59	G	C6-N1	-5.84	1.35	1.39
38	BE	139	U	C1'-N1	-5.84	1.38	1.46
40	BG	155	A	C2'-C1'	-5.84	1.47	1.53
85	AA	211	C	C1'-N1	-5.84	1.38	1.46
85	AA	663	C	P-O5'	-5.84	1.53	1.59
85	AA	1215	A	C5'-C4'	-5.84	1.44	1.51
85	AA	1450	U	C4'-C3'	-5.84	1.46	1.52
85	AA	1792	C	C2'-C1'	-5.84	1.47	1.53
85	AA	1925	A	P-O5'	-5.84	1.53	1.59
85	AA	1961	U	C3'-C2'	-5.84	1.46	1.52
85	AA	1974	C	N1-C6	5.84	1.40	1.37
85	AA	2207	A	P-O5'	-5.84	1.53	1.59
86	AB	10	G	C2'-C1'	-5.84	1.47	1.53
34	BA	569	C	C2'-C1'	-5.83	1.47	1.53
34	BA	1798	G	N9-C8	-5.83	1.33	1.37
35	BB	119	G	C2'-C1'	-5.83	1.47	1.53
35	BB	538	A	O3'-P	-5.83	1.54	1.61
35	BB	1500	U	N1-C2	-5.83	1.33	1.38
85	AA	1492	U	C2-N3	-5.83	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1992	A	N9-C4	-5.83	1.34	1.37
34	BA	1033	G	C5-C4	-5.83	1.34	1.38
34	BA	1150	A	C6-N1	-5.83	1.31	1.35
34	BA	1403	G	C2'-C1'	-5.83	1.47	1.53
34	BA	1718	C	C4'-O4'	-5.83	1.38	1.45
34	BA	1781	A	P-O5'	-5.83	1.53	1.59
34	BA	1800	G	C8-N7	-5.83	1.27	1.30
35	BB	1306	G	N9-C4	-5.83	1.33	1.38
35	BB	1466	A	P-O5'	-5.83	1.53	1.59
36	BC	123	G	C5-C4	-5.83	1.34	1.38
37	BD	35	C	O3'-P	-5.83	1.54	1.61
85	AA	545	A	N3-C4	-5.83	1.31	1.34
85	AA	555	C	C2-N3	-5.83	1.31	1.35
85	AA	665	A	C6-N1	-5.83	1.31	1.35
85	AA	2120	C	C2'-C1'	-5.83	1.47	1.53
85	AA	2127	G	C2-N2	-5.83	1.28	1.34
34	BA	66	C	C2'-C1'	-5.83	1.47	1.53
34	BA	115	U	N3-C4	-5.83	1.33	1.38
34	BA	248	G	C8-N7	-5.83	1.27	1.30
34	BA	604	G	C2'-C1'	-5.83	1.47	1.53
34	BA	1223	C	C2'-O2'	-5.83	1.34	1.41
34	BA	1411	C	C4'-C3'	-5.83	1.46	1.52
35	BB	372	U	C1'-N1	-5.83	1.38	1.46
35	BB	639	A	N7-C5	-5.83	1.35	1.39
37	BD	12	U	C1'-N1	-5.83	1.38	1.46
38	BE	92	C	C2-N3	-5.83	1.31	1.35
38	BE	164	C	O3'-P	-5.83	1.54	1.61
40	BG	35	G	N9-C4	-5.83	1.33	1.38
85	AA	1276	A	N9-C4	-5.83	1.34	1.37
85	AA	1510	A	O3'-P	-5.83	1.54	1.61
34	BA	721	A	N9-C4	-5.83	1.34	1.37
35	BB	73	G	N1-C2	-5.83	1.33	1.37
35	BB	788	U	O4'-C1'	-5.83	1.34	1.41
85	AA	1167	G	C1'-N9	-5.83	1.38	1.46
34	BA	626	G	N9-C4	5.83	1.42	1.38
34	BA	825	G	N9-C4	-5.83	1.33	1.38
34	BA	882	G	C6-N1	5.83	1.43	1.39
34	BA	1148	U	C5'-C4'	-5.83	1.44	1.51
34	BA	1671	A	C1'-N9	-5.83	1.38	1.46
35	BB	785	G	N1-C2	-5.83	1.33	1.37
35	BB	1036	G	C2'-C1'	-5.83	1.47	1.53
37	BD	50	A	C2'-C1'	-5.83	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	204	U	C4'-C3'	-5.83	1.46	1.52
74	Bo	78	THR	CA-C	-5.83	1.37	1.52
85	AA	928	U	N1-C6	-5.83	1.32	1.38
34	BA	367	G	C3'-O3'	5.83	1.50	1.42
35	BB	396	C	C2'-C1'	-5.83	1.47	1.53
38	BE	168	C	O3'-P	-5.83	1.54	1.61
85	AA	710	A	N9-C4	-5.83	1.34	1.37
85	AA	944	C	C2'-C1'	-5.83	1.47	1.53
85	AA	1182	A	C1'-N9	-5.83	1.38	1.46
85	AA	1531	G	N7-C5	-5.83	1.35	1.39
34	BA	201	A	C4'-C3'	-5.83	1.46	1.52
34	BA	203	U	N3-C4	-5.83	1.33	1.38
34	BA	416	A	C1'-N9	-5.83	1.38	1.46
34	BA	1019	C	P-O5'	-5.83	1.53	1.59
34	BA	1555	G	C6-N1	-5.83	1.35	1.39
34	BA	1610	A	C5'-C4'	-5.83	1.44	1.51
35	BB	1431	G	N1-C2	-5.83	1.33	1.37
35	BB	1485	G	C6-N1	-5.83	1.35	1.39
36	BC	155	C	N1-C2	-5.83	1.34	1.40
40	BG	129	G	P-O5'	-5.83	1.53	1.59
85	AA	565	G	C2'-C1'	-5.83	1.47	1.53
34	BA	30	A	N9-C4	-5.82	1.34	1.37
34	BA	87	G	C1'-N9	-5.82	1.38	1.46
34	BA	258	C	C4-N4	-5.82	1.28	1.33
34	BA	431	A	P-O5'	-5.82	1.53	1.59
34	BA	462	C	C3'-C2'	-5.82	1.46	1.52
34	BA	474	A	C5-C4	-5.82	1.34	1.38
35	BB	473	U	C4'-C3'	-5.82	1.46	1.52
35	BB	664	A	C5'-C4'	-5.82	1.44	1.51
85	AA	244	G	C1'-N9	-5.82	1.38	1.46
85	AA	500	C	C1'-N1	-5.82	1.38	1.46
85	AA	1668	G	N9-C8	-5.82	1.33	1.37
34	BA	499	C	C1'-N1	-5.82	1.38	1.46
34	BA	1683	C	C4-N4	-5.82	1.28	1.33
35	BB	574	G	C4'-C3'	5.82	1.59	1.53
36	BC	45	C	C2-N3	-5.82	1.31	1.35
36	BC	48	A	C5-C4	-5.82	1.34	1.38
40	BG	180	C	O3'-P	-5.82	1.54	1.61
85	AA	358	U	P-O5'	-5.82	1.53	1.59
85	AA	1865	C	P-O5'	-5.82	1.53	1.59
85	AA	1972	A	O3'-P	-5.82	1.54	1.61
85	AA	2132	A	C5'-C4'	-5.82	1.44	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	547	C	O4'-C1'	-5.82	1.34	1.41
34	BA	1189	A	C2'-C1'	-5.82	1.47	1.53
34	BA	1221	A	C6-N1	-5.82	1.31	1.35
34	BA	1718	C	C1'-N1	-5.82	1.38	1.46
35	BB	460	C	C3'-C2'	-5.82	1.46	1.52
35	BB	552	C	C2-N3	-5.82	1.31	1.35
35	BB	666	A	C1'-N9	-5.82	1.38	1.46
35	BB	707	G	C5'-C4'	5.82	1.58	1.51
35	BB	767	A	C4'-C3'	-5.82	1.46	1.52
37	BD	105	G	N9-C8	-5.82	1.33	1.37
38	BE	130	G	C3'-C2'	-5.82	1.46	1.52
39	BF	34	C	O4'-C1'	-5.82	1.34	1.41
85	AA	622	G	N1-C2	-5.82	1.33	1.37
85	AA	1117	G	C5-C4	-5.82	1.34	1.38
85	AA	1531	G	C5-C6	-5.82	1.36	1.42
85	AA	2073	U	P-O5'	-5.82	1.53	1.59
34	BA	1211	G	P-O5'	5.82	1.65	1.59
34	BA	1412	G	C8-N7	-5.82	1.27	1.30
35	BB	1269	A	P-O5'	-5.82	1.53	1.59
36	BC	103	A	C1'-N9	-5.82	1.38	1.46
37	BD	97	U	C4'-C3'	-5.82	1.46	1.52
37	BD	111	U	C3'-C2'	-5.82	1.46	1.52
38	BE	6	A	N3-C4	-5.82	1.31	1.34
41	BH	27	A	N3-C4	-5.82	1.31	1.34
85	AA	501	A	C3'-C2'	-5.82	1.46	1.52
85	AA	532	G	C2'-C1'	-5.82	1.47	1.53
85	AA	754	C	O3'-P	-5.82	1.54	1.61
34	BA	794	G	C3'-C2'	-5.82	1.46	1.52
34	BA	1592	U	N3-C4	-5.82	1.33	1.38
34	BA	1703	A	C4'-O4'	-5.82	1.38	1.45
35	BB	5	A	C5'-C4'	5.82	1.58	1.51
35	BB	60	A	O3'-P	-5.82	1.54	1.61
35	BB	87	G	C3'-C2'	-5.82	1.46	1.52
35	BB	347	G	P-O5'	-5.82	1.53	1.59
35	BB	427	U	C2'-C1'	-5.82	1.47	1.53
35	BB	597	C	N1-C6	-5.82	1.33	1.37
35	BB	1332	G	N7-C5	-5.82	1.35	1.39
35	BB	1333	U	C2'-C1'	-5.82	1.47	1.53
36	BC	9	G	N9-C8	-5.82	1.33	1.37
38	BE	149	A	P-O5'	-5.82	1.53	1.59
39	BF	33	C	P-O5'	-5.82	1.53	1.59
41	BH	99	G	C2-N3	-5.82	1.28	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
79	Bt	39	GLY	CA-C	-5.82	1.42	1.51
85	AA	46	U	O3'-P	-5.82	1.54	1.61
85	AA	154	U	O3'-P	-5.82	1.54	1.61
85	AA	257	U	N3-C4	-5.82	1.33	1.38
85	AA	428	G	C3'-C2'	-5.82	1.46	1.52
85	AA	1158	U	C3'-C2'	-5.82	1.46	1.52
85	AA	1681	G	C4'-O4'	-5.82	1.38	1.45
85	AA	1993	C	C3'-O3'	5.82	1.50	1.42
34	BA	948	C	C4'-C3'	-5.82	1.46	1.52
34	BA	1013	A	N3-C4	-5.82	1.31	1.34
34	BA	1331	G	C3'-C2'	-5.82	1.46	1.52
34	BA	1371	U	P-O5'	-5.82	1.53	1.59
34	BA	1564	A	C5'-C4'	-5.82	1.44	1.51
34	BA	1585	A	C2'-C1'	-5.82	1.47	1.53
34	BA	1685	C	O4'-C1'	-5.82	1.34	1.41
35	BB	694	C	C3'-C2'	-5.82	1.46	1.52
37	BD	107	G	C6-N1	-5.82	1.35	1.39
85	AA	105	A	C5-C4	-5.82	1.34	1.38
85	AA	154	U	C2'-C1'	-5.82	1.47	1.53
85	AA	244	G	C2'-C1'	-5.82	1.47	1.53
85	AA	293	A	O3'-P	-5.82	1.54	1.61
85	AA	614	U	O4'-C1'	-5.82	1.34	1.41
85	AA	1549	G	N3-C4	-5.82	1.31	1.35
85	AA	1901	G	C6-N1	-5.82	1.35	1.39
86	AB	19	G	N3-C4	-5.82	1.31	1.35
34	BA	58	A	N3-C4	-5.81	1.31	1.34
34	BA	476	U	N3-C4	-5.81	1.33	1.38
34	BA	566	G	N3-C4	-5.81	1.31	1.35
34	BA	678	C	C4-C5	-5.81	1.38	1.43
34	BA	854	A	N9-C4	-5.81	1.34	1.37
85	AA	2043	A	P-O5'	-5.81	1.53	1.59
34	BA	520	G	C6-N1	-5.81	1.35	1.39
34	BA	523	A	C3'-C2'	-5.81	1.46	1.52
34	BA	1016	A	C1'-N9	-5.81	1.38	1.46
34	BA	1050	A	C5-C4	-5.81	1.34	1.38
34	BA	1399	A	C1'-N9	-5.81	1.38	1.46
34	BA	1409	A	C1'-N9	-5.81	1.38	1.46
34	BA	1539	A	N7-C5	-5.81	1.35	1.39
34	BA	1594	G	P-O5'	-5.81	1.53	1.59
35	BB	90	G	C5-C4	-5.81	1.34	1.38
38	BE	2	G	C2'-C1'	-5.81	1.47	1.53
41	BH	68	G	C6-N1	-5.81	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	724	A	P-O5'	-5.81	1.53	1.59
85	AA	942	A	O3'-P	-5.81	1.54	1.61
85	AA	1010	U	P-O5'	-5.81	1.53	1.59
85	AA	1243	G	N7-C5	-5.81	1.35	1.39
85	AA	1731	G	C2-N2	-5.81	1.28	1.34
34	BA	410	G	C5-C4	-5.81	1.34	1.38
34	BA	721	A	C5-C4	-5.81	1.34	1.38
34	BA	1070	G	C6-N1	-5.81	1.35	1.39
35	BB	550	G	C6-N1	-5.81	1.35	1.39
35	BB	956	G	C2'-C1'	-5.81	1.47	1.53
36	BC	121	G	C2-N2	-5.81	1.28	1.34
40	BG	57	A	O3'-P	-5.81	1.54	1.61
85	AA	447	C	C4-N4	-5.81	1.28	1.33
85	AA	533	C	N1-C2	-5.81	1.34	1.40
85	AA	654	A	C4'-O4'	-5.81	1.38	1.45
34	BA	63	A	C6-N6	-5.81	1.29	1.33
34	BA	1064	A	C2'-C1'	-5.81	1.47	1.53
34	BA	1198	U	C5'-C4'	5.81	1.58	1.51
34	BA	1258	G	P-O5'	-5.81	1.53	1.59
35	BB	102	G	C2-N2	-5.81	1.28	1.34
35	BB	623	A	C5-C4	-5.81	1.34	1.38
36	BC	78	G	C1'-N9	-5.81	1.38	1.46
36	BC	126	G	N7-C5	-5.81	1.35	1.39
39	BF	52	A	C1'-N9	-5.81	1.38	1.46
40	BG	88	G	N1-C2	-5.81	1.33	1.37
40	BG	92	U	O3'-P	-5.81	1.54	1.61
41	BH	7	C	C4-N4	-5.81	1.28	1.33
85	AA	460	U	N3-C4	-5.81	1.33	1.38
85	AA	2073	U	N3-C4	-5.81	1.33	1.38
34	BA	1467	U	N3-C4	-5.81	1.33	1.38
34	BA	1673	G	C1'-N9	-5.81	1.38	1.46
35	BB	373	C	C4-N4	-5.81	1.28	1.33
36	BC	28	C	C1'-N1	-5.81	1.38	1.46
36	BC	91	G	C6-N1	-5.81	1.35	1.39
38	BE	160	C	C3'-C2'	-5.81	1.46	1.52
41	BH	122	U	C2'-C1'	-5.81	1.47	1.53
85	AA	418	G	N9-C8	-5.81	1.33	1.37
85	AA	527	A	N9-C8	-5.81	1.33	1.37
85	AA	692	U	C2-N3	-5.81	1.33	1.37
85	AA	1373	U	C3'-C2'	-5.81	1.46	1.52
85	AA	1971	G	C2'-C1'	-5.81	1.47	1.53
34	BA	1291	A	C3'-C2'	-5.81	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	88	U	N3-C4	-5.81	1.33	1.38
35	BB	651	G	P-O5'	-5.81	1.53	1.59
36	BC	143	C	P-O5'	-5.81	1.53	1.59
85	AA	706	U	O3'-P	-5.81	1.54	1.61
85	AA	1289	U	O3'-P	-5.81	1.54	1.61
34	BA	71	G	N3-C4	-5.80	1.31	1.35
34	BA	341	U	N3-C4	-5.80	1.33	1.38
34	BA	804	G	N1-C2	-5.80	1.33	1.37
34	BA	1235	C	C4-N4	-5.80	1.28	1.33
34	BA	1778	U	P-O5'	-5.80	1.53	1.59
34	BA	1833	G	C5-C4	-5.80	1.34	1.38
34	BA	1846	G	C6-N1	-5.80	1.35	1.39
35	BB	99	G	C2'-C1'	-5.80	1.47	1.53
35	BB	783	U	C3'-C2'	-5.80	1.46	1.52
35	BB	833	G	N7-C5	-5.80	1.35	1.39
35	BB	975	G	N1-C2	-5.80	1.33	1.37
35	BB	1456	G	N9-C8	-5.80	1.33	1.37
36	BC	110	A	P-O5'	-5.80	1.53	1.59
38	BE	140	G	C2-N2	-5.80	1.28	1.34
39	BF	23	G	P-O5'	5.80	1.65	1.59
71	Bl	111	ARG	CD-NE	5.80	1.56	1.46
85	AA	36	U	N3-C4	-5.80	1.33	1.38
85	AA	363	A	C5'-C4'	-5.80	1.44	1.51
85	AA	622	G	C5-C4	-5.80	1.34	1.38
85	AA	1117	G	C6-N1	-5.80	1.35	1.39
85	AA	1295	G	C6-N1	-5.80	1.35	1.39
85	AA	1735	U	N1-C2	5.80	1.43	1.38
85	AA	2088	U	O3'-P	-5.80	1.54	1.61
34	BA	144	C	C1'-N1	-5.80	1.38	1.46
35	BB	1131	C	C4'-C3'	-5.80	1.46	1.52
37	BD	4	U	N3-C4	-5.80	1.33	1.38
85	AA	151	A	N3-C4	-5.80	1.31	1.34
34	BA	588	C	O4'-C1'	-5.80	1.34	1.41
34	BA	1225	A	P-O5'	-5.80	1.53	1.59
34	BA	1301	G	P-O5'	-5.80	1.53	1.59
34	BA	1728	G	C8-N7	-5.80	1.27	1.30
34	BA	1808	A	C5-C4	-5.80	1.34	1.38
35	BB	68	G	C4'-O4'	-5.80	1.38	1.45
35	BB	560	C	C3'-C2'	-5.80	1.46	1.52
35	BB	642	G	C1'-N9	-5.80	1.38	1.46
35	BB	1148	U	C5'-C4'	-5.80	1.44	1.51
35	BB	1268	C	C2-N3	-5.80	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	4	G	C1'-N9	-5.80	1.38	1.46
36	BC	121	G	P-O5'	-5.80	1.53	1.59
38	BE	47	U	C2'-C1'	-5.80	1.47	1.53
39	BF	32	G	N9-C4	-5.80	1.33	1.38
85	AA	110	U	C2-N3	-5.80	1.33	1.37
85	AA	238	C	O3'-P	-5.80	1.54	1.61
85	AA	265	A	C4'-C3'	5.80	1.59	1.53
85	AA	585	G	P-O5'	-5.80	1.53	1.59
85	AA	670	C	P-O5'	-5.80	1.53	1.59
85	AA	860	C	C2'-C1'	-5.80	1.47	1.53
34	BA	15	G	C2-N3	-5.80	1.28	1.32
34	BA	261	A	O3'-P	-5.80	1.54	1.61
34	BA	456	G	C6-N1	-5.80	1.35	1.39
34	BA	982	A	C8-N7	-5.80	1.27	1.31
34	BA	1202	G	C5'-C4'	-5.80	1.44	1.51
34	BA	1827	C	C2-N3	-5.80	1.31	1.35
36	BC	54	G	O3'-P	-5.80	1.54	1.61
37	BD	53	U	P-O5'	-5.80	1.53	1.59
37	BD	54	A	C2'-C1'	-5.80	1.47	1.53
37	BD	76	U	C2-N3	-5.80	1.33	1.37
38	BE	26	G	N7-C5	-5.80	1.35	1.39
40	BG	61	A	C5-C4	-5.80	1.34	1.38
85	AA	392	G	N1-C2	-5.80	1.33	1.37
85	AA	1513	U	O3'-P	-5.80	1.54	1.61
85	AA	1924	C	O3'-P	-5.80	1.54	1.61
34	BA	1020	A	P-O5'	-5.80	1.53	1.59
34	BA	1338	G	C2'-C1'	-5.80	1.47	1.53
34	BA	1437	G	N9-C4	-5.80	1.33	1.38
35	BB	104	G	N9-C4	-5.80	1.33	1.38
35	BB	434	A	N3-C4	-5.80	1.31	1.34
35	BB	1334	C	N3-C4	-5.80	1.29	1.33
35	BB	1524	G	C1'-N9	-5.80	1.38	1.46
36	BC	37	U	C4'-C3'	-5.80	1.46	1.52
37	BD	30	A	P-O5'	-5.80	1.53	1.59
34	BA	93	A	O4'-C1'	-5.80	1.34	1.41
34	BA	183	G	N1-C2	-5.80	1.33	1.37
34	BA	716	C	C3'-C2'	-5.80	1.46	1.52
34	BA	1039	G	C5'-C4'	-5.80	1.44	1.51
34	BA	1102	A	C5-C4	-5.80	1.34	1.38
34	BA	1189	A	C6-N1	-5.80	1.31	1.35
34	BA	1602	A	P-O5'	-5.80	1.53	1.59
34	BA	1603	A	N9-C8	-5.80	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1645	C	C3'-C2'	-5.80	1.46	1.52
35	BB	11	A	C3'-C2'	-5.80	1.46	1.52
35	BB	31	U	P-O5'	-5.80	1.53	1.59
35	BB	54	U	C2'-C1'	-5.80	1.47	1.53
35	BB	392	G	C3'-C2'	-5.80	1.46	1.52
35	BB	706	G	C5-C4	-5.80	1.34	1.38
35	BB	1049	G	C5-C4	-5.80	1.34	1.38
35	BB	1233	U	O3'-P	-5.80	1.54	1.61
35	BB	1424	G	O4'-C1'	-5.80	1.34	1.41
40	BG	17	A	C1'-N9	-5.80	1.38	1.46
40	BG	62	C	C3'-C2'	-5.80	1.46	1.52
85	AA	157	G	N9-C8	-5.80	1.33	1.37
85	AA	1472	G	C3'-C2'	-5.80	1.46	1.52
34	BA	1138	C	P-O5'	-5.79	1.53	1.59
34	BA	1451	A	C6-N6	-5.79	1.29	1.33
36	BC	169	G	N9-C4	-5.79	1.33	1.38
39	BF	53	G	C4'-O4'	-5.79	1.38	1.45
34	BA	557	U	C5-C6	-5.79	1.28	1.34
34	BA	720	A	C5-C6	-5.79	1.35	1.41
35	BB	417	A	O3'-P	-5.79	1.54	1.61
35	BB	448	G	C5-C4	-5.79	1.34	1.38
35	BB	487	A	C3'-C2'	-5.79	1.46	1.52
35	BB	518	G	C2'-C1'	-5.79	1.47	1.53
35	BB	520	G	O3'-P	-5.79	1.54	1.61
35	BB	1431	G	C1'-N9	-5.79	1.38	1.46
85	AA	28	A	C2'-C1'	-5.79	1.47	1.53
85	AA	133	G	C2'-C1'	-5.79	1.47	1.53
85	AA	422	G	C2-N2	-5.79	1.28	1.34
85	AA	972	G	C4'-C3'	5.79	1.59	1.53
85	AA	2006	G	C2-N2	-5.79	1.28	1.34
85	AA	2177	C	C1'-N1	-5.79	1.38	1.46
34	BA	1325	G	C5-C4	-5.79	1.34	1.38
35	BB	1263	A	C2'-C1'	-5.79	1.47	1.53
35	BB	1416	A	C2'-C1'	-5.79	1.47	1.53
35	BB	1511	U	C2'-C1'	-5.79	1.47	1.53
36	BC	95	A	C4'-C3'	-5.79	1.46	1.52
38	BE	61	A	C5-C4	-5.79	1.34	1.38
41	BH	114	G	N3-C4	-5.79	1.31	1.35
85	AA	408	C	C3'-C2'	-5.79	1.46	1.52
85	AA	441	C	C4-N4	-5.79	1.28	1.33
85	AA	819	G	O3'-P	-5.79	1.54	1.61
85	AA	830	A	C1'-N9	-5.79	1.38	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	877	G	C6-O6	-5.79	1.19	1.24
85	AA	1162	A	C3'-C2'	-5.79	1.46	1.52
85	AA	1273	C	P-O5'	-5.79	1.53	1.59
85	AA	2233	A	O3'-P	-5.79	1.54	1.61
34	BA	68	A	N9-C8	-5.79	1.33	1.37
34	BA	116	G	O3'-P	-5.79	1.54	1.61
34	BA	1315	C	C3'-C2'	-5.79	1.46	1.52
38	BE	163	A	C6-N1	-5.79	1.31	1.35
40	BG	44	G	C2-N2	-5.79	1.28	1.34
85	AA	92	G	C4'-C3'	-5.79	1.46	1.52
85	AA	1549	G	O4'-C1'	-5.79	1.34	1.41
85	AA	1614	G	O3'-P	-5.79	1.54	1.61
34	BA	103	G	N1-C2	-5.79	1.33	1.37
34	BA	124	G	C2'-C1'	-5.79	1.47	1.53
34	BA	215	C	P-O5'	-5.79	1.53	1.59
34	BA	220	U	O3'-P	-5.79	1.54	1.61
34	BA	565	U	C1'-N1	-5.79	1.38	1.46
34	BA	805	A	N7-C5	-5.79	1.35	1.39
34	BA	939	C	C4-C5	-5.79	1.38	1.43
34	BA	985	C	C2'-C1'	-5.79	1.47	1.53
34	BA	1456	C	C4'-O4'	-5.79	1.38	1.45
34	BA	1557	G	N1-C2	-5.79	1.33	1.37
34	BA	1693	U	C2-N3	-5.79	1.33	1.37
35	BB	641	C	P-O5'	-5.79	1.53	1.59
35	BB	1398	A	N3-C4	-5.79	1.31	1.34
85	AA	269	G	N7-C5	-5.79	1.35	1.39
85	AA	2131	C	C2-N3	-5.79	1.31	1.35
85	AA	629	A	N7-C5	-5.79	1.35	1.39
85	AA	861	G	N3-C4	-5.79	1.31	1.35
85	AA	1217	U	P-O5'	-5.79	1.53	1.59
85	AA	1243	G	C2-N2	-5.79	1.28	1.34
85	AA	2169	C	N3-C4	-5.79	1.29	1.33
34	BA	819	G	O3'-P	-5.79	1.54	1.61
34	BA	1088	G	N9-C8	-5.79	1.33	1.37
34	BA	1164	C	C3'-C2'	-5.79	1.46	1.52
34	BA	1213	A	C5-C6	-5.79	1.35	1.41
34	BA	1409	A	N3-C4	-5.79	1.31	1.34
34	BA	1497	A	C3'-C2'	5.79	1.59	1.52
34	BA	1670	A	C2'-C1'	-5.79	1.47	1.53
34	BA	1675	C	C5'-C4'	-5.79	1.44	1.51
35	BB	1198	C	C2-N3	-5.79	1.31	1.35
38	BE	55	C	C3'-O3'	5.79	1.50	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	10	U	O3'-P	-5.79	1.54	1.61
41	BH	129	G	O3'-P	-5.79	1.54	1.61
85	AA	693	A	P-O5'	-5.79	1.53	1.59
85	AA	1856	G	C3'-C2'	-5.79	1.46	1.52
34	BA	116	G	C2'-C1'	-5.78	1.47	1.53
34	BA	388	A	N3-C4	-5.78	1.31	1.34
34	BA	513	U	C1'-N1	-5.78	1.38	1.46
34	BA	703	U	N3-C4	-5.78	1.33	1.38
34	BA	795	G	C5-C6	-5.78	1.36	1.42
34	BA	1533	G	C2-N2	-5.78	1.28	1.34
34	BA	1603	A	P-O5'	-5.78	1.53	1.59
35	BB	625	A	N7-C5	-5.78	1.35	1.39
35	BB	1203	C	P-O5'	-5.78	1.53	1.59
35	BB	1248	A	C4'-C3'	-5.78	1.46	1.52
35	BB	1318	U	P-O5'	-5.78	1.53	1.59
38	BE	10	G	C4'-O4'	5.78	1.53	1.45
38	BE	37	C	C2'-C1'	-5.78	1.47	1.53
38	BE	49	A	N7-C5	-5.78	1.35	1.39
40	BG	1	G	C5-C6	-5.78	1.36	1.42
40	BG	86	U	C4-C5	-5.78	1.38	1.43
41	BH	33	G	C5-C4	-5.78	1.34	1.38
85	AA	111	A	N3-C4	-5.78	1.31	1.34
85	AA	362	G	C3'-C2'	-5.78	1.46	1.52
85	AA	682	C	C4'-C3'	-5.78	1.46	1.52
85	AA	887	A	O4'-C1'	-5.78	1.34	1.41
85	AA	1271	U	C2-N3	-5.78	1.33	1.37
34	BA	1492	G	C6-N1	-5.78	1.35	1.39
35	BB	813	C	O3'-P	-5.78	1.54	1.61
35	BB	1461	C	C4'-O4'	-5.78	1.38	1.45
40	BG	65	C	C4'-O4'	-5.78	1.38	1.45
85	AA	1249	U	O3'-P	-5.78	1.54	1.61
85	AA	1581	C	O3'-P	-5.78	1.54	1.61
85	AA	2094	U	C3'-C2'	-5.78	1.46	1.52
34	BA	142	A	C8-N7	-5.78	1.27	1.31
34	BA	580	U	C4'-O4'	-5.78	1.38	1.45
34	BA	966	G	C2-N2	-5.78	1.28	1.34
34	BA	1005	C	P-O5'	-5.78	1.53	1.59
34	BA	1078	U	N1-C2	-5.78	1.33	1.38
34	BA	1203	G	C5'-C4'	-5.78	1.44	1.51
34	BA	1514	A	N9-C4	-5.78	1.34	1.37
34	BA	1655	G	C2'-C1'	-5.78	1.47	1.53
34	BA	1706	A	O3'-P	-5.78	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1802	C	C3'-C2'	-5.78	1.46	1.52
35	BB	19	C	C4'-O4'	-5.78	1.38	1.45
35	BB	115	A	C6-N1	5.78	1.39	1.35
35	BB	979	G	C2-N2	-5.78	1.28	1.34
35	BB	1097	U	N3-C4	-5.78	1.33	1.38
40	BG	134	U	C3'-C2'	-5.78	1.46	1.52
71	Bl	42	GLY	CA-C	-5.78	1.42	1.51
85	AA	433	U	O3'-P	-5.78	1.54	1.61
85	AA	622	G	O3'-P	-5.78	1.54	1.61
85	AA	704	A	C5-C4	-5.78	1.34	1.38
85	AA	843	U	O3'-P	-5.78	1.54	1.61
85	AA	1144	G	N7-C5	-5.78	1.35	1.39
85	AA	1190	G	O4'-C1'	-5.78	1.34	1.41
85	AA	2089	G	C5-C4	-5.78	1.34	1.38
85	AA	2174	G	N3-C4	-5.78	1.31	1.35
34	BA	434	U	C2'-C1'	-5.78	1.47	1.53
34	BA	482	C	C1'-N1	-5.78	1.38	1.46
34	BA	1290	A	C2'-C1'	-5.78	1.47	1.53
35	BB	1292	G	C3'-C2'	-5.78	1.46	1.52
38	BE	142	A	N9-C8	-5.78	1.33	1.37
85	AA	165	C	C5'-C4'	-5.78	1.44	1.51
85	AA	1675	U	P-O5'	-5.78	1.53	1.59
34	BA	33	C	P-O5'	-5.78	1.53	1.59
34	BA	375	C	C2'-C1'	-5.78	1.47	1.53
34	BA	517	A	C2'-C1'	-5.78	1.47	1.53
34	BA	720	A	N3-C4	-5.78	1.31	1.34
34	BA	823	G	C2'-C1'	-5.78	1.47	1.53
35	BB	765	G	O3'-P	-5.78	1.54	1.61
85	AA	1172	A	O3'-P	-5.78	1.54	1.61
85	AA	1498	C	O3'-P	-5.78	1.54	1.61
85	AA	1689	G	N9-C4	-5.78	1.33	1.38
85	AA	1711	C	C1'-N1	-5.78	1.38	1.46
34	BA	94	G	P-O5'	-5.78	1.53	1.59
34	BA	326	A	O4'-C1'	-5.78	1.34	1.41
34	BA	583	G	O3'-P	-5.78	1.54	1.61
34	BA	955	G	N7-C5	-5.78	1.35	1.39
34	BA	1023	G	C5-C4	-5.78	1.34	1.38
34	BA	1203	G	C8-N7	-5.78	1.27	1.30
35	BB	1224	C	C2'-C1'	-5.78	1.47	1.53
35	BB	1465	U	C3'-C2'	-5.78	1.46	1.52
38	BE	48	G	C2-N2	-5.78	1.28	1.34
40	BG	35	G	C2-N2	-5.78	1.28	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	125	A	N3-C4	-5.78	1.31	1.34
85	AA	210	G	C6-O6	-5.78	1.19	1.24
85	AA	493	A	C1'-N9	-5.78	1.38	1.46
85	AA	680	U	N3-C4	-5.78	1.33	1.38
85	AA	1242	A	N3-C4	-5.78	1.31	1.34
85	AA	1948	A	P-O5'	-5.78	1.53	1.59
85	AA	2085	C	C5'-C4'	-5.78	1.44	1.51
34	BA	14	G	C2'-C1'	-5.77	1.47	1.53
34	BA	261	A	P-O5'	-5.77	1.53	1.59
34	BA	728	A	C5-C4	-5.77	1.34	1.38
34	BA	768	G	C2'-C1'	-5.77	1.47	1.53
34	BA	1077	G	N7-C5	-5.77	1.35	1.39
35	BB	817	C	C5'-C4'	5.77	1.58	1.51
35	BB	1068	G	C2-N2	-5.77	1.28	1.34
85	AA	670	C	O3'-P	-5.77	1.54	1.61
85	AA	1462	A	O3'-P	-5.77	1.54	1.61
34	BA	809	U	C4'-C3'	-5.77	1.46	1.52
34	BA	1025	A	C2'-C1'	-5.77	1.47	1.53
34	BA	1092	U	C4'-C3'	-5.77	1.46	1.52
34	BA	1194	G	C1'-N9	-5.77	1.38	1.46
34	BA	1202	G	C2'-C1'	-5.77	1.47	1.53
34	BA	1713	U	N1-C2	-5.77	1.33	1.38
35	BB	629	C	C2'-C1'	-5.77	1.47	1.53
35	BB	697	G	C5-C4	-5.77	1.34	1.38
35	BB	1276	U	O3'-P	-5.77	1.54	1.61
37	BD	56	G	C2'-C1'	-5.77	1.47	1.53
85	AA	327	G	C4'-C3'	5.77	1.59	1.53
85	AA	636	G	O3'-P	-5.77	1.54	1.61
85	AA	925	G	C2-N2	-5.77	1.28	1.34
85	AA	2146	G	C2-N3	-5.77	1.28	1.32
34	BA	230	A	P-O5'	-5.77	1.53	1.59
34	BA	452	A	N3-C4	-5.77	1.31	1.34
34	BA	1600	G	O4'-C1'	-5.77	1.34	1.41
34	BA	1623	U	C2'-C1'	-5.77	1.47	1.53
34	BA	1685	C	C5'-C4'	-5.77	1.44	1.51
35	BB	4	C	C1'-N1	5.77	1.57	1.48
35	BB	283	A	P-O5'	-5.77	1.53	1.59
35	BB	1538	G	P-O5'	-5.77	1.53	1.59
85	AA	2221	A	O3'-P	-5.77	1.54	1.61
34	BA	791	A	N3-C4	-5.77	1.31	1.34
34	BA	966	G	C6-N1	-5.77	1.35	1.39
34	BA	1013	A	N7-C5	-5.77	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1268	C	O3'-P	-5.77	1.54	1.61
35	BB	1235	A	C5-C4	-5.77	1.34	1.38
40	BG	170	G	O3'-P	-5.77	1.54	1.61
41	BH	42	U	O3'-P	-5.77	1.54	1.61
85	AA	540	A	O3'-P	-5.77	1.54	1.61
85	AA	565	G	N7-C5	-5.77	1.35	1.39
85	AA	690	G	C5'-C4'	-5.77	1.44	1.51
85	AA	1156	A	C8-N7	-5.77	1.27	1.31
85	AA	1541	G	C2-N2	-5.77	1.28	1.34
34	BA	467	A	C8-N7	-5.77	1.27	1.31
34	BA	714	G	C2-N2	-5.77	1.28	1.34
34	BA	1420	A	N7-C5	-5.77	1.35	1.39
35	BB	421	U	C3'-C2'	-5.77	1.46	1.52
35	BB	858	U	P-O5'	-5.77	1.53	1.59
35	BB	1263	A	O3'-P	-5.77	1.54	1.61
38	BE	108	U	C3'-O3'	-5.77	1.34	1.42
85	AA	914	U	C2'-C1'	-5.77	1.47	1.53
85	AA	1440	C	C3'-C2'	-5.77	1.46	1.52
34	BA	263	G	C2'-C1'	-5.77	1.47	1.53
34	BA	517	A	C8-N7	-5.77	1.27	1.31
34	BA	1672	C	N3-C4	-5.77	1.29	1.33
35	BB	832	C	C2-N3	-5.77	1.31	1.35
35	BB	1004	A	N7-C5	-5.77	1.35	1.39
35	BB	1040	C	C5'-C4'	5.77	1.58	1.51
35	BB	1106	G	N1-C2	-5.77	1.33	1.37
36	BC	42	G	C5-C4	-5.77	1.34	1.38
49	BP	19	PRO	N-CD	-5.77	1.39	1.47
66	Bg	81	TYR	CB-CG	-5.77	1.43	1.51
35	BB	779	C	C2'-C1'	-5.76	1.47	1.53
35	BB	1193	G	C3'-C2'	-5.76	1.46	1.52
40	BG	176	G	N9-C8	-5.76	1.33	1.37
85	AA	373	G	C4'-C3'	-5.76	1.46	1.52
85	AA	621	U	P-O5'	-5.76	1.53	1.59
85	AA	1061	C	O3'-P	-5.76	1.54	1.61
85	AA	1293	U	O3'-P	-5.76	1.54	1.61
85	AA	2140	U	N3-C4	-5.76	1.33	1.38
34	BA	257	G	C5-C6	-5.76	1.36	1.42
34	BA	702	G	N1-C2	-5.76	1.33	1.37
34	BA	1709	A	C1'-N9	-5.76	1.38	1.46
37	BD	110	G	N1-C2	-5.76	1.33	1.37
41	BH	59	G	N9-C4	-5.76	1.33	1.38
85	AA	648	G	C3'-C2'	-5.76	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1673	A	C1'-N9	-5.76	1.38	1.46
85	AA	1929	G	C4'-C3'	-5.76	1.46	1.52
85	AA	2105	G	C5-C4	-5.76	1.34	1.38
34	BA	191	G	C3'-C2'	-5.76	1.46	1.52
34	BA	331	G	C2-N2	-5.76	1.28	1.34
34	BA	1560	U	O4'-C1'	-5.76	1.34	1.41
35	BB	113	C	C5'-C4'	-5.76	1.44	1.51
35	BB	399	A	C2'-C1'	-5.76	1.47	1.53
38	BE	187	G	O3'-P	-5.76	1.54	1.61
40	BG	73	U	C3'-C2'	-5.76	1.46	1.52
40	BG	176	G	C2-N2	-5.76	1.28	1.34
41	BH	11	C	C2-N3	-5.76	1.31	1.35
85	AA	515	C	C4-N4	-5.76	1.28	1.33
85	AA	1253	G	N9-C8	-5.76	1.33	1.37
85	AA	1778	C	P-O5'	-5.76	1.53	1.59
85	AA	1850	G	C2'-C1'	-5.76	1.47	1.53
34	BA	844	U	C4'-C3'	-5.76	1.46	1.52
34	BA	1692	U	C2'-C1'	-5.76	1.47	1.53
34	BA	1782	C	C3'-C2'	-5.76	1.46	1.52
35	BB	391	G	O3'-P	-5.76	1.54	1.61
35	BB	1059	U	C3'-C2'	-5.76	1.46	1.52
35	BB	1070	G	C5-C4	-5.76	1.34	1.38
35	BB	1292	G	N7-C5	-5.76	1.35	1.39
36	BC	39	G	C2-N2	-5.76	1.28	1.34
36	BC	61	A	C1'-N9	-5.76	1.38	1.46
40	BG	124	A	C3'-C2'	-5.76	1.46	1.52
41	BH	21	G	C5'-C4'	-5.76	1.44	1.51
85	AA	386	G	N9-C8	-5.76	1.33	1.37
85	AA	544	A	C3'-C2'	-5.76	1.46	1.52
85	AA	705	G	N9-C4	-5.76	1.33	1.38
85	AA	934	A	C2'-C1'	-5.76	1.47	1.53
85	AA	1893	G	P-O5'	-5.76	1.53	1.59
85	AA	1996	A	C5'-C4'	5.76	1.58	1.51
34	BA	49	A	N9-C8	-5.76	1.33	1.37
34	BA	458	G	O3'-P	-5.76	1.54	1.61
34	BA	1117	G	N1-C2	-5.76	1.33	1.37
34	BA	1315	C	C1'-N1	-5.76	1.38	1.46
34	BA	1818	A	N3-C4	-5.76	1.31	1.34
35	BB	390	G	P-O5'	-5.76	1.53	1.59
36	BC	12	A	N3-C4	-5.76	1.31	1.34
85	AA	54	C	O3'-P	-5.76	1.54	1.61
34	BA	210	G	C5-C4	-5.76	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1648	G	N9-C4	-5.76	1.33	1.38
34	BA	1805	C	C2'-C1'	-5.76	1.47	1.53
35	BB	1036	G	C2-N3	-5.76	1.28	1.32
35	BB	1134	G	C8-N7	-5.76	1.27	1.30
35	BB	1397	G	C1'-N9	-5.76	1.38	1.46
35	BB	1478	G	C6-N1	-5.76	1.35	1.39
36	BC	15	G	N3-C4	-5.76	1.31	1.35
38	BE	146	U	C1'-N1	-5.76	1.38	1.46
39	BF	70	A	C2'-C1'	-5.76	1.47	1.53
85	AA	261	U	C1'-N1	-5.76	1.38	1.46
85	AA	338	G	C2'-C1'	-5.76	1.47	1.53
85	AA	499	G	C1'-N9	-5.76	1.38	1.46
85	AA	1240	A	C2'-C1'	-5.76	1.47	1.53
85	AA	1289	U	N3-C4	-5.76	1.33	1.38
85	AA	1850	G	P-O5'	-5.76	1.53	1.59
34	BA	441	A	C1'-N9	-5.75	1.38	1.46
34	BA	792	A	C1'-N9	-5.75	1.38	1.46
34	BA	820	C	C4-N4	-5.75	1.28	1.33
34	BA	1052	G	C3'-C2'	-5.75	1.46	1.52
34	BA	1780	U	P-O5'	-5.75	1.53	1.59
35	BB	488	G	C1'-N9	-5.75	1.38	1.46
35	BB	1020	U	C1'-N1	-5.75	1.38	1.46
36	BC	55	U	C4'-C3'	-5.75	1.46	1.52
36	BC	97	U	C4'-C3'	-5.75	1.46	1.52
40	BG	34	A	C1'-N9	-5.75	1.38	1.46
85	AA	2002	A	N7-C5	-5.75	1.35	1.39
85	AA	2218	G	C4'-O4'	-5.75	1.38	1.45
34	BA	116	G	C5'-C4'	-5.75	1.44	1.51
34	BA	951	C	C2'-C1'	-5.75	1.47	1.53
35	BB	61	A	P-O5'	-5.75	1.53	1.59
35	BB	127	U	C2'-O2'	-5.75	1.34	1.41
35	BB	1057	G	C6-N1	-5.75	1.35	1.39
35	BB	1118	G	C6-N1	-5.75	1.35	1.39
35	BB	1162	A	N7-C5	-5.75	1.35	1.39
35	BB	1421	C	C2'-C1'	-5.75	1.47	1.53
36	BC	67	U	C3'-C2'	-5.75	1.46	1.52
41	BH	14	C	C2'-C1'	-5.75	1.47	1.53
41	BH	129	G	N3-C4	-5.75	1.31	1.35
85	AA	383	C	C4-N4	-5.75	1.28	1.33
85	AA	532	G	C6-N1	-5.75	1.35	1.39
85	AA	669	G	P-O5'	-5.75	1.53	1.59
85	AA	700	U	C2'-C1'	-5.75	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1621	U	C4'-C3'	-5.75	1.46	1.52
85	AA	1873	U	C5'-C4'	5.75	1.58	1.51
85	AA	1974	C	P-O5'	-5.75	1.53	1.59
85	AA	2038	C	C3'-C2'	-5.75	1.46	1.52
34	BA	976	C	O3'-P	-5.75	1.54	1.61
34	BA	1105	A	N3-C4	-5.75	1.31	1.34
34	BA	1707	C	C1'-N1	-5.75	1.38	1.46
35	BB	401	U	C4'-C3'	-5.75	1.46	1.52
35	BB	615	A	C3'-C2'	-5.75	1.46	1.52
36	BC	159	U	O3'-P	-5.75	1.54	1.61
85	AA	103	U	C3'-C2'	-5.75	1.46	1.52
85	AA	471	U	C4'-C3'	-5.75	1.46	1.52
85	AA	588	G	C2-N2	-5.75	1.28	1.34
85	AA	631	G	N9-C4	-5.75	1.33	1.38
85	AA	927	A	C6-N6	-5.75	1.29	1.33
85	AA	1146	C	O3'-P	-5.75	1.54	1.61
85	AA	1825	A	N3-C4	-5.75	1.31	1.34
85	AA	2244	G	N9-C4	-5.75	1.33	1.38
34	BA	45	A	N9-C8	-5.75	1.33	1.37
34	BA	538	G	O3'-P	-5.75	1.54	1.61
34	BA	1484	A	C2'-C1'	-5.75	1.47	1.53
35	BB	425	G	N7-C5	-5.75	1.35	1.39
35	BB	897	C	P-O5'	-5.75	1.53	1.59
36	BC	66	G	P-O5'	-5.75	1.54	1.59
41	BH	31	A	O4'-C1'	-5.75	1.34	1.41
85	AA	365	G	O4'-C1'	-5.75	1.34	1.41
85	AA	1154	A	O3'-P	-5.75	1.54	1.61
34	BA	108	A	C3'-C2'	-5.75	1.46	1.52
34	BA	202	A	C1'-N9	-5.75	1.38	1.46
34	BA	260	A	N7-C5	-5.75	1.35	1.39
34	BA	397	A	C3'-C2'	-5.75	1.46	1.52
34	BA	467	A	C1'-N9	-5.75	1.38	1.46
34	BA	521	C	C4-N4	-5.75	1.28	1.33
34	BA	1355	G	C8-N7	-5.75	1.27	1.30
34	BA	1464	C	C2-N3	-5.75	1.31	1.35
34	BA	1847	G	C6-N1	-5.75	1.35	1.39
35	BB	890	U	P-O5'	-5.75	1.54	1.59
35	BB	960	C	P-O5'	-5.75	1.54	1.59
38	BE	87	U	C4-C5	-5.75	1.38	1.43
40	BG	81	G	N9-C4	-5.75	1.33	1.38
65	Bf	280	HIS	CB-CG	-5.75	1.39	1.50
85	AA	911	A	O3'-P	-5.75	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1217	U	C3'-C2'	-5.75	1.46	1.52
85	AA	2091	C	O3'-P	-5.75	1.54	1.61
34	BA	118	C	N3-C4	-5.75	1.29	1.33
34	BA	990	G	N7-C5	-5.75	1.35	1.39
34	BA	1653	G	N9-C8	-5.75	1.33	1.37
35	BB	585	U	N3-C4	-5.75	1.33	1.38
35	BB	1306	G	N9-C8	-5.75	1.33	1.37
35	BB	1319	U	C2'-C1'	-5.75	1.47	1.53
35	BB	1327	U	N1-C6	-5.75	1.32	1.38
39	BF	28	C	P-O5'	-5.75	1.54	1.59
40	BG	6	A	O3'-P	-5.75	1.54	1.61
34	BA	433	G	C4'-C3'	5.75	1.59	1.53
34	BA	1287	G	N7-C5	-5.75	1.35	1.39
34	BA	1787	U	N3-C4	-5.75	1.33	1.38
85	AA	964	C	C3'-O3'	5.75	1.50	1.42
85	AA	1940	A	P-O5'	-5.75	1.54	1.59
34	BA	8	G	C5-C4	-5.74	1.34	1.38
34	BA	1289	C	C1'-N1	-5.74	1.38	1.46
35	BB	26	C	C4'-C3'	-5.74	1.46	1.52
35	BB	1315	C	N1-C6	-5.74	1.33	1.37
37	BD	11	A	C6-N6	-5.74	1.29	1.33
38	BE	124	G	N7-C5	-5.74	1.35	1.39
41	BH	120	C	O3'-P	-5.74	1.54	1.61
85	AA	394	C	O3'-P	-5.74	1.54	1.61
85	AA	605	A	C3'-C2'	-5.74	1.46	1.52
85	AA	1641	A	C2'-C1'	-5.74	1.47	1.53
34	BA	332	U	P-O5'	-5.74	1.54	1.59
34	BA	367	G	P-O5'	-5.74	1.54	1.59
34	BA	697	A	N9-C4	-5.74	1.34	1.37
34	BA	1534	U	C1'-N1	-5.74	1.38	1.46
77	Br	133	PRO	CA-C	-5.74	1.41	1.52
85	AA	1285	C	C2-N3	-5.74	1.31	1.35
85	AA	1464	G	N1-C2	-5.74	1.33	1.37
34	BA	50	G	C2'-C1'	-5.74	1.47	1.53
34	BA	627	U	P-O5'	-5.74	1.54	1.59
34	BA	728	A	C3'-C2'	-5.74	1.46	1.52
34	BA	1004	U	N3-C4	-5.74	1.33	1.38
34	BA	1580	U	C2'-C1'	-5.74	1.47	1.53
35	BB	707	G	C2'-C1'	-5.74	1.47	1.53
38	BE	123	A	C2'-C1'	-5.74	1.47	1.53
40	BG	168	A	P-O5'	-5.74	1.54	1.59
41	BH	84	A	N3-C4	-5.74	1.31	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
65	Bf	295	PRO	CA-C	-5.74	1.41	1.52
85	AA	110	U	P-O5'	-5.74	1.54	1.59
85	AA	355	G	C3'-C2'	-5.74	1.46	1.52
85	AA	391	G	P-O5'	-5.74	1.54	1.59
85	AA	916	A	C2'-C1'	-5.74	1.47	1.53
85	AA	1143	C	O4'-C1'	-5.74	1.34	1.41
85	AA	1565	G	C2-N2	-5.74	1.28	1.34
85	AA	1960	C	O3'-P	-5.74	1.54	1.61
34	BA	666	C	C2-N3	-5.74	1.31	1.35
34	BA	956	G	C2-N2	-5.74	1.28	1.34
34	BA	1013	A	C8-N7	-5.74	1.27	1.31
34	BA	1163	G	P-O5'	-5.74	1.54	1.59
35	BB	31	U	C2'-C1'	-5.74	1.47	1.53
35	BB	664	A	N9-C8	-5.74	1.33	1.37
35	BB	1493	A	C3'-C2'	-5.74	1.46	1.52
36	BC	36	G	C2-N3	-5.74	1.28	1.32
40	BG	180	C	C4'-C3'	-5.74	1.46	1.52
85	AA	422	G	C6-N1	-5.74	1.35	1.39
85	AA	749	C	C2-N3	-5.74	1.31	1.35
85	AA	1584	U	C2'-C1'	-5.74	1.47	1.53
85	AA	2081	A	N9-C4	-5.74	1.34	1.37
34	BA	236	A	C8-N7	-5.74	1.27	1.31
35	BB	128	C	C3'-C2'	-5.74	1.46	1.52
35	BB	394	A	C3'-C2'	-5.74	1.46	1.52
35	BB	399	A	C4'-C3'	-5.74	1.46	1.52
35	BB	604	C	C5'-C4'	-5.74	1.44	1.51
35	BB	762	C	P-O5'	-5.74	1.54	1.59
35	BB	1048	A	C3'-C2'	-5.74	1.46	1.52
37	BD	108	G	O4'-C1'	-5.74	1.34	1.41
34	BA	255	G	C1'-N9	-5.74	1.38	1.46
34	BA	439	A	C1'-N9	-5.74	1.38	1.46
34	BA	799	A	C5'-C4'	5.74	1.58	1.51
34	BA	799	A	C8-N7	-5.74	1.27	1.31
34	BA	960	C	C4'-C3'	-5.74	1.46	1.52
34	BA	1176	C	O4'-C1'	-5.74	1.34	1.41
34	BA	1228	G	N7-C5	-5.74	1.35	1.39
35	BB	399	A	N9-C8	-5.74	1.33	1.37
35	BB	446	U	O4'-C1'	-5.74	1.34	1.41
35	BB	622	G	C1'-N9	-5.74	1.38	1.46
35	BB	679	G	C3'-C2'	-5.74	1.46	1.52
35	BB	811	C	C2-N3	-5.74	1.31	1.35
38	BE	45	G	C3'-C2'	-5.74	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	509	C	C3'-C2'	-5.74	1.46	1.52
85	AA	754	C	C2'-C1'	-5.74	1.47	1.53
85	AA	975	G	C1'-N9	-5.74	1.38	1.46
85	AA	1561	A	C1'-N9	-5.74	1.38	1.46
85	AA	1702	G	C2-N2	-5.74	1.28	1.34
85	AA	1858	G	C2'-C1'	-5.74	1.47	1.53
85	AA	2013	A	C3'-C2'	-5.74	1.46	1.52
34	BA	1033	G	C3'-C2'	-5.73	1.46	1.52
35	BB	1221	G	C6-N1	-5.73	1.35	1.39
35	BB	1378	U	C4'-C3'	-5.73	1.46	1.52
38	BE	82	C	C3'-O3'	5.73	1.50	1.42
85	AA	1307	U	P-O5'	-5.73	1.54	1.59
85	AA	1462	A	C6-N1	-5.73	1.31	1.35
34	BA	155	U	C2'-C1'	-5.73	1.47	1.53
34	BA	243	C	O3'-P	-5.73	1.54	1.61
35	BB	568	A	O3'-P	-5.73	1.54	1.61
37	BD	2	G	C6-N1	-5.73	1.35	1.39
37	BD	7	G	C4'-C3'	-5.73	1.46	1.52
38	BE	20	C	N3-C4	-5.73	1.29	1.33
40	BG	92	U	P-O5'	-5.73	1.54	1.59
41	BH	10	U	C5'-C4'	-5.73	1.44	1.51
85	AA	380	C	O3'-P	-5.73	1.54	1.61
85	AA	504	U	C4'-C3'	-5.73	1.46	1.52
85	AA	799	G	C2'-C1'	-5.73	1.47	1.53
85	AA	897	A	C4'-C3'	-5.73	1.46	1.52
85	AA	1292	A	C2'-C1'	-5.73	1.47	1.53
85	AA	1540	A	N3-C4	-5.73	1.31	1.34
85	AA	2054	G	C4'-C3'	-5.73	1.46	1.52
34	BA	365	A	C3'-C2'	-5.73	1.46	1.52
34	BA	1119	A	C2'-C1'	-5.73	1.47	1.53
34	BA	1585	A	N3-C4	-5.73	1.31	1.34
34	BA	1702	G	N9-C4	-5.73	1.33	1.38
35	BB	376	A	C6-N6	-5.73	1.29	1.33
35	BB	1296	A	C4'-C3'	-5.73	1.46	1.52
35	BB	1350	A	C4'-C3'	-5.73	1.46	1.52
37	BD	44	U	C2'-C1'	-5.73	1.47	1.53
34	BA	36	A	C5'-C4'	-5.73	1.44	1.51
34	BA	461	A	C8-N7	-5.73	1.27	1.31
34	BA	967	C	P-O5'	-5.73	1.54	1.59
34	BA	1510	C	C1'-N1	-5.73	1.38	1.46
85	AA	450	A	C8-N7	-5.73	1.27	1.31
85	AA	684	G	C2-N2	-5.73	1.28	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1984	A	C8-N7	-5.73	1.27	1.31
34	BA	1007	G	N9-C8	-5.73	1.33	1.37
34	BA	1832	A	C5-C4	-5.73	1.34	1.38
35	BB	132	G	O4'-C1'	-5.73	1.34	1.41
35	BB	844	G	C6-N1	-5.73	1.35	1.39
37	BD	95	G	C1'-N9	-5.73	1.38	1.46
40	BG	23	C	O5'-C5'	-5.73	1.33	1.42
85	AA	9	U	O3'-P	-5.73	1.54	1.61
85	AA	153	C	O3'-P	-5.73	1.54	1.61
85	AA	728	U	O3'-P	-5.73	1.54	1.61
85	AA	1671	G	C3'-C2'	-5.73	1.46	1.52
34	BA	680	C	C2-N3	-5.73	1.31	1.35
34	BA	1114	G	C2-N2	-5.73	1.28	1.34
34	BA	1379	G	P-O5'	-5.73	1.54	1.59
34	BA	1620	U	O3'-P	-5.73	1.54	1.61
35	BB	20	U	C2'-C1'	-5.73	1.47	1.53
35	BB	1495	U	C3'-C2'	-5.73	1.46	1.52
35	BB	1516	C	O3'-P	-5.73	1.54	1.61
38	BE	127	G	N9-C8	-5.73	1.33	1.37
85	AA	283	A	P-O5'	-5.73	1.54	1.59
85	AA	895	C	C4'-C3'	-5.73	1.46	1.52
85	AA	2193	A	C4'-O4'	-5.73	1.38	1.45
34	BA	557	U	C3'-O3'	5.72	1.50	1.42
34	BA	1061	A	C2'-C1'	-5.72	1.47	1.53
34	BA	1208	U	C3'-C2'	-5.72	1.46	1.52
34	BA	1817	G	C2'-C1'	-5.72	1.47	1.53
40	BG	155	A	C5-C4	-5.72	1.34	1.38
41	BH	120	C	O4'-C1'	-5.72	1.34	1.41
85	AA	1852	U	C5'-C4'	5.72	1.58	1.51
34	BA	195	G	C4'-C3'	-5.72	1.46	1.52
34	BA	202	A	C5-C4	-5.72	1.34	1.38
34	BA	465	A	N3-C4	-5.72	1.31	1.34
34	BA	979	G	N9-C8	-5.72	1.33	1.37
34	BA	1195	G	C3'-C2'	-5.72	1.46	1.52
35	BB	439	G	C6-N1	-5.72	1.35	1.39
35	BB	448	G	C6-N1	-5.72	1.35	1.39
35	BB	1019	C	N1-C6	-5.72	1.33	1.37
35	BB	1041	A	N3-C4	-5.72	1.31	1.34
36	BC	59	A	C3'-C2'	-5.72	1.46	1.52
40	BG	61	A	C6-N1	5.72	1.39	1.35
40	BG	147	U	O3'-P	-5.72	1.54	1.61
41	BH	54	U	C3'-C2'	-5.72	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	735	G	P-O5'	-5.72	1.54	1.59
85	AA	1130	G	O4'-C1'	-5.72	1.34	1.41
85	AA	2229	G	C2-N2	-5.72	1.28	1.34
34	BA	346	A	C2'-C1'	-5.72	1.47	1.53
34	BA	649	A	N7-C5	-5.72	1.35	1.39
34	BA	710	A	C3'-C2'	-5.72	1.46	1.52
34	BA	1152	A	C4'-O4'	-5.72	1.38	1.45
34	BA	1788	U	C3'-C2'	-5.72	1.46	1.52
35	BB	1272	G	C1'-N9	-5.72	1.38	1.46
41	BH	39	G	C2-N2	-5.72	1.28	1.34
85	AA	49	C	C2-N3	-5.72	1.31	1.35
85	AA	430	G	C2-N2	-5.72	1.28	1.34
85	AA	721	C	P-O5'	-5.72	1.54	1.59
85	AA	861	G	N9-C4	-5.72	1.33	1.38
85	AA	1595	G	N9-C4	-5.72	1.33	1.38
34	BA	214	A	O3'-P	-5.72	1.54	1.61
34	BA	252	A	C5-C4	-5.72	1.34	1.38
34	BA	272	A	P-O5'	-5.72	1.54	1.59
35	BB	139	G	O3'-P	-5.72	1.54	1.61
35	BB	404	A	N7-C5	-5.72	1.35	1.39
35	BB	586	U	O4'-C1'	-5.72	1.34	1.41
35	BB	1123	A	O3'-P	-5.72	1.54	1.61
35	BB	1282	G	C5'-C4'	-5.72	1.44	1.51
85	AA	164	G	C2-N2	-5.72	1.28	1.34
85	AA	622	G	C3'-C2'	-5.72	1.46	1.52
85	AA	1689	G	C2'-C1'	-5.72	1.47	1.53
34	BA	1521	C	C2'-C1'	-5.72	1.47	1.53
38	BE	166	G	N9-C8	-5.72	1.33	1.37
40	BG	182	G	C2'-C1'	-5.72	1.47	1.53
85	AA	191	C	C2-N3	-5.72	1.31	1.35
85	AA	2048	C	C3'-C2'	-5.72	1.46	1.52
34	BA	99	G	O4'-C1'	-5.72	1.34	1.41
34	BA	267	G	C5-C4	-5.72	1.34	1.38
34	BA	387	A	N3-C4	-5.72	1.31	1.34
34	BA	406	G	O3'-P	-5.72	1.54	1.61
34	BA	424	U	O3'-P	-5.72	1.54	1.61
34	BA	727	G	C2-N2	-5.72	1.28	1.34
34	BA	889	U	N1-C6	-5.72	1.32	1.38
34	BA	1215	U	C3'-C2'	-5.72	1.46	1.52
34	BA	1219	G	C5-C4	-5.72	1.34	1.38
34	BA	1335	A	C5'-C4'	-5.72	1.44	1.51
34	BA	1546	C	C4'-C3'	-5.72	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1681	U	C2-N3	-5.72	1.33	1.37
35	BB	57	G	C2'-C1'	-5.72	1.47	1.53
35	BB	93	A	C8-N7	-5.72	1.27	1.31
35	BB	577	U	N3-C4	-5.72	1.33	1.38
35	BB	794	G	N1-C2	-5.72	1.33	1.37
35	BB	1086	G	C3'-C2'	-5.72	1.46	1.52
35	BB	1335	G	N3-C4	-5.72	1.31	1.35
36	BC	88	A	C1'-N9	-5.72	1.38	1.46
37	BD	100	A	O3'-P	-5.72	1.54	1.61
40	BG	95	U	O3'-P	-5.72	1.54	1.61
41	BH	54	U	C2-N3	-5.72	1.33	1.37
85	AA	164	G	C1'-N9	-5.72	1.38	1.46
85	AA	176	C	C3'-C2'	-5.72	1.46	1.52
85	AA	189	G	N1-C2	-5.72	1.33	1.37
85	AA	1139	G	C5-C4	-5.72	1.34	1.38
85	AA	1181	U	C2'-C1'	-5.72	1.47	1.53
85	AA	1587	C	O3'-P	-5.72	1.54	1.61
85	AA	1762	G	O3'-P	-5.72	1.54	1.61
34	BA	188	C	O3'-P	-5.71	1.54	1.61
34	BA	488	C	C5'-C4'	5.71	1.58	1.51
34	BA	1395	C	O3'-P	-5.71	1.54	1.61
34	BA	1669	C	N1-C6	-5.71	1.33	1.37
35	BB	1401	G	C1'-N9	-5.71	1.38	1.46
36	BC	46	G	C1'-N9	-5.71	1.38	1.46
37	BD	17	G	N9-C4	-5.71	1.33	1.38
37	BD	68	C	C3'-C2'	-5.71	1.46	1.52
39	BF	32	G	C5-C6	-5.71	1.36	1.42
40	BG	9	G	N9-C8	-5.71	1.33	1.37
85	AA	57	G	N9-C4	5.71	1.42	1.38
85	AA	76	G	C2'-C1'	-5.71	1.47	1.53
85	AA	247	G	C8-N7	-5.71	1.27	1.30
85	AA	1202	G	P-O5'	-5.71	1.54	1.59
85	AA	1612	C	O3'-P	-5.71	1.54	1.61
85	AA	2227	A	C3'-C2'	-5.71	1.46	1.52
85	AA	2248	A	P-O5'	-5.71	1.54	1.59
34	BA	409	A	O4'-C1'	-5.71	1.34	1.41
34	BA	784	C	C4'-O4'	-5.71	1.38	1.45
34	BA	1448	G	C1'-N9	-5.71	1.38	1.46
35	BB	18	A	C3'-C2'	-5.71	1.46	1.52
35	BB	1062	G	C1'-N9	-5.71	1.38	1.46
40	BG	178	G	C2-N2	-5.71	1.28	1.34
85	AA	84	C	C3'-C2'	-5.71	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1794	U	O4'-C1'	-5.71	1.34	1.41
85	AA	2134	U	O3'-P	-5.71	1.54	1.61
34	BA	97	A	C8-N7	-5.71	1.27	1.31
34	BA	111	U	C2-N3	-5.71	1.33	1.37
34	BA	383	G	C5-C4	-5.71	1.34	1.38
34	BA	493	G	N1-C2	-5.71	1.33	1.37
34	BA	668	G	N7-C5	-5.71	1.35	1.39
34	BA	938	C	O3'-P	-5.71	1.54	1.61
34	BA	960	C	N1-C6	-5.71	1.33	1.37
34	BA	1073	G	O3'-P	-5.71	1.54	1.61
34	BA	1490	U	O4'-C1'	-5.71	1.34	1.41
34	BA	1647	G	C6-N1	-5.71	1.35	1.39
34	BA	1655	G	N1-C2	-5.71	1.33	1.37
35	BB	372	U	P-O5'	-5.71	1.54	1.59
35	BB	522	A	N9-C4	-5.71	1.34	1.37
35	BB	545	C	O3'-P	-5.71	1.54	1.61
35	BB	1176	G	C3'-C2'	-5.71	1.46	1.52
36	BC	123	G	C3'-C2'	-5.71	1.46	1.52
37	BD	15	U	C2-N3	-5.71	1.33	1.37
40	BG	130	G	C4'-C3'	-5.71	1.46	1.52
85	AA	233	C	C2'-C1'	-5.71	1.47	1.53
85	AA	275	A	O3'-P	-5.71	1.54	1.61
85	AA	328	U	C2'-C1'	-5.71	1.47	1.53
85	AA	782	G	N9-C4	-5.71	1.33	1.38
85	AA	2002	A	P-O5'	-5.71	1.54	1.59
85	AA	2137	A	C2'-C1'	-5.71	1.47	1.53
34	BA	1077	G	O3'-P	-5.71	1.54	1.61
34	BA	1434	U	C4'-C3'	-5.71	1.46	1.52
36	BC	123	G	C6-N1	5.71	1.43	1.39
39	BF	54	U	C3'-O3'	5.71	1.50	1.42
85	AA	1828	C	C2'-C1'	-5.71	1.47	1.53
85	AA	1892	G	C1'-N9	-5.71	1.38	1.46
34	BA	171	U	C2'-C1'	-5.71	1.47	1.53
34	BA	212	A	O3'-P	-5.71	1.54	1.61
34	BA	595	U	C5'-C4'	5.71	1.58	1.51
34	BA	797	A	O3'-P	-5.71	1.54	1.61
34	BA	908	G	C1'-N9	-5.71	1.38	1.46
34	BA	951	C	O3'-P	-5.71	1.54	1.61
34	BA	1062	G	C8-N7	-5.71	1.27	1.30
34	BA	1070	G	N9-C8	-5.71	1.33	1.37
34	BA	1253	G	C2'-C1'	-5.71	1.47	1.53
34	BA	1581	G	N9-C8	-5.71	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	63	A	C1'-N9	-5.71	1.38	1.46
35	BB	102	G	C4'-C3'	-5.71	1.46	1.52
35	BB	1145	G	C6-N1	-5.71	1.35	1.39
35	BB	1156	U	C2-N3	-5.71	1.33	1.37
35	BB	1259	A	C8-N7	-5.71	1.27	1.31
35	BB	1289	G	C3'-C2'	-5.71	1.46	1.52
35	BB	1329	G	N7-C5	-5.71	1.35	1.39
35	BB	1390	U	P-O5'	-5.71	1.54	1.59
35	BB	1449	G	C5-C6	-5.71	1.36	1.42
35	BB	1466	A	C8-N7	-5.71	1.27	1.31
36	BC	13	U	O3'-P	-5.71	1.54	1.61
39	BF	4	A	C5-C4	-5.71	1.34	1.38
85	AA	982	G	C4'-C3'	-5.71	1.46	1.52
85	AA	1676	G	C2-N2	-5.71	1.28	1.34
34	BA	28	C	C2'-C1'	-5.71	1.47	1.53
34	BA	243	C	C2'-C1'	-5.71	1.47	1.53
34	BA	790	G	N3-C4	-5.71	1.31	1.35
34	BA	1203	G	C3'-C2'	-5.71	1.46	1.52
35	BB	392	G	N9-C8	-5.71	1.33	1.37
35	BB	457	U	C3'-C2'	-5.71	1.46	1.52
35	BB	663	G	C8-N7	-5.71	1.27	1.30
35	BB	773	G	O4'-C1'	-5.71	1.34	1.41
35	BB	798	A	N9-C4	-5.71	1.34	1.37
36	BC	10	C	C5'-C4'	5.71	1.58	1.51
36	BC	62	A	C2'-C1'	-5.71	1.47	1.53
85	AA	125	A	P-O5'	-5.71	1.54	1.59
85	AA	885	A	C5'-C4'	5.71	1.58	1.51
85	AA	1094	G	C2'-C1'	-5.71	1.47	1.53
85	AA	1123	C	C3'-C2'	-5.71	1.46	1.52
85	AA	1464	G	O3'-P	-5.71	1.54	1.61
85	AA	1473	U	C3'-C2'	-5.71	1.46	1.52
34	BA	245	U	C2'-C1'	-5.71	1.47	1.53
34	BA	876	C	C2-N3	-5.71	1.31	1.35
34	BA	1839	G	C5-C4	-5.71	1.34	1.38
40	BG	79	U	N3-C4	-5.71	1.33	1.38
40	BG	181	C	C3'-C2'	-5.71	1.46	1.52
85	AA	698	G	N3-C4	-5.71	1.31	1.35
85	AA	931	G	N9-C8	-5.71	1.33	1.37
85	AA	1123	C	C2'-C1'	-5.71	1.47	1.53
85	AA	2011	C	C1'-N1	-5.71	1.38	1.46
34	BA	67	A	C5-C4	-5.70	1.34	1.38
34	BA	205	G	C8-N7	-5.70	1.27	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	330	A	C3'-C2'	-5.70	1.46	1.52
34	BA	886	G	C6-N1	-5.70	1.35	1.39
34	BA	1018	U	C2'-C1'	-5.70	1.47	1.53
34	BA	1036	G	C5-C4	-5.70	1.34	1.38
35	BB	439	G	C3'-C2'	-5.70	1.46	1.52
35	BB	700	C	C3'-C2'	-5.70	1.46	1.52
35	BB	1086	G	C5-C6	-5.70	1.36	1.42
35	BB	1153	G	C8-N7	-5.70	1.27	1.30
35	BB	1362	G	N3-C4	-5.70	1.31	1.35
35	BB	1454	G	O3'-P	-5.70	1.54	1.61
35	BB	1526	C	C2-N3	-5.70	1.31	1.35
36	BC	80	A	C2'-C1'	-5.70	1.47	1.53
37	BD	118	C	C3'-C2'	-5.70	1.46	1.52
40	BG	175	G	C2-N2	-5.70	1.28	1.34
85	AA	618	A	P-O5'	-5.70	1.54	1.59
85	AA	2098	A	C8-N7	-5.70	1.27	1.31
34	BA	242	U	O3'-P	-5.70	1.54	1.61
35	BB	539	G	P-O5'	-5.70	1.54	1.59
35	BB	1322	A	N7-C5	-5.70	1.35	1.39
85	AA	451	G	C6-N1	-5.70	1.35	1.39
34	BA	324	C	P-O5'	-5.70	1.54	1.59
34	BA	460	G	C6-N1	-5.70	1.35	1.39
34	BA	503	C	P-O5'	-5.70	1.54	1.59
34	BA	763	U	C4-C5	-5.70	1.38	1.43
34	BA	1418	G	C2'-C1'	-5.70	1.47	1.53
34	BA	1807	G	C3'-C2'	-5.70	1.46	1.52
35	BB	1080	U	C3'-C2'	-5.70	1.46	1.52
36	BC	5	U	C4'-C3'	-5.70	1.46	1.52
36	BC	58	G	C5-C4	-5.70	1.34	1.38
41	BH	32	U	N3-C4	-5.70	1.33	1.38
85	AA	258	G	N1-C2	-5.70	1.33	1.37
85	AA	707	U	C2'-C1'	-5.70	1.47	1.53
85	AA	880	A	P-O5'	-5.70	1.54	1.59
85	AA	1695	G	C5-C4	-5.70	1.34	1.38
85	AA	2145	G	N9-C4	-5.70	1.33	1.38
85	AA	2249	U	O3'-P	-5.70	1.54	1.61
34	BA	265	A	O3'-P	-5.70	1.54	1.61
34	BA	631	G	O3'-P	-5.70	1.54	1.61
34	BA	1442	A	N3-C4	5.70	1.38	1.34
34	BA	1557	G	O4'-C1'	-5.70	1.34	1.41
34	BA	1735	G	N1-C2	-5.70	1.33	1.37
34	BA	1748	G	P-O5'	-5.70	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1842	U	C2-N3	-5.70	1.33	1.37
35	BB	1271	A	C5-C4	-5.70	1.34	1.38
35	BB	1343	C	C2-N3	-5.70	1.31	1.35
35	BB	1346	A	C3'-C2'	-5.70	1.46	1.52
35	BB	1366	C	P-O5'	-5.70	1.54	1.59
59	BZ	60	PHE	CB-CG	-5.70	1.41	1.51
85	AA	442	G	C5-C6	-5.70	1.36	1.42
85	AA	1469	G	C6-N1	5.70	1.43	1.39
85	AA	1814	U	P-O5'	-5.70	1.54	1.59
85	AA	2061	C	O3'-P	-5.70	1.54	1.61
85	AA	2209	U	C2-N3	-5.70	1.33	1.37
85	AA	2228	G	C5-C4	-5.70	1.34	1.38
34	BA	672	G	C2-N2	-5.70	1.28	1.34
35	BB	9	G	O3'-P	-5.70	1.54	1.61
35	BB	619	A	C1'-N9	-5.70	1.38	1.46
35	BB	876	G	O3'-P	-5.70	1.54	1.61
35	BB	878	G	N9-C4	-5.70	1.33	1.38
35	BB	1054	G	C1'-N9	-5.70	1.38	1.46
35	BB	1210	U	O3'-P	-5.70	1.54	1.61
35	BB	1254	G	O3'-P	-5.70	1.54	1.61
36	BC	5	U	N1-C2	-5.70	1.33	1.38
37	BD	1	G	N1-C2	-5.70	1.33	1.37
41	BH	52	G	C4'-C3'	-5.70	1.46	1.52
85	AA	687	G	C2-N2	-5.70	1.28	1.34
85	AA	929	G	C8-N7	-5.70	1.27	1.30
34	BA	1481	U	P-O5'	-5.70	1.54	1.59
34	BA	1699	A	C3'-C2'	-5.70	1.46	1.52
35	BB	713	U	C2'-C1'	-5.70	1.47	1.53
35	BB	1314	G	C2'-C1'	-5.70	1.47	1.53
35	BB	1398	A	C2'-C1'	-5.70	1.47	1.53
38	BE	199	A	N9-C8	-5.70	1.33	1.37
40	BG	51	U	O3'-P	-5.70	1.54	1.61
85	AA	367	A	C8-N7	-5.70	1.27	1.31
85	AA	543	A	C1'-N9	-5.70	1.38	1.46
85	AA	1675	U	C2'-C1'	-5.70	1.47	1.53
85	AA	2187	G	C1'-N9	-5.70	1.38	1.46
34	BA	1302	C	C4-N4	-5.69	1.28	1.33
34	BA	1413	G	C2-N2	-5.69	1.28	1.34
34	BA	1543	A	N3-C4	-5.69	1.31	1.34
35	BB	134	G	C5-C4	-5.69	1.34	1.38
35	BB	663	G	C5-C4	-5.69	1.34	1.38
85	AA	672	U	C3'-C2'	-5.69	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	993	G	C5-C4	-5.69	1.34	1.38
85	AA	1246	G	C5-C4	-5.69	1.34	1.38
85	AA	1263	G	C8-N7	-5.69	1.27	1.30
85	AA	2006	G	C3'-C2'	-5.69	1.46	1.52
34	BA	941	G	C4'-C3'	-5.69	1.46	1.52
34	BA	1023	G	C8-N7	-5.69	1.27	1.30
34	BA	1039	G	N1-C2	-5.69	1.33	1.37
34	BA	1211	G	C5-C6	-5.69	1.36	1.42
34	BA	1286	C	C4-N4	-5.69	1.28	1.33
35	BB	1341	U	C3'-C2'	-5.69	1.46	1.52
35	BB	1489	A	N7-C5	-5.69	1.35	1.39
36	BC	63	G	C2'-C1'	-5.69	1.47	1.53
38	BE	196	C	C3'-C2'	-5.69	1.46	1.52
40	BG	148	C	C2-N3	-5.69	1.31	1.35
41	BH	57	A	O3'-P	-5.69	1.54	1.61
41	BH	107	A	C6-N1	-5.69	1.31	1.35
85	AA	435	A	N3-C4	-5.69	1.31	1.34
85	AA	2096	G	C1'-N9	-5.69	1.38	1.46
34	BA	519	G	N9-C8	-5.69	1.33	1.37
34	BA	525	A	P-O5'	-5.69	1.54	1.59
34	BA	801	U	N3-C4	-5.69	1.33	1.38
34	BA	1006	G	O4'-C1'	-5.69	1.34	1.41
34	BA	1449	U	N1-C2	-5.69	1.33	1.38
34	BA	1808	A	C5'-C4'	-5.69	1.44	1.51
35	BB	463	C	C2-N3	-5.69	1.31	1.35
35	BB	584	A	C3'-C2'	-5.69	1.46	1.52
35	BB	786	A	N3-C4	-5.69	1.31	1.34
35	BB	855	G	P-O5'	-5.69	1.54	1.59
35	BB	1257	A	N7-C5	-5.69	1.35	1.39
36	BC	26	U	N3-C4	-5.69	1.33	1.38
36	BC	160	C	C1'-N1	-5.69	1.38	1.46
38	BE	147	G	C2'-C1'	-5.69	1.47	1.53
41	BH	40	C	C4-N4	-5.69	1.28	1.33
65	Bf	139	GLY	CA-C	-5.69	1.42	1.51
85	AA	94	C	N1-C6	-5.69	1.33	1.37
85	AA	156	G	O3'-P	-5.69	1.54	1.61
85	AA	1549	G	C2-N2	-5.69	1.28	1.34
85	AA	2238	C	C1'-N1	-5.69	1.38	1.46
34	BA	233	U	C3'-C2'	-5.69	1.46	1.52
34	BA	783	U	C3'-C2'	-5.69	1.46	1.52
34	BA	1643	U	N1-C2	-5.69	1.33	1.38
34	BA	1814	U	C4'-C3'	-5.69	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	509	A	C2'-C1'	-5.69	1.47	1.53
40	BG	177	U	C1'-N1	-5.69	1.38	1.46
85	AA	130	G	C6-N1	-5.69	1.35	1.39
85	AA	166	C	C3'-C2'	-5.69	1.46	1.52
34	BA	167	U	C3'-C2'	-5.69	1.46	1.52
34	BA	762	A	C5-C4	-5.69	1.34	1.38
34	BA	941	G	N1-C2	-5.69	1.33	1.37
34	BA	1035	A	C2'-C1'	-5.69	1.47	1.53
34	BA	1141	C	C3'-C2'	-5.69	1.46	1.52
34	BA	1313	U	C4'-C3'	-5.69	1.46	1.52
34	BA	1661	U	C2'-C1'	-5.69	1.47	1.53
35	BB	79	U	N3-C4	-5.69	1.33	1.38
35	BB	402	G	C3'-C2'	-5.69	1.46	1.52
35	BB	470	C	P-O5'	-5.69	1.54	1.59
35	BB	1061	G	C2-N2	-5.69	1.28	1.34
36	BC	31	A	N3-C4	-5.69	1.31	1.34
36	BC	103	A	N7-C5	-5.69	1.35	1.39
37	BD	98	G	C2-N2	-5.69	1.28	1.34
40	BG	150	A	C5-C4	-5.69	1.34	1.38
85	AA	252	G	C5-C4	-5.69	1.34	1.38
85	AA	678	A	O3'-P	-5.69	1.54	1.61
85	AA	708	G	N3-C4	-5.69	1.31	1.35
85	AA	984	A	C3'-C2'	-5.69	1.46	1.52
85	AA	1199	C	C3'-C2'	-5.69	1.46	1.52
34	BA	478	G	C5-C4	-5.69	1.34	1.38
34	BA	1116	G	N1-C2	-5.69	1.33	1.37
34	BA	1651	C	C4'-C3'	-5.69	1.46	1.52
35	BB	99	G	O3'-P	-5.69	1.54	1.61
35	BB	671	A	C5'-C4'	5.69	1.58	1.51
36	BC	9	G	N9-C4	-5.69	1.33	1.38
85	AA	625	G	C3'-C2'	-5.69	1.46	1.52
85	AA	1633	A	C1'-N9	-5.69	1.38	1.46
85	AA	2135	A	C1'-N9	-5.69	1.38	1.46
34	BA	228	A	O3'-P	-5.68	1.54	1.61
34	BA	388	A	C6-N1	-5.68	1.31	1.35
34	BA	967	C	C2-N3	-5.68	1.31	1.35
34	BA	1048	C	C3'-C2'	-5.68	1.46	1.52
34	BA	1578	A	O4'-C1'	-5.68	1.34	1.41
34	BA	1692	U	N3-C4	-5.68	1.33	1.38
35	BB	11	A	C5-C4	-5.68	1.34	1.38
35	BB	43	G	C8-N7	-5.68	1.27	1.30
35	BB	421	U	N3-C4	-5.68	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	459	U	C2-N3	-5.68	1.33	1.37
35	BB	484	G	N1-C2	-5.68	1.33	1.37
35	BB	656	A	O3'-P	-5.68	1.54	1.61
35	BB	699	U	O3'-P	-5.68	1.54	1.61
35	BB	830	G	C6-N1	-5.68	1.35	1.39
35	BB	1068	G	C1'-N9	-5.68	1.38	1.46
37	BD	8	A	C5-C4	-5.68	1.34	1.38
38	BE	140	G	N7-C5	-5.68	1.35	1.39
40	BG	143	C	N1-C6	-5.68	1.33	1.37
85	AA	622	G	P-O5'	-5.68	1.54	1.59
85	AA	976	G	C5-C4	-5.68	1.34	1.38
34	BA	90	G	N9-C4	-5.68	1.33	1.38
34	BA	547	C	C2'-O2'	-5.68	1.34	1.41
34	BA	1547	G	C2-N2	-5.68	1.28	1.34
34	BA	1605	G	O4'-C1'	-5.68	1.34	1.41
35	BB	20	U	C3'-C2'	-5.68	1.46	1.52
35	BB	475	A	O3'-P	-5.68	1.54	1.61
35	BB	658	G	C2-N2	-5.68	1.28	1.34
35	BB	830	G	C4'-C3'	-5.68	1.46	1.52
35	BB	1453	G	N9-C8	-5.68	1.33	1.37
38	BE	43	A	O3'-P	-5.68	1.54	1.61
41	BH	59	G	C2-N2	-5.68	1.28	1.34
85	AA	68	A	C1'-N9	-5.68	1.38	1.46
85	AA	515	C	C4'-O4'	-5.68	1.38	1.45
85	AA	579	U	P-O5'	-5.68	1.54	1.59
85	AA	678	A	N9-C4	-5.68	1.34	1.37
85	AA	932	A	C2'-C1'	-5.68	1.47	1.53
85	AA	1178	A	P-O5'	-5.68	1.54	1.59
85	AA	1621	U	O3'-P	-5.68	1.54	1.61
85	AA	2114	U	C2-N3	-5.68	1.33	1.37
34	BA	925	G	C2'-C1'	-5.68	1.47	1.53
35	BB	67	A	N9-C4	-5.68	1.34	1.37
35	BB	377	A	C5-C4	-5.68	1.34	1.38
36	BC	53	A	C4'-C3'	-5.68	1.46	1.52
40	BG	178	G	C1'-N9	-5.68	1.38	1.46
85	AA	639	C	C4'-O4'	-5.68	1.38	1.45
85	AA	709	A	N9-C8	-5.68	1.33	1.37
85	AA	1062	U	P-O5'	-5.68	1.54	1.59
34	BA	250	G	C5'-C4'	5.68	1.58	1.51
34	BA	414	A	C6-N6	-5.68	1.29	1.33
34	BA	564	C	N1-C6	-5.68	1.33	1.37
34	BA	767	U	N1-C2	-5.68	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1072	U	C1'-N1	-5.68	1.38	1.46
34	BA	1301	G	N1-C2	-5.68	1.33	1.37
34	BA	1435	A	O4'-C1'	-5.68	1.34	1.41
35	BB	1053	G	N1-C2	-5.68	1.33	1.37
35	BB	1291	G	C2'-C1'	-5.68	1.47	1.53
36	BC	52	A	C3'-O3'	-5.68	1.34	1.42
36	BC	103	A	P-O5'	-5.68	1.54	1.59
38	BE	161	G	O3'-P	-5.68	1.54	1.61
85	AA	70	U	C1'-N1	-5.68	1.38	1.46
85	AA	1189	A	C5-C4	-5.68	1.34	1.38
85	AA	1458	G	C5-C4	-5.68	1.34	1.38
34	BA	36	A	C5-C4	-5.68	1.34	1.38
35	BB	527	U	C4-C5	-5.68	1.38	1.43
35	BB	1344	U	N3-C4	-5.68	1.33	1.38
85	AA	1111	A	P-O5'	-5.68	1.54	1.59
85	AA	1174	G	C2'-C1'	-5.68	1.47	1.53
34	BA	50	G	O3'-P	-5.68	1.54	1.61
34	BA	245	U	N3-C4	-5.68	1.33	1.38
34	BA	398	G	C1'-N9	-5.68	1.39	1.46
34	BA	1364	G	P-O5'	-5.68	1.54	1.59
34	BA	1425	G	C8-N7	-5.68	1.27	1.30
35	BB	416	U	C2'-C1'	-5.68	1.47	1.53
35	BB	568	A	C1'-N9	-5.68	1.39	1.46
35	BB	1103	A	N9-C4	-5.68	1.34	1.37
35	BB	1119	G	C4'-C3'	-5.68	1.46	1.52
38	BE	96	G	N9-C8	-5.68	1.33	1.37
38	BE	138	U	C2'-C1'	-5.68	1.47	1.53
40	BG	111	C	C2-N3	-5.68	1.31	1.35
85	AA	910	G	N1-C2	-5.68	1.33	1.37
85	AA	1532	G	C2'-C1'	-5.68	1.47	1.53
85	AA	1652	A	P-O5'	-5.68	1.54	1.59
34	BA	471	U	C5'-C4'	-5.67	1.44	1.51
34	BA	734	G	C1'-N9	-5.67	1.39	1.46
34	BA	820	C	C1'-N1	-5.67	1.39	1.46
34	BA	1556	A	N7-C5	-5.67	1.35	1.39
34	BA	1847	G	N7-C5	-5.67	1.35	1.39
35	BB	620	G	C2-N2	-5.67	1.28	1.34
35	BB	1247	C	O3'-P	-5.67	1.54	1.61
35	BB	1409	G	C1'-N9	-5.67	1.39	1.46
36	BC	67	U	N3-C4	-5.67	1.33	1.38
37	BD	57	C	C3'-C2'	-5.67	1.46	1.52
38	BE	7	U	N1-C6	-5.67	1.32	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	BH	25	A	N7-C5	-5.67	1.35	1.39
85	AA	637	U	N1-C2	-5.67	1.33	1.38
85	AA	993	G	P-O5'	-5.67	1.54	1.59
85	AA	1496	U	C2'-C1'	-5.67	1.47	1.53
85	AA	1586	C	O3'-P	-5.67	1.54	1.61
85	AA	1995	U	O3'-P	-5.67	1.54	1.61
85	AA	2163	G	C2'-C1'	-5.67	1.47	1.53
34	BA	1290	A	O3'-P	-5.67	1.54	1.61
34	BA	320	G	C1'-N9	-5.67	1.39	1.46
34	BA	457	A	N7-C5	-5.67	1.35	1.39
34	BA	1013	A	P-O5'	-5.67	1.54	1.59
34	BA	1229	G	N3-C4	-5.67	1.31	1.35
34	BA	1323	G	N9-C8	-5.67	1.33	1.37
34	BA	1581	G	C3'-C2'	5.67	1.59	1.52
34	BA	1653	G	C6-N1	-5.67	1.35	1.39
35	BB	1251	G	C3'-C2'	-5.67	1.46	1.52
38	BE	90	G	C2-N2	-5.67	1.28	1.34
40	BG	137	G	C2-N2	-5.67	1.28	1.34
40	BG	173	C	C3'-C2'	-5.67	1.46	1.52
85	AA	82	A	C3'-C2'	-5.67	1.46	1.52
85	AA	418	G	N9-C4	-5.67	1.33	1.38
85	AA	565	G	O3'-P	-5.67	1.54	1.61
85	AA	926	C	C2-N3	-5.67	1.31	1.35
85	AA	1719	C	C4-N4	-5.67	1.28	1.33
85	AA	1911	A	N9-C4	-5.67	1.34	1.37
34	BA	8	G	C6-N1	-5.67	1.35	1.39
34	BA	1475	G	N9-C8	-5.67	1.33	1.37
34	BA	1658	G	C1'-N9	-5.67	1.39	1.46
35	BB	562	A	C4'-O4'	-5.67	1.38	1.45
35	BB	1136	G	O4'-C1'	-5.67	1.34	1.41
40	BG	40	G	C5-C4	-5.67	1.34	1.38
85	AA	642	G	C2'-C1'	-5.67	1.47	1.53
85	AA	1897	A	N9-C4	5.67	1.41	1.37
34	BA	293	A	N3-C4	-5.67	1.31	1.34
34	BA	375	C	C2-N3	-5.67	1.31	1.35
34	BA	1323	G	N3-C4	-5.67	1.31	1.35
34	BA	1817	G	C2-N2	-5.67	1.28	1.34
35	BB	473	U	P-O5'	-5.67	1.54	1.59
35	BB	707	G	C6-N1	-5.67	1.35	1.39
35	BB	842	G	C6-N1	-5.67	1.35	1.39
35	BB	1116	U	O3'-P	-5.67	1.54	1.61
35	BB	1364	C	C1'-N1	-5.67	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1500	U	C2-N3	-5.67	1.33	1.37
35	BB	1509	G	C4'-C3'	-5.67	1.46	1.52
38	BE	192	A	N3-C4	-5.67	1.31	1.34
85	AA	195	C	C2'-C1'	-5.67	1.47	1.53
85	AA	1435	C	C4-N4	-5.67	1.28	1.33
85	AA	2040	A	N9-C4	-5.67	1.34	1.37
85	AA	2083	G	C4'-O4'	-5.67	1.38	1.45
85	AA	2113	U	O3'-P	-5.67	1.54	1.61
34	BA	144	C	N1-C6	-5.67	1.33	1.37
34	BA	673	U	C4'-C3'	5.67	1.59	1.53
34	BA	702	G	C1'-N9	-5.67	1.39	1.46
34	BA	928	C	O3'-P	-5.67	1.54	1.61
34	BA	983	A	C5-C4	-5.67	1.34	1.38
34	BA	1604	A	C5-C4	-5.67	1.34	1.38
34	BA	1691	G	C2'-C1'	-5.67	1.47	1.53
34	BA	1847	G	N9-C4	-5.67	1.33	1.38
35	BB	562	A	C1'-N9	-5.67	1.39	1.46
35	BB	751	A	P-O5'	-5.67	1.54	1.59
35	BB	765	G	C2'-C1'	-5.67	1.47	1.53
35	BB	1061	G	N1-C2	-5.67	1.33	1.37
35	BB	1154	C	C2'-C1'	-5.67	1.47	1.53
35	BB	1179	C	C2'-C1'	-5.67	1.47	1.53
35	BB	1388	A	C5'-C4'	-5.67	1.44	1.51
40	BG	66	C	C4'-O4'	-5.67	1.38	1.45
42	BI	49	HIS	CB-CG	-5.67	1.39	1.50
85	AA	487	G	C2-N2	-5.67	1.28	1.34
85	AA	625	G	O3'-P	-5.67	1.54	1.61
85	AA	1298	G	C2'-C1'	-5.67	1.47	1.53
85	AA	2051	G	O3'-P	-5.67	1.54	1.61
85	AA	2196	G	C2-N2	-5.67	1.28	1.34
34	BA	810	A	C5-C4	-5.67	1.34	1.38
34	BA	1260	G	C5'-C4'	-5.67	1.44	1.51
40	BG	115	C	C2'-C1'	-5.67	1.47	1.53
85	AA	257	U	C2'-C1'	-5.67	1.47	1.53
85	AA	1848	G	C2-N2	-5.67	1.28	1.34
7	A6	18	PHE	CB-CG	-5.66	1.41	1.51
34	BA	278	U	C3'-C2'	-5.66	1.46	1.52
34	BA	649	A	N9-C4	-5.66	1.34	1.37
34	BA	1097	G	N9-C8	-5.66	1.33	1.37
34	BA	1103	G	N9-C8	-5.66	1.33	1.37
34	BA	1643	U	C3'-C2'	-5.66	1.46	1.52
34	BA	1706	A	C4'-C3'	5.66	1.59	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1739	G	P-O5'	-5.66	1.54	1.59
35	BB	649	A	C3'-C2'	-5.66	1.46	1.52
35	BB	1095	G	N1-C2	-5.66	1.33	1.37
35	BB	1361	A	O3'-P	-5.66	1.54	1.61
35	BB	1408	G	N3-C4	-5.66	1.31	1.35
36	BC	109	A	C2'-C1'	-5.66	1.47	1.53
36	BC	115	G	N1-C2	-5.66	1.33	1.37
37	BD	119	U	P-O5'	-5.66	1.54	1.59
41	BH	48	G	C6-N1	-5.66	1.35	1.39
41	BH	75	G	N3-C4	-5.66	1.31	1.35
85	AA	16	G	N9-C4	-5.66	1.33	1.38
85	AA	76	G	N9-C4	-5.66	1.33	1.38
85	AA	247	G	N7-C5	-5.66	1.35	1.39
85	AA	267	U	C5'-C4'	5.66	1.58	1.51
85	AA	299	A	C5-C4	-5.66	1.34	1.38
85	AA	646	C	C2-N3	-5.66	1.31	1.35
85	AA	808	A	N7-C5	-5.66	1.35	1.39
34	BA	3	G	C2-N2	-5.66	1.28	1.34
34	BA	32	A	C1'-N9	-5.66	1.39	1.46
34	BA	108	A	C5-C4	-5.66	1.34	1.38
34	BA	183	G	C5-C4	-5.66	1.34	1.38
34	BA	381	A	C2'-C1'	-5.66	1.47	1.53
34	BA	1056	C	C2'-C1'	-5.66	1.47	1.53
34	BA	1138	C	C3'-C2'	-5.66	1.46	1.52
34	BA	1274	A	C5-C6	-5.66	1.35	1.41
34	BA	1365	U	O3'-P	-5.66	1.54	1.61
35	BB	425	G	P-O5'	-5.66	1.54	1.59
35	BB	716	G	N9-C4	5.66	1.42	1.38
37	BD	32	A	C2'-C1'	-5.66	1.47	1.53
85	AA	1102	C	C2'-C1'	-5.66	1.47	1.53
85	AA	1467	U	N1-C2	-5.66	1.33	1.38
85	AA	1505	G	O3'-P	-5.66	1.54	1.61
34	BA	255	G	C6-N1	-5.66	1.35	1.39
34	BA	366	G	C2-N2	-5.66	1.28	1.34
34	BA	803	U	N3-C4	-5.66	1.33	1.38
34	BA	1001	G	C3'-C2'	-5.66	1.46	1.52
35	BB	130	G	C2-N2	-5.66	1.28	1.34
35	BB	585	U	C3'-C2'	-5.66	1.46	1.52
35	BB	1286	G	N3-C4	-5.66	1.31	1.35
85	AA	229	U	C2-N3	-5.66	1.33	1.37
85	AA	691	U	C3'-C2'	-5.66	1.46	1.52
85	AA	986	U	C3'-C2'	-5.66	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1925	A	N3-C4	-5.66	1.31	1.34
34	BA	146	G	N9-C8	-5.66	1.33	1.37
34	BA	213	A	C5-C4	-5.66	1.34	1.38
34	BA	339	G	N9-C8	-5.66	1.33	1.37
34	BA	409	A	C1'-N9	-5.66	1.39	1.46
34	BA	571	G	N9-C8	5.66	1.41	1.37
34	BA	682	A	C2'-C1'	-5.66	1.47	1.53
34	BA	850	C	C4-N4	-5.66	1.28	1.33
34	BA	953	G	O4'-C1'	-5.66	1.34	1.41
34	BA	996	U	P-O5'	-5.66	1.54	1.59
34	BA	1589	U	C2'-C1'	-5.66	1.47	1.53
34	BA	1646	U	C4'-O4'	-5.66	1.38	1.45
35	BB	402	G	N1-C2	-5.66	1.33	1.37
35	BB	1021	C	C2-N3	-5.66	1.31	1.35
35	BB	1287	U	O3'-P	-5.66	1.54	1.61
35	BB	1450	G	C1'-N9	-5.66	1.39	1.46
36	BC	64	U	C3'-C2'	-5.66	1.46	1.52
39	BF	45	G	N1-C2	-5.66	1.33	1.37
40	BG	142	A	N9-C8	-5.66	1.33	1.37
41	BH	70	U	C2'-C1'	-5.66	1.47	1.53
85	AA	348	G	P-O5'	-5.66	1.54	1.59
85	AA	532	G	C2-N3	-5.66	1.28	1.32
85	AA	1809	G	C2-N2	-5.66	1.28	1.34
34	BA	1044	A	C5-C4	-5.66	1.34	1.38
34	BA	1172	C	C1'-N1	-5.66	1.39	1.46
34	BA	1302	C	C2-N3	-5.66	1.31	1.35
35	BB	58	G	N7-C5	-5.66	1.35	1.39
35	BB	584	A	N3-C4	-5.66	1.31	1.34
36	BC	160	C	C4'-O4'	-5.66	1.38	1.45
85	AA	1490	A	C4'-O4'	-5.66	1.38	1.45
85	AA	2175	U	C4'-C3'	-5.66	1.46	1.52
34	BA	197	A	N7-C5	-5.66	1.35	1.39
34	BA	1704	G	C2-N2	-5.66	1.28	1.34
34	BA	1841	A	C5-C4	-5.66	1.34	1.38
35	BB	255	A	C5'-C4'	5.66	1.58	1.51
35	BB	646	U	C2-N3	-5.66	1.33	1.37
35	BB	828	G	C5-C4	-5.66	1.34	1.38
35	BB	1155	U	C1'-N1	-5.66	1.39	1.46
35	BB	1189	C	N1-C6	-5.66	1.33	1.37
36	BC	67	U	C2'-C1'	-5.66	1.47	1.53
38	BE	88	G	C1'-N9	-5.66	1.39	1.46
38	BE	129	G	N9-C8	-5.66	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	59	G	C8-N7	-5.66	1.27	1.30
64	Be	113	VAL	CA-C	-5.66	1.38	1.52
85	AA	749	C	C2'-C1'	-5.66	1.47	1.53
85	AA	773	G	N9-C8	-5.66	1.33	1.37
85	AA	1596	A	O3'-P	-5.66	1.54	1.61
85	AA	1999	C	O3'-P	-5.66	1.54	1.61
85	AA	2235	C	C3'-C2'	-5.66	1.46	1.52
34	BA	268	U	C2-N3	-5.65	1.33	1.37
34	BA	454	G	N3-C4	-5.65	1.31	1.35
34	BA	868	C	C4'-O4'	5.65	1.52	1.45
35	BB	1336	G	O3'-P	-5.65	1.54	1.61
36	BC	118	U	C2'-C1'	-5.65	1.47	1.53
40	BG	164	U	C4'-C3'	-5.65	1.46	1.52
85	AA	21	U	C3'-C2'	-5.65	1.46	1.52
85	AA	1185	G	C2-N2	-5.65	1.28	1.34
34	BA	15	G	C2'-C1'	-5.65	1.47	1.53
34	BA	63	A	P-O5'	-5.65	1.54	1.59
34	BA	155	U	N3-C4	-5.65	1.33	1.38
34	BA	273	G	C6-N1	-5.65	1.35	1.39
34	BA	986	G	C2-N2	-5.65	1.28	1.34
34	BA	1723	U	N3-C4	-5.65	1.33	1.38
35	BB	1108	G	C2-N2	-5.65	1.28	1.34
37	BD	99	G	P-O5'	-5.65	1.54	1.59
38	BE	85	G	C2'-C1'	-5.65	1.47	1.53
39	BF	32	G	N9-C8	-5.65	1.33	1.37
41	BH	118	U	C4'-O4'	-5.65	1.38	1.45
85	AA	20	G	C5'-C4'	-5.65	1.44	1.51
85	AA	316	C	C2'-C1'	-5.65	1.47	1.53
85	AA	442	G	P-O5'	-5.65	1.54	1.59
85	AA	1241	A	N3-C4	-5.65	1.31	1.34
85	AA	1357	U	C2'-C1'	-5.65	1.47	1.53
85	AA	1795	C	P-O5'	-5.65	1.54	1.59
85	AA	2062	U	C4'-O4'	-5.65	1.38	1.45
34	BA	314	A	C2'-C1'	5.65	1.59	1.53
34	BA	542	A	C4'-C3'	-5.65	1.46	1.52
34	BA	979	G	C5-C4	-5.65	1.34	1.38
34	BA	1011	G	C3'-C2'	-5.65	1.46	1.52
34	BA	1145	U	C3'-C2'	-5.65	1.46	1.52
34	BA	1261	G	O3'-P	-5.65	1.54	1.61
36	BC	7	U	O3'-P	-5.65	1.54	1.61
37	BD	95	G	N9-C8	-5.65	1.33	1.37
38	BE	60	C	C4'-C3'	-5.65	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	109	C	C5'-C4'	5.65	1.58	1.51
38	BE	149	A	C2'-C1'	-5.65	1.47	1.53
38	BE	154	A	C2'-C1'	-5.65	1.47	1.53
38	BE	208	G	C5'-C4'	5.65	1.58	1.51
39	BF	48	G	C3'-C2'	-5.65	1.46	1.52
41	BH	2	U	C2-N3	-5.65	1.33	1.37
63	Bd	17	HIS	CB-CG	-5.65	1.39	1.50
85	AA	181	A	C8-N7	-5.65	1.27	1.31
85	AA	456	A	C4'-O4'	-5.65	1.38	1.45
85	AA	493	A	O4'-C1'	-5.65	1.34	1.41
85	AA	1192	C	O3'-P	-5.65	1.54	1.61
85	AA	1199	C	C1'-N1	-5.65	1.39	1.46
85	AA	1504	A	N3-C4	-5.65	1.31	1.34
85	AA	1520	A	C8-N7	-5.65	1.27	1.31
85	AA	2086	C	C4-C5	-5.65	1.38	1.43
85	AA	2138	G	C3'-C2'	-5.65	1.46	1.52
34	BA	234	A	C5-C4	-5.65	1.34	1.38
34	BA	1239	G	N7-C5	-5.65	1.35	1.39
34	BA	1286	C	N3-C4	-5.65	1.29	1.33
35	BB	373	C	N3-C4	-5.65	1.29	1.33
40	BG	1	G	C5-C4	-5.65	1.34	1.38
41	BH	74	G	N7-C5	5.65	1.42	1.39
85	AA	1557	U	O4'-C1'	-5.65	1.34	1.41
85	AA	1846	G	O3'-P	-5.65	1.54	1.61
34	BA	1050	A	N3-C4	-5.65	1.31	1.34
34	BA	1560	U	C4'-C3'	-5.65	1.47	1.52
34	BA	1640	G	N7-C5	-5.65	1.35	1.39
35	BB	58	G	C2-N2	-5.65	1.28	1.34
35	BB	1108	G	P-O5'	-5.65	1.54	1.59
36	BC	94	C	C2'-C1'	-5.65	1.47	1.53
36	BC	149	A	C2'-C1'	-5.65	1.47	1.53
38	BE	203	C	C2-N3	-5.65	1.31	1.35
39	BF	23	G	C6-N1	-5.65	1.35	1.39
40	BG	46	G	C5-C4	-5.65	1.34	1.38
47	BN	113	GLY	CA-C	-5.65	1.42	1.51
85	AA	278	C	O3'-P	-5.65	1.54	1.61
85	AA	1092	G	P-O5'	-5.65	1.54	1.59
85	AA	1134	G	C2-N2	-5.65	1.28	1.34
85	AA	1672	G	C3'-C2'	-5.65	1.46	1.52
34	BA	91	C	C1'-N1	-5.65	1.39	1.46
34	BA	387	A	C6-N1	-5.65	1.31	1.35
34	BA	1123	G	C6-N1	-5.65	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1562	G	C2-N2	-5.65	1.28	1.34
34	BA	1639	U	O4'-C1'	-5.65	1.34	1.41
40	BG	132	U	N1-C2	-5.65	1.33	1.38
85	AA	1517	G	N7-C5	-5.65	1.35	1.39
34	BA	88	C	N3-C4	-5.64	1.29	1.33
34	BA	957	A	C3'-C2'	-5.64	1.46	1.52
34	BA	1378	A	N9-C4	-5.64	1.34	1.37
34	BA	1784	G	C2'-C1'	-5.64	1.47	1.53
35	BB	449	C	O3'-P	-5.64	1.54	1.61
35	BB	487	A	O4'-C1'	-5.64	1.34	1.41
35	BB	548	A	N3-C4	-5.64	1.31	1.34
35	BB	826	G	C3'-C2'	-5.64	1.46	1.52
35	BB	997	G	C2'-C1'	-5.64	1.47	1.53
35	BB	1095	G	O3'-P	-5.64	1.54	1.61
35	BB	1221	G	N3-C4	-5.64	1.31	1.35
35	BB	1313	C	N1-C6	-5.64	1.33	1.37
36	BC	18	G	C1'-N9	-5.64	1.39	1.46
36	BC	19	A	C8-N7	-5.64	1.27	1.31
38	BE	202	C	C2-N3	-5.64	1.31	1.35
39	BF	56	C	C5'-C4'	-5.64	1.44	1.51
40	BG	14	G	O4'-C1'	-5.64	1.34	1.41
85	AA	93	G	C2'-C1'	-5.64	1.47	1.53
85	AA	887	A	O3'-P	-5.64	1.54	1.61
85	AA	1683	U	O3'-P	-5.64	1.54	1.61
85	AA	1879	U	C2-N3	-5.64	1.33	1.37
85	AA	1913	G	N7-C5	-5.64	1.35	1.39
34	BA	805	A	N9-C8	-5.64	1.33	1.37
34	BA	970	U	C3'-C2'	-5.64	1.46	1.52
34	BA	1182	U	P-O5'	-5.64	1.54	1.59
34	BA	1806	A	N7-C5	-5.64	1.35	1.39
35	BB	565	U	C2-N3	-5.64	1.33	1.37
35	BB	1161	G	C3'-C2'	-5.64	1.46	1.52
35	BB	1423	U	N3-C4	-5.64	1.33	1.38
35	BB	1514	G	C3'-C2'	-5.64	1.46	1.52
35	BB	1518	U	C2'-C1'	-5.64	1.47	1.53
36	BC	148	C	C2'-C1'	-5.64	1.47	1.53
37	BD	93	G	N9-C8	-5.64	1.33	1.37
40	BG	56	G	N7-C5	-5.64	1.35	1.39
41	BH	15	A	C1'-N9	-5.64	1.39	1.46
85	AA	440	U	C1'-N1	-5.64	1.39	1.46
85	AA	465	A	O3'-P	-5.64	1.54	1.61
85	AA	649	C	C3'-C2'	-5.64	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	865	G	C1'-N9	-5.64	1.39	1.46
85	AA	1529	A	P-O5'	-5.64	1.54	1.59
85	AA	2053	A	N3-C4	-5.64	1.31	1.34
85	AA	2193	A	C5-C4	-5.64	1.34	1.38
34	BA	622	G	O3'-P	-5.64	1.54	1.61
34	BA	930	A	C5-C4	-5.64	1.34	1.38
34	BA	1607	U	C5'-C4'	-5.64	1.44	1.51
34	BA	1616	A	C6-N1	5.64	1.39	1.35
35	BB	61	A	C4'-C3'	-5.64	1.47	1.52
35	BB	1157	G	C2'-C1'	-5.64	1.47	1.53
41	BH	59	G	N7-C5	-5.64	1.35	1.39
85	AA	452	A	O4'-C1'	-5.64	1.34	1.41
85	AA	958	C	O3'-P	-5.64	1.54	1.61
85	AA	1148	G	O3'-P	-5.64	1.54	1.61
85	AA	2125	A	N3-C4	-5.64	1.31	1.34
34	BA	37	A	N7-C5	-5.64	1.35	1.39
34	BA	437	G	C4'-C3'	-5.64	1.47	1.52
34	BA	1193	A	C5-C4	-5.64	1.34	1.38
34	BA	1239	G	C2-N2	-5.64	1.28	1.34
34	BA	1413	G	P-O5'	-5.64	1.54	1.59
34	BA	1601	C	C1'-N1	-5.64	1.39	1.46
35	BB	610	U	O4'-C1'	-5.64	1.34	1.41
36	BC	109	A	C5-C4	-5.64	1.34	1.38
38	BE	12	A	P-O5'	-5.64	1.54	1.59
40	BG	72	G	C2-N2	-5.64	1.28	1.34
40	BG	87	G	N9-C8	-5.64	1.33	1.37
40	BG	110	U	C5'-C4'	5.64	1.58	1.51
85	AA	13	U	O3'-P	-5.64	1.54	1.61
85	AA	418	G	C2-N2	-5.64	1.28	1.34
85	AA	442	G	O4'-C1'	-5.64	1.34	1.41
85	AA	631	G	O4'-C1'	-5.64	1.34	1.41
85	AA	1096	G	O3'-P	-5.64	1.54	1.61
85	AA	1187	G	C3'-C2'	-5.64	1.46	1.52
85	AA	1570	A	C2'-C1'	-5.64	1.47	1.53
35	BB	1033	U	N3-C4	-5.64	1.33	1.38
40	BG	24	A	C5'-C4'	-5.64	1.44	1.51
85	AA	382	G	C6-N1	-5.64	1.35	1.39
85	AA	819	G	N9-C4	5.64	1.42	1.38
34	BA	1804	A	C5-C6	-5.64	1.35	1.41
35	BB	1119	G	C2-N2	-5.64	1.28	1.34
35	BB	1379	U	C2-N3	-5.64	1.33	1.37
39	BF	31	U	C3'-C2'	-5.64	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	157	A	C6-N6	-5.64	1.29	1.33
41	BH	43	G	C6-N1	-5.64	1.35	1.39
85	AA	149	A	C5-C4	-5.64	1.34	1.38
85	AA	421	G	C1'-N9	-5.64	1.39	1.46
85	AA	506	G	N7-C5	-5.64	1.35	1.39
85	AA	1107	A	C3'-C2'	-5.64	1.46	1.52
85	AA	1109	G	C2-N2	-5.64	1.28	1.34
85	AA	1879	U	C3'-C2'	-5.64	1.46	1.52
34	BA	189	G	O3'-P	-5.63	1.54	1.61
34	BA	363	G	C6-N1	-5.63	1.35	1.39
34	BA	768	G	O3'-P	-5.63	1.54	1.61
34	BA	801	U	C3'-C2'	-5.63	1.46	1.52
34	BA	925	G	C3'-C2'	-5.63	1.46	1.52
34	BA	1490	U	C2-N3	-5.63	1.33	1.37
34	BA	1674	G	C6-N1	-5.63	1.35	1.39
35	BB	487	A	N9-C8	-5.63	1.33	1.37
35	BB	493	U	O3'-P	-5.63	1.54	1.61
35	BB	550	G	N7-C5	-5.63	1.35	1.39
35	BB	614	U	C4'-O4'	-5.63	1.38	1.45
35	BB	1109	A	C3'-C2'	-5.63	1.46	1.52
35	BB	1194	A	N3-C4	-5.63	1.31	1.34
36	BC	136	G	C5-C4	-5.63	1.34	1.38
38	BE	12	A	N9-C8	-5.63	1.33	1.37
39	BF	31	U	C4'-C3'	-5.63	1.47	1.52
40	BG	158	A	N7-C5	-5.63	1.35	1.39
41	BH	123	G	C5-C4	-5.63	1.34	1.38
85	AA	1363	U	P-O5'	-5.63	1.54	1.59
85	AA	2056	C	C2-N3	-5.63	1.31	1.35
85	AA	2150	G	N3-C4	-5.63	1.31	1.35
34	BA	89	G	C1'-N9	-5.63	1.39	1.46
34	BA	896	U	C1'-N1	5.63	1.57	1.48
34	BA	1439	C	C2-N3	-5.63	1.31	1.35
34	BA	1502	G	C8-N7	-5.63	1.27	1.30
34	BA	1602	A	C1'-N9	-5.63	1.39	1.46
35	BB	506	G	N1-C2	-5.63	1.33	1.37
35	BB	797	C	C4'-C3'	5.63	1.59	1.53
35	BB	1000	U	P-O5'	-5.63	1.54	1.59
35	BB	1021	C	C3'-C2'	-5.63	1.46	1.52
35	BB	1180	G	C6-N1	-5.63	1.35	1.39
36	BC	38	U	C2-N3	-5.63	1.33	1.37
39	BF	14	C	C4'-O4'	-5.63	1.38	1.45
85	AA	1230	U	C1'-N1	-5.63	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	568	G	C6-N1	-5.63	1.35	1.39
34	BA	773	A	N3-C4	-5.63	1.31	1.34
34	BA	773	A	N7-C5	-5.63	1.35	1.39
35	BB	463	C	C3'-C2'	-5.63	1.46	1.52
35	BB	1254	G	C5-C4	-5.63	1.34	1.38
35	BB	1374	U	C2-N3	-5.63	1.33	1.37
36	BC	19	A	C4'-C3'	-5.63	1.47	1.52
36	BC	118	U	P-O5'	-5.63	1.54	1.59
38	BE	129	G	C2'-C1'	-5.63	1.47	1.53
40	BG	25	G	C2-N2	-5.63	1.28	1.34
40	BG	152	G	C1'-N9	-5.63	1.39	1.46
41	BH	114	G	C5-C4	-5.63	1.34	1.38
85	AA	257	U	O3'-P	-5.63	1.54	1.61
85	AA	371	C	C2-N3	-5.63	1.31	1.35
85	AA	544	A	N9-C4	-5.63	1.34	1.37
85	AA	1040	U	O3'-P	-5.63	1.54	1.61
85	AA	1849	A	P-O5'	-5.63	1.54	1.59
34	BA	681	G	C3'-C2'	5.63	1.59	1.52
35	BB	1216	G	P-O5'	-5.63	1.54	1.59
35	BB	1494	G	N7-C5	-5.63	1.35	1.39
56	BW	73	ARG	CD-NE	5.63	1.56	1.46
85	AA	2042	G	C3'-C2'	-5.63	1.46	1.52
85	AA	2130	G	C4'-C3'	-5.63	1.47	1.52
34	BA	291	C	N1-C6	-5.63	1.33	1.37
34	BA	592	G	C5'-C4'	5.63	1.58	1.51
34	BA	707	C	N1-C6	-5.63	1.33	1.37
34	BA	1215	U	C4'-O4'	-5.63	1.38	1.45
34	BA	1512	C	C4'-O4'	-5.63	1.38	1.45
34	BA	1585	A	N7-C5	-5.63	1.35	1.39
35	BB	69	A	N7-C5	-5.63	1.35	1.39
35	BB	549	U	C4'-C3'	-5.63	1.47	1.52
35	BB	1487	G	O3'-P	-5.63	1.54	1.61
37	BD	58	G	C3'-C2'	-5.63	1.46	1.52
38	BE	43	A	C1'-N9	-5.63	1.39	1.46
85	AA	602	U	N1-C2	-5.63	1.33	1.38
85	AA	683	U	N1-C2	-5.63	1.33	1.38
85	AA	1273	C	C1'-N1	5.63	1.57	1.48
34	BA	184	C	C2-N3	-5.63	1.31	1.35
34	BA	1205	A	C4'-O4'	-5.63	1.38	1.45
34	BA	1382	G	P-O5'	-5.63	1.54	1.59
34	BA	1679	C	C4-N4	-5.63	1.28	1.33
35	BB	87	G	C4'-C3'	-5.63	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	369	A	C2'-C1'	-5.63	1.47	1.53
35	BB	370	A	C2'-C1'	-5.63	1.47	1.53
35	BB	659	C	N1-C6	-5.63	1.33	1.37
35	BB	662	G	C2-N2	-5.63	1.28	1.34
35	BB	1523	U	N3-C4	-5.63	1.33	1.38
85	AA	1112	G	N3-C4	-5.63	1.31	1.35
85	AA	1435	C	P-O5'	-5.63	1.54	1.59
85	AA	1734	A	C5'-C4'	5.63	1.58	1.51
85	AA	2049	U	O3'-P	-5.63	1.54	1.61
85	AA	2200	A	C5-C4	-5.63	1.34	1.38
34	BA	369	A	C4'-O4'	-5.62	1.38	1.45
35	BB	385	C	C2'-C1'	-5.62	1.47	1.53
35	BB	1348	C	C2'-C1'	-5.62	1.47	1.53
85	AA	34	G	C1'-N9	-5.62	1.39	1.46
85	AA	227	A	O3'-P	-5.62	1.54	1.61
34	BA	19	G	O3'-P	-5.62	1.54	1.61
34	BA	57	A	N9-C8	-5.62	1.33	1.37
34	BA	218	G	N9-C4	-5.62	1.33	1.38
34	BA	355	U	N3-C4	-5.62	1.33	1.38
34	BA	714	G	N7-C5	-5.62	1.35	1.39
34	BA	983	A	P-O5'	-5.62	1.54	1.59
34	BA	1194	G	N9-C8	-5.62	1.33	1.37
35	BB	32	C	C4'-O4'	-5.62	1.38	1.45
35	BB	33	A	C8-N7	-5.62	1.27	1.31
35	BB	94	A	C6-N1	-5.62	1.31	1.35
35	BB	102	G	P-O5'	-5.62	1.54	1.59
35	BB	112	G	C2-N2	-5.62	1.28	1.34
35	BB	553	U	C2'-C1'	-5.62	1.47	1.53
35	BB	574	G	N1-C2	-5.62	1.33	1.37
36	BC	98	C	C2'-C1'	-5.62	1.47	1.53
40	BG	30	C	C5'-C4'	-5.62	1.44	1.51
40	BG	162	A	C5-C6	-5.62	1.35	1.41
85	AA	428	G	C8-N7	-5.62	1.27	1.30
85	AA	513	G	C1'-N9	-5.62	1.39	1.46
85	AA	790	A	C1'-N9	-5.62	1.39	1.46
85	AA	1512	U	C2-N3	-5.62	1.33	1.37
85	AA	1542	A	C5-C4	-5.62	1.34	1.38
85	AA	1604	A	P-O5'	-5.62	1.54	1.59
85	AA	2081	A	N9-C8	-5.62	1.33	1.37
34	BA	378	C	C2'-C1'	-5.62	1.47	1.53
34	BA	386	A	C8-N7	-5.62	1.27	1.31
34	BA	482	C	C4'-C3'	-5.62	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	567	U	C3'-C2'	-5.62	1.46	1.52
34	BA	816	G	N7-C5	-5.62	1.35	1.39
34	BA	817	U	C1'-N1	-5.62	1.39	1.46
35	BB	1283	C	C2'-C1'	-5.62	1.47	1.53
37	BD	75	G	C6-N1	-5.62	1.35	1.39
38	BE	64	A	C1'-N9	-5.62	1.39	1.46
85	AA	187	C	N3-C4	5.62	1.37	1.33
85	AA	579	U	C3'-C2'	-5.62	1.46	1.52
85	AA	965	G	O3'-P	-5.62	1.54	1.61
85	AA	977	U	C2-N3	-5.62	1.33	1.37
85	AA	1194	U	N1-C2	-5.62	1.33	1.38
85	AA	1551	G	N9-C4	-5.62	1.33	1.38
85	AA	2108	C	C3'-C2'	-5.62	1.46	1.52
34	BA	284	U	C5'-C4'	5.62	1.58	1.51
34	BA	501	U	C3'-O3'	5.62	1.50	1.42
34	BA	1131	G	O3'-P	-5.62	1.54	1.61
35	BB	423	G	C5-C4	-5.62	1.34	1.38
35	BB	566	A	C6-N1	-5.62	1.31	1.35
35	BB	638	G	C4'-O4'	-5.62	1.38	1.45
35	BB	1050	A	C2'-C1'	-5.62	1.47	1.53
35	BB	1157	G	C5-C4	-5.62	1.34	1.38
35	BB	1464	G	C3'-O3'	5.62	1.50	1.42
85	AA	665	A	C2'-C1'	-5.62	1.47	1.53
85	AA	1121	U	N3-C4	-5.62	1.33	1.38
34	BA	38	G	N1-C2	-5.62	1.33	1.37
34	BA	354	G	C2-N2	-5.62	1.28	1.34
34	BA	480	G	C2-N2	-5.62	1.28	1.34
34	BA	664	C	N1-C6	-5.62	1.33	1.37
34	BA	686	U	N3-C4	-5.62	1.33	1.38
34	BA	1287	G	C6-N1	-5.62	1.35	1.39
34	BA	1465	C	C4-N4	-5.62	1.28	1.33
34	BA	1832	A	C6-N6	-5.62	1.29	1.33
35	BB	82	G	C2-N2	-5.62	1.28	1.34
35	BB	771	U	P-O5'	-5.62	1.54	1.59
35	BB	1103	A	O4'-C1'	-5.62	1.34	1.41
36	BC	109	A	N3-C4	-5.62	1.31	1.34
36	BC	116	C	C5'-C4'	-5.62	1.44	1.51
39	BF	2	G	N7-C5	-5.62	1.35	1.39
64	Be	230	PRO	CA-C	-5.62	1.41	1.52
85	AA	448	G	C2'-C1'	-5.62	1.47	1.53
85	AA	636	G	P-O5'	-5.62	1.54	1.59
85	AA	735	G	C2-N3	5.62	1.37	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1254	A	O3'-P	-5.62	1.54	1.61
85	AA	2060	G	C3'-C2'	-5.62	1.46	1.52
34	BA	403	A	N9-C8	-5.62	1.33	1.37
34	BA	825	G	N9-C8	-5.62	1.33	1.37
34	BA	1776	G	C3'-C2'	-5.62	1.46	1.52
35	BB	20	U	O3'-P	-5.62	1.54	1.61
35	BB	1530	U	N3-C4	-5.62	1.33	1.38
38	BE	47	U	P-O5'	-5.62	1.54	1.59
85	AA	184	A	O3'-P	-5.62	1.54	1.61
34	BA	95	C	N3-C4	-5.62	1.30	1.33
34	BA	241	U	C2'-C1'	-5.62	1.47	1.53
34	BA	800	G	N9-C4	5.62	1.42	1.38
34	BA	1076	U	N1-C2	-5.62	1.33	1.38
34	BA	1448	G	P-O5'	-5.62	1.54	1.59
35	BB	580	A	N3-C4	-5.62	1.31	1.34
35	BB	636	G	N1-C2	-5.62	1.33	1.37
35	BB	1043	C	P-O5'	-5.62	1.54	1.59
35	BB	1513	U	O3'-P	-5.62	1.54	1.61
36	BC	51	A	N3-C4	-5.62	1.31	1.34
36	BC	117	A	P-O5'	-5.62	1.54	1.59
39	BF	64	U	C2'-C1'	-5.62	1.47	1.53
41	BH	39	G	C3'-C2'	-5.62	1.46	1.52
85	AA	487	G	C3'-C2'	-5.62	1.46	1.52
85	AA	1002	G	N9-C4	-5.62	1.33	1.38
85	AA	1162	A	N9-C8	-5.62	1.33	1.37
34	BA	14	G	N1-C2	-5.61	1.33	1.37
34	BA	307	C	C1'-N1	-5.61	1.39	1.46
34	BA	1553	G	C2-N2	-5.61	1.28	1.34
35	BB	591	A	N7-C5	-5.61	1.35	1.39
36	BC	168	C	C3'-C2'	-5.61	1.46	1.52
37	BD	101	A	C3'-C2'	-5.61	1.46	1.52
38	BE	52	U	C3'-C2'	-5.61	1.46	1.52
38	BE	117	A	O4'-C1'	-5.61	1.34	1.41
41	BH	47	G	N1-C2	-5.61	1.33	1.37
41	BH	130	G	N1-C2	-5.61	1.33	1.37
85	AA	378	A	O3'-P	-5.61	1.54	1.61
85	AA	629	A	N3-C4	-5.61	1.31	1.34
85	AA	1224	C	C1'-N1	-5.61	1.39	1.46
85	AA	1923	A	C8-N7	-5.61	1.27	1.31
85	AA	2056	C	C2'-C1'	-5.61	1.47	1.53
34	BA	262	A	O3'-P	-5.61	1.54	1.61
34	BA	332	U	C2-N3	-5.61	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	995	A	O3'-P	-5.61	1.54	1.61
34	BA	1061	A	C1'-N9	-5.61	1.39	1.46
35	BB	27	C	C1'-N1	-5.61	1.39	1.46
35	BB	650	A	O3'-P	-5.61	1.54	1.61
35	BB	779	C	C3'-C2'	-5.61	1.46	1.52
35	BB	1239	A	C5-C4	-5.61	1.34	1.38
41	BH	13	C	C1'-N1	-5.61	1.39	1.46
85	AA	160	A	N9-C8	-5.61	1.33	1.37
85	AA	630	A	N3-C4	-5.61	1.31	1.34
85	AA	1508	A	C6-N6	-5.61	1.29	1.33
34	BA	92	G	N7-C5	-5.61	1.35	1.39
34	BA	107	C	P-O5'	-5.61	1.54	1.59
34	BA	191	G	C5-C4	-5.61	1.34	1.38
34	BA	378	C	N1-C6	-5.61	1.33	1.37
34	BA	400	A	O3'-P	-5.61	1.54	1.61
34	BA	925	G	C6-N1	-5.61	1.35	1.39
34	BA	1026	C	C4-N4	-5.61	1.28	1.33
34	BA	1268	C	C2-N3	-5.61	1.31	1.35
35	BB	48	G	C6-N1	-5.61	1.35	1.39
35	BB	996	G	C2'-C1'	-5.61	1.47	1.53
35	BB	1121	A	O3'-P	-5.61	1.54	1.61
35	BB	1342	C	C4-N4	-5.61	1.28	1.33
36	BC	67	U	O3'-P	-5.61	1.54	1.61
36	BC	108	A	C5-C4	-5.61	1.34	1.38
38	BE	93	U	N3-C4	-5.61	1.33	1.38
38	BE	183	C	C4-C5	-5.61	1.38	1.43
85	AA	105	A	P-O5'	-5.61	1.54	1.59
85	AA	359	A	C2'-C1'	-5.61	1.47	1.53
85	AA	393	C	O4'-C1'	-5.61	1.34	1.41
85	AA	1239	C	O3'-P	-5.61	1.54	1.61
85	AA	1564	U	C2'-C1'	-5.61	1.47	1.53
85	AA	1681	G	N7-C5	-5.61	1.35	1.39
85	AA	1845	G	C1'-N9	-5.61	1.39	1.46
85	AA	1924	C	C2'-C1'	-5.61	1.47	1.53
85	AA	2053	A	P-O5'	-5.61	1.54	1.59
34	BA	701	G	C1'-N9	-5.61	1.39	1.46
34	BA	937	G	N7-C5	-5.61	1.35	1.39
34	BA	1667	G	N1-C2	-5.61	1.33	1.37
34	BA	1809	G	N7-C5	-5.61	1.35	1.39
35	BB	1410	G	C2-N2	-5.61	1.28	1.34
37	BD	80	G	N1-C2	-5.61	1.33	1.37
85	AA	94	C	C2'-C1'	-5.61	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	329	G	N1-C2	-5.61	1.33	1.37
34	BA	498	A	C4'-O4'	-5.61	1.38	1.45
34	BA	682	A	O3'-P	-5.61	1.54	1.61
34	BA	772	G	C4'-O4'	5.61	1.52	1.45
34	BA	1615	A	C2'-C1'	-5.61	1.47	1.53
34	BA	1719	G	C1'-N9	-5.61	1.39	1.46
35	BB	1049	G	C1'-N9	-5.61	1.39	1.46
35	BB	1310	C	N1-C6	-5.61	1.33	1.37
36	BC	2	A	N3-C4	-5.61	1.31	1.34
36	BC	97	U	N1-C6	-5.61	1.32	1.38
38	BE	194	A	N3-C4	-5.61	1.31	1.34
41	BH	105	U	O4'-C1'	-5.61	1.34	1.41
85	AA	132	G	O3'-P	-5.61	1.54	1.61
85	AA	505	U	C1'-N1	-5.61	1.39	1.46
85	AA	926	C	C2'-C1'	-5.61	1.47	1.53
85	AA	993	G	C4'-C3'	-5.61	1.47	1.52
86	AB	48	C	P-O5'	-5.61	1.54	1.59
34	BA	81	C	C2'-C1'	-5.61	1.47	1.53
34	BA	121	A	N7-C5	-5.61	1.35	1.39
34	BA	225	A	N7-C5	-5.61	1.35	1.39
34	BA	475	A	C8-N7	-5.61	1.27	1.31
34	BA	798	G	C5-C6	-5.61	1.36	1.42
34	BA	843	G	C5'-C4'	-5.61	1.44	1.51
34	BA	916	A	N7-C5	-5.61	1.35	1.39
34	BA	1221	A	N3-C4	-5.61	1.31	1.34
34	BA	1543	A	C5-C4	-5.61	1.34	1.38
34	BA	1575	U	C2'-C1'	-5.61	1.47	1.53
34	BA	1710	C	C2'-C1'	-5.61	1.47	1.53
35	BB	698	C	C1'-N1	-5.61	1.39	1.46
35	BB	778	A	O4'-C1'	-5.61	1.34	1.41
35	BB	802	G	C2'-C1'	-5.61	1.47	1.53
35	BB	984	U	C4'-C3'	-5.61	1.47	1.52
35	BB	1351	G	C6-N1	-5.61	1.35	1.39
36	BC	28	C	C2'-C1'	-5.61	1.47	1.53
37	BD	55	A	C3'-C2'	-5.61	1.46	1.52
38	BE	113	C	C4-C5	-5.61	1.38	1.43
40	BG	105	A	N9-C8	-5.61	1.33	1.37
40	BG	113	G	N3-C4	-5.61	1.31	1.35
85	AA	293	A	C2'-C1'	-5.61	1.47	1.53
85	AA	631	G	N1-C2	-5.61	1.33	1.37
85	AA	675	A	C3'-C2'	-5.61	1.46	1.52
85	AA	1237	A	C2'-C1'	-5.61	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1489	G	C2'-C1'	-5.61	1.47	1.53
85	AA	1961	U	C2'-C1'	-5.61	1.47	1.53
86	AB	10	G	O3'-P	-5.61	1.54	1.61
34	BA	437	G	N9-C8	-5.60	1.33	1.37
34	BA	900	A	C5-C6	-5.60	1.36	1.41
34	BA	945	A	P-O5'	-5.60	1.54	1.59
34	BA	1037	C	C2-N3	-5.60	1.31	1.35
41	BH	16	A	C8-N7	-5.60	1.27	1.31
41	BH	125	U	O3'-P	-5.60	1.54	1.61
51	BR	83	TRP	C-N	-5.60	1.23	1.34
85	AA	350	U	O3'-P	-5.60	1.54	1.61
85	AA	843	U	P-O5'	-5.60	1.54	1.59
34	BA	933	U	N1-C6	-5.60	1.32	1.38
34	BA	1334	G	C5-C4	-5.60	1.34	1.38
34	BA	1511	C	N1-C6	-5.60	1.33	1.37
35	BB	547	A	N3-C4	-5.60	1.31	1.34
35	BB	867	C	P-O5'	-5.60	1.54	1.59
35	BB	1055	G	C2'-C1'	-5.60	1.47	1.53
35	BB	1436	U	C5'-C4'	5.60	1.58	1.51
85	AA	917	A	N3-C4	-5.60	1.31	1.34
85	AA	1220	A	C3'-C2'	-5.60	1.46	1.52
85	AA	1531	G	N9-C4	-5.60	1.33	1.38
85	AA	2033	C	O3'-P	-5.60	1.54	1.61
34	BA	447	U	C2'-C1'	-5.60	1.47	1.53
34	BA	1301	G	N7-C5	-5.60	1.35	1.39
34	BA	1442	A	C2-N3	5.60	1.38	1.33
36	BC	138	C	C2'-C1'	-5.60	1.47	1.53
38	BE	161	G	C6-N1	-5.60	1.35	1.39
85	AA	860	C	P-O5'	-5.60	1.54	1.59
85	AA	1264	U	C4'-C3'	-5.60	1.47	1.52
85	AA	2000	C	P-O5'	-5.60	1.54	1.59
34	BA	56	G	C5-C6	-5.60	1.36	1.42
34	BA	186	G	C6-N1	-5.60	1.35	1.39
34	BA	474	A	N3-C4	-5.60	1.31	1.34
34	BA	1192	A	C5-C4	-5.60	1.34	1.38
34	BA	1333	G	N1-C2	-5.60	1.33	1.37
34	BA	1455	C	C3'-C2'	-5.60	1.46	1.52
34	BA	1793	G	N9-C8	-5.60	1.33	1.37
35	BB	824	C	C5'-C4'	-5.60	1.44	1.51
35	BB	1322	A	C5-C4	-5.60	1.34	1.38
35	BB	1435	G	C2-N2	-5.60	1.28	1.34
40	BG	141	A	C5-C6	-5.60	1.36	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	471	U	C4'-O4'	-5.60	1.38	1.45
85	AA	626	G	C1'-N9	-5.60	1.39	1.46
85	AA	883	A	O4'-C1'	-5.60	1.34	1.41
85	AA	1156	A	P-O5'	-5.60	1.54	1.59
85	AA	2031	C	C3'-C2'	-5.60	1.46	1.52
34	BA	178	C	O3'-P	-5.60	1.54	1.61
34	BA	739	A	O3'-P	-5.60	1.54	1.61
34	BA	785	G	N9-C8	-5.60	1.33	1.37
34	BA	890	G	O4'-C1'	-5.60	1.34	1.41
34	BA	1201	G	C2-N2	-5.60	1.28	1.34
35	BB	750	G	P-O5'	-5.60	1.54	1.59
35	BB	996	G	C3'-C2'	-5.60	1.46	1.52
35	BB	1394	A	P-O5'	-5.60	1.54	1.59
36	BC	159	U	O4'-C1'	-5.60	1.34	1.41
37	BD	72	U	C4'-C3'	-5.60	1.47	1.52
38	BE	124	G	C8-N7	-5.60	1.27	1.30
38	BE	184	G	C5'-C4'	5.60	1.58	1.51
39	BF	26	U	P-O5'	-5.60	1.54	1.59
40	BG	100	G	C6-N1	-5.60	1.35	1.39
34	BA	1517	U	C2'-C1'	-5.60	1.47	1.53
35	BB	396	C	C3'-C2'	-5.60	1.46	1.52
38	BE	127	G	C6-N1	-5.60	1.35	1.39
85	AA	1148	G	N9-C4	-5.60	1.33	1.38
34	BA	244	A	C3'-C2'	-5.59	1.46	1.52
34	BA	427	G	C2-N2	-5.59	1.28	1.34
34	BA	734	G	C5'-C4'	-5.59	1.44	1.51
34	BA	835	U	C4'-C3'	-5.59	1.47	1.52
34	BA	892	C	C2'-C1'	-5.59	1.47	1.53
34	BA	1262	A	C4'-O4'	-5.59	1.38	1.45
35	BB	113	C	C3'-C2'	-5.59	1.46	1.52
35	BB	117	A	C1'-N9	-5.59	1.39	1.46
35	BB	587	A	O3'-P	-5.59	1.54	1.61
35	BB	1191	G	C2'-C1'	-5.59	1.47	1.53
38	BE	2	G	C2-N2	-5.59	1.28	1.34
40	BG	27	C	C4'-C3'	-5.59	1.47	1.52
40	BG	56	G	C2'-C1'	-5.59	1.47	1.53
40	BG	159	A	C2'-C1'	-5.59	1.47	1.53
41	BH	113	G	N7-C5	-5.59	1.35	1.39
64	Be	153	GLY	CA-C	-5.59	1.42	1.51
85	AA	107	A	C4'-C3'	-5.59	1.47	1.52
85	AA	402	G	C5-C6	-5.59	1.36	1.42
85	AA	1119	A	C2'-C1'	-5.59	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1678	U	N1-C2	-5.59	1.33	1.38
85	AA	2060	G	C2-N2	-5.59	1.28	1.34
34	BA	1578	A	O3'-P	-5.59	1.54	1.61
34	BA	1607	U	C2-N3	-5.59	1.33	1.37
35	BB	802	G	C4'-C3'	-5.59	1.47	1.52
35	BB	1271	A	N3-C4	-5.59	1.31	1.34
37	BD	117	U	C3'-C2'	-5.59	1.46	1.52
40	BG	1	G	C2'-C1'	-5.59	1.47	1.53
85	AA	133	G	O3'-P	-5.59	1.54	1.61
85	AA	1671	G	C4'-C3'	-5.59	1.47	1.52
34	BA	4	A	N3-C4	-5.59	1.31	1.34
34	BA	1349	A	O3'-P	-5.59	1.54	1.61
34	BA	1561	C	O3'-P	-5.59	1.54	1.61
35	BB	386	G	C5-C4	-5.59	1.34	1.38
35	BB	639	A	O4'-C1'	-5.59	1.34	1.41
35	BB	1239	A	C3'-C2'	-5.59	1.46	1.52
35	BB	1521	G	C1'-N9	-5.59	1.39	1.46
36	BC	134	G	N7-C5	-5.59	1.35	1.39
37	BD	13	A	C3'-C2'	-5.59	1.46	1.52
41	BH	33	G	C5-C6	-5.59	1.36	1.42
85	AA	481	A	C5-C4	-5.59	1.34	1.38
85	AA	728	U	C2-N3	-5.59	1.33	1.37
85	AA	1219	A	C2'-C1'	-5.59	1.47	1.53
85	AA	2033	C	C2'-C1'	-5.59	1.47	1.53
34	BA	569	C	C2-N3	-5.59	1.31	1.35
34	BA	747	G	C8-N7	-5.59	1.27	1.30
34	BA	819	G	N9-C8	-5.59	1.33	1.37
35	BB	635	A	C4'-C3'	-5.59	1.47	1.52
35	BB	636	G	C6-N1	-5.59	1.35	1.39
35	BB	816	U	C5'-C4'	-5.59	1.44	1.51
35	BB	843	G	O3'-P	-5.59	1.54	1.61
35	BB	1092	G	O3'-P	-5.59	1.54	1.61
35	BB	1275	A	N3-C4	-5.59	1.31	1.34
35	BB	1336	G	C8-N7	-5.59	1.27	1.30
85	AA	495	G	N9-C4	-5.59	1.33	1.38
85	AA	1267	A	C8-N7	-5.59	1.27	1.31
85	AA	1371	C	C2'-C1'	-5.59	1.47	1.53
85	AA	1478	G	P-O5'	-5.59	1.54	1.59
85	AA	1549	G	C4'-C3'	-5.59	1.47	1.52
85	AA	1715	C	P-O5'	-5.59	1.54	1.59
85	AA	1999	C	P-O5'	-5.59	1.54	1.59
34	BA	436	U	C5'-C4'	-5.59	1.44	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	22	A	C5-C4	-5.59	1.34	1.38
37	BD	81	C	N1-C6	-5.59	1.33	1.37
85	AA	304	G	C3'-O3'	5.59	1.50	1.42
34	BA	521	C	C1'-N1	-5.59	1.39	1.46
34	BA	1215	U	C1'-N1	-5.59	1.39	1.46
34	BA	1629	A	P-O5'	-5.59	1.54	1.59
35	BB	110	U	N3-C4	-5.59	1.33	1.38
35	BB	478	G	C3'-C2'	-5.59	1.46	1.52
35	BB	1037	A	C3'-C2'	-5.59	1.46	1.52
37	BD	101	A	O3'-P	-5.59	1.54	1.61
38	BE	194	A	P-O5'	-5.59	1.54	1.59
40	BG	6	A	N3-C4	-5.59	1.31	1.34
40	BG	167	C	P-O5'	-5.59	1.54	1.59
85	AA	391	G	C3'-C2'	-5.59	1.46	1.52
85	AA	866	U	C2-N3	-5.59	1.33	1.37
85	AA	1579	A	N9-C4	-5.59	1.34	1.37
85	AA	2083	G	C5-C4	-5.59	1.34	1.38
34	BA	1053	U	C3'-C2'	-5.58	1.46	1.52
35	BB	690	C	C2'-C1'	-5.58	1.47	1.53
85	AA	1997	G	C4'-C3'	5.58	1.59	1.53
85	AA	2089	G	C1'-N9	-5.58	1.39	1.46
85	AA	2208	G	C6-N1	-5.58	1.35	1.39
35	BB	517	G	C2'-C1'	-5.58	1.47	1.53
35	BB	604	C	O3'-P	-5.58	1.54	1.61
35	BB	1458	U	O3'-P	-5.58	1.54	1.61
37	BD	106	G	C2-N2	-5.58	1.28	1.34
38	BE	201	A	N3-C4	-5.58	1.31	1.34
85	AA	650	G	C2-N2	-5.58	1.28	1.34
85	AA	2040	A	P-O5'	-5.58	1.54	1.59
34	BA	93	A	C2'-C1'	-5.58	1.47	1.53
34	BA	470	C	C2-N3	-5.58	1.31	1.35
34	BA	514	U	O3'-P	-5.58	1.54	1.61
34	BA	956	G	C2'-C1'	-5.58	1.47	1.53
34	BA	1051	A	O3'-P	-5.58	1.54	1.61
34	BA	1197	U	C5'-C4'	-5.58	1.44	1.51
34	BA	1240	G	C2-N2	-5.58	1.28	1.34
34	BA	1779	U	C2-N3	-5.58	1.33	1.37
34	BA	1824	U	C2'-C1'	-5.58	1.47	1.53
35	BB	809	U	C3'-C2'	-5.58	1.46	1.52
35	BB	1371	G	C4'-C3'	-5.58	1.47	1.52
36	BC	46	G	N7-C5	-5.58	1.35	1.39
36	BC	56	G	C6-N1	-5.58	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	93	U	C3'-C2'	-5.58	1.46	1.52
40	BG	105	A	O4'-C1'	-5.58	1.34	1.41
41	BH	13	C	C4'-O4'	-5.58	1.38	1.45
85	AA	639	C	C4-C5	-5.58	1.38	1.43
85	AA	1185	G	N9-C8	-5.58	1.33	1.37
85	AA	1194	U	N3-C4	-5.58	1.33	1.38
85	AA	1221	G	C3'-C2'	-5.58	1.46	1.52
85	AA	1256	C	C3'-C2'	-5.58	1.46	1.52
85	AA	1267	A	N7-C5	-5.58	1.35	1.39
85	AA	1668	G	N7-C5	-5.58	1.35	1.39
85	AA	1679	U	N1-C6	-5.58	1.32	1.38
34	BA	94	G	C1'-N9	-5.58	1.39	1.46
34	BA	223	U	C2-N3	-5.58	1.33	1.37
34	BA	1369	C	C4-N4	-5.58	1.28	1.33
34	BA	1557	G	C5-C4	-5.58	1.34	1.38
34	BA	1608	C	C3'-C2'	-5.58	1.46	1.52
35	BB	112	G	C5-C4	-5.58	1.34	1.38
35	BB	1081	U	C2-N3	-5.58	1.33	1.37
85	AA	116	G	C5-C4	-5.58	1.34	1.38
85	AA	1986	G	C2-N2	-5.58	1.28	1.34
34	BA	14	G	O4'-C1'	-5.58	1.34	1.41
34	BA	109	A	C4'-C3'	-5.58	1.47	1.52
34	BA	255	G	N1-C2	-5.58	1.33	1.37
34	BA	384	U	C2-N3	-5.58	1.33	1.37
34	BA	398	G	C5-C4	-5.58	1.34	1.38
34	BA	965	A	C3'-C2'	-5.58	1.46	1.52
34	BA	1098	G	C2'-C1'	-5.58	1.47	1.53
35	BB	573	C	C2'-C1'	-5.58	1.47	1.53
35	BB	622	G	C2-N2	-5.58	1.28	1.34
35	BB	665	A	C5-C6	-5.58	1.36	1.41
35	BB	1286	G	N1-C2	-5.58	1.33	1.37
35	BB	1296	A	C1'-N9	-5.58	1.39	1.46
38	BE	188	C	O3'-P	-5.58	1.54	1.61
40	BG	26	G	C8-N7	-5.58	1.27	1.30
40	BG	160	C	C3'-C2'	-5.58	1.46	1.52
85	AA	598	C	C4'-C3'	-5.58	1.47	1.52
85	AA	691	U	O3'-P	-5.58	1.54	1.61
85	AA	874	A	N9-C4	-5.58	1.34	1.37
85	AA	960	G	C8-N7	-5.58	1.27	1.30
85	AA	1636	C	C2'-C1'	-5.58	1.47	1.53
85	AA	1647	G	C2'-C1'	-5.58	1.47	1.53
34	BA	54	A	C4'-C3'	-5.58	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	378	C	C1'-N1	-5.58	1.39	1.46
34	BA	458	G	C8-N7	-5.58	1.27	1.30
34	BA	1491	U	C3'-C2'	5.58	1.59	1.52
36	BC	66	G	C8-N7	-5.58	1.27	1.30
85	AA	1490	A	N9-C8	-5.58	1.33	1.37
85	AA	2133	A	N3-C4	-5.58	1.31	1.34
34	BA	38	G	C3'-C2'	-5.58	1.46	1.52
34	BA	332	U	C2'-C1'	-5.58	1.47	1.53
34	BA	860	G	C3'-C2'	-5.58	1.46	1.52
35	BB	1169	A	N9-C4	-5.58	1.34	1.37
35	BB	1538	G	C1'-N9	-5.58	1.39	1.46
38	BE	32	U	C5'-C4'	-5.58	1.44	1.51
38	BE	63	C	C4'-O4'	-5.58	1.38	1.45
38	BE	110	U	C4'-O4'	-5.58	1.38	1.45
38	BE	117	A	C8-N7	-5.58	1.27	1.31
38	BE	194	A	C6-N6	-5.58	1.29	1.33
40	BG	16	G	C6-N1	-5.58	1.35	1.39
41	BH	40	C	C2-N3	-5.58	1.31	1.35
41	BH	113	G	C1'-N9	-5.58	1.39	1.46
85	AA	1117	G	C2'-C1'	-5.58	1.47	1.53
85	AA	2240	G	N9-C8	-5.58	1.33	1.37
86	AB	68	C	C2'-C1'	-5.58	1.47	1.53
34	BA	178	C	C1'-N1	-5.57	1.39	1.46
34	BA	256	A	O3'-P	-5.57	1.54	1.61
34	BA	588	C	C2'-C1'	-5.57	1.47	1.53
34	BA	1007	G	N7-C5	-5.57	1.35	1.39
34	BA	1201	G	N1-C2	-5.57	1.33	1.37
34	BA	1564	A	C2'-C1'	-5.57	1.47	1.53
35	BB	54	U	C3'-C2'	-5.57	1.46	1.52
35	BB	736	G	N3-C4	-5.57	1.31	1.35
40	BG	101	G	C2'-C1'	-5.57	1.47	1.53
40	BG	131	U	O3'-P	-5.57	1.54	1.61
85	AA	817	G	N9-C8	-5.57	1.33	1.37
85	AA	1116	G	C6-N1	-5.57	1.35	1.39
34	BA	1631	U	P-O5'	5.57	1.65	1.59
35	BB	587	A	N9-C8	-5.57	1.33	1.37
35	BB	1337	C	C2-N3	-5.57	1.31	1.35
36	BC	49	G	C1'-N9	-5.57	1.39	1.46
37	BD	110	G	P-O5'	-5.57	1.54	1.59
38	BE	97	G	N7-C5	-5.57	1.35	1.39
40	BG	4	A	N9-C8	-5.57	1.33	1.37
85	AA	448	G	C5-C4	-5.57	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	536	C	O4'-C1'	-5.57	1.34	1.41
34	BA	1221	A	C6-N6	-5.57	1.29	1.33
34	BA	1295	U	C5'-C4'	5.57	1.58	1.51
34	BA	1311	G	C2-N2	-5.57	1.28	1.34
34	BA	1700	C	C2'-C1'	-5.57	1.47	1.53
35	BB	1041	A	O3'-P	-5.57	1.54	1.61
35	BB	1279	C	C2'-C1'	-5.57	1.47	1.53
85	AA	384	C	N1-C6	-5.57	1.33	1.37
85	AA	838	G	O3'-P	-5.57	1.54	1.61
85	AA	1838	C	O3'-P	-5.57	1.54	1.61
85	AA	1905	A	N9-C4	5.57	1.41	1.37
85	AA	2009	A	N7-C5	-5.57	1.35	1.39
34	BA	257	G	C1'-N9	-5.57	1.39	1.46
34	BA	368	U	P-O5'	-5.57	1.54	1.59
34	BA	567	U	C2-N3	-5.57	1.33	1.37
34	BA	912	G	C1'-N9	-5.57	1.39	1.46
34	BA	1475	G	C8-N7	-5.57	1.27	1.30
34	BA	1640	G	C4'-O4'	-5.57	1.38	1.45
35	BB	450	A	C5-C6	-5.57	1.36	1.41
35	BB	503	G	C3'-C2'	-5.57	1.46	1.52
35	BB	1356	G	O3'-P	-5.57	1.54	1.61
85	AA	791	C	P-O5'	-5.57	1.54	1.59
85	AA	2094	U	P-O5'	-5.57	1.54	1.59
34	BA	165	C	C4-C5	-5.57	1.38	1.43
34	BA	800	G	O3'-P	-5.57	1.54	1.61
34	BA	902	C	C3'-C2'	-5.57	1.46	1.52
34	BA	1255	G	C6-N1	-5.57	1.35	1.39
34	BA	1593	U	N3-C4	-5.57	1.33	1.38
34	BA	1732	A	C2'-C1'	-5.57	1.47	1.53
35	BB	457	U	P-O5'	-5.57	1.54	1.59
35	BB	1048	A	N9-C8	-5.57	1.33	1.37
36	BC	41	A	N3-C4	-5.57	1.31	1.34
37	BD	81	C	C1'-N1	-5.57	1.39	1.46
41	BH	5	G	N9-C4	-5.57	1.33	1.38
47	BN	80	PHE	CB-CG	-5.57	1.41	1.51
85	AA	569	A	C2'-C1'	-5.57	1.47	1.53
85	AA	1250	A	O3'-P	-5.57	1.54	1.61
85	AA	2165	C	C2-N3	-5.57	1.31	1.35
34	BA	87	G	N9-C8	-5.57	1.33	1.37
34	BA	196	A	N1-C2	-5.57	1.29	1.34
34	BA	454	G	C5-C4	-5.57	1.34	1.38
34	BA	946	A	C8-N7	-5.57	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1363	A	O3'-P	-5.57	1.54	1.61
34	BA	1605	G	P-O5'	-5.57	1.54	1.59
35	BB	138	A	N9-C4	5.57	1.41	1.37
35	BB	582	G	N9-C4	-5.57	1.33	1.38
35	BB	783	U	N1-C6	-5.57	1.32	1.38
35	BB	986	C	C4'-O4'	-5.57	1.38	1.45
85	AA	642	G	C3'-C2'	-5.57	1.46	1.52
85	AA	1154	A	C4'-C3'	-5.57	1.47	1.52
85	AA	2122	A	C2'-O2'	-5.57	1.34	1.41
85	AA	2142	A	C5-C4	-5.57	1.34	1.38
34	BA	1219	G	C1'-N9	-5.56	1.39	1.46
35	BB	388	C	C2'-C1'	-5.56	1.47	1.53
85	AA	41	G	C2'-C1'	-5.56	1.47	1.53
85	AA	255	A	C1'-N9	-5.56	1.39	1.46
85	AA	458	C	P-O5'	-5.56	1.54	1.59
85	AA	521	A	C8-N7	-5.56	1.27	1.31
85	AA	1121	U	C3'-C2'	-5.56	1.46	1.52
85	AA	1350	A	P-O5'	-5.56	1.54	1.59
34	BA	90	G	C1'-N9	-5.56	1.39	1.46
34	BA	613	A	C2'-C1'	-5.56	1.47	1.53
34	BA	682	A	C3'-C2'	-5.56	1.46	1.52
34	BA	1193	A	N9-C4	-5.56	1.34	1.37
34	BA	1796	A	C3'-C2'	-5.56	1.46	1.52
35	BB	372	U	O4'-C1'	-5.56	1.34	1.41
35	BB	814	A	C5-C4	-5.56	1.34	1.38
35	BB	872	A	C2'-C1'	-5.56	1.47	1.53
35	BB	1066	G	N9-C4	-5.56	1.33	1.38
35	BB	1371	G	C2-N2	-5.56	1.28	1.34
35	BB	1404	A	C6-N1	-5.56	1.31	1.35
37	BD	78	C	C3'-C2'	-5.56	1.46	1.52
38	BE	154	A	N3-C4	-5.56	1.31	1.34
85	AA	654	A	O4'-C1'	-5.56	1.34	1.41
85	AA	1507	G	C2-N2	-5.56	1.28	1.34
85	AA	1703	A	C1'-N9	-5.56	1.39	1.46
34	BA	296	G	C3'-O3'	5.56	1.50	1.42
34	BA	1246	G	N9-C8	-5.56	1.33	1.37
34	BA	1548	A	C4'-O4'	-5.56	1.38	1.45
35	BB	404	A	N3-C4	-5.56	1.31	1.34
35	BB	469	G	C2'-C1'	-5.56	1.47	1.53
35	BB	1343	C	C3'-C2'	-5.56	1.46	1.52
36	BC	108	A	N7-C5	-5.56	1.35	1.39
38	BE	10	G	N9-C4	-5.56	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	145	A	P-O5'	-5.56	1.54	1.59
41	BH	42	U	C2'-C1'	-5.56	1.47	1.53
85	AA	393	C	C2'-C1'	-5.56	1.47	1.53
85	AA	486	G	C5'-C4'	-5.56	1.44	1.51
85	AA	1812	C	O4'-C1'	-5.56	1.34	1.41
85	AA	2204	A	C4'-C3'	-5.56	1.47	1.52
34	BA	791	A	C2'-C1'	-5.56	1.47	1.53
34	BA	1125	G	O3'-P	-5.56	1.54	1.61
34	BA	1159	A	O3'-P	-5.56	1.54	1.61
34	BA	1413	G	C1'-N9	-5.56	1.39	1.46
35	BB	9	G	C5-C4	-5.56	1.34	1.38
35	BB	136	A	C1'-N9	-5.56	1.39	1.46
35	BB	415	A	C4'-C3'	-5.56	1.47	1.52
35	BB	1023	G	C1'-N9	-5.56	1.39	1.46
35	BB	1333	U	C1'-N1	5.56	1.57	1.48
35	BB	1489	A	C1'-N9	-5.56	1.39	1.46
40	BG	14	G	N7-C5	-5.56	1.35	1.39
40	BG	77	U	O3'-P	-5.56	1.54	1.61
85	AA	886	A	C2'-C1'	-5.56	1.47	1.53
85	AA	1676	G	C4'-O4'	-5.56	1.38	1.45
85	AA	2016	A	O3'-P	-5.56	1.54	1.61
85	AA	2078	A	C5'-C4'	5.56	1.58	1.51
85	AA	2182	A	C1'-N9	-5.56	1.39	1.46
34	BA	15	G	C2-N2	-5.56	1.28	1.34
34	BA	52	G	N7-C5	-5.56	1.35	1.39
34	BA	599	U	P-O5'	-5.56	1.54	1.59
34	BA	1119	A	N3-C4	-5.56	1.31	1.34
34	BA	1260	G	C2-N3	-5.56	1.28	1.32
34	BA	1496	G	P-O5'	-5.56	1.54	1.59
34	BA	1663	U	C2-N3	-5.56	1.33	1.37
35	BB	322	G	O3'-P	-5.56	1.54	1.61
35	BB	403	U	N3-C4	-5.56	1.33	1.38
35	BB	507	G	C6-N1	-5.56	1.35	1.39
35	BB	530	C	C4-N4	-5.56	1.28	1.33
35	BB	582	G	C5-C4	-5.56	1.34	1.38
35	BB	1062	G	N9-C8	-5.56	1.33	1.37
35	BB	1532	C	C2-N3	-5.56	1.31	1.35
35	BB	1544	A	C3'-C2'	-5.56	1.46	1.52
36	BC	36	G	N7-C5	-5.56	1.35	1.39
37	BD	85	C	O3'-P	-5.56	1.54	1.61
38	BE	121	G	C4'-C3'	-5.56	1.47	1.52
40	BG	44	G	C1'-N9	-5.56	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	104	A	O3'-P	-5.56	1.54	1.61
41	BH	32	U	C4-O4	-5.56	1.19	1.23
41	BH	107	A	C6-N6	-5.56	1.29	1.33
85	AA	2	A	C5-C4	-5.56	1.34	1.38
85	AA	2024	U	P-O5'	-5.56	1.54	1.59
85	AA	2039	G	O4'-C1'	-5.56	1.34	1.41
34	BA	235	C	N1-C6	-5.56	1.33	1.37
34	BA	266	G	C2'-C1'	-5.56	1.47	1.53
34	BA	315	U	C2'-C1'	-5.56	1.47	1.53
34	BA	360	C	P-O5'	-5.56	1.54	1.59
34	BA	1274	A	O3'-P	-5.56	1.54	1.61
35	BB	95	A	C5-C4	-5.56	1.34	1.38
35	BB	500	C	N1-C6	-5.56	1.33	1.37
35	BB	798	A	C5-C6	-5.56	1.36	1.41
35	BB	963	G	C2'-C1'	-5.56	1.47	1.53
35	BB	1153	G	C5-C6	-5.56	1.36	1.42
35	BB	1491	G	O3'-P	-5.56	1.54	1.61
85	AA	1431	U	O3'-P	-5.56	1.54	1.61
85	AA	2007	G	C2-N2	-5.56	1.28	1.34
34	BA	53	G	C2-N2	-5.55	1.28	1.34
34	BA	455	A	C4'-C3'	-5.55	1.47	1.52
34	BA	546	U	C3'-C2'	-5.55	1.46	1.52
34	BA	632	U	C2'-C1'	-5.55	1.47	1.53
34	BA	744	G	P-O5'	-5.55	1.54	1.59
34	BA	809	U	C3'-C2'	-5.55	1.46	1.52
34	BA	1246	G	C2-N2	-5.55	1.28	1.34
35	BB	838	G	C4'-C3'	-5.55	1.47	1.52
35	BB	1369	A	N9-C8	-5.55	1.33	1.37
37	BD	51	G	C5'-C4'	-5.55	1.44	1.51
38	BE	128	G	C2'-C1'	-5.55	1.47	1.53
85	AA	39	A	C1'-N9	-5.55	1.39	1.46
85	AA	282	C	O4'-C1'	-5.55	1.34	1.41
85	AA	402	G	C3'-C2'	-5.55	1.46	1.52
85	AA	533	C	C4-N4	-5.55	1.28	1.33
85	AA	854	A	O3'-P	-5.55	1.54	1.61
85	AA	1466	U	C2'-C1'	-5.55	1.47	1.53
13	AE	24	TYR	CB-CG	-5.55	1.43	1.51
34	BA	566	G	C2-N2	-5.55	1.28	1.34
34	BA	690	G	N9-C8	-5.55	1.33	1.37
34	BA	1475	G	N1-C2	-5.55	1.33	1.37
35	BB	1219	A	C1'-N9	-5.55	1.39	1.46
35	BB	1244	U	C1'-N1	-5.55	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	22	A	C3'-C2'	-5.55	1.46	1.52
34	BA	1070	G	C4'-C3'	-5.55	1.47	1.52
34	BA	1218	G	C1'-N9	-5.55	1.39	1.46
34	BA	1243	A	N9-C8	-5.55	1.33	1.37
34	BA	1732	A	C1'-N9	-5.55	1.39	1.46
35	BB	100	A	C1'-N9	-5.55	1.39	1.46
35	BB	476	A	C2'-C1'	-5.55	1.47	1.53
35	BB	481	A	N7-C5	-5.55	1.35	1.39
35	BB	1365	G	C6-N1	-5.55	1.35	1.39
35	BB	1470	G	C8-N7	-5.55	1.27	1.30
37	BD	26	C	O3'-P	-5.55	1.54	1.61
37	BD	82	G	C1'-N9	-5.55	1.39	1.46
38	BE	171	U	C2-N3	-5.55	1.33	1.37
85	AA	307	G	C4'-C3'	5.55	1.59	1.53
85	AA	396	U	O4'-C1'	-5.55	1.34	1.41
85	AA	460	U	P-O5'	-5.55	1.54	1.59
85	AA	501	A	N7-C5	-5.55	1.35	1.39
85	AA	915	G	C2'-C1'	-5.55	1.47	1.53
85	AA	999	A	N7-C5	5.55	1.42	1.39
85	AA	1368	G	C2-N2	-5.55	1.28	1.34
85	AA	1519	A	C6-N6	-5.55	1.29	1.33
85	AA	1688	U	C2-N3	-5.55	1.33	1.37
34	BA	992	A	C6-N6	-5.55	1.29	1.33
34	BA	1066	A	C5-C4	-5.55	1.34	1.38
34	BA	1591	G	C4'-C3'	-5.55	1.47	1.52
35	BB	524	C	C2'-C1'	-5.55	1.47	1.53
35	BB	642	G	C2'-C1'	-5.55	1.47	1.53
35	BB	1055	G	N1-C2	-5.55	1.33	1.37
35	BB	1062	G	N3-C4	-5.55	1.31	1.35
35	BB	1485	G	C3'-C2'	-5.55	1.46	1.52
36	BC	23	G	N1-C2	-5.55	1.33	1.37
38	BE	107	U	N1-C2	-5.55	1.33	1.38
40	BG	17	A	O4'-C1'	-5.55	1.34	1.41
44	BK	46	PHE	C-N	-5.55	1.23	1.34
85	AA	616	A	O3'-P	-5.55	1.54	1.61
85	AA	633	C	P-O5'	-5.55	1.54	1.59
85	AA	662	U	C1'-N1	-5.55	1.39	1.46
85	AA	695	A	C1'-N9	-5.55	1.39	1.46
85	AA	788	G	C2'-C1'	-5.55	1.47	1.53
85	AA	1283	C	O4'-C1'	-5.55	1.34	1.41
85	AA	1977	G	C2'-C1'	-5.55	1.47	1.53
85	AA	2028	G	C2'-C1'	-5.55	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	440	A	C8-N7	-5.55	1.27	1.31
34	BA	451	A	C3'-C2'	-5.55	1.46	1.52
34	BA	949	C	C2'-C1'	-5.55	1.47	1.53
34	BA	1462	U	P-O5'	-5.55	1.54	1.59
34	BA	1615	A	N3-C4	-5.55	1.31	1.34
36	BC	104	A	C2'-C1'	-5.55	1.47	1.53
85	AA	117	C	C4-C5	-5.55	1.38	1.43
85	AA	392	G	C6-N1	-5.55	1.35	1.39
85	AA	763	U	P-O5'	-5.55	1.54	1.59
85	AA	932	A	C8-N7	-5.55	1.27	1.31
85	AA	993	G	C2-N2	-5.55	1.29	1.34
85	AA	1387	C	P-O5'	-5.55	1.54	1.59
85	AA	1978	G	C8-N7	-5.55	1.27	1.30
34	BA	135	G	N7-C5	-5.55	1.35	1.39
34	BA	334	G	C5-C4	-5.55	1.34	1.38
34	BA	997	U	C2'-C1'	-5.55	1.47	1.53
34	BA	1046	G	C1'-N9	-5.55	1.39	1.46
34	BA	1280	A	N3-C4	-5.55	1.31	1.34
34	BA	1326	U	O3'-P	-5.55	1.54	1.61
34	BA	1327	G	C3'-C2'	-5.55	1.46	1.52
34	BA	1513	G	N1-C2	-5.55	1.33	1.37
34	BA	1553	G	N1-C2	-5.55	1.33	1.37
34	BA	1729	G	N9-C8	-5.55	1.33	1.37
35	BB	521	U	C3'-C2'	-5.55	1.46	1.52
35	BB	976	U	C1'-N1	-5.55	1.39	1.46
35	BB	1377	A	C4'-C3'	-5.55	1.47	1.52
37	BD	21	G	N1-C2	-5.55	1.33	1.37
40	BG	37	G	N1-C2	-5.55	1.33	1.37
85	AA	409	C	N3-C4	-5.55	1.30	1.33
85	AA	1163	G	O3'-P	-5.55	1.54	1.61
85	AA	1916	A	C8-N7	-5.55	1.27	1.31
34	BA	44	U	O4'-C1'	-5.54	1.34	1.41
34	BA	83	G	C2-N2	-5.54	1.29	1.34
34	BA	305	C	C3'-C2'	-5.54	1.46	1.52
34	BA	1025	A	O3'-P	-5.54	1.54	1.61
34	BA	1086	A	C6-N1	-5.54	1.31	1.35
34	BA	1710	C	C4-N4	-5.54	1.28	1.33
35	BB	666	A	C4'-C3'	-5.54	1.47	1.52
35	BB	1134	G	N3-C4	-5.54	1.31	1.35
35	BB	1517	G	N9-C8	-5.54	1.33	1.37
36	BC	59	A	O3'-P	-5.54	1.54	1.61
40	BG	8	U	N1-C2	5.54	1.43	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	19	A	C3'-C2'	-5.54	1.46	1.52
85	AA	195	C	N1-C6	-5.54	1.33	1.37
85	AA	534	A	O3'-P	-5.54	1.54	1.61
85	AA	773	G	N9-C4	-5.54	1.33	1.38
85	AA	1484	G	C3'-C2'	-5.54	1.46	1.52
85	AA	1630	U	P-O5'	-5.54	1.54	1.59
85	AA	1652	A	N7-C5	-5.54	1.35	1.39
34	BA	390	A	C8-N7	-5.54	1.27	1.31
34	BA	920	U	N3-C4	-5.54	1.33	1.38
34	BA	1334	G	C6-O6	-5.54	1.19	1.24
34	BA	1412	G	N1-C2	-5.54	1.33	1.37
34	BA	1530	G	C2-N2	-5.54	1.29	1.34
35	BB	54	U	C1'-N1	-5.54	1.39	1.46
35	BB	988	G	O3'-P	-5.54	1.54	1.61
35	BB	1395	G	C2-N2	-5.54	1.29	1.34
36	BC	95	A	N7-C5	-5.54	1.35	1.39
37	BD	28	C	C2-N3	-5.54	1.31	1.35
38	BE	110	U	C4'-C3'	-5.54	1.47	1.52
40	BG	133	C	C4'-C3'	-5.54	1.47	1.52
41	BH	96	G	C2-N3	5.54	1.37	1.32
85	AA	294	G	N9-C4	5.54	1.42	1.38
85	AA	428	G	N7-C5	-5.54	1.35	1.39
85	AA	484	G	N9-C8	-5.54	1.33	1.37
85	AA	511	A	N9-C4	-5.54	1.34	1.37
85	AA	709	A	C3'-C2'	-5.54	1.46	1.52
85	AA	939	A	C1'-N9	-5.54	1.39	1.46
85	AA	2018	U	O3'-P	-5.54	1.54	1.61
85	AA	2089	G	C4'-C3'	5.54	1.59	1.53
85	AA	2097	U	P-O5'	-5.54	1.54	1.59
34	BA	94	G	C3'-C2'	-5.54	1.46	1.52
34	BA	347	A	N7-C5	-5.54	1.35	1.39
34	BA	1082	U	O3'-P	-5.54	1.54	1.61
34	BA	1246	G	P-O5'	-5.54	1.54	1.59
34	BA	1531	G	C5-C4	-5.54	1.34	1.38
34	BA	1662	U	P-O5'	-5.54	1.54	1.59
35	BB	73	G	C4'-O4'	-5.54	1.38	1.45
35	BB	93	A	C3'-C2'	-5.54	1.46	1.52
35	BB	684	U	O3'-P	-5.54	1.54	1.61
35	BB	764	C	C3'-C2'	-5.54	1.46	1.52
35	BB	1474	A	C5'-C4'	5.54	1.57	1.51
40	BG	167	C	C2-N3	-5.54	1.31	1.35
85	AA	532	G	N1-C2	-5.54	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1556	G	P-O5'	-5.54	1.54	1.59
34	BA	596	G	C3'-O3'	5.54	1.50	1.42
34	BA	1011	G	N9-C8	-5.54	1.33	1.37
34	BA	1539	A	C3'-C2'	-5.54	1.46	1.52
34	BA	1789	A	P-O5'	-5.54	1.54	1.59
35	BB	266	C	O3'-P	-5.54	1.54	1.61
35	BB	388	C	P-O5'	-5.54	1.54	1.59
35	BB	438	G	O3'-P	-5.54	1.54	1.61
35	BB	1400	C	C4'-C3'	-5.54	1.47	1.52
40	BG	179	C	C4-N4	-5.54	1.28	1.33
41	BH	4	U	C1'-N1	-5.54	1.39	1.46
41	BH	105	U	C3'-C2'	-5.54	1.46	1.52
85	AA	683	U	N3-C4	-5.54	1.33	1.38
85	AA	1968	A	C2'-C1'	-5.54	1.47	1.53
85	AA	2121	G	C2'-C1'	-5.54	1.47	1.53
34	BA	396	U	O3'-P	-5.54	1.54	1.61
34	BA	398	G	P-O5'	-5.54	1.54	1.59
34	BA	596	G	C2-N2	-5.54	1.29	1.34
34	BA	1116	G	C6-N1	-5.54	1.35	1.39
34	BA	1286	C	N1-C2	-5.54	1.34	1.40
35	BB	78	C	C2-N3	-5.54	1.31	1.35
35	BB	831	C	C3'-C2'	-5.54	1.46	1.52
35	BB	899	C	C2'-C1'	-5.54	1.47	1.53
35	BB	1216	G	O3'-P	-5.54	1.54	1.61
35	BB	1375	G	N9-C8	-5.54	1.33	1.37
39	BF	60	C	P-O5'	-5.54	1.54	1.59
41	BH	107	A	C2'-C1'	-5.54	1.47	1.53
85	AA	503	A	O3'-P	-5.54	1.54	1.61
85	AA	671	G	C2-N2	-5.54	1.29	1.34
85	AA	986	U	O3'-P	-5.54	1.54	1.61
85	AA	1156	A	N9-C4	-5.54	1.34	1.37
85	AA	1214	C	O4'-C1'	-5.54	1.34	1.41
85	AA	1240	A	C5-C4	-5.54	1.34	1.38
85	AA	1790	G	N9-C4	-5.54	1.33	1.38
85	AA	2050	C	C2'-C1'	-5.54	1.47	1.53
34	BA	204	U	C2'-C1'	-5.54	1.47	1.53
34	BA	1222	C	C3'-C2'	-5.54	1.46	1.52
34	BA	1299	G	N1-C2	-5.54	1.33	1.37
34	BA	1425	G	N9-C4	-5.54	1.33	1.38
34	BA	1687	A	O4'-C1'	-5.54	1.34	1.41
35	BB	404	A	O3'-P	-5.54	1.54	1.61
35	BB	638	G	N9-C8	-5.54	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	676	G	C2-N2	-5.54	1.29	1.34
40	BG	40	G	N1-C2	-5.54	1.33	1.37
49	BP	127	ILE	CA-CB	-5.54	1.42	1.54
85	AA	1140	G	C6-N1	-5.54	1.35	1.39
85	AA	1241	A	N7-C5	-5.54	1.35	1.39
85	AA	1298	G	O3'-P	-5.54	1.54	1.61
85	AA	2089	G	N9-C4	-5.54	1.33	1.38
34	BA	58	A	N9-C8	-5.54	1.33	1.37
34	BA	109	A	C5-C4	-5.54	1.34	1.38
34	BA	405	C	C1'-N1	-5.54	1.39	1.46
34	BA	922	C	C3'-O3'	-5.54	1.34	1.42
34	BA	1052	G	C8-N7	-5.54	1.27	1.30
34	BA	1237	U	C2'-C1'	-5.54	1.47	1.53
34	BA	1248	A	O3'-P	-5.54	1.54	1.61
34	BA	1526	C	O3'-P	-5.54	1.54	1.61
34	BA	1683	C	C2'-C1'	-5.54	1.47	1.53
34	BA	1796	A	N7-C5	-5.54	1.35	1.39
35	BB	544	C	N3-C4	-5.54	1.30	1.33
35	BB	567	G	C2'-C1'	-5.54	1.47	1.53
35	BB	701	U	O3'-P	-5.54	1.54	1.61
35	BB	1136	G	N3-C4	-5.54	1.31	1.35
35	BB	1147	G	N9-C8	-5.54	1.33	1.37
35	BB	1490	G	C6-N1	-5.54	1.35	1.39
40	BG	121	C	C3'-C2'	-5.54	1.46	1.52
40	BG	135	C	O3'-P	-5.54	1.54	1.61
85	AA	923	A	C2'-C1'	-5.54	1.47	1.53
85	AA	962	U	C1'-N1	-5.54	1.39	1.46
85	AA	1161	U	C2-N3	-5.54	1.33	1.37
85	AA	2120	C	N1-C2	-5.54	1.34	1.40
34	BA	491	U	C2'-C1'	-5.53	1.47	1.53
34	BA	1304	C	O3'-P	-5.53	1.54	1.61
34	BA	1399	A	C4'-C3'	-5.53	1.47	1.52
34	BA	1408	C	P-O5'	-5.53	1.54	1.59
35	BB	44	C	N1-C6	-5.53	1.33	1.37
35	BB	1375	G	N3-C4	-5.53	1.31	1.35
38	BE	61	A	C3'-C2'	-5.53	1.46	1.52
38	BE	210	G	C2'-C1'	-5.53	1.47	1.53
41	BH	33	G	C3'-O3'	-5.53	1.34	1.42
85	AA	483	G	N1-C2	-5.53	1.33	1.37
85	AA	575	G	C2'-C1'	-5.53	1.47	1.53
85	AA	681	G	C8-N7	-5.53	1.27	1.30
85	AA	920	A	C4'-C3'	5.53	1.59	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1128	G	C2-N2	-5.53	1.29	1.34
85	AA	1688	U	N1-C2	-5.53	1.33	1.38
85	AA	2241	C	C5'-C4'	-5.53	1.44	1.51
34	BA	13	U	C4'-C3'	-5.53	1.47	1.52
34	BA	977	G	C6-N1	-5.53	1.35	1.39
34	BA	1496	G	N3-C4	-5.53	1.31	1.35
35	BB	1014	U	O3'-P	-5.53	1.54	1.61
35	BB	1510	G	C5-C4	-5.53	1.34	1.38
35	BB	1539	C	C3'-C2'	-5.53	1.46	1.52
41	BH	47	G	C2'-C1'	-5.53	1.47	1.53
85	AA	585	G	N3-C4	-5.53	1.31	1.35
34	BA	205	G	N9-C4	-5.53	1.33	1.38
34	BA	445	C	C2-N3	-5.53	1.31	1.35
34	BA	904	G	N7-C5	-5.53	1.35	1.39
34	BA	955	G	C6-N1	-5.53	1.35	1.39
34	BA	1656	A	C1'-N9	-5.53	1.39	1.46
34	BA	1674	G	C1'-N9	-5.53	1.39	1.46
35	BB	48	G	C2-N2	-5.53	1.29	1.34
35	BB	368	C	C4'-C3'	-5.53	1.47	1.52
35	BB	1295	A	C4'-C3'	-5.53	1.47	1.52
35	BB	1384	A	C4'-C3'	-5.53	1.47	1.52
35	BB	1389	C	C3'-C2'	-5.53	1.46	1.52
36	BC	60	U	C4'-C3'	-5.53	1.47	1.52
37	BD	98	G	N7-C5	-5.53	1.35	1.39
38	BE	27	A	C5-C4	-5.53	1.34	1.38
85	AA	536	C	P-O5'	-5.53	1.54	1.59
85	AA	879	G	N9-C8	-5.53	1.33	1.37
85	AA	1484	G	N3-C4	-5.53	1.31	1.35
85	AA	1949	U	O3'-P	-5.53	1.54	1.61
85	AA	2069	A	C3'-C2'	-5.53	1.46	1.52
34	BA	1018	U	P-O5'	-5.53	1.54	1.59
34	BA	1324	G	C4'-C3'	-5.53	1.47	1.52
35	BB	1294	C	C4'-O4'	-5.53	1.38	1.45
35	BB	1369	A	C1'-N9	-5.53	1.39	1.46
35	BB	1464	G	N7-C5	-5.53	1.35	1.39
85	AA	143	U	C2'-C1'	-5.53	1.47	1.53
85	AA	2128	G	C6-N1	-5.53	1.35	1.39
34	BA	94	G	O3'-P	-5.53	1.54	1.61
34	BA	104	A	C2'-C1'	-5.53	1.47	1.53
34	BA	185	A	N7-C5	-5.53	1.35	1.39
34	BA	199	U	P-O5'	-5.53	1.54	1.59
34	BA	389	U	O3'-P	-5.53	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	427	G	N9-C8	-5.53	1.33	1.37
34	BA	476	U	C3'-C2'	-5.53	1.46	1.52
34	BA	620	C	O3'-P	-5.53	1.54	1.61
34	BA	733	G	C3'-C2'	-5.53	1.46	1.52
34	BA	965	A	N3-C4	-5.53	1.31	1.34
34	BA	1139	G	P-O5'	-5.53	1.54	1.59
35	BB	48	G	O3'-P	-5.53	1.54	1.61
35	BB	437	U	C3'-C2'	-5.53	1.46	1.52
35	BB	438	G	C2-N2	-5.53	1.29	1.34
35	BB	1013	U	P-O5'	-5.53	1.54	1.59
35	BB	1341	U	C4'-C3'	-5.53	1.47	1.52
36	BC	15	G	C5-C4	-5.53	1.34	1.38
37	BD	52	U	O3'-P	-5.53	1.54	1.61
40	BG	105	A	C6-N6	-5.53	1.29	1.33
68	Bi	27	PRO	CA-C	-5.53	1.41	1.52
85	AA	20	G	N9-C8	-5.53	1.33	1.37
85	AA	110	U	C2'-C1'	-5.53	1.47	1.53
85	AA	455	G	C3'-C2'	-5.53	1.46	1.52
85	AA	867	G	N1-C2	-5.53	1.33	1.37
85	AA	1291	A	O3'-P	-5.53	1.54	1.61
85	AA	1515	A	C5'-C4'	-5.53	1.44	1.51
34	BA	288	U	C1'-N1	-5.53	1.39	1.46
34	BA	498	A	O4'-C1'	-5.53	1.34	1.41
34	BA	905	A	C1'-N9	-5.53	1.39	1.46
34	BA	1046	G	C8-N7	-5.53	1.27	1.30
34	BA	1046	G	N9-C4	-5.53	1.33	1.38
34	BA	1156	U	O4'-C1'	-5.53	1.34	1.41
34	BA	1468	U	C3'-C2'	-5.53	1.46	1.52
34	BA	1519	G	N9-C8	-5.53	1.33	1.37
35	BB	590	G	P-O5'	-5.53	1.54	1.59
35	BB	634	A	C8-N7	-5.53	1.27	1.31
35	BB	1431	G	C2-N2	-5.53	1.29	1.34
36	BC	104	A	N7-C5	-5.53	1.35	1.39
37	BD	3	G	C2-N2	-5.53	1.29	1.34
85	AA	393	C	C2-N3	-5.53	1.31	1.35
85	AA	689	U	C2-N3	-5.53	1.33	1.37
85	AA	1012	C	O3'-P	-5.53	1.54	1.61
85	AA	1546	G	C6-N1	-5.53	1.35	1.39
85	AA	1942	U	P-O5'	-5.53	1.54	1.59
85	AA	2201	A	P-O5'	-5.53	1.54	1.59
34	BA	713	C	N1-C6	-5.52	1.33	1.37
34	BA	1308	C	P-O5'	-5.52	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	152	C	C2-N3	-5.52	1.31	1.35
37	BD	100	A	C1'-N9	-5.52	1.39	1.46
44	BK	85	PHE	N-CA	-5.52	1.35	1.46
85	AA	252	G	N7-C5	-5.52	1.35	1.39
85	AA	308	U	C4'-C3'	-5.52	1.47	1.52
85	AA	1666	U	C3'-C2'	-5.52	1.46	1.52
15	AG	136	PRO	C-N	-5.52	1.23	1.34
34	BA	195	G	N1-C2	-5.52	1.33	1.37
34	BA	668	G	C6-N1	-5.52	1.35	1.39
34	BA	733	G	C1'-N9	-5.52	1.39	1.46
34	BA	1097	G	C1'-N9	-5.52	1.39	1.46
34	BA	1130	U	C2'-C1'	-5.52	1.47	1.53
34	BA	1179	U	C4'-C3'	-5.52	1.47	1.52
34	BA	1738	G	C3'-C2'	-5.52	1.46	1.52
35	BB	381	C	C2-N3	-5.52	1.31	1.35
36	BC	25	C	C4'-C3'	-5.52	1.47	1.52
36	BC	28	C	P-O5'	-5.52	1.54	1.59
36	BC	33	U	C2'-C1'	-5.52	1.47	1.53
36	BC	125	A	N9-C8	-5.52	1.33	1.37
38	BE	6	A	C5'-C4'	-5.52	1.44	1.51
38	BE	114	G	O4'-C1'	-5.52	1.34	1.41
40	BG	90	G	C2-N2	-5.52	1.29	1.34
85	AA	68	A	N9-C4	-5.52	1.34	1.37
85	AA	463	G	O3'-P	-5.52	1.54	1.61
85	AA	500	C	C4'-O4'	-5.52	1.38	1.45
85	AA	1163	G	N7-C5	-5.52	1.35	1.39
85	AA	1200	A	N3-C4	-5.52	1.31	1.34
85	AA	1497	U	C1'-N1	-5.52	1.39	1.46
85	AA	1531	G	C1'-N9	-5.52	1.39	1.46
85	AA	1923	A	C4'-C3'	-5.52	1.47	1.52
2	A1	96	PHE	CB-CG	-5.52	1.42	1.51
34	BA	962	U	N3-C4	-5.52	1.33	1.38
35	BB	362	A	N9-C4	-5.52	1.34	1.37
35	BB	365	U	N3-C4	-5.52	1.33	1.38
35	BB	1227	G	C4'-O4'	-5.52	1.38	1.45
35	BB	1508	G	N7-C5	-5.52	1.35	1.39
38	BE	53	U	P-O5'	-5.52	1.54	1.59
34	BA	321	G	C2'-C1'	-5.52	1.47	1.53
34	BA	330	A	N9-C8	-5.52	1.33	1.37
34	BA	400	A	C5-C4	-5.52	1.34	1.38
34	BA	408	U	C2-N3	-5.52	1.33	1.37
35	BB	426	A	C1'-N9	-5.52	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	121	G	C6-N1	-5.52	1.35	1.39
36	BC	164	G	C4'-C3'	-5.52	1.47	1.52
38	BE	49	A	C5-C6	-5.52	1.36	1.41
40	BG	171	A	C6-N6	-5.52	1.29	1.33
85	AA	342	C	C4'-C3'	-5.52	1.47	1.52
85	AA	413	G	N9-C4	-5.52	1.33	1.38
85	AA	1476	C	C3'-C2'	-5.52	1.46	1.52
85	AA	1505	G	C6-N1	-5.52	1.35	1.39
85	AA	1622	G	N7-C5	-5.52	1.35	1.39
85	AA	1843	A	C1'-N9	-5.52	1.39	1.46
85	AA	2169	C	C3'-C2'	-5.52	1.46	1.52
34	BA	36	A	O4'-C1'	-5.52	1.34	1.41
34	BA	1098	G	C5-C4	-5.52	1.34	1.38
34	BA	1816	G	N1-C2	-5.52	1.33	1.37
34	BA	1833	G	O3'-P	-5.52	1.54	1.61
35	BB	1054	G	C2'-C1'	-5.52	1.47	1.53
35	BB	1431	G	O4'-C1'	-5.52	1.34	1.41
36	BC	110	A	C1'-N9	-5.52	1.39	1.46
36	BC	117	A	N3-C4	-5.52	1.31	1.34
37	BD	26	C	C2'-C1'	-5.52	1.47	1.53
37	BD	55	A	N9-C4	-5.52	1.34	1.37
37	BD	98	G	C5-C6	-5.52	1.36	1.42
38	BE	8	G	C5-C4	-5.52	1.34	1.38
74	Bo	22	PRO	CA-C	-5.52	1.41	1.52
85	AA	598	C	C2'-C1'	-5.52	1.47	1.53
85	AA	624	A	N3-C4	-5.52	1.31	1.34
85	AA	979	U	O3'-P	-5.52	1.54	1.61
85	AA	1597	C	P-O5'	-5.52	1.54	1.59
85	AA	2165	C	C2'-C1'	-5.52	1.47	1.53
34	BA	415	C	O3'-P	-5.52	1.54	1.61
34	BA	714	G	C5-C6	-5.52	1.36	1.42
35	BB	40	C	C3'-C2'	-5.52	1.46	1.52
35	BB	84	G	N1-C2	-5.52	1.33	1.37
35	BB	411	A	C2'-C1'	-5.52	1.47	1.53
35	BB	1195	A	N3-C4	-5.52	1.31	1.34
35	BB	1360	A	C2'-C1'	-5.52	1.47	1.53
36	BC	93	C	C2-N3	-5.52	1.31	1.35
37	BD	92	G	N7-C5	-5.52	1.35	1.39
40	BG	160	C	C2-N3	-5.52	1.31	1.35
41	BH	54	U	O3'-P	-5.52	1.54	1.61
69	Bj	80	GLY	CA-C	-5.52	1.43	1.51
71	Bl	119	SER	CA-CB	5.52	1.61	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	353	G	C5-C4	-5.52	1.34	1.38
85	AA	377	U	C2-N3	-5.52	1.33	1.37
85	AA	481	A	O3'-P	-5.52	1.54	1.61
85	AA	565	G	P-O5'	-5.52	1.54	1.59
85	AA	608	A	O4'-C1'	-5.52	1.34	1.41
85	AA	668	A	O4'-C1'	-5.52	1.34	1.41
34	BA	246	G	N1-C2	-5.51	1.33	1.37
34	BA	1466	U	C1'-N1	-5.51	1.39	1.46
34	BA	1637	G	C2-N2	-5.51	1.29	1.34
34	BA	1691	G	N1-C2	-5.51	1.33	1.37
35	BB	108	G	C1'-N9	-5.51	1.39	1.46
35	BB	619	A	C4'-C3'	-5.51	1.47	1.52
35	BB	787	A	O4'-C1'	-5.51	1.34	1.41
35	BB	1422	G	P-O5'	-5.51	1.54	1.59
36	BC	110	A	N9-C4	-5.51	1.34	1.37
37	BD	79	G	N9-C8	-5.51	1.33	1.37
40	BG	29	U	O3'-P	-5.51	1.54	1.61
40	BG	111	C	C4-N4	-5.51	1.28	1.33
85	AA	55	A	O3'-P	-5.51	1.54	1.61
85	AA	64	A	C3'-O3'	5.51	1.49	1.42
85	AA	106	G	N7-C5	-5.51	1.35	1.39
85	AA	125	A	C4'-C3'	-5.51	1.47	1.52
85	AA	941	C	C2-N3	-5.51	1.31	1.35
34	BA	22	C	O3'-P	-5.51	1.54	1.61
34	BA	144	C	C2-N3	-5.51	1.31	1.35
34	BA	1106	A	C4'-O4'	-5.51	1.38	1.45
35	BB	550	G	O4'-C1'	-5.51	1.34	1.41
35	BB	644	A	O4'-C1'	-5.51	1.34	1.41
36	BC	11	G	N9-C8	-5.51	1.33	1.37
37	BD	60	C	C4'-C3'	-5.51	1.47	1.52
85	AA	463	G	C8-N7	-5.51	1.27	1.30
85	AA	528	U	O4'-C1'	-5.51	1.34	1.41
85	AA	961	U	C2'-C1'	-5.51	1.47	1.53
85	AA	2120	C	O3'-P	-5.51	1.54	1.61
34	BA	432	A	N7-C5	-5.51	1.35	1.39
34	BA	844	U	C2'-C1'	-5.51	1.47	1.53
34	BA	963	G	N7-C5	-5.51	1.35	1.39
34	BA	1598	U	C4'-O4'	-5.51	1.38	1.45
35	BB	1521	G	C2-N2	-5.51	1.29	1.34
38	BE	9	C	N3-C4	-5.51	1.30	1.33
40	BG	49	A	P-O5'	-5.51	1.54	1.59
40	BG	101	G	N1-C2	-5.51	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	17	C	C4'-C3'	-5.51	1.47	1.52
85	AA	526	G	C6-N1	-5.51	1.35	1.39
85	AA	959	C	C2-N3	-5.51	1.31	1.35
85	AA	1580	A	P-O5'	-5.51	1.54	1.59
85	AA	1691	U	C2'-C1'	-5.51	1.47	1.53
34	BA	25	C	C2'-C1'	-5.51	1.47	1.53
34	BA	475	A	O3'-P	-5.51	1.54	1.61
34	BA	681	G	C6-N1	-5.51	1.35	1.39
34	BA	796	G	C3'-C2'	-5.51	1.46	1.52
34	BA	1534	U	P-O5'	-5.51	1.54	1.59
34	BA	1680	G	P-O5'	-5.51	1.54	1.59
35	BB	607	G	O3'-P	-5.51	1.54	1.61
35	BB	1017	U	C3'-C2'	-5.51	1.46	1.52
35	BB	1071	G	C2-N2	-5.51	1.29	1.34
35	BB	1320	U	C4'-O4'	-5.51	1.38	1.45
41	BH	14	C	C3'-C2'	-5.51	1.46	1.52
85	AA	397	G	C5'-C4'	-5.51	1.44	1.51
85	AA	1122	U	C2-N3	-5.51	1.33	1.37
85	AA	1294	U	C1'-N1	-5.51	1.39	1.46
85	AA	1460	G	C6-N1	-5.51	1.35	1.39
85	AA	1769	A	P-O5'	-5.51	1.54	1.59
85	AA	2210	C	O3'-P	-5.51	1.54	1.61
34	BA	202	A	C5'-C4'	-5.51	1.44	1.51
34	BA	633	G	P-O5'	-5.51	1.54	1.59
34	BA	707	C	C2-N3	-5.51	1.31	1.35
34	BA	1049	G	C5-C6	-5.51	1.36	1.42
34	BA	1319	A	O3'-P	-5.51	1.54	1.61
34	BA	1341	A	N3-C4	-5.51	1.31	1.34
34	BA	1632	G	O3'-P	-5.51	1.54	1.61
85	AA	799	G	N9-C4	-5.51	1.33	1.38
85	AA	1187	G	C1'-N9	-5.51	1.39	1.46
85	AA	1295	G	C2-N2	-5.51	1.29	1.34
85	AA	1844	A	C5-C4	-5.51	1.34	1.38
2	A1	190	GLY	N-CA	-5.51	1.37	1.46
34	BA	463	A	O4'-C1'	-5.51	1.34	1.41
34	BA	726	G	C5-C4	-5.51	1.34	1.38
34	BA	792	A	N3-C4	-5.51	1.31	1.34
34	BA	805	A	N9-C4	-5.51	1.34	1.37
34	BA	1215	U	C2'-C1'	-5.51	1.47	1.53
34	BA	1311	G	N9-C4	-5.51	1.33	1.38
35	BB	489	A	C2'-C1'	-5.51	1.47	1.53
35	BB	826	G	C1'-N9	-5.51	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	53	C	C2-N3	-5.51	1.31	1.35
40	BG	58	G	N9-C8	-5.51	1.33	1.37
85	AA	361	U	C5'-C4'	5.51	1.57	1.51
85	AA	645	C	C1'-N1	-5.51	1.39	1.46
85	AA	931	G	C3'-C2'	-5.51	1.46	1.52
85	AA	1517	G	C5-C4	-5.51	1.34	1.38
85	AA	1548	A	C3'-C2'	-5.51	1.46	1.52
85	AA	2115	G	N9-C4	-5.51	1.33	1.38
34	BA	543	A	N3-C4	-5.50	1.31	1.34
34	BA	809	U	N1-C2	-5.50	1.33	1.38
35	BB	370	A	C3'-C2'	-5.50	1.46	1.52
35	BB	1174	C	C1'-N1	-5.50	1.39	1.46
35	BB	1346	A	C5-C4	-5.50	1.34	1.38
38	BE	188	C	N1-C6	-5.50	1.33	1.37
85	AA	1622	G	N1-C2	-5.50	1.33	1.37
85	AA	2008	G	N1-C2	-5.50	1.33	1.37
34	BA	263	G	C1'-N9	-5.50	1.39	1.46
34	BA	475	A	O4'-C1'	-5.50	1.34	1.41
34	BA	611	A	C6-N1	-5.50	1.31	1.35
34	BA	670	U	C1'-N1	-5.50	1.39	1.46
34	BA	1246	G	C1'-N9	-5.50	1.39	1.46
35	BB	91	G	N9-C4	-5.50	1.33	1.38
35	BB	116	G	N1-C2	-5.50	1.33	1.37
35	BB	357	C	O3'-P	-5.50	1.54	1.61
35	BB	486	G	N9-C4	-5.50	1.33	1.38
35	BB	651	G	N1-C2	-5.50	1.33	1.37
35	BB	905	C	P-O5'	-5.50	1.54	1.59
35	BB	1322	A	N3-C4	-5.50	1.31	1.34
35	BB	1399	A	N3-C4	-5.50	1.31	1.34
35	BB	1530	U	O4'-C1'	-5.50	1.34	1.41
85	AA	351	C	O3'-P	-5.50	1.54	1.61
85	AA	816	A	C4'-O4'	5.50	1.52	1.45
85	AA	996	A	C1'-N9	-5.50	1.39	1.46
85	AA	1585	A	C3'-C2'	-5.50	1.46	1.52
85	AA	1988	A	N3-C4	-5.50	1.31	1.34
34	BA	324	C	C1'-N1	-5.50	1.39	1.46
34	BA	473	A	C3'-C2'	-5.50	1.46	1.52
34	BA	498	A	C6-N6	-5.50	1.29	1.33
34	BA	1193	A	N9-C8	-5.50	1.33	1.37
34	BA	1324	G	C5-C6	-5.50	1.36	1.42
35	BB	33	A	C5-C6	-5.50	1.36	1.41
35	BB	1334	C	C1'-N1	-5.50	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	166	C	N3-C4	-5.50	1.30	1.33
41	BH	121	A	C8-N7	-5.50	1.27	1.31
85	AA	1105	G	N7-C5	-5.50	1.35	1.39
85	AA	1422	A	O3'-P	-5.50	1.54	1.61
85	AA	1812	C	C3'-C2'	-5.50	1.46	1.52
85	AA	2089	G	C5'-C4'	5.50	1.57	1.51
34	BA	350	C	C2'-C1'	-5.50	1.47	1.53
34	BA	739	A	P-O5'	-5.50	1.54	1.59
34	BA	1623	U	O3'-P	-5.50	1.54	1.61
35	BB	1476	C	C1'-N1	-5.50	1.39	1.46
85	AA	373	G	N1-C2	-5.50	1.33	1.37
85	AA	453	G	C6-N1	-5.50	1.35	1.39
85	AA	454	G	N9-C4	-5.50	1.33	1.38
85	AA	687	G	C6-O6	-5.50	1.19	1.24
34	BA	148	G	P-O5'	-5.50	1.54	1.59
34	BA	257	G	N7-C5	-5.50	1.35	1.39
34	BA	337	C	C4-N4	-5.50	1.29	1.33
34	BA	522	C	N1-C6	-5.50	1.33	1.37
34	BA	921	G	N1-C2	-5.50	1.33	1.37
34	BA	1565	U	P-O5'	-5.50	1.54	1.59
34	BA	1594	G	C1'-N9	-5.50	1.39	1.46
34	BA	1610	A	C4'-C3'	-5.50	1.47	1.52
34	BA	1669	C	O3'-P	-5.50	1.54	1.61
35	BB	550	G	C1'-N9	-5.50	1.39	1.46
36	BC	126	G	O3'-P	-5.50	1.54	1.61
68	Bi	58	TYR	CB-CG	-5.50	1.43	1.51
75	Bp	27	HIS	CB-CG	-5.50	1.40	1.50
85	AA	308	U	C1'-N1	-5.50	1.39	1.46
85	AA	630	A	C1'-N9	-5.50	1.39	1.46
85	AA	684	G	O3'-P	-5.50	1.54	1.61
85	AA	1695	G	C2-N2	-5.50	1.29	1.34
34	BA	426	A	N3-C4	-5.50	1.31	1.34
34	BA	454	G	N1-C2	-5.50	1.33	1.37
34	BA	454	G	N7-C5	-5.50	1.35	1.39
34	BA	694	G	O3'-P	-5.50	1.54	1.61
34	BA	994	G	N3-C4	-5.50	1.31	1.35
34	BA	1015	G	N7-C5	-5.50	1.35	1.39
34	BA	1052	G	N7-C5	-5.50	1.35	1.39
34	BA	1117	G	C2-N2	-5.50	1.29	1.34
34	BA	1419	A	N7-C5	-5.50	1.35	1.39
34	BA	1538	G	P-O5'	-5.50	1.54	1.59
35	BB	1108	G	N9-C4	5.50	1.42	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1147	G	N3-C4	-5.50	1.31	1.35
35	BB	1512	C	C4-C5	-5.50	1.38	1.43
36	BC	91	G	C2-N2	-5.50	1.29	1.34
37	BD	3	G	C2'-C1'	-5.50	1.47	1.53
42	BI	29	LEU	CA-C	-5.50	1.38	1.52
80	Bu	194	ASP	N-CA	-5.50	1.35	1.46
85	AA	475	A	N3-C4	-5.50	1.31	1.34
85	AA	487	G	C1'-N9	-5.50	1.39	1.46
85	AA	602	U	O3'-P	-5.50	1.54	1.61
34	BA	777	C	C2-N3	-5.50	1.31	1.35
34	BA	1094	U	N1-C6	-5.50	1.33	1.38
34	BA	1270	G	C5-C4	-5.50	1.34	1.38
34	BA	1338	G	C1'-N9	-5.50	1.39	1.46
35	BB	1238	A	C1'-N9	-5.50	1.39	1.46
35	BB	1435	G	N1-C2	-5.50	1.33	1.37
37	BD	107	G	P-O5'	-5.50	1.54	1.59
41	BH	4	U	C4'-O4'	-5.50	1.38	1.45
85	AA	245	A	P-O5'	-5.50	1.54	1.59
86	AB	70	G	C6-N1	-5.50	1.35	1.39
34	BA	4	A	C8-N7	-5.49	1.27	1.31
34	BA	819	G	C2-N2	-5.49	1.29	1.34
34	BA	1516	G	C5-C4	-5.49	1.34	1.38
35	BB	126	C	O3'-P	-5.49	1.54	1.61
35	BB	555	G	C6-N1	-5.49	1.35	1.39
35	BB	745	C	C4'-C3'	-5.49	1.47	1.52
35	BB	960	C	C2'-C1'	-5.49	1.47	1.53
35	BB	1103	A	C2'-C1'	-5.49	1.47	1.53
35	BB	1189	C	C4'-C3'	-5.49	1.47	1.52
36	BC	74	U	N1-C2	-5.49	1.33	1.38
85	AA	577	U	N3-C4	-5.49	1.33	1.38
85	AA	650	G	C3'-C2'	-5.49	1.46	1.52
85	AA	1132	A	C2'-C1'	-5.49	1.47	1.53
85	AA	1170	C	C4'-C3'	-5.49	1.47	1.52
85	AA	2048	C	P-O5'	-5.49	1.54	1.59
34	BA	410	G	C6-N1	-5.49	1.35	1.39
35	BB	566	A	N3-C4	-5.49	1.31	1.34
35	BB	1391	G	C2'-C1'	-5.49	1.47	1.53
35	BB	1426	G	C1'-N9	-5.49	1.39	1.46
39	BF	38	C	N1-C6	-5.49	1.33	1.37
85	AA	443	A	C4'-O4'	-5.49	1.38	1.45
85	AA	1097	G	C2'-C1'	-5.49	1.47	1.53
85	AA	1126	G	C2'-C1'	-5.49	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2193	A	C1'-N9	-5.49	1.39	1.46
35	BB	633	C	C4-N4	-5.49	1.29	1.33
41	BH	129	G	C6-N1	-5.49	1.35	1.39
74	Bo	56	ARG	CD-NE	5.49	1.55	1.46
85	AA	583	U	C2'-C1'	-5.49	1.47	1.53
85	AA	632	U	O3'-P	-5.49	1.54	1.61
85	AA	650	G	O3'-P	-5.49	1.54	1.61
85	AA	1528	A	O3'-P	-5.49	1.54	1.61
85	AA	1619	A	N7-C5	-5.49	1.35	1.39
85	AA	1659	C	C2'-C1'	-5.49	1.47	1.53
85	AA	2105	G	N3-C4	-5.49	1.31	1.35
34	BA	51	C	P-O5'	-5.49	1.54	1.59
34	BA	219	U	C4'-C3'	-5.49	1.47	1.52
34	BA	371	U	C4'-C3'	-5.49	1.47	1.52
34	BA	504	A	C4'-C3'	-5.49	1.47	1.52
34	BA	730	C	C2-N3	-5.49	1.31	1.35
34	BA	1428	G	N9-C4	-5.49	1.33	1.38
35	BB	72	G	N7-C5	-5.49	1.35	1.39
35	BB	118	A	C8-N7	-5.49	1.27	1.31
35	BB	491	A	C5-C6	-5.49	1.36	1.41
35	BB	852	G	N9-C4	5.49	1.42	1.38
35	BB	1278	A	C2'-C1'	-5.49	1.47	1.53
36	BC	131	C	P-O5'	-5.49	1.54	1.59
39	BF	1	C	O3'-P	-5.49	1.54	1.61
41	BH	17	A	C1'-N9	-5.49	1.39	1.46
85	AA	33	U	O3'-P	-5.49	1.54	1.61
85	AA	291	G	C1'-N9	-5.49	1.39	1.46
85	AA	679	A	N7-C5	-5.49	1.35	1.39
85	AA	1539	A	N3-C4	-5.49	1.31	1.34
85	AA	2174	G	C5'-C4'	-5.49	1.44	1.51
85	AA	2199	G	N1-C2	-5.49	1.33	1.37
34	BA	1474	G	C2-N2	-5.49	1.29	1.34
35	BB	494	C	O3'-P	-5.49	1.54	1.61
35	BB	1212	C	N3-C4	-5.49	1.30	1.33
36	BC	95	A	C4'-O4'	-5.49	1.38	1.45
85	AA	1084	A	P-O5'	-5.49	1.54	1.59
85	AA	1175	A	P-O5'	-5.49	1.54	1.59
34	BA	59	A	C2'-C1'	-5.49	1.47	1.53
34	BA	811	C	C2-N3	-5.49	1.31	1.35
34	BA	821	G	C5-C4	-5.49	1.34	1.38
34	BA	1502	G	N1-C2	-5.49	1.33	1.37
34	BA	1659	G	C3'-C2'	-5.49	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	47	C	C3'-C2'	-5.49	1.46	1.52
35	BB	647	U	C2'-C1'	-5.49	1.47	1.53
35	BB	667	G	N3-C4	-5.49	1.31	1.35
35	BB	824	C	C3'-C2'	-5.49	1.46	1.52
35	BB	975	G	C3'-C2'	-5.49	1.46	1.52
38	BE	162	U	C2'-C1'	-5.49	1.47	1.53
40	BG	161	C	P-O5'	-5.49	1.54	1.59
41	BH	66	G	O3'-P	-5.49	1.54	1.61
41	BH	107	A	P-O5'	-5.49	1.54	1.59
85	AA	607	U	P-O5'	-5.49	1.54	1.59
85	AA	681	G	N3-C4	-5.49	1.31	1.35
85	AA	1351	U	P-O5'	-5.49	1.54	1.59
85	AA	1668	G	O3'-P	-5.49	1.54	1.61
85	AA	1906	C	P-O5'	-5.49	1.54	1.59
85	AA	2202	G	C1'-N9	-5.49	1.39	1.46
34	BA	405	C	O3'-P	-5.48	1.54	1.61
34	BA	1530	G	N7-C5	-5.48	1.35	1.39
35	BB	53	C	C4'-O4'	-5.48	1.38	1.45
35	BB	400	C	C3'-C2'	-5.48	1.46	1.52
35	BB	1129	C	C2-N3	-5.48	1.31	1.35
38	BE	189	A	C5-C4	-5.48	1.34	1.38
85	AA	965	G	O4'-C1'	-5.48	1.34	1.41
85	AA	1177	G	P-O5'	-5.48	1.54	1.59
85	AA	1676	G	C2-N3	-5.48	1.28	1.32
85	AA	2016	A	C5'-C4'	5.48	1.57	1.51
85	AA	2146	G	N1-C2	-5.48	1.33	1.37
34	BA	183	G	C6-N1	-5.48	1.35	1.39
34	BA	203	U	C3'-C2'	-5.48	1.46	1.52
34	BA	251	U	C4'-C3'	-5.48	1.47	1.52
34	BA	803	U	O4'-C1'	-5.48	1.34	1.41
34	BA	1274	A	C5-C4	-5.48	1.34	1.38
34	BA	1586	U	C2'-C1'	-5.48	1.47	1.53
35	BB	73	G	O3'-P	-5.48	1.54	1.61
35	BB	1295	A	O3'-P	-5.48	1.54	1.61
35	BB	1346	A	N3-C4	-5.48	1.31	1.34
38	BE	54	U	P-O5'	-5.48	1.54	1.59
40	BG	42	A	C2'-C1'	-5.48	1.47	1.53
40	BG	95	U	C5'-C4'	-5.48	1.44	1.51
85	AA	971	U	N3-C4	-5.48	1.33	1.38
85	AA	1487	G	C6-N1	-5.48	1.35	1.39
85	AA	1559	U	C2'-C1'	-5.48	1.47	1.53
85	AA	1697	C	C4'-C3'	-5.48	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1840	C	O3'-P	-5.48	1.54	1.61
85	AA	1894	G	C5-C6	-5.48	1.36	1.42
34	BA	180	G	C2-N3	5.48	1.37	1.32
34	BA	479	U	C2'-C1'	-5.48	1.47	1.53
34	BA	1033	G	P-O5'	-5.48	1.54	1.59
35	BB	359	A	C4'-C3'	-5.48	1.47	1.52
35	BB	517	G	C1'-N9	-5.48	1.39	1.46
35	BB	564	U	C2'-C1'	-5.48	1.47	1.53
35	BB	795	A	C6-N1	-5.48	1.31	1.35
35	BB	850	U	O3'-P	-5.48	1.54	1.61
35	BB	1238	A	C3'-C2'	-5.48	1.46	1.52
35	BB	1440	A	C6-N6	-5.48	1.29	1.33
36	BC	62	A	N7-C5	-5.48	1.35	1.39
38	BE	8	G	C2-N2	-5.48	1.29	1.34
41	BH	129	G	C2-N2	-5.48	1.29	1.34
85	AA	121	C	C4'-C3'	5.48	1.59	1.53
85	AA	530	A	N9-C4	5.48	1.41	1.37
85	AA	1697	C	C3'-C2'	-5.48	1.46	1.52
34	BA	50	G	P-O5'	-5.48	1.54	1.59
34	BA	1362	A	N9-C4	-5.48	1.34	1.37
34	BA	1453	U	C2'-C1'	-5.48	1.47	1.53
35	BB	607	G	C5-C4	-5.48	1.34	1.38
35	BB	650	A	C1'-N9	-5.48	1.39	1.46
35	BB	1057	G	C5'-C4'	-5.48	1.44	1.51
35	BB	1316	U	N3-C4	-5.48	1.33	1.38
36	BC	72	A	P-O5'	-5.48	1.54	1.59
37	BD	28	C	P-O5'	-5.48	1.54	1.59
37	BD	86	A	O4'-C1'	-5.48	1.34	1.41
40	BG	150	A	N9-C4	-5.48	1.34	1.37
85	AA	1808	G	C8-N7	-5.48	1.27	1.30
85	AA	2076	C	O4'-C1'	-5.48	1.34	1.41
34	BA	142	A	C4'-C3'	-5.48	1.47	1.52
34	BA	272	A	C2'-C1'	-5.48	1.47	1.53
34	BA	342	U	C2-N3	-5.48	1.33	1.37
34	BA	768	G	C6-N1	-5.48	1.35	1.39
34	BA	846	U	C2'-C1'	-5.48	1.47	1.53
34	BA	892	C	C4'-O4'	-5.48	1.38	1.45
34	BA	1020	A	O3'-P	-5.48	1.54	1.61
34	BA	1242	A	C3'-C2'	-5.48	1.46	1.52
34	BA	1443	U	N3-C4	-5.48	1.33	1.38
35	BB	751	A	O3'-P	-5.48	1.54	1.61
35	BB	1306	G	O3'-P	-5.48	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1534	U	C4'-O4'	-5.48	1.38	1.45
36	BC	76	C	C3'-C2'	-5.48	1.46	1.52
38	BE	10	G	N7-C5	-5.48	1.35	1.39
39	BF	28	C	C2'-C1'	-5.48	1.47	1.53
85	AA	145	C	P-O5'	-5.48	1.54	1.59
85	AA	589	A	C4'-C3'	5.48	1.59	1.53
85	AA	936	C	C2'-C1'	-5.48	1.47	1.53
85	AA	978	U	C1'-N1	-5.48	1.39	1.46
85	AA	1472	G	C2'-C1'	-5.48	1.47	1.53
85	AA	1572	C	O3'-P	-5.48	1.54	1.61
85	AA	1576	G	N9-C4	-5.48	1.33	1.38
85	AA	1625	C	P-O5'	-5.48	1.54	1.59
34	BA	11	U	C5'-C4'	-5.48	1.44	1.51
34	BA	1174	A	C2'-C1'	-5.48	1.47	1.53
34	BA	1271	C	C2-N3	-5.48	1.31	1.35
34	BA	1285	G	N9-C4	-5.48	1.33	1.38
34	BA	1723	U	C4'-O4'	5.48	1.52	1.45
35	BB	10	C	O3'-P	-5.48	1.54	1.61
35	BB	1153	G	C2'-C1'	-5.48	1.47	1.53
35	BB	1468	A	N9-C4	-5.48	1.34	1.37
36	BC	29	C	P-O5'	-5.48	1.54	1.59
36	BC	36	G	C5-C4	-5.48	1.34	1.38
39	BF	22	U	N1-C2	5.48	1.43	1.38
85	AA	486	G	C6-N1	5.48	1.43	1.39
85	AA	616	A	C1'-N9	-5.48	1.39	1.46
85	AA	1503	G	C3'-C2'	-5.48	1.46	1.52
85	AA	1624	U	C2-N3	-5.48	1.33	1.37
34	BA	65	A	O3'-P	-5.47	1.54	1.61
34	BA	760	G	C5-C4	-5.47	1.34	1.38
34	BA	996	U	C2'-C1'	-5.47	1.47	1.53
34	BA	1604	A	O4'-C1'	-5.47	1.34	1.41
35	BB	764	C	O3'-P	-5.47	1.54	1.61
35	BB	1094	A	C5'-C4'	-5.47	1.44	1.51
35	BB	1181	A	N9-C8	-5.47	1.33	1.37
35	BB	1303	A	N1-C2	-5.47	1.29	1.34
35	BB	1435	G	C2'-C1'	-5.47	1.47	1.53
36	BC	53	A	C5-C4	-5.47	1.34	1.38
41	BH	22	A	C5-C4	-5.47	1.34	1.38
85	AA	251	A	N7-C5	-5.47	1.35	1.39
85	AA	862	U	N1-C2	5.47	1.43	1.38
85	AA	874	A	C1'-N9	-5.47	1.39	1.46
85	AA	1300	A	C2'-C1'	-5.47	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1712	A	O3'-P	-5.47	1.54	1.61
34	BA	27	G	C2'-C1'	-5.47	1.47	1.53
34	BA	517	A	O5'-C5'	-5.47	1.34	1.42
34	BA	825	G	C1'-N9	-5.47	1.39	1.46
35	BB	535	U	C2'-C1'	-5.47	1.47	1.53
35	BB	1191	G	C5-C4	-5.47	1.34	1.38
35	BB	1284	U	C4'-C3'	-5.47	1.47	1.52
35	BB	1449	G	C2-N3	-5.47	1.28	1.32
35	BB	1491	G	C2-N2	-5.47	1.29	1.34
36	BC	117	A	C8-N7	-5.47	1.27	1.31
39	BF	4	A	O3'-P	-5.47	1.54	1.61
40	BG	169	A	C2'-C1'	-5.47	1.47	1.53
85	AA	202	U	O3'-P	-5.47	1.54	1.61
85	AA	1509	A	O3'-P	-5.47	1.54	1.61
85	AA	1547	G	O3'-P	-5.47	1.54	1.61
85	AA	2168	C	O3'-P	-5.47	1.54	1.61
34	BA	413	A	P-O5'	-5.47	1.54	1.59
34	BA	738	C	C2'-C1'	-5.47	1.47	1.53
85	AA	130	G	N7-C5	-5.47	1.35	1.39
85	AA	368	C	C4'-C3'	5.47	1.59	1.53
85	AA	985	G	O3'-P	-5.47	1.54	1.61
85	AA	2218	G	C2'-C1'	-5.47	1.47	1.53
34	BA	174	A	C3'-O3'	-5.47	1.34	1.42
34	BA	255	G	N9-C8	-5.47	1.34	1.37
34	BA	309	U	C2'-C1'	-5.47	1.47	1.53
34	BA	857	C	C4'-O4'	-5.47	1.38	1.45
34	BA	1006	G	C2'-C1'	-5.47	1.47	1.53
34	BA	1623	U	C4'-O4'	-5.47	1.38	1.45
35	BB	1189	C	C4-N4	-5.47	1.29	1.33
38	BE	138	U	C4'-C3'	-5.47	1.47	1.52
39	BF	4	A	C8-N7	-5.47	1.27	1.31
40	BG	2	U	N1-C2	-5.47	1.33	1.38
40	BG	56	G	C8-N7	-5.47	1.27	1.30
40	BG	129	G	N1-C2	-5.47	1.33	1.37
85	AA	53	G	C3'-C2'	-5.47	1.46	1.52
85	AA	116	G	P-O5'	-5.47	1.54	1.59
85	AA	164	G	N7-C5	-5.47	1.35	1.39
85	AA	386	G	C4'-C3'	-5.47	1.47	1.52
85	AA	388	G	C4'-C3'	-5.47	1.47	1.52
85	AA	492	C	C4-N4	-5.47	1.29	1.33
85	AA	531	G	C1'-N9	-5.47	1.39	1.46
85	AA	881	C	C3'-C2'	-5.47	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1193	A	C5'-C4'	5.47	1.57	1.51
85	AA	2057	G	C5-C4	-5.47	1.34	1.38
85	AA	2102	A	O3'-P	-5.47	1.54	1.61
34	BA	1461	A	C2'-C1'	-5.47	1.47	1.53
35	BB	798	A	C5-C4	-5.47	1.34	1.38
35	BB	1492	C	C2-N3	-5.47	1.31	1.35
36	BC	38	U	O4'-C1'	-5.47	1.34	1.41
85	AA	157	G	N1-C2	-5.47	1.33	1.37
86	AB	65	G	O3'-P	-5.47	1.54	1.61
34	BA	816	G	C5'-C4'	5.47	1.57	1.51
34	BA	1550	G	C6-N1	-5.47	1.35	1.39
35	BB	815	G	C3'-C2'	-5.47	1.46	1.52
35	BB	975	G	C1'-N9	-5.47	1.39	1.46
35	BB	1389	C	C2-N3	-5.47	1.31	1.35
36	BC	112	G	N9-C8	-5.47	1.34	1.37
36	BC	135	A	C3'-C2'	-5.47	1.46	1.52
39	BF	57	C	C2'-C1'	-5.47	1.47	1.53
85	AA	159	G	N3-C4	-5.47	1.31	1.35
85	AA	428	G	C2-N3	-5.47	1.28	1.32
85	AA	805	A	P-O5'	-5.47	1.54	1.59
85	AA	907	G	C2'-C1'	-5.47	1.47	1.53
85	AA	956	C	O3'-P	-5.47	1.54	1.61
85	AA	2141	G	O4'-C1'	-5.47	1.34	1.41
34	BA	723	C	C1'-N1	-5.46	1.39	1.46
34	BA	818	G	O3'-P	-5.46	1.54	1.61
34	BA	1166	A	C4'-O4'	-5.46	1.38	1.45
34	BA	1246	G	N3-C4	-5.46	1.31	1.35
34	BA	1527	G	C2'-C1'	-5.46	1.47	1.53
35	BB	805	G	P-O5'	-5.46	1.54	1.59
38	BE	158	U	P-O5'	-5.46	1.54	1.59
85	AA	383	C	O3'-P	-5.46	1.54	1.61
85	AA	415	G	C2-N3	-5.46	1.28	1.32
85	AA	475	A	N9-C8	-5.46	1.33	1.37
85	AA	485	A	O3'-P	-5.46	1.54	1.61
85	AA	1016	G	P-O5'	-5.46	1.54	1.59
85	AA	1804	U	O3'-P	-5.46	1.54	1.61
85	AA	1937	G	O3'-P	-5.46	1.54	1.61
85	AA	2125	A	C1'-N9	-5.46	1.39	1.46
34	BA	1000	G	N9-C8	-5.46	1.34	1.37
34	BA	1413	G	C2'-C1'	-5.46	1.47	1.53
35	BB	1392	A	C3'-C2'	-5.46	1.46	1.52
36	BC	126	G	N1-C2	-5.46	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	594	C	P-O5'	-5.46	1.54	1.59
85	AA	1681	G	C8-N7	-5.46	1.27	1.30
34	BA	92	G	O3'-P	-5.46	1.54	1.61
34	BA	94	G	N9-C8	-5.46	1.34	1.37
34	BA	1088	G	C5-C6	-5.46	1.36	1.42
34	BA	1098	G	C2-N2	-5.46	1.29	1.34
34	BA	1192	A	C1'-N9	-5.46	1.39	1.46
34	BA	1508	C	C4-N4	-5.46	1.29	1.33
34	BA	1549	U	C3'-C2'	-5.46	1.46	1.52
35	BB	553	U	C1'-N1	-5.46	1.39	1.46
35	BB	769	C	C4-N4	-5.46	1.29	1.33
35	BB	1249	G	N9-C4	-5.46	1.33	1.38
35	BB	1294	C	C2'-C1'	-5.46	1.47	1.53
35	BB	1329	G	C5-C6	-5.46	1.36	1.42
37	BD	112	U	C2'-C1'	-5.46	1.47	1.53
38	BE	17	U	O3'-P	-5.46	1.54	1.61
85	AA	428	G	C6-N1	-5.46	1.35	1.39
85	AA	620	U	N3-C4	-5.46	1.33	1.38
85	AA	650	G	C5-C4	-5.46	1.34	1.38
85	AA	867	G	C6-N1	-5.46	1.35	1.39
85	AA	1530	U	O3'-P	-5.46	1.54	1.61
85	AA	1812	C	C4'-C3'	-5.46	1.47	1.52
85	AA	2085	C	C4'-O4'	-5.46	1.38	1.45
85	AA	2123	U	N3-C4	-5.46	1.33	1.38
85	AA	2224	U	O3'-P	-5.46	1.54	1.61
34	BA	224	G	N9-C4	-5.46	1.33	1.38
34	BA	355	U	C2'-C1'	-5.46	1.47	1.53
34	BA	692	U	C4'-O4'	-5.46	1.38	1.45
34	BA	883	C	O3'-P	-5.46	1.54	1.61
34	BA	1224	A	N3-C4	-5.46	1.31	1.34
34	BA	1331	G	C2-N2	-5.46	1.29	1.34
34	BA	1714	A	N3-C4	-5.46	1.31	1.34
35	BB	260	A	N9-C4	-5.46	1.34	1.37
35	BB	425	G	O4'-C1'	-5.46	1.34	1.41
85	AA	363	A	N7-C5	-5.46	1.35	1.39
85	AA	365	G	N1-C2	-5.46	1.33	1.37
85	AA	931	G	O3'-P	-5.46	1.54	1.61
85	AA	1487	G	N9-C8	-5.46	1.34	1.37
85	AA	1578	G	C4'-C3'	5.46	1.59	1.53
34	BA	238	C	O3'-P	-5.46	1.54	1.61
34	BA	557	U	C5'-C4'	5.46	1.57	1.51
34	BA	604	G	C4'-O4'	-5.46	1.38	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	808	U	O3'-P	-5.46	1.54	1.61
34	BA	847	U	C4'-C3'	5.46	1.59	1.53
34	BA	1062	G	N9-C8	-5.46	1.34	1.37
34	BA	1112	U	N3-C4	-5.46	1.33	1.38
34	BA	1707	C	C4-N4	-5.46	1.29	1.33
35	BB	14	C	C2-N3	-5.46	1.31	1.35
35	BB	41	A	C4'-C3'	-5.46	1.47	1.52
35	BB	787	A	C4'-C3'	-5.46	1.47	1.52
35	BB	853	U	C2'-C1'	-5.46	1.47	1.53
35	BB	1193	G	C2'-C1'	-5.46	1.47	1.53
38	BE	11	A	N3-C4	-5.46	1.31	1.34
40	BG	89	A	C8-N7	-5.46	1.27	1.31
40	BG	151	A	C4'-C3'	-5.46	1.47	1.52
85	AA	10	G	N7-C5	-5.46	1.35	1.39
85	AA	66	U	C4'-C3'	5.46	1.59	1.53
85	AA	167	A	C5-C4	-5.46	1.34	1.38
34	BA	167	U	O3'-P	-5.46	1.54	1.61
34	BA	946	A	C5-C4	-5.46	1.34	1.38
34	BA	982	A	O3'-P	-5.46	1.54	1.61
34	BA	1265	G	C2'-C1'	-5.46	1.47	1.53
34	BA	1549	U	C5'-C4'	-5.46	1.44	1.51
35	BB	17	U	N3-C4	-5.46	1.33	1.38
35	BB	22	A	N3-C4	-5.46	1.31	1.34
35	BB	67	A	C3'-C2'	-5.46	1.46	1.52
35	BB	411	A	C1'-N9	-5.46	1.39	1.46
35	BB	456	A	C5-C4	-5.46	1.34	1.38
35	BB	480	C	P-O5'	-5.46	1.54	1.59
35	BB	1091	C	C2'-C1'	-5.46	1.47	1.53
35	BB	1122	C	C2'-C1'	-5.46	1.47	1.53
35	BB	1377	A	C5-C4	-5.46	1.34	1.38
35	BB	1389	C	C4'-C3'	-5.46	1.47	1.52
36	BC	4	G	C2-N3	-5.46	1.28	1.32
37	BD	96	C	C1'-N1	-5.46	1.39	1.46
38	BE	116	U	C2'-C1'	-5.46	1.47	1.53
38	BE	175	U	C5'-C4'	5.46	1.57	1.51
40	BG	74	G	N3-C4	-5.46	1.31	1.35
85	AA	2045	U	C2'-C1'	-5.46	1.47	1.53
34	BA	377	G	O3'-P	-5.46	1.54	1.61
34	BA	1413	G	O3'-P	-5.46	1.54	1.61
35	BB	266	C	P-O5'	-5.46	1.54	1.59
35	BB	645	C	C2-N3	-5.46	1.31	1.35
35	BB	1361	A	C1'-N9	-5.46	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	28	C	C2-N3	-5.46	1.31	1.35
40	BG	52	A	C4'-C3'	-5.46	1.47	1.52
40	BG	136	G	N9-C8	-5.46	1.34	1.37
41	BH	108	U	C2'-C1'	-5.46	1.47	1.53
85	AA	748	C	C2'-C1'	-5.46	1.47	1.53
85	AA	1467	U	C2'-C1'	-5.46	1.47	1.53
34	BA	82	A	C5-C4	-5.45	1.34	1.38
34	BA	373	G	C5-C6	-5.45	1.36	1.42
34	BA	408	U	C1'-N1	-5.45	1.39	1.46
34	BA	499	C	C4-N4	-5.45	1.29	1.33
34	BA	648	C	C2'-C1'	-5.45	1.47	1.53
34	BA	976	C	C2'-C1'	-5.45	1.47	1.53
34	BA	1049	G	N3-C4	-5.45	1.31	1.35
34	BA	1293	A	C1'-N9	-5.45	1.39	1.46
34	BA	1427	U	C1'-N1	-5.45	1.39	1.46
34	BA	1542	A	C4'-O4'	-5.45	1.38	1.45
34	BA	1666	U	O3'-P	-5.45	1.54	1.61
36	BC	169	G	C2-N3	-5.45	1.28	1.32
38	BE	62	C	C2'-C1'	-5.45	1.47	1.53
39	BF	2	G	C2'-C1'	-5.45	1.47	1.53
40	BG	149	U	C3'-C2'	-5.45	1.46	1.52
40	BG	174	G	C8-N7	-5.45	1.27	1.30
41	BH	127	A	C5-C4	-5.45	1.34	1.38
85	AA	30	G	C5-C4	-5.45	1.34	1.38
85	AA	756	G	C6-N1	-5.45	1.35	1.39
85	AA	766	G	N9-C8	-5.45	1.34	1.37
85	AA	862	U	C3'-C2'	-5.45	1.46	1.52
85	AA	974	U	P-O5'	5.45	1.65	1.59
85	AA	1903	G	C2-N2	-5.45	1.29	1.34
85	AA	2150	G	O3'-P	-5.45	1.54	1.61
85	AA	2240	G	P-O5'	-5.45	1.54	1.59
34	BA	1202	G	N7-C5	-5.45	1.35	1.39
34	BA	1451	A	C4'-O4'	-5.45	1.38	1.45
34	BA	1661	U	C2-N3	-5.45	1.33	1.37
34	BA	1680	G	C5-C6	-5.45	1.36	1.42
35	BB	1042	U	C3'-C2'	-5.45	1.46	1.52
35	BB	1213	U	C2-N3	-5.45	1.33	1.37
36	BC	105	C	C2-N3	-5.45	1.31	1.35
40	BG	37	G	C6-N1	-5.45	1.35	1.39
85	AA	20	G	O3'-P	-5.45	1.54	1.61
85	AA	509	C	C3'-O3'	-5.45	1.34	1.42
85	AA	623	G	C5-C4	-5.45	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	978	U	C2-N3	-5.45	1.33	1.37
85	AA	2170	G	C1'-N9	-5.45	1.39	1.46
34	BA	65	A	N9-C8	-5.45	1.33	1.37
34	BA	78	U	C1'-N1	-5.45	1.39	1.46
34	BA	238	C	C1'-N1	-5.45	1.39	1.46
34	BA	450	G	C2-N2	-5.45	1.29	1.34
34	BA	677	U	C2'-C1'	-5.45	1.47	1.53
34	BA	909	G	C2-N2	-5.45	1.29	1.34
34	BA	1041	U	N1-C2	-5.45	1.33	1.38
34	BA	1506	C	C4'-O4'	-5.45	1.38	1.45
35	BB	7	C	O4'-C1'	-5.45	1.34	1.41
36	BC	38	U	P-O5'	-5.45	1.54	1.59
40	BG	156	G	C2'-C1'	-5.45	1.47	1.53
85	AA	206	U	O3'-P	-5.45	1.54	1.61
85	AA	422	G	N9-C8	-5.45	1.34	1.37
85	AA	515	C	C1'-N1	-5.45	1.39	1.46
85	AA	691	U	C5'-C4'	5.45	1.57	1.51
85	AA	886	A	P-O5'	-5.45	1.54	1.59
85	AA	1150	G	C1'-N9	-5.45	1.39	1.46
34	BA	207	A	N7-C5	-5.45	1.35	1.39
34	BA	563	A	C2'-C1'	-5.45	1.47	1.53
34	BA	932	G	N1-C2	-5.45	1.33	1.37
34	BA	1101	A	N3-C4	-5.45	1.31	1.34
34	BA	1201	G	O3'-P	-5.45	1.54	1.61
34	BA	1658	G	C6-N1	-5.45	1.35	1.39
34	BA	1696	G	C8-N7	-5.45	1.27	1.30
34	BA	1724	G	C5-C4	-5.45	1.34	1.38
35	BB	728	A	N3-C4	-5.45	1.31	1.34
35	BB	1509	G	C1'-N9	-5.45	1.39	1.46
37	BD	52	U	C2'-C1'	-5.45	1.47	1.53
39	BF	9	C	O4'-C1'	-5.45	1.34	1.41
39	BF	30	C	C1'-N1	-5.45	1.39	1.46
85	AA	14	C	P-O5'	-5.45	1.54	1.59
85	AA	382	G	C2-N2	-5.45	1.29	1.34
85	AA	2174	G	O4'-C1'	-5.45	1.34	1.41
85	AA	2179	C	C1'-N1	-5.45	1.39	1.46
34	BA	424	U	C4'-C3'	-5.45	1.47	1.52
34	BA	567	U	O3'-P	-5.45	1.54	1.61
35	BB	94	A	C8-N7	-5.45	1.27	1.31
35	BB	525	U	O3'-P	-5.45	1.54	1.61
85	AA	318	A	O4'-C1'	-5.45	1.34	1.41
31	AX	182	GLY	CA-C	-5.45	1.43	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	299	C	O3'-P	-5.45	1.54	1.61
34	BA	596	G	C8-N7	-5.45	1.27	1.30
34	BA	1433	U	P-O5'	-5.45	1.54	1.59
34	BA	1595	G	C2'-C1'	-5.45	1.47	1.53
35	BB	596	C	C4-N4	-5.45	1.29	1.33
35	BB	618	U	C2'-C1'	-5.45	1.47	1.53
35	BB	689	C	C2-N3	-5.45	1.31	1.35
38	BE	127	G	O3'-P	-5.45	1.54	1.61
38	BE	150	G	C2-N3	5.45	1.37	1.32
40	BG	31	G	C8-N7	-5.45	1.27	1.30
41	BH	26	C	N1-C2	-5.45	1.34	1.40
85	AA	267	U	C2-N3	-5.45	1.33	1.37
85	AA	1225	C	C5'-C4'	-5.45	1.44	1.51
85	AA	1351	U	O3'-P	-5.45	1.54	1.61
85	AA	1589	G	N7-C5	-5.45	1.35	1.39
85	AA	1869	U	C3'-C2'	-5.45	1.46	1.52
85	AA	2138	G	N9-C4	-5.45	1.33	1.38
34	BA	490	A	N3-C4	-5.44	1.31	1.34
34	BA	846	U	P-O5'	-5.44	1.54	1.59
34	BA	1070	G	O3'-P	-5.44	1.54	1.61
34	BA	1205	A	P-O5'	-5.44	1.54	1.59
85	AA	65	A	N7-C5	-5.44	1.35	1.39
85	AA	346	U	C2'-C1'	-5.44	1.47	1.53
85	AA	529	G	N9-C4	-5.44	1.33	1.38
85	AA	651	G	C2'-C1'	-5.44	1.47	1.53
85	AA	787	U	N1-C2	-5.44	1.33	1.38
85	AA	1320	G	O3'-P	-5.44	1.54	1.61
34	BA	631	G	C2'-C1'	-5.44	1.47	1.53
34	BA	914	G	C2'-C1'	-5.44	1.47	1.53
34	BA	1252	G	C2'-C1'	-5.44	1.47	1.53
34	BA	1415	C	C2-N3	-5.44	1.31	1.35
34	BA	1498	A	O3'-P	-5.44	1.54	1.61
37	BD	51	G	O3'-P	-5.44	1.54	1.61
40	BG	101	G	N9-C4	-5.44	1.33	1.38
85	AA	153	C	P-O5'	-5.44	1.54	1.59
85	AA	675	A	P-O5'	-5.44	1.54	1.59
85	AA	1478	G	N7-C5	-5.44	1.35	1.39
34	BA	36	A	N9-C8	-5.44	1.33	1.37
34	BA	86	A	C3'-C2'	-5.44	1.46	1.52
34	BA	94	G	N7-C5	-5.44	1.35	1.39
34	BA	269	G	N9-C8	-5.44	1.34	1.37
34	BA	373	G	C1'-N9	-5.44	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	465	A	C5-C4	-5.44	1.34	1.38
34	BA	773	A	N9-C4	-5.44	1.34	1.37
34	BA	1191	C	N1-C6	-5.44	1.33	1.37
34	BA	1461	A	C1'-N9	-5.44	1.39	1.46
35	BB	1277	A	O3'-P	-5.44	1.54	1.61
35	BB	1351	G	N1-C2	-5.44	1.33	1.37
36	BC	72	A	C1'-N9	-5.44	1.39	1.46
36	BC	148	C	C3'-C2'	-5.44	1.46	1.52
41	BH	30	C	N1-C6	-5.44	1.33	1.37
41	BH	35	G	C2-N2	-5.44	1.29	1.34
85	AA	243	A	O3'-P	-5.44	1.54	1.61
85	AA	1243	G	C3'-C2'	-5.44	1.46	1.52
85	AA	1277	C	O3'-P	-5.44	1.54	1.61
85	AA	1862	C	C3'-C2'	-5.44	1.46	1.52
85	AA	2182	A	N3-C4	-5.44	1.31	1.34
34	BA	101	G	C1'-N9	-5.44	1.39	1.46
34	BA	258	C	C2-N3	-5.44	1.31	1.35
34	BA	329	G	C2'-C1'	-5.44	1.47	1.53
34	BA	1593	U	O3'-P	-5.44	1.54	1.61
34	BA	1821	A	C1'-N9	-5.44	1.39	1.46
35	BB	1199	A	C2'-C1'	-5.44	1.47	1.53
35	BB	1308	G	N9-C8	-5.44	1.34	1.37
35	BB	1350	A	C1'-N9	-5.44	1.39	1.46
35	BB	1387	C	C2'-C1'	-5.44	1.47	1.53
38	BE	45	G	C5-C4	-5.44	1.34	1.38
38	BE	152	U	O3'-P	-5.44	1.54	1.61
38	BE	189	A	C1'-N9	-5.44	1.39	1.46
85	AA	628	C	N1-C6	-5.44	1.33	1.37
85	AA	1240	A	C5-C6	-5.44	1.36	1.41
85	AA	1668	G	C3'-C2'	-5.44	1.46	1.52
34	BA	687	G	C5'-C4'	-5.44	1.44	1.51
34	BA	750	C	O3'-P	-5.44	1.54	1.61
34	BA	1040	G	C3'-C2'	-5.44	1.46	1.52
34	BA	1451	A	C1'-N9	-5.44	1.39	1.46
35	BB	267	C	O3'-P	-5.44	1.54	1.61
35	BB	706	G	C2'-C1'	-5.44	1.47	1.53
35	BB	1187	G	N9-C4	5.44	1.42	1.38
35	BB	1513	U	C3'-C2'	-5.44	1.46	1.52
36	BC	79	A	P-O5'	-5.44	1.54	1.59
37	BD	7	G	N3-C4	-5.44	1.31	1.35
38	BE	25	U	C4-C5	-5.44	1.38	1.43
41	BH	31	A	N9-C8	-5.44	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	429	G	C2-N2	-5.44	1.29	1.34
85	AA	1146	C	C3'-C2'	-5.44	1.46	1.52
85	AA	1196	C	C4'-O4'	-5.44	1.38	1.45
85	AA	1329	U	O3'-P	-5.44	1.54	1.61
85	AA	1450	U	C4-C5	-5.44	1.38	1.43
34	BA	331	G	N3-C4	-5.44	1.31	1.35
34	BA	787	A	O3'-P	-5.44	1.54	1.61
34	BA	914	G	N9-C4	-5.44	1.33	1.38
34	BA	1569	C	O3'-P	-5.44	1.54	1.61
35	BB	1482	A	C4'-C3'	-5.44	1.47	1.52
39	BF	72	A	C3'-C2'	-5.44	1.46	1.52
41	BH	112	U	C2'-C1'	-5.44	1.47	1.53
85	AA	408	C	O3'-P	-5.44	1.54	1.61
85	AA	1172	A	N9-C4	-5.44	1.34	1.37
85	AA	1981	A	N3-C4	-5.44	1.31	1.34
27	AT	36	GLY	CA-C	-5.43	1.43	1.51
34	BA	478	G	N9-C4	-5.43	1.33	1.38
34	BA	1008	A	C1'-N9	-5.43	1.39	1.46
34	BA	1196	C	O4'-C1'	-5.43	1.34	1.41
34	BA	1253	G	C2-N2	-5.43	1.29	1.34
34	BA	1497	A	N7-C5	-5.43	1.35	1.39
34	BA	1542	A	N7-C5	-5.43	1.35	1.39
34	BA	1730	A	N9-C4	5.43	1.41	1.37
35	BB	50	A	C3'-C2'	-5.43	1.46	1.52
35	BB	396	C	P-O5'	-5.43	1.54	1.59
35	BB	529	A	C6-N1	5.43	1.39	1.35
35	BB	1349	U	C3'-C2'	-5.43	1.46	1.52
36	BC	40	A	N7-C5	-5.43	1.35	1.39
36	BC	63	G	C6-N1	-5.43	1.35	1.39
38	BE	48	G	N7-C5	-5.43	1.35	1.39
38	BE	189	A	N7-C5	-5.43	1.35	1.39
85	AA	309	G	C1'-N9	-5.43	1.39	1.46
85	AA	446	C	C4-N4	-5.43	1.29	1.33
85	AA	609	U	N3-C4	-5.43	1.33	1.38
85	AA	1926	A	O3'-P	-5.43	1.54	1.61
85	AA	2077	G	C8-N7	-5.43	1.27	1.30
34	BA	849	G	O4'-C1'	-5.43	1.34	1.41
34	BA	1588	U	C4'-C3'	-5.43	1.47	1.52
35	BB	114	A	N3-C4	-5.43	1.31	1.34
35	BB	405	U	O4'-C1'	-5.43	1.34	1.41
35	BB	490	G	C4'-C3'	-5.43	1.47	1.52
35	BB	691	A	P-O5'	-5.43	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	796	C	C4'-C3'	-5.43	1.47	1.52
35	BB	1110	G	C1'-N9	-5.43	1.39	1.46
35	BB	1253	U	N3-C4	-5.43	1.33	1.38
85	AA	657	C	O3'-P	-5.43	1.54	1.61
85	AA	680	U	O4'-C1'	-5.43	1.34	1.41
85	AA	1203	G	C2'-C1'	-5.43	1.47	1.53
85	AA	2180	C	C2-N3	-5.43	1.31	1.35
34	BA	451	A	C2'-C1'	-5.43	1.47	1.53
34	BA	1483	U	N3-C4	-5.43	1.33	1.38
35	BB	258	C	P-O5'	-5.43	1.54	1.59
40	BG	50	G	N9-C4	-5.43	1.33	1.38
85	AA	308	U	O3'-P	-5.43	1.54	1.61
85	AA	806	G	N7-C5	-5.43	1.35	1.39
85	AA	976	G	N1-C2	-5.43	1.33	1.37
85	AA	995	G	C6-N1	-5.43	1.35	1.39
34	BA	932	G	C1'-N9	-5.43	1.39	1.46
34	BA	939	C	C4-N4	-5.43	1.29	1.33
34	BA	952	G	N1-C2	-5.43	1.33	1.37
34	BA	1204	U	N3-C4	-5.43	1.33	1.38
34	BA	1414	C	C4-N4	-5.43	1.29	1.33
34	BA	1742	G	C6-N1	-5.43	1.35	1.39
35	BB	380	G	N1-C2	-5.43	1.33	1.37
35	BB	483	C	O3'-P	-5.43	1.54	1.61
35	BB	602	G	C2-N2	-5.43	1.29	1.34
35	BB	639	A	C4'-O4'	-5.43	1.38	1.45
35	BB	1173	C	P-O5'	-5.43	1.54	1.59
35	BB	1227	G	C5'-C4'	-5.43	1.44	1.51
35	BB	1387	C	C4-N4	-5.43	1.29	1.33
38	BE	205	G	N9-C4	-5.43	1.33	1.38
40	BG	27	C	C1'-N1	-5.43	1.39	1.46
85	AA	107	A	C8-N7	-5.43	1.27	1.31
85	AA	147	G	O3'-P	-5.43	1.54	1.61
85	AA	1188	A	C2'-C1'	-5.43	1.47	1.53
85	AA	1215	A	N3-C4	-5.43	1.31	1.34
85	AA	1263	G	C5-C6	-5.43	1.36	1.42
85	AA	1279	A	N9-C8	-5.43	1.33	1.37
85	AA	1368	G	N7-C5	-5.43	1.35	1.39
34	BA	71	G	C2-N2	-5.43	1.29	1.34
34	BA	100	A	C5-C4	-5.43	1.34	1.38
34	BA	711	C	C4'-C3'	5.43	1.59	1.53
34	BA	930	A	C3'-C2'	-5.43	1.46	1.52
34	BA	1245	C	C4'-C3'	-5.43	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1566	G	C2-N2	-5.43	1.29	1.34
36	BC	31	A	C3'-C2'	-5.43	1.46	1.52
85	AA	2041	G	C6-N1	-5.43	1.35	1.39
85	AA	2043	A	N7-C5	-5.43	1.35	1.39
34	BA	182	U	O3'-P	-5.43	1.54	1.61
34	BA	216	C	C4'-C3'	-5.43	1.47	1.52
34	BA	584	A	P-O5'	-5.43	1.54	1.59
34	BA	887	U	N3-C4	-5.43	1.33	1.38
34	BA	901	C	C3'-C2'	-5.43	1.46	1.52
34	BA	1641	G	C5'-C4'	-5.43	1.44	1.51
35	BB	392	G	C2-N2	-5.43	1.29	1.34
35	BB	790	A	O3'-P	-5.43	1.54	1.61
35	BB	1213	U	P-O5'	-5.43	1.54	1.59
35	BB	1414	A	O3'-P	-5.43	1.54	1.61
35	BB	1429	A	N9-C4	-5.43	1.34	1.37
35	BB	1434	G	N9-C8	-5.43	1.34	1.37
38	BE	92	C	C2'-C1'	-5.43	1.47	1.53
38	BE	95	G	C3'-C2'	-5.43	1.46	1.52
38	BE	187	G	N7-C5	-5.43	1.35	1.39
39	BF	5	U	P-O5'	-5.43	1.54	1.59
39	BF	23	G	N9-C4	-5.43	1.33	1.38
40	BG	38	A	C4'-O4'	-5.43	1.38	1.45
41	BH	34	G	N9-C8	-5.43	1.34	1.37
85	AA	168	A	O3'-P	-5.43	1.54	1.61
85	AA	341	C	C2-N3	-5.43	1.31	1.35
85	AA	384	C	C4'-C3'	-5.43	1.47	1.52
85	AA	438	G	N9-C8	-5.43	1.34	1.37
85	AA	616	A	N9-C8	-5.43	1.33	1.37
85	AA	1022	G	O3'-P	-5.43	1.54	1.61
85	AA	1478	G	C3'-C2'	-5.43	1.46	1.52
85	AA	1855	U	N3-C4	-5.43	1.33	1.38
85	AA	2067	A	O3'-P	-5.43	1.54	1.61
85	AA	2126	U	C1'-N1	-5.43	1.39	1.46
34	BA	148	G	C5-C4	-5.42	1.34	1.38
34	BA	687	G	C2-N2	-5.42	1.29	1.34
34	BA	941	G	C2'-C1'	-5.42	1.47	1.53
34	BA	1667	G	C3'-C2'	-5.42	1.46	1.52
35	BB	437	U	O3'-P	-5.42	1.54	1.61
35	BB	598	C	P-O5'	-5.42	1.54	1.59
35	BB	1215	U	O3'-P	-5.42	1.54	1.61
35	BB	1267	C	C5'-C4'	5.42	1.57	1.51
35	BB	1310	C	C1'-N1	-5.42	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	77	A	N3-C4	-5.42	1.31	1.34
41	BH	17	A	C3'-C2'	-5.42	1.46	1.52
85	AA	247	G	N1-C2	-5.42	1.33	1.37
85	AA	312	G	N1-C2	-5.42	1.33	1.37
85	AA	385	A	C5-C6	-5.42	1.36	1.41
85	AA	769	C	O4'-C1'	-5.42	1.34	1.41
34	BA	770	G	C4'-C3'	-5.42	1.47	1.52
34	BA	1455	C	C4'-C3'	-5.42	1.47	1.52
34	BA	1659	G	C2'-C1'	-5.42	1.47	1.53
34	BA	1825	U	O3'-P	-5.42	1.54	1.61
35	BB	592	G	N1-C2	-5.42	1.33	1.37
35	BB	1304	U	C4'-C3'	-5.42	1.47	1.52
35	BB	1440	A	C5-C6	-5.42	1.36	1.41
37	BD	26	C	O5'-C5'	-5.42	1.34	1.42
85	AA	89	C	C2'-C1'	-5.42	1.47	1.53
85	AA	1260	G	N3-C4	-5.42	1.31	1.35
34	BA	130	U	C3'-C2'	-5.42	1.46	1.52
34	BA	167	U	C4'-C3'	-5.42	1.47	1.52
34	BA	455	A	C2'-C1'	-5.42	1.47	1.53
34	BA	920	U	C3'-C2'	-5.42	1.46	1.52
34	BA	935	A	C8-N7	-5.42	1.27	1.31
34	BA	1245	C	N3-C4	-5.42	1.30	1.33
34	BA	1299	G	C2-N2	-5.42	1.29	1.34
35	BB	1306	G	C5-C4	-5.42	1.34	1.38
35	BB	1430	G	C4'-O4'	-5.42	1.38	1.45
35	BB	1533	U	O3'-P	-5.42	1.54	1.61
37	BD	25	G	N3-C4	-5.42	1.31	1.35
40	BG	112	C	C1'-N1	-5.42	1.39	1.46
85	AA	573	U	C2-N3	-5.42	1.33	1.37
85	AA	745	C	O3'-P	-5.42	1.54	1.61
85	AA	1292	A	C5-C6	-5.42	1.36	1.41
85	AA	1456	A	C5-C4	-5.42	1.34	1.38
85	AA	2181	G	N9-C8	-5.42	1.34	1.37
85	AA	2228	G	C5'-C4'	-5.42	1.44	1.51
34	BA	412	G	C5-C6	-5.42	1.36	1.42
34	BA	1656	A	C5-C6	-5.42	1.36	1.41
35	BB	543	G	C2-N2	-5.42	1.29	1.34
35	BB	571	C	C2'-C1'	-5.42	1.47	1.53
35	BB	610	U	C3'-C2'	-5.42	1.46	1.52
37	BD	116	C	C2'-C1'	-5.42	1.47	1.53
85	AA	3	U	C2'-C1'	-5.42	1.47	1.53
85	AA	1233	G	N3-C4	-5.42	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	16	C	C2-N3	-5.42	1.31	1.35
34	BA	307	C	C2-N3	-5.42	1.31	1.35
34	BA	387	A	O3'-P	-5.42	1.54	1.61
34	BA	670	U	N3-C4	-5.42	1.33	1.38
34	BA	1513	G	C2-N2	-5.42	1.29	1.34
35	BB	364	U	O3'-P	-5.42	1.54	1.61
35	BB	523	A	C5'-C4'	5.42	1.57	1.51
35	BB	1065	G	N3-C4	-5.42	1.31	1.35
35	BB	1066	G	C5-C6	-5.42	1.36	1.42
35	BB	1100	C	N3-C4	-5.42	1.30	1.33
35	BB	1124	G	C2-N2	-5.42	1.29	1.34
35	BB	1249	G	C8-N7	-5.42	1.27	1.30
38	BE	46	G	N9-C8	-5.42	1.34	1.37
38	BE	123	A	C5-C4	-5.42	1.34	1.38
38	BE	130	G	N1-C2	-5.42	1.33	1.37
40	BG	46	G	N9-C8	-5.42	1.34	1.37
40	BG	171	A	C5-C4	-5.42	1.34	1.38
85	AA	89	C	P-O5'	-5.42	1.54	1.59
85	AA	1137	C	N1-C6	-5.42	1.33	1.37
85	AA	1708	A	C3'-C2'	-5.42	1.46	1.52
85	AA	2228	G	C4'-C3'	-5.42	1.47	1.52
34	BA	35	U	C1'-N1	-5.42	1.39	1.46
34	BA	226	A	P-O5'	-5.42	1.54	1.59
34	BA	814	C	C2'-C1'	-5.42	1.47	1.53
34	BA	860	G	N9-C4	-5.42	1.33	1.38
34	BA	969	A	O3'-P	-5.42	1.54	1.61
34	BA	1658	G	C2-N2	-5.42	1.29	1.34
35	BB	35	G	O3'-P	-5.42	1.54	1.61
35	BB	598	C	C4-N4	-5.42	1.29	1.33
35	BB	827	U	C2-N3	-5.42	1.33	1.37
35	BB	883	G	C5'-C4'	5.42	1.57	1.51
35	BB	1252	G	C2-N2	-5.42	1.29	1.34
35	BB	1262	A	C3'-C2'	-5.42	1.46	1.52
35	BB	1329	G	O3'-P	-5.42	1.54	1.61
35	BB	1362	G	C6-N1	-5.42	1.35	1.39
35	BB	1479	C	O3'-P	-5.42	1.54	1.61
38	BE	126	G	P-O5'	-5.42	1.54	1.59
38	BE	160	C	C1'-N1	-5.42	1.39	1.46
85	AA	100	A	C6-N6	-5.42	1.29	1.33
85	AA	521	A	C3'-C2'	-5.42	1.46	1.52
85	AA	719	C	C3'-C2'	-5.42	1.46	1.52
85	AA	1827	U	P-O5'	-5.42	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1891	U	C2'-C1'	-5.42	1.47	1.53
85	AA	2244	G	C5-C6	-5.42	1.36	1.42
34	BA	362	G	C5-C4	-5.42	1.34	1.38
34	BA	1049	G	C6-N1	-5.42	1.35	1.39
34	BA	1195	G	C6-N1	-5.42	1.35	1.39
34	BA	1469	G	C5-C4	-5.42	1.34	1.38
35	BB	392	G	C2'-C1'	-5.42	1.47	1.53
35	BB	484	G	C5-C4	-5.42	1.34	1.38
85	AA	1526	G	C4'-O4'	-5.42	1.38	1.45
85	AA	1925	A	C2'-C1'	-5.42	1.47	1.53
86	AB	1	G	N9-C4	-5.42	1.33	1.38
34	BA	336	A	C3'-C2'	-5.41	1.46	1.52
34	BA	1213	A	O3'-P	-5.41	1.54	1.61
34	BA	1435	A	C3'-C2'	-5.41	1.46	1.52
35	BB	18	A	N3-C4	-5.41	1.31	1.34
35	BB	267	C	P-O5'	-5.41	1.54	1.59
35	BB	568	A	N9-C8	-5.41	1.33	1.37
35	BB	1404	A	C4'-C3'	-5.41	1.47	1.52
36	BC	36	G	C4'-O4'	-5.41	1.38	1.45
85	AA	167	A	C2'-C1'	-5.41	1.47	1.53
85	AA	1130	G	N3-C4	-5.41	1.31	1.35
85	AA	1818	C	O3'-P	-5.41	1.54	1.61
85	AA	2094	U	C5'-C4'	-5.41	1.44	1.51
34	BA	55	G	C1'-N9	-5.41	1.39	1.46
34	BA	107	C	C2'-C1'	-5.41	1.47	1.53
34	BA	811	C	C3'-C2'	-5.41	1.46	1.52
34	BA	1636	C	C3'-C2'	-5.41	1.46	1.52
34	BA	1832	A	C6-N1	-5.41	1.31	1.35
35	BB	817	C	C4-C5	-5.41	1.38	1.43
35	BB	1291	G	O3'-P	-5.41	1.54	1.61
37	BD	13	A	C1'-N9	-5.41	1.39	1.46
38	BE	206	G	C4'-C3'	-5.41	1.47	1.52
85	AA	1283	C	C4-N4	-5.41	1.29	1.33
34	BA	18	G	C2'-C1'	-5.41	1.47	1.53
34	BA	137	C	C2-N3	-5.41	1.31	1.35
34	BA	312	U	C2'-C1'	-5.41	1.47	1.53
34	BA	626	G	N7-C5	-5.41	1.36	1.39
34	BA	694	G	C5-C6	-5.41	1.36	1.42
34	BA	1246	G	N7-C5	-5.41	1.36	1.39
34	BA	1782	C	P-O5'	-5.41	1.54	1.59
35	BB	552	C	N1-C2	-5.41	1.34	1.40
35	BB	805	G	C3'-C2'	-5.41	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1244	U	N3-C4	-5.41	1.33	1.38
41	BH	19	G	N9-C8	-5.41	1.34	1.37
85	AA	339	A	C2'-C1'	-5.41	1.47	1.53
85	AA	688	C	N3-C4	-5.41	1.30	1.33
85	AA	755	G	N7-C5	-5.41	1.36	1.39
85	AA	1722	G	P-O5'	-5.41	1.54	1.59
34	BA	221	G	C2-N3	-5.41	1.28	1.32
34	BA	410	G	N7-C5	-5.41	1.36	1.39
34	BA	800	G	C2-N2	-5.41	1.29	1.34
34	BA	1152	A	C2'-C1'	-5.41	1.47	1.53
34	BA	1815	G	C1'-N9	-5.41	1.39	1.46
35	BB	511	A	C6-N1	-5.41	1.31	1.35
35	BB	648	G	C3'-C2'	-5.41	1.46	1.52
37	BD	64	A	C1'-N9	-5.41	1.39	1.46
85	AA	234	G	O3'-P	-5.41	1.54	1.61
85	AA	995	G	N1-C2	-5.41	1.33	1.37
37	BD	10	C	C2'-C1'	-5.41	1.47	1.53
41	BH	66	G	C3'-C2'	-5.41	1.46	1.52
85	AA	116	G	N3-C4	-5.41	1.31	1.35
85	AA	763	U	O4'-C1'	-5.41	1.34	1.41
85	AA	853	G	P-O5'	-5.41	1.54	1.59
34	BA	40	A	C2'-C1'	-5.41	1.47	1.53
34	BA	52	G	C8-N7	-5.41	1.27	1.30
34	BA	582	U	C2'-C1'	-5.41	1.47	1.53
34	BA	859	G	N7-C5	-5.41	1.36	1.39
34	BA	912	G	C2'-C1'	-5.41	1.47	1.53
34	BA	1348	G	P-O5'	-5.41	1.54	1.59
34	BA	1613	G	C2'-C1'	-5.41	1.47	1.53
34	BA	1652	G	C6-N1	-5.41	1.35	1.39
35	BB	539	G	C3'-C2'	-5.41	1.46	1.52
35	BB	1025	A	N7-C5	-5.41	1.36	1.39
35	BB	1385	C	C5'-C4'	-5.41	1.44	1.51
39	BF	14	C	N3-C4	5.41	1.37	1.33
39	BF	56	C	O4'-C1'	-5.41	1.34	1.41
41	BH	23	G	P-O5'	-5.41	1.54	1.59
41	BH	120	C	P-O5'	-5.41	1.54	1.59
81	Bv	75	ARG	CD-NE	5.41	1.55	1.46
85	AA	678	A	N9-C8	-5.41	1.33	1.37
85	AA	1406	U	O3'-P	-5.41	1.54	1.61
85	AA	1458	G	C5'-C4'	-5.41	1.44	1.51
85	AA	1885	A	C3'-C2'	-5.41	1.46	1.52
34	BA	295	G	C2-N2	-5.40	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1124	U	O3'-P	-5.40	1.54	1.61
34	BA	1530	G	N1-C2	-5.40	1.33	1.37
35	BB	1096	G	C2-N2	-5.40	1.29	1.34
37	BD	62	A	C2'-C1'	-5.40	1.47	1.53
37	BD	102	C	P-O5'	-5.40	1.54	1.59
85	AA	881	C	O3'-P	-5.40	1.54	1.61
85	AA	1853	U	O3'-P	-5.40	1.54	1.61
85	AA	1861	A	P-O5'	-5.40	1.54	1.59
34	BA	373	G	C5-C4	-5.40	1.34	1.38
34	BA	660	C	C2'-C1'	-5.40	1.47	1.53
34	BA	709	C	C2-N3	-5.40	1.31	1.35
34	BA	1507	C	N1-C6	-5.40	1.33	1.37
34	BA	1590	G	C3'-C2'	-5.40	1.46	1.52
35	BB	397	C	P-O5'	-5.40	1.54	1.59
35	BB	440	U	C2-N3	-5.40	1.33	1.37
35	BB	843	G	N9-C8	-5.40	1.34	1.37
35	BB	1100	C	C2-N3	-5.40	1.31	1.35
36	BC	40	A	N9-C4	-5.40	1.34	1.37
36	BC	66	G	N3-C4	-5.40	1.31	1.35
37	BD	107	G	C2'-C1'	-5.40	1.47	1.53
40	BG	169	A	N7-C5	-5.40	1.36	1.39
85	AA	270	A	N7-C5	-5.40	1.36	1.39
85	AA	1149	A	C5-C4	-5.40	1.34	1.38
85	AA	1892	G	C4'-C3'	-5.40	1.47	1.52
85	AA	2052	U	C2'-C1'	-5.40	1.47	1.53
85	AA	2223	C	C4-N4	-5.40	1.29	1.33
34	BA	428	C	C2-N3	-5.40	1.31	1.35
34	BA	1812	C	C3'-C2'	-5.40	1.46	1.52
35	BB	311	C	P-O5'	-5.40	1.54	1.59
35	BB	676	G	N9-C8	-5.40	1.34	1.37
35	BB	681	G	C3'-C2'	-5.40	1.46	1.52
35	BB	834	U	C4-C5	-5.40	1.38	1.43
37	BD	17	G	C2'-C1'	-5.40	1.47	1.53
37	BD	69	U	C2'-C1'	-5.40	1.47	1.53
38	BE	91	G	C2'-C1'	-5.40	1.47	1.53
41	BH	7	C	C2-N3	-5.40	1.31	1.35
85	AA	98	U	C2'-C1'	-5.40	1.47	1.53
85	AA	421	G	C3'-C2'	-5.40	1.46	1.52
85	AA	497	G	N9-C4	-5.40	1.33	1.38
85	AA	1111	A	N7-C5	-5.40	1.36	1.39
85	AA	1245	U	O3'-P	-5.40	1.54	1.61
85	AA	1438	C	O3'-P	-5.40	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1444	U	N3-C4	-5.40	1.33	1.38
85	AA	1894	G	C1'-N9	-5.40	1.39	1.46
85	AA	2176	U	C2'-C1'	-5.40	1.47	1.53
34	BA	510	U	C5'-C4'	5.40	1.57	1.51
34	BA	1238	C	C3'-C2'	-5.40	1.46	1.52
35	BB	489	A	P-O5'	-5.40	1.54	1.59
39	BF	9	C	N3-C4	-5.40	1.30	1.33
85	AA	367	A	P-O5'	-5.40	1.54	1.59
85	AA	484	G	O3'-P	-5.40	1.54	1.61
34	BA	716	C	C4'-O4'	-5.40	1.38	1.45
35	BB	408	U	C2'-C1'	-5.40	1.47	1.53
35	BB	649	A	C2'-C1'	-5.40	1.47	1.53
35	BB	1226	G	C5-C6	-5.40	1.36	1.42
38	BE	115	U	O3'-P	-5.40	1.54	1.61
38	BE	117	A	N3-C4	-5.40	1.31	1.34
40	BG	96	C	O3'-P	-5.40	1.54	1.61
40	BG	109	C	N1-C6	-5.40	1.33	1.37
85	AA	179	G	C4'-O4'	-5.40	1.38	1.45
85	AA	394	C	C4-N4	-5.40	1.29	1.33
85	AA	888	A	N7-C5	-5.40	1.36	1.39
85	AA	912	C	O3'-P	-5.40	1.54	1.61
85	AA	1356	U	C2-N3	-5.40	1.33	1.37
85	AA	2107	C	C3'-C2'	-5.40	1.46	1.52
34	BA	470	C	C3'-C2'	-5.40	1.46	1.52
34	BA	851	C	C2-N3	-5.40	1.31	1.35
34	BA	921	G	C1'-N9	-5.40	1.39	1.46
34	BA	1282	G	C2-N3	-5.40	1.28	1.32
34	BA	1414	C	C2-N3	-5.40	1.31	1.35
34	BA	1795	A	N9-C8	-5.40	1.33	1.37
35	BB	1427	A	N9-C4	-5.40	1.34	1.37
36	BC	108	A	N9-C8	-5.40	1.33	1.37
85	AA	719	C	C2-N3	-5.40	1.31	1.35
85	AA	2150	G	N7-C5	-5.40	1.36	1.39
86	AB	15	G	C6-N1	-5.40	1.35	1.39
34	BA	32	A	C8-N7	-5.39	1.27	1.31
34	BA	543	A	N7-C5	-5.39	1.36	1.39
34	BA	933	U	C4'-C3'	-5.39	1.47	1.52
34	BA	942	G	C2'-C1'	-5.39	1.47	1.53
34	BA	1045	C	C2-N3	-5.39	1.31	1.35
35	BB	574	G	C5-C4	-5.39	1.34	1.38
35	BB	1059	U	O3'-P	-5.39	1.54	1.61
36	BC	88	A	C2'-C1'	-5.39	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	119	G	O4'-C1'	-5.39	1.34	1.41
40	BG	9	G	C2-N2	-5.39	1.29	1.34
40	BG	126	G	C5-C6	-5.39	1.36	1.42
65	Bf	93	HIS	N-CA	-5.39	1.35	1.46
85	AA	637	U	O4'-C1'	-5.39	1.34	1.41
85	AA	1471	G	C2-N2	-5.39	1.29	1.34
85	AA	1790	G	P-O5'	-5.39	1.54	1.59
34	BA	515	U	O3'-P	-5.39	1.54	1.61
34	BA	1270	G	C4'-C3'	-5.39	1.47	1.52
34	BA	1323	G	C6-N1	-5.39	1.35	1.39
34	BA	1552	C	C3'-C2'	-5.39	1.46	1.52
34	BA	1667	G	C1'-N9	-5.39	1.39	1.46
34	BA	1686	G	C5-C6	-5.39	1.36	1.42
35	BB	58	G	C2'-C1'	-5.39	1.47	1.53
35	BB	766	G	C4'-C3'	-5.39	1.47	1.52
35	BB	1065	G	P-O5'	-5.39	1.54	1.59
35	BB	1094	A	O4'-C1'	-5.39	1.34	1.41
37	BD	98	G	C2'-C1'	-5.39	1.47	1.53
85	AA	339	A	P-O5'	-5.39	1.54	1.59
85	AA	359	A	O3'-P	-5.39	1.54	1.61
85	AA	1236	G	C2-N2	-5.39	1.29	1.34
85	AA	2013	A	O3'-P	-5.39	1.54	1.61
85	AA	2238	C	O3'-P	-5.39	1.54	1.61
34	BA	900	A	C4'-C3'	-5.39	1.47	1.52
35	BB	1055	G	C1'-N9	-5.39	1.39	1.46
35	BB	1418	C	C4-N4	-5.39	1.29	1.33
64	Be	83	PHE	CB-CG	-5.39	1.42	1.51
85	AA	84	C	C2'-C1'	-5.39	1.47	1.53
34	BA	167	U	N3-C4	-5.39	1.33	1.38
34	BA	328	A	P-O5'	-5.39	1.54	1.59
34	BA	894	G	C5-C6	-5.39	1.36	1.42
34	BA	1071	G	C5-C4	-5.39	1.34	1.38
34	BA	1709	A	C6-N6	-5.39	1.29	1.33
35	BB	804	U	O4'-C1'	-5.39	1.34	1.41
35	BB	1103	A	C5-C6	-5.39	1.36	1.41
35	BB	1422	G	C4'-C3'	-5.39	1.47	1.52
36	BC	49	G	C5-C4	-5.39	1.34	1.38
37	BD	86	A	C5-C4	-5.39	1.34	1.38
38	BE	93	U	C4-C5	-5.39	1.38	1.43
85	AA	11	A	C5-C4	-5.39	1.34	1.38
85	AA	767	A	C4'-C3'	5.39	1.59	1.53
85	AA	1232	U	C2-N3	-5.39	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1242	A	C3'-C2'	-5.39	1.46	1.52
85	AA	1528	A	C3'-C2'	-5.39	1.46	1.52
34	BA	1437	G	C4'-C3'	-5.39	1.47	1.52
34	BA	1508	C	O3'-P	-5.39	1.54	1.61
35	BB	441	G	C4'-O4'	-5.39	1.38	1.45
35	BB	1234	G	C4'-C3'	-5.39	1.47	1.52
35	BB	1538	G	N9-C8	-5.39	1.34	1.37
36	BC	147	G	C5-C4	-5.39	1.34	1.38
38	BE	65	U	O3'-P	-5.39	1.54	1.61
85	AA	38	C	C3'-C2'	-5.39	1.46	1.52
85	AA	488	G	N9-C4	5.39	1.42	1.38
85	AA	858	G	P-O5'	-5.39	1.54	1.59
85	AA	1531	G	C2'-C1'	-5.39	1.47	1.53
34	BA	84	U	C5'-C4'	-5.39	1.44	1.51
34	BA	178	C	P-O5'	-5.39	1.54	1.59
34	BA	216	C	C2-N3	-5.39	1.31	1.35
34	BA	520	G	P-O5'	-5.39	1.54	1.59
34	BA	538	G	N7-C5	-5.39	1.36	1.39
34	BA	621	G	O3'-P	-5.39	1.54	1.61
34	BA	883	C	P-O5'	-5.39	1.54	1.59
34	BA	1322	A	P-O5'	-5.39	1.54	1.59
35	BB	53	C	C1'-N1	-5.39	1.39	1.46
35	BB	104	G	C2-N2	-5.39	1.29	1.34
35	BB	109	U	C3'-C2'	-5.39	1.46	1.52
35	BB	789	G	C6-N1	-5.39	1.35	1.39
36	BC	13	U	C2-N3	-5.39	1.33	1.37
37	BD	74	A	C8-N7	-5.39	1.27	1.31
40	BG	119	A	N9-C4	-5.39	1.34	1.37
41	BH	75	G	O3'-P	-5.39	1.54	1.61
85	AA	694	A	C4'-C3'	-5.39	1.47	1.52
85	AA	869	A	O4'-C1'	-5.39	1.34	1.41
85	AA	1058	G	P-O5'	-5.39	1.54	1.59
85	AA	1195	U	C3'-C2'	-5.39	1.46	1.52
85	AA	1370	G	N7-C5	-5.39	1.36	1.39
85	AA	1486	G	C2-N2	-5.39	1.29	1.34
85	AA	2134	U	C2-N3	-5.39	1.33	1.37
34	BA	412	G	C6-N1	-5.38	1.35	1.39
34	BA	478	G	C5'-C4'	5.38	1.57	1.51
34	BA	821	G	C3'-O3'	5.38	1.49	1.42
34	BA	861	C	C2'-C1'	-5.38	1.47	1.53
34	BA	1177	C	C5'-C4'	-5.38	1.44	1.51
34	BA	1327	G	N3-C4	-5.38	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1648	G	C4'-C3'	-5.38	1.47	1.52
34	BA	1691	G	O4'-C1'	-5.38	1.34	1.41
34	BA	1825	U	C4'-O4'	-5.38	1.38	1.45
35	BB	19	C	C1'-N1	-5.38	1.39	1.46
35	BB	50	A	C5-C4	-5.38	1.34	1.38
35	BB	751	A	C2'-C1'	-5.38	1.47	1.53
35	BB	1220	A	C3'-O3'	-5.38	1.34	1.42
37	BD	100	A	C2'-C1'	-5.38	1.47	1.53
37	BD	104	C	P-O5'	-5.38	1.54	1.59
38	BE	51	C	C4-N4	-5.38	1.29	1.33
39	BF	42	G	N1-C2	-5.38	1.33	1.37
40	BG	76	C	C3'-C2'	-5.38	1.46	1.52
41	BH	92	A	N3-C4	5.38	1.38	1.34
85	AA	2	A	C5'-C4'	5.38	1.57	1.51
85	AA	50	C	C2-N3	-5.38	1.31	1.35
85	AA	318	A	P-O5'	-5.38	1.54	1.59
85	AA	598	C	C5'-C4'	5.38	1.57	1.51
85	AA	690	G	C3'-C2'	-5.38	1.46	1.52
85	AA	1288	A	C3'-C2'	-5.38	1.46	1.52
85	AA	1378	U	O3'-P	-5.38	1.54	1.61
85	AA	1951	U	C2'-C1'	-5.38	1.47	1.53
34	BA	757	G	C3'-O3'	-5.38	1.34	1.42
35	BB	666	A	N7-C5	-5.38	1.36	1.39
38	BE	121	G	C3'-C2'	-5.38	1.46	1.52
85	AA	393	C	C4'-O4'	-5.38	1.38	1.45
85	AA	802	A	C2'-C1'	-5.38	1.47	1.53
34	BA	166	G	C1'-N9	-5.38	1.39	1.46
34	BA	366	G	C4'-O4'	-5.38	1.38	1.45
34	BA	702	G	C5'-C4'	5.38	1.57	1.51
34	BA	956	G	N3-C4	-5.38	1.31	1.35
34	BA	992	A	C3'-C2'	-5.38	1.46	1.52
34	BA	1261	G	C2-N2	-5.38	1.29	1.34
34	BA	1516	G	N7-C5	-5.38	1.36	1.39
35	BB	132	G	N9-C4	-5.38	1.33	1.38
35	BB	539	G	C2-N3	-5.38	1.28	1.32
35	BB	708	C	C2'-C1'	-5.38	1.47	1.53
35	BB	1089	A	C3'-C2'	-5.38	1.46	1.52
35	BB	1405	G	C6-N1	-5.38	1.35	1.39
37	BD	5	A	P-O5'	-5.38	1.54	1.59
38	BE	180	G	N7-C5	-5.38	1.36	1.39
85	AA	53	G	N1-C2	-5.38	1.33	1.37
85	AA	66	U	C5'-C4'	5.38	1.57	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	181	A	C4'-C3'	-5.38	1.47	1.52
85	AA	585	G	O3'-P	-5.38	1.54	1.61
85	AA	1522	U	O3'-P	-5.38	1.54	1.61
85	AA	1665	G	C2-N2	-5.38	1.29	1.34
85	AA	2089	G	C3'-C2'	-5.38	1.46	1.52
34	BA	758	G	C2'-C1'	-5.38	1.47	1.53
34	BA	1170	A	C5-C4	-5.38	1.34	1.38
35	BB	568	A	N3-C4	-5.38	1.31	1.34
35	BB	1158	C	C2-N3	-5.38	1.31	1.35
36	BC	67	U	C2-N3	-5.38	1.33	1.37
39	BF	2	G	P-O5'	-5.38	1.54	1.59
85	AA	335	G	C6-N1	-5.38	1.35	1.39
85	AA	1116	G	C5-C4	-5.38	1.34	1.38
34	BA	364	C	P-O5'	-5.38	1.54	1.59
34	BA	767	U	C2'-C1'	-5.38	1.47	1.53
34	BA	892	C	C4'-C3'	-5.38	1.47	1.52
34	BA	1201	G	C1'-N9	-5.38	1.39	1.46
35	BB	13	A	C3'-C2'	-5.38	1.46	1.52
35	BB	1394	A	O3'-P	-5.38	1.54	1.61
35	BB	1403	G	C2'-C1'	-5.38	1.47	1.53
85	AA	457	G	O4'-C1'	-5.38	1.34	1.41
85	AA	756	G	C2'-C1'	-5.38	1.47	1.53
85	AA	877	G	C3'-C2'	-5.38	1.46	1.52
85	AA	1009	G	C2-N2	-5.38	1.29	1.34
85	AA	1484	G	N9-C8	-5.38	1.34	1.37
34	BA	17	A	N7-C5	-5.38	1.36	1.39
34	BA	55	G	P-O5'	-5.38	1.54	1.59
34	BA	912	G	C3'-C2'	-5.38	1.46	1.52
34	BA	1214	U	C2'-C1'	-5.38	1.47	1.53
34	BA	1307	U	C1'-N1	5.38	1.56	1.48
34	BA	1486	U	N1-C2	5.38	1.43	1.38
34	BA	1577	U	O3'-P	-5.38	1.54	1.61
34	BA	1596	C	C1'-N1	-5.38	1.39	1.46
35	BB	1154	C	C2-N3	-5.38	1.31	1.35
38	BE	124	G	N3-C4	-5.38	1.31	1.35
40	BG	81	G	C2-N2	-5.38	1.29	1.34
40	BG	89	A	C6-N6	-5.38	1.29	1.33
85	AA	880	A	N7-C5	-5.38	1.36	1.39
85	AA	1104	G	C4'-C3'	-5.38	1.47	1.52
85	AA	1441	G	N3-C4	-5.38	1.31	1.35
85	AA	1457	C	N1-C6	5.38	1.40	1.37
85	AA	1489	G	N9-C4	-5.38	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1501	A	C3'-C2'	-5.38	1.46	1.52
85	AA	1540	A	C8-N7	-5.38	1.27	1.31
34	BA	1098	G	C3'-C2'	-5.38	1.46	1.52
34	BA	1321	A	C6-N1	5.38	1.39	1.35
34	BA	1490	U	C3'-O3'	5.38	1.49	1.42
35	BB	474	G	C2-N2	-5.38	1.29	1.34
35	BB	550	G	C2'-C1'	-5.38	1.47	1.53
35	BB	669	A	P-O5'	-5.38	1.54	1.59
35	BB	993	A	O3'-P	-5.38	1.54	1.61
35	BB	1198	C	C2'-C1'	-5.38	1.47	1.53
41	BH	48	G	N7-C5	-5.38	1.36	1.39
85	AA	779	G	O3'-P	-5.38	1.54	1.61
85	AA	822	U	C3'-C2'	-5.38	1.46	1.52
85	AA	1230	U	C2-N3	-5.38	1.33	1.37
34	BA	135	G	C5-C6	-5.37	1.36	1.42
34	BA	320	G	C2-N2	-5.37	1.29	1.34
34	BA	453	A	C5-C4	-5.37	1.34	1.38
34	BA	1516	G	N3-C4	-5.37	1.31	1.35
35	BB	41	A	N9-C4	-5.37	1.34	1.37
35	BB	997	G	C6-N1	-5.37	1.35	1.39
36	BC	8	C	C2-N3	-5.37	1.31	1.35
38	BE	98	C	P-O5'	-5.37	1.54	1.59
40	BG	50	G	N3-C4	-5.37	1.31	1.35
41	BH	19	G	C6-N1	-5.37	1.35	1.39
67	Bh	64	ARG	CA-C	-5.37	1.39	1.52
85	AA	312	G	C1'-N9	-5.37	1.39	1.46
85	AA	524	A	N9-C4	-5.37	1.34	1.37
85	AA	1490	A	O4'-C1'	-5.37	1.34	1.41
34	BA	198	U	C2-N3	-5.37	1.33	1.37
34	BA	276	C	C1'-N1	-5.37	1.39	1.46
34	BA	448	U	C3'-C2'	-5.37	1.46	1.52
34	BA	486	G	O4'-C1'	-5.37	1.34	1.41
34	BA	609	G	C4'-C3'	-5.37	1.47	1.52
34	BA	795	G	N3-C4	-5.37	1.31	1.35
34	BA	908	G	N1-C2	-5.37	1.33	1.37
34	BA	1613	G	C2-N2	-5.37	1.29	1.34
35	BB	609	G	C5-C4	-5.37	1.34	1.38
35	BB	956	G	O3'-P	-5.37	1.54	1.61
35	BB	1075	A	C5-C4	-5.37	1.34	1.38
35	BB	1142	C	P-O5'	-5.37	1.54	1.59
35	BB	1409	G	C5-C4	-5.37	1.34	1.38
36	BC	139	A	O3'-P	-5.37	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	44	G	N1-C2	-5.37	1.33	1.37
40	BG	58	G	C5'-C4'	-5.37	1.45	1.51
41	BH	64	U	P-O5'	-5.37	1.54	1.59
85	AA	1454	U	N1-C6	-5.37	1.33	1.38
34	BA	415	C	C2-N3	-5.37	1.31	1.35
34	BA	1097	G	C2-N2	-5.37	1.29	1.34
34	BA	1303	U	C2-N3	-5.37	1.33	1.37
34	BA	1327	G	C2-N2	-5.37	1.29	1.34
34	BA	1457	C	C1'-N1	-5.37	1.39	1.46
35	BB	742	G	C2'-C1'	-5.37	1.47	1.53
35	BB	1261	U	C2'-C1'	-5.37	1.47	1.53
36	BC	91	G	N1-C2	-5.37	1.33	1.37
85	AA	160	A	C6-N1	-5.37	1.31	1.35
34	BA	525	A	C3'-C2'	-5.37	1.46	1.52
34	BA	873	G	C2'-C1'	-5.37	1.47	1.53
34	BA	1181	G	C2'-C1'	-5.37	1.47	1.53
35	BB	425	G	N3-C4	-5.37	1.31	1.35
35	BB	845	C	C3'-C2'	-5.37	1.46	1.52
35	BB	1059	U	C2-N3	-5.37	1.33	1.37
35	BB	1365	G	C2-N2	-5.37	1.29	1.34
35	BB	1374	U	N1-C6	-5.37	1.33	1.38
35	BB	1424	G	C2-N3	-5.37	1.28	1.32
40	BG	16	G	N1-C2	-5.37	1.33	1.37
40	BG	149	U	N1-C6	-5.37	1.33	1.38
48	BO	35	GLY	CA-C	-5.37	1.43	1.51
74	Bo	79	VAL	CA-CB	-5.37	1.43	1.54
85	AA	779	G	N9-C8	-5.37	1.34	1.37
85	AA	934	A	N7-C5	-5.37	1.36	1.39
85	AA	1842	C	P-O5'	-5.37	1.54	1.59
34	BA	46	C	P-O5'	-5.37	1.54	1.59
34	BA	86	A	N3-C4	-5.37	1.31	1.34
34	BA	1213	A	N9-C8	-5.37	1.33	1.37
41	BH	72	G	O4'-C1'	-5.37	1.34	1.41
85	AA	66	U	C4'-O4'	-5.37	1.38	1.45
85	AA	439	U	C1'-N1	-5.37	1.39	1.46
85	AA	1447	U	P-O5'	-5.37	1.54	1.59
85	AA	1760	C	P-O5'	-5.37	1.54	1.59
85	AA	2128	G	O4'-C1'	-5.37	1.34	1.41
34	BA	162	G	N7-C5	-5.37	1.36	1.39
34	BA	363	G	C5-C6	-5.37	1.36	1.42
34	BA	455	A	C5-C4	-5.37	1.34	1.38
34	BA	1046	G	N1-C2	-5.37	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1166	A	C4'-C3'	-5.37	1.47	1.52
34	BA	1580	U	C4'-O4'	-5.37	1.38	1.45
35	BB	16	G	O4'-C1'	-5.37	1.34	1.41
35	BB	1087	A	C5-C4	-5.37	1.34	1.38
38	BE	50	G	P-O5'	-5.37	1.54	1.59
41	BH	18	C	C4'-C3'	-5.37	1.47	1.52
85	AA	189	G	N9-C8	-5.37	1.34	1.37
85	AA	250	C	O3'-P	-5.37	1.54	1.61
85	AA	267	U	P-O5'	-5.37	1.54	1.59
85	AA	712	U	O3'-P	-5.37	1.54	1.61
85	AA	1507	G	P-O5'	-5.37	1.54	1.59
85	AA	2014	G	P-O5'	-5.37	1.54	1.59
34	BA	26	C	C4-N4	-5.36	1.29	1.33
34	BA	292	C	C2-N3	-5.36	1.31	1.35
34	BA	754	G	C6-N1	-5.36	1.35	1.39
34	BA	919	A	N3-C4	-5.36	1.31	1.34
34	BA	1839	G	C4'-C3'	-5.36	1.47	1.52
35	BB	739	C	P-O5'	-5.36	1.54	1.59
36	BC	40	A	C3'-C2'	-5.36	1.46	1.52
36	BC	104	A	C5'-C4'	-5.36	1.45	1.51
38	BE	146	U	O3'-P	-5.36	1.54	1.61
85	AA	411	U	O4'-C1'	-5.36	1.34	1.41
85	AA	651	G	C3'-C2'	-5.36	1.46	1.52
85	AA	883	A	C5-C4	-5.36	1.34	1.38
85	AA	1019	U	O3'-P	-5.36	1.54	1.61
85	AA	1295	G	O3'-P	-5.36	1.54	1.61
85	AA	1456	A	C4'-O4'	5.36	1.52	1.45
85	AA	1935	G	N7-C5	-5.36	1.36	1.39
34	BA	1249	G	C2-N2	-5.36	1.29	1.34
34	BA	1567	G	C2-N2	-5.36	1.29	1.34
35	BB	49	A	C5-C4	-5.36	1.34	1.38
35	BB	840	C	C2'-C1'	-5.36	1.47	1.53
35	BB	1299	G	C2'-C1'	-5.36	1.47	1.53
35	BB	1351	G	C2'-C1'	-5.36	1.47	1.53
35	BB	1419	G	C2-N2	-5.36	1.29	1.34
61	Bb	71	PRO	N-CA	-5.36	1.38	1.47
85	AA	820	G	C6-N1	-5.36	1.35	1.39
85	AA	900	G	N1-C2	-5.36	1.33	1.37
85	AA	1512	U	C2'-C1'	-5.36	1.47	1.53
85	AA	1708	A	O4'-C1'	-5.36	1.34	1.41
27	AT	117	ARG	CD-NE	5.36	1.55	1.46
34	BA	359	G	C2'-C1'	-5.36	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	933	U	C3'-C2'	-5.36	1.46	1.52
34	BA	986	G	O3'-P	-5.36	1.54	1.61
34	BA	1363	A	C1'-N9	-5.36	1.39	1.46
34	BA	1428	G	C5-C6	-5.36	1.36	1.42
34	BA	1439	C	C4'-C3'	-5.36	1.47	1.52
34	BA	1546	C	O3'-P	-5.36	1.54	1.61
35	BB	664	A	O3'-P	-5.36	1.54	1.61
85	AA	506	G	P-O5'	-5.36	1.54	1.59
85	AA	1508	A	P-O5'	-5.36	1.54	1.59
85	AA	1727	U	C4'-O4'	-5.36	1.38	1.45
85	AA	2041	G	N9-C8	-5.36	1.34	1.37
34	BA	1163	G	C3'-C2'	-5.36	1.46	1.52
34	BA	1448	G	O3'-P	-5.36	1.54	1.61
34	BA	1565	U	C5'-C4'	5.36	1.57	1.51
34	BA	1706	A	C5-C4	-5.36	1.34	1.38
40	BG	117	C	N1-C6	-5.36	1.33	1.37
40	BG	172	C	O4'-C1'	-5.36	1.34	1.41
41	BH	106	G	N9-C4	-5.36	1.33	1.38
34	BA	77	C	C4-N4	-5.36	1.29	1.33
34	BA	146	G	C5-C4	-5.36	1.34	1.38
34	BA	292	C	N1-C6	-5.36	1.33	1.37
34	BA	331	G	N1-C2	-5.36	1.33	1.37
34	BA	696	A	N9-C8	-5.36	1.33	1.37
34	BA	823	G	C6-N1	-5.36	1.35	1.39
34	BA	852	C	C4-N4	-5.36	1.29	1.33
34	BA	1729	G	N7-C5	-5.36	1.36	1.39
35	BB	115	A	C1'-N9	-5.36	1.39	1.46
35	BB	134	G	P-O5'	-5.36	1.54	1.59
35	BB	366	G	C1'-N9	-5.36	1.39	1.46
35	BB	428	G	C4'-C3'	-5.36	1.47	1.52
35	BB	586	U	C1'-N1	-5.36	1.39	1.46
35	BB	1414	A	C5-C4	-5.36	1.35	1.38
35	BB	1504	U	C1'-N1	-5.36	1.39	1.46
36	BC	160	C	C5'-C4'	-5.36	1.45	1.51
85	AA	574	U	O3'-P	-5.36	1.54	1.61
85	AA	688	C	C1'-N1	-5.36	1.39	1.46
85	AA	1244	A	C1'-N9	-5.36	1.39	1.46
85	AA	1608	U	P-O5'	-5.36	1.54	1.59
85	AA	1700	C	C1'-N1	-5.36	1.39	1.46
85	AA	2190	U	O3'-P	-5.36	1.54	1.61
34	BA	262	A	C6-N6	-5.36	1.29	1.33
34	BA	359	G	O3'-P	-5.36	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	726	G	C3'-C2'	-5.36	1.46	1.52
34	BA	1253	G	N9-C8	-5.36	1.34	1.37
34	BA	1354	G	C1'-N9	-5.36	1.39	1.46
34	BA	1709	A	C2'-C1'	-5.36	1.47	1.53
35	BB	370	A	C4'-O4'	-5.36	1.38	1.45
35	BB	555	G	N9-C4	-5.36	1.33	1.38
35	BB	702	G	C6-N1	-5.36	1.35	1.39
35	BB	1342	C	O4'-C1'	-5.36	1.34	1.41
38	BE	26	G	C8-N7	-5.36	1.27	1.30
38	BE	61	A	C1'-N9	-5.36	1.39	1.46
41	BH	74	G	C2-N3	5.36	1.37	1.32
85	AA	243	A	C6-N6	-5.36	1.29	1.33
85	AA	1506	U	C2-N3	-5.36	1.34	1.37
85	AA	2187	G	C5'-C4'	-5.36	1.45	1.51
34	BA	1264	U	C3'-C2'	-5.35	1.46	1.52
34	BA	1405	A	N3-C4	-5.35	1.31	1.34
34	BA	1435	A	C8-N7	-5.35	1.27	1.31
35	BB	520	G	C5-C4	-5.35	1.34	1.38
35	BB	896	C	C4-N4	-5.35	1.29	1.33
36	BC	56	G	C5-C4	-5.35	1.34	1.38
37	BD	86	A	C1'-N9	-5.35	1.39	1.46
56	BW	109	GLY	CA-C	-5.35	1.43	1.51
85	AA	1515	A	C4'-O4'	-5.35	1.38	1.45
34	BA	143	A	O3'-P	-5.35	1.54	1.61
34	BA	774	A	N3-C4	-5.35	1.31	1.34
34	BA	845	U	P-O5'	-5.35	1.54	1.59
35	BB	590	G	C2'-C1'	-5.35	1.47	1.53
35	BB	657	A	C5'-C4'	-5.35	1.45	1.51
35	BB	1308	G	N3-C4	-5.35	1.31	1.35
38	BE	99	C	C4'-C3'	-5.35	1.47	1.52
41	BH	3	U	P-O5'	-5.35	1.54	1.59
41	BH	23	G	C4'-O4'	-5.35	1.38	1.45
41	BH	74	G	C5'-C4'	5.35	1.57	1.51
85	AA	879	G	N3-C4	-5.35	1.31	1.35
85	AA	882	C	P-O5'	-5.35	1.54	1.59
85	AA	1009	G	C2'-C1'	-5.35	1.47	1.53
85	AA	1196	C	C4'-C3'	-5.35	1.47	1.52
85	AA	1822	G	P-O5'	-5.35	1.54	1.59
85	AA	1864	G	C5-C4	-5.35	1.34	1.38
34	BA	540	G	N7-C5	-5.35	1.36	1.39
34	BA	1741	G	C1'-N9	-5.35	1.39	1.46
35	BB	515	C	C4-N4	-5.35	1.29	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	805	G	C2-N2	-5.35	1.29	1.34
37	BD	30	A	C1'-N9	-5.35	1.39	1.46
40	BG	174	G	C1'-N9	-5.35	1.39	1.46
85	AA	162	A	C8-N7	-5.35	1.27	1.31
85	AA	995	G	N9-C8	-5.35	1.34	1.37
85	AA	1288	A	N9-C4	-5.35	1.34	1.37
34	BA	73	G	N9-C8	-5.35	1.34	1.37
34	BA	364	C	C4'-O4'	-5.35	1.38	1.45
34	BA	776	U	C4'-O4'	-5.35	1.38	1.45
34	BA	850	C	O3'-P	-5.35	1.54	1.61
34	BA	977	G	C8-N7	-5.35	1.27	1.30
34	BA	1028	A	C8-N7	-5.35	1.27	1.31
34	BA	1522	G	N9-C8	-5.35	1.34	1.37
35	BB	28	G	N9-C8	-5.35	1.34	1.37
35	BB	594	U	C1'-N1	-5.35	1.39	1.46
35	BB	879	G	C5'-C4'	5.35	1.57	1.51
35	BB	1003	G	C6-N1	-5.35	1.35	1.39
35	BB	1210	U	N1-C2	-5.35	1.33	1.38
35	BB	1314	G	N7-C5	-5.35	1.36	1.39
35	BB	1435	G	C8-N7	-5.35	1.27	1.30
35	BB	1449	G	C2-N2	-5.35	1.29	1.34
36	BC	59	A	N3-C4	-5.35	1.31	1.34
36	BC	121	G	C1'-N9	-5.35	1.39	1.46
37	BD	60	C	N1-C6	-5.35	1.33	1.37
38	BE	202	C	N3-C4	-5.35	1.30	1.33
39	BF	27	G	O3'-P	-5.35	1.54	1.61
83	Bx	50	GLY	CA-C	5.35	1.60	1.51
85	AA	48	G	P-O5'	-5.35	1.54	1.59
85	AA	109	G	C2-N2	-5.35	1.29	1.34
85	AA	208	U	P-O5'	-5.35	1.54	1.59
85	AA	575	G	C5-C4	-5.35	1.34	1.38
85	AA	676	U	C2-N3	-5.35	1.34	1.37
85	AA	1227	A	C2'-C1'	-5.35	1.47	1.53
34	BA	124	G	C4'-C3'	-5.35	1.47	1.52
34	BA	242	U	P-O5'	-5.35	1.54	1.59
34	BA	378	C	C2-N3	-5.35	1.31	1.35
34	BA	1007	G	P-O5'	-5.35	1.54	1.59
34	BA	1065	U	C4'-C3'	-5.35	1.47	1.52
34	BA	1151	A	C2'-C1'	-5.35	1.47	1.53
34	BA	1273	U	O4'-C1'	-5.35	1.34	1.41
35	BB	973	G	O3'-P	-5.35	1.54	1.61
35	BB	1123	A	O4'-C1'	-5.35	1.34	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1231	U	C5'-C4'	5.35	1.57	1.51
36	BC	60	U	C4-C5	-5.35	1.38	1.43
37	BD	43	U	C2-N3	-5.35	1.34	1.37
39	BF	55	A	C4'-C3'	5.35	1.59	1.53
41	BH	17	A	N3-C4	-5.35	1.31	1.34
85	AA	51	A	N9-C4	-5.35	1.34	1.37
85	AA	86	G	C2'-C1'	-5.35	1.47	1.53
85	AA	446	C	C2'-C1'	-5.35	1.47	1.53
85	AA	471	U	N3-C4	-5.35	1.33	1.38
85	AA	939	A	C5-C6	-5.35	1.36	1.41
85	AA	1002	G	O3'-P	-5.35	1.54	1.61
85	AA	1252	A	N7-C5	-5.35	1.36	1.39
34	BA	1677	C	C4-N4	-5.35	1.29	1.33
35	BB	523	A	O3'-P	-5.35	1.54	1.61
35	BB	859	U	C2'-C1'	-5.35	1.47	1.53
35	BB	1288	G	C2-N2	-5.35	1.29	1.34
35	BB	1351	G	C2-N2	-5.35	1.29	1.34
36	BC	28	C	C2-O2	-5.35	1.19	1.24
37	BD	18	G	C3'-C2'	-5.35	1.46	1.52
85	AA	118	C	C4-N4	-5.35	1.29	1.33
85	AA	2092	A	C1'-N9	-5.35	1.39	1.46
85	AA	2208	G	P-O5'	-5.35	1.54	1.59
34	BA	235	C	C4-C5	-5.34	1.38	1.43
34	BA	726	G	C5-C6	-5.34	1.37	1.42
34	BA	1160	U	C3'-C2'	-5.34	1.46	1.52
34	BA	1520	A	C5-C4	-5.34	1.35	1.38
35	BB	494	C	P-O5'	-5.34	1.54	1.59
35	BB	1066	G	N9-C8	-5.34	1.34	1.37
35	BB	1104	A	C1'-N9	-5.34	1.39	1.46
35	BB	1154	C	C1'-N1	-5.34	1.39	1.46
35	BB	1249	G	N9-C8	-5.34	1.34	1.37
35	BB	1275	A	C2'-C1'	-5.34	1.47	1.53
37	BD	19	C	O3'-P	-5.34	1.54	1.61
37	BD	79	G	C2-N3	-5.34	1.28	1.32
38	BE	9	C	C3'-C2'	-5.34	1.46	1.52
38	BE	159	A	O3'-P	-5.34	1.54	1.61
38	BE	164	C	C2-N3	-5.34	1.31	1.35
40	BG	66	C	C4'-C3'	-5.34	1.47	1.52
85	AA	10	G	N9-C4	-5.34	1.33	1.38
85	AA	300	C	O3'-P	-5.34	1.54	1.61
85	AA	1034	U	C2'-C1'	-5.34	1.47	1.53
85	AA	1151	G	C1'-N9	-5.34	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	2077	G	C3'-O3'	5.34	1.49	1.42
85	AA	2135	A	C5-C4	-5.34	1.35	1.38
13	AE	153	TYR	CB-CG	-5.34	1.43	1.51
34	BA	211	C	C2'-C1'	-5.34	1.47	1.53
34	BA	1198	U	O3'-P	-5.34	1.54	1.61
34	BA	1773	U	P-O5'	-5.34	1.54	1.59
35	BB	567	G	N9-C8	-5.34	1.34	1.37
41	BH	3	U	C2-N3	-5.34	1.34	1.37
41	BH	118	U	C4'-C3'	-5.34	1.47	1.52
85	AA	1566	A	C1'-N9	-5.34	1.39	1.46
86	AB	15	G	N1-C2	-5.34	1.33	1.37
34	BA	331	G	N9-C4	-5.34	1.33	1.38
34	BA	719	G	N9-C4	-5.34	1.33	1.38
34	BA	824	C	C3'-C2'	-5.34	1.46	1.52
34	BA	1139	G	C6-N1	-5.34	1.35	1.39
34	BA	1230	G	N9-C8	-5.34	1.34	1.37
34	BA	1333	G	C8-N7	-5.34	1.27	1.30
34	BA	1551	G	C2-N2	-5.34	1.29	1.34
34	BA	1639	U	C3'-C2'	-5.34	1.46	1.52
38	BE	52	U	C4'-O4'	-5.34	1.38	1.45
85	AA	1246	G	N9-C4	5.34	1.42	1.38
85	AA	1490	A	C8-N7	-5.34	1.27	1.31
85	AA	1507	G	C6-N1	-5.34	1.35	1.39
85	AA	2185	U	C1'-N1	-5.34	1.39	1.46
34	BA	542	A	C5-C4	-5.34	1.35	1.38
34	BA	593	G	C2-N2	-5.34	1.29	1.34
34	BA	829	U	C3'-C2'	-5.34	1.46	1.52
34	BA	1034	U	N1-C2	-5.34	1.33	1.38
34	BA	1159	A	C3'-C2'	-5.34	1.46	1.52
34	BA	1228	G	C2'-C1'	-5.34	1.47	1.53
34	BA	1522	G	C2-N2	-5.34	1.29	1.34
34	BA	1651	C	N1-C6	-5.34	1.33	1.37
35	BB	53	C	C3'-C2'	-5.34	1.46	1.52
35	BB	702	G	C8-N7	-5.34	1.27	1.30
35	BB	1087	A	N9-C8	-5.34	1.33	1.37
35	BB	1108	G	N9-C8	-5.34	1.34	1.37
35	BB	1207	C	N1-C2	-5.34	1.34	1.40
35	BB	1218	G	C2-N2	-5.34	1.29	1.34
36	BC	21	U	C2'-C1'	-5.34	1.47	1.53
36	BC	22	U	P-O5'	-5.34	1.54	1.59
38	BE	43	A	N7-C5	-5.34	1.36	1.39
38	BE	185	G	N7-C5	-5.34	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	18	U	C1'-N1	-5.34	1.39	1.46
40	BG	66	C	P-O5'	-5.34	1.54	1.59
85	AA	578	U	P-O5'	-5.34	1.54	1.59
85	AA	1470	A	N3-C4	-5.34	1.31	1.34
85	AA	2101	C	P-O5'	-5.34	1.54	1.59
85	AA	2146	G	N7-C5	-5.34	1.36	1.39
34	BA	262	A	C5-C4	-5.34	1.35	1.38
34	BA	687	G	C4'-C3'	-5.34	1.47	1.52
35	BB	412	A	C5-C6	-5.34	1.36	1.41
35	BB	517	G	N1-C2	-5.34	1.33	1.37
35	BB	770	G	P-O5'	-5.34	1.54	1.59
38	BE	193	A	C2'-C1'	-5.34	1.47	1.53
40	BG	106	G	N7-C5	-5.34	1.36	1.39
41	BH	45	G	C5-C4	-5.34	1.34	1.38
85	AA	132	G	C2'-C1'	-5.34	1.47	1.53
85	AA	372	U	C4'-C3'	-5.34	1.47	1.52
85	AA	1001	G	C6-N1	-5.34	1.35	1.39
85	AA	1242	A	O3'-P	-5.34	1.54	1.61
34	BA	251	U	P-O5'	-5.34	1.54	1.59
34	BA	719	G	O3'-P	-5.34	1.54	1.61
34	BA	1447	C	O4'-C1'	-5.34	1.34	1.41
35	BB	477	U	P-O5'	-5.34	1.54	1.59
37	BD	58	G	P-O5'	-5.34	1.54	1.59
37	BD	65	G	C3'-C2'	-5.34	1.46	1.52
34	BA	425	G	C6-N1	-5.33	1.35	1.39
34	BA	1001	G	N3-C4	-5.33	1.31	1.35
37	BD	13	A	C2'-C1'	-5.33	1.47	1.53
85	AA	494	G	C2'-C1'	-5.33	1.47	1.53
85	AA	558	U	O3'-P	-5.33	1.54	1.61
85	AA	1111	A	O3'-P	-5.33	1.54	1.61
85	AA	1254	A	N7-C5	-5.33	1.36	1.39
34	BA	164	C	O3'-P	-5.33	1.54	1.61
34	BA	329	G	P-O5'	-5.33	1.54	1.59
34	BA	1100	A	C5-C4	-5.33	1.35	1.38
35	BB	584	A	C4'-C3'	-5.33	1.47	1.52
35	BB	1299	G	N9-C8	-5.33	1.34	1.37
35	BB	1498	G	C5-C4	-5.33	1.34	1.38
36	BC	57	C	C3'-C2'	-5.33	1.46	1.52
36	BC	142	C	C2-N3	-5.33	1.31	1.35
37	BD	108	G	C8-N7	-5.33	1.27	1.30
38	BE	42	C	P-O5'	-5.33	1.54	1.59
53	BT	100	ARG	N-CA	-5.33	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	40	A	N9-C4	-5.33	1.34	1.37
85	AA	79	G	N9-C8	-5.33	1.34	1.37
85	AA	305	A	C2'-C1'	-5.33	1.47	1.53
85	AA	990	U	O3'-P	-5.33	1.54	1.61
85	AA	1127	G	P-O5'	-5.33	1.54	1.59
85	AA	1585	A	N7-C5	-5.33	1.36	1.39
85	AA	1611	A	C1'-N9	-5.33	1.39	1.46
85	AA	2178	A	P-O5'	-5.33	1.54	1.59
34	BA	199	U	C1'-N1	-5.33	1.39	1.46
35	BB	12	G	C6-N1	-5.33	1.35	1.39
35	BB	133	G	N9-C8	-5.33	1.34	1.37
35	BB	471	U	O4'-C1'	-5.33	1.34	1.41
35	BB	583	G	C3'-C2'	-5.33	1.46	1.52
35	BB	772	U	C2-N3	-5.33	1.34	1.37
35	BB	795	A	C6-N6	-5.33	1.29	1.33
37	BD	46	G	N3-C4	-5.33	1.31	1.35
38	BE	74	U	O3'-P	-5.33	1.54	1.61
40	BG	81	G	N9-C8	-5.33	1.34	1.37
85	AA	102	A	C8-N7	-5.33	1.27	1.31
85	AA	204	U	C2'-C1'	-5.33	1.47	1.53
85	AA	294	G	C5'-C4'	5.33	1.57	1.51
85	AA	309	G	N9-C4	-5.33	1.33	1.38
85	AA	457	G	O3'-P	-5.33	1.54	1.61
85	AA	584	G	N1-C2	-5.33	1.33	1.37
85	AA	871	U	C2-N3	-5.33	1.34	1.37
85	AA	2082	C	C4-N4	-5.33	1.29	1.33
35	BB	588	A	C1'-N9	-5.33	1.39	1.46
35	BB	1155	U	C4'-C3'	-5.33	1.47	1.52
35	BB	1516	C	C3'-C2'	-5.33	1.46	1.52
85	AA	935	A	O3'-P	-5.33	1.54	1.61
85	AA	1624	U	C2'-C1'	-5.33	1.47	1.53
85	AA	2032	G	C6-N1	-5.33	1.35	1.39
86	AB	63	G	C3'-C2'	-5.33	1.46	1.52
34	BA	1626	U	C3'-C2'	5.33	1.58	1.52
35	BB	866	A	C2'-C1'	-5.33	1.47	1.53
35	BB	1322	A	O4'-C1'	-5.33	1.34	1.41
36	BC	42	G	C2-N3	-5.33	1.28	1.32
37	BD	11	A	C2'-C1'	-5.33	1.47	1.53
40	BG	175	G	N1-C2	-5.33	1.33	1.37
85	AA	27	U	C3'-C2'	-5.33	1.46	1.52
85	AA	386	G	N1-C2	-5.33	1.33	1.37
85	AA	1231	G	N1-C2	-5.33	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1554	C	C1'-N1	-5.33	1.39	1.46
85	AA	2215	C	O3'-P	-5.33	1.54	1.61
34	BA	517	A	C3'-C2'	-5.33	1.46	1.52
34	BA	863	G	C8-N7	5.33	1.34	1.30
34	BA	1276	G	C2-N2	-5.33	1.29	1.34
35	BB	438	G	O4'-C1'	-5.33	1.34	1.41
35	BB	452	A	C3'-O3'	5.33	1.49	1.42
38	BE	198	A	C3'-C2'	-5.33	1.46	1.52
85	AA	453	G	P-O5'	-5.33	1.54	1.59
85	AA	1176	C	N1-C6	-5.33	1.33	1.37
85	AA	1502	A	N9-C4	-5.33	1.34	1.37
34	BA	248	G	N3-C4	-5.33	1.31	1.35
34	BA	261	A	C2'-C1'	-5.33	1.47	1.53
34	BA	288	U	C3'-O3'	5.33	1.49	1.42
34	BA	825	G	C8-N7	-5.33	1.27	1.30
34	BA	990	G	N9-C8	-5.33	1.34	1.37
34	BA	1009	G	N9-C8	-5.33	1.34	1.37
34	BA	1275	G	C5-C4	-5.33	1.34	1.38
34	BA	1343	A	O3'-P	-5.33	1.54	1.61
34	BA	1735	G	O3'-P	-5.33	1.54	1.61
34	BA	1832	A	C4'-C3'	-5.33	1.47	1.52
35	BB	441	G	N9-C8	-5.33	1.34	1.37
35	BB	803	U	C1'-N1	-5.33	1.39	1.46
40	BG	76	C	C4'-C3'	-5.33	1.47	1.52
85	AA	203	C	C2'-C1'	-5.33	1.47	1.53
85	AA	553	G	N7-C5	-5.33	1.36	1.39
85	AA	729	U	P-O5'	-5.33	1.54	1.59
85	AA	781	G	P-O5'	-5.33	1.54	1.59
85	AA	809	A	O3'-P	-5.33	1.54	1.61
85	AA	929	G	C5-C4	-5.33	1.34	1.38
85	AA	2241	C	C2-N3	-5.33	1.31	1.35
34	BA	327	G	C5-C4	-5.32	1.34	1.38
34	BA	880	G	C3'-C2'	-5.32	1.46	1.52
34	BA	1404	A	O3'-P	-5.32	1.54	1.61
34	BA	1407	C	C2-N3	-5.32	1.31	1.35
34	BA	1654	G	N3-C4	-5.32	1.31	1.35
34	BA	1839	G	C2-N2	-5.32	1.29	1.34
35	BB	71	A	C6-N6	-5.32	1.29	1.33
35	BB	315	C	P-O5'	-5.32	1.54	1.59
35	BB	1162	A	C4'-C3'	-5.32	1.47	1.52
36	BC	87	C	C4'-O4'	-5.32	1.38	1.45
38	BE	69	C	C3'-C2'	-5.32	1.46	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	176	G	C2-N2	-5.32	1.29	1.34
38	BE	193	A	C1'-N9	-5.32	1.39	1.46
41	BH	135	U	C2-N3	-5.32	1.34	1.37
85	AA	238	C	P-O5'	-5.32	1.54	1.59
85	AA	315	U	C3'-C2'	-5.32	1.46	1.52
85	AA	424	A	P-O5'	-5.32	1.54	1.59
85	AA	1503	G	O4'-C1'	-5.32	1.34	1.41
85	AA	1872	G	O3'-P	-5.32	1.54	1.61
85	AA	1873	U	O3'-P	-5.32	1.54	1.61
85	AA	2166	G	C2'-C1'	-5.32	1.47	1.53
85	AA	2213	A	C1'-N9	-5.32	1.39	1.46
34	BA	121	A	C8-N7	-5.32	1.27	1.31
34	BA	207	A	C5-C6	-5.32	1.36	1.41
34	BA	230	A	O3'-P	-5.32	1.54	1.61
34	BA	800	G	C5-C4	-5.32	1.34	1.38
34	BA	1469	G	C2-N2	-5.32	1.29	1.34
35	BB	622	G	C3'-C2'	-5.32	1.46	1.52
35	BB	1069	C	O3'-P	-5.32	1.54	1.61
85	AA	371	C	C2'-C1'	-5.32	1.47	1.53
85	AA	2131	C	C4-N4	-5.32	1.29	1.33
34	BA	20	A	N9-C8	-5.32	1.33	1.37
34	BA	141	G	C3'-C2'	-5.32	1.46	1.52
34	BA	474	A	P-O5'	-5.32	1.54	1.59
34	BA	516	U	C5'-C4'	5.32	1.57	1.51
34	BA	1067	G	O3'-P	-5.32	1.54	1.61
34	BA	1226	G	N1-C2	-5.32	1.33	1.37
34	BA	1229	G	C2-N3	-5.32	1.28	1.32
35	BB	90	G	C1'-N9	-5.32	1.39	1.46
35	BB	282	A	P-O5'	-5.32	1.54	1.59
35	BB	1031	G	C4'-O4'	-5.32	1.38	1.45
35	BB	1046	C	C2-N3	-5.32	1.31	1.35
35	BB	1166	A	C5-C4	-5.32	1.35	1.38
35	BB	1251	G	C2-N2	-5.32	1.29	1.34
85	AA	687	G	N9-C8	-5.32	1.34	1.37
85	AA	995	G	C3'-C2'	-5.32	1.46	1.52
85	AA	1142	G	O3'-P	-5.32	1.54	1.61
85	AA	1475	A	N7-C5	-5.32	1.36	1.39
85	AA	1529	A	C3'-C2'	-5.32	1.46	1.52
85	AA	1682	U	C3'-O3'	-5.32	1.34	1.42
85	AA	1839	G	C4'-C3'	-5.32	1.47	1.52
85	AA	2125	A	C4'-C3'	-5.32	1.47	1.52
34	BA	321	G	C2-N2	-5.32	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	330	A	P-O5'	-5.32	1.54	1.59
34	BA	1846	G	C6-O6	-5.32	1.19	1.24
40	BG	31	G	N9-C8	-5.32	1.34	1.37
85	AA	60	U	P-O5'	-5.32	1.54	1.59
34	BA	340	U	N1-C6	-5.32	1.33	1.38
34	BA	584	A	C1'-N9	-5.32	1.39	1.46
34	BA	1230	G	C2'-C1'	-5.32	1.47	1.53
34	BA	1406	U	C3'-C2'	-5.32	1.47	1.52
34	BA	1672	C	C2-N3	-5.32	1.31	1.35
35	BB	639	A	N9-C8	-5.32	1.33	1.37
35	BB	662	G	C8-N7	-5.32	1.27	1.30
35	BB	1026	G	N9-C8	-5.32	1.34	1.37
35	BB	1119	G	C6-N1	-5.32	1.35	1.39
36	BC	43	A	N9-C8	-5.32	1.33	1.37
36	BC	59	A	C6-N1	-5.32	1.31	1.35
36	BC	149	A	C6-N1	-5.32	1.31	1.35
37	BD	86	A	C4'-O4'	-5.32	1.38	1.45
38	BE	48	G	C5'-C4'	-5.32	1.45	1.51
85	AA	85	U	N1-C2	-5.32	1.33	1.38
85	AA	141	A	C5-C4	-5.32	1.35	1.38
85	AA	552	C	P-O5'	-5.32	1.54	1.59
85	AA	719	C	O3'-P	-5.32	1.54	1.61
85	AA	1254	A	C5-C4	-5.32	1.35	1.38
85	AA	1757	C	C2'-C1'	-5.32	1.47	1.53
34	BA	596	G	C6-O6	-5.32	1.19	1.24
34	BA	666	C	P-O5'	-5.32	1.54	1.59
34	BA	852	C	C2'-C1'	-5.32	1.47	1.53
34	BA	860	G	C5-C4	-5.32	1.34	1.38
34	BA	902	C	C4'-C3'	-5.32	1.47	1.52
34	BA	973	U	O4'-C1'	-5.32	1.34	1.41
35	BB	490	G	O3'-P	-5.32	1.54	1.61
35	BB	590	G	C2-N2	-5.32	1.29	1.34
35	BB	694	C	P-O5'	-5.32	1.54	1.59
35	BB	901	U	C2'-C1'	-5.32	1.47	1.53
35	BB	1193	G	C4'-O4'	-5.32	1.38	1.45
38	BE	177	U	C1'-N1	-5.32	1.39	1.46
38	BE	199	A	C4'-C3'	-5.32	1.47	1.52
40	BG	22	G	N9-C4	-5.32	1.33	1.38
40	BG	39	A	N7-C5	-5.32	1.36	1.39
40	BG	131	U	C3'-C2'	-5.32	1.47	1.52
85	AA	265	A	C2'-C1'	-5.32	1.47	1.53
85	AA	395	G	C2-N2	-5.32	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	438	G	N7-C5	-5.32	1.36	1.39
85	AA	494	G	C6-N1	-5.32	1.35	1.39
85	AA	852	C	C2'-C1'	-5.32	1.47	1.53
85	AA	1880	C	C4'-C3'	5.32	1.58	1.53
34	BA	216	C	C4-C5	-5.31	1.38	1.43
34	BA	1324	G	N1-C2	-5.31	1.33	1.37
34	BA	1422	A	C2'-C1'	-5.31	1.47	1.53
34	BA	1568	A	N3-C4	-5.31	1.31	1.34
35	BB	1544	A	N7-C5	-5.31	1.36	1.39
41	BH	65	G	P-O5'	-5.31	1.54	1.59
41	BH	117	U	O3'-P	-5.31	1.54	1.61
85	AA	791	C	C4'-O4'	-5.31	1.38	1.45
85	AA	1923	A	C2'-O2'	-5.31	1.34	1.41
34	BA	218	G	O3'-P	-5.31	1.54	1.61
34	BA	248	G	C2-N2	-5.31	1.29	1.34
34	BA	483	A	C5-C4	-5.31	1.35	1.38
34	BA	596	G	C2-N3	-5.31	1.28	1.32
34	BA	1086	A	N9-C8	-5.31	1.33	1.37
35	BB	704	G	C1'-N9	-5.31	1.39	1.46
35	BB	801	G	O3'-P	-5.31	1.54	1.61
35	BB	1448	U	N3-C4	-5.31	1.33	1.38
36	BC	139	A	P-O5'	-5.31	1.54	1.59
37	BD	107	G	N9-C4	-5.31	1.33	1.38
40	BG	89	A	C1'-N9	-5.31	1.39	1.46
41	BH	103	C	O3'-P	-5.31	1.54	1.61
41	BH	130	G	C5-C4	-5.31	1.34	1.38
85	AA	513	G	C4'-C3'	-5.31	1.47	1.52
85	AA	568	C	C2'-C1'	-5.31	1.47	1.53
85	AA	996	A	C2'-C1'	-5.31	1.47	1.53
85	AA	1542	A	C4'-O4'	-5.31	1.38	1.45
85	AA	2062	U	N3-C4	-5.31	1.33	1.38
85	AA	2115	G	C3'-C2'	-5.31	1.47	1.52
85	AA	2163	G	N9-C4	5.31	1.42	1.38
34	BA	608	G	C2'-C1'	-5.31	1.47	1.53
35	BB	353	G	P-O5'	-5.31	1.54	1.59
35	BB	568	A	C2'-C1'	-5.31	1.47	1.53
35	BB	643	G	N1-C2	-5.31	1.33	1.37
35	BB	1147	G	C4'-C3'	-5.31	1.47	1.52
35	BB	1525	G	C6-N1	-5.31	1.35	1.39
85	AA	826	C	P-O5'	-5.31	1.54	1.59
85	AA	1544	G	C5-C4	-5.31	1.34	1.38
34	BA	134	U	O3'-P	-5.31	1.54	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	939	C	C2-N3	-5.31	1.31	1.35
34	BA	998	U	P-O5'	-5.31	1.54	1.59
34	BA	1580	U	C4-C5	-5.31	1.38	1.43
35	BB	479	U	C2'-C1'	-5.31	1.47	1.53
35	BB	688	U	C4'-C3'	-5.31	1.47	1.52
35	BB	1098	G	C4'-O4'	-5.31	1.38	1.45
35	BB	1110	G	C3'-C2'	-5.31	1.47	1.52
36	BC	52	A	C3'-C2'	-5.31	1.47	1.52
36	BC	94	C	C4-N4	-5.31	1.29	1.33
40	BG	175	G	C3'-C2'	-5.31	1.47	1.52
41	BH	66	G	C1'-N9	-5.31	1.39	1.46
85	AA	100	A	O3'-P	-5.31	1.54	1.61
85	AA	650	G	C6-N1	-5.31	1.35	1.39
85	AA	1109	G	N7-C5	-5.31	1.36	1.39
85	AA	1445	C	C4'-O4'	-5.31	1.38	1.45
34	BA	61	G	N3-C4	-5.31	1.31	1.35
34	BA	1053	U	O3'-P	-5.31	1.54	1.61
34	BA	1093	G	N9-C4	-5.31	1.33	1.38
34	BA	1145	U	C2-N3	-5.31	1.34	1.37
34	BA	1277	G	O3'-P	-5.31	1.54	1.61
34	BA	1436	A	O3'-P	-5.31	1.54	1.61
34	BA	1556	A	C5'-C4'	-5.31	1.45	1.51
34	BA	1735	G	C3'-C2'	-5.31	1.47	1.52
34	BA	1793	G	N7-C5	-5.31	1.36	1.39
35	BB	139	G	C4'-C3'	-5.31	1.47	1.52
35	BB	520	G	N1-C2	-5.31	1.33	1.37
35	BB	996	G	C6-N1	-5.31	1.35	1.39
35	BB	1018	U	C2-N3	-5.31	1.34	1.37
35	BB	1106	G	C2'-C1'	-5.31	1.47	1.53
35	BB	1345	A	O3'-P	-5.31	1.54	1.61
35	BB	1408	G	C1'-N9	-5.31	1.39	1.46
35	BB	1530	U	C4'-C3'	-5.31	1.47	1.52
36	BC	52	A	C5'-C4'	-5.31	1.45	1.51
38	BE	178	G	C2-N3	-5.31	1.28	1.32
40	BG	108	G	N9-C4	-5.31	1.33	1.38
85	AA	81	A	N9-C8	-5.31	1.33	1.37
85	AA	172	A	N3-C4	-5.31	1.31	1.34
85	AA	667	A	P-O5'	-5.31	1.54	1.59
85	AA	1917	G	C5-C4	-5.31	1.34	1.38
85	AA	2021	A	O3'-P	-5.31	1.54	1.61
34	BA	1378	A	C2'-C1'	-5.31	1.47	1.53
35	BB	894	A	C5'-C4'	5.31	1.57	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1122	C	N1-C6	-5.31	1.33	1.37
35	BB	1272	G	N9-C8	-5.31	1.34	1.37
40	BG	56	G	N9-C4	-5.31	1.33	1.38
85	AA	63	G	C2-N3	-5.31	1.28	1.32
85	AA	351	C	N1-C6	-5.31	1.33	1.37
85	AA	937	G	C1'-N9	-5.31	1.39	1.46
85	AA	1215	A	C5-C4	-5.31	1.35	1.38
85	AA	1923	A	C4'-O4'	-5.31	1.38	1.45
34	BA	234	A	O4'-C1'	-5.30	1.34	1.41
34	BA	583	G	C5-C4	-5.30	1.34	1.38
34	BA	1293	A	O3'-P	-5.30	1.54	1.61
34	BA	1519	G	N3-C4	-5.30	1.31	1.35
34	BA	1566	G	C8-N7	-5.30	1.27	1.30
34	BA	1685	C	C4-N4	-5.30	1.29	1.33
35	BB	419	G	C3'-C2'	-5.30	1.47	1.52
35	BB	1378	U	C3'-C2'	-5.30	1.47	1.52
35	BB	1434	G	C5-C6	-5.30	1.37	1.42
36	BC	24	G	N1-C2	-5.30	1.33	1.37
40	BG	165	C	C3'-C2'	-5.30	1.47	1.52
85	AA	260	A	C5-C4	-5.30	1.35	1.38
85	AA	277	G	O3'-P	-5.30	1.54	1.61
85	AA	767	A	C3'-C2'	-5.30	1.47	1.52
85	AA	1314	C	P-O5'	-5.30	1.54	1.59
85	AA	1728	G	C4'-C3'	5.30	1.58	1.53
34	BA	422	C	N1-C6	-5.30	1.33	1.37
34	BA	563	A	C5'-C4'	-5.30	1.45	1.51
35	BB	28	G	C5-C6	-5.30	1.37	1.42
35	BB	488	G	C5-C4	-5.30	1.34	1.38
35	BB	671	A	O3'-P	-5.30	1.54	1.61
35	BB	1450	G	C4'-C3'	-5.30	1.47	1.52
37	BD	64	A	C5-C4	-5.30	1.35	1.38
40	BG	150	A	N7-C5	-5.30	1.36	1.39
85	AA	676	U	N3-C4	-5.30	1.33	1.38
85	AA	864	C	O3'-P	-5.30	1.54	1.61
85	AA	1202	G	N1-C2	-5.30	1.33	1.37
85	AA	1585	A	P-O5'	-5.30	1.54	1.59
85	AA	2183	U	C2-N3	-5.30	1.34	1.37
34	BA	320	G	N3-C4	-5.30	1.31	1.35
34	BA	676	G	P-O5'	-5.30	1.54	1.59
34	BA	1086	A	C4'-C3'	-5.30	1.47	1.52
35	BB	34	G	C2'-C1'	-5.30	1.47	1.53
35	BB	565	U	P-O5'	-5.30	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1272	G	N3-C4	-5.30	1.31	1.35
35	BB	1489	A	O3'-P	-5.30	1.54	1.61
38	BE	18	U	N3-C4	-5.30	1.33	1.38
38	BE	112	G	P-O5'	-5.30	1.54	1.59
39	BF	14	C	C4'-C3'	-5.30	1.47	1.52
40	BG	36	G	C2'-C1'	-5.30	1.47	1.53
40	BG	65	C	O4'-C1'	-5.30	1.34	1.41
40	BG	162	A	N1-C2	-5.30	1.29	1.34
77	Br	232	SER	N-CA	-5.30	1.35	1.46
85	AA	23	G	O3'-P	-5.30	1.54	1.61
85	AA	72	C	P-O5'	-5.30	1.54	1.59
85	AA	1176	C	C3'-C2'	-5.30	1.47	1.52
85	AA	1652	A	N3-C4	-5.30	1.31	1.34
85	AA	1670	U	C5'-C4'	-5.30	1.45	1.51
85	AA	1830	U	O3'-P	-5.30	1.54	1.61
34	BA	727	G	N3-C4	-5.30	1.31	1.35
34	BA	1139	G	O4'-C1'	-5.30	1.34	1.41
34	BA	1275	G	C2-N2	-5.30	1.29	1.34
34	BA	1676	A	C5-C4	-5.30	1.35	1.38
34	BA	1835	A	C5-C4	-5.30	1.35	1.38
35	BB	130	G	N9-C8	-5.30	1.34	1.37
35	BB	700	C	C4-C5	-5.30	1.38	1.43
35	BB	726	A	O3'-P	-5.30	1.54	1.61
35	BB	1069	C	C2'-C1'	-5.30	1.47	1.53
35	BB	1226	G	N3-C4	-5.30	1.31	1.35
35	BB	1289	G	C2-N2	-5.30	1.29	1.34
36	BC	121	G	C8-N7	-5.30	1.27	1.30
37	BD	56	G	C4'-C3'	-5.30	1.47	1.52
37	BD	106	G	C1'-N9	-5.30	1.39	1.46
85	AA	250	C	C4-C5	-5.30	1.38	1.43
85	AA	536	C	C2'-C1'	-5.30	1.47	1.53
85	AA	602	U	P-O5'	-5.30	1.54	1.59
85	AA	1006	C	P-O5'	-5.30	1.54	1.59
85	AA	1269	A	N9-C8	-5.30	1.33	1.37
85	AA	1453	U	C2'-C1'	-5.30	1.47	1.53
86	AB	14	A	C2'-C1'	-5.30	1.47	1.53
6	A5	205	GLY	CA-C	-5.30	1.43	1.51
34	BA	363	G	C2-N2	-5.30	1.29	1.34
34	BA	734	G	C6-N1	-5.30	1.35	1.39
35	BB	1286	G	N9-C8	-5.30	1.34	1.37
35	BB	1468	A	C4'-O4'	-5.30	1.38	1.45
85	AA	360	C	C3'-O3'	5.30	1.49	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	404	A	N7-C5	-5.30	1.36	1.39
85	AA	1185	G	N7-C5	-5.30	1.36	1.39
85	AA	1991	C	C2'-C1'	-5.30	1.47	1.53
34	BA	230	A	C3'-C2'	-5.30	1.47	1.52
34	BA	323	C	O3'-P	-5.30	1.54	1.61
34	BA	425	G	O3'-P	-5.30	1.54	1.61
34	BA	564	C	C4-C5	-5.30	1.38	1.43
34	BA	765	U	C3'-O3'	5.30	1.49	1.42
34	BA	992	A	N9-C8	-5.30	1.33	1.37
34	BA	1001	G	C6-N1	-5.30	1.35	1.39
34	BA	1012	A	N9-C4	-5.30	1.34	1.37
34	BA	1218	G	P-O5'	-5.30	1.54	1.59
34	BA	1234	U	C5'-C4'	-5.30	1.45	1.51
34	BA	1695	G	C2-N2	-5.30	1.29	1.34
35	BB	57	G	N7-C5	-5.30	1.36	1.39
35	BB	627	G	O3'-P	-5.30	1.54	1.61
37	BD	115	A	C2'-C1'	-5.30	1.47	1.53
41	BH	24	U	O3'-P	-5.30	1.54	1.61
85	AA	48	G	C3'-C2'	-5.30	1.47	1.52
85	AA	690	G	C1'-N9	-5.30	1.39	1.46
85	AA	722	G	O3'-P	-5.30	1.54	1.61
85	AA	820	G	C1'-N9	-5.30	1.39	1.46
85	AA	1808	G	O3'-P	-5.30	1.54	1.61
85	AA	2083	G	N9-C4	5.30	1.42	1.38
86	AB	9	A	P-O5'	-5.30	1.54	1.59
34	BA	52	G	N9-C8	-5.29	1.34	1.37
34	BA	276	C	O3'-P	-5.29	1.54	1.61
34	BA	605	G	C8-N7	-5.29	1.27	1.30
34	BA	1069	U	C4'-C3'	-5.29	1.47	1.52
34	BA	1512	C	O4'-C1'	-5.29	1.34	1.41
35	BB	425	G	C2'-C1'	-5.29	1.47	1.53
35	BB	468	U	C5'-C4'	-5.29	1.45	1.51
35	BB	628	A	N9-C8	-5.29	1.33	1.37
35	BB	1119	G	O4'-C1'	-5.29	1.34	1.41
36	BC	48	A	N9-C4	-5.29	1.34	1.37
85	AA	472	A	C2'-C1'	-5.29	1.47	1.53
85	AA	1165	C	C2-N3	-5.29	1.31	1.35
85	AA	1249	U	N3-C4	-5.29	1.33	1.38
85	AA	1257	A	N9-C8	-5.29	1.33	1.37
34	BA	113	G	C5-C4	-5.29	1.34	1.38
34	BA	313	C	C3'-C2'	-5.29	1.47	1.52
34	BA	478	G	C5-C6	-5.29	1.37	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	908	G	C2-N2	-5.29	1.29	1.34
34	BA	1284	G	N9-C8	-5.29	1.34	1.37
34	BA	1453	U	O3'-P	-5.29	1.54	1.61
34	BA	1501	U	N1-C6	-5.29	1.33	1.38
34	BA	1585	A	C4'-C3'	-5.29	1.47	1.52
34	BA	1597	G	O3'-P	-5.29	1.54	1.61
34	BA	1825	U	N1-C6	-5.29	1.33	1.38
35	BB	410	A	O3'-P	-5.29	1.54	1.61
35	BB	1066	G	P-O5'	-5.29	1.54	1.59
35	BB	1067	G	C4'-C3'	-5.29	1.47	1.52
35	BB	1231	U	P-O5'	-5.29	1.54	1.59
35	BB	1494	G	N9-C4	-5.29	1.33	1.38
85	AA	264	A	C8-N7	-5.29	1.27	1.31
85	AA	1911	A	C3'-C2'	-5.29	1.47	1.52
85	AA	2122	A	O4'-C1'	-5.29	1.34	1.41
34	BA	125	G	C5-C4	-5.29	1.34	1.38
34	BA	497	U	C3'-C2'	-5.29	1.47	1.52
34	BA	603	U	P-O5'	-5.29	1.54	1.59
35	BB	636	G	C5-C4	-5.29	1.34	1.38
35	BB	816	U	C1'-N1	-5.29	1.39	1.46
35	BB	988	G	N3-C4	-5.29	1.31	1.35
35	BB	1337	C	C4-N4	-5.29	1.29	1.33
35	BB	1449	G	O4'-C1'	-5.29	1.34	1.41
35	BB	1478	G	N1-C2	-5.29	1.33	1.37
35	BB	1524	G	P-O5'	-5.29	1.54	1.59
39	BF	11	C	C4'-O4'	5.29	1.52	1.45
39	BF	59	U	O3'-P	-5.29	1.54	1.61
40	BG	153	C	O4'-C1'	-5.29	1.34	1.41
41	BH	17	A	N7-C5	-5.29	1.36	1.39
85	AA	438	G	O3'-P	-5.29	1.54	1.61
85	AA	1424	G	C5'-C4'	5.29	1.57	1.51
85	AA	1878	C	O3'-P	-5.29	1.54	1.61
85	AA	2134	U	C4'-C3'	-5.29	1.47	1.52
34	BA	654	C	C4-N4	-5.29	1.29	1.33
34	BA	674	G	P-O5'	-5.29	1.54	1.59
34	BA	969	A	C5-C6	-5.29	1.36	1.41
34	BA	988	U	C2-N3	-5.29	1.34	1.37
35	BB	567	G	C1'-N9	-5.29	1.39	1.46
35	BB	576	A	C6-N1	-5.29	1.31	1.35
35	BB	1050	A	C5-C6	-5.29	1.36	1.41
40	BG	59	G	C5-C6	-5.29	1.37	1.42
41	BH	35	G	N9-C4	-5.29	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	407	G	N9-C8	-5.29	1.34	1.37
85	AA	637	U	C4'-O4'	-5.29	1.38	1.45
85	AA	1496	U	N1-C6	-5.29	1.33	1.38
34	BA	666	C	N1-C2	-5.29	1.34	1.40
34	BA	745	A	N9-C4	5.29	1.41	1.37
34	BA	930	A	C1'-N9	-5.29	1.39	1.46
34	BA	1063	G	C2-N2	-5.29	1.29	1.34
34	BA	1191	C	C4'-C3'	-5.29	1.47	1.52
34	BA	1698	C	C1'-N1	-5.29	1.39	1.46
34	BA	1704	G	C6-N1	-5.29	1.35	1.39
35	BB	136	A	N3-C4	-5.29	1.31	1.34
35	BB	349	U	P-O5'	-5.29	1.54	1.59
35	BB	365	U	C3'-C2'	-5.29	1.47	1.52
35	BB	583	G	C5-C6	-5.29	1.37	1.42
35	BB	839	G	C1'-N9	-5.29	1.39	1.46
35	BB	1434	G	N9-C4	-5.29	1.33	1.38
40	BG	23	C	O4'-C1'	-5.29	1.34	1.41
85	AA	590	U	C2-N3	-5.29	1.34	1.37
85	AA	1112	G	O3'-P	-5.29	1.54	1.61
85	AA	1547	G	C3'-C2'	-5.29	1.47	1.52
85	AA	1902	C	O3'-P	-5.29	1.54	1.61
85	AA	2022	A	C5-C6	-5.29	1.36	1.41
85	AA	2030	U	O3'-P	-5.29	1.54	1.61
34	BA	1010	C	C2'-C1'	-5.29	1.47	1.53
34	BA	1040	G	C6-N1	-5.29	1.35	1.39
35	BB	1280	U	C2'-C1'	-5.29	1.47	1.53
36	BC	108	A	C2'-C1'	-5.29	1.47	1.53
38	BE	25	U	O4'-C1'	-5.29	1.34	1.41
38	BE	149	A	C8-N7	-5.29	1.27	1.31
85	AA	1955	U	N3-C4	-5.29	1.33	1.38
34	BA	78	U	N1-C6	-5.29	1.33	1.38
34	BA	386	A	C5-C6	-5.29	1.36	1.41
34	BA	395	G	C2'-C1'	-5.29	1.47	1.53
34	BA	931	G	C2-N2	-5.29	1.29	1.34
34	BA	1083	A	N9-C8	-5.29	1.33	1.37
34	BA	1152	A	C8-N7	-5.29	1.27	1.31
34	BA	1328	U	C4'-C3'	-5.29	1.47	1.52
35	BB	55	C	C3'-C2'	-5.29	1.47	1.52
35	BB	578	G	C1'-N9	-5.29	1.39	1.46
35	BB	695	U	N1-C2	-5.29	1.33	1.38
35	BB	1293	C	C4-N4	-5.29	1.29	1.33
36	BC	70	C	C3'-O3'	5.29	1.49	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	14	C	C3'-C2'	-5.29	1.47	1.52
37	BD	29	C	C2'-C1'	-5.29	1.47	1.53
40	BG	80	G	P-O5'	-5.29	1.54	1.59
41	BH	126	C	C1'-N1	-5.29	1.39	1.46
61	Bb	132	LYS	N-CA	-5.29	1.35	1.46
85	AA	313	A	O4'-C1'	-5.29	1.34	1.41
85	AA	554	A	N9-C4	-5.29	1.34	1.37
85	AA	687	G	N9-C4	-5.29	1.33	1.38
85	AA	1210	U	C3'-O3'	5.29	1.49	1.42
20	AL	5	ARG	CD-NE	5.28	1.55	1.46
34	BA	49	A	C4'-C3'	-5.28	1.47	1.52
34	BA	422	C	C2'-C1'	-5.28	1.47	1.53
34	BA	764	G	N9-C4	5.28	1.42	1.38
34	BA	946	A	N3-C4	-5.28	1.31	1.34
34	BA	1108	U	C4'-C3'	-5.28	1.47	1.52
34	BA	1124	U	O4'-C1'	-5.28	1.34	1.41
35	BB	8	U	C4-C5	-5.28	1.38	1.43
35	BB	104	G	N9-C8	-5.28	1.34	1.37
35	BB	1057	G	C2-N2	-5.28	1.29	1.34
35	BB	1077	C	N3-C4	-5.28	1.30	1.33
35	BB	1524	G	C5-C4	-5.28	1.34	1.38
38	BE	18	U	C2'-C1'	-5.28	1.47	1.53
40	BG	49	A	N3-C4	-5.28	1.31	1.34
40	BG	166	C	C4-N4	-5.28	1.29	1.33
41	BH	43	G	N7-C5	-5.28	1.36	1.39
85	AA	698	G	N9-C8	-5.28	1.34	1.37
85	AA	855	G	C4'-O4'	-5.28	1.38	1.45
85	AA	914	U	P-O5'	-5.28	1.54	1.59
85	AA	921	C	C2'-C1'	-5.28	1.47	1.53
85	AA	1443	U	O3'-P	-5.28	1.54	1.61
85	AA	2128	G	C2-N3	-5.28	1.28	1.32
34	BA	1	C	N1-C2	-5.28	1.34	1.40
34	BA	418	G	C2'-C1'	-5.28	1.47	1.53
34	BA	919	A	C1'-N9	-5.28	1.39	1.46
34	BA	926	A	N9-C8	-5.28	1.33	1.37
34	BA	1644	A	C1'-N9	-5.28	1.39	1.46
35	BB	52	G	C6-N1	-5.28	1.35	1.39
85	AA	308	U	N1-C6	-5.28	1.33	1.38
34	BA	9	A	N3-C4	-5.28	1.31	1.34
34	BA	87	G	C2-N2	-5.28	1.29	1.34
34	BA	129	U	C4-C5	-5.28	1.38	1.43
34	BA	295	G	N7-C5	-5.28	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	503	C	C4'-C3'	-5.28	1.47	1.52
34	BA	796	G	N3-C4	-5.28	1.31	1.35
34	BA	1448	G	N9-C4	-5.28	1.33	1.38
35	BB	111	C	P-O5'	-5.28	1.54	1.59
35	BB	522	A	N7-C5	-5.28	1.36	1.39
36	BC	135	A	N7-C5	-5.28	1.36	1.39
40	BG	32	U	C3'-C2'	-5.28	1.47	1.52
40	BG	102	G	C1'-N9	-5.28	1.39	1.46
40	BG	113	G	C3'-C2'	-5.28	1.47	1.52
40	BG	124	A	C5-C4	-5.28	1.35	1.38
41	BH	34	G	C3'-C2'	-5.28	1.47	1.52
85	AA	5	U	C3'-C2'	-5.28	1.47	1.52
85	AA	437	G	C8-N7	-5.28	1.27	1.30
85	AA	922	A	C1'-N9	-5.28	1.39	1.46
85	AA	1125	G	C5-C6	-5.28	1.37	1.42
85	AA	1203	G	N7-C5	-5.28	1.36	1.39
85	AA	1208	C	O3'-P	-5.28	1.54	1.61
85	AA	1585	A	O3'-P	-5.28	1.54	1.61
85	AA	1937	G	C2'-C1'	-5.28	1.47	1.53
2	A1	133	ILE	C-N	-5.28	1.24	1.34
34	BA	55	G	C5-C4	-5.28	1.34	1.38
34	BA	426	A	C5-C6	-5.28	1.36	1.41
34	BA	760	G	C5'-C4'	-5.28	1.45	1.51
35	BB	971	A	N9-C8	-5.28	1.33	1.37
36	BC	24	G	N3-C4	-5.28	1.31	1.35
37	BD	44	U	O3'-P	-5.28	1.54	1.61
34	BA	267	G	N3-C4	-5.28	1.31	1.35
34	BA	268	U	N3-C4	-5.28	1.33	1.38
34	BA	1219	G	C2-N2	-5.28	1.29	1.34
34	BA	1425	G	N3-C4	-5.28	1.31	1.35
34	BA	1513	G	C1'-N9	-5.28	1.39	1.46
35	BB	34	G	C5-C4	-5.28	1.34	1.38
35	BB	485	U	N3-C4	-5.28	1.33	1.38
35	BB	1422	G	C2-N2	-5.28	1.29	1.34
35	BB	1524	G	N1-C2	-5.28	1.33	1.37
34	BA	186	G	C1'-N9	-5.28	1.39	1.46
34	BA	1000	G	N1-C2	-5.28	1.33	1.37
35	BB	546	A	C1'-N9	-5.28	1.39	1.46
35	BB	808	U	C4-C5	-5.28	1.38	1.43
35	BB	1130	U	C5'-C4'	-5.28	1.45	1.51
35	BB	1316	U	C2'-C1'	-5.28	1.47	1.53
36	BC	25	C	C5-C6	-5.28	1.30	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	160	C	N1-C6	-5.28	1.33	1.37
37	BD	61	C	C4-N4	-5.28	1.29	1.33
40	BG	77	U	C3'-C2'	-5.28	1.47	1.52
85	AA	527	A	C5-C4	-5.28	1.35	1.38
85	AA	716	G	C3'-C2'	-5.28	1.47	1.52
85	AA	750	A	N3-C4	-5.28	1.31	1.34
85	AA	862	U	C3'-O3'	-5.28	1.34	1.42
85	AA	1168	C	C3'-C2'	-5.28	1.47	1.52
85	AA	1227	A	C4'-C3'	-5.28	1.47	1.52
85	AA	1672	G	C1'-N9	-5.28	1.39	1.46
85	AA	1687	U	C3'-C2'	-5.28	1.47	1.52
34	BA	56	G	N1-C2	-5.27	1.33	1.37
34	BA	506	U	C2-N3	-5.27	1.34	1.37
34	BA	1422	A	P-O5'	-5.27	1.54	1.59
35	BB	125	G	C3'-C2'	-5.27	1.47	1.52
35	BB	1498	G	C1'-N9	-5.27	1.39	1.46
85	AA	644	A	C3'-C2'	-5.27	1.47	1.52
85	AA	814	G	C2-N2	-5.27	1.29	1.34
85	AA	1422	A	P-O5'	-5.27	1.54	1.59
85	AA	2057	G	C2'-C1'	-5.27	1.47	1.53
13	AE	86	ILE	CA-CB	-5.27	1.42	1.54
34	BA	99	G	N7-C5	-5.27	1.36	1.39
34	BA	277	A	C1'-N9	-5.27	1.39	1.46
34	BA	359	G	C3'-C2'	-5.27	1.47	1.52
34	BA	477	C	C3'-C2'	-5.27	1.47	1.52
34	BA	624	G	N9-C4	5.27	1.42	1.38
34	BA	755	G	N7-C5	-5.27	1.36	1.39
34	BA	1170	A	O3'-P	-5.27	1.54	1.61
34	BA	1256	A	N9-C8	-5.27	1.33	1.37
34	BA	1703	A	C3'-C2'	-5.27	1.47	1.52
35	BB	30	A	C5-C4	-5.27	1.35	1.38
35	BB	701	U	N3-C4	-5.27	1.33	1.38
35	BB	1000	U	C4'-C3'	-5.27	1.47	1.52
35	BB	1309	A	C5-C4	-5.27	1.35	1.38
36	BC	136	G	P-O5'	-5.27	1.54	1.59
37	BD	32	A	N7-C5	-5.27	1.36	1.39
38	BE	169	C	P-O5'	-5.27	1.54	1.59
41	BH	107	A	C5-C4	-5.27	1.35	1.38
85	AA	97	A	N3-C4	-5.27	1.31	1.34
85	AA	574	U	C3'-C2'	-5.27	1.47	1.52
85	AA	587	G	N9-C4	-5.27	1.33	1.38
85	AA	1301	C	P-O5'	-5.27	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1677	A	C5-C4	-5.27	1.35	1.38
34	BA	1604	A	C3'-C2'	-5.27	1.47	1.52
35	BB	313	C	P-O5'	-5.27	1.54	1.59
35	BB	983	C	C2'-C1'	-5.27	1.47	1.53
35	BB	1410	G	P-O5'	-5.27	1.54	1.59
35	BB	1466	A	C1'-N9	-5.27	1.39	1.46
41	BH	1	U	C3'-O3'	5.27	1.49	1.42
85	AA	40	A	P-O5'	-5.27	1.54	1.59
85	AA	476	C	N3-C4	-5.27	1.30	1.33
85	AA	1615	A	O3'-P	-5.27	1.54	1.61
85	AA	1672	G	N9-C4	-5.27	1.33	1.38
85	AA	1793	A	N9-C4	5.27	1.41	1.37
86	AB	39	U	C2'-C1'	-5.27	1.47	1.53
34	BA	801	U	C4'-C3'	-5.27	1.47	1.52
34	BA	1024	A	N3-C4	-5.27	1.31	1.34
34	BA	1302	C	C2'-C1'	-5.27	1.47	1.53
34	BA	1403	G	C6-N1	-5.27	1.35	1.39
34	BA	1418	G	C8-N7	-5.27	1.27	1.30
35	BB	9	G	N9-C8	-5.27	1.34	1.37
35	BB	66	G	O4'-C1'	-5.27	1.34	1.41
35	BB	93	A	C5-C4	-5.27	1.35	1.38
35	BB	1031	G	C6-N1	-5.27	1.35	1.39
35	BB	1076	U	C1'-N1	-5.27	1.39	1.46
35	BB	1149	A	N7-C5	-5.27	1.36	1.39
35	BB	1177	U	N1-C6	-5.27	1.33	1.38
39	BF	3	A	C6-N1	-5.27	1.31	1.35
85	AA	173	A	P-O5'	-5.27	1.54	1.59
85	AA	202	U	N1-C2	-5.27	1.33	1.38
85	AA	503	A	N3-C4	-5.27	1.31	1.34
85	AA	869	A	C8-N7	-5.27	1.27	1.31
85	AA	911	A	C2'-C1'	-5.27	1.47	1.53
85	AA	1097	G	O3'-P	-5.27	1.54	1.61
85	AA	1439	A	C2'-C1'	-5.27	1.47	1.53
34	BA	52	G	C6-O6	-5.27	1.19	1.24
34	BA	482	C	O4'-C1'	-5.27	1.34	1.41
34	BA	741	A	O3'-P	-5.27	1.54	1.61
34	BA	794	G	C2'-C1'	-5.27	1.47	1.53
35	BB	128	C	O3'-P	-5.27	1.54	1.61
35	BB	272	C	P-O5'	-5.27	1.54	1.59
35	BB	363	A	O3'-P	-5.27	1.54	1.61
35	BB	377	A	N3-C4	-5.27	1.31	1.34
35	BB	627	G	N1-C2	-5.27	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	997	G	C2-N2	-5.27	1.29	1.34
35	BB	1278	A	N9-C8	-5.27	1.33	1.37
37	BD	72	U	C2'-C1'	-5.27	1.47	1.53
38	BE	95	G	O4'-C1'	-5.27	1.34	1.41
40	BG	127	G	N1-C2	-5.27	1.33	1.37
40	BG	130	G	N3-C4	-5.27	1.31	1.35
85	AA	541	A	C5'-C4'	-5.27	1.45	1.51
85	AA	586	G	C5-C4	-5.27	1.34	1.38
85	AA	1452	C	C3'-C2'	-5.27	1.47	1.52
85	AA	1711	C	C3'-C2'	-5.27	1.47	1.52
85	AA	1899	A	P-O5'	-5.27	1.54	1.59
85	AA	2080	U	P-O5'	-5.27	1.54	1.59
85	AA	2124	G	C5-C4	-5.27	1.34	1.38
85	AA	2247	C	C5'-C4'	5.27	1.57	1.51
34	BA	572	G	C4'-C3'	5.27	1.58	1.53
34	BA	584	A	N9-C4	5.27	1.41	1.37
34	BA	1607	U	P-O5'	-5.27	1.54	1.59
35	BB	425	G	C2-N2	-5.27	1.29	1.34
35	BB	629	C	C3'-C2'	-5.27	1.47	1.52
35	BB	1031	G	P-O5'	-5.27	1.54	1.59
39	BF	6	C	P-O5'	-5.27	1.54	1.59
85	AA	427	G	C1'-N9	-5.27	1.39	1.46
85	AA	2041	G	N7-C5	-5.27	1.36	1.39
34	BA	76	U	N1-C2	-5.26	1.33	1.38
34	BA	85	C	C3'-C2'	-5.26	1.47	1.52
34	BA	86	A	C2'-C1'	-5.26	1.47	1.53
34	BA	199	U	C3'-C2'	-5.26	1.47	1.52
34	BA	308	C	C4'-C3'	-5.26	1.47	1.52
34	BA	321	G	N3-C4	-5.26	1.31	1.35
34	BA	707	C	P-O5'	-5.26	1.54	1.59
34	BA	825	G	N3-C4	-5.26	1.31	1.35
34	BA	1277	G	C5-C4	-5.26	1.34	1.38
34	BA	1409	A	C6-N6	-5.26	1.29	1.33
34	BA	1413	G	N3-C4	-5.26	1.31	1.35
34	BA	1556	A	C5-C6	-5.26	1.36	1.41
34	BA	1558	C	C4-N4	-5.26	1.29	1.33
34	BA	1720	U	C4-O4	-5.26	1.19	1.23
35	BB	557	C	C4-N4	-5.26	1.29	1.33
35	BB	606	C	C2'-C1'	-5.26	1.47	1.53
35	BB	1167	C	P-O5'	-5.26	1.54	1.59
85	AA	670	C	C4'-C3'	-5.26	1.47	1.52
85	AA	883	A	C2'-C1'	-5.26	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1174	G	O3'-P	-5.26	1.54	1.61
85	AA	1183	C	P-O5'	-5.26	1.54	1.59
85	AA	1514	A	O3'-P	-5.26	1.54	1.61
85	AA	1976	G	C4'-C3'	-5.26	1.47	1.52
34	BA	532	C	N1-C2	-5.26	1.34	1.40
34	BA	614	A	N7-C5	-5.26	1.36	1.39
34	BA	1021	U	O3'-P	-5.26	1.54	1.61
34	BA	1454	G	N7-C5	-5.26	1.36	1.39
34	BA	1839	G	N9-C4	-5.26	1.33	1.38
36	BC	25	C	N1-C6	-5.26	1.33	1.37
36	BC	80	A	O3'-P	-5.26	1.54	1.61
38	BE	202	C	O3'-P	-5.26	1.54	1.61
85	AA	537	G	C1'-N9	-5.26	1.39	1.46
85	AA	817	G	C3'-C2'	-5.26	1.47	1.52
85	AA	961	U	C3'-C2'	-5.26	1.47	1.52
85	AA	1098	C	C3'-C2'	-5.26	1.47	1.52
34	BA	126	G	C5-C6	-5.26	1.37	1.42
34	BA	1152	A	C5-C6	-5.26	1.36	1.41
34	BA	1160	U	C1'-N1	-5.26	1.39	1.46
34	BA	1262	A	C3'-C2'	-5.26	1.47	1.52
34	BA	1679	C	O3'-P	-5.26	1.54	1.61
35	BB	260	A	P-O5'	-5.26	1.54	1.59
35	BB	589	U	C2-N3	-5.26	1.34	1.37
35	BB	1239	A	N9-C4	-5.26	1.34	1.37
36	BC	73	U	C2-N3	-5.26	1.34	1.37
40	BG	157	A	C4'-C3'	-5.26	1.47	1.52
85	AA	324	U	C2'-C1'	-5.26	1.47	1.53
85	AA	448	G	N7-C5	-5.26	1.36	1.39
85	AA	519	A	C3'-C2'	-5.26	1.47	1.52
85	AA	689	U	C5'-C4'	-5.26	1.45	1.51
85	AA	781	G	C6-N1	-5.26	1.35	1.39
85	AA	908	C	P-O5'	-5.26	1.54	1.59
85	AA	968	U	O3'-P	-5.26	1.54	1.61
85	AA	1016	G	C2'-C1'	-5.26	1.47	1.53
85	AA	1263	G	C3'-C2'	-5.26	1.47	1.52
85	AA	1272	G	C5-C4	-5.26	1.34	1.38
85	AA	1442	U	O3'-P	-5.26	1.54	1.61
85	AA	1531	G	N9-C8	-5.26	1.34	1.37
85	AA	1878	C	C2'-C1'	-5.26	1.47	1.53
13	AE	46	GLY	CA-C	-5.26	1.43	1.51
34	BA	183	G	C5-C6	-5.26	1.37	1.42
34	BA	257	G	C2-N3	-5.26	1.28	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1012	A	O3'-P	-5.26	1.54	1.61
34	BA	1116	G	C2-N2	-5.26	1.29	1.34
34	BA	1269	C	C1'-N1	-5.26	1.39	1.46
34	BA	1509	U	C1'-N1	-5.26	1.39	1.46
35	BB	366	G	C3'-C2'	-5.26	1.47	1.52
35	BB	537	A	N3-C4	-5.26	1.31	1.34
35	BB	995	C	P-O5'	-5.26	1.54	1.59
35	BB	1330	A	N7-C5	-5.26	1.36	1.39
35	BB	1335	G	N9-C8	-5.26	1.34	1.37
35	BB	1337	C	O3'-P	-5.26	1.54	1.61
35	BB	1469	A	O4'-C1'	-5.26	1.34	1.41
35	BB	1475	U	C4'-O4'	5.26	1.52	1.45
38	BE	126	G	C2-N2	-5.26	1.29	1.34
39	BF	62	U	P-O5'	-5.26	1.54	1.59
41	BH	36	C	O4'-C1'	-5.26	1.34	1.41
85	AA	342	C	O3'-P	-5.26	1.54	1.61
85	AA	452	A	N9-C8	-5.26	1.33	1.37
85	AA	490	A	C6-N1	5.26	1.39	1.35
85	AA	570	U	P-O5'	-5.26	1.54	1.59
85	AA	756	G	O4'-C1'	-5.26	1.34	1.41
85	AA	860	C	C1'-N1	-5.26	1.39	1.46
85	AA	1462	A	O4'-C1'	-5.26	1.34	1.41
85	AA	1565	G	C2'-C1'	-5.26	1.47	1.53
85	AA	1578	G	P-O5'	5.26	1.65	1.59
85	AA	1649	U	P-O5'	-5.26	1.54	1.59
85	AA	1961	U	C2-N3	-5.26	1.34	1.37
85	AA	2028	G	C2-N3	-5.26	1.28	1.32
85	AA	2188	C	C1'-N1	-5.26	1.39	1.46
34	BA	1074	C	C4-N4	-5.26	1.29	1.33
35	BB	116	G	N3-C4	-5.26	1.31	1.35
40	BG	173	C	P-O5'	-5.26	1.54	1.59
85	AA	1439	A	C3'-C2'	-5.26	1.47	1.52
34	BA	20	A	C6-N1	-5.26	1.31	1.35
34	BA	74	A	N3-C4	-5.26	1.31	1.34
34	BA	103	G	P-O5'	-5.26	1.54	1.59
34	BA	1222	C	C4-C5	-5.26	1.38	1.43
34	BA	1227	U	N3-C4	-5.26	1.33	1.38
34	BA	1256	A	C5-C4	-5.26	1.35	1.38
34	BA	1583	A	N9-C8	-5.26	1.33	1.37
35	BB	534	C	C2-N3	-5.26	1.31	1.35
35	BB	547	A	C4'-O4'	-5.26	1.38	1.45
37	BD	7	G	C6-N1	-5.26	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	106	G	C2'-C1'	-5.26	1.47	1.53
85	AA	83	U	C2'-C1'	-5.26	1.47	1.53
85	AA	152	A	C1'-N9	-5.26	1.39	1.46
85	AA	158	C	N1-C6	-5.26	1.33	1.37
85	AA	352	G	C8-N7	-5.26	1.27	1.30
85	AA	557	G	C4'-C3'	-5.26	1.47	1.52
86	AB	18	G	O3'-P	-5.26	1.54	1.61
34	BA	412	G	N3-C4	-5.25	1.31	1.35
34	BA	1082	U	P-O5'	-5.25	1.54	1.59
35	BB	509	A	C5-C4	-5.25	1.35	1.38
35	BB	1020	U	C2'-C1'	-5.25	1.47	1.53
35	BB	1197	G	C1'-N9	-5.25	1.39	1.46
35	BB	1392	A	C2'-C1'	-5.25	1.47	1.53
38	BE	28	C	C4-N4	-5.25	1.29	1.33
41	BH	74	G	N3-C4	-5.25	1.31	1.35
85	AA	1146	C	C2-N3	-5.25	1.31	1.35
85	AA	1865	C	C3'-C2'	-5.25	1.47	1.52
34	BA	162	G	C2-N2	-5.25	1.29	1.34
34	BA	229	C	C3'-C2'	-5.25	1.47	1.52
34	BA	966	G	P-O5'	-5.25	1.54	1.59
34	BA	1166	A	C5-C4	-5.25	1.35	1.38
34	BA	1227	U	C4'-C3'	5.25	1.58	1.53
35	BB	483	C	C2-N3	-5.25	1.31	1.35
35	BB	970	C	P-O5'	-5.25	1.54	1.59
35	BB	1338	U	P-O5'	-5.25	1.54	1.59
35	BB	1524	G	C3'-C2'	-5.25	1.47	1.52
38	BE	44	C	N1-C2	-5.25	1.34	1.40
85	AA	747	U	N3-C4	-5.25	1.33	1.38
85	AA	1196	C	O4'-C1'	-5.25	1.34	1.41
85	AA	1285	C	P-O5'	-5.25	1.54	1.59
85	AA	2221	A	C2'-C1'	-5.25	1.47	1.53
34	BA	12	G	N9-C8	-5.25	1.34	1.37
34	BA	129	U	C3'-C2'	-5.25	1.47	1.52
34	BA	349	G	N7-C5	-5.25	1.36	1.39
34	BA	671	C	C2'-C1'	-5.25	1.47	1.53
34	BA	751	A	C3'-C2'	-5.25	1.47	1.52
34	BA	813	C	N1-C6	-5.25	1.33	1.37
34	BA	844	U	C2-N3	-5.25	1.34	1.37
35	BB	465	C	C3'-C2'	-5.25	1.47	1.52
35	BB	511	A	C6-N6	-5.25	1.29	1.33
35	BB	1094	A	C3'-C2'	-5.25	1.47	1.52
38	BE	88	G	C8-N7	-5.25	1.27	1.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	319	U	C2'-C1'	-5.25	1.47	1.53
85	AA	359	A	C1'-N9	-5.25	1.39	1.46
85	AA	429	G	C8-N7	-5.25	1.27	1.30
85	AA	698	G	C5-C4	-5.25	1.34	1.38
85	AA	736	U	C5'-C4'	5.25	1.57	1.51
85	AA	1207	C	C3'-C2'	-5.25	1.47	1.52
85	AA	1650	G	N9-C8	-5.25	1.34	1.37
85	AA	1861	A	C4'-C3'	-5.25	1.47	1.52
85	AA	2191	C	C4-N4	-5.25	1.29	1.33
85	AA	2221	A	C1'-N9	-5.25	1.39	1.46
34	BA	114	U	O4'-C1'	-5.25	1.34	1.41
34	BA	192	G	N7-C5	-5.25	1.36	1.39
34	BA	667	U	C2-N3	-5.25	1.34	1.37
34	BA	1103	G	C4'-O4'	-5.25	1.38	1.45
35	BB	73	G	P-O5'	-5.25	1.54	1.59
35	BB	1394	A	C5'-C4'	-5.25	1.45	1.51
35	BB	1456	G	P-O5'	-5.25	1.54	1.59
85	AA	320	U	P-O5'	-5.25	1.54	1.59
85	AA	936	C	C4'-O4'	-5.25	1.38	1.45
34	BA	753	G	C1'-N9	-5.25	1.39	1.46
34	BA	1192	A	C3'-O3'	-5.25	1.34	1.42
34	BA	1324	G	C3'-C2'	-5.25	1.47	1.52
34	BA	1657	A	C3'-C2'	-5.25	1.47	1.52
34	BA	1779	U	C2'-C1'	-5.25	1.47	1.53
35	BB	93	A	N3-C4	-5.25	1.31	1.34
35	BB	490	G	C1'-N9	-5.25	1.39	1.46
35	BB	1299	G	C1'-N9	-5.25	1.39	1.46
35	BB	1343	C	N1-C6	-5.25	1.34	1.37
37	BD	74	A	C4'-O4'	-5.25	1.38	1.45
38	BE	168	C	C3'-C2'	-5.25	1.47	1.52
38	BE	203	C	C1'-N1	-5.25	1.39	1.46
39	BF	54	U	N1-C6	-5.25	1.33	1.38
41	BH	32	U	C5'-C4'	-5.25	1.45	1.51
85	AA	116	G	N7-C5	-5.25	1.36	1.39
34	BA	254	U	P-O5'	-5.25	1.54	1.59
34	BA	461	A	C2'-C1'	-5.25	1.47	1.53
34	BA	734	G	N1-C2	-5.25	1.33	1.37
34	BA	1340	G	O3'-P	-5.25	1.54	1.61
34	BA	1618	A	C8-N7	-5.25	1.27	1.31
34	BA	1739	G	C3'-C2'	-5.25	1.47	1.52
34	BA	1814	U	P-O5'	-5.25	1.54	1.59
35	BB	34	G	N3-C4	-5.25	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	386	G	C5-C6	-5.25	1.37	1.42
35	BB	1488	G	C3'-C2'	-5.25	1.47	1.52
40	BG	45	G	C3'-C2'	-5.25	1.47	1.52
85	AA	496	C	N3-C4	-5.25	1.30	1.33
85	AA	970	U	C2'-C1'	5.25	1.59	1.53
85	AA	983	A	O3'-P	-5.25	1.54	1.61
85	AA	1577	G	C2-N2	-5.25	1.29	1.34
85	AA	1660	U	N3-C4	-5.25	1.33	1.38
85	AA	2028	G	C2-N2	-5.25	1.29	1.34
85	AA	2036	A	C2'-C1'	-5.25	1.47	1.53
85	AA	2039	G	N9-C8	-5.25	1.34	1.37
34	BA	621	G	N7-C5	-5.25	1.36	1.39
34	BA	974	G	N9-C8	-5.25	1.34	1.37
34	BA	1090	A	N3-C4	-5.25	1.31	1.34
35	BB	822	G	C6-N1	-5.25	1.35	1.39
39	BF	52	A	C2'-C1'	-5.25	1.47	1.53
85	AA	529	G	N3-C4	-5.25	1.31	1.35
85	AA	596	A	C5-C4	-5.25	1.35	1.38
85	AA	833	U	C3'-C2'	-5.25	1.47	1.52
6	A5	78	ILE	CA-C	-5.24	1.39	1.52
34	BA	144	C	C2'-C1'	-5.24	1.47	1.53
34	BA	593	G	N7-C5	-5.24	1.36	1.39
34	BA	672	G	C6-N1	-5.24	1.35	1.39
34	BA	789	U	C2'-C1'	-5.24	1.47	1.53
34	BA	1010	C	N1-C6	-5.24	1.34	1.37
34	BA	1029	C	C4'-C3'	-5.24	1.47	1.52
34	BA	1576	C	C2'-C1'	-5.24	1.47	1.53
34	BA	1808	A	N9-C8	-5.24	1.33	1.37
35	BB	492	U	C3'-C2'	-5.24	1.47	1.52
35	BB	607	G	C1'-N9	-5.24	1.39	1.46
35	BB	1335	G	C3'-C2'	-5.24	1.47	1.52
38	BE	94	U	C5'-C4'	-5.24	1.45	1.51
39	BF	36	G	C2-N2	-5.24	1.29	1.34
85	AA	363	A	O4'-C1'	-5.24	1.34	1.41
85	AA	364	C	C1'-N1	-5.24	1.39	1.46
85	AA	994	A	C5-C4	-5.24	1.35	1.38
85	AA	1441	G	C2-N2	-5.24	1.29	1.34
85	AA	1457	C	O4'-C1'	-5.24	1.34	1.41
85	AA	1469	G	P-O5'	-5.24	1.54	1.59
85	AA	2199	G	C1'-N9	-5.24	1.39	1.46
34	BA	22	C	C4'-C3'	-5.24	1.47	1.52
37	BD	77	A	C8-N7	-5.24	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	495	G	C8-N7	-5.24	1.27	1.30
85	AA	1157	U	C4'-C3'	-5.24	1.47	1.52
85	AA	1184	A	C2'-C1'	-5.24	1.47	1.53
85	AA	1569	C	O3'-P	-5.24	1.54	1.61
85	AA	1587	C	C3'-C2'	-5.24	1.47	1.52
85	AA	1589	G	C3'-C2'	-5.24	1.47	1.52
25	AR	14	VAL	N-CA	-5.24	1.35	1.46
34	BA	248	G	C5-C6	-5.24	1.37	1.42
34	BA	857	C	N1-C6	-5.24	1.34	1.37
34	BA	934	G	C1'-N9	-5.24	1.39	1.46
34	BA	945	A	O3'-P	-5.24	1.54	1.61
34	BA	1064	A	N3-C4	-5.24	1.31	1.34
35	BB	445	G	C6-N1	-5.24	1.35	1.39
35	BB	543	G	N7-C5	-5.24	1.36	1.39
35	BB	1374	U	C2'-C1'	-5.24	1.47	1.53
35	BB	1397	G	N1-C2	-5.24	1.33	1.37
35	BB	1416	A	C1'-N9	-5.24	1.39	1.46
36	BC	42	G	C3'-C2'	-5.24	1.47	1.52
40	BG	9	G	C2-N3	-5.24	1.28	1.32
40	BG	34	A	O4'-C1'	-5.24	1.34	1.41
41	BH	17	A	O4'-C1'	-5.24	1.34	1.41
85	AA	431	G	C2-N2	-5.24	1.29	1.34
85	AA	500	C	P-O5'	-5.24	1.54	1.59
85	AA	1352	U	C5'-C4'	5.24	1.57	1.51
85	AA	1754	G	O3'-P	-5.24	1.54	1.61
85	AA	2127	G	C4'-C3'	-5.24	1.47	1.52
85	AA	2225	G	C2-N2	-5.24	1.29	1.34
34	BA	401	A	C3'-C2'	-5.24	1.47	1.52
34	BA	1029	C	C2'-C1'	-5.24	1.47	1.53
34	BA	1142	C	C3'-C2'	-5.24	1.47	1.52
34	BA	1707	C	O4'-C1'	-5.24	1.34	1.41
35	BB	60	A	C5'-C4'	-5.24	1.45	1.51
35	BB	1184	C	O3'-P	-5.24	1.54	1.61
35	BB	1355	C	C2'-C1'	-5.24	1.47	1.53
35	BB	1480	G	O4'-C1'	-5.24	1.34	1.41
36	BC	6	G	C2-N3	-5.24	1.28	1.32
38	BE	121	G	N3-C4	-5.24	1.31	1.35
38	BE	178	G	C2'-C1'	-5.24	1.47	1.53
41	BH	45	G	C5-C6	-5.24	1.37	1.42
85	AA	1134	G	C2'-C1'	-5.24	1.47	1.53
85	AA	1134	G	C5-C4	-5.24	1.34	1.38
85	AA	1485	G	P-O5'	-5.24	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1732	G	N9-C4	-5.24	1.33	1.38
34	BA	362	G	C3'-C2'	-5.24	1.47	1.52
34	BA	454	G	C2'-C1'	-5.24	1.47	1.53
34	BA	1028	A	C5-C6	-5.24	1.36	1.41
35	BB	1020	U	N3-C4	-5.24	1.33	1.38
36	BC	119	G	C5-C6	-5.24	1.37	1.42
40	BG	17	A	N3-C4	-5.24	1.31	1.34
85	AA	173	A	C4'-O4'	-5.24	1.38	1.45
85	AA	1236	G	N3-C4	-5.24	1.31	1.35
85	AA	2083	G	C2'-C1'	-5.24	1.47	1.53
85	AA	2143	U	N3-C4	-5.24	1.33	1.38
34	BA	62	A	C2'-C1'	-5.24	1.47	1.53
34	BA	458	G	C2'-C1'	-5.24	1.47	1.53
34	BA	628	U	O3'-P	-5.24	1.54	1.61
34	BA	1182	U	C2'-C1'	-5.24	1.47	1.53
34	BA	1451	A	N3-C4	-5.24	1.31	1.34
34	BA	1502	G	C3'-C2'	-5.24	1.47	1.52
35	BB	428	G	C5-C4	-5.24	1.34	1.38
35	BB	651	G	C5-C4	-5.24	1.34	1.38
35	BB	1416	A	O3'-P	-5.24	1.54	1.61
36	BC	14	G	C5-C4	-5.24	1.34	1.38
37	BD	84	U	O3'-P	-5.24	1.54	1.61
37	BD	101	A	C8-N7	-5.24	1.27	1.31
38	BE	195	G	C5-C4	-5.24	1.34	1.38
39	BF	65	U	C3'-C2'	-5.24	1.47	1.52
40	BG	47	G	N9-C4	-5.24	1.33	1.38
85	AA	303	A	C2'-C1'	-5.24	1.47	1.53
85	AA	624	A	N1-C2	-5.24	1.29	1.34
85	AA	2092	A	C3'-C2'	-5.24	1.47	1.52
85	AA	2175	U	O4'-C1'	-5.24	1.34	1.41
85	AA	2245	A	C2'-C1'	-5.24	1.47	1.53
34	BA	236	A	C2'-O2'	-5.23	1.34	1.41
34	BA	1016	A	C6-N6	-5.23	1.29	1.33
35	BB	568	A	C8-N7	-5.23	1.27	1.31
37	BD	72	U	C4'-O4'	-5.23	1.38	1.45
40	BG	73	U	O4'-C1'	-5.23	1.34	1.41
85	AA	307	G	N3-C4	-5.23	1.31	1.35
85	AA	2144	C	O3'-P	-5.23	1.54	1.61
86	AB	67	C	P-O5'	-5.23	1.54	1.59
33	AZ	41	GLY	CA-C	-5.23	1.43	1.51
34	BA	112	C	P-O5'	-5.23	1.54	1.59
34	BA	474	A	C8-N7	-5.23	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	815	C	C4'-C3'	-5.23	1.47	1.52
34	BA	850	C	C2-N3	-5.23	1.31	1.35
34	BA	1093	G	C2-N2	-5.23	1.29	1.34
34	BA	1165	A	C8-N7	-5.23	1.27	1.31
34	BA	1227	U	C2'-C1'	-5.23	1.47	1.53
34	BA	1413	G	C3'-C2'	-5.23	1.47	1.52
34	BA	1598	U	C2'-C1'	-5.23	1.47	1.53
35	BB	269	A	O3'-P	-5.23	1.54	1.61
35	BB	795	A	C2'-C1'	-5.23	1.47	1.53
35	BB	1099	U	P-O5'	-5.23	1.54	1.59
35	BB	1160	U	C4'-C3'	-5.23	1.47	1.52
35	BB	1257	A	P-O5'	-5.23	1.54	1.59
35	BB	1352	C	O3'-P	-5.23	1.54	1.61
36	BC	4	G	C2-N2	-5.23	1.29	1.34
36	BC	91	G	N7-C5	-5.23	1.36	1.39
38	BE	31	A	O3'-P	-5.23	1.54	1.61
38	BE	204	U	P-O5'	-5.23	1.54	1.59
85	AA	15	U	O3'-P	-5.23	1.54	1.61
85	AA	404	A	C3'-C2'	-5.23	1.47	1.52
85	AA	542	G	N7-C5	-5.23	1.36	1.39
34	BA	83	G	N1-C2	-5.23	1.33	1.37
34	BA	118	C	C4-N4	-5.23	1.29	1.33
34	BA	433	G	N9-C4	-5.23	1.33	1.38
34	BA	537	C	C3'-C2'	-5.23	1.47	1.52
34	BA	689	C	C4-N4	-5.23	1.29	1.33
34	BA	932	G	C2-N2	-5.23	1.29	1.34
34	BA	1440	C	N3-C4	-5.23	1.30	1.33
34	BA	1658	G	C2'-C1'	-5.23	1.47	1.53
35	BB	4	C	C3'-O3'	5.23	1.49	1.42
35	BB	41	A	C6-N1	5.23	1.39	1.35
35	BB	429	C	C2-N3	-5.23	1.31	1.35
35	BB	1351	G	N9-C8	-5.23	1.34	1.37
35	BB	1385	C	C2-N3	-5.23	1.31	1.35
40	BG	29	U	C2-N3	-5.23	1.34	1.37
41	BH	11	C	C2'-C1'	-5.23	1.47	1.53
85	AA	390	U	C2-N3	-5.23	1.34	1.37
85	AA	808	A	C4'-O4'	5.23	1.52	1.45
85	AA	1624	U	C1'-N1	-5.23	1.39	1.46
34	BA	130	U	N3-C4	-5.23	1.33	1.38
34	BA	238	C	C5-C6	-5.23	1.30	1.34
34	BA	763	U	C2'-C1'	5.23	1.59	1.53
34	BA	926	A	O4'-C1'	-5.23	1.34	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1560	U	P-O5'	-5.23	1.54	1.59
37	BD	16	U	N3-C4	-5.23	1.33	1.38
38	BE	121	G	C2-N2	-5.23	1.29	1.34
39	BF	36	G	C5-C6	-5.23	1.37	1.42
40	BG	18	U	C2'-C1'	-5.23	1.47	1.53
40	BG	52	A	C8-N7	-5.23	1.27	1.31
41	BH	6	U	N3-C4	-5.23	1.33	1.38
85	AA	157	G	O4'-C1'	-5.23	1.34	1.41
34	BA	73	G	C5-C4	-5.23	1.34	1.38
34	BA	296	G	C1'-N9	5.23	1.56	1.48
34	BA	478	G	N1-C2	-5.23	1.33	1.37
34	BA	774	A	C4'-O4'	-5.23	1.38	1.45
34	BA	818	G	C6-N1	-5.23	1.35	1.39
34	BA	841	G	C2'-C1'	-5.23	1.47	1.53
34	BA	853	A	N1-C2	-5.23	1.29	1.34
34	BA	861	C	O3'-P	-5.23	1.54	1.61
34	BA	956	G	C1'-N9	-5.23	1.39	1.46
34	BA	1090	A	N7-C5	-5.23	1.36	1.39
34	BA	1409	A	C6-N1	-5.23	1.31	1.35
34	BA	1595	G	C5-C4	-5.23	1.34	1.38
35	BB	9	G	C8-N7	-5.23	1.27	1.30
35	BB	121	A	N1-C2	-5.23	1.29	1.34
35	BB	1351	G	N9-C4	-5.23	1.33	1.38
35	BB	1363	A	C2'-C1'	-5.23	1.47	1.53
36	BC	109	A	N9-C8	-5.23	1.33	1.37
37	BD	54	A	C5-C4	-5.23	1.35	1.38
37	BD	69	U	C1'-N1	-5.23	1.39	1.46
38	BE	13	A	P-O5'	5.23	1.65	1.59
40	BG	129	G	C5'-C4'	-5.23	1.45	1.51
85	AA	539	A	N7-C5	-5.23	1.36	1.39
85	AA	1689	G	P-O5'	-5.23	1.54	1.59
85	AA	1872	G	C5-C6	-5.23	1.37	1.42
34	BA	36	A	C2'-C1'	-5.23	1.47	1.53
34	BA	859	G	C2-N3	-5.23	1.28	1.32
35	BB	572	G	N3-C4	-5.23	1.31	1.35
85	AA	1516	A	C3'-O3'	-5.23	1.34	1.42
85	AA	1904	C	C4'-C3'	-5.23	1.47	1.52
34	BA	36	A	C1'-N9	-5.22	1.39	1.46
34	BA	454	G	N9-C4	-5.22	1.33	1.38
34	BA	1276	G	C6-N1	-5.22	1.35	1.39
34	BA	1433	U	O3'-P	-5.22	1.54	1.61
34	BA	1442	A	C2'-C1'	-5.22	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1444	G	C4'-C3'	5.22	1.58	1.53
34	BA	1481	U	O3'-P	-5.22	1.54	1.61
34	BA	1688	G	N3-C4	-5.22	1.31	1.35
35	BB	85	A	N3-C4	-5.22	1.31	1.34
35	BB	111	C	C3'-C2'	-5.22	1.47	1.52
35	BB	375	G	C3'-C2'	-5.22	1.47	1.52
35	BB	692	G	C2-N2	-5.22	1.29	1.34
35	BB	875	G	C5'-C4'	5.22	1.57	1.51
35	BB	1515	C	N1-C6	-5.22	1.34	1.37
38	BE	85	G	C5-C4	-5.22	1.34	1.38
40	BG	90	G	C3'-C2'	-5.22	1.47	1.52
41	BH	115	A	C8-N7	-5.22	1.27	1.31
61	Bb	106	TYR	CB-CG	-5.22	1.43	1.51
85	AA	416	U	N3-C4	-5.22	1.33	1.38
85	AA	436	G	C2'-C1'	-5.22	1.47	1.53
85	AA	764	U	N1-C2	-5.22	1.33	1.38
85	AA	802	A	C1'-N9	-5.22	1.39	1.46
85	AA	1197	U	C3'-C2'	-5.22	1.47	1.52
34	BA	575	U	C1'-N1	-5.22	1.39	1.46
34	BA	584	A	C5-C4	-5.22	1.35	1.38
34	BA	588	C	N1-C6	-5.22	1.34	1.37
34	BA	1704	G	N1-C2	-5.22	1.33	1.37
35	BB	673	C	C4'-C3'	-5.22	1.47	1.52
35	BB	1536	G	O3'-P	-5.22	1.54	1.61
36	BC	33	U	C1'-N1	5.22	1.56	1.48
54	BU	8	LYS	N-CA	-5.22	1.35	1.46
85	AA	395	G	C1'-N9	-5.22	1.39	1.46
85	AA	1517	G	C4'-C3'	-5.22	1.47	1.52
85	AA	1701	G	C5-C4	-5.22	1.34	1.38
85	AA	2034	G	O3'-P	-5.22	1.54	1.61
34	BA	192	G	C2'-C1'	-5.22	1.47	1.53
34	BA	619	U	C2'-C1'	-5.22	1.47	1.53
34	BA	1250	C	N3-C4	-5.22	1.30	1.33
35	BB	481	A	P-O5'	-5.22	1.54	1.59
36	BC	60	U	C2'-C1'	-5.22	1.47	1.53
36	BC	106	G	N9-C8	-5.22	1.34	1.37
36	BC	154	A	N7-C5	-5.22	1.36	1.39
36	BC	163	A	N9-C8	-5.22	1.33	1.37
40	BG	70	C	N3-C4	-5.22	1.30	1.33
41	BH	38	G	C4'-C3'	-5.22	1.47	1.52
85	AA	246	C	C2'-C1'	-5.22	1.47	1.53
85	AA	506	G	C2-N2	-5.22	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	513	G	C6-N1	-5.22	1.35	1.39
85	AA	913	U	N1-C2	-5.22	1.33	1.38
85	AA	1650	G	N9-C4	-5.22	1.33	1.38
85	AA	1698	A	C8-N7	-5.22	1.27	1.31
34	BA	1	C	O3'-P	-5.22	1.54	1.61
34	BA	439	A	P-O5'	-5.22	1.54	1.59
34	BA	719	G	N1-C2	-5.22	1.33	1.37
34	BA	1528	U	O3'-P	-5.22	1.54	1.61
34	BA	1535	G	O3'-P	-5.22	1.54	1.61
35	BB	467	G	C6-N1	-5.22	1.35	1.39
35	BB	590	G	N9-C4	-5.22	1.33	1.38
35	BB	679	G	C2-N2	-5.22	1.29	1.34
35	BB	954	G	C2'-C1'	-5.22	1.47	1.53
35	BB	1045	G	N9-C4	-5.22	1.33	1.38
35	BB	1313	C	C4-N4	-5.22	1.29	1.33
35	BB	1503	U	C5'-C4'	5.22	1.57	1.51
36	BC	54	G	C2-N2	-5.22	1.29	1.34
40	BG	58	G	C2-N2	-5.22	1.29	1.34
85	AA	19	A	C8-N7	-5.22	1.27	1.31
85	AA	779	G	C4'-C3'	-5.22	1.47	1.52
85	AA	1220	A	C2'-C1'	-5.22	1.47	1.53
34	BA	303	C	C4'-C3'	-5.22	1.47	1.52
34	BA	496	G	N1-C2	-5.22	1.33	1.37
34	BA	1563	G	O3'-P	-5.22	1.54	1.61
35	BB	1071	G	C2'-C1'	-5.22	1.47	1.53
35	BB	1455	A	N9-C8	-5.22	1.33	1.37
39	BF	72	A	N9-C4	-5.22	1.34	1.37
40	BG	54	G	C2-N2	-5.22	1.29	1.34
40	BG	100	G	C2-N2	-5.22	1.29	1.34
85	AA	93	G	N3-C4	-5.22	1.31	1.35
85	AA	1100	U	N1-C2	5.22	1.43	1.38
85	AA	1128	G	C5-C4	-5.22	1.34	1.38
34	BA	352	G	C5-C4	-5.22	1.34	1.38
34	BA	910	U	C2'-C1'	-5.22	1.47	1.53
34	BA	1234	U	C1'-N1	-5.22	1.39	1.46
34	BA	1330	G	C4'-C3'	5.22	1.58	1.53
34	BA	1711	G	C2'-C1'	-5.22	1.47	1.53
35	BB	681	G	C2-N2	-5.22	1.29	1.34
35	BB	830	G	P-O5'	-5.22	1.54	1.59
35	BB	1038	G	C5-C6	-5.22	1.37	1.42
35	BB	1108	G	C2'-C1'	-5.22	1.47	1.53
35	BB	1391	G	C5-C4	-5.22	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	30	A	O3'-P	-5.22	1.54	1.61
40	BG	134	U	N1-C6	-5.22	1.33	1.38
40	BG	150	A	C5-C6	-5.22	1.36	1.41
85	AA	606	A	O3'-P	-5.22	1.54	1.61
85	AA	642	G	C1'-N9	-5.22	1.39	1.46
85	AA	644	A	C2'-C1'	-5.22	1.47	1.53
85	AA	1245	U	C2-N3	-5.22	1.34	1.37
85	AA	2046	G	C6-N1	-5.22	1.35	1.39
34	BA	315	U	C3'-C2'	-5.21	1.47	1.52
34	BA	894	G	C3'-C2'	-5.21	1.47	1.52
34	BA	1192	A	C4'-C3'	-5.21	1.47	1.52
34	BA	1511	C	C4'-C3'	-5.21	1.47	1.52
34	BA	1637	G	N9-C4	5.21	1.42	1.38
34	BA	1787	U	C2-N3	-5.21	1.34	1.37
35	BB	833	G	C1'-N9	-5.21	1.39	1.46
35	BB	1380	G	O3'-P	-5.21	1.54	1.61
35	BB	1469	A	P-O5'	-5.21	1.54	1.59
36	BC	63	G	C3'-C2'	-5.21	1.47	1.52
38	BE	59	U	C4'-C3'	-5.21	1.47	1.52
38	BE	60	C	C4-N4	-5.21	1.29	1.33
39	BF	48	G	C5-C6	-5.21	1.37	1.42
41	BH	106	G	C2-N2	-5.21	1.29	1.34
63	Bd	36	GLY	C-N	5.21	1.46	1.34
85	AA	62	A	O3'-P	-5.21	1.54	1.61
85	AA	123	A	O3'-P	-5.21	1.54	1.61
85	AA	126	U	N1-C2	-5.21	1.33	1.38
85	AA	269	G	C5'-C4'	-5.21	1.45	1.51
85	AA	435	A	C1'-N9	-5.21	1.39	1.46
85	AA	528	U	C4'-O4'	-5.21	1.38	1.45
85	AA	1983	C	O3'-P	-5.21	1.54	1.61
85	AA	2017	U	C4'-C3'	-5.21	1.47	1.52
86	AB	59	U	O3'-P	-5.21	1.54	1.61
34	BA	878	G	C4'-O4'	-5.21	1.38	1.45
35	BB	703	U	C1'-N1	-5.21	1.39	1.46
35	BB	1028	C	C4'-O4'	-5.21	1.38	1.45
35	BB	1334	C	C4-N4	-5.21	1.29	1.33
37	BD	58	G	N1-C2	-5.21	1.33	1.37
85	AA	157	G	C5'-C4'	-5.21	1.45	1.51
85	AA	394	C	C2'-C1'	-5.21	1.47	1.53
85	AA	797	C	P-O5'	-5.21	1.54	1.59
85	AA	1672	G	C5-C4	-5.21	1.34	1.38
85	AA	2112	G	N7-C5	-5.21	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	2	A	C4'-C3'	-5.21	1.47	1.52
34	BA	460	G	N9-C4	-5.21	1.33	1.38
34	BA	798	G	C4'-C3'	-5.21	1.47	1.52
34	BA	1008	A	N7-C5	-5.21	1.36	1.39
35	BB	102	G	C5-C6	-5.21	1.37	1.42
35	BB	378	C	C2'-C1'	-5.21	1.47	1.53
35	BB	593	A	N3-C4	-5.21	1.31	1.34
35	BB	811	C	O3'-P	-5.21	1.54	1.61
35	BB	1512	C	C1'-N1	5.21	1.56	1.48
41	BH	91	G	C8-N7	5.21	1.34	1.30
85	AA	276	C	C4-N4	-5.21	1.29	1.33
85	AA	692	U	C5'-C4'	-5.21	1.45	1.51
85	AA	2136	C	O3'-P	-5.21	1.54	1.61
34	BA	403	A	N9-C4	-5.21	1.34	1.37
34	BA	546	U	C2-N3	-5.21	1.34	1.37
35	BB	109	U	N1-C6	-5.21	1.33	1.38
35	BB	406	A	P-O5'	-5.21	1.54	1.59
35	BB	1509	G	C2-N2	-5.21	1.29	1.34
38	BE	175	U	C4'-C3'	5.21	1.58	1.53
41	BH	26	C	O3'-P	-5.21	1.54	1.61
85	AA	488	G	C1'-N9	-5.21	1.39	1.46
34	BA	375	C	C3'-C2'	-5.21	1.47	1.52
34	BA	889	U	C4'-C3'	-5.21	1.47	1.52
34	BA	999	G	C2-N2	-5.21	1.29	1.34
34	BA	1238	C	C2'-C1'	-5.21	1.47	1.53
34	BA	1512	C	C2-N3	-5.21	1.31	1.35
34	BA	1574	C	C4-N4	-5.21	1.29	1.33
35	BB	101	U	O4'-C1'	-5.21	1.34	1.41
35	BB	841	U	N1-C6	-5.21	1.33	1.38
35	BB	1187	G	P-O5'	-5.21	1.54	1.59
35	BB	1220	A	N9-C8	-5.21	1.33	1.37
35	BB	1430	G	O4'-C1'	-5.21	1.34	1.41
37	BD	69	U	N1-C6	-5.21	1.33	1.38
38	BE	163	A	C5-C6	-5.21	1.36	1.41
39	BF	56	C	N3-C4	5.21	1.37	1.33
40	BG	135	C	C4'-C3'	-5.21	1.47	1.52
41	BH	54	U	N3-C4	-5.21	1.33	1.38
41	BH	68	G	C2-N2	-5.21	1.29	1.34
85	AA	457	G	C2'-C1'	-5.21	1.47	1.53
85	AA	648	G	N9-C8	-5.21	1.34	1.37
85	AA	732	G	P-O5'	-5.21	1.54	1.59
85	AA	1918	U	C2'-C1'	-5.21	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	181	G	N3-C4	-5.21	1.31	1.35
34	BA	540	G	C2'-C1'	-5.21	1.47	1.53
34	BA	654	C	O3'-P	-5.21	1.54	1.61
34	BA	689	C	C1'-N1	-5.21	1.39	1.46
34	BA	755	G	C5'-C4'	-5.21	1.45	1.51
34	BA	807	U	C2-N3	-5.21	1.34	1.37
34	BA	1007	G	C2'-C1'	-5.21	1.47	1.53
34	BA	1227	U	P-O5'	-5.21	1.54	1.59
34	BA	1509	U	C2'-C1'	-5.21	1.47	1.53
34	BA	1597	G	C8-N7	-5.21	1.27	1.30
35	BB	32	C	C3'-C2'	-5.21	1.47	1.52
35	BB	68	G	N7-C5	-5.21	1.36	1.39
35	BB	348	G	P-O5'	-5.21	1.54	1.59
35	BB	1224	C	N1-C6	-5.21	1.34	1.37
35	BB	1440	A	N3-C4	-5.21	1.31	1.34
35	BB	1529	G	N7-C5	-5.21	1.36	1.39
36	BC	69	U	C1'-N1	-5.21	1.39	1.46
40	BG	78	C	N1-C6	-5.21	1.34	1.37
40	BG	180	C	C4'-O4'	-5.21	1.38	1.45
56	BW	97	PHE	CB-CG	-5.21	1.42	1.51
85	AA	11	A	C6-N1	-5.21	1.31	1.35
85	AA	49	C	C3'-C2'	-5.21	1.47	1.52
85	AA	119	G	N9-C8	-5.21	1.34	1.37
85	AA	611	G	N9-C4	5.21	1.42	1.38
85	AA	995	G	C4'-C3'	-5.21	1.47	1.52
85	AA	1508	A	C3'-C2'	-5.21	1.47	1.52
85	AA	1536	C	O5'-C5'	5.21	1.52	1.44
85	AA	1788	U	P-O5'	-5.21	1.54	1.59
85	AA	1854	U	P-O5'	-5.21	1.54	1.59
34	BA	19	G	N1-C2	-5.21	1.33	1.37
34	BA	371	U	N3-C4	-5.21	1.33	1.38
35	BB	108	G	C2'-C1'	-5.21	1.47	1.53
36	BC	70	C	P-O5'	-5.21	1.54	1.59
61	Bb	133	LYS	CA-C	-5.21	1.39	1.52
85	AA	323	U	C3'-C2'	-5.21	1.47	1.52
85	AA	968	U	C3'-C2'	-5.21	1.47	1.52
85	AA	1877	G	C6-N1	-5.21	1.35	1.39
85	AA	2187	G	N1-C2	-5.21	1.33	1.37
34	BA	386	A	C2'-C1'	-5.20	1.47	1.53
34	BA	1094	U	C2'-C1'	-5.20	1.47	1.53
34	BA	1248	A	C2'-C1'	-5.20	1.47	1.53
34	BA	1252	G	C4'-C3'	-5.20	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1674	G	C2-N2	-5.20	1.29	1.34
35	BB	66	G	C5-C6	-5.20	1.37	1.42
35	BB	514	G	O3'-P	-5.20	1.54	1.61
35	BB	1123	A	C5-C4	-5.20	1.35	1.38
40	BG	61	A	O3'-P	-5.20	1.54	1.61
85	AA	19	A	N3-C4	-5.20	1.31	1.34
85	AA	938	A	C8-N7	-5.20	1.27	1.31
85	AA	1305	A	C2'-C1'	-5.20	1.47	1.53
85	AA	2072	G	C4'-C3'	-5.20	1.47	1.52
34	BA	291	C	C4'-C3'	-5.20	1.47	1.52
34	BA	828	A	C3'-C2'	-5.20	1.47	1.52
34	BA	1159	A	N9-C4	-5.20	1.34	1.37
36	BC	98	C	O3'-P	-5.20	1.54	1.61
85	AA	1223	A	C3'-C2'	-5.20	1.47	1.52
85	AA	2028	G	C6-N1	-5.20	1.35	1.39
85	AA	2069	A	C6-N6	-5.20	1.29	1.33
34	BA	12	G	C6-N1	-5.20	1.35	1.39
34	BA	666	C	C4-C5	-5.20	1.38	1.43
34	BA	1253	G	C1'-N9	-5.20	1.39	1.46
34	BA	1273	U	C4'-O4'	-5.20	1.38	1.45
35	BB	101	U	C1'-N1	-5.20	1.39	1.46
35	BB	374	A	C6-N6	-5.20	1.29	1.33
35	BB	1185	G	O4'-C1'	-5.20	1.34	1.41
35	BB	1524	G	C6-N1	-5.20	1.35	1.39
37	BD	65	G	C1'-N9	-5.20	1.39	1.46
39	BF	32	G	C2-N2	-5.20	1.29	1.34
85	AA	317	A	N7-C5	-5.20	1.36	1.39
85	AA	671	G	C2'-C1'	-5.20	1.47	1.53
85	AA	1143	C	P-O5'	-5.20	1.54	1.59
85	AA	2240	G	C5-C4	-5.20	1.34	1.38
34	BA	38	G	N9-C4	-5.20	1.33	1.38
34	BA	320	G	C6-N1	-5.20	1.35	1.39
34	BA	472	G	C5-C6	-5.20	1.37	1.42
34	BA	1287	G	C4'-C3'	-5.20	1.47	1.52
34	BA	1310	C	C3'-C2'	-5.20	1.47	1.52
34	BA	1406	U	N3-C4	-5.20	1.33	1.38
34	BA	1665	G	C3'-C2'	-5.20	1.47	1.52
35	BB	103	C	C2'-C1'	-5.20	1.47	1.53
35	BB	830	G	C2-N2	-5.20	1.29	1.34
35	BB	1181	A	N3-C4	-5.20	1.31	1.34
35	BB	1222	A	C8-N7	-5.20	1.27	1.31
35	BB	1365	G	C5-C4	-5.20	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1501	U	P-O5'	-5.20	1.54	1.59
39	BF	57	C	N1-C2	-5.20	1.34	1.40
40	BG	7	U	C2'-C1'	-5.20	1.47	1.53
40	BG	144	G	C2'-C1'	-5.20	1.47	1.53
41	BH	114	G	C5-C6	-5.20	1.37	1.42
85	AA	57	G	O3'-P	-5.20	1.54	1.61
85	AA	182	C	C4'-C3'	-5.20	1.47	1.52
85	AA	430	G	C5-C6	-5.20	1.37	1.42
85	AA	875	C	O3'-P	-5.20	1.54	1.61
85	AA	1172	A	C2'-C1'	-5.20	1.47	1.53
85	AA	1485	G	C3'-C2'	-5.20	1.47	1.52
85	AA	1921	G	C6-N1	5.20	1.43	1.39
34	BA	486	G	O3'-P	-5.20	1.54	1.61
34	BA	1063	G	N9-C4	-5.20	1.33	1.38
34	BA	1269	C	C4-N4	-5.20	1.29	1.33
34	BA	1276	G	N1-C2	-5.20	1.33	1.37
34	BA	1687	A	C5-C4	-5.20	1.35	1.38
35	BB	623	A	O3'-P	-5.20	1.54	1.61
85	AA	549	A	C2'-C1'	-5.20	1.47	1.53
85	AA	1007	G	P-O5'	-5.20	1.54	1.59
85	AA	1234	G	C2'-C1'	-5.20	1.47	1.53
85	AA	1564	U	C2-N3	-5.20	1.34	1.37
85	AA	2044	A	N3-C4	-5.20	1.31	1.34
85	AA	2044	A	P-O5'	-5.20	1.54	1.59
85	AA	2170	G	C3'-C2'	-5.20	1.47	1.52
34	BA	63	A	C1'-N9	-5.20	1.39	1.46
34	BA	289	A	N1-C2	-5.20	1.29	1.34
34	BA	461	A	O3'-P	-5.20	1.54	1.61
34	BA	466	G	N7-C5	-5.20	1.36	1.39
34	BA	684	G	N9-C4	5.20	1.42	1.38
34	BA	704	G	C2-N2	-5.20	1.29	1.34
34	BA	1498	A	C4'-C3'	-5.20	1.47	1.52
35	BB	25	A	O4'-C1'	-5.20	1.34	1.41
35	BB	529	A	C5'-C4'	5.20	1.57	1.51
35	BB	632	U	N3-C4	-5.20	1.33	1.38
35	BB	1003	G	C3'-C2'	-5.20	1.47	1.52
35	BB	1027	U	C3'-C2'	-5.20	1.47	1.52
35	BB	1159	U	C4'-O4'	-5.20	1.38	1.45
35	BB	1476	C	C2-N3	-5.20	1.31	1.35
85	AA	665	A	C5-C4	-5.20	1.35	1.38
85	AA	687	G	N3-C4	-5.20	1.31	1.35
85	AA	1629	C	C2'-C1'	-5.20	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1864	G	C2'-C1'	-5.20	1.47	1.53
34	BA	1277	G	C2-N2	-5.19	1.29	1.34
35	BB	560	C	C2-N3	-5.19	1.31	1.35
35	BB	1028	C	P-O5'	-5.19	1.54	1.59
35	BB	1186	A	C2'-C1'	-5.19	1.47	1.53
42	BI	177	GLY	CA-C	-5.19	1.43	1.51
52	BS	21	ASN	C-N	-5.19	1.24	1.34
52	BS	44	TRP	CB-CG	-5.19	1.41	1.50
85	AA	188	G	C1'-N9	-5.19	1.39	1.46
85	AA	391	G	C6-N1	-5.19	1.35	1.39
34	BA	87	G	O3'-P	-5.19	1.54	1.61
34	BA	194	G	P-O5'	-5.19	1.54	1.59
34	BA	269	G	C5-C4	-5.19	1.34	1.38
34	BA	412	G	C8-N7	-5.19	1.27	1.30
34	BA	812	A	C1'-N9	-5.19	1.39	1.46
34	BA	1095	G	C2'-C1'	-5.19	1.47	1.53
35	BB	23	U	O3'-P	-5.19	1.54	1.61
35	BB	386	G	C2-N2	-5.19	1.29	1.34
35	BB	1256	C	P-O5'	-5.19	1.54	1.59
40	BG	15	G	C4'-C3'	-5.19	1.47	1.52
40	BG	139	U	C4'-C3'	-5.19	1.47	1.52
85	AA	453	G	C2'-C1'	-5.19	1.47	1.53
85	AA	814	G	C2'-C1'	-5.19	1.47	1.53
85	AA	1429	U	O3'-P	-5.19	1.54	1.61
5	A4	68	TYR	CB-CG	-5.19	1.43	1.51
34	BA	37	A	N3-C4	-5.19	1.31	1.34
34	BA	515	U	N1-C6	-5.19	1.33	1.38
34	BA	627	U	C2'-C1'	-5.19	1.47	1.53
34	BA	1095	G	N1-C2	-5.19	1.33	1.37
34	BA	1142	C	P-O5'	-5.19	1.54	1.59
35	BB	637	G	N1-C2	-5.19	1.33	1.37
35	BB	793	A	N7-C5	-5.19	1.36	1.39
35	BB	1186	A	C5-C4	-5.19	1.35	1.38
36	BC	66	G	N7-C5	-5.19	1.36	1.39
85	AA	53	G	C6-N1	-5.19	1.35	1.39
85	AA	129	U	N3-C4	-5.19	1.33	1.38
85	AA	510	A	C5'-C4'	5.19	1.57	1.51
85	AA	1467	U	O4'-C1'	-5.19	1.34	1.41
85	AA	1802	U	O3'-P	-5.19	1.54	1.61
85	AA	2074	G	C2'-C1'	-5.19	1.47	1.53
85	AA	2074	G	C3'-C2'	-5.19	1.47	1.52
35	BB	94	A	P-O5'	-5.19	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	61	A	N7-C5	-5.19	1.36	1.39
40	BG	36	G	C3'-C2'	-5.19	1.47	1.52
85	AA	188	G	O3'-P	-5.19	1.54	1.61
85	AA	713	G	C4'-C3'	-5.19	1.47	1.52
34	BA	920	U	O3'-P	-5.19	1.54	1.61
34	BA	921	G	C2'-C1'	-5.19	1.47	1.53
34	BA	1219	G	N1-C2	-5.19	1.33	1.37
34	BA	1357	C	C2'-C1'	-5.19	1.47	1.53
34	BA	1684	A	C5-C4	-5.19	1.35	1.38
35	BB	67	A	C1'-N9	-5.19	1.39	1.46
35	BB	1271	A	N9-C4	-5.19	1.34	1.37
35	BB	1360	A	C3'-C2'	-5.19	1.47	1.52
35	BB	1392	A	C4'-O4'	-5.19	1.38	1.45
37	BD	42	A	C3'-C2'	-5.19	1.47	1.52
37	BD	44	U	C3'-C2'	-5.19	1.47	1.52
38	BE	67	A	N9-C8	-5.19	1.33	1.37
85	AA	19	A	N7-C5	-5.19	1.36	1.39
85	AA	122	A	C5-C4	-5.19	1.35	1.38
85	AA	428	G	C5-C6	-5.19	1.37	1.42
85	AA	615	A	P-O5'	-5.19	1.54	1.59
85	AA	864	C	C2-N3	-5.19	1.31	1.35
85	AA	1187	G	C5-C4	-5.19	1.34	1.38
85	AA	1866	A	P-O5'	-5.19	1.54	1.59
85	AA	1900	C	C4'-C3'	-5.19	1.47	1.52
86	AB	6	G	C6-N1	-5.19	1.35	1.39
34	BA	533	U	C1'-N1	-5.19	1.39	1.46
34	BA	1256	A	C3'-C2'	-5.19	1.47	1.52
34	BA	1614	G	C6-O6	-5.19	1.19	1.24
35	BB	376	A	N7-C5	-5.19	1.36	1.39
35	BB	829	C	C1'-N1	-5.19	1.39	1.46
35	BB	864	U	P-O5'	-5.19	1.54	1.59
74	Bo	21	ASN	C-N	-5.19	1.24	1.34
85	AA	374	C	O5'-C5'	-5.19	1.34	1.42
11	AC	124	PHE	CB-CG	-5.18	1.42	1.51
34	BA	382	G	N7-C5	-5.18	1.36	1.39
34	BA	906	A	C1'-N9	-5.18	1.39	1.46
34	BA	1090	A	N9-C8	-5.18	1.33	1.37
34	BA	1170	A	O4'-C1'	-5.18	1.34	1.41
35	BB	121	A	N9-C4	-5.18	1.34	1.37
35	BB	430	A	N3-C4	-5.18	1.31	1.34
35	BB	825	U	C3'-C2'	-5.18	1.47	1.52
35	BB	1134	G	N1-C2	-5.18	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1428	C	C4'-C3'	-5.18	1.47	1.52
35	BB	1474	A	C3'-O3'	5.18	1.49	1.42
37	BD	66	G	C6-N1	-5.18	1.35	1.39
37	BD	71	G	C3'-C2'	-5.18	1.47	1.52
37	BD	107	G	C5-C4	-5.18	1.34	1.38
38	BE	2	G	N9-C8	-5.18	1.34	1.37
39	BF	4	A	C2'-C1'	-5.18	1.47	1.53
41	BH	3	U	O4'-C1'	-5.18	1.34	1.41
41	BH	23	G	N9-C4	-5.18	1.33	1.38
85	AA	596	A	C4'-C3'	-5.18	1.47	1.52
85	AA	1942	U	O3'-P	-5.18	1.54	1.61
85	AA	2002	A	C3'-C2'	-5.18	1.47	1.52
85	AA	2121	G	C2-N2	-5.18	1.29	1.34
85	AA	2125	A	C2'-O2'	-5.18	1.34	1.41
34	BA	483	A	O5'-C5'	-5.18	1.34	1.42
34	BA	613	A	C5-C4	-5.18	1.35	1.38
34	BA	696	A	O4'-C1'	-5.18	1.34	1.41
34	BA	723	C	C3'-C2'	-5.18	1.47	1.52
34	BA	1255	G	C2'-C1'	-5.18	1.47	1.53
34	BA	1349	A	P-O5'	-5.18	1.54	1.59
34	BA	1769	U	P-O5'	-5.18	1.54	1.59
34	BA	1770	U	P-O5'	-5.18	1.54	1.59
35	BB	514	G	C2-N2	-5.18	1.29	1.34
35	BB	561	C	C4-N4	-5.18	1.29	1.33
35	BB	1419	G	N3-C4	-5.18	1.31	1.35
36	BC	27	U	C4'-C3'	-5.18	1.47	1.52
36	BC	38	U	C3'-C2'	-5.18	1.47	1.52
36	BC	168	C	O4'-C1'	-5.18	1.34	1.41
37	BD	66	G	P-O5'	-5.18	1.54	1.59
40	BG	118	U	N3-C4	-5.18	1.33	1.38
85	AA	156	G	C3'-C2'	-5.18	1.47	1.52
85	AA	355	G	N9-C4	-5.18	1.33	1.38
85	AA	367	A	C1'-N9	-5.18	1.39	1.46
85	AA	753	U	O3'-P	-5.18	1.54	1.61
85	AA	1792	C	C3'-C2'	-5.18	1.47	1.52
85	AA	2203	C	C1'-N1	-5.18	1.39	1.46
34	BA	8	G	N1-C2	-5.18	1.33	1.37
34	BA	951	C	C1'-N1	-5.18	1.39	1.46
34	BA	1201	G	N9-C8	-5.18	1.34	1.37
34	BA	1534	U	C2'-C1'	-5.18	1.47	1.53
35	BB	576	A	N9-C8	-5.18	1.33	1.37
36	BC	47	C	C4-N4	-5.18	1.29	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	73	U	C3'-C2'	-5.18	1.47	1.52
85	AA	2054	G	O4'-C1'	-5.18	1.34	1.41
34	BA	1041	U	N1-C6	-5.18	1.33	1.38
34	BA	1424	G	O3'-P	-5.18	1.54	1.61
34	BA	1546	C	C4'-O4'	-5.18	1.38	1.45
35	BB	440	U	O3'-P	-5.18	1.54	1.61
35	BB	486	G	C5-C6	-5.18	1.37	1.42
35	BB	1264	U	C5'-C4'	-5.18	1.45	1.51
35	BB	1332	G	N9-C4	-5.18	1.33	1.38
36	BC	88	A	C5'-C4'	-5.18	1.45	1.51
38	BE	176	G	N1-C2	5.18	1.41	1.37
85	AA	81	A	N7-C5	-5.18	1.36	1.39
85	AA	94	C	C1'-N1	-5.18	1.39	1.46
85	AA	340	G	C6-N1	-5.18	1.35	1.39
85	AA	923	A	P-O5'	-5.18	1.54	1.59
85	AA	1227	A	N7-C5	-5.18	1.36	1.39
2	A1	190	GLY	CA-C	-5.18	1.43	1.51
34	BA	328	A	N9-C4	-5.18	1.34	1.37
34	BA	959	G	O3'-P	-5.18	1.54	1.61
34	BA	1086	A	N9-C4	5.18	1.41	1.37
34	BA	1322	A	C1'-N9	-5.18	1.39	1.46
34	BA	1545	C	N1-C6	-5.18	1.34	1.37
35	BB	380	G	C5-C4	-5.18	1.34	1.38
35	BB	600	C	C3'-C2'	-5.18	1.47	1.52
85	AA	425	G	P-O5'	-5.18	1.54	1.59
85	AA	480	U	C2-N3	-5.18	1.34	1.37
85	AA	1491	G	N7-C5	-5.18	1.36	1.39
34	BA	183	G	C8-N7	-5.18	1.27	1.30
34	BA	289	A	O3'-P	-5.18	1.54	1.61
34	BA	518	C	C4'-O4'	-5.18	1.38	1.45
34	BA	600	G	C5-C4	-5.18	1.34	1.38
34	BA	706	C	C1'-N1	-5.18	1.39	1.46
34	BA	1832	A	O5'-C5'	-5.18	1.34	1.42
35	BB	590	G	O3'-P	-5.18	1.54	1.61
35	BB	616	U	C1'-N1	-5.18	1.39	1.46
35	BB	1042	U	C2-N3	-5.18	1.34	1.37
36	BC	77	A	C3'-C2'	-5.18	1.47	1.52
37	BD	113	G	C2'-C1'	-5.18	1.47	1.53
85	AA	533	C	C1'-N1	-5.18	1.39	1.46
85	AA	638	G	N7-C5	-5.18	1.36	1.39
85	AA	749	C	N3-C4	-5.18	1.30	1.33
85	AA	867	G	C5-C6	-5.18	1.37	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	911	A	C6-N1	-5.18	1.31	1.35
85	AA	2128	G	N7-C5	-5.18	1.36	1.39
34	BA	310	C	C2'-C1'	-5.17	1.47	1.53
34	BA	924	U	C3'-C2'	-5.17	1.47	1.52
34	BA	952	G	C2'-C1'	-5.17	1.47	1.53
34	BA	1155	U	O3'-P	-5.17	1.54	1.61
34	BA	1401	C	C3'-C2'	-5.17	1.47	1.52
34	BA	1516	G	C1'-N9	-5.17	1.39	1.46
35	BB	371	C	C1'-N1	-5.17	1.39	1.46
36	BC	124	A	C2'-C1'	-5.17	1.47	1.53
38	BE	199	A	C3'-C2'	-5.17	1.47	1.52
40	BG	82	U	N3-C4	-5.17	1.33	1.38
40	BG	89	A	N7-C5	-5.17	1.36	1.39
85	AA	138	C	O3'-P	-5.17	1.54	1.61
85	AA	874	A	C2'-C1'	-5.17	1.47	1.53
85	AA	1128	G	N9-C4	5.17	1.42	1.38
85	AA	1526	G	O4'-C1'	-5.17	1.34	1.41
85	AA	1693	C	C2-N3	-5.17	1.31	1.35
85	AA	1808	G	O4'-C1'	-5.17	1.34	1.41
85	AA	2138	G	C8-N7	-5.17	1.27	1.30
85	AA	2153	G	C5-C6	-5.17	1.37	1.42
34	BA	430	A	C5-C4	-5.17	1.35	1.38
34	BA	1450	G	N7-C5	-5.17	1.36	1.39
34	BA	1573	C	C3'-C2'	-5.17	1.47	1.52
35	BB	140	U	C2'-C1'	-5.17	1.47	1.53
35	BB	640	A	N3-C4	-5.17	1.31	1.34
35	BB	993	A	C4'-C3'	-5.17	1.47	1.52
35	BB	1173	C	C4'-O4'	-5.17	1.38	1.45
40	BG	107	U	O3'-P	-5.17	1.54	1.61
85	AA	108	C	C4'-C3'	-5.17	1.47	1.52
85	AA	710	A	C4'-C3'	-5.17	1.47	1.52
85	AA	805	A	C3'-C2'	-5.17	1.47	1.52
34	BA	64	A	C5-C4	-5.17	1.35	1.38
34	BA	524	G	C5-C6	-5.17	1.37	1.42
34	BA	573	U	C4'-O4'	-5.17	1.38	1.45
34	BA	704	G	C5-C4	-5.17	1.34	1.38
34	BA	921	G	C5'-C4'	-5.17	1.45	1.51
34	BA	1456	C	C4-N4	-5.17	1.29	1.33
34	BA	1840	C	C4'-O4'	-5.17	1.38	1.45
35	BB	1136	G	C5-C4	-5.17	1.34	1.38
35	BB	1207	C	C3'-C2'	-5.17	1.47	1.52
35	BB	1395	G	C4'-C3'	-5.17	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	90	A	C5-C4	-5.17	1.35	1.38
41	BH	35	G	C6-N1	-5.17	1.35	1.39
85	AA	31	C	C4-N4	-5.17	1.29	1.33
85	AA	792	A	C6-N1	-5.17	1.31	1.35
85	AA	1253	G	O3'-P	-5.17	1.54	1.61
85	AA	1533	C	C2'-C1'	-5.17	1.47	1.53
85	AA	1934	A	C6-N6	-5.17	1.29	1.33
34	BA	14	G	P-O5'	-5.17	1.54	1.59
34	BA	24	C	C4-C5	-5.17	1.38	1.43
34	BA	129	U	C2-N3	-5.17	1.34	1.37
34	BA	720	A	C8-N7	-5.17	1.27	1.31
34	BA	1027	C	P-O5'	-5.17	1.54	1.59
34	BA	1145	U	C4'-C3'	-5.17	1.47	1.52
35	BB	438	G	C5-C4	-5.17	1.34	1.38
35	BB	827	U	C1'-N1	-5.17	1.39	1.46
35	BB	1379	U	O3'-P	-5.17	1.54	1.61
85	AA	410	A	C6-N1	-5.17	1.31	1.35
85	AA	1222	A	C5-C4	-5.17	1.35	1.38
34	BA	13	U	N3-C4	-5.17	1.33	1.38
34	BA	121	A	O3'-P	-5.17	1.54	1.61
34	BA	136	A	C1'-N9	-5.17	1.39	1.46
34	BA	315	U	C2-N3	-5.17	1.34	1.37
34	BA	325	A	C4'-C3'	-5.17	1.47	1.52
34	BA	374	U	N3-C4	-5.17	1.33	1.38
34	BA	678	C	N1-C2	5.17	1.45	1.40
34	BA	816	G	C2-N2	-5.17	1.29	1.34
34	BA	1180	A	N9-C4	-5.17	1.34	1.37
34	BA	1502	G	C6-N1	-5.17	1.35	1.39
34	BA	1554	C	C5'-C4'	-5.17	1.45	1.51
34	BA	1562	G	C3'-C2'	-5.17	1.47	1.52
35	BB	540	G	C8-N7	-5.17	1.27	1.30
35	BB	582	G	C2-N2	-5.17	1.29	1.34
35	BB	591	A	C8-N7	-5.17	1.27	1.31
35	BB	1029	U	N3-C4	-5.17	1.33	1.38
35	BB	1050	A	C3'-C2'	-5.17	1.47	1.52
35	BB	1176	G	N3-C4	-5.17	1.31	1.35
35	BB	1329	G	N9-C4	-5.17	1.33	1.38
41	BH	90	C	P-O5'	-5.17	1.54	1.59
85	AA	786	G	O3'-P	-5.17	1.54	1.61
85	AA	1919	G	C2-N2	-5.17	1.29	1.34
85	AA	2215	C	C2-N3	-5.17	1.31	1.35
34	BA	691	A	C8-N7	-5.17	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	733	G	P-O5'	-5.17	1.54	1.59
34	BA	1144	A	N1-C2	-5.17	1.29	1.34
34	BA	1554	C	C4'-O4'	-5.17	1.38	1.45
34	BA	1616	A	C4'-C3'	-5.17	1.47	1.52
34	BA	1701	U	N1-C6	-5.17	1.33	1.38
35	BB	649	A	N1-C2	-5.17	1.29	1.34
35	BB	1078	U	C2-N3	-5.17	1.34	1.37
35	BB	1362	G	N9-C4	-5.17	1.33	1.38
35	BB	1369	A	N7-C5	-5.17	1.36	1.39
36	BC	14	G	N7-C5	-5.17	1.36	1.39
36	BC	115	G	C2-N2	-5.17	1.29	1.34
37	BD	38	U	N1-C2	-5.17	1.33	1.38
38	BE	64	A	N9-C4	-5.17	1.34	1.37
41	BH	12	U	C2'-C1'	-5.17	1.47	1.53
85	AA	402	G	N3-C4	-5.17	1.31	1.35
85	AA	1689	G	N9-C8	-5.17	1.34	1.37
34	BA	922	C	C1'-N1	-5.17	1.39	1.46
34	BA	1612	C	C4-N4	-5.17	1.29	1.33
85	AA	397	G	N3-C4	-5.17	1.31	1.35
85	AA	429	G	C3'-C2'	-5.17	1.47	1.52
85	AA	522	A	C2'-C1'	-5.17	1.47	1.53
85	AA	958	C	C2'-C1'	-5.17	1.47	1.53
85	AA	1471	G	C2'-C1'	-5.17	1.47	1.53
34	BA	55	G	N9-C8	-5.16	1.34	1.37
34	BA	96	G	N3-C4	-5.16	1.31	1.35
34	BA	506	U	P-O5'	-5.16	1.54	1.59
34	BA	757	G	C5-C4	-5.16	1.34	1.38
34	BA	849	G	C5-C4	-5.16	1.34	1.38
34	BA	897	U	C2-N3	-5.16	1.34	1.37
34	BA	1479	G	C1'-N9	-5.16	1.39	1.46
34	BA	1547	G	O4'-C1'	-5.16	1.34	1.41
35	BB	537	A	C2'-C1'	-5.16	1.47	1.53
35	BB	991	C	C2'-C1'	-5.16	1.47	1.53
35	BB	1274	G	C5-C4	-5.16	1.34	1.38
35	BB	1426	G	O3'-P	-5.16	1.54	1.61
39	BF	63	U	C2-N3	-5.16	1.34	1.37
41	BH	15	A	N3-C4	-5.16	1.31	1.34
85	AA	313	A	C4'-C3'	-5.16	1.47	1.52
85	AA	352	G	N9-C8	-5.16	1.34	1.37
85	AA	407	G	C2-N2	-5.16	1.29	1.34
85	AA	1169	A	C1'-N9	-5.16	1.39	1.46
85	AA	1251	G	C5-C4	-5.16	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1441	G	C5'-C4'	-5.16	1.45	1.51
85	AA	1451	U	C4-O4	-5.16	1.19	1.23
85	AA	1475	A	O3'-P	-5.16	1.54	1.61
85	AA	2182	A	C5-C4	-5.16	1.35	1.38
1	A0	136	GLY	CA-C	-5.16	1.43	1.51
34	BA	327	G	C1'-N9	-5.16	1.39	1.46
35	BB	63	A	C2'-C1'	-5.16	1.47	1.53
85	AA	773	G	O3'-P	-5.16	1.54	1.61
85	AA	797	C	O3'-P	-5.16	1.54	1.61
85	AA	1251	G	C3'-C2'	-5.16	1.47	1.52
85	AA	2186	U	C4'-C3'	-5.16	1.47	1.52
34	BA	355	U	C1'-N1	-5.16	1.39	1.46
34	BA	407	A	N9-C4	-5.16	1.34	1.37
34	BA	410	G	C1'-N9	-5.16	1.39	1.46
34	BA	668	G	P-O5'	-5.16	1.54	1.59
34	BA	732	A	C1'-N9	-5.16	1.39	1.46
34	BA	871	G	C6-N1	-5.16	1.35	1.39
34	BA	925	G	C2-N3	-5.16	1.28	1.32
34	BA	1045	C	C1'-N1	-5.16	1.39	1.46
34	BA	1337	A	N9-C4	-5.16	1.34	1.37
34	BA	1668	C	C2-N3	-5.16	1.31	1.35
34	BA	1711	G	O4'-C1'	-5.16	1.34	1.41
35	BB	32	C	C1'-N1	-5.16	1.39	1.46
35	BB	855	G	C2'-C1'	-5.16	1.47	1.53
35	BB	1063	C	C3'-C2'	-5.16	1.47	1.52
35	BB	1438	U	C3'-C2'	-5.16	1.47	1.52
36	BC	87	C	C4'-C3'	-5.16	1.47	1.52
38	BE	28	C	C4-C5	-5.16	1.38	1.43
39	BF	63	U	N1-C2	-5.16	1.33	1.38
85	AA	10	G	C2-N2	-5.16	1.29	1.34
85	AA	362	G	C5-C4	-5.16	1.34	1.38
85	AA	493	A	N9-C4	-5.16	1.34	1.37
85	AA	1442	U	C2'-C1'	-5.16	1.47	1.53
85	AA	1728	G	C2'-C1'	-5.16	1.47	1.53
34	BA	20	A	C3'-C2'	-5.16	1.47	1.52
34	BA	59	A	C4'-C3'	-5.16	1.47	1.52
34	BA	785	G	N9-C4	-5.16	1.33	1.38
34	BA	852	C	N1-C6	-5.16	1.34	1.37
34	BA	855	C	C1'-N1	-5.16	1.39	1.46
34	BA	1040	G	N1-C2	-5.16	1.33	1.37
34	BA	1301	G	O4'-C1'	-5.16	1.34	1.41
35	BB	517	G	C2-N2	-5.16	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	894	A	C4'-C3'	5.16	1.58	1.53
35	BB	1291	G	C2-N2	-5.16	1.29	1.34
36	BC	37	U	N1-C6	-5.16	1.33	1.38
38	BE	50	G	C2'-C1'	-5.16	1.47	1.53
41	BH	77	G	C8-N7	5.16	1.34	1.30
41	BH	113	G	O4'-C1'	-5.16	1.34	1.41
82	Bw	66	TYR	CB-CG	-5.16	1.44	1.51
85	AA	457	G	N1-C2	-5.16	1.33	1.37
85	AA	532	G	O3'-P	-5.16	1.54	1.61
85	AA	591	A	N9-C8	-5.16	1.33	1.37
85	AA	1488	G	N9-C8	-5.16	1.34	1.37
85	AA	1589	G	O3'-P	-5.16	1.54	1.61
85	AA	1688	U	C3'-C2'	-5.16	1.47	1.52
85	AA	1856	G	O3'-P	-5.16	1.54	1.61
85	AA	2149	C	N1-C6	-5.16	1.34	1.37
85	AA	2194	U	N1-C6	-5.16	1.33	1.38
86	AB	21	A	C2'-C1'	-5.16	1.47	1.53
34	BA	61	G	C4'-C3'	-5.16	1.47	1.52
34	BA	291	C	C4-N4	-5.16	1.29	1.33
34	BA	1519	G	C3'-C2'	-5.16	1.47	1.52
35	BB	435	A	C5-C6	-5.16	1.36	1.41
35	BB	519	A	P-O5'	-5.16	1.54	1.59
35	BB	703	U	C3'-C2'	-5.16	1.47	1.52
35	BB	1115	G	C1'-N9	-5.16	1.39	1.46
35	BB	1396	G	C8-N7	-5.16	1.27	1.30
36	BC	53	A	N9-C8	-5.16	1.33	1.37
38	BE	6	A	O4'-C1'	-5.16	1.34	1.41
40	BG	71	C	C4-C5	-5.16	1.38	1.43
85	AA	472	A	C3'-C2'	-5.16	1.47	1.52
85	AA	1515	A	N7-C5	-5.16	1.36	1.39
85	AA	2015	U	O3'-P	-5.16	1.54	1.61
34	BA	405	C	C2'-C1'	-5.16	1.47	1.53
34	BA	461	A	C5-C6	-5.16	1.36	1.41
34	BA	849	G	C2'-C1'	-5.16	1.47	1.53
34	BA	932	G	N9-C8	-5.16	1.34	1.37
35	BB	582	G	C1'-N9	-5.16	1.39	1.46
35	BB	815	G	N9-C8	-5.16	1.34	1.37
35	BB	970	C	C3'-O3'	-5.16	1.34	1.42
35	BB	1098	G	C2-N2	-5.16	1.29	1.34
35	BB	1321	G	C4'-O4'	-5.16	1.38	1.45
35	BB	1505	U	C4'-C3'	-5.16	1.47	1.52
36	BC	11	G	N7-C5	-5.16	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	BC	157	U	C5'-C4'	5.16	1.57	1.51
38	BE	8	G	O4'-C1'	5.16	1.48	1.41
38	BE	111	C	C1'-N1	5.16	1.56	1.48
39	BF	72	A	N3-C4	-5.16	1.31	1.34
40	BG	54	G	C4'-C3'	-5.16	1.47	1.52
40	BG	139	U	P-O5'	-5.16	1.54	1.59
41	BH	67	G	N9-C4	5.16	1.42	1.38
85	AA	37	U	C2-N3	-5.16	1.34	1.37
85	AA	404	A	C5-C4	-5.16	1.35	1.38
85	AA	442	G	N1-C2	-5.16	1.33	1.37
85	AA	1044	G	P-O5'	-5.16	1.54	1.59
85	AA	1206	A	C1'-N9	-5.16	1.39	1.46
85	AA	1481	U	C2'-C1'	-5.16	1.47	1.53
85	AA	1547	G	C5-C4	-5.16	1.34	1.38
85	AA	1640	G	P-O5'	-5.16	1.54	1.59
85	AA	1976	G	N1-C2	-5.16	1.33	1.37
34	BA	55	G	O3'-P	-5.15	1.54	1.61
34	BA	80	U	C3'-C2'	-5.15	1.47	1.52
34	BA	353	U	C1'-N1	-5.15	1.39	1.46
34	BA	697	A	C8-N7	-5.15	1.27	1.31
34	BA	1653	G	O4'-C1'	-5.15	1.34	1.41
38	BE	45	G	N1-C2	-5.15	1.33	1.37
64	Be	154	GLN	N-CA	-5.15	1.36	1.46
76	Bq	25	TYR	CB-CG	-5.15	1.44	1.51
85	AA	180	A	N7-C5	-5.15	1.36	1.39
85	AA	801	U	O3'-P	-5.15	1.54	1.61
34	BA	63	A	C5'-C4'	-5.15	1.45	1.51
34	BA	189	G	N9-C4	-5.15	1.33	1.38
34	BA	279	U	O3'-P	-5.15	1.54	1.61
34	BA	333	A	C4'-C3'	-5.15	1.47	1.52
34	BA	600	G	N9-C8	-5.15	1.34	1.37
34	BA	861	C	P-O5'	-5.15	1.54	1.59
34	BA	1250	C	C4-N4	-5.15	1.29	1.33
34	BA	1618	A	O3'-P	-5.15	1.54	1.61
34	BA	1671	A	N9-C8	-5.15	1.33	1.37
35	BB	41	A	N3-C4	-5.15	1.31	1.34
35	BB	562	A	N3-C4	-5.15	1.31	1.34
35	BB	769	C	C2'-C1'	-5.15	1.47	1.53
35	BB	810	G	C2-N2	-5.15	1.29	1.34
35	BB	1216	G	N9-C4	-5.15	1.33	1.38
36	BC	16	A	N3-C4	-5.15	1.31	1.34
36	BC	87	C	C1'-N1	-5.15	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	60	C	C5'-C4'	-5.15	1.45	1.51
40	BG	166	C	N1-C6	-5.15	1.34	1.37
66	Bg	54	VAL	CA-CB	-5.15	1.44	1.54
85	AA	302	C	C3'-O3'	5.15	1.49	1.42
85	AA	820	G	N7-C5	-5.15	1.36	1.39
85	AA	836	A	C2'-C1'	-5.15	1.47	1.53
85	AA	941	C	C4'-O4'	-5.15	1.38	1.45
85	AA	1293	U	N3-C4	-5.15	1.33	1.38
85	AA	1452	C	C2'-C1'	-5.15	1.47	1.53
85	AA	1913	G	C2'-C1'	-5.15	1.47	1.53
85	AA	2186	U	O4'-C1'	-5.15	1.34	1.41
34	BA	39	C	C4-C5	-5.15	1.38	1.43
34	BA	609	G	C1'-N9	-5.15	1.39	1.46
34	BA	1056	C	P-O5'	-5.15	1.54	1.59
34	BA	1385	U	O3'-P	-5.15	1.54	1.61
34	BA	1632	G	C5-C6	-5.15	1.37	1.42
35	BB	108	G	N7-C5	-5.15	1.36	1.39
35	BB	423	G	C3'-C2'	-5.15	1.47	1.52
35	BB	453	C	N1-C6	-5.15	1.34	1.37
35	BB	702	G	N9-C8	-5.15	1.34	1.37
35	BB	998	G	C6-N1	-5.15	1.35	1.39
35	BB	1287	U	C2'-C1'	-5.15	1.47	1.53
37	BD	8	A	C1'-N9	-5.15	1.39	1.46
40	BG	31	G	C4'-C3'	-5.15	1.47	1.52
40	BG	44	G	C3'-C2'	-5.15	1.47	1.52
40	BG	126	G	N1-C2	-5.15	1.33	1.37
85	AA	211	C	C2-N3	-5.15	1.31	1.35
85	AA	702	G	C2-N2	-5.15	1.29	1.34
85	AA	771	A	C2'-C1'	-5.15	1.47	1.53
85	AA	993	G	N3-C4	-5.15	1.31	1.35
85	AA	1018	G	C2-N2	-5.15	1.29	1.34
85	AA	1203	G	N3-C4	-5.15	1.31	1.35
1	A0	104	GLY	CA-C	-5.15	1.43	1.51
34	BA	162	G	C5'-C4'	5.15	1.57	1.51
34	BA	443	U	P-O5'	-5.15	1.54	1.59
34	BA	497	U	P-O5'	-5.15	1.54	1.59
40	BG	18	U	C2-N3	-5.15	1.34	1.37
34	BA	67	A	O4'-C1'	-5.15	1.34	1.41
34	BA	212	A	C2'-C1'	-5.15	1.47	1.53
34	BA	368	U	N1-C6	-5.15	1.33	1.38
34	BA	491	U	O4'-C1'	-5.15	1.34	1.41
34	BA	956	G	N9-C8	-5.15	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1015	G	C2-N2	-5.15	1.29	1.34
34	BA	1306	U	O3'-P	-5.15	1.54	1.61
34	BA	1558	C	C3'-C2'	-5.15	1.47	1.52
34	BA	1578	A	P-O5'	-5.15	1.54	1.59
35	BB	120	C	C2'-C1'	-5.15	1.47	1.53
35	BB	521	U	N3-C4	-5.15	1.33	1.38
35	BB	1114	A	O3'-P	-5.15	1.54	1.61
35	BB	1272	G	C5-C6	-5.15	1.37	1.42
35	BB	1427	A	N7-C5	-5.15	1.36	1.39
40	BG	17	A	N7-C5	-5.15	1.36	1.39
85	AA	13	U	C2'-C1'	-5.15	1.47	1.53
85	AA	15	U	O4'-C1'	-5.15	1.34	1.41
85	AA	525	C	C4'-C3'	-5.15	1.47	1.52
85	AA	1589	G	N1-C2	-5.15	1.33	1.37
85	AA	1644	G	C6-N1	-5.15	1.35	1.39
85	AA	1831	U	C2-N3	-5.15	1.34	1.37
85	AA	2109	G	N3-C4	-5.15	1.31	1.35
34	BA	227	C	C5'-C4'	-5.15	1.45	1.51
34	BA	859	G	N9-C8	-5.15	1.34	1.37
34	BA	1498	A	N9-C4	-5.15	1.34	1.37
34	BA	1731	A	N3-C4	-5.15	1.31	1.34
35	BB	42	A	C4'-O4'	-5.15	1.38	1.45
35	BB	534	C	O3'-P	-5.15	1.54	1.61
35	BB	1036	G	C5-C4	-5.15	1.34	1.38
35	BB	1168	G	C5-C4	-5.15	1.34	1.38
42	BI	140	GLY	CA-C	-5.15	1.43	1.51
85	AA	37	U	C3'-C2'	-5.15	1.47	1.52
85	AA	97	A	O3'-P	-5.15	1.54	1.61
85	AA	152	A	C5-C6	-5.15	1.36	1.41
85	AA	159	G	C2-N2	-5.15	1.29	1.34
85	AA	1219	A	N3-C4	-5.15	1.31	1.34
85	AA	1558	U	C1'-N1	-5.15	1.39	1.46
34	BA	99	G	C1'-N9	-5.14	1.39	1.46
34	BA	149	G	C2-N2	-5.14	1.29	1.34
34	BA	532	C	C1'-N1	-5.14	1.39	1.46
34	BA	1499	A	C3'-C2'	-5.14	1.47	1.52
35	BB	59	U	C4-O4	-5.14	1.19	1.23
35	BB	98	A	C3'-C2'	-5.14	1.47	1.52
35	BB	351	G	P-O5'	-5.14	1.54	1.59
35	BB	550	G	P-O5'	-5.14	1.54	1.59
35	BB	789	G	C4'-O4'	-5.14	1.38	1.45
35	BB	1016	C	C2-N3	-5.14	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1297	G	C3'-C2'	-5.14	1.47	1.52
39	BF	6	C	C4'-O4'	-5.14	1.38	1.45
39	BF	60	C	O3'-P	-5.14	1.54	1.61
40	BG	39	A	C8-N7	-5.14	1.27	1.31
41	BH	107	A	C8-N7	-5.14	1.27	1.31
85	AA	71	G	C2-N2	-5.14	1.29	1.34
85	AA	351	C	C1'-N1	-5.14	1.39	1.46
85	AA	403	G	C2-N2	-5.14	1.29	1.34
85	AA	491	G	N1-C2	-5.14	1.33	1.37
85	AA	989	U	N1-C2	-5.14	1.33	1.38
85	AA	1466	U	C4-C5	-5.14	1.39	1.43
85	AA	1520	A	O4'-C1'	-5.14	1.34	1.41
85	AA	1658	G	N9-C8	-5.14	1.34	1.37
85	AA	2127	G	N3-C4	-5.14	1.31	1.35
34	BA	354	G	N1-C2	-5.14	1.33	1.37
34	BA	376	U	C2'-C1'	-5.14	1.47	1.53
34	BA	421	G	C2-N2	-5.14	1.29	1.34
34	BA	1238	C	P-O5'	-5.14	1.54	1.59
34	BA	1452	U	C2'-C1'	-5.14	1.47	1.53
35	BB	52	G	N1-C2	-5.14	1.33	1.37
35	BB	1067	G	C3'-C2'	-5.14	1.47	1.52
35	BB	1296	A	C5-C6	-5.14	1.36	1.41
35	BB	1327	U	N1-C2	-5.14	1.33	1.38
35	BB	1409	G	C3'-C2'	-5.14	1.47	1.52
36	BC	92	C	C1'-N1	-5.14	1.39	1.46
38	BE	189	A	O3'-P	-5.14	1.54	1.61
38	BE	198	A	C4'-C3'	-5.14	1.47	1.52
40	BG	36	G	C2-N2	-5.14	1.29	1.34
85	AA	275	A	C1'-N9	-5.14	1.39	1.46
85	AA	395	G	C3'-C2'	-5.14	1.47	1.52
85	AA	1466	U	C5'-C4'	5.14	1.57	1.51
85	AA	1495	G	O5'-C5'	-5.14	1.34	1.42
85	AA	1706	A	O3'-P	-5.14	1.54	1.61
85	AA	1949	U	P-O5'	-5.14	1.54	1.59
85	AA	2095	U	N3-C4	-5.14	1.33	1.38
34	BA	260	A	N3-C4	-5.14	1.31	1.34
34	BA	414	A	O4'-C1'	-5.14	1.34	1.41
34	BA	886	G	C5-C4	-5.14	1.34	1.38
35	BB	42	A	C2'-C1'	-5.14	1.47	1.53
35	BB	549	U	P-O5'	-5.14	1.54	1.59
35	BB	957	A	C2'-C1'	-5.14	1.47	1.53
36	BC	38	U	N3-C4	-5.14	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
37	BD	112	U	O4'-C1'	-5.14	1.34	1.41
39	BF	35	C	C4'-C3'	-5.14	1.47	1.52
85	AA	1159	C	N1-C6	5.14	1.40	1.37
85	AA	2151	U	N3-C4	-5.14	1.33	1.38
34	BA	1507	C	C3'-C2'	-5.14	1.47	1.52
34	BA	1572	G	N1-C2	-5.14	1.33	1.37
34	BA	1688	G	C2'-C1'	-5.14	1.47	1.53
34	BA	1810	A	N9-C4	5.14	1.41	1.37
35	BB	95	A	C3'-C2'	-5.14	1.47	1.52
35	BB	570	A	C6-N1	-5.14	1.31	1.35
35	BB	860	U	P-O5'	-5.14	1.54	1.59
35	BB	861	C	P-O5'	-5.14	1.54	1.59
35	BB	1062	G	O4'-C1'	-5.14	1.34	1.41
35	BB	1206	G	C2'-C1'	-5.14	1.47	1.53
36	BC	68	A	P-O5'	-5.14	1.54	1.59
36	BC	160	C	C4'-C3'	-5.14	1.47	1.52
37	BD	104	C	C1'-N1	-5.14	1.39	1.46
40	BG	157	A	N3-C4	-5.14	1.31	1.34
85	AA	450	A	C5-C4	-5.14	1.35	1.38
85	AA	1522	U	C1'-N1	-5.14	1.39	1.46
85	AA	1808	G	P-O5'	-5.14	1.54	1.59
34	BA	110	C	N1-C6	-5.14	1.34	1.37
34	BA	541	C	C2-N3	-5.14	1.31	1.35
34	BA	774	A	C5'-C4'	-5.14	1.45	1.51
34	BA	982	A	O4'-C1'	-5.14	1.34	1.41
34	BA	1091	U	C1'-N1	-5.14	1.39	1.46
34	BA	1119	A	C6-N6	-5.14	1.29	1.33
34	BA	1531	G	C4'-C3'	-5.14	1.47	1.52
34	BA	1629	A	C3'-C2'	-5.14	1.47	1.52
34	BA	1723	U	O3'-P	-5.14	1.54	1.61
35	BB	24	C	O3'-P	-5.14	1.54	1.61
35	BB	602	G	C6-N1	-5.14	1.35	1.39
40	BG	1	G	C2-N2	-5.14	1.29	1.34
41	BH	33	G	C8-N7	-5.14	1.27	1.30
85	AA	1615	A	P-O5'	-5.14	1.54	1.59
34	BA	74	A	P-O5'	-5.14	1.54	1.59
34	BA	579	U	P-O5'	-5.14	1.54	1.59
34	BA	1077	G	C6-N1	-5.14	1.35	1.39
34	BA	1116	G	O3'-P	-5.14	1.54	1.61
34	BA	1482	A	C3'-C2'	-5.14	1.47	1.52
34	BA	1727	A	C2'-C1'	-5.14	1.47	1.53
34	BA	1790	U	C3'-C2'	-5.14	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	500	C	C4-C5	-5.14	1.38	1.43
35	BB	526	A	C2'-C1'	-5.14	1.47	1.53
35	BB	1132	A	C1'-N9	-5.14	1.39	1.46
36	BC	156	A	O3'-P	-5.14	1.54	1.61
40	BG	11	G	C5'-C4'	5.14	1.57	1.51
41	BH	111	U	N1-C2	-5.14	1.33	1.38
77	Br	284	SER	CA-C	-5.14	1.39	1.52
85	AA	436	G	C2-N3	-5.14	1.28	1.32
85	AA	1345	C	O3'-P	-5.14	1.54	1.61
34	BA	492	G	N1-C2	-5.13	1.33	1.37
34	BA	660	C	C2-N3	-5.13	1.31	1.35
34	BA	766	A	O3'-P	-5.13	1.54	1.61
34	BA	1158	A	N9-C8	-5.13	1.33	1.37
34	BA	1476	G	C2-N2	-5.13	1.29	1.34
34	BA	1691	G	C2-N2	-5.13	1.29	1.34
35	BB	1410	G	N9-C4	-5.13	1.33	1.38
38	BE	198	A	C8-N7	-5.13	1.27	1.31
40	BG	60	A	C2'-C1'	-5.13	1.47	1.53
85	AA	303	A	C4'-C3'	5.13	1.58	1.53
85	AA	312	G	C5-C4	-5.13	1.34	1.38
85	AA	2041	G	C1'-N9	-5.13	1.39	1.46
34	BA	135	G	C1'-N9	-5.13	1.39	1.46
34	BA	201	A	N7-C5	-5.13	1.36	1.39
34	BA	880	G	N7-C5	-5.13	1.36	1.39
34	BA	1235	C	C3'-C2'	-5.13	1.47	1.52
85	AA	601	A	C4'-C3'	-5.13	1.47	1.52
85	AA	1039	U	O3'-P	-5.13	1.54	1.61
34	BA	472	G	C2'-C1'	-5.13	1.47	1.53
34	BA	1226	G	C5'-C4'	-5.13	1.45	1.51
35	BB	47	C	P-O5'	-5.13	1.54	1.59
35	BB	134	G	O3'-P	-5.13	1.54	1.61
35	BB	639	A	C8-N7	-5.13	1.27	1.31
35	BB	664	A	C5-C4	-5.13	1.35	1.38
35	BB	741	A	O3'-P	-5.13	1.54	1.61
36	BC	68	A	C2'-C1'	-5.13	1.47	1.53
36	BC	160	C	O4'-C1'	-5.13	1.34	1.41
37	BD	118	C	C4'-C3'	-5.13	1.47	1.52
40	BG	55	A	C3'-C2'	-5.13	1.47	1.52
41	BH	34	G	O3'-P	-5.13	1.54	1.61
85	AA	16	G	N1-C2	-5.13	1.33	1.37
85	AA	419	A	C5-C6	-5.13	1.36	1.41
85	AA	480	U	C4'-O4'	-5.13	1.38	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	561	C	N3-C4	-5.13	1.30	1.33
85	AA	1140	G	N9-C8	-5.13	1.34	1.37
85	AA	1214	C	C4-C5	-5.13	1.38	1.43
85	AA	1532	G	C5-C4	-5.13	1.34	1.38
85	AA	1703	A	C5-C4	-5.13	1.35	1.38
85	AA	2237	G	C2'-C1'	-5.13	1.47	1.53
34	BA	447	U	N3-C4	-5.13	1.33	1.38
34	BA	1028	A	N7-C5	-5.13	1.36	1.39
34	BA	1710	C	O3'-P	-5.13	1.54	1.61
35	BB	499	A	C2'-C1'	-5.13	1.47	1.53
35	BB	571	C	P-O5'	-5.13	1.54	1.59
85	AA	435	A	N9-C8	-5.13	1.33	1.37
85	AA	1127	G	C4'-O4'	-5.13	1.38	1.45
85	AA	1177	G	N1-C2	-5.13	1.33	1.37
85	AA	1726	G	C3'-C2'	-5.13	1.47	1.52
86	AB	44	G	P-O5'	-5.13	1.54	1.59
34	BA	186	G	C2-N2	-5.13	1.29	1.34
34	BA	387	A	N9-C4	-5.13	1.34	1.37
34	BA	526	C	O3'-P	-5.13	1.54	1.61
34	BA	600	G	C2-N3	-5.13	1.28	1.32
34	BA	1088	G	C2-N2	-5.13	1.29	1.34
34	BA	1490	U	C5'-C4'	5.13	1.57	1.51
34	BA	1547	G	C4'-C3'	-5.13	1.47	1.52
34	BA	1614	G	N9-C8	-5.13	1.34	1.37
35	BB	40	C	C2'-C1'	-5.13	1.47	1.53
35	BB	438	G	N9-C4	-5.13	1.33	1.38
35	BB	689	C	C5'-C4'	-5.13	1.45	1.51
35	BB	1401	G	C5-C6	-5.13	1.37	1.42
39	BF	13	U	N3-C4	-5.13	1.33	1.38
39	BF	32	G	C8-N7	-5.13	1.27	1.30
40	BG	45	G	C2-N2	-5.13	1.29	1.34
85	AA	29	U	C1'-N1	-5.13	1.39	1.46
85	AA	111	A	C2'-C1'	-5.13	1.47	1.53
85	AA	487	G	O4'-C1'	-5.13	1.34	1.41
85	AA	1093	C	C2'-C1'	-5.13	1.47	1.53
85	AA	1189	A	O3'-P	-5.13	1.54	1.61
85	AA	1268	C	C1'-N1	-5.13	1.39	1.46
85	AA	1274	A	N9-C4	-5.13	1.34	1.37
85	AA	1593	C	N3-C4	-5.13	1.30	1.33
85	AA	2198	G	N9-C4	-5.13	1.33	1.38
31	AX	155	TYR	CB-CG	-5.13	1.44	1.51
34	BA	588	C	C4-N4	-5.13	1.29	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	656	U	P-O5'	-5.13	1.54	1.59
34	BA	1016	A	C6-N1	-5.13	1.31	1.35
34	BA	1581	G	N7-C5	-5.13	1.36	1.39
34	BA	1825	U	C2'-C1'	-5.13	1.47	1.53
35	BB	441	G	C2-N2	-5.13	1.29	1.34
35	BB	1397	G	C2-N2	-5.13	1.29	1.34
36	BC	97	U	C2'-C1'	-5.13	1.47	1.53
36	BC	162	C	C3'-C2'	-5.13	1.47	1.52
39	BF	61	A	C5'-C4'	5.13	1.57	1.51
40	BG	154	C	C1'-N1	-5.13	1.39	1.46
85	AA	324	U	P-O5'	-5.13	1.54	1.59
85	AA	792	A	C1'-N9	-5.13	1.39	1.46
85	AA	877	G	C2-N2	-5.13	1.29	1.34
85	AA	1260	G	N7-C5	-5.13	1.36	1.39
85	AA	1544	G	C2'-C1'	-5.13	1.47	1.53
34	BA	265	A	N9-C4	-5.12	1.34	1.37
34	BA	529	A	C6-N6	-5.12	1.29	1.33
35	BB	552	C	N1-C6	-5.12	1.34	1.37
35	BB	1024	G	C3'-O3'	5.12	1.49	1.42
35	BB	1337	C	C1'-N1	-5.12	1.39	1.46
35	BB	1343	C	C4-N4	-5.12	1.29	1.33
85	AA	615	A	C5'-C4'	5.12	1.57	1.51
85	AA	709	A	P-O5'	-5.12	1.54	1.59
85	AA	1243	G	C6-N1	-5.12	1.35	1.39
86	AB	5	G	C2-N2	-5.12	1.29	1.34
34	BA	854	A	C4'-O4'	-5.12	1.38	1.45
34	BA	876	C	C4'-O4'	-5.12	1.38	1.45
34	BA	1571	C	O3'-P	-5.12	1.55	1.61
34	BA	1812	C	P-O5'	-5.12	1.54	1.59
35	BB	528	G	C2'-C1'	-5.12	1.47	1.53
35	BB	1085	C	O3'-P	-5.12	1.55	1.61
35	BB	1105	G	C2-N2	-5.12	1.29	1.34
35	BB	1178	A	N9-C8	-5.12	1.33	1.37
37	BD	11	A	C5-C4	-5.12	1.35	1.38
84	By	184	ASN	CA-C	-5.12	1.39	1.52
85	AA	438	G	C2'-C1'	-5.12	1.47	1.53
85	AA	514	U	C2'-C1'	-5.12	1.47	1.53
85	AA	668	A	N7-C5	-5.12	1.36	1.39
85	AA	973	U	C4'-C3'	5.12	1.58	1.53
85	AA	1125	G	N7-C5	-5.12	1.36	1.39
85	AA	1220	A	N3-C4	-5.12	1.31	1.34
85	AA	1261	U	O3'-P	-5.12	1.55	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1540	A	C2'-C1'	-5.12	1.47	1.53
85	AA	2064	A	O3'-P	-5.12	1.55	1.61
85	AA	2239	A	N3-C4	-5.12	1.31	1.34
34	BA	363	G	C3'-C2'	-5.12	1.47	1.52
34	BA	388	A	C1'-N9	-5.12	1.39	1.46
34	BA	442	G	N9-C8	-5.12	1.34	1.37
34	BA	444	A	P-O5'	-5.12	1.54	1.59
34	BA	454	G	C2-N2	-5.12	1.29	1.34
34	BA	562	C	O3'-P	-5.12	1.55	1.61
35	BB	567	G	N7-C5	-5.12	1.36	1.39
35	BB	1403	G	C1'-N9	-5.12	1.39	1.46
35	BB	1413	U	C2-N3	-5.12	1.34	1.37
37	BD	89	G	N3-C4	-5.12	1.31	1.35
38	BE	33	C	N1-C6	-5.12	1.34	1.37
40	BG	106	G	C2-N2	-5.12	1.29	1.34
41	BH	121	A	C1'-N9	-5.12	1.39	1.46
85	AA	534	A	C2'-C1'	-5.12	1.47	1.53
85	AA	1419	U	P-O5'	-5.12	1.54	1.59
85	AA	1783	G	P-O5'	-5.12	1.54	1.59
85	AA	1785	U	C5'-C4'	5.12	1.57	1.51
85	AA	2125	A	C3'-C2'	-5.12	1.47	1.52
85	AA	2189	U	C4'-C3'	-5.12	1.47	1.52
34	BA	24	C	P-O5'	-5.12	1.54	1.59
34	BA	1603	A	C5-C6	-5.12	1.36	1.41
35	BB	1272	G	C2-N2	-5.12	1.29	1.34
39	BF	48	G	N1-C2	-5.12	1.33	1.37
85	AA	80	G	P-O5'	-5.12	1.54	1.59
85	AA	272	C	C4'-O4'	5.12	1.52	1.45
85	AA	810	C	C4-N4	-5.12	1.29	1.33
85	AA	1547	G	C6-N1	-5.12	1.35	1.39
34	BA	294	C	C4'-C3'	-5.12	1.47	1.52
34	BA	354	G	C5-C4	-5.12	1.34	1.38
34	BA	696	A	P-O5'	-5.12	1.54	1.59
34	BA	803	U	C2'-C1'	-5.12	1.47	1.53
34	BA	920	U	N1-C6	-5.12	1.33	1.38
34	BA	1030	C	C1'-N1	-5.12	1.39	1.46
34	BA	1077	G	C5-C6	-5.12	1.37	1.42
34	BA	1282	G	C2-N2	-5.12	1.29	1.34
34	BA	1550	G	N9-C8	-5.12	1.34	1.37
35	BB	95	A	C5-C6	-5.12	1.36	1.41
35	BB	593	A	C2'-C1'	-5.12	1.47	1.53
35	BB	661	G	C5-C4	-5.12	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1267	C	N1-C2	-5.12	1.35	1.40
35	BB	1455	A	C5-C6	-5.12	1.36	1.41
36	BC	143	C	C2'-C1'	-5.12	1.47	1.53
37	BD	108	G	N3-C4	-5.12	1.31	1.35
40	BG	18	U	C4'-C3'	-5.12	1.47	1.52
85	AA	23	G	P-O5'	-5.12	1.54	1.59
85	AA	1131	A	N7-C5	-5.12	1.36	1.39
85	AA	1563	U	C3'-C2'	-5.12	1.47	1.52
85	AA	1859	C	C4'-O4'	-5.12	1.38	1.45
34	BA	726	G	C1'-N9	-5.12	1.39	1.46
34	BA	1063	G	C5-C4	-5.12	1.34	1.38
34	BA	1139	G	C3'-C2'	-5.12	1.47	1.52
34	BA	1633	C	P-O5'	-5.12	1.54	1.59
35	BB	1423	U	N1-C2	-5.12	1.33	1.38
36	BC	96	A	O4'-C1'	-5.12	1.34	1.41
36	BC	121	G	C5-C4	-5.12	1.34	1.38
38	BE	179	A	P-O5'	-5.12	1.54	1.59
85	AA	229	U	P-O5'	-5.12	1.54	1.59
85	AA	735	G	O3'-P	-5.12	1.55	1.61
85	AA	1293	U	C2'-C1'	-5.12	1.47	1.53
34	BA	237	A	O3'-P	-5.12	1.55	1.61
34	BA	1326	U	C5'-C4'	5.12	1.57	1.51
34	BA	1500	G	N1-C2	-5.12	1.33	1.37
34	BA	1644	A	C4'-C3'	-5.12	1.47	1.52
34	BA	1676	A	N9-C8	-5.12	1.33	1.37
35	BB	697	G	C3'-C2'	-5.12	1.47	1.52
35	BB	1172	U	C2-N3	-5.12	1.34	1.37
35	BB	1175	A	C2'-C1'	-5.12	1.47	1.53
35	BB	1234	G	N1-C2	-5.12	1.33	1.37
38	BE	49	A	C4'-C3'	-5.12	1.47	1.52
41	BH	44	A	C3'-C2'	-5.12	1.47	1.52
41	BH	134	U	P-O5'	-5.12	1.54	1.59
65	Bf	416	PHE	CB-CG	-5.12	1.42	1.51
85	AA	114	C	C2-N3	-5.12	1.31	1.35
85	AA	1047	G	P-O5'	-5.12	1.54	1.59
34	BA	205	G	N9-C8	-5.11	1.34	1.37
34	BA	462	C	C5'-C4'	-5.11	1.45	1.51
34	BA	714	G	N1-C2	-5.11	1.33	1.37
34	BA	928	C	O4'-C1'	-5.11	1.35	1.41
34	BA	1537	G	N3-C4	-5.11	1.31	1.35
35	BB	116	G	C5-C6	-5.11	1.37	1.42
35	BB	390	G	C2-N2	-5.11	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	402	G	N3-C4	-5.11	1.31	1.35
35	BB	1061	G	N7-C5	-5.11	1.36	1.39
38	BE	180	G	O4'-C1'	-5.11	1.35	1.41
40	BG	69	G	C3'-C2'	-5.11	1.47	1.52
40	BG	105	A	N3-C4	-5.11	1.31	1.34
85	AA	158	C	C2-N3	-5.11	1.31	1.35
85	AA	351	C	C2-N3	-5.11	1.31	1.35
85	AA	358	U	C4-C5	-5.11	1.39	1.43
85	AA	463	G	O4'-C1'	-5.11	1.35	1.41
85	AA	934	A	N9-C4	-5.11	1.34	1.37
85	AA	1285	C	O3'-P	-5.11	1.55	1.61
85	AA	1473	U	N1-C2	-5.11	1.33	1.38
34	BA	465	A	N9-C8	-5.11	1.33	1.37
34	BA	998	U	C4'-O4'	-5.11	1.39	1.45
34	BA	1001	G	C5-C6	-5.11	1.37	1.42
34	BA	1411	C	N1-C6	-5.11	1.34	1.37
34	BA	1836	A	P-O5'	-5.11	1.54	1.59
35	BB	778	A	C8-N7	-5.11	1.27	1.31
35	BB	1423	U	C4'-C3'	-5.11	1.47	1.52
36	BC	16	A	P-O5'	-5.11	1.54	1.59
37	BD	114	U	O3'-P	-5.11	1.55	1.61
40	BG	35	G	O4'-C1'	-5.11	1.35	1.41
85	AA	1597	C	C4'-C3'	5.11	1.58	1.53
85	AA	1991	C	C4'-C3'	-5.11	1.47	1.52
6	A5	39	GLY	C-N	-5.11	1.24	1.34
34	BA	197	A	C3'-O3'	5.11	1.49	1.42
34	BA	222	C	N1-C6	-5.11	1.34	1.37
34	BA	886	G	C1'-N9	-5.11	1.39	1.46
34	BA	1170	A	C3'-C2'	-5.11	1.47	1.52
34	BA	1470	G	C6-N1	-5.11	1.35	1.39
34	BA	1609	U	C4'-C3'	-5.11	1.47	1.52
34	BA	1658	G	C5-C4	-5.11	1.34	1.38
35	BB	58	G	C5-C6	-5.11	1.37	1.42
35	BB	108	G	C3'-C2'	-5.11	1.47	1.52
35	BB	592	G	C2-N2	-5.11	1.29	1.34
35	BB	1250	A	N1-C2	-5.11	1.29	1.34
35	BB	1274	G	C2-N2	-5.11	1.29	1.34
35	BB	1432	U	C1'-N1	-5.11	1.39	1.46
35	BB	1531	G	C5-C4	-5.11	1.34	1.38
37	BD	52	U	C4'-C3'	-5.11	1.47	1.52
41	BH	39	G	N1-C2	-5.11	1.33	1.37
78	Bs	35	ARG	CD-NE	5.11	1.55	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	84	C	N1-C6	-5.11	1.34	1.37
85	AA	128	U	P-O5'	-5.11	1.54	1.59
85	AA	535	G	C1'-N9	-5.11	1.39	1.46
85	AA	1575	G	N7-C5	-5.11	1.36	1.39
85	AA	1638	C	C4'-C3'	-5.11	1.47	1.52
85	AA	1674	G	C3'-C2'	-5.11	1.47	1.52
85	AA	1694	C	C2-N3	-5.11	1.31	1.35
85	AA	1697	C	P-O5'	-5.11	1.54	1.59
34	BA	1541	G	O3'-P	-5.11	1.55	1.61
34	BA	1784	G	C1'-N9	-5.11	1.39	1.46
35	BB	338	C	P-O5'	-5.11	1.54	1.59
35	BB	451	A	N9-C8	-5.11	1.33	1.37
35	BB	876	G	P-O5'	-5.11	1.54	1.59
35	BB	1202	G	C2-N3	5.11	1.36	1.32
36	BC	77	A	N3-C4	-5.11	1.31	1.34
40	BG	138	C	N1-C6	-5.11	1.34	1.37
40	BG	176	G	C4'-C3'	-5.11	1.47	1.52
40	BG	178	G	C4'-O4'	-5.11	1.39	1.45
41	BH	132	C	N1-C6	-5.11	1.34	1.37
85	AA	992	G	C2'-C1'	-5.11	1.47	1.53
85	AA	1892	G	C3'-C2'	-5.11	1.47	1.52
85	AA	2150	G	C2-N3	-5.11	1.28	1.32
34	BA	34	U	P-O5'	-5.11	1.54	1.59
34	BA	125	G	C6-O6	-5.11	1.19	1.24
34	BA	292	C	C1'-N1	-5.11	1.39	1.46
34	BA	665	C	C3'-C2'	-5.11	1.47	1.52
34	BA	903	C	C1'-N1	-5.11	1.39	1.46
34	BA	1527	G	N9-C8	-5.11	1.34	1.37
34	BA	1594	G	C2-N2	-5.11	1.29	1.34
35	BB	592	G	C3'-C2'	-5.11	1.47	1.52
35	BB	1250	A	C6-N1	-5.11	1.31	1.35
35	BB	1282	G	C2-N2	-5.11	1.29	1.34
36	BC	36	G	C3'-C2'	-5.11	1.47	1.52
36	BC	62	A	C8-N7	-5.11	1.27	1.31
37	BD	46	G	N7-C5	-5.11	1.36	1.39
38	BE	97	G	C2-N3	-5.11	1.28	1.32
40	BG	72	G	N7-C5	-5.11	1.36	1.39
40	BG	158	A	C5-C4	-5.11	1.35	1.38
41	BH	102	C	C2'-C1'	-5.11	1.47	1.53
41	BH	108	U	C3'-O3'	-5.11	1.34	1.42
85	AA	165	C	C2'-C1'	-5.11	1.47	1.53
85	AA	424	A	C8-N7	-5.11	1.27	1.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	668	A	P-O5'	-5.11	1.54	1.59
85	AA	975	G	C6-N1	-5.11	1.35	1.39
85	AA	1660	U	C1'-N1	-5.11	1.39	1.46
85	AA	1805	A	O3'-P	-5.11	1.55	1.61
85	AA	1830	U	C3'-C2'	-5.11	1.47	1.52
85	AA	1994	G	C2'-C1'	-5.11	1.47	1.53
85	AA	2080	U	N3-C4	-5.11	1.33	1.38
85	AA	2090	C	C2'-C1'	-5.11	1.47	1.53
85	AA	2216	A	O4'-C1'	-5.11	1.35	1.41
34	BA	52	G	C5-C6	-5.11	1.37	1.42
34	BA	280	A	C5-C4	-5.11	1.35	1.38
34	BA	389	U	C3'-C2'	-5.11	1.47	1.52
34	BA	401	A	N7-C5	-5.11	1.36	1.39
34	BA	667	U	N1-C2	-5.11	1.33	1.38
34	BA	936	A	C2'-C1'	-5.11	1.47	1.53
34	BA	1007	G	C3'-C2'	-5.11	1.47	1.52
34	BA	1052	G	N9-C8	-5.11	1.34	1.37
34	BA	1195	G	N1-C2	-5.11	1.33	1.37
34	BA	1577	U	C5'-C4'	-5.11	1.45	1.51
35	BB	359	A	N3-C4	-5.11	1.31	1.34
35	BB	382	U	N3-C4	-5.11	1.33	1.38
35	BB	453	C	C1'-N1	-5.11	1.39	1.46
35	BB	634	A	N7-C5	-5.11	1.36	1.39
35	BB	1071	G	C6-N1	-5.11	1.35	1.39
35	BB	1501	U	O3'-P	-5.11	1.55	1.61
36	BC	22	U	O3'-P	-5.11	1.55	1.61
37	BD	41	G	C5-C4	-5.11	1.34	1.38
38	BE	124	G	C5-C6	-5.11	1.37	1.42
40	BG	64	C	O3'-P	-5.11	1.55	1.61
40	BG	108	G	C5-C4	-5.11	1.34	1.38
41	BH	133	U	C2'-C1'	-5.11	1.47	1.53
85	AA	323	U	O3'-P	-5.11	1.55	1.61
85	AA	930	G	C4'-C3'	-5.11	1.47	1.52
85	AA	1155	A	O4'-C1'	-5.11	1.35	1.41
85	AA	1204	A	N3-C4	-5.11	1.31	1.34
85	AA	1239	C	C2'-C1'	-5.11	1.47	1.53
85	AA	1506	U	C4'-C3'	-5.11	1.47	1.52
85	AA	1638	C	P-O5'	-5.11	1.54	1.59
85	AA	1649	U	C2-N3	-5.11	1.34	1.37
85	AA	1654	G	C3'-C2'	-5.11	1.47	1.52
35	BB	502	C	N1-C6	-5.10	1.34	1.37
35	BB	1119	G	N1-C2	-5.10	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1427	A	N3-C4	-5.10	1.31	1.34
40	BG	163	G	C4'-O4'	-5.10	1.39	1.45
85	AA	66	U	C2'-C1'	-5.10	1.47	1.53
85	AA	1929	G	O4'-C1'	-5.10	1.35	1.41
34	BA	19	G	O4'-C1'	-5.10	1.35	1.41
34	BA	73	G	N1-C2	-5.10	1.33	1.37
34	BA	516	U	O3'-P	-5.10	1.55	1.61
34	BA	758	G	C8-N7	-5.10	1.27	1.30
34	BA	1223	C	C5'-C4'	-5.10	1.45	1.51
34	BA	1253	G	N1-C2	-5.10	1.33	1.37
35	BB	43	G	C3'-C2'	-5.10	1.47	1.52
35	BB	275	A	P-O5'	-5.10	1.54	1.59
35	BB	304	U	P-O5'	-5.10	1.54	1.59
35	BB	716	G	O3'-P	-5.10	1.55	1.61
35	BB	978	C	O3'-P	-5.10	1.55	1.61
36	BC	14	G	N3-C4	-5.10	1.31	1.35
39	BF	47	C	C4-N4	-5.10	1.29	1.33
85	AA	189	G	C2'-C1'	-5.10	1.47	1.53
85	AA	405	C	C2-N3	-5.10	1.31	1.35
85	AA	635	G	C5'-C4'	-5.10	1.45	1.51
85	AA	940	G	N7-C5	-5.10	1.36	1.39
85	AA	1666	U	C2'-C1'	-5.10	1.47	1.53
85	AA	2188	C	N1-C6	-5.10	1.34	1.37
86	AB	9	A	O4'-C1'	-5.10	1.35	1.41
34	BA	921	G	C3'-C2'	-5.10	1.47	1.52
35	BB	60	A	P-O5'	-5.10	1.54	1.59
85	AA	463	G	C5'-C4'	-5.10	1.45	1.51
85	AA	2150	G	C3'-C2'	-5.10	1.47	1.52
85	AA	2218	G	C5-C6	-5.10	1.37	1.42
34	BA	297	A	C8-N7	-5.10	1.27	1.31
34	BA	470	C	O3'-P	-5.10	1.55	1.61
34	BA	528	C	N1-C6	-5.10	1.34	1.37
34	BA	593	G	N3-C4	-5.10	1.31	1.35
34	BA	1117	G	C6-N1	-5.10	1.35	1.39
34	BA	1254	C	C3'-C2'	-5.10	1.47	1.52
34	BA	1337	A	C5-C4	-5.10	1.35	1.38
34	BA	1450	G	C1'-N9	-5.10	1.39	1.46
34	BA	1538	G	C2-N2	-5.10	1.29	1.34
35	BB	66	G	P-O5'	-5.10	1.54	1.59
35	BB	697	G	C2-N2	-5.10	1.29	1.34
36	BC	133	C	C3'-C2'	-5.10	1.47	1.52
36	BC	168	C	N1-C2	-5.10	1.35	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
40	BG	172	C	C4'-O4'	-5.10	1.39	1.45
85	AA	355	G	C2'-C1'	-5.10	1.47	1.53
85	AA	666	A	C5-C4	-5.10	1.35	1.38
85	AA	697	G	C1'-N9	-5.10	1.39	1.46
85	AA	1454	U	O3'-P	-5.10	1.55	1.61
85	AA	1677	A	N3-C4	-5.10	1.31	1.34
34	BA	340	U	C1'-N1	-5.10	1.39	1.46
34	BA	528	C	C4-C5	-5.10	1.38	1.43
34	BA	850	C	N1-C6	-5.10	1.34	1.37
34	BA	900	A	C3'-C2'	-5.10	1.47	1.52
34	BA	969	A	C3'-C2'	-5.10	1.47	1.52
34	BA	1059	U	C2-N3	-5.10	1.34	1.37
34	BA	1410	C	C3'-C2'	-5.10	1.47	1.52
34	BA	1472	G	N9-C8	-5.10	1.34	1.37
34	BA	1545	C	C1'-N1	-5.10	1.39	1.46
34	BA	1561	C	C2-N3	-5.10	1.31	1.35
35	BB	9	G	N1-C2	-5.10	1.33	1.37
35	BB	535	U	N1-C6	-5.10	1.33	1.38
35	BB	580	A	C1'-N9	-5.10	1.39	1.46
35	BB	793	A	C2'-C1'	-5.10	1.47	1.53
35	BB	1146	C	C4-N4	-5.10	1.29	1.33
35	BB	1308	G	N7-C5	-5.10	1.36	1.39
35	BB	1424	G	C4'-O4'	-5.10	1.39	1.45
35	BB	1476	C	C3'-C2'	-5.10	1.47	1.52
36	BC	20	C	C2'-C1'	-5.10	1.47	1.53
36	BC	44	A	C3'-C2'	-5.10	1.47	1.52
38	BE	125	C	O3'-P	-5.10	1.55	1.61
39	BF	17	U	C1'-N1	-5.10	1.39	1.46
40	BG	174	G	O4'-C1'	-5.10	1.35	1.41
85	AA	131	C	C4'-C3'	-5.10	1.47	1.52
85	AA	640	C	C5'-C4'	-5.10	1.45	1.51
85	AA	903	G	N9-C8	-5.10	1.34	1.37
85	AA	1244	A	C3'-C2'	-5.10	1.47	1.52
85	AA	1263	G	C2-N2	-5.10	1.29	1.34
85	AA	2059	A	C2'-C1'	-5.10	1.47	1.53
34	BA	64	A	O3'-P	-5.10	1.55	1.61
34	BA	92	G	C6-N1	-5.10	1.35	1.39
34	BA	96	G	C6-N1	-5.10	1.35	1.39
34	BA	697	A	N9-C8	-5.10	1.33	1.37
34	BA	1139	G	C4'-O4'	-5.10	1.39	1.45
35	BB	61	A	C3'-O3'	-5.10	1.35	1.42
35	BB	1362	G	C5-C6	-5.10	1.37	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	482	C	C4-C5	-5.10	1.38	1.43
85	AA	575	G	N7-C5	-5.10	1.36	1.39
34	BA	542	A	C3'-C2'	-5.09	1.47	1.52
34	BA	743	A	C1'-N9	-5.09	1.39	1.46
34	BA	785	G	C3'-C2'	-5.09	1.47	1.52
34	BA	792	A	C2'-C1'	-5.09	1.47	1.53
34	BA	877	U	O3'-P	-5.09	1.55	1.61
34	BA	1841	A	C3'-C2'	-5.09	1.47	1.52
35	BB	680	A	N9-C8	-5.09	1.33	1.37
35	BB	1210	U	C1'-N1	-5.09	1.39	1.46
36	BC	4	G	O4'-C1'	-5.09	1.35	1.41
38	BE	85	G	O4'-C1'	-5.09	1.35	1.41
38	BE	184	G	C6-N1	-5.09	1.35	1.39
85	AA	451	G	C2'-C1'	-5.09	1.47	1.53
85	AA	560	C	O3'-P	-5.09	1.55	1.61
85	AA	600	C	C5'-C4'	5.09	1.57	1.51
85	AA	654	A	C1'-N9	-5.09	1.39	1.46
85	AA	993	G	N9-C4	-5.09	1.33	1.38
85	AA	1370	G	C6-N1	-5.09	1.35	1.39
85	AA	1732	G	O3'-P	-5.09	1.55	1.61
34	BA	278	U	C2-N3	-5.09	1.34	1.37
34	BA	473	A	N7-C5	-5.09	1.36	1.39
34	BA	1064	A	N9-C8	-5.09	1.33	1.37
34	BA	1482	A	N9-C8	-5.09	1.33	1.37
34	BA	1799	G	C5-C4	-5.09	1.34	1.38
35	BB	470	C	C2'-C1'	-5.09	1.47	1.53
35	BB	486	G	C2'-C1'	-5.09	1.47	1.53
35	BB	1082	A	N3-C4	-5.09	1.31	1.34
36	BC	166	G	N3-C4	-5.09	1.31	1.35
40	BG	136	G	C6-N1	-5.09	1.35	1.39
85	AA	2208	G	C2-N2	-5.09	1.29	1.34
34	BA	267	G	C2'-C1'	-5.09	1.47	1.53
34	BA	677	U	O3'-P	-5.09	1.55	1.61
34	BA	1220	C	O4'-C1'	-5.09	1.35	1.41
34	BA	1340	G	C2-N2	-5.09	1.29	1.34
34	BA	1609	U	C2'-C1'	-5.09	1.47	1.53
34	BA	1674	G	P-O5'	-5.09	1.54	1.59
34	BA	1738	G	P-O5'	-5.09	1.54	1.59
35	BB	121	A	C8-N7	-5.09	1.27	1.31
35	BB	398	A	C6-N1	-5.09	1.31	1.35
35	BB	776	U	C3'-C2'	-5.09	1.47	1.52
35	BB	790	A	C5-C6	-5.09	1.36	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	873	C	C3'-C2'	-5.09	1.47	1.52
35	BB	1151	A	C1'-N9	-5.09	1.39	1.46
35	BB	1242	C	C1'-N1	-5.09	1.39	1.46
35	BB	1455	A	C8-N7	-5.09	1.27	1.31
35	BB	1539	C	C1'-N1	-5.09	1.39	1.46
36	BC	84	U	O3'-P	-5.09	1.55	1.61
37	BD	106	G	C3'-C2'	-5.09	1.47	1.52
41	BH	113	G	P-O5'	-5.09	1.54	1.59
85	AA	112	A	C4'-O4'	-5.09	1.39	1.45
85	AA	256	A	C5-C4	-5.09	1.35	1.38
85	AA	717	G	C2'-C1'	-5.09	1.47	1.53
85	AA	738	C	C3'-C2'	5.09	1.58	1.52
85	AA	1447	U	N3-C4	-5.09	1.33	1.38
85	AA	1716	U	C2-N3	-5.09	1.34	1.37
85	AA	2192	A	N9-C4	-5.09	1.34	1.37
34	BA	425	G	C5-C4	-5.09	1.34	1.38
34	BA	528	C	C3'-C2'	-5.09	1.47	1.52
34	BA	906	A	C2'-C1'	-5.09	1.47	1.53
34	BA	1122	G	C5-C6	-5.09	1.37	1.42
34	BA	1581	G	N1-C2	-5.09	1.33	1.37
34	BA	1596	C	N3-C4	-5.09	1.30	1.33
35	BB	643	G	C1'-N9	-5.09	1.39	1.46
35	BB	806	U	P-O5'	-5.09	1.54	1.59
35	BB	988	G	C2-N2	-5.09	1.29	1.34
35	BB	1206	G	O3'-P	-5.09	1.55	1.61
35	BB	1523	U	C3'-C2'	-5.09	1.47	1.52
36	BC	2	A	C8-N7	-5.09	1.27	1.31
36	BC	63	G	C2-N2	-5.09	1.29	1.34
38	BE	147	G	C4'-C3'	-5.09	1.47	1.52
40	BG	56	G	P-O5'	-5.09	1.54	1.59
40	BG	139	U	C4-O4	-5.09	1.19	1.23
85	AA	72	C	O3'-P	-5.09	1.55	1.61
85	AA	365	G	O3'-P	-5.09	1.55	1.61
85	AA	482	C	C2'-C1'	-5.09	1.47	1.53
85	AA	544	A	C5-C4	-5.09	1.35	1.38
85	AA	1252	A	P-O5'	-5.09	1.54	1.59
34	BA	979	G	C2-N3	-5.09	1.28	1.32
34	BA	1011	G	C2'-C1'	-5.09	1.47	1.53
34	BA	1677	C	C2'-C1'	-5.09	1.47	1.53
35	BB	489	A	O3'-P	-5.09	1.55	1.61
35	BB	793	A	C5-C4	-5.09	1.35	1.38
35	BB	1175	A	C1'-N9	-5.09	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1266	A	N9-C4	-5.09	1.34	1.37
35	BB	1396	G	C5-C6	-5.09	1.37	1.42
85	AA	89	C	N1-C6	-5.09	1.34	1.37
85	AA	1106	A	C4'-O4'	-5.09	1.39	1.45
86	AB	45	U	C2'-C1'	-5.09	1.47	1.53
34	BA	45	A	C3'-C2'	-5.09	1.47	1.52
34	BA	139	U	C5'-C4'	5.09	1.57	1.51
34	BA	306	G	N1-C2	-5.09	1.33	1.37
34	BA	381	A	C8-N7	-5.09	1.27	1.31
34	BA	429	G	C2-N2	-5.09	1.29	1.34
34	BA	1109	G	C2-N2	-5.09	1.29	1.34
34	BA	1488	C	C2'-C1'	-5.09	1.47	1.53
34	BA	1493	U	P-O5'	5.09	1.64	1.59
34	BA	1699	A	C6-N1	-5.09	1.31	1.35
37	BD	3	G	C6-N1	-5.09	1.35	1.39
37	BD	28	C	C3'-C2'	-5.09	1.47	1.52
85	AA	734	C	O3'-P	-5.09	1.55	1.61
85	AA	880	A	C5-C4	-5.09	1.35	1.38
85	AA	907	G	P-O5'	-5.09	1.54	1.59
85	AA	2145	G	C5-C6	-5.09	1.37	1.42
34	BA	1278	A	C5'-C4'	-5.08	1.45	1.51
35	BB	641	C	O3'-P	-5.08	1.55	1.61
35	BB	1259	A	N9-C8	-5.08	1.33	1.37
38	BE	59	U	C5'-C4'	-5.08	1.45	1.51
85	AA	374	C	C4'-C3'	-5.08	1.47	1.52
85	AA	378	A	N1-C2	-5.08	1.29	1.34
85	AA	879	G	C2-N2	-5.08	1.29	1.34
85	AA	1503	G	C6-N1	-5.08	1.35	1.39
34	BA	79	C	N3-C4	-5.08	1.30	1.33
34	BA	397	A	C5-C4	-5.08	1.35	1.38
34	BA	977	G	C3'-C2'	-5.08	1.47	1.52
35	BB	42	A	C5-C4	-5.08	1.35	1.38
35	BB	498	G	C1'-N9	-5.08	1.39	1.46
35	BB	540	G	N9-C8	-5.08	1.34	1.37
35	BB	1009	U	O3'-P	-5.08	1.55	1.61
35	BB	1272	G	C3'-C2'	-5.08	1.47	1.52
35	BB	1490	G	C4'-O4'	5.08	1.52	1.45
37	BD	18	G	C6-N1	-5.08	1.35	1.39
40	BG	153	C	C4'-C3'	-5.08	1.47	1.52
41	BH	25	A	P-O5'	-5.08	1.54	1.59
85	AA	162	A	N9-C8	-5.08	1.33	1.37
85	AA	277	G	C6-N1	-5.08	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1110	A	N1-C2	-5.08	1.29	1.34
85	AA	1982	C	P-O5'	-5.08	1.54	1.59
85	AA	2004	U	C4'-C3'	-5.08	1.47	1.52
85	AA	2012	G	C2-N2	-5.08	1.29	1.34
85	AA	2037	A	C3'-C2'	-5.08	1.47	1.52
34	BA	188	C	C2'-C1'	-5.08	1.47	1.53
34	BA	417	A	C3'-C2'	-5.08	1.47	1.52
34	BA	684	G	O5'-C5'	5.08	1.52	1.44
34	BA	1240	G	C5-C4	-5.08	1.34	1.38
34	BA	1493	U	C1'-N1	5.08	1.56	1.48
34	BA	1566	G	N3-C4	-5.08	1.31	1.35
34	BA	1691	G	O3'-P	-5.08	1.55	1.61
34	BA	1833	G	C4'-C3'	-5.08	1.47	1.52
35	BB	854	G	C2'-C1'	-5.08	1.47	1.53
35	BB	1310	C	C4-N4	-5.08	1.29	1.33
35	BB	1315	C	C3'-C2'	-5.08	1.47	1.52
35	BB	1376	G	N9-C4	-5.08	1.33	1.38
36	BC	18	G	N9-C8	-5.08	1.34	1.37
36	BC	20	C	O4'-C1'	-5.08	1.35	1.41
37	BD	39	C	O3'-P	-5.08	1.55	1.61
40	BG	32	U	N3-C4	-5.08	1.33	1.38
40	BG	76	C	C1'-N1	-5.08	1.39	1.46
85	AA	577	U	C3'-O3'	5.08	1.49	1.42
85	AA	660	G	C1'-N9	-5.08	1.39	1.46
85	AA	708	G	C2-N3	-5.08	1.28	1.32
85	AA	820	G	N1-C2	-5.08	1.33	1.37
85	AA	931	G	C5-C4	-5.08	1.34	1.38
85	AA	1167	G	C2-N2	-5.08	1.29	1.34
85	AA	1528	A	C5-C4	-5.08	1.35	1.38
85	AA	1845	G	C5-C4	-5.08	1.34	1.38
34	BA	607	C	C2'-C1'	-5.08	1.47	1.53
34	BA	794	G	C5'-C4'	-5.08	1.45	1.51
34	BA	971	G	C2-N2	-5.08	1.29	1.34
34	BA	1335	A	O4'-C1'	-5.08	1.35	1.41
34	BA	1341	A	N9-C8	-5.08	1.33	1.37
35	BB	1406	C	C4'-O4'	-5.08	1.39	1.45
85	AA	171	U	C4'-O4'	-5.08	1.39	1.45
85	AA	936	C	C4'-C3'	-5.08	1.47	1.52
85	AA	1046	C	P-O5'	-5.08	1.54	1.59
34	BA	48	C	C4-N4	-5.08	1.29	1.33
34	BA	67	A	P-O5'	-5.08	1.54	1.59
34	BA	221	G	N7-C5	-5.08	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	269	G	C1'-N9	-5.08	1.39	1.46
34	BA	787	A	C6-N6	-5.08	1.29	1.33
34	BA	1009	G	C4'-C3'	-5.08	1.47	1.52
34	BA	1075	U	C4'-C3'	-5.08	1.47	1.52
34	BA	1529	G	C1'-N9	-5.08	1.39	1.46
34	BA	1647	G	C5'-C4'	-5.08	1.45	1.51
35	BB	416	U	C1'-N1	-5.08	1.39	1.46
35	BB	569	G	C1'-N9	-5.08	1.39	1.46
35	BB	773	G	N9-C4	-5.08	1.33	1.38
35	BB	1189	C	C2-N3	-5.08	1.31	1.35
35	BB	1434	G	C8-N7	-5.08	1.27	1.30
35	BB	1541	G	P-O5'	-5.08	1.54	1.59
36	BC	117	A	C5-C4	-5.08	1.35	1.38
37	BD	64	A	C3'-C2'	-5.08	1.47	1.52
85	AA	16	G	P-O5'	-5.08	1.54	1.59
85	AA	1722	G	C2'-C1'	-5.08	1.47	1.53
85	AA	2242	U	N3-C4	-5.08	1.33	1.38
34	BA	93	A	N9-C8	-5.08	1.33	1.37
34	BA	1490	U	C2'-C1'	-5.08	1.47	1.53
35	BB	1055	G	C2-N2	-5.08	1.29	1.34
37	BD	108	G	C2-N2	-5.08	1.29	1.34
38	BE	8	G	C5'-C4'	5.08	1.57	1.51
39	BF	44	C	C2'-C1'	-5.08	1.47	1.53
85	AA	297	A	P-O5'	-5.08	1.54	1.59
34	BA	42	A	C2'-C1'	-5.08	1.47	1.53
34	BA	68	A	N3-C4	-5.08	1.31	1.34
34	BA	380	A	C3'-C2'	-5.08	1.47	1.52
34	BA	452	A	N7-C5	-5.08	1.36	1.39
34	BA	516	U	C1'-N1	-5.08	1.39	1.46
34	BA	680	C	N3-C4	-5.08	1.30	1.33
34	BA	717	U	C2-N3	-5.08	1.34	1.37
34	BA	985	C	C1'-N1	-5.08	1.39	1.46
34	BA	1102	A	N9-C4	-5.08	1.34	1.37
34	BA	1231	C	N3-C4	-5.08	1.30	1.33
34	BA	1249	G	C3'-C2'	-5.08	1.47	1.52
34	BA	1331	G	C6-N1	-5.08	1.35	1.39
34	BA	1465	C	N1-C6	-5.08	1.34	1.37
35	BB	34	G	N1-C2	-5.08	1.33	1.37
35	BB	391	G	C1'-N9	-5.08	1.39	1.46
35	BB	422	U	P-O5'	-5.08	1.54	1.59
35	BB	502	C	O3'-P	-5.08	1.55	1.61
35	BB	1128	U	C3'-C2'	-5.08	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1365	G	C2'-C1'	-5.08	1.47	1.53
40	BG	44	G	C6-N1	-5.08	1.35	1.39
41	BH	116	A	P-O5'	-5.08	1.54	1.59
85	AA	50	C	N3-C4	-5.08	1.30	1.33
85	AA	288	G	O3'-P	-5.08	1.55	1.61
85	AA	475	A	N7-C5	-5.08	1.36	1.39
85	AA	1590	A	O3'-P	-5.08	1.55	1.61
86	AB	51	U	P-O5'	-5.08	1.54	1.59
34	BA	252	A	P-O5'	-5.07	1.54	1.59
34	BA	466	G	N9-C8	-5.07	1.34	1.37
34	BA	1030	C	C4-N4	-5.07	1.29	1.33
34	BA	1587	C	C1'-N1	-5.07	1.39	1.46
35	BB	659	C	C3'-C2'	-5.07	1.47	1.52
35	BB	1405	G	C4'-O4'	-5.07	1.39	1.45
35	BB	1407	U	N3-C4	-5.07	1.33	1.38
38	BE	95	G	C2-N2	-5.07	1.29	1.34
41	BH	15	A	N9-C8	-5.07	1.33	1.37
85	AA	175	A	N9-C4	-5.07	1.34	1.37
85	AA	557	G	O5'-C5'	-5.07	1.34	1.42
85	AA	581	A	C2'-C1'	-5.07	1.47	1.53
85	AA	597	A	N3-C4	-5.07	1.31	1.34
85	AA	1365	U	P-O5'	-5.07	1.54	1.59
85	AA	1854	U	C2-N3	-5.07	1.34	1.37
35	BB	833	G	C5-C4	-5.07	1.34	1.38
35	BB	904	C	P-O5'	-5.07	1.54	1.59
35	BB	1213	U	C3'-C2'	-5.07	1.47	1.52
35	BB	1416	A	N3-C4	-5.07	1.31	1.34
35	BB	1460	G	P-O5'	-5.07	1.54	1.59
37	BD	73	U	C2'-C1'	-5.07	1.47	1.53
40	BG	93	U	O4'-C1'	-5.07	1.35	1.41
85	AA	277	G	N9-C4	-5.07	1.33	1.38
85	AA	1868	G	N7-C5	-5.07	1.36	1.39
34	BA	197	A	C8-N7	-5.07	1.28	1.31
34	BA	517	A	O4'-C1'	-5.07	1.35	1.41
34	BA	935	A	C1'-N9	-5.07	1.39	1.46
34	BA	1135	U	C2'-C1'	-5.07	1.47	1.53
34	BA	1211	G	O4'-C1'	-5.07	1.35	1.41
35	BB	18	A	C4'-C3'	-5.07	1.47	1.52
35	BB	113	C	C4'-C3'	-5.07	1.47	1.52
35	BB	581	U	O3'-P	-5.07	1.55	1.61
35	BB	636	G	C2-N2	-5.07	1.29	1.34
35	BB	1130	U	C4'-C3'	-5.07	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1271	A	N9-C8	-5.07	1.33	1.37
39	BF	23	G	N3-C4	-5.07	1.31	1.35
74	Bo	79	VAL	CA-C	-5.07	1.39	1.52
85	AA	43	A	P-O5'	-5.07	1.54	1.59
85	AA	375	C	C4-N4	-5.07	1.29	1.33
85	AA	525	C	C1'-N1	-5.07	1.39	1.46
85	AA	740	A	N7-C5	-5.07	1.36	1.39
85	AA	1233	G	C6-N1	-5.07	1.36	1.39
85	AA	1561	A	O3'-P	-5.07	1.55	1.61
85	AA	2022	A	C8-N7	-5.07	1.28	1.31
34	BA	1146	U	C2-N3	-5.07	1.34	1.37
34	BA	1171	C	C3'-C2'	-5.07	1.47	1.52
35	BB	1115	G	C2-N2	-5.07	1.29	1.34
36	BC	74	U	C4-C5	-5.07	1.39	1.43
38	BE	8	G	C2-N3	5.07	1.36	1.32
85	AA	857	G	C2'-C1'	-5.07	1.47	1.53
85	AA	924	A	N3-C4	-5.07	1.31	1.34
85	AA	1185	G	C1'-N9	-5.07	1.39	1.46
85	AA	1223	A	O3'-P	-5.07	1.55	1.61
34	BA	178	C	O4'-C1'	-5.07	1.35	1.41
34	BA	269	G	N1-C2	-5.07	1.33	1.37
34	BA	491	U	P-O5'	-5.07	1.54	1.59
34	BA	492	G	C3'-C2'	-5.07	1.47	1.52
34	BA	954	U	C3'-C2'	-5.07	1.47	1.52
34	BA	1476	G	N1-C2	-5.07	1.33	1.37
34	BA	1516	G	P-O5'	-5.07	1.54	1.59
34	BA	1809	G	O4'-C1'	-5.07	1.35	1.41
35	BB	22	A	C6-N6	-5.07	1.29	1.33
35	BB	367	C	O3'-P	-5.07	1.55	1.61
35	BB	630	A	C5'-C4'	-5.07	1.45	1.51
35	BB	651	G	N3-C4	-5.07	1.31	1.35
35	BB	678	U	N3-C4	-5.07	1.33	1.38
35	BB	1187	G	O3'-P	-5.07	1.55	1.61
36	BC	156	A	N7-C5	-5.07	1.36	1.39
37	BD	10	C	C1'-N1	-5.07	1.39	1.46
38	BE	151	C	C4'-C3'	-5.07	1.47	1.52
40	BG	7	U	C2-N3	-5.07	1.34	1.37
85	AA	376	C	C2-N3	-5.07	1.31	1.35
85	AA	629	A	C3'-C2'	-5.07	1.47	1.52
85	AA	683	U	P-O5'	-5.07	1.54	1.59
85	AA	727	U	C5'-C4'	5.07	1.57	1.51
85	AA	1594	G	N9-C4	-5.07	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	1922	A	O4'-C1'	-5.07	1.35	1.41
34	BA	340	U	C2'-C1'	-5.07	1.47	1.53
34	BA	665	C	O3'-P	-5.07	1.55	1.61
34	BA	856	G	N1-C2	-5.07	1.33	1.37
34	BA	967	C	C4'-C3'	-5.07	1.47	1.52
34	BA	1822	U	C4'-C3'	-5.07	1.47	1.52
35	BB	51	U	P-O5'	-5.07	1.54	1.59
35	BB	73	G	N9-C8	-5.07	1.34	1.37
35	BB	375	G	P-O5'	-5.07	1.54	1.59
35	BB	549	U	C2'-C1'	-5.07	1.47	1.53
35	BB	1520	C	O3'-P	-5.07	1.55	1.61
37	BD	20	C	C3'-C2'	-5.07	1.47	1.52
37	BD	37	G	C1'-N9	-5.07	1.39	1.46
37	BD	105	G	C6-N1	-5.07	1.36	1.39
40	BG	45	G	N1-C2	-5.07	1.33	1.37
85	AA	545	A	C3'-C2'	-5.07	1.47	1.52
85	AA	1279	A	C5-C4	-5.07	1.35	1.38
85	AA	1488	G	C2-N2	-5.07	1.29	1.34
85	AA	1563	U	C2'-C1'	-5.07	1.47	1.53
85	AA	2034	G	C2-N2	-5.07	1.29	1.34
85	AA	2193	A	N3-C4	-5.07	1.31	1.34
34	BA	146	G	C2-N3	-5.06	1.28	1.32
34	BA	524	G	N3-C4	-5.06	1.31	1.35
34	BA	837	U	C3'-O3'	-5.06	1.35	1.42
35	BB	988	G	C1'-N9	-5.06	1.39	1.46
35	BB	1120	A	C2'-C1'	-5.06	1.47	1.53
35	BB	1127	A	C2'-C1'	-5.06	1.47	1.53
35	BB	1295	A	C5-C4	-5.06	1.35	1.38
40	BG	54	G	O4'-C1'	-5.06	1.35	1.41
85	AA	30	G	N1-C2	-5.06	1.33	1.37
85	AA	448	G	C1'-N9	-5.06	1.39	1.46
85	AA	1049	G	N9-C4	-5.06	1.33	1.38
85	AA	1713	A	O3'-P	-5.06	1.55	1.61
85	AA	1850	G	N7-C5	-5.06	1.36	1.39
34	BA	295	G	C6-N1	-5.06	1.36	1.39
34	BA	1329	U	N1-C2	5.06	1.43	1.38
34	BA	1671	A	C5-C4	-5.06	1.35	1.38
35	BB	1155	U	N1-C6	-5.06	1.33	1.38
35	BB	1185	G	C4'-O4'	-5.06	1.39	1.45
35	BB	1454	G	C2-N2	-5.06	1.29	1.34
36	BC	113	G	N3-C4	-5.06	1.31	1.35
38	BE	49	A	C2'-C1'	-5.06	1.47	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	191	U	C2'-C1'	-5.06	1.47	1.53
62	Bc	13	ARG	CA-CB	5.06	1.65	1.53
85	AA	312	G	C2-N2	-5.06	1.29	1.34
85	AA	506	G	C3'-C2'	-5.06	1.47	1.52
85	AA	933	U	C2-N3	-5.06	1.34	1.37
85	AA	1113	G	C4'-C3'	-5.06	1.47	1.52
85	AA	1679	U	O3'-P	-5.06	1.55	1.61
85	AA	2143	U	O3'-P	-5.06	1.55	1.61
34	BA	333	A	C3'-C2'	-5.06	1.47	1.52
34	BA	1001	G	N7-C5	-5.06	1.36	1.39
34	BA	1285	G	C2'-C1'	-5.06	1.47	1.53
34	BA	1782	C	C4'-C3'	-5.06	1.47	1.52
35	BB	53	C	O4'-C1'	-5.06	1.35	1.41
40	BG	130	G	C8-N7	-5.06	1.27	1.30
85	AA	134	U	C2'-C1'	-5.06	1.47	1.53
85	AA	733	C	C2'-C1'	-5.06	1.47	1.53
85	AA	836	A	C1'-N9	-5.06	1.39	1.46
85	AA	879	G	C3'-C2'	-5.06	1.47	1.52
34	BA	37	A	C2'-C1'	-5.06	1.47	1.53
34	BA	608	G	C3'-O3'	5.06	1.49	1.42
34	BA	1163	G	N3-C4	-5.06	1.31	1.35
34	BA	1284	G	C5-C4	-5.06	1.34	1.38
34	BA	1300	G	C2-N3	-5.06	1.28	1.32
34	BA	1840	C	C1'-N1	-5.06	1.39	1.46
35	BB	48	G	C2'-C1'	-5.06	1.47	1.53
35	BB	509	A	C6-N6	-5.06	1.29	1.33
35	BB	1026	G	N3-C4	-5.06	1.31	1.35
36	BC	2	A	O4'-C1'	-5.06	1.35	1.41
39	BF	25	G	C8-N7	-5.06	1.27	1.30
41	BH	101	A	C2'-C1'	-5.06	1.47	1.53
85	AA	599	C	C4'-O4'	-5.06	1.39	1.45
85	AA	1179	A	O3'-P	-5.06	1.55	1.61
85	AA	1875	A	N3-C4	-5.06	1.31	1.34
85	AA	2128	G	P-O5'	-5.06	1.54	1.59
85	AA	2142	A	P-O5'	-5.06	1.54	1.59
34	BA	342	U	C3'-C2'	-5.06	1.47	1.52
34	BA	480	G	C2-N3	-5.06	1.28	1.32
34	BA	911	G	C3'-C2'	-5.06	1.47	1.52
34	BA	1045	C	C2'-C1'	-5.06	1.47	1.53
34	BA	1690	U	C2'-C1'	-5.06	1.47	1.53
35	BB	273	G	P-O5'	-5.06	1.54	1.59
35	BB	382	U	C2-N3	5.06	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	394	A	N9-C4	-5.06	1.34	1.37
35	BB	482	A	P-O5'	-5.06	1.54	1.59
35	BB	682	U	N1-C2	-5.06	1.33	1.38
35	BB	1097	U	C2'-C1'	-5.06	1.47	1.53
35	BB	1117	G	C5-C4	-5.06	1.34	1.38
35	BB	1263	A	C5'-C4'	-5.06	1.45	1.51
41	BH	126	C	C3'-C2'	-5.06	1.47	1.52
85	AA	30	G	C6-N1	-5.06	1.36	1.39
85	AA	177	A	C5-C4	-5.06	1.35	1.38
85	AA	577	U	C2-N3	-5.06	1.34	1.37
85	AA	630	A	C2'-C1'	-5.06	1.47	1.53
85	AA	682	C	C2'-C1'	-5.06	1.47	1.53
85	AA	1238	U	C3'-C2'	-5.06	1.47	1.52
85	AA	2223	C	O3'-P	-5.06	1.55	1.61
85	AA	2226	U	N3-C4	-5.06	1.33	1.38
34	BA	961	C	P-O5'	-5.06	1.54	1.59
34	BA	1232	C	C2'-C1'	-5.06	1.47	1.53
34	BA	1311	G	P-O5'	-5.06	1.54	1.59
34	BA	1719	G	C3'-C2'	-5.06	1.47	1.52
34	BA	1811	A	P-O5'	-5.06	1.54	1.59
41	BH	32	U	C5-C6	-5.06	1.29	1.34
85	AA	332	A	N9-C4	-5.06	1.34	1.37
85	AA	663	C	C2-N3	-5.06	1.31	1.35
85	AA	1279	A	O3'-P	-5.06	1.55	1.61
85	AA	1367	C	O3'-P	-5.06	1.55	1.61
34	BA	452	A	C2'-C1'	-5.05	1.47	1.53
34	BA	460	G	C2-N2	-5.05	1.29	1.34
34	BA	731	A	P-O5'	-5.05	1.54	1.59
34	BA	775	C	N1-C2	-5.05	1.35	1.40
34	BA	850	C	C5'-C4'	-5.05	1.45	1.51
34	BA	966	G	C2'-C1'	-5.05	1.47	1.53
34	BA	1457	C	N1-C6	-5.05	1.34	1.37
34	BA	1503	U	C3'-O3'	-5.05	1.35	1.42
35	BB	460	C	C5'-C4'	5.05	1.57	1.51
35	BB	498	G	C3'-C2'	-5.05	1.47	1.52
35	BB	575	C	C3'-C2'	-5.05	1.47	1.52
35	BB	900	C	O3'-P	-5.05	1.55	1.61
35	BB	1136	G	N9-C8	-5.05	1.34	1.37
35	BB	1266	A	C2'-C1'	-5.05	1.47	1.53
38	BE	183	C	C3'-C2'	5.05	1.58	1.52
85	AA	338	G	C6-N1	-5.05	1.36	1.39
85	AA	420	C	C1'-N1	-5.05	1.39	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	755	G	C4'-C3'	-5.05	1.47	1.52
85	AA	1455	C	C4'-C3'	-5.05	1.47	1.52
85	AA	1525	C	C1'-N1	-5.05	1.39	1.46
34	BA	32	A	N9-C8	-5.05	1.33	1.37
34	BA	1599	A	N9-C8	-5.05	1.33	1.37
34	BA	1630	A	P-O5'	-5.05	1.54	1.59
35	BB	799	A	N9-C4	-5.05	1.34	1.37
35	BB	1380	G	P-O5'	-5.05	1.54	1.59
35	BB	1542	C	O3'-P	-5.05	1.55	1.61
85	AA	192	G	N9-C4	5.05	1.42	1.38
85	AA	1207	C	C4'-O4'	-5.05	1.39	1.45
85	AA	1762	G	P-O5'	-5.05	1.54	1.59
5	A4	69	PRO	CA-C	-5.05	1.42	1.52
34	BA	77	C	C2'-C1'	-5.05	1.47	1.53
34	BA	315	U	C4'-O4'	-5.05	1.39	1.45
34	BA	735	A	C4'-C3'	-5.05	1.47	1.52
34	BA	861	C	C1'-N1	-5.05	1.39	1.46
34	BA	1213	A	C5'-C4'	-5.05	1.45	1.51
34	BA	1303	U	C4'-C3'	-5.05	1.47	1.52
34	BA	1529	G	N9-C8	-5.05	1.34	1.37
34	BA	1695	G	C5-C6	-5.05	1.37	1.42
34	BA	1710	C	C5'-C4'	-5.05	1.45	1.51
35	BB	71	A	N3-C4	-5.05	1.31	1.34
35	BB	74	U	C4'-C3'	-5.05	1.47	1.52
35	BB	125	G	C2'-C1'	-5.05	1.47	1.53
35	BB	611	U	N3-C4	-5.05	1.33	1.38
35	BB	805	G	C1'-N9	-5.05	1.39	1.46
35	BB	1457	A	N7-C5	-5.05	1.36	1.39
40	BG	78	C	C4'-C3'	-5.05	1.47	1.52
40	BG	179	C	C2'-C1'	-5.05	1.47	1.53
85	AA	316	C	C2-N3	-5.05	1.31	1.35
85	AA	560	C	C2-N3	-5.05	1.31	1.35
85	AA	723	U	O3'-P	-5.05	1.55	1.61
85	AA	752	C	P-O5'	-5.05	1.54	1.59
85	AA	879	G	C5-C6	-5.05	1.37	1.42
85	AA	1298	G	C1'-N9	-5.05	1.39	1.46
85	AA	1472	G	C1'-N9	-5.05	1.39	1.46
85	AA	1506	U	C1'-N1	-5.05	1.39	1.46
85	AA	1524	A	C5-C4	-5.05	1.35	1.38
85	AA	1541	G	C1'-N9	-5.05	1.39	1.46
85	AA	2064	A	N3-C4	-5.05	1.31	1.34
34	BA	81	C	C3'-C2'	-5.05	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	724	A	N3-C4	-5.05	1.31	1.34
34	BA	1001	G	C8-N7	-5.05	1.27	1.30
34	BA	1046	G	O4'-C1'	-5.05	1.35	1.41
35	BB	362	A	P-O5'	-5.05	1.54	1.59
35	BB	507	G	C2-N2	-5.05	1.29	1.34
35	BB	518	G	C5-C4	-5.05	1.34	1.38
35	BB	1046	C	N1-C6	-5.05	1.34	1.37
40	BG	38	A	O4'-C1'	-5.05	1.35	1.41
85	AA	189	G	C3'-C2'	-5.05	1.47	1.52
85	AA	573	U	C4'-C3'	-5.05	1.47	1.52
85	AA	1671	G	N3-C4	-5.05	1.31	1.35
85	AA	2194	U	C3'-C2'	-5.05	1.47	1.52
85	AA	2196	G	N9-C8	-5.05	1.34	1.37
34	BA	753	G	P-O5'	-5.05	1.54	1.59
34	BA	1498	A	C2'-C1'	-5.05	1.47	1.53
34	BA	1544	G	N1-C2	-5.05	1.33	1.37
34	BA	1828	A	C5-C4	-5.05	1.35	1.38
35	BB	66	G	N1-C2	-5.05	1.33	1.37
35	BB	76	C	C2-N3	-5.05	1.31	1.35
35	BB	1146	C	C3'-C2'	-5.05	1.47	1.52
36	BC	146	U	N3-C4	-5.05	1.33	1.38
37	BD	62	A	C5-C4	-5.05	1.35	1.38
85	AA	260	A	C1'-N9	-5.05	1.39	1.46
85	AA	761	G	C6-N1	-5.05	1.36	1.39
34	BA	213	A	O3'-P	-5.05	1.55	1.61
34	BA	821	G	C2-N2	-5.05	1.29	1.34
34	BA	1614	G	C6-N1	-5.05	1.36	1.39
35	BB	1422	G	C3'-C2'	-5.05	1.47	1.52
35	BB	1536	G	C2'-C1'	-5.05	1.47	1.53
36	BC	24	G	C6-N1	-5.05	1.36	1.39
40	BG	96	C	O4'-C1'	-5.05	1.35	1.41
85	AA	1148	G	C4'-C3'	-5.05	1.47	1.52
85	AA	1825	A	C2'-C1'	-5.05	1.47	1.53
85	AA	2200	A	P-O5'	-5.05	1.54	1.59
34	BA	329	G	O3'-P	-5.04	1.55	1.61
35	BB	1417	C	P-O5'	-5.04	1.54	1.59
36	BC	15	G	N7-C5	-5.04	1.36	1.39
40	BG	6	A	C8-N7	-5.04	1.28	1.31
85	AA	1112	G	C2'-C1'	-5.04	1.47	1.53
85	AA	1925	A	N9-C8	-5.04	1.33	1.37
34	BA	110	C	C3'-O3'	-5.04	1.35	1.42
34	BA	243	C	C4'-C3'	-5.04	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	732	A	N9-C8	-5.04	1.33	1.37
34	BA	1175	G	C5-C6	-5.04	1.37	1.42
34	BA	1292	A	C5'-C4'	-5.04	1.45	1.51
34	BA	1541	G	C2-N2	-5.04	1.29	1.34
35	BB	543	G	C3'-C2'	-5.04	1.47	1.52
35	BB	677	U	C2'-C1'	-5.04	1.47	1.53
35	BB	723	A	P-O5'	-5.04	1.54	1.59
35	BB	1053	G	N9-C8	-5.04	1.34	1.37
35	BB	1221	G	N9-C8	-5.04	1.34	1.37
38	BE	34	C	O3'-P	-5.04	1.55	1.61
40	BG	12	A	C4'-C3'	-5.04	1.47	1.52
40	BG	160	C	C2'-C1'	-5.04	1.47	1.53
85	AA	204	U	C2-N3	-5.04	1.34	1.37
85	AA	271	A	C2'-C1'	-5.04	1.47	1.53
85	AA	277	G	C2-N2	-5.04	1.29	1.34
85	AA	644	A	C5-C4	-5.04	1.35	1.38
86	AB	1	G	C2'-C1'	-5.04	1.47	1.53
86	AB	53	G	P-O5'	-5.04	1.54	1.59
34	BA	194	G	O3'-P	-5.04	1.55	1.61
34	BA	497	U	O3'-P	-5.04	1.55	1.61
34	BA	672	G	C8-N7	-5.04	1.27	1.30
34	BA	676	G	N3-C4	-5.04	1.31	1.35
34	BA	1223	C	C4'-O4'	-5.04	1.39	1.45
34	BA	1455	C	C5'-C4'	5.04	1.57	1.51
34	BA	1475	G	C2-N2	-5.04	1.29	1.34
34	BA	1657	A	C6-N1	-5.04	1.32	1.35
35	BB	672	C	O3'-P	-5.04	1.55	1.61
35	BB	826	G	C2-N2	-5.04	1.29	1.34
35	BB	1397	G	C5-C4	-5.04	1.34	1.38
35	BB	1447	U	C2-N3	-5.04	1.34	1.37
36	BC	146	U	C1'-N1	-5.04	1.39	1.46
38	BE	3	G	N7-C5	-5.04	1.36	1.39
40	BG	19	C	C4'-C3'	-5.04	1.47	1.52
85	AA	594	C	N1-C6	-5.04	1.34	1.37
85	AA	1542	A	N3-C4	-5.04	1.31	1.34
34	BA	25	C	C4-N4	-5.04	1.29	1.33
34	BA	1156	U	C3'-C2'	-5.04	1.47	1.52
34	BA	1231	C	C2-N3	-5.04	1.31	1.35
34	BA	1615	A	C5-C4	-5.04	1.35	1.38
34	BA	10	G	C6-N1	-5.04	1.36	1.39
34	BA	354	G	C1'-N9	-5.04	1.39	1.46
34	BA	433	G	N9-C8	-5.04	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1275	G	C4'-O4'	-5.04	1.39	1.45
34	BA	1449	U	P-O5'	-5.04	1.54	1.59
34	BA	1786	C	O3'-P	-5.04	1.55	1.61
34	BA	1845	G	C3'-C2'	-5.04	1.47	1.52
35	BB	1196	A	N3-C4	-5.04	1.31	1.34
35	BB	1408	G	C5-C4	-5.04	1.34	1.38
36	BC	61	A	C5-C4	-5.04	1.35	1.38
38	BE	4	A	N3-C4	-5.04	1.31	1.34
85	AA	465	A	C4'-C3'	-5.04	1.47	1.52
85	AA	603	C	O3'-P	-5.04	1.55	1.61
85	AA	683	U	C1'-N1	-5.04	1.39	1.46
85	AA	798	A	C2'-C1'	-5.04	1.47	1.53
85	AA	1212	C	C4-N4	-5.04	1.29	1.33
85	AA	1409	U	P-O5'	-5.04	1.54	1.59
34	BA	570	G	O3'-P	-5.04	1.55	1.61
34	BA	1161	G	C2'-C1'	-5.04	1.47	1.53
34	BA	1244	G	C1'-N9	-5.04	1.39	1.46
35	BB	1477	C	C2-N3	-5.04	1.31	1.35
38	BE	23	G	C5-C4	-5.04	1.34	1.38
41	BH	69	C	C3'-C2'	-5.04	1.47	1.52
85	AA	731	U	C2'-C1'	-5.04	1.47	1.53
85	AA	1492	U	P-O5'	-5.04	1.54	1.59
34	BA	68	A	C8-N7	-5.04	1.28	1.31
34	BA	172	A	C8-N7	-5.04	1.28	1.31
34	BA	276	C	C4-N4	-5.04	1.29	1.33
35	BB	312	U	P-O5'	-5.04	1.54	1.59
35	BB	599	U	C3'-C2'	-5.04	1.47	1.52
35	BB	1270	C	C2-N3	-5.04	1.31	1.35
35	BB	1534	U	O4'-C1'	-5.04	1.35	1.41
35	BB	1536	G	P-O5'	-5.04	1.54	1.59
36	BC	40	A	C1'-N9	-5.04	1.39	1.46
36	BC	68	A	C6-N1	-5.04	1.32	1.35
37	BD	50	A	C3'-C2'	-5.04	1.47	1.52
38	BE	147	G	C3'-C2'	-5.04	1.47	1.52
38	BE	179	A	N3-C4	-5.04	1.31	1.34
85	AA	158	C	C3'-C2'	-5.04	1.47	1.52
85	AA	365	G	C2-N2	-5.04	1.29	1.34
85	AA	504	U	C5'-C4'	-5.04	1.45	1.51
85	AA	582	A	C2'-C1'	-5.04	1.47	1.53
85	AA	1171	C	C4'-C3'	-5.04	1.47	1.52
85	AA	1582	U	C2'-C1'	-5.04	1.47	1.53
85	AA	2116	U	C4'-C3'	-5.04	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
6	A5	200	GLY	N-CA	-5.03	1.38	1.46
34	BA	142	A	C3'-C2'	-5.03	1.47	1.52
34	BA	447	U	O4'-C1'	-5.03	1.35	1.41
34	BA	1054	U	N3-C4	-5.03	1.33	1.38
34	BA	1419	A	N9-C8	-5.03	1.33	1.37
35	BB	662	G	N9-C8	-5.03	1.34	1.37
35	BB	1120	A	C5'-C4'	-5.03	1.45	1.51
35	BB	1365	G	N1-C2	-5.03	1.33	1.37
36	BC	163	A	C5-C6	-5.03	1.36	1.41
38	BE	77	C	C5'-C4'	5.03	1.57	1.51
38	BE	119	U	N1-C2	-5.03	1.34	1.38
40	BG	46	G	O3'-P	-5.03	1.55	1.61
85	AA	62	A	O4'-C1'	-5.03	1.35	1.41
85	AA	343	U	P-O5'	-5.03	1.54	1.59
85	AA	380	C	C2-N3	-5.03	1.31	1.35
85	AA	686	U	C4'-C3'	-5.03	1.47	1.52
85	AA	1683	U	C2'-C1'	-5.03	1.47	1.53
85	AA	1828	C	C2-N3	-5.03	1.31	1.35
7	A6	116	GLY	CA-C	-5.03	1.43	1.51
34	BA	468	A	C3'-O3'	5.03	1.49	1.42
34	BA	615	A	C5-C4	-5.03	1.35	1.38
34	BA	684	G	N7-C5	-5.03	1.36	1.39
35	BB	510	A	N9-C8	-5.03	1.33	1.37
38	BE	193	A	C3'-C2'	-5.03	1.47	1.52
39	BF	8	C	C4'-C3'	-5.03	1.47	1.52
40	BG	37	G	C5-C4	-5.03	1.34	1.38
85	AA	646	C	O3'-P	-5.03	1.55	1.61
34	BA	160	G	C1'-N9	-5.03	1.39	1.46
34	BA	697	A	C4'-C3'	-5.03	1.47	1.52
34	BA	785	G	C2-N3	-5.03	1.28	1.32
34	BA	1431	G	N1-C2	-5.03	1.33	1.37
34	BA	1604	A	N3-C4	-5.03	1.31	1.34
35	BB	1461	C	C1'-N1	-5.03	1.39	1.46
38	BE	16	C	O4'-C1'	5.03	1.48	1.41
41	BH	13	C	N1-C6	-5.03	1.34	1.37
85	AA	357	C	C4-C5	-5.03	1.39	1.43
85	AA	548	G	C1'-N9	-5.03	1.39	1.46
85	AA	1164	A	C5-C4	-5.03	1.35	1.38
85	AA	1253	G	N9-C4	-5.03	1.33	1.38
85	AA	1522	U	C3'-C2'	-5.03	1.47	1.52
85	AA	1796	C	O3'-P	-5.03	1.55	1.61
85	AA	1991	C	O4'-C1'	-5.03	1.35	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
6	A5	112	TRP	CB-CG	-5.03	1.41	1.50
34	BA	1091	U	C4'-C3'	-5.03	1.47	1.52
35	BB	1254	G	C2'-C1'	-5.03	1.47	1.53
35	BB	1384	A	C1'-N9	-5.03	1.39	1.46
38	BE	133	C	C2-N3	-5.03	1.31	1.35
38	BE	189	A	C6-N6	-5.03	1.29	1.33
41	BH	20	A	O4'-C1'	-5.03	1.35	1.41
85	AA	1182	A	N7-C5	-5.03	1.36	1.39
34	BA	148	G	C6-N1	-5.03	1.36	1.39
34	BA	356	C	C1'-N1	-5.03	1.39	1.46
34	BA	410	G	N9-C8	-5.03	1.34	1.37
34	BA	697	A	C4'-O4'	-5.03	1.39	1.45
34	BA	698	U	O3'-P	-5.03	1.55	1.61
34	BA	792	A	C5-C4	-5.03	1.35	1.38
34	BA	937	G	P-O5'	-5.03	1.54	1.59
34	BA	1266	A	O3'-P	-5.03	1.55	1.61
34	BA	1478	G	C5'-C4'	5.03	1.57	1.51
34	BA	1632	G	N3-C4	-5.03	1.31	1.35
35	BB	16	G	P-O5'	-5.03	1.54	1.59
35	BB	444	U	C3'-C2'	-5.03	1.47	1.52
35	BB	584	A	C6-N1	-5.03	1.32	1.35
35	BB	744	U	C4'-C3'	-5.03	1.47	1.52
35	BB	976	U	C2-N3	-5.03	1.34	1.37
35	BB	1079	G	C4'-C3'	-5.03	1.47	1.52
77	Br	277	LEU	C-N	-5.03	1.24	1.34
85	AA	266	U	C5'-C4'	5.03	1.57	1.51
85	AA	273	C	N1-C6	-5.03	1.34	1.37
85	AA	364	C	C2-N3	-5.03	1.31	1.35
85	AA	420	C	N1-C2	-5.03	1.35	1.40
85	AA	854	A	O4'-C1'	5.03	1.48	1.41
85	AA	894	A	O3'-P	-5.03	1.55	1.61
85	AA	911	A	N7-C5	-5.03	1.36	1.39
85	AA	928	U	C1'-N1	-5.03	1.39	1.46
85	AA	1148	G	N3-C4	-5.03	1.31	1.35
85	AA	1483	A	N1-C2	-5.03	1.29	1.34
85	AA	1493	A	C3'-C2'	5.03	1.58	1.52
85	AA	2063	C	O3'-P	-5.03	1.55	1.61
85	AA	2097	U	C2-N3	-5.03	1.34	1.37
85	AA	2176	U	C4'-C3'	-5.03	1.47	1.52
34	BA	113	G	C8-N7	-5.03	1.27	1.30
34	BA	278	U	N3-C4	-5.03	1.33	1.38
34	BA	621	G	C2-N2	-5.03	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	714	G	C6-N1	-5.03	1.36	1.39
34	BA	1266	A	C5'-C4'	5.03	1.57	1.51
34	BA	1610	A	C1'-N9	-5.03	1.39	1.46
34	BA	1839	G	O4'-C1'	-5.03	1.35	1.41
35	BB	367	C	C2'-C1'	-5.03	1.47	1.53
35	BB	536	U	O3'-P	-5.03	1.55	1.61
35	BB	1056	A	C5-C4	-5.03	1.35	1.38
35	BB	1070	G	C6-N1	-5.03	1.36	1.39
35	BB	1091	C	C4'-C3'	-5.03	1.47	1.52
35	BB	1363	A	C5-C4	-5.03	1.35	1.38
35	BB	1448	U	C5'-C4'	-5.03	1.45	1.51
36	BC	95	A	C3'-C2'	-5.03	1.47	1.52
36	BC	165	U	N1-C2	-5.03	1.34	1.38
38	BE	12	A	C3'-C2'	-5.03	1.47	1.52
40	BG	174	G	N3-C4	-5.03	1.31	1.35
85	AA	177	A	C4'-C3'	-5.03	1.47	1.52
85	AA	248	U	C1'-N1	-5.03	1.39	1.46
85	AA	327	G	C2'-C1'	-5.03	1.47	1.53
85	AA	698	G	C6-N1	-5.03	1.36	1.39
85	AA	1169	A	O3'-P	-5.03	1.55	1.61
85	AA	1624	U	O3'-P	-5.03	1.55	1.61
85	AA	1859	C	P-O5'	-5.03	1.54	1.59
85	AA	1936	C	C2'-C1'	-5.03	1.47	1.53
7	A6	157	PHE	CB-CG	-5.02	1.42	1.51
34	BA	322	U	C4-O4	-5.02	1.19	1.23
34	BA	583	G	C6-N1	-5.02	1.36	1.39
34	BA	790	G	C5-C4	-5.02	1.34	1.38
34	BA	1176	C	C5'-C4'	-5.02	1.45	1.51
34	BA	1382	G	O3'-P	-5.02	1.55	1.61
34	BA	1523	U	P-O5'	-5.02	1.54	1.59
34	BA	1813	C	C4-N4	-5.02	1.29	1.33
34	BA	1832	A	C3'-C2'	-5.02	1.47	1.52
35	BB	503	G	C4'-C3'	-5.02	1.47	1.52
35	BB	586	U	C3'-C2'	-5.02	1.47	1.52
35	BB	637	G	C6-N1	-5.02	1.36	1.39
36	BC	24	G	O4'-C1'	-5.02	1.35	1.41
37	BD	27	A	C1'-N9	-5.02	1.39	1.46
85	AA	496	C	C1'-N1	-5.02	1.39	1.46
34	BA	861	C	C4-N4	-5.02	1.29	1.33
34	BA	1315	C	P-O5'	-5.02	1.54	1.59
34	BA	1725	U	C4'-O4'	-5.02	1.39	1.45
35	BB	444	U	O3'-P	-5.02	1.55	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	BB	1243	A	C8-N7	-5.02	1.28	1.31
35	BB	1440	A	N7-C5	-5.02	1.36	1.39
35	BB	1508	G	C2-N2	-5.02	1.29	1.34
38	BE	119	U	C3'-C2'	-5.02	1.47	1.52
38	BE	128	G	C2-N2	-5.02	1.29	1.34
41	BH	54	U	N1-C2	5.02	1.43	1.38
85	AA	36	U	C4'-C3'	-5.02	1.47	1.52
85	AA	534	A	C5-C4	-5.02	1.35	1.38
85	AA	630	A	C6-N1	-5.02	1.32	1.35
85	AA	1690	A	C2'-C1'	-5.02	1.47	1.53
85	AA	2138	G	C5-C6	-5.02	1.37	1.42
34	BA	1133	A	C2'-C1'	-5.02	1.47	1.53
36	BC	52	A	C6-N1	-5.02	1.32	1.35
65	Bf	165	PHE	CB-CG	-5.02	1.42	1.51
85	AA	133	G	C2-N3	-5.02	1.28	1.32
85	AA	466	A	N7-C5	-5.02	1.36	1.39
85	AA	794	A	C5-C6	-5.02	1.36	1.41
85	AA	905	C	O3'-P	-5.02	1.55	1.61
85	AA	995	G	N3-C4	-5.02	1.31	1.35
15	AG	37	VAL	CA-C	-5.02	1.39	1.52
34	BA	146	G	C4'-O4'	-5.02	1.39	1.45
34	BA	249	A	N7-C5	-5.02	1.36	1.39
34	BA	417	A	N3-C4	-5.02	1.31	1.34
34	BA	800	G	C2-N3	5.02	1.36	1.32
34	BA	1085	G	P-O5'	-5.02	1.54	1.59
34	BA	1095	G	C1'-N9	-5.02	1.39	1.46
34	BA	1476	G	C6-N1	-5.02	1.36	1.39
34	BA	1709	A	C5-C6	-5.02	1.36	1.41
35	BB	82	G	O3'-P	-5.02	1.55	1.61
35	BB	435	A	N9-C8	-5.02	1.33	1.37
35	BB	1067	G	C2-N2	-5.02	1.29	1.34
35	BB	1449	G	N3-C4	-5.02	1.31	1.35
85	AA	464	A	C4'-C3'	-5.02	1.47	1.52
34	BA	401	A	N1-C2	-5.02	1.29	1.34
34	BA	861	C	C2-N3	-5.02	1.31	1.35
34	BA	959	G	C1'-N9	-5.02	1.39	1.46
34	BA	1232	C	C4-N4	-5.02	1.29	1.33
34	BA	1708	A	C4'-C3'	-5.02	1.47	1.52
35	BB	1053	G	O4'-C1'	-5.02	1.35	1.41
85	AA	499	G	N3-C4	-5.02	1.31	1.35
85	AA	541	A	N7-C5	-5.02	1.36	1.39
85	AA	872	U	P-O5'	-5.02	1.54	1.59

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
85	AA	974	U	C4'-C3'	5.02	1.58	1.53
85	AA	1374	A	O3'-P	-5.02	1.55	1.61
34	BA	1212	A	N7-C5	-5.02	1.36	1.39
38	BE	83	U	C4'-C3'	-5.02	1.47	1.52
85	AA	471	U	C1'-N1	-5.02	1.39	1.46
85	AA	627	A	C5-C4	-5.02	1.35	1.38
85	AA	1104	G	C4'-O4'	-5.02	1.39	1.45
85	AA	1150	G	N7-C5	-5.02	1.36	1.39
34	BA	504	A	C4'-O4'	-5.01	1.39	1.45
34	BA	1008	A	N3-C4	-5.01	1.31	1.34
34	BA	1096	C	C1'-N1	-5.01	1.39	1.46
34	BA	1288	U	C4'-O4'	-5.01	1.39	1.45
34	BA	1525	G	N3-C4	-5.01	1.31	1.35
34	BA	1833	G	C2'-C1'	-5.01	1.47	1.53
35	BB	276	U	P-O5'	-5.01	1.54	1.59
35	BB	547	A	C3'-C2'	-5.01	1.47	1.52
35	BB	705	C	C3'-C2'	-5.01	1.47	1.52
35	BB	1211	C	C3'-C2'	-5.01	1.47	1.52
35	BB	1336	G	C6-N1	-5.01	1.36	1.39
40	BG	65	C	C2'-C1'	-5.01	1.47	1.53
85	AA	39	A	N7-C5	-5.01	1.36	1.39
85	AA	679	A	C8-N7	-5.01	1.28	1.31
85	AA	906	U	P-O5'	-5.01	1.54	1.59
85	AA	1158	U	P-O5'	-5.01	1.54	1.59
85	AA	1519	A	C3'-C2'	-5.01	1.47	1.52
85	AA	1863	A	C3'-C2'	-5.01	1.47	1.52
85	AA	2204	A	N9-C4	-5.01	1.34	1.37
34	BA	773	A	C4'-C3'	-5.01	1.47	1.52
34	BA	1225	A	C5-C4	-5.01	1.35	1.38
35	BB	78	C	C2'-C1'	-5.01	1.47	1.53
35	BB	412	A	C5-C4	-5.01	1.35	1.38
35	BB	1063	C	C2-N3	-5.01	1.31	1.35
35	BB	1487	G	N7-C5	-5.01	1.36	1.39
37	BD	95	G	O3'-P	-5.01	1.55	1.61
85	AA	605	A	C5-C4	-5.01	1.35	1.38
34	BA	377	G	C3'-C2'	-5.01	1.47	1.52
34	BA	393	G	P-O5'	-5.01	1.54	1.59
34	BA	992	A	N3-C4	-5.01	1.31	1.34
35	BB	103	C	C3'-C2'	-5.01	1.47	1.52
35	BB	383	U	P-O5'	-5.01	1.54	1.59
37	BD	56	G	C1'-N9	-5.01	1.39	1.46
37	BD	60	C	C4-N4	-5.01	1.29	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	171	U	C4'-C3'	-5.01	1.47	1.52
56	BW	128	TRP	C-N	-5.01	1.24	1.34
85	AA	54	C	C4-N4	-5.01	1.29	1.33
85	AA	436	G	C2-N2	-5.01	1.29	1.34
85	AA	851	G	P-O5'	-5.01	1.54	1.59
85	AA	1579	A	O3'-P	-5.01	1.55	1.61
85	AA	1698	A	N9-C8	-5.01	1.33	1.37
85	AA	1856	G	N1-C2	-5.01	1.33	1.37
85	AA	1923	A	O4'-C1'	-5.01	1.35	1.41
34	BA	260	A	O3'-P	-5.01	1.55	1.61
34	BA	504	A	O4'-C1'	-5.01	1.35	1.41
34	BA	523	A	N9-C4	5.01	1.40	1.37
34	BA	1104	C	C3'-C2'	-5.01	1.47	1.52
34	BA	1331	G	P-O5'	-5.01	1.54	1.59
34	BA	1553	G	C3'-O3'	5.01	1.49	1.42
34	BA	1679	C	N1-C6	-5.01	1.34	1.37
34	BA	1799	G	N1-C2	-5.01	1.33	1.37
35	BB	89	C	C3'-C2'	-5.01	1.47	1.52
35	BB	306	U	P-O5'	-5.01	1.54	1.59
35	BB	449	C	C3'-C2'	-5.01	1.47	1.52
35	BB	578	G	N3-C4	-5.01	1.31	1.35
35	BB	698	C	C4'-C3'	-5.01	1.47	1.52
35	BB	1170	U	C1'-N1	-5.01	1.39	1.46
35	BB	1309	A	N3-C4	-5.01	1.31	1.34
35	BB	1508	G	N1-C2	-5.01	1.33	1.37
37	BD	86	A	N9-C8	-5.01	1.33	1.37
37	BD	107	G	N1-C2	-5.01	1.33	1.37
38	BE	18	U	O3'-P	-5.01	1.55	1.61
38	BE	146	U	O4'-C1'	-5.01	1.35	1.41
39	BF	25	G	C2'-C1'	-5.01	1.47	1.53
56	BW	134	HIS	CA-C	-5.01	1.40	1.52
85	AA	61	C	C2-N3	-5.01	1.31	1.35
85	AA	774	C	O3'-P	-5.01	1.55	1.61
85	AA	1701	G	O4'-C1'	-5.01	1.35	1.41
85	AA	1977	G	O3'-P	-5.01	1.55	1.61
85	AA	1979	A	C2'-C1'	-5.01	1.47	1.53
85	AA	2247	C	O3'-P	-5.01	1.55	1.61
34	BA	28	C	N3-C4	-5.01	1.30	1.33
34	BA	1098	G	N7-C5	-5.01	1.36	1.39
34	BA	1429	A	C2'-C1'	-5.01	1.47	1.53
34	BA	1812	C	C2-N3	-5.01	1.31	1.35
35	BB	293	G	O3'-P	-5.01	1.55	1.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	BE	185	G	C5'-C4'	-5.01	1.45	1.51
40	BG	32	U	C2'-C1'	-5.01	1.47	1.53
34	BA	15	G	C6-N1	-5.01	1.36	1.39
34	BA	323	C	C4'-O4'	-5.01	1.39	1.45
34	BA	366	G	C5'-C4'	-5.01	1.45	1.51
34	BA	748	C	N1-C6	-5.01	1.34	1.37
34	BA	1826	C	C2'-C1'	-5.01	1.47	1.53
35	BB	572	G	C5-C4	-5.01	1.34	1.38
35	BB	1480	G	C8-N7	-5.01	1.27	1.30
35	BB	1521	G	C4'-C3'	-5.01	1.47	1.52
40	BG	31	G	C2-N3	-5.01	1.28	1.32
41	BH	47	G	N9-C8	-5.01	1.34	1.37
41	BH	105	U	N3-C4	-5.01	1.33	1.38
41	BH	107	A	N3-C4	-5.01	1.31	1.34
41	BH	113	G	C2-N2	-5.01	1.29	1.34
74	Bo	79	VAL	N-CA	-5.01	1.36	1.46
85	AA	732	G	N7-C5	-5.01	1.36	1.39
85	AA	741	G	C3'-C2'	-5.01	1.47	1.52
85	AA	939	A	C6-N6	-5.01	1.29	1.33
85	AA	1539	A	N7-C5	-5.01	1.36	1.39
34	BA	457	A	C1'-N9	-5.00	1.39	1.46
34	BA	1146	U	C5'-C4'	-5.00	1.45	1.51
34	BA	1158	A	C4'-C3'	-5.00	1.47	1.52
34	BA	1816	G	N7-C5	-5.00	1.36	1.39
35	BB	1260	A	N7-C5	-5.00	1.36	1.39
37	BD	41	G	C6-N1	-5.00	1.36	1.39
37	BD	48	G	C3'-C2'	-5.00	1.47	1.52
85	AA	293	A	N7-C5	-5.00	1.36	1.39
85	AA	406	U	C4'-C3'	-5.00	1.47	1.52
85	AA	2025	A	C1'-N9	-5.00	1.39	1.46
34	BA	35	U	O3'-P	-5.00	1.55	1.61
34	BA	263	G	N1-C2	-5.00	1.33	1.37
34	BA	454	G	C4'-C3'	-5.00	1.47	1.52
34	BA	568	G	C1'-N9	-5.00	1.39	1.46
34	BA	744	G	C2-N2	-5.00	1.29	1.34
34	BA	893	U	C1'-N1	-5.00	1.39	1.46
34	BA	1008	A	C2'-C1'	-5.00	1.47	1.53
34	BA	1161	G	P-O5'	-5.00	1.54	1.59
34	BA	1225	A	C4'-C3'	-5.00	1.47	1.52
34	BA	1244	G	N9-C4	-5.00	1.33	1.38
34	BA	1293	A	O4'-C1'	-5.00	1.35	1.41
34	BA	1379	G	C2-N2	-5.00	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
34	BA	1465	C	C4'-C3'	-5.00	1.47	1.52
34	BA	1469	G	O3'-P	-5.00	1.55	1.61
34	BA	1806	A	C3'-C2'	-5.00	1.47	1.52
35	BB	825	U	C2-N3	-5.00	1.34	1.37
35	BB	982	A	P-O5'	-5.00	1.54	1.59
35	BB	1377	A	C3'-C2'	-5.00	1.47	1.52
36	BC	88	A	C3'-C2'	-5.00	1.47	1.52
36	BC	120	G	C2-N2	-5.00	1.29	1.34
38	BE	96	G	C5'-C4'	-5.00	1.45	1.51
40	BG	126	G	C4'-C3'	-5.00	1.47	1.52
58	BY	8	PHE	CB-CG	-5.00	1.42	1.51
85	AA	4	C	N3-C4	5.00	1.37	1.33
85	AA	152	A	O4'-C1'	-5.00	1.35	1.41
85	AA	402	G	C2-N2	-5.00	1.29	1.34
85	AA	443	A	C5-C4	-5.00	1.35	1.38
85	AA	561	C	C2-N3	-5.00	1.31	1.35
85	AA	698	G	O3'-P	-5.00	1.55	1.61
85	AA	2096	G	N7-C5	-5.00	1.36	1.39
85	AA	2101	C	O3'-P	-5.00	1.55	1.61
34	BA	100	A	C8-N7	-5.00	1.28	1.31
34	BA	694	G	P-O5'	-5.00	1.54	1.59
34	BA	1050	A	O3'-P	-5.00	1.55	1.61
34	BA	1438	C	C4-N4	-5.00	1.29	1.33
34	BA	1622	U	C4'-C3'	-5.00	1.47	1.52
35	BB	8	U	C5'-C4'	-5.00	1.45	1.51
35	BB	757	C	O3'-P	-5.00	1.55	1.61
35	BB	1176	G	C8-N7	-5.00	1.27	1.30
35	BB	1434	G	N7-C5	-5.00	1.36	1.39
36	BC	53	A	C8-N7	-5.00	1.28	1.31
36	BC	105	C	C4'-O4'	-5.00	1.39	1.45
40	BG	117	C	C3'-C2'	-5.00	1.47	1.52
85	AA	23	G	C2-N2	-5.00	1.29	1.34
85	AA	539	A	C5-C6	-5.00	1.36	1.41
85	AA	542	G	C3'-C2'	-5.00	1.47	1.52
85	AA	1442	U	P-O5'	-5.00	1.54	1.59
85	AA	1894	G	N9-C4	-5.00	1.33	1.38
85	AA	2108	C	O3'-P	-5.00	1.55	1.61

All (28260) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	469	G	N9-C4-C5	-49.23	85.71	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	32	G	P-O3'-C3'	41.15	169.07	119.70
35	BB	1212	C	C6-N1-C2	-40.14	104.24	120.30
85	AA	769	C	C6-N1-C2	-39.18	104.63	120.30
34	BA	692	U	P-O3'-C3'	38.13	165.46	119.70
38	BE	203	C	C6-N1-C2	-37.90	105.14	120.30
85	AA	469	G	C8-N9-C4	37.72	121.49	106.40
38	BE	175	U	P-O3'-C3'	35.72	162.56	119.70
85	AA	469	G	N3-C4-N9	35.43	147.26	126.00
34	BA	605	G	C8-N9-C4	-34.02	92.79	106.40
34	BA	647	U	P-O3'-C3'	-33.75	79.20	119.70
34	BA	743	A	C4-C5-N7	33.48	127.44	110.70
34	BA	1523	U	P-O3'-C3'	33.29	159.65	119.70
40	BG	9	G	P-O3'-C3'	33.23	159.58	119.70
85	AA	24	U	P-O3'-C3'	32.89	159.17	119.70
39	BF	65	U	C2-N3-C4	-32.08	107.75	127.00
34	BA	633	G	P-O3'-C3'	31.36	157.33	119.70
85	AA	1371	C	P-O3'-C3'	31.14	157.07	119.70
34	BA	743	A	N9-C4-C5	-30.20	93.72	105.80
35	BB	1024	G	P-O3'-C3'	29.96	155.65	119.70
34	BA	855	C	C6-N1-C2	-29.92	108.33	120.30
85	AA	150	U	C2-N3-C4	-29.81	109.12	127.00
35	BB	61	A	P-O3'-C3'	29.68	155.32	119.70
85	AA	1973	G	P-O3'-C3'	29.62	155.24	119.70
34	BA	896	U	C2-N3-C4	-29.21	109.47	127.00
35	BB	835	C	C6-N1-C2	-29.17	108.63	120.30
40	BG	22	G	P-O3'-C3'	28.94	154.43	119.70
34	BA	578	C	P-O3'-C3'	28.41	153.80	119.70
85	AA	1294	U	C2-N3-C4	-28.17	110.10	127.00
85	AA	330	C	C6-N1-C2	-28.03	109.09	120.30
35	BB	972	C	P-O3'-C3'	27.92	153.21	119.70
34	BA	557	U	C4'-C3'-C2'	-27.90	74.70	102.60
34	BA	1561	C	C6-N1-C2	-27.68	109.23	120.30
34	BA	517	A	P-O5'-C5'	27.48	164.87	120.90
38	BE	18	U	P-O3'-C3'	27.02	152.12	119.70
38	BE	183	C	C6-N1-C2	-26.98	109.51	120.30
85	AA	1816	C	C6-N1-C2	-26.98	109.51	120.30
34	BA	768	G	P-O3'-C3'	26.85	151.92	119.70
35	BB	1494	G	P-O3'-C3'	26.59	151.61	119.70
35	BB	838	G	P-O3'-C3'	26.53	151.54	119.70
34	BA	689	C	P-O3'-C3'	26.51	151.51	119.70
37	BD	83	A	P-O3'-C3'	26.34	151.31	119.70
34	BA	214	A	C2-N3-C4	26.26	123.73	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1280	U	C2-N3-C4	-26.14	111.31	127.00
34	BA	513	U	C2-N3-C4	-26.05	111.37	127.00
85	AA	742	U	C2-N3-C4	-26.03	111.38	127.00
34	BA	1494	G	C5-C6-O6	-25.92	113.05	128.60
37	BD	118	C	C6-N1-C2	-25.84	109.97	120.30
34	BA	205	G	P-O3'-C3'	25.73	150.58	119.70
34	BA	112	C	C6-N1-C2	-25.66	110.03	120.30
34	BA	1787	U	P-O3'-C3'	25.55	150.37	119.70
34	BA	606	G	P-O3'-C3'	25.46	150.26	119.70
40	BG	24	A	C4-C5-C6	-25.42	104.29	117.00
85	AA	654	A	N1-C6-N6	-25.16	103.51	118.60
34	BA	598	G	P-O3'-C3'	24.78	149.44	119.70
34	BA	743	A	C4-C5-C6	-24.71	104.64	117.00
34	BA	605	G	P-O5'-C5'	24.68	160.38	120.90
35	BB	2	C	P-O5'-C5'	24.66	160.36	120.90
34	BA	777	C	P-O5'-C5'	24.63	160.31	120.90
34	BA	766	A	P-O5'-C5'	24.43	159.99	120.90
34	BA	1831	A	P-O3'-C3'	24.43	149.01	119.70
34	BA	1494	G	P-O3'-C3'	24.40	148.99	119.70
34	BA	771	A	P-O3'-C3'	24.37	148.94	119.70
34	BA	280	A	P-O5'-C5'	24.27	159.74	120.90
85	AA	164	G	P-O5'-C5'	24.16	159.56	120.90
34	BA	678	C	C6-N1-C2	-24.07	110.67	120.30
85	AA	862	U	P-O3'-C3'	24.03	148.53	119.70
38	BE	23	G	P-O5'-C5'	23.94	159.21	120.90
34	BA	743	A	C8-N9-C4	-23.91	96.24	105.80
34	BA	763	U	C2-N3-C4	-23.88	112.67	127.00
40	BG	24	A	P-O3'-C3'	23.77	148.23	119.70
34	BA	605	G	C4-N9-C1'	23.66	157.26	126.50
40	BG	23	C	P-O5'-C5'	23.55	158.58	120.90
85	AA	1535	C	P-O3'-C3'	23.52	147.92	119.70
85	AA	529	G	P-O3'-C3'	23.44	147.83	119.70
85	AA	98	U	C2-N3-C4	-23.29	113.03	127.00
34	BA	147	U	C2-N3-C4	-23.27	113.04	127.00
34	BA	870	C	O5'-P-OP2	-23.05	83.04	110.70
35	BB	797	C	P-O3'-C3'	22.90	147.18	119.70
85	AA	300	C	P-O3'-C3'	22.89	147.17	119.70
35	BB	1438	U	C2-N3-C4	-22.89	113.27	127.00
85	AA	2244	G	P-O3'-C3'	22.74	146.99	119.70
85	AA	80	G	P-O3'-C3'	22.61	146.83	119.70
85	AA	1921	G	C5'-C4'-C3'	22.57	152.11	116.00
34	BA	570	G	P-O3'-C3'	22.57	146.78	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	890	U	P-O3'-C3'	22.39	146.56	119.70
85	AA	867	G	P-O3'-C3'	22.37	146.54	119.70
34	BA	384	U	C2-N3-C4	-22.29	113.62	127.00
41	BH	71	C	P-O3'-C3'	22.24	146.39	119.70
36	BC	123	G	P-O3'-C3'	22.23	146.38	119.70
35	BB	970	C	P-O3'-C3'	22.23	146.37	119.70
39	BF	51	C	P-O3'-C3'	22.06	146.17	119.70
35	BB	1337	C	C6-N1-C2	-22.03	111.49	120.30
85	AA	1301	C	P-O3'-C3'	21.85	145.92	119.70
41	BH	26	C	P-O3'-C3'	21.84	145.91	119.70
85	AA	735	G	P-O3'-C3'	21.76	145.81	119.70
34	BA	1722	U	C2-N3-C4	-21.73	113.96	127.00
38	BE	134	A	P-O3'-C3'	21.73	145.78	119.70
85	AA	45	U	P-O3'-C3'	21.73	145.77	119.70
35	BB	1202	G	O4'-C1'-N9	21.71	125.56	108.20
85	AA	381	A	N1-C6-N6	-21.58	105.65	118.60
35	BB	614	U	C2-N3-C4	-21.57	114.06	127.00
34	BA	91	C	O4'-C1'-N1	21.56	125.45	108.20
85	AA	1210	U	C6-N1-C2	-21.52	108.09	121.00
35	BB	1167	C	P-O3'-C3'	21.50	145.50	119.70
35	BB	3	C	O4'-C1'-N1	21.43	125.34	108.20
85	AA	744	C	C6-N1-C2	-21.42	111.73	120.30
34	BA	80	U	C2-N3-C4	-21.27	114.24	127.00
35	BB	795	A	P-O3'-C3'	21.27	145.22	119.70
85	AA	469	G	N7-C8-N9	-21.22	102.49	113.10
34	BA	871	G	C5-C6-O6	20.97	141.18	128.60
34	BA	532	C	C6-N1-C2	-20.92	111.93	120.30
34	BA	1307	U	P-O3'-C3'	20.89	144.77	119.70
34	BA	871	G	N1-C6-O6	-20.89	107.37	119.90
34	BA	605	G	N7-C8-N9	20.83	123.52	113.10
85	AA	4	C	C6-N1-C2	-20.70	112.02	120.30
34	BA	1488	C	C6-N1-C2	-20.69	112.02	120.30
85	AA	787	U	P-O5'-C5'	20.62	153.89	120.90
85	AA	902	A	P-O5'-C5'	20.58	153.82	120.90
41	BH	74	G	N1-C6-O6	20.54	132.23	119.90
34	BA	1422	A	P-O3'-C3'	20.54	144.35	119.70
35	BB	5	A	P-O5'-C5'	20.48	153.67	120.90
85	AA	710	A	P-O3'-C3'	20.47	144.26	119.70
85	AA	1458	G	N1-C6-O6	-20.42	107.65	119.90
85	AA	1018	G	P-O5'-C5'	20.42	153.57	120.90
34	BA	954	U	C2-N3-C4	-20.36	114.78	127.00
41	BH	74	G	C5-C6-O6	-20.33	116.40	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	936	C	C6-N1-C2	-20.30	112.18	120.30
85	AA	1535	C	C6-N1-C2	-20.27	112.19	120.30
34	BA	487	A	N1-C6-N6	-20.27	106.44	118.60
34	BA	827	A	P-O3'-C3'	20.27	144.02	119.70
34	BA	590	U	C2-N3-C4	-20.20	114.88	127.00
34	BA	896	U	C6-N1-C2	-20.20	108.88	121.00
35	BB	870	C	C6-N1-C2	-20.16	112.24	120.30
34	BA	315	U	P-O5'-C5'	20.15	153.14	120.90
85	AA	47	A	P-O3'-C3'	20.14	143.87	119.70
41	BH	128	G	C5-C6-O6	-20.14	116.51	128.60
85	AA	1090	A	P-O3'-C3'	20.11	143.84	119.70
85	AA	56	U	P-O3'-C3'	20.07	143.78	119.70
85	AA	1159	C	C6-N1-C2	-20.06	112.28	120.30
36	BC	148	C	P-O5'-C5'	20.02	152.93	120.90
35	BB	652	G	P-O3'-C3'	-19.93	95.79	119.70
36	BC	81	U	P-O5'-C5'	19.89	152.72	120.90
85	AA	2216	A	P-O3'-C3'	19.89	143.56	119.70
85	AA	1978	G	C5-C6-O6	-19.87	116.68	128.60
34	BA	961	C	C6-N1-C2	-19.78	112.39	120.30
34	BA	206	C	P-O3'-C3'	19.77	143.42	119.70
34	BA	769	U	P-O3'-C3'	19.73	143.38	119.70
38	BE	20	C	O4'-C1'-N1	19.48	123.78	108.20
34	BA	296	G	C5'-C4'-C3'	19.41	147.05	116.00
34	BA	162	G	N1-C6-O6	19.39	131.54	119.90
34	BA	712	C	P-O3'-C3'	19.38	142.96	119.70
34	BA	1202	G	N1-C6-O6	-19.27	108.34	119.90
40	BG	168	A	N1-C6-N6	-19.24	107.05	118.60
85	AA	1183	C	P-O3'-C3'	19.23	142.78	119.70
35	BB	1379	U	C2-N3-C4	-19.22	115.47	127.00
85	AA	523	U	P-O5'-C5'	19.21	151.64	120.90
85	AA	738	C	C6-N1-C2	-19.19	112.62	120.30
35	BB	1512	C	C6-N1-C2	-19.14	112.64	120.30
85	AA	2031	C	C6-N1-C2	-19.14	112.64	120.30
35	BB	894	A	P-O3'-C3'	19.13	142.66	119.70
35	BB	766	G	P-O5'-C5'	18.99	151.28	120.90
34	BA	625	U	P-O3'-C3'	18.93	142.42	119.70
85	AA	1396	C	P-O3'-C3'	18.91	142.39	119.70
34	BA	646	C	P-O3'-C3'	18.91	142.39	119.70
34	BA	557	U	C4'-C3'-O3'	18.86	150.71	113.00
85	AA	1208	C	P-O3'-C3'	18.82	142.28	119.70
85	AA	2022	A	N1-C6-N6	-18.81	107.31	118.60
34	BA	1494	G	N1-C6-O6	18.80	131.18	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	778	U	P-O3'-C3'	18.79	142.25	119.70
34	BA	1215	U	C2-N3-C4	-18.76	115.74	127.00
34	BA	711	C	P-O3'-C3'	18.75	142.21	119.70
34	BA	862	C	P-O3'-C3'	18.69	142.13	119.70
85	AA	1876	U	C2-N3-C4	-18.68	115.79	127.00
85	AA	674	U	C2-N3-C4	-18.67	115.80	127.00
85	AA	1067	G	P-O3'-C3'	18.66	142.09	119.70
34	BA	162	G	C5-C6-O6	-18.57	117.46	128.60
34	BA	896	U	C2-N1-C1'	18.54	139.95	117.70
40	BG	24	A	N1-C6-N6	-18.53	107.48	118.60
85	AA	1091	C	C6-N1-C2	-18.52	112.89	120.30
85	AA	267	U	C2-N1-C1'	-18.50	95.50	117.70
38	BE	183	C	C2-N1-C1'	18.50	139.15	118.80
85	AA	368	C	C6-N1-C2	-18.45	112.92	120.30
85	AA	1492	U	C6-N1-C2	-18.41	109.95	121.00
34	BA	870	C	O5'-P-OP1	-18.38	88.64	110.70
37	BD	95	G	C8-N9-C4	-18.38	99.05	106.40
34	BA	680	C	C6-N1-C2	-18.36	112.96	120.30
34	BA	591	G	P-O5'-C5'	18.36	150.27	120.90
35	BB	1439	U	C2-N3-C4	-18.29	116.03	127.00
40	BG	105	A	N1-C6-N6	-18.28	107.63	118.60
34	BA	214	A	N9-C4-C5	-18.20	98.52	105.80
34	BA	251	U	C2-N3-C4	-18.19	116.09	127.00
35	BB	5	A	C5'-C4'-C3'	18.18	145.09	116.00
40	BG	21	C	O4'-C1'-N1	18.17	122.74	108.20
35	BB	561	C	C1'-O4'-C4'	-18.15	95.38	109.90
36	BC	31	A	P-O3'-C3'	18.15	141.48	119.70
56	BW	87	TRP	CB-CG-CD2	-18.15	103.01	126.60
34	BA	214	A	N1-C2-N3	-18.04	120.28	129.30
85	AA	901	C	P-O3'-C3'	18.04	141.35	119.70
85	AA	327	G	O4'-C1'-N9	18.03	122.62	108.20
85	AA	1277	C	P-O3'-C3'	18.00	141.30	119.70
34	BA	1225	A	N1-C6-N6	-17.98	107.81	118.60
35	BB	59	U	P-O3'-C3'	17.97	141.26	119.70
85	AA	250	C	C6-N1-C2	-17.96	113.12	120.30
34	BA	629	G	C5-C6-O6	17.92	139.35	128.60
34	BA	588	C	P-O3'-C3'	17.91	141.20	119.70
34	BA	607	C	C6-N1-C2	-17.88	113.15	120.30
34	BA	1562	G	P-O3'-C3'	17.85	141.12	119.70
85	AA	207	G	O4'-C1'-N9	17.83	122.47	108.20
34	BA	797	A	P-O3'-C3'	17.83	141.10	119.70
85	AA	1825	A	P-O3'-C3'	17.78	141.04	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	882	G	P-O3'-C3'	17.78	141.03	119.70
34	BA	1222	C	C6-N1-C2	-17.77	113.19	120.30
34	BA	1204	U	P-O3'-C3'	17.75	140.99	119.70
34	BA	1700	C	C6-N1-C2	-17.74	113.20	120.30
85	AA	1514	A	P-O3'-C3'	17.73	140.98	119.70
34	BA	557	U	O4'-C1'-N1	17.73	122.38	108.20
34	BA	1005	C	C6-N1-C2	-17.73	113.21	120.30
35	BB	1063	C	C6-N1-C2	-17.71	113.22	120.30
34	BA	589	A	P-O3'-C3'	17.70	140.94	119.70
34	BA	1501	U	C2-N3-C4	-17.69	116.39	127.00
34	BA	1577	U	C2-N3-C4	-17.68	116.39	127.00
34	BA	1775	U	P-O3'-C3'	17.68	140.91	119.70
34	BA	528	C	C6-N1-C2	-17.66	113.24	120.30
34	BA	300	C	O4'-C1'-N1	17.63	122.30	108.20
34	BA	605	G	N3-C4-C5	-17.62	119.79	128.60
34	BA	7	U	C2-N3-C4	-17.60	116.44	127.00
85	AA	1454	U	P-O5'-C5'	17.57	149.02	120.90
40	BG	106	G	O4'-C1'-N9	17.54	122.23	108.20
35	BB	363	A	O4'-C1'-N9	17.54	122.23	108.20
34	BA	629	G	N1-C6-O6	-17.53	109.38	119.90
41	BH	128	G	P-O3'-C3'	17.51	140.72	119.70
34	BA	598	G	P-O5'-C5'	17.51	148.91	120.90
34	BA	1176	C	C6-N1-C2	-17.50	113.30	120.30
85	AA	1459	C	C6-N1-C2	-17.50	113.30	120.30
36	BC	81	U	P-O3'-C3'	17.47	140.66	119.70
34	BA	559	C	OP1-P-OP2	-17.44	93.44	119.60
38	BE	183	C	C6-N1-C1'	-17.44	99.88	120.80
85	AA	837	C	P-O3'-C3'	17.38	140.56	119.70
38	BE	103	C	P-O3'-C3'	17.37	140.55	119.70
35	BB	14	C	C6-N1-C2	-17.36	113.36	120.30
34	BA	800	G	C6-N1-C2	-17.34	114.69	125.10
34	BA	1724	G	C4-N9-C1'	-17.34	103.95	126.50
41	BH	48	G	P-O3'-C3'	17.34	140.50	119.70
34	BA	501	U	O4'-C1'-N1	17.33	122.06	108.20
34	BA	745	A	N1-C6-N6	-17.31	108.21	118.60
34	BA	1330	G	P-O3'-C3'	17.31	140.47	119.70
34	BA	1699	A	O4'-C1'-N9	17.30	122.04	108.20
35	BB	897	C	C6-N1-C2	-17.29	113.38	120.30
35	BB	683	U	C2-N3-C4	-17.28	116.63	127.00
41	BH	101	A	P-O5'-C5'	17.27	148.53	120.90
34	BA	281	C	P-O3'-C3'	17.27	140.42	119.70
85	AA	1789	C	P-O3'-C3'	17.26	140.41	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1604	A	P-O3'-C3'	17.23	140.38	119.70
40	BG	150	A	N1-C6-N6	17.22	128.93	118.60
38	BE	25	U	C2-N3-C4	-17.21	116.67	127.00
35	BB	25	A	N1-C6-N6	-17.18	108.29	118.60
85	AA	654	A	C5-C6-N6	17.13	137.41	123.70
35	BB	561	C	O4'-C1'-N1	17.12	121.90	108.20
34	BA	289	A	C4-C5-C6	-17.11	108.44	117.00
35	BB	1490	G	O4'-C1'-N9	17.11	121.89	108.20
34	BA	89	G	P-O3'-C3'	17.10	140.22	119.70
35	BB	1502	U	P-O3'-C3'	17.10	140.22	119.70
34	BA	1475	G	P-O3'-C3'	17.10	140.22	119.70
36	BC	31	A	O4'-C1'-N9	17.09	121.88	108.20
34	BA	122	U	C2-N3-C4	-17.09	116.75	127.00
85	AA	466	A	N3-C4-N9	-17.06	113.75	127.40
37	BD	48	G	C8-N9-C4	17.05	113.22	106.40
34	BA	1149	C	C6-N1-C2	-17.04	113.48	120.30
85	AA	556	C	C6-N1-C2	-17.01	113.49	120.30
85	AA	414	C	C6-N1-C2	-17.00	113.50	120.30
85	AA	509	C	P-O3'-C3'	16.99	140.09	119.70
36	BC	131	C	P-O5'-C5'	16.98	148.07	120.90
41	BH	72	G	P-O3'-C3'	16.98	140.08	119.70
34	BA	1656	A	C5-C6-N6	-16.98	110.12	123.70
34	BA	1282	G	C5-C6-O6	-16.95	118.43	128.60
34	BA	765	U	O4'-C1'-N1	16.95	121.76	108.20
34	BA	84	U	P-O3'-C3'	16.94	140.03	119.70
34	BA	1809	G	O4'-C1'-N9	16.94	121.75	108.20
34	BA	1506	C	P-O5'-C5'	16.93	147.99	120.90
41	BH	33	G	C5-C6-O6	-16.93	118.44	128.60
34	BA	1809	G	C1'-O4'-C4'	-16.91	96.37	109.90
35	BB	899	C	C6-N1-C2	-16.91	113.54	120.30
34	BA	1211	G	C8-N9-C4	-16.89	99.64	106.40
85	AA	115	U	C2-N3-C4	-16.88	116.87	127.00
40	BG	57	A	N1-C6-N6	16.87	128.72	118.60
34	BA	174	A	P-O3'-C3'	16.86	139.93	119.70
35	BB	784	C	P-O5'-C5'	16.84	147.84	120.90
85	AA	1691	U	P-O3'-C3'	16.83	139.90	119.70
85	AA	606	A	P-O3'-C3'	16.83	139.89	119.70
35	BB	868	C	C6-N1-C2	-16.82	113.57	120.30
85	AA	1455	C	P-O3'-C3'	-16.80	99.54	119.70
34	BA	1832	A	P-O5'-C5'	16.80	147.78	120.90
34	BA	597	C	P-O5'-C5'	16.80	147.78	120.90
38	BE	31	A	C5'-C4'-C3'	-16.79	89.14	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	641	C	P-O5'-C5'	16.78	147.75	120.90
34	BA	1636	C	P-O3'-C3'	16.77	139.83	119.70
38	BE	127	G	C4-N9-C1'	-16.74	104.73	126.50
34	BA	316	G	P-O5'-C5'	16.74	147.68	120.90
35	BB	870	C	P-O5'-C5'	16.73	147.67	120.90
85	AA	2058	C	C6-N1-C2	-16.73	113.61	120.30
85	AA	1134	G	C5-C6-O6	-16.72	118.57	128.60
38	BE	133	C	P-O3'-C3'	16.71	139.75	119.70
85	AA	603	C	P-O3'-C3'	16.71	139.76	119.70
85	AA	454	G	P-O3'-C3'	16.71	139.75	119.70
35	BB	1498	G	P-O3'-C3'	16.70	139.74	119.70
34	BA	141	G	N3-C2-N2	-16.69	108.22	119.90
85	AA	1421	U	P-O3'-C3'	16.68	139.71	119.70
39	BF	32	G	C5'-C4'-C3'	-16.68	89.32	116.00
85	AA	1271	U	P-O3'-C3'	16.67	139.70	119.70
85	AA	1107	A	P-O5'-C5'	16.66	147.55	120.90
41	BH	99	G	N1-C6-O6	-16.65	109.91	119.90
35	BB	1506	C	C6-N1-C2	-16.65	113.64	120.30
34	BA	1487	U	P-O3'-C3'	16.64	139.67	119.70
36	BC	125	A	P-O5'-C5'	16.62	147.49	120.90
34	BA	1696	G	P-O3'-C3'	16.61	139.64	119.70
35	BB	839	G	P-O5'-C5'	16.60	147.46	120.90
85	AA	1466	U	P-O3'-C3'	16.60	139.62	119.70
85	AA	136	U	P-O5'-C5'	16.59	147.44	120.90
35	BB	799	A	P-O3'-C3'	16.57	139.59	119.70
85	AA	862	U	C2-N1-C1'	-16.55	97.84	117.70
34	BA	312	U	C2-N3-C4	-16.55	117.07	127.00
36	BC	10	C	O4'-C1'-N1	16.52	121.42	108.20
34	BA	1319	A	P-O3'-C3'	16.48	139.48	119.70
34	BA	56	G	P-O3'-C3'	16.48	139.47	119.70
34	BA	1697	U	P-O3'-C3'	16.48	139.47	119.70
35	BB	1445	A	N1-C6-N6	-16.48	108.71	118.60
38	BE	190	U	P-O3'-C3'	16.44	139.43	119.70
34	BA	780	U	O4'-C1'-N1	16.40	121.32	108.20
38	BE	108	U	C5'-C4'-C3'	16.40	142.24	116.00
38	BE	117	A	N1-C2-N3	-16.39	121.10	129.30
34	BA	1443	U	P-O3'-C3'	16.38	139.35	119.70
35	BB	3	C	C6-N1-C2	-16.35	113.76	120.30
39	BF	33	C	P-O3'-C3'	16.33	139.30	119.70
34	BA	848	U	P-O3'-C3'	16.33	139.30	119.70
34	BA	1639	U	C6-N1-C2	-16.33	111.20	121.00
85	AA	860	C	C2-N1-C1'	-16.33	100.84	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	818	U	P-O3'-C3'	16.33	139.29	119.70
34	BA	91	C	C1'-O4'-C4'	-16.32	96.85	109.90
41	BH	33	G	N1-C6-O6	16.32	129.69	119.90
85	AA	1789	C	C5'-C4'-C3'	16.32	142.11	116.00
38	BE	9	C	P-O3'-C3'	16.32	139.28	119.70
86	AB	3	C	O4'-C1'-N1	16.32	121.25	108.20
85	AA	617	C	C6-N1-C2	-16.30	113.78	120.30
34	BA	813	C	C6-N1-C1'	16.30	140.36	120.80
35	BB	1102	U	O4'-C1'-N1	16.30	121.24	108.20
35	BB	1063	C	P-O3'-C3'	16.30	139.26	119.70
34	BA	685	C	P-O3'-C3'	-16.29	100.16	119.70
37	BD	70	C	C6-N1-C2	-16.27	113.79	120.30
35	BB	995	C	P-O5'-C5'	16.26	146.92	120.90
41	BH	85	C	P-O3'-C3'	16.26	139.21	119.70
85	AA	2004	U	P-O3'-C3'	16.26	139.21	119.70
34	BA	279	U	P-O3'-C3'	16.25	139.21	119.70
35	BB	981	A	P-O3'-C3'	16.24	139.19	119.70
39	BF	39	C	C6-N1-C2	-16.24	113.81	120.30
64	Be	70	ARG	NE-CZ-NH1	16.23	128.42	120.30
34	BA	743	A	N1-C6-N6	16.22	128.33	118.60
85	AA	1110	A	P-O5'-C5'	16.20	146.82	120.90
35	BB	472	C	C6-N1-C2	-16.18	113.83	120.30
35	BB	768	A	O4'-C1'-N9	16.18	121.15	108.20
85	AA	267	U	O4'-C1'-N1	16.18	121.14	108.20
38	BE	135	A	P-O3'-C3'	16.16	139.09	119.70
41	BH	102	C	O3'-P-O5'	16.15	134.68	104.00
37	BD	88	U	O4'-C1'-N1	16.14	121.11	108.20
35	BB	905	C	P-O3'-C3'	16.14	139.07	119.70
85	AA	360	C	P-O5'-C5'	16.13	146.71	120.90
34	BA	516	U	C5'-C4'-C3'	16.12	141.79	116.00
34	BA	1656	A	N1-C6-N6	16.12	128.27	118.60
85	AA	181	A	P-O3'-C3'	16.11	139.04	119.70
34	BA	156	U	P-O3'-C3'	16.08	139.00	119.70
85	AA	330	C	C2-N1-C1'	16.07	136.47	118.80
34	BA	1735	G	C5-C6-O6	-16.05	118.97	128.60
85	AA	1495	G	P-O5'-C5'	16.04	146.56	120.90
85	AA	1168	C	C6-N1-C2	-16.04	113.89	120.30
85	AA	1153	G	C5-C6-O6	-16.03	118.98	128.60
39	BF	21	C	P-O3'-C3'	-16.03	100.47	119.70
34	BA	1414	C	C6-N1-C2	-16.02	113.89	120.30
35	BB	1202	G	C5'-C4'-O4'	16.02	128.32	109.10
35	BB	4	C	P-O3'-C3'	16.01	138.91	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	99	G	C5-C6-O6	16.01	138.21	128.60
35	BB	1423	U	O4'-C1'-N1	16.00	121.00	108.20
37	BD	88	U	C1'-O4'-C4'	-15.98	97.11	109.90
86	AB	4	C	P-O3'-C3'	15.98	138.88	119.70
85	AA	913	U	P-O5'-C5'	15.98	146.47	120.90
85	AA	1211	C	P-O3'-C3'	-15.96	100.55	119.70
34	BA	214	A	C8-N9-C4	15.95	112.18	105.80
85	AA	708	G	P-O3'-C3'	15.95	138.84	119.70
34	BA	594	G	O4'-C1'-N9	15.93	120.94	108.20
40	BG	169	A	P-O3'-C3'	15.92	138.80	119.70
35	BB	822	G	C8-N9-C4	-15.90	100.04	106.40
38	BE	31	A	P-O5'-C5'	15.89	146.33	120.90
85	AA	1458	G	C5-C6-O6	15.89	138.14	128.60
40	BG	85	C	P-O3'-C3'	15.89	138.77	119.70
85	AA	760	U	O4'-C1'-N1	15.87	120.90	108.20
85	AA	895	C	O4'-C1'-N1	15.86	120.89	108.20
38	BE	127	G	N1-C6-O6	-15.86	110.39	119.90
34	BA	1614	G	C5-C6-O6	-15.85	119.09	128.60
34	BA	574	U	P-O3'-C3'	15.84	138.71	119.70
85	AA	117	C	C6-N1-C2	-15.84	113.96	120.30
85	AA	1562	U	P-O3'-C3'	15.84	138.71	119.70
36	BC	33	U	O4'-C1'-N1	15.84	120.87	108.20
35	BB	22	A	N1-C6-N6	-15.83	109.10	118.60
38	BE	207	G	C5-C6-O6	-15.82	119.11	128.60
34	BA	888	G	P-O5'-C5'	15.80	146.19	120.90
85	AA	830	A	P-O3'-C3'	15.77	138.62	119.70
85	AA	2063	C	P-O5'-C5'	15.77	146.12	120.90
85	AA	192	G	P-O3'-C3'	15.73	138.58	119.70
34	BA	740	A	C6-N1-C2	-15.72	109.17	118.60
85	AA	685	U	C6-N1-C2	-15.72	111.57	121.00
35	BB	878	G	P-O5'-C5'	15.71	146.04	120.90
35	BB	1507	U	O4'-C1'-N1	15.71	120.77	108.20
40	BG	171	A	O4'-C1'-N9	15.72	120.77	108.20
36	BC	38	U	C2-N3-C4	-15.70	117.58	127.00
35	BB	976	U	P-O3'-C3'	15.70	138.54	119.70
34	BA	3	G	C8-N9-C1'	15.70	147.41	127.00
38	BE	16	C	P-O5'-C5'	15.70	146.02	120.90
34	BA	743	A	C5-N7-C8	15.68	111.74	103.90
34	BA	568	G	P-O3'-C3'	15.67	138.51	119.70
34	BA	1485	U	P-O3'-C3'	15.67	138.51	119.70
34	BA	164	C	O4'-C1'-N1	15.64	120.72	108.20
34	BA	187	G	P-O3'-C3'	15.64	138.47	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	809	A	P-O5'-C5'	15.63	145.91	120.90
34	BA	1822	U	C2-N3-C4	-15.63	117.62	127.00
35	BB	685	G	P-O3'-C3'	15.63	138.45	119.70
34	BA	281	C	C6-N1-C2	-15.60	114.06	120.30
85	AA	821	U	C6-N1-C2	-15.59	111.64	121.00
34	BA	546	U	P-O3'-C3'	15.58	138.40	119.70
34	BA	209	A	C8-N9-C4	-15.58	99.57	105.80
85	AA	1662	U	P-O3'-C3'	15.56	138.37	119.70
34	BA	1795	A	P-O3'-C3'	15.55	138.36	119.70
37	BD	21	G	P-O3'-C3'	15.55	138.36	119.70
85	AA	1000	U	C2-N3-C4	-15.55	117.67	127.00
35	BB	901	U	O4'-C1'-N1	15.54	120.63	108.20
85	AA	1526	G	P-O3'-C3'	15.54	138.35	119.70
85	AA	1367	C	O4'-C1'-N1	15.54	120.63	108.20
35	BB	23	U	C2-N3-C4	-15.54	117.68	127.00
34	BA	372	U	C2-N1-C1'	-15.53	99.06	117.70
36	BC	13	U	C2-N3-C4	-15.53	117.68	127.00
36	BC	139	A	P-O3'-C3'	15.53	138.34	119.70
41	BH	81	U	P-O3'-C3'	15.52	138.32	119.70
35	BB	1231	U	P-O3'-C3'	15.52	138.32	119.70
85	AA	65	A	C5'-C4'-C3'	15.51	140.82	116.00
85	AA	1368	G	P-O5'-C5'	15.51	145.72	120.90
35	BB	1308	G	P-O3'-C3'	15.51	138.31	119.70
85	AA	488	G	P-O3'-C3'	15.48	138.27	119.70
40	BG	181	C	O4'-C1'-N1	15.45	120.56	108.20
34	BA	664	C	C6-N1-C2	-15.45	114.12	120.30
85	AA	504	U	P-O3'-C3'	15.43	138.22	119.70
85	AA	2127	G	C5-C6-O6	-15.43	119.34	128.60
85	AA	884	A	N1-C6-N6	-15.42	109.35	118.60
34	BA	185	A	P-O5'-C5'	15.40	145.55	120.90
85	AA	773	G	C5-C6-O6	-15.40	119.36	128.60
85	AA	174	U	P-O3'-C3'	15.39	138.16	119.70
34	BA	862	C	OP1-P-O3'	15.37	139.02	105.20
85	AA	2225	G	C5-C6-O6	-15.36	119.38	128.60
35	BB	1059	U	C2-N3-C4	-15.35	117.79	127.00
85	AA	1964	A	O4'-C1'-N9	15.32	120.46	108.20
34	BA	401	A	N1-C6-N6	-15.32	109.41	118.60
40	BG	174	G	C5-C6-O6	-15.32	119.41	128.60
85	AA	1116	G	N1-C6-O6	-15.31	110.71	119.90
34	BA	875	G	P-O5'-C5'	15.30	145.39	120.90
85	AA	150	U	C6-N1-C2	-15.30	111.82	121.00
35	BB	578	G	C5-C6-O6	-15.28	119.43	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1921	G	P-O5'-C5'	-15.28	96.46	120.90
41	BH	58	C	P-O3'-C3'	15.27	138.03	119.70
34	BA	692	U	P-O5'-C5'	15.26	145.31	120.90
36	BC	113	G	P-O5'-C5'	15.25	145.30	120.90
85	AA	286	C	C6-N1-C2	-15.24	114.20	120.30
35	BB	29	C	P-O3'-C3'	15.22	137.96	119.70
40	BG	113	G	P-O3'-C3'	15.21	137.95	119.70
34	BA	246	G	P-O3'-C3'	15.20	137.94	119.70
34	BA	1619	U	P-O3'-C3'	15.17	137.91	119.70
39	BF	63	U	P-O3'-C3'	15.16	137.89	119.70
35	BB	561	C	N3-C4-N4	15.16	128.61	118.00
34	BA	1723	U	O4'-C1'-N1	15.15	120.32	108.20
34	BA	1112	U	C2-N3-C4	-15.14	117.92	127.00
37	BD	93	G	N1-C6-O6	-15.14	110.82	119.90
85	AA	265	A	P-O3'-C3'	15.11	137.83	119.70
36	BC	108	A	P-O5'-C5'	15.10	145.06	120.90
34	BA	1138	C	C6-N1-C2	-15.10	114.26	120.30
85	AA	1533	C	C6-N1-C2	-15.10	114.26	120.30
85	AA	603	C	P-O5'-C5'	15.09	145.03	120.90
35	BB	314	A	P-O3'-C3'	15.06	137.78	119.70
34	BA	801	U	P-O5'-C5'	15.05	144.99	120.90
38	BE	24	G	O5'-P-OP2	-15.05	92.15	105.70
65	Bf	157	TRP	CB-CG-CD2	-15.05	107.03	126.60
41	BH	101	A	O5'-P-OP2	-15.05	92.16	105.70
34	BA	1790	U	P-O5'-C5'	15.04	144.97	120.90
34	BA	1626	U	C6-N1-C2	-15.04	111.98	121.00
35	BB	1044	U	P-O3'-C3'	15.04	137.75	119.70
85	AA	1735	U	C6-N1-C2	-15.03	111.98	121.00
85	AA	1235	G	C5-C6-O6	-15.03	119.58	128.60
85	AA	1275	A	P-O5'-C5'	15.03	144.94	120.90
77	Br	217	ARG	NE-CZ-NH1	15.02	127.81	120.30
85	AA	682	C	P-O3'-C3'	15.01	137.71	119.70
37	BD	94	C	C6-N1-C2	-15.01	114.30	120.30
39	BF	11	C	P-O3'-C3'	15.01	137.71	119.70
41	BH	104	U	C2-N3-C4	-15.01	118.00	127.00
35	BB	653	G	O4'-C1'-N9	15.00	120.20	108.20
35	BB	261	C	P-O3'-C3'	15.00	137.70	119.70
35	BB	1534	U	C2-N3-C4	-14.99	118.00	127.00
35	BB	431	U	C2-N1-C1'	-14.99	99.71	117.70
85	AA	1795	C	C6-N1-C2	-14.98	114.31	120.30
85	AA	252	G	P-O3'-C3'	14.98	137.67	119.70
85	AA	1896	G	P-O3'-C3'	14.96	137.66	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	267	U	P-O5'-C5'	14.96	144.84	120.90
34	BA	1732	A	C8-N9-C4	-14.96	99.82	105.80
35	BB	896	C	P-O3'-C3'	14.95	137.64	119.70
40	BG	150	A	C5-C6-N6	-14.94	111.75	123.70
40	BG	172	C	C2-N3-C4	-14.94	112.43	119.90
35	BB	975	G	P-O5'-C5'	14.93	144.79	120.90
41	BH	128	G	N1-C6-O6	14.93	128.86	119.90
85	AA	196	U	C2-N3-C4	-14.93	118.04	127.00
34	BA	800	G	C8-N9-C4	-14.91	100.43	106.40
34	BA	117	C	O4'-C1'-N1	14.91	120.13	108.20
34	BA	547	C	C4'-C3'-C2'	-14.91	87.69	102.60
35	BB	147	C	P-O3'-C3'	14.90	137.58	119.70
35	BB	1454	G	N1-C6-O6	14.90	128.84	119.90
85	AA	1921	G	C5'-C4'-O4'	-14.89	91.23	109.10
35	BB	816	U	C2-N3-C4	-14.88	118.07	127.00
66	Bg	96	PHE	CB-CG-CD2	-14.88	110.38	120.80
85	AA	859	G	C4-N9-C1'	-14.88	107.16	126.50
85	AA	1237	A	O4'-C1'-N9	14.87	120.10	108.20
85	AA	251	A	O4'-C1'-N9	14.87	120.10	108.20
85	AA	76	G	P-O3'-C3'	-14.87	101.86	119.70
37	BD	64	A	P-O5'-C5'	14.86	144.67	120.90
34	BA	1801	G	C5'-C4'-C3'	14.85	139.76	116.00
85	AA	1209	U	P-O5'-C5'	14.85	144.66	120.90
34	BA	1295	U	C5'-C4'-C3'	14.84	139.75	116.00
35	BB	561	C	C5-C4-N4	-14.84	109.81	120.20
38	BE	117	A	N9-C4-C5	-14.82	99.87	105.80
35	BB	68	G	P-O3'-C3'	14.82	137.48	119.70
34	BA	240	C	C6-N1-C2	-14.81	114.37	120.30
35	BB	778	A	C5'-C4'-C3'	-14.81	92.31	116.00
85	AA	1153	G	N1-C6-O6	14.80	128.78	119.90
38	BE	1	U	C2-N3-C4	-14.79	118.13	127.00
34	BA	1303	U	C2-N3-C4	-14.78	118.13	127.00
34	BA	1477	C	C1'-O4'-C4'	-14.78	98.07	109.90
85	AA	100	A	P-O5'-C5'	14.78	144.54	120.90
38	BE	100	U	O4'-C1'-N1	14.77	120.02	108.20
85	AA	743	C	P-O3'-C3'	14.77	137.43	119.70
34	BA	579	U	P-O3'-C3'	14.76	137.42	119.70
36	BC	116	C	P-O3'-C3'	14.76	137.41	119.70
85	AA	252	G	C5-C6-O6	-14.76	119.74	128.60
34	BA	471	U	N1-C2-O2	-14.76	112.47	122.80
34	BA	757	G	P-O3'-C3'	-14.76	101.99	119.70
34	BA	774	A	P-O5'-C5'	14.76	144.51	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1057	G	P-O3'-C3'	14.75	137.40	119.70
85	AA	1235	G	C6-N1-C2	-14.74	116.25	125.10
35	BB	651	G	P-O3'-C3'	14.74	137.38	119.70
35	BB	1464	G	N1-C6-O6	-14.74	111.06	119.90
34	BA	230	A	C5'-C4'-C3'	14.73	139.57	116.00
35	BB	973	G	P-O3'-C3'	14.73	137.38	119.70
35	BB	1024	G	C8-N9-C4	-14.73	100.51	106.40
85	AA	2218	G	C5'-C4'-C3'	14.73	139.56	116.00
85	AA	179	G	C5-C6-O6	-14.73	119.76	128.60
85	AA	1362	A	C5'-C4'-C3'	14.72	139.56	116.00
34	BA	865	C	O5'-P-OP2	-14.72	92.45	105.70
34	BA	593	G	P-O5'-C5'	14.72	144.45	120.90
85	AA	900	G	P-O3'-C3'	14.72	137.36	119.70
35	BB	1199	A	N1-C6-N6	-14.71	109.78	118.60
35	BB	1490	G	O4'-C4'-C3'	-14.70	89.30	104.00
85	AA	630	A	P-O3'-C3'	14.69	137.33	119.70
85	AA	374	C	O4'-C1'-N1	14.68	119.94	108.20
34	BA	587	U	C5'-C4'-C3'	-14.67	92.53	116.00
40	BG	169	A	O4'-C1'-N9	14.67	119.94	108.20
35	BB	764	C	C6-N1-C2	-14.66	114.44	120.30
85	AA	2119	C	C2-N3-C4	-14.65	112.58	119.90
85	AA	1016	G	P-O3'-C3'	14.59	137.21	119.70
34	BA	138	C	C6-N1-C2	-14.57	114.47	120.30
85	AA	469	G	C6-N1-C2	14.57	133.84	125.10
41	BH	47	G	N1-C6-O6	-14.57	111.16	119.90
85	AA	381	A	C5-C6-N6	14.56	135.35	123.70
85	AA	894	A	O4'-C1'-N9	14.56	119.85	108.20
34	BA	1724	G	N1-C6-O6	-14.56	111.17	119.90
85	AA	1465	C	C6-N1-C2	-14.55	114.48	120.30
35	BB	657	A	N1-C6-N6	-14.55	109.87	118.60
85	AA	2249	U	O4'-C1'-N1	14.54	119.83	108.20
41	BH	128	G	C8-N9-C4	14.53	112.21	106.40
85	AA	1035	C	O4'-C1'-N1	14.53	119.82	108.20
37	BD	84	U	C6-N1-C2	-14.52	112.29	121.00
34	BA	793	A	P-O5'-C5'	14.52	144.12	120.90
82	Bw	62	ARG	NE-CZ-NH1	14.50	127.55	120.30
34	BA	706	C	C6-N1-C2	-14.49	114.50	120.30
85	AA	997	U	C2-N3-C4	-14.48	118.31	127.00
85	AA	2121	G	C8-N9-C4	-14.47	100.61	106.40
34	BA	548	G	P-O5'-C5'	14.46	144.03	120.90
65	Bf	391	ARG	NE-CZ-NH1	14.46	127.53	120.30
34	BA	780	U	P-O5'-C5'	14.46	144.03	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	31	A	C5'-C4'-C3'	14.44	139.11	116.00
85	AA	599	C	P-O3'-C3'	14.44	137.02	119.70
85	AA	1989	A	P-O3'-C3'	14.43	137.02	119.70
49	BP	46	MET	CG-SD-CE	-14.42	77.12	100.20
34	BA	813	C	C2-N1-C1'	-14.41	102.94	118.80
34	BA	1425	G	O4'-C1'-N9	14.41	119.73	108.20
40	BG	16	G	N1-C6-O6	-14.41	111.25	119.90
38	BE	108	U	O4'-C1'-N1	14.41	119.73	108.20
41	BH	33	G	P-O3'-C3'	14.39	136.97	119.70
41	BH	32	U	C5'-C4'-C3'	-14.39	92.98	116.00
35	BB	957	A	P-O3'-C3'	14.38	136.96	119.70
36	BC	23	G	N1-C6-O6	-14.39	111.27	119.90
38	BE	25	U	P-O5'-C5'	14.38	143.91	120.90
85	AA	1860	A	P-O3'-C3'	14.38	136.96	119.70
36	BC	100	U	C2-N3-C4	-14.37	118.38	127.00
34	BA	690	G	P-O3'-C3'	14.37	136.94	119.70
54	BU	7	TYR	CB-CG-CD1	14.36	129.62	121.00
85	AA	1493	A	C4'-C3'-C2'	-14.36	88.24	102.60
85	AA	1923	A	C1'-O4'-C4'	-14.36	98.41	109.90
34	BA	1299	G	C8-N9-C1'	14.35	145.66	127.00
35	BB	68	G	C5-C6-O6	-14.35	119.99	128.60
40	BG	174	G	N1-C6-O6	14.35	128.51	119.90
34	BA	433	G	C5'-C4'-C3'	14.34	138.94	116.00
40	BG	84	U	C2-N3-C4	-14.34	118.40	127.00
34	BA	449	G	C4-N9-C1'	-14.33	107.87	126.50
85	AA	1539	A	P-O3'-C3'	14.33	136.90	119.70
85	AA	1960	C	C6-N1-C2	-14.32	114.57	120.30
35	BB	838	G	O5'-P-OP1	-14.31	92.82	105.70
85	AA	1718	C	P-O3'-C3'	14.31	136.87	119.70
85	AA	2107	C	C6-N1-C2	-14.30	114.58	120.30
34	BA	3	G	C4-N9-C1'	-14.30	107.91	126.50
41	BH	75	G	C5-C6-O6	-14.30	120.02	128.60
85	AA	26	A	P-O5'-C5'	14.30	143.78	120.90
35	BB	1436	U	P-O3'-C3'	14.30	136.85	119.70
35	BB	132	G	P-O3'-C3'	14.29	136.85	119.70
85	AA	745	C	P-O5'-C5'	14.29	143.76	120.90
85	AA	171	U	C6-N1-C2	-14.27	112.44	121.00
34	BA	1846	G	C6-N1-C2	-14.26	116.54	125.10
85	AA	895	C	C1'-O4'-C4'	-14.26	98.49	109.90
85	AA	766	G	C5'-C4'-C3'	-14.25	93.20	116.00
34	BA	570	G	O4'-C1'-N9	14.24	119.59	108.20
35	BB	1152	U	C2-N3-C4	-14.24	118.46	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	611	G	C5-C6-O6	-14.23	120.06	128.60
85	AA	1921	G	C8-N9-C1'	14.23	145.50	127.00
85	AA	1447	U	P-O3'-C3'	14.23	136.77	119.70
36	BC	33	U	P-O3'-C3'	14.22	136.77	119.70
34	BA	231	U	O3'-P-O5'	14.22	131.01	104.00
35	BB	995	C	P-O3'-C3'	14.22	136.76	119.70
34	BA	222	C	C6-N1-C1'	14.21	137.85	120.80
34	BA	1477	C	O4'-C1'-N1	14.21	119.56	108.20
40	BG	163	G	C5-C6-O6	-14.19	120.09	128.60
34	BA	1305	A	N1-C6-N6	-14.18	110.09	118.60
34	BA	858	C	C6-N1-C2	-14.18	114.63	120.30
85	AA	108	C	O4'-C1'-N1	14.18	119.54	108.20
85	AA	1978	G	N1-C6-O6	14.18	128.41	119.90
34	BA	871	G	O4'-C1'-N9	14.17	119.54	108.20
34	BA	896	U	C4'-C3'-C2'	-14.17	88.43	102.60
38	BE	20	C	C1'-O4'-C4'	-14.16	98.57	109.90
34	BA	1486	U	C2-N1-C1'	-14.16	100.71	117.70
40	BG	122	G	C5-C6-O6	-14.15	120.11	128.60
34	BA	513	U	N3-C2-O2	-14.14	112.30	122.20
34	BA	515	U	P-O5'-C5'	14.14	143.52	120.90
34	BA	1200	U	O4'-C1'-N1	14.13	119.51	108.20
34	BA	1299	G	C4-N9-C1'	-14.13	108.13	126.50
85	AA	1797	U	C2-N3-C4	-14.13	118.52	127.00
34	BA	215	C	C6-N1-C2	-14.12	114.65	120.30
38	BE	155	C	O4'-C1'-N1	14.12	119.49	108.20
38	BE	170	U	P-O3'-C3'	-14.12	102.76	119.70
38	BE	91	G	C4-N9-C1'	-14.11	108.15	126.50
38	BE	111	C	P-O3'-C3'	14.11	136.63	119.70
34	BA	484	A	P-O3'-C3'	14.10	136.62	119.70
85	AA	861	G	P-O3'-C3'	14.10	136.62	119.70
35	BB	802	G	O4'-C1'-N9	14.10	119.48	108.20
34	BA	683	C	C6-N1-C2	-14.09	114.66	120.30
36	BC	110	A	P-O3'-C3'	14.09	136.60	119.70
38	BE	202	C	O4'-C1'-N1	14.09	119.47	108.20
34	BA	211	C	C6-N1-C2	-14.09	114.67	120.30
34	BA	546	U	P-O5'-C5'	14.08	143.43	120.90
38	BE	117	A	C8-N9-C1'	-14.08	102.35	127.70
38	BE	107	U	C5'-C4'-C3'	-14.08	93.47	116.00
38	BE	207	G	N1-C6-O6	14.07	128.34	119.90
85	AA	1235	G	N1-C6-O6	14.07	128.34	119.90
35	BB	1201	G	C4'-C3'-C2'	-14.06	88.54	102.60
34	BA	1318	G	O4'-C1'-N9	14.06	119.45	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	269	G	P-O5'-C5'	14.06	143.40	120.90
85	AA	1373	U	C6-N1-C2	-14.06	112.56	121.00
85	AA	1644	G	C8-N9-C4	-14.06	100.78	106.40
85	AA	1959	G	C5-C6-O6	-14.06	120.17	128.60
85	AA	1466	U	C5'-C4'-C3'	14.06	138.49	116.00
36	BC	33	U	C5'-C4'-C3'	-14.05	93.52	116.00
38	BE	107	U	P-O5'-C5'	14.05	143.37	120.90
85	AA	65	A	P-O5'-C5'	14.05	143.37	120.90
35	BB	1251	G	C4-N9-C1'	-14.04	108.25	126.50
85	AA	1011	G	P-O3'-C3'	14.04	136.55	119.70
34	BA	1205	A	N1-C6-N6	-14.04	110.18	118.60
34	BA	847	U	O4'-C1'-N1	14.04	119.43	108.20
34	BA	205	G	C5-C6-O6	-14.03	120.18	128.60
35	BB	1283	C	C6-N1-C2	-14.03	114.69	120.30
85	AA	1238	U	C2-N3-C4	-14.03	118.58	127.00
40	BG	64	C	C6-N1-C2	-14.03	114.69	120.30
40	BG	78	C	C6-N1-C2	-14.02	114.69	120.30
40	BG	25	G	O4'-C1'-N9	14.01	119.41	108.20
85	AA	1473	U	P-O3'-C3'	14.01	136.51	119.70
39	BF	38	C	C6-N1-C2	-14.01	114.70	120.30
35	BB	1503	U	P-O3'-C3'	14.00	136.50	119.70
34	BA	1299	G	C4-C5-C6	-14.00	110.40	118.80
85	AA	314	C	P-O5'-C5'	14.00	143.30	120.90
34	BA	507	U	C5'-C4'-C3'	13.99	138.38	116.00
51	BR	56	ARG	NE-CZ-NH1	13.99	127.30	120.30
38	BE	208	G	P-O3'-C3'	13.98	136.48	119.70
38	BE	26	G	O5'-P-OP2	-13.98	93.12	105.70
34	BA	1565	U	C5'-C4'-C3'	13.97	138.36	116.00
34	BA	1816	G	C5-C6-O6	-13.97	120.22	128.60
85	AA	486	G	C4-N9-C1'	-13.96	108.35	126.50
85	AA	280	U	P-O3'-C3'	13.96	136.46	119.70
34	BA	1699	A	N1-C6-N6	-13.95	110.23	118.60
85	AA	1292	A	N1-C6-N6	13.95	126.97	118.60
85	AA	1701	G	N3-C2-N2	-13.95	110.13	119.90
34	BA	605	G	C6-N1-C2	-13.94	116.74	125.10
35	BB	529	A	N1-C6-N6	13.94	126.96	118.60
34	BA	472	G	C4-N9-C1'	13.94	144.62	126.50
85	AA	1535	C	O4'-C1'-N1	13.93	119.34	108.20
35	BB	60	A	P-O3'-C3'	13.93	136.41	119.70
34	BA	556	A	P-O3'-C3'	-13.92	103.00	119.70
34	BA	685	C	O4'-C1'-N1	13.92	119.33	108.20
34	BA	896	U	C6-N1-C1'	-13.91	101.73	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1894	G	C5-C6-O6	-13.90	120.26	128.60
34	BA	1566	G	C5-C6-O6	-13.89	120.26	128.60
38	BE	72	C	P-O5'-C5'	13.88	143.11	120.90
85	AA	1759	U	P-O3'-C3'	13.88	136.35	119.70
34	BA	665	C	C6-N1-C2	-13.87	114.75	120.30
34	BA	765	U	C6-N1-C2	-13.87	112.68	121.00
85	AA	388	G	P-O3'-C3'	13.87	136.34	119.70
35	BB	1512	C	O4'-C1'-N1	13.86	119.29	108.20
36	BC	84	U	P-O3'-C3'	13.85	136.32	119.70
34	BA	687	G	P-O3'-C3'	13.85	136.32	119.70
34	BA	238	C	C6-N1-C2	-13.85	114.76	120.30
36	BC	80	A	P-O3'-C3'	13.84	136.31	119.70
35	BB	2	C	O5'-C5'-C4'	-13.84	85.41	111.70
34	BA	618	G	C8-N9-C1'	13.83	144.98	127.00
34	BA	605	G	C5'-C4'-C3'	-13.83	93.88	116.00
34	BA	8	G	C4-N9-C1'	-13.82	108.53	126.50
34	BA	828	A	N1-C6-N6	-13.82	110.31	118.60
85	AA	25	C	P-O3'-C3'	13.82	136.28	119.70
85	AA	628	C	C6-N1-C2	-13.82	114.77	120.30
34	BA	1181	G	P-O5'-C5'	13.81	143.00	120.90
35	BB	1393	C	P-O3'-C3'	13.81	136.27	119.70
34	BA	756	A	P-O5'-C5'	13.81	142.99	120.90
86	AB	56	C	P-O3'-C3'	13.80	136.26	119.70
34	BA	866	C	O4'-C1'-N1	13.80	119.24	108.20
35	BB	622	G	C5-C6-O6	-13.80	120.32	128.60
34	BA	1200	U	P-O3'-C3'	13.79	136.25	119.70
40	BG	163	G	N1-C6-O6	13.79	128.17	119.90
41	BH	27	A	P-O3'-C3'	13.79	136.24	119.70
35	BB	824	C	O4'-C1'-N1	13.78	119.23	108.20
34	BA	488	C	P-O3'-C3'	13.77	136.23	119.70
85	AA	1654	G	C5-C6-O6	-13.77	120.34	128.60
34	BA	804	G	N1-C6-O6	-13.76	111.64	119.90
35	BB	1317	U	C2-N3-C4	-13.76	118.74	127.00
36	BC	86	U	O4'-C1'-N1	13.76	119.21	108.20
85	AA	1824	G	P-O3'-C3'	13.76	136.21	119.70
34	BA	1493	U	O4'-C1'-N1	13.75	119.20	108.20
85	AA	1355	U	C5'-C4'-C3'	13.74	137.99	116.00
85	AA	313	A	C4-C5-C6	-13.74	110.13	117.00
35	BB	878	G	C5'-C4'-C3'	13.74	137.98	116.00
34	BA	1282	G	N1-C6-O6	13.73	128.13	119.90
35	BB	798	A	N1-C6-N6	13.72	126.83	118.60
34	BA	796	G	C5-C6-O6	-13.72	120.37	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	546	A	C5-C6-N6	-13.71	112.73	123.70
45	BL	56	ARG	NE-CZ-NH1	13.71	127.16	120.30
85	AA	985	G	C5'-C4'-C3'	-13.71	94.07	116.00
85	AA	1898	C	P-O3'-C3'	13.69	136.13	119.70
34	BA	1052	G	P-O3'-C3'	13.69	136.13	119.70
85	AA	2197	A	C5'-C4'-C3'	13.68	137.89	116.00
34	BA	1634	A	P-O3'-C3'	13.67	136.11	119.70
35	BB	380	G	P-O3'-C3'	13.67	136.10	119.70
85	AA	1976	G	C5'-C4'-C3'	-13.66	94.14	116.00
41	BH	116	A	P-O3'-C3'	13.65	136.08	119.70
34	BA	189	G	C5'-C4'-C3'	-13.65	94.16	116.00
85	AA	1440	C	P-O3'-C3'	13.63	136.06	119.70
35	BB	1532	C	C2-N1-C1'	-13.63	103.81	118.80
85	AA	185	A	P-O5'-C5'	13.63	142.70	120.90
85	AA	2001	C	P-O3'-C3'	13.63	136.06	119.70
34	BA	572	G	P-O3'-C3'	13.62	136.05	119.70
85	AA	273	C	C6-N1-C2	-13.62	114.85	120.30
85	AA	991	G	C5-C6-O6	-13.62	120.43	128.60
7	A6	163	PHE	CB-CG-CD2	-13.61	111.27	120.80
35	BB	54	U	C2-N3-C4	-13.61	118.83	127.00
37	BD	48	G	C5-C6-O6	-13.61	120.43	128.60
85	AA	755	G	C8-N9-C4	-13.60	100.96	106.40
35	BB	1202	G	C1'-O4'-C4'	-13.60	99.02	109.90
85	AA	924	A	C5'-C4'-C3'	-13.60	94.24	116.00
85	AA	1301	C	C2-N3-C4	-13.59	113.11	119.90
85	AA	1457	C	C6-N1-C2	-13.58	114.87	120.30
35	BB	964	G	C4-N9-C1'	-13.57	108.86	126.50
85	AA	1716	U	P-O3'-C3'	13.56	135.97	119.70
34	BA	930	A	N1-C6-N6	-13.56	110.47	118.60
38	BE	29	C	C6-N1-C2	-13.55	114.88	120.30
34	BA	800	G	C5-C6-O6	-13.55	120.47	128.60
34	BA	111	U	P-O3'-C3'	13.54	135.95	119.70
85	AA	859	G	C8-N9-C1'	13.54	144.61	127.00
85	AA	730	G	C6-N1-C2	-13.54	116.98	125.10
35	BB	1492	C	C6-N1-C1'	13.54	137.04	120.80
85	AA	58	C	P-O3'-C3'	13.53	135.94	119.70
85	AA	854	A	C5'-C4'-O4'	13.53	125.34	109.10
40	BG	13	A	P-O3'-C3'	13.53	135.93	119.70
34	BA	1440	C	C5'-C4'-C3'	-13.52	94.37	116.00
35	BB	1150	A	N1-C6-N6	-13.52	110.49	118.60
27	AT	111	ARG	NE-CZ-NH2	13.52	127.06	120.30
34	BA	334	G	N1-C6-O6	-13.52	111.79	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	31	G	C5-C6-O6	-13.52	120.49	128.60
34	BA	429	G	C6-C5-N7	-13.51	122.30	130.40
35	BB	1359	G	C5-C6-O6	-13.51	120.50	128.60
34	BA	1676	A	P-O5'-C5'	13.50	142.50	120.90
34	BA	1055	U	C2-N3-C4	-13.50	118.90	127.00
34	BA	131	A	P-O3'-C3'	13.49	135.90	119.70
39	BF	16	C	P-O5'-C5'	13.49	142.49	120.90
34	BA	1217	A	P-O5'-C5'	13.49	142.49	120.90
37	BD	14	C	P-O5'-C5'	13.49	142.49	120.90
85	AA	1886	U	P-O3'-C3'	13.49	135.89	119.70
85	AA	165	C	P-O5'-C5'	-13.49	99.32	120.90
85	AA	930	G	C5-C6-O6	-13.48	120.51	128.60
85	AA	1022	G	P-O3'-C3'	13.48	135.88	119.70
85	AA	2117	U	C2-N3-C4	-13.48	118.91	127.00
34	BA	1561	C	O5'-P-OP1	-13.47	93.58	105.70
38	BE	108	U	O4'-C4'-C3'	-13.47	90.53	104.00
34	BA	1478	G	C5'-C4'-C3'	13.46	137.54	116.00
34	BA	572	G	C8-N9-C4	-13.46	101.02	106.40
34	BA	874	G	C5-C6-O6	-13.46	120.53	128.60
34	BA	1227	U	C5'-C4'-C3'	13.46	137.53	116.00
85	AA	740	A	N1-C6-N6	-13.45	110.53	118.60
38	BE	136	G	C5-C6-O6	-13.44	120.54	128.60
85	AA	23	G	P-O3'-C3'	13.44	135.83	119.70
85	AA	807	A	O4'-C1'-N9	13.44	118.95	108.20
41	BH	129	G	C5'-C4'-C3'	-13.43	94.51	116.00
34	BA	1003	A	C4'-C3'-C2'	13.42	116.02	102.60
85	AA	569	A	N1-C6-N6	-13.42	110.55	118.60
34	BA	1592	U	C5'-C4'-C3'	-13.42	94.53	116.00
85	AA	1482	C	C6-N1-C2	-13.42	114.93	120.30
85	AA	1756	C	C6-N1-C2	-13.41	114.94	120.30
34	BA	1735	G	N9-C4-C5	-13.39	100.04	105.40
85	AA	881	C	C6-N1-C2	-13.38	114.95	120.30
39	BF	51	C	C6-N1-C2	-13.38	114.95	120.30
34	BA	665	C	C6-N1-C1'	13.38	136.85	120.80
34	BA	674	G	C1'-O4'-C4'	-13.37	99.20	109.90
35	BB	487	A	C5'-C4'-C3'	13.37	137.40	116.00
34	BA	559	C	P-O3'-C3'	13.37	135.75	119.70
38	BE	195	G	C5-C6-N1	13.36	118.18	111.50
38	BE	147	G	C5-C6-O6	-13.36	120.59	128.60
38	BE	95	G	N1-C6-O6	-13.35	111.89	119.90
85	AA	1469	G	P-O3'-C3'	13.34	135.71	119.70
34	BA	548	G	C5'-C4'-C3'	13.33	137.33	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2234	C	P-O5'-C5'	13.33	142.22	120.90
35	BB	578	G	N1-C6-O6	13.32	127.89	119.90
34	BA	1538	G	O3'-P-O5'	13.32	129.31	104.00
35	BB	353	G	P-O3'-C3'	13.32	135.68	119.70
54	BU	7	TYR	CB-CG-CD2	-13.32	113.01	121.00
85	AA	1728	G	P-O3'-C3'	13.32	135.69	119.70
85	AA	2078	A	P-O3'-C3'	13.32	135.69	119.70
86	AB	56	C	C6-N1-C2	-13.32	114.97	120.30
57	BX	82	PHE	CB-CG-CD1	13.31	130.12	120.80
85	AA	179	G	N1-C6-O6	13.31	127.88	119.90
85	AA	1991	C	C6-N1-C2	-13.31	114.98	120.30
35	BB	837	A	P-O5'-C5'	13.30	142.18	120.90
38	BE	27	A	N1-C6-N6	13.30	126.58	118.60
39	BF	33	C	O3'-P-O5'	13.30	129.26	104.00
85	AA	2074	G	P-O3'-C3'	13.30	135.66	119.70
85	AA	805	A	O4'-C1'-N9	13.29	118.83	108.20
85	AA	569	A	O4'-C1'-N9	13.28	118.83	108.20
35	BB	1025	A	O4'-C1'-N9	13.28	118.82	108.20
34	BA	757	G	C3'-C2'-C1'	-13.28	90.88	101.50
85	AA	509	C	O4'-C1'-N1	13.28	118.82	108.20
34	BA	453	A	C5'-C4'-C3'	-13.26	94.78	116.00
34	BA	673	U	C5'-C4'-C3'	13.26	137.22	116.00
17	AI	104	TYR	CB-CG-CD2	-13.26	113.05	121.00
35	BB	1227	G	P-O3'-C3'	13.25	135.60	119.70
85	AA	308	U	P-O3'-C3'	-13.25	103.80	119.70
34	BA	777	C	P-O3'-C3'	13.25	135.59	119.70
41	BH	24	U	C1'-O4'-C4'	-13.24	99.31	109.90
37	BD	82	G	C4-N9-C1'	-13.24	109.28	126.50
85	AA	1878	C	C6-N1-C2	-13.24	115.00	120.30
34	BA	972	C	P-O3'-C3'	13.24	135.59	119.70
35	BB	1480	G	C4-N9-C1'	-13.24	109.29	126.50
35	BB	818	U	C2-N1-C1'	-13.23	101.82	117.70
34	BA	561	U	O3'-P-O5'	13.23	129.13	104.00
37	BD	87	G	C4-N9-C1'	-13.23	109.30	126.50
38	BE	104	G	O4'-C1'-N9	13.22	118.78	108.20
35	BB	405	U	C2-N3-C4	-13.21	119.07	127.00
40	BG	1	G	C5-C6-O6	-13.21	120.67	128.60
34	BA	83	G	O4'-C1'-N9	13.21	118.77	108.20
35	BB	1052	G	C5-C6-O6	-13.21	120.67	128.60
34	BA	1320	A	C4-C5-C6	-13.21	110.39	117.00
41	BH	28	U	C2-N3-C4	-13.21	119.07	127.00
36	BC	124	A	C5-C6-N6	-13.21	113.13	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	30	C	C1'-O4'-C4'	-13.21	99.34	109.90
56	BW	44	TYR	CB-CG-CD2	-13.20	113.08	121.00
41	BH	113	G	C5-C6-O6	-13.20	120.68	128.60
85	AA	2150	G	C8-N9-C1'	13.19	144.15	127.00
38	BE	195	G	C6-N1-C2	-13.19	117.19	125.10
35	BB	1226	G	C5-C6-O6	-13.18	120.69	128.60
34	BA	674	G	O4'-C1'-N9	13.17	118.74	108.20
34	BA	7	U	C2-N1-C1'	-13.17	101.90	117.70
34	BA	1454	G	P-O3'-C3'	13.16	135.50	119.70
85	AA	1900	C	O5'-P-OP1	-13.16	93.85	105.70
34	BA	1202	G	C5-C6-O6	13.16	136.50	128.60
40	BG	9	G	C4-N9-C1'	-13.16	109.39	126.50
34	BA	1723	U	O4'-C1'-C2'	-13.16	92.64	105.80
85	AA	830	A	C5'-C4'-C3'	13.16	137.05	116.00
85	AA	84	C	C5'-C4'-C3'	-13.15	94.96	116.00
85	AA	1911	A	P-O3'-C3'	13.14	135.47	119.70
34	BA	692	U	C5'-C4'-O4'	13.14	124.87	109.10
34	BA	1052	G	C5-C6-O6	-13.14	120.72	128.60
41	BH	22	A	P-O3'-C3'	13.14	135.47	119.70
34	BA	1787	U	O3'-P-O5'	13.14	128.96	104.00
36	BC	111	C	P-O5'-C5'	13.13	141.91	120.90
85	AA	850	U	C5'-C4'-C3'	-13.12	95.00	116.00
34	BA	1792	U	C2-N1-C1'	-13.12	101.96	117.70
20	AL	29	TYR	CB-CG-CD1	13.12	128.87	121.00
34	BA	289	A	O3'-P-O5'	13.12	128.92	104.00
34	BA	1792	U	C6-N1-C1'	13.12	139.56	121.20
85	AA	1287	C	C6-N1-C2	-13.12	115.05	120.30
34	BA	605	G	C8-N9-C1'	-13.12	109.95	127.00
35	BB	1464	G	C5-C6-O6	13.11	136.47	128.60
34	BA	260	A	P-O3'-C3'	-13.10	103.98	119.70
34	BA	599	U	C5'-C4'-O4'	13.10	124.82	109.10
85	AA	2058	C	C3'-C2'-C1'	-13.10	91.02	101.50
34	BA	171	U	C2-N3-C4	-13.09	119.15	127.00
35	BB	875	G	C5'-C4'-C3'	13.08	136.93	116.00
38	BE	85	G	C4-N9-C1'	-13.08	109.49	126.50
38	BE	91	G	C5-C6-O6	-13.08	120.75	128.60
41	BH	30	C	O4'-C1'-N1	13.08	118.66	108.20
85	AA	608	A	C1'-O4'-C4'	-13.07	99.44	109.90
34	BA	1334	G	C5-C6-O6	-13.07	120.76	128.60
40	BG	170	G	O4'-C1'-N9	13.06	118.65	108.20
37	BD	95	G	N3-C4-C5	-13.04	122.08	128.60
34	BA	1480	C	P-O3'-C3'	13.04	135.34	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	772	U	P-O5'-C5'	13.03	141.75	120.90
36	BC	141	C	P-O5'-C5'	13.03	141.75	120.90
85	AA	2122	A	C8-N9-C4	-13.03	100.59	105.80
34	BA	1005	C	P-O3'-C3'	-13.03	104.07	119.70
85	AA	709	A	P-O3'-C3'	-13.02	104.07	119.70
35	BB	1229	A	P-O5'-C5'	13.02	141.73	120.90
34	BA	399	G	N1-C6-O6	13.01	127.71	119.90
34	BA	13	U	P-O5'-C5'	13.01	141.71	120.90
34	BA	683	C	P-O3'-C3'	13.01	135.31	119.70
34	BA	1294	C	C1'-O4'-C4'	-13.00	99.50	109.90
34	BA	515	U	O3'-P-O5'	12.99	128.69	104.00
38	BE	136	G	P-O5'-C5'	12.99	141.69	120.90
85	AA	469	G	N3-C4-C5	-12.99	122.11	128.60
35	BB	546	A	N1-C6-N6	12.99	126.39	118.60
34	BA	222	C	C2-N1-C1'	-12.98	104.52	118.80
34	BA	1211	G	N1-C6-O6	12.98	127.69	119.90
34	BA	1211	G	C5-C6-O6	-12.97	120.81	128.60
40	BG	98	A	C5'-C4'-C3'	-12.97	95.25	116.00
34	BA	1735	G	C6-C5-N7	-12.97	122.62	130.40
85	AA	605	A	N1-C6-N6	12.96	126.38	118.60
35	BB	802	G	C5'-C4'-C3'	-12.96	95.26	116.00
35	BB	802	G	C4'-C3'-C2'	-12.96	89.64	102.60
34	BA	535	G	N1-C6-O6	-12.95	112.13	119.90
34	BA	547	C	C4'-C3'-O3'	12.95	138.90	113.00
85	AA	44	C	C6-N1-C2	-12.95	115.12	120.30
34	BA	1716	A	O4'-C1'-N9	12.94	118.55	108.20
34	BA	563	A	N1-C6-N6	-12.94	110.84	118.60
34	BA	606	G	P-O5'-C5'	12.93	141.59	120.90
36	BC	139	A	C8-N9-C4	-12.93	100.63	105.80
34	BA	1724	G	C5'-C4'-C3'	12.93	136.68	116.00
34	BA	1505	G	P-O3'-C3'	12.92	135.20	119.70
85	AA	469	G	C4-C5-C6	-12.92	111.05	118.80
30	AW	48	TYR	CB-CG-CD2	-12.92	113.25	121.00
34	BA	305	C	P-O5'-C5'	12.91	141.56	120.90
35	BB	1492	C	C2-N1-C1'	-12.91	104.59	118.80
34	BA	1649	A	C5'-C4'-C3'	-12.91	95.34	116.00
85	AA	68	A	P-O3'-C3'	-12.90	104.22	119.70
35	BB	1187	G	N3-C2-N2	-12.90	110.87	119.90
36	BC	123	G	C5-C6-O6	-12.90	120.86	128.60
85	AA	97	A	C8-N9-C4	12.90	110.96	105.80
34	BA	1792	U	C6-N1-C2	-12.89	113.27	121.00
85	AA	327	G	C4-N9-C1'	12.89	143.26	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	468	A	P-O3'-C3'	12.89	135.16	119.70
34	BA	388	A	P-O3'-C3'	12.88	135.16	119.70
35	BB	440	U	C2-N3-C4	-12.88	119.28	127.00
34	BA	1297	G	C5-C6-O6	-12.86	120.88	128.60
34	BA	303	C	P-O3'-C3'	12.86	135.13	119.70
35	BB	365	U	P-O5'-C5'	12.86	141.47	120.90
35	BB	831	C	C5'-C4'-C3'	-12.85	95.44	116.00
85	AA	514	U	P-O5'-C5'	12.85	141.46	120.90
34	BA	618	G	C4-N9-C1'	-12.84	109.80	126.50
34	BA	1068	C	C6-N1-C2	-12.84	115.16	120.30
34	BA	680	C	P-O3'-C3'	12.84	135.11	119.70
35	BB	957	A	O3'-P-O5'	-12.84	79.60	104.00
60	Ba	50	ARG	NE-CZ-NH1	12.84	126.72	120.30
85	AA	1374	A	P-O5'-C5'	12.84	141.44	120.90
85	AA	2196	G	C8-N9-C1'	12.84	143.69	127.00
34	BA	219	U	P-O3'-C3'	-12.84	104.30	119.70
34	BA	1570	C	C6-N1-C2	-12.84	115.17	120.30
85	AA	1505	G	N1-C6-O6	-12.83	112.20	119.90
85	AA	368	C	C2-N3-C4	-12.82	113.49	119.90
34	BA	557	U	C1'-O4'-C4'	-12.81	99.65	109.90
35	BB	1472	U	P-O3'-C3'	12.81	135.07	119.70
34	BA	559	C	O5'-P-OP1	12.81	126.07	110.70
41	BH	63	G	C5-C6-O6	-12.81	120.92	128.60
34	BA	10	G	C4-N9-C1'	-12.80	109.86	126.50
79	Bt	16	ARG	NE-CZ-NH1	12.80	126.70	120.30
38	BE	82	C	O4'-C1'-N1	12.79	118.44	108.20
85	AA	1955	U	P-O3'-C3'	-12.79	104.35	119.70
34	BA	323	C	O4'-C1'-N1	12.79	118.43	108.20
85	AA	2237	G	N1-C6-O6	-12.79	112.22	119.90
35	BB	1124	G	C5-C6-O6	-12.79	120.93	128.60
85	AA	2150	G	C4-N9-C1'	-12.79	109.88	126.50
35	BB	384	A	C5'-C4'-C3'	-12.79	95.54	116.00
85	AA	84	C	C6-N1-C2	-12.78	115.19	120.30
35	BB	1203	C	O4'-C1'-N1	12.76	118.41	108.20
36	BC	157	U	P-O3'-C3'	12.76	135.01	119.70
34	BA	1110	A	P-O3'-C3'	12.75	135.00	119.70
85	AA	615	A	P-O5'-C5'	12.75	141.31	120.90
34	BA	1723	U	C6-N1-C2	-12.75	113.35	121.00
36	BC	129	C	O4'-C1'-N1	12.74	118.39	108.20
85	AA	252	G	N1-C6-O6	12.74	127.55	119.90
86	AB	62	C	C6-N1-C2	-12.74	115.20	120.30
65	Bf	406	ARG	NE-CZ-NH1	12.74	126.67	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1204	A	N1-C6-N6	-12.74	110.96	118.60
34	BA	584	A	C5'-C4'-C3'	12.73	136.37	116.00
36	BC	23	G	C5'-C4'-C3'	-12.73	95.63	116.00
85	AA	471	U	C2-N3-C4	-12.72	119.36	127.00
38	BE	25	U	C2-N1-C1'	-12.72	102.44	117.70
34	BA	556	A	N1-C6-N6	12.71	126.23	118.60
36	BC	31	A	C1'-O4'-C4'	-12.71	99.73	109.90
85	AA	589	A	O4'-C1'-N9	12.71	118.37	108.20
85	AA	370	A	C5'-C4'-C3'	12.70	136.32	116.00
38	BE	174	U	C6-N1-C2	-12.69	113.39	121.00
85	AA	1684	U	O4'-C1'-N1	12.69	118.35	108.20
34	BA	842	U	C2-N3-C4	-12.69	119.39	127.00
37	BD	69	U	C2-N3-C4	-12.69	119.39	127.00
85	AA	1102	C	P-O3'-C3'	12.68	134.92	119.70
34	BA	1620	U	P-O3'-C3'	12.68	134.91	119.70
38	BE	107	U	C6-N1-C2	-12.67	113.40	121.00
35	BB	850	U	P-O5'-C5'	12.67	141.17	120.90
35	BB	1480	G	C5-C6-O6	-12.67	121.00	128.60
34	BA	539	C	C6-N1-C2	-12.66	115.24	120.30
34	BA	1468	U	C2-N3-C4	-12.65	119.41	127.00
34	BA	520	G	C5-C6-O6	-12.64	121.02	128.60
85	AA	1982	C	P-O3'-C3'	12.64	134.87	119.70
35	BB	498	G	C5-C6-O6	-12.63	121.02	128.60
85	AA	327	G	C8-N9-C1'	-12.63	110.58	127.00
34	BA	1197	U	P-O3'-C3'	12.62	134.85	119.70
34	BA	399	G	C5-C6-O6	-12.62	121.03	128.60
34	BA	297	A	P-O5'-C5'	12.62	141.09	120.90
34	BA	609	G	O4'-C1'-N9	12.61	118.29	108.20
85	AA	1361	A	N1-C6-N6	12.61	126.17	118.60
35	BB	797	C	C5'-C4'-C3'	12.61	136.18	116.00
85	AA	305	A	C5'-C4'-C3'	12.61	136.18	116.00
85	AA	1592	C	C6-N1-C2	-12.61	115.26	120.30
35	BB	62	C	P-O3'-C3'	12.60	134.82	119.70
37	BD	40	C	P-O3'-C3'	12.60	134.82	119.70
85	AA	387	U	O4'-C1'-N1	12.60	118.28	108.20
85	AA	374	C	C5'-C4'-C3'	-12.60	95.85	116.00
85	AA	814	G	C5-C6-O6	-12.60	121.04	128.60
35	BB	1023	G	P-O3'-C3'	12.59	134.81	119.70
38	BE	8	G	C4'-C3'-C2'	12.58	115.18	102.60
85	AA	1974	C	C5-C4-N4	-12.58	111.40	120.20
38	BE	198	A	C8-N9-C4	12.57	110.83	105.80
85	AA	526	G	P-O5'-C5'	12.57	141.01	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	539	A	N1-C6-N6	12.57	126.14	118.60
52	BS	69	ARG	NE-CZ-NH1	12.56	126.58	120.30
34	BA	497	U	C5'-C4'-C3'	12.56	136.10	116.00
35	BB	808	U	C6-N1-C2	-12.56	113.47	121.00
40	BG	86	U	P-O3'-C3'	-12.56	104.63	119.70
85	AA	2149	C	C6-N1-C2	-12.55	115.28	120.30
35	BB	730	G	P-O5'-C5'	12.55	140.98	120.90
35	BB	518	G	C5'-C4'-C3'	-12.55	95.92	116.00
36	BC	138	C	C6-N1-C2	-12.55	115.28	120.30
36	BC	139	A	O3'-P-O5'	12.54	127.83	104.00
34	BA	1632	G	O4'-C1'-N9	12.54	118.23	108.20
34	BA	1800	G	P-O3'-C3'	-12.54	104.65	119.70
85	AA	206	U	P-O3'-C3'	12.53	134.74	119.70
35	BB	569	G	N1-C6-O6	-12.53	112.38	119.90
35	BB	868	C	O4'-C1'-N1	12.53	118.22	108.20
41	BH	24	U	O4'-C1'-N1	12.51	118.21	108.20
34	BA	1626	U	C5'-C4'-C3'	12.50	135.99	116.00
35	BB	1215	U	C6-N1-C2	-12.50	113.50	121.00
38	BE	188	C	O4'-C1'-N1	12.50	118.20	108.20
39	BF	60	C	O4'-C1'-N1	12.50	118.20	108.20
37	BD	83	A	O3'-P-O5'	-12.49	80.26	104.00
39	BF	34	C	C5'-C4'-C3'	12.49	135.99	116.00
72	Bm	6	ARG	NE-CZ-NH1	12.49	126.55	120.30
40	BG	122	G	N1-C6-O6	12.49	127.39	119.90
41	BH	39	G	N1-C6-O6	-12.48	112.41	119.90
85	AA	2078	A	C5'-C4'-C3'	12.48	135.97	116.00
34	BA	472	G	C5-C6-O6	-12.48	121.11	128.60
34	BA	896	U	O4'-C1'-C2'	-12.48	93.32	105.80
34	BA	327	G	P-O3'-C3'	12.48	134.67	119.70
35	BB	504	C	C5'-C4'-C3'	-12.48	96.04	116.00
38	BE	45	G	C8-N9-C1'	12.48	143.22	127.00
35	BB	631	G	C5-C6-O6	-12.47	121.12	128.60
35	BB	1218	G	P-O3'-C3'	-12.46	104.74	119.70
38	BE	123	A	N1-C6-N6	12.46	126.08	118.60
85	AA	83	U	O4'-C1'-N1	12.46	118.16	108.20
85	AA	1320	G	P-O3'-C3'	12.45	134.64	119.70
35	BB	267	C	P-O3'-C3'	12.45	134.64	119.70
85	AA	403	G	C6-N1-C2	-12.45	117.63	125.10
39	BF	54	U	P-O3'-C3'	12.44	134.63	119.70
85	AA	8	U	C2-N3-C4	-12.44	119.53	127.00
85	AA	860	C	N3-C2-O2	-12.44	113.19	121.90
38	BE	37	C	C6-N1-C2	-12.44	115.33	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	2	G	N1-C6-O6	-12.43	112.44	119.90
34	BA	583	G	P-O3'-C3'	12.43	134.62	119.70
13	AE	97	ARG	NE-CZ-NH2	-12.43	114.09	120.30
85	AA	276	C	C5'-C4'-C3'	12.43	135.88	116.00
85	AA	967	C	C6-N1-C2	-12.43	115.33	120.30
34	BA	1442	A	N1-C6-N6	-12.42	111.15	118.60
35	BB	1454	G	O4'-C1'-N9	12.42	118.14	108.20
85	AA	2155	U	O4'-C1'-N1	12.42	118.13	108.20
38	BE	203	C	C5-C6-N1	12.41	127.21	121.00
36	BC	3	C	C2-N3-C4	-12.41	113.70	119.90
34	BA	186	G	P-O5'-C5'	12.41	140.75	120.90
34	BA	1699	A	C3'-C2'-C1'	-12.41	91.58	101.50
35	BB	1306	G	N1-C6-O6	-12.41	112.46	119.90
34	BA	535	G	P-O3'-C3'	12.40	134.58	119.70
38	BE	110	U	C5'-C4'-C3'	-12.40	96.16	116.00
35	BB	1202	G	C5'-C4'-C3'	-12.39	96.17	116.00
85	AA	1812	C	C6-N1-C2	-12.39	115.34	120.30
85	AA	1225	C	P-O3'-C3'	12.39	134.57	119.70
35	BB	1454	G	C5-C6-O6	-12.39	121.17	128.60
34	BA	1194	G	C4'-C3'-C2'	12.38	114.98	102.60
85	AA	926	C	P-O3'-C3'	12.38	134.56	119.70
85	AA	2077	G	N1-C6-O6	12.38	127.33	119.90
85	AA	1354	A	O3'-P-O5'	12.38	127.51	104.00
34	BA	891	C	P-O5'-C5'	12.37	140.70	120.90
29	AV	45	ARG	NE-CZ-NH1	12.37	126.48	120.30
34	BA	1541	G	C5-C6-O6	-12.37	121.18	128.60
35	BB	1329	G	N1-C6-O6	12.37	127.32	119.90
85	AA	982	G	C8-N9-C4	12.37	111.35	106.40
85	AA	2119	C	C5'-C4'-C3'	12.37	135.80	116.00
34	BA	1699	A	O4'-C4'-C3'	-12.37	91.63	104.00
35	BB	1024	G	O3'-P-O5'	12.37	127.50	104.00
38	BE	89	G	N1-C6-O6	-12.37	112.48	119.90
34	BA	855	C	C2-N3-C4	-12.36	113.72	119.90
35	BB	1329	G	C5-C6-O6	-12.36	121.18	128.60
85	AA	687	G	C5-C6-O6	-12.36	121.18	128.60
85	AA	266	U	O4'-C1'-N1	12.36	118.08	108.20
85	AA	1704	C	C2-N3-C4	-12.36	113.72	119.90
85	AA	1485	G	C8-N9-C1'	12.35	143.06	127.00
5	A4	145	TRP	CB-CG-CD1	12.35	143.05	127.00
34	BA	383	G	C8-N9-C1'	12.35	143.05	127.00
38	BE	102	U	P-O3'-C3'	12.35	134.51	119.70
38	BE	176	G	C8-N9-C4	-12.34	101.46	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	486	G	C8-N9-C1'	12.34	143.04	127.00
34	BA	678	C	C2-N1-C1'	12.33	132.36	118.80
40	BG	61	A	N1-C6-N6	12.32	126.00	118.60
85	AA	1157	U	P-O3'-C3'	12.32	134.48	119.70
35	BB	711	C	C6-N1-C2	-12.31	115.38	120.30
35	BB	1032	U	O4'-C1'-N1	12.31	118.05	108.20
36	BC	124	A	N1-C6-N6	12.31	125.99	118.60
34	BA	1489	U	P-O5'-C5'	12.30	140.59	120.90
34	BA	268	U	P-O3'-C3'	12.30	134.46	119.70
35	BB	287	C	P-O3'-C3'	12.30	134.46	119.70
85	AA	264	A	P-O5'-C5'	12.30	140.58	120.90
34	BA	1321	A	N1-C6-N6	12.29	125.98	118.60
85	AA	1052	C	P-O5'-C5'	12.29	140.57	120.90
34	BA	127	U	P-O3'-C3'	12.29	134.45	119.70
34	BA	115	U	C4'-C3'-C2'	-12.29	90.31	102.60
35	BB	845	C	P-O3'-C3'	12.29	134.45	119.70
85	AA	1325	C	P-O3'-C3'	12.29	134.44	119.70
85	AA	334	A	P-O5'-C5'	12.29	140.56	120.90
85	AA	1229	G	C8-N9-C4	-12.28	101.49	106.40
77	Br	32	ARG	NE-CZ-NH2	-12.28	114.16	120.30
34	BA	1656	A	C8-N9-C4	12.28	110.71	105.80
34	BA	270	U	O4'-C1'-N1	12.27	118.02	108.20
85	AA	989	U	P-O5'-C5'	12.27	140.53	120.90
35	BB	816	U	N3-C2-O2	-12.27	113.61	122.20
85	AA	1246	G	C5-C6-O6	-12.27	121.24	128.60
34	BA	178	C	C6-N1-C2	-12.26	115.40	120.30
86	AB	16	U	O4'-C1'-N1	12.26	118.01	108.20
35	BB	822	G	N3-C4-C5	-12.25	122.47	128.60
39	BF	22	U	C2-N1-C1'	-12.25	103.00	117.70
85	AA	736	U	P-O3'-C3'	-12.25	105.00	119.70
39	BF	10	A	O3'-P-O5'	12.25	127.28	104.00
41	BH	75	G	N1-C6-O6	12.25	127.25	119.90
35	BB	5	A	N1-C6-N6	-12.25	111.25	118.60
85	AA	2154	C	O4'-C1'-N1	12.25	118.00	108.20
35	BB	68	G	N1-C6-O6	12.24	127.24	119.90
35	BB	475	A	N1-C6-N6	-12.24	111.25	118.60
34	BA	1820	G	C5-C6-O6	-12.24	121.26	128.60
85	AA	208	U	P-O3'-C3'	12.24	134.38	119.70
36	BC	60	U	P-O3'-C3'	12.23	134.38	119.70
37	BD	82	G	C4'-C3'-C2'	12.23	114.83	102.60
85	AA	1964	A	C1'-O4'-C4'	-12.23	100.11	109.90
85	AA	8	U	C6-N1-C2	-12.23	113.66	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1097	G	P-O3'-C3'	12.22	134.37	119.70
40	BG	146	C	C6-N1-C2	-12.22	115.41	120.30
34	BA	1571	C	C6-N1-C2	-12.22	115.41	120.30
35	BB	653	G	C5-C6-O6	-12.22	121.27	128.60
85	AA	769	C	C2-N1-C1'	-12.21	105.36	118.80
34	BA	18	G	C5-C6-O6	-12.21	121.27	128.60
6	A5	197	ARG	NE-CZ-NH1	12.21	126.41	120.30
35	BB	798	A	C5-C6-N6	-12.21	113.94	123.70
34	BA	882	G	C5-C6-O6	-12.20	121.28	128.60
35	BB	869	G	O4'-C1'-N9	12.20	117.96	108.20
34	BA	1711	G	C1'-O4'-C4'	-12.20	100.14	109.90
34	BA	8	G	C8-N9-C1'	12.20	142.86	127.00
34	BA	1295	U	C2-N1-C1'	-12.20	103.06	117.70
35	BB	797	C	O4'-C1'-N1	12.19	117.95	108.20
34	BA	1491	U	O4'-C1'-N1	12.19	117.95	108.20
34	BA	880	G	C5-C6-O6	-12.19	121.29	128.60
34	BA	1549	U	C5'-C4'-C3'	-12.19	96.50	116.00
85	AA	1974	C	N3-C4-N4	12.19	126.53	118.00
35	BB	1091	C	C6-N1-C2	-12.18	115.43	120.30
38	BE	36	U	N3-C2-O2	-12.18	113.67	122.20
34	BA	1809	G	P-O5'-C5'	12.18	140.39	120.90
35	BB	692	G	N1-C6-O6	-12.18	112.59	119.90
34	BA	559	C	C6-N1-C2	-12.17	115.43	120.30
34	BA	603	U	C4'-C3'-C2'	-12.17	90.43	102.60
35	BB	139	G	O4'-C1'-N9	12.17	117.94	108.20
35	BB	622	G	N1-C6-O6	12.17	127.20	119.90
85	AA	1897	A	N1-C6-N6	-12.17	111.30	118.60
85	AA	860	C	P-O3'-C3'	12.17	134.30	119.70
34	BA	1293	A	C6-N1-C2	-12.16	111.30	118.60
38	BE	77	C	C6-N1-C2	-12.16	115.44	120.30
4	A3	7	TYR	CB-CG-CD2	-12.16	113.70	121.00
34	BA	869	C	O4'-C1'-N1	12.16	117.93	108.20
85	AA	619	A	O4'-C1'-N9	12.16	117.93	108.20
35	BB	1294	C	C6-N1-C1'	-12.15	106.21	120.80
85	AA	815	G	O4'-C1'-N9	12.15	117.92	108.20
35	BB	768	A	P-O3'-C3'	12.15	134.28	119.70
85	AA	1179	A	P-O3'-C3'	12.15	134.28	119.70
34	BA	1033	G	C5-C6-O6	-12.14	121.31	128.60
34	BA	635	G	P-O5'-C5'	12.14	140.32	120.90
85	AA	2127	G	N1-C6-O6	12.14	127.18	119.90
85	AA	864	C	O4'-C1'-N1	12.14	117.91	108.20
34	BA	1674	G	C1'-O4'-C4'	-12.12	100.20	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	976	U	C2-N3-C4	-12.12	119.73	127.00
41	BH	63	G	C8-N9-C4	-12.12	101.55	106.40
85	AA	1457	C	P-O3'-C3'	12.12	134.24	119.70
35	BB	877	A	C4'-C3'-C2'	12.11	114.71	102.60
85	AA	1197	U	C5'-C4'-C3'	-12.11	96.63	116.00
37	BD	77	A	P-O5'-C5'	12.11	140.27	120.90
85	AA	1535	C	O3'-P-O5'	12.10	127.00	104.00
35	BB	1140	C	P-O3'-C3'	12.10	134.22	119.70
85	AA	307	G	O4'-C1'-N9	12.10	117.88	108.20
34	BA	472	G	N1-C6-O6	12.10	127.16	119.90
34	BA	871	G	O5'-P-OP1	-12.09	94.82	105.70
85	AA	1292	A	C5-C6-N6	-12.09	114.03	123.70
34	BA	94	G	P-O5'-C5'	-12.09	101.56	120.90
39	BF	23	G	O5'-C5'-C4'	12.09	134.67	111.70
85	AA	414	C	C6-N1-C1'	12.09	135.31	120.80
85	AA	1719	C	P-O5'-C5'	12.09	140.24	120.90
85	AA	696	G	C5-C6-O6	-12.09	121.35	128.60
85	AA	2121	G	P-O3'-C3'	12.09	134.20	119.70
85	AA	47	A	C5'-C4'-C3'	-12.08	96.67	116.00
85	AA	1010	U	C6-N1-C2	-12.08	113.75	121.00
34	BA	1505	G	N9-C4-C5	-12.08	100.57	105.40
35	BB	315	C	C6-N1-C2	-12.08	115.47	120.30
85	AA	1921	G	C4-N9-C1'	-12.08	110.80	126.50
34	BA	128	C	C6-N1-C1'	12.07	135.29	120.80
85	AA	445	U	C1'-O4'-C4'	-12.07	100.25	109.90
85	AA	1123	C	C6-N1-C2	-12.07	115.47	120.30
85	AA	1699	A	P-O5'-C5'	12.07	140.21	120.90
34	BA	328	A	P-O3'-C3'	-12.06	105.22	119.70
34	BA	615	A	C5'-C4'-C3'	-12.06	96.70	116.00
85	AA	603	C	C6-N1-C2	-12.06	115.48	120.30
26	AS	131	TYR	CB-CG-CD2	-12.05	113.77	121.00
34	BA	1616	A	P-O3'-C3'	12.05	134.16	119.70
35	BB	50	A	C8-N9-C4	12.05	110.62	105.80
35	BB	712	U	P-O3'-C3'	12.05	134.16	119.70
34	BA	802	G	O4'-C1'-N9	12.04	117.84	108.20
41	BH	110	C	C6-N1-C2	-12.04	115.48	120.30
85	AA	61	C	C6-N1-C2	-12.04	115.48	120.30
85	AA	28	A	C5'-C4'-O4'	-12.04	94.66	109.10
34	BA	878	G	C5-C6-O6	-12.03	121.38	128.60
34	BA	1412	G	N1-C6-O6	-12.03	112.68	119.90
35	BB	1494	G	C8-N9-C1'	12.03	142.64	127.00
85	AA	1430	A	P-O5'-C5'	12.02	140.14	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	8	U	C2-N3-C4	-12.02	119.79	127.00
85	AA	284	C	O4'-C1'-N1	12.02	117.82	108.20
29	AV	100	TYR	CB-CG-CD2	-12.02	113.79	121.00
37	BD	49	A	C5'-C4'-C3'	-12.02	96.77	116.00
85	AA	1485	G	C4-N9-C1'	-12.01	110.88	126.50
35	BB	997	G	C5-C6-O6	-12.01	121.39	128.60
37	BD	84	U	P-O5'-C5'	12.01	140.12	120.90
34	BA	205	G	N1-C6-O6	12.01	127.11	119.90
38	BE	20	C	C5'-C4'-O4'	12.01	123.51	109.10
38	BE	149	A	C2-N3-C4	-12.01	104.60	110.60
35	BB	850	U	P-O3'-C3'	12.00	134.10	119.70
85	AA	1828	C	P-O5'-C5'	12.00	140.10	120.90
85	AA	2074	G	C5-C6-O6	-12.00	121.40	128.60
85	AA	1796	C	O4'-C1'-N1	12.00	117.80	108.20
35	BB	436	G	N1-C6-O6	11.99	127.09	119.90
34	BA	593	G	C5'-C4'-O4'	11.99	123.49	109.10
85	AA	535	G	C5-C6-O6	-11.99	121.41	128.60
85	AA	920	A	P-O5'-C5'	11.99	140.08	120.90
40	BG	76	C	C2-N3-C4	-11.98	113.91	119.90
85	AA	696	G	C4-N9-C1'	-11.98	110.92	126.50
85	AA	1730	C	P-O3'-C3'	11.98	134.07	119.70
38	BE	129	G	O4'-C1'-N9	11.98	117.78	108.20
34	BA	672	G	P-O3'-C3'	11.97	134.07	119.70
85	AA	770	C	C6-N1-C2	-11.97	115.51	120.30
85	AA	741	G	P-O3'-C3'	-11.97	105.34	119.70
34	BA	319	C	C6-N1-C2	-11.97	115.51	120.30
34	BA	363	G	C5-C6-O6	-11.96	121.42	128.60
34	BA	382	G	N1-C6-O6	-11.96	112.72	119.90
85	AA	599	C	O3'-P-O5'	11.96	126.73	104.00
85	AA	1053	A	P-O3'-C3'	11.96	134.06	119.70
85	AA	1372	C	P-O3'-C3'	-11.96	105.35	119.70
85	AA	2145	G	O4'-C1'-N9	11.96	117.77	108.20
40	BG	171	A	C8-N9-C4	-11.95	101.02	105.80
34	BA	520	G	N1-C6-O6	11.95	127.07	119.90
34	BA	817	U	C5'-C4'-C3'	-11.95	96.88	116.00
34	BA	993	C	O4'-C1'-N1	11.95	117.76	108.20
85	AA	525	C	O3'-P-O5'	-11.95	81.29	104.00
34	BA	21	C	P-O5'-C5'	11.95	140.01	120.90
85	AA	1784	G	P-O3'-C3'	11.95	134.03	119.70
35	BB	1480	G	C8-N9-C1'	11.94	142.52	127.00
38	BE	144	A	P-O5'-C5'	11.94	140.00	120.90
38	BE	176	G	C6-N1-C2	-11.94	117.94	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1490	A	C6-C5-N7	-11.94	123.94	132.30
39	BF	50	C	C2-N1-C1'	-11.94	105.67	118.80
35	BB	816	U	C2-N1-C1'	-11.93	103.38	117.70
52	BS	137	ARG	NE-CZ-NH1	11.93	126.27	120.30
34	BA	1537	G	P-O5'-C5'	11.93	139.99	120.90
54	BU	12	ARG	NE-CZ-NH1	11.93	126.27	120.30
39	BF	10	A	P-O3'-C3'	-11.93	105.39	119.70
34	BA	804	G	C5-C6-O6	11.93	135.76	128.60
85	AA	1492	U	O4'-C1'-N1	11.92	117.74	108.20
34	BA	874	G	N1-C6-O6	11.92	127.05	119.90
85	AA	461	G	C5-C6-O6	-11.92	121.45	128.60
40	BG	33	G	N1-C6-O6	11.91	127.05	119.90
40	BG	67	A	O4'-C1'-N9	11.91	117.73	108.20
85	AA	992	G	P-O5'-C5'	-11.91	101.85	120.90
41	BH	26	C	O4'-C1'-N1	11.90	117.72	108.20
34	BA	683	C	C2-N1-C1'	11.90	131.89	118.80
34	BA	1041	U	P-O3'-C3'	11.90	133.98	119.70
50	BQ	55	ARG	NE-CZ-NH1	11.90	126.25	120.30
34	BA	201	A	N1-C6-N6	-11.89	111.46	118.60
34	BA	1648	G	C5'-C4'-C3'	-11.89	96.97	116.00
85	AA	1577	G	P-O3'-C3'	11.89	133.97	119.70
34	BA	1191	C	O4'-C1'-N1	11.89	117.71	108.20
34	BA	113	G	C5'-C4'-C3'	11.89	135.02	116.00
85	AA	863	C	O4'-C1'-N1	11.88	117.71	108.20
35	BB	436	G	C5-C6-O6	-11.88	121.47	128.60
85	AA	1485	G	C5-C6-O6	-11.88	121.47	128.60
35	BB	1540	U	P-O3'-C3'	-11.88	105.45	119.70
85	AA	991	G	N1-C6-O6	11.88	127.03	119.90
34	BA	1442	A	P-O3'-C3'	11.88	133.95	119.70
34	BA	1658	G	N1-C6-O6	-11.88	112.77	119.90
85	AA	443	A	O3'-P-O5'	-11.88	81.44	104.00
34	BA	691	A	N1-C6-N6	-11.87	111.48	118.60
85	AA	1226	A	P-O5'-C5'	11.87	139.90	120.90
34	BA	692	U	O4'-C1'-N1	-11.87	98.71	108.20
35	BB	798	A	C5'-C4'-C3'	11.86	134.98	116.00
38	BE	87	U	C6-N1-C2	-11.86	113.89	121.00
39	BF	20	U	P-O3'-C3'	11.86	133.93	119.70
85	AA	696	G	C8-N9-C1'	11.86	142.41	127.00
34	BA	1641	G	C4-N9-C1'	-11.85	111.09	126.50
35	BB	1294	C	C2-N1-C1'	11.85	131.84	118.80
35	BB	1515	C	C6-N1-C1'	11.85	135.02	120.80
85	AA	1236	G	C6-N1-C2	-11.84	118.00	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	875	G	C8-N9-C4	11.84	111.14	106.40
38	BE	171	U	C5'-C4'-C3'	-11.84	97.06	116.00
85	AA	890	U	N1-C2-O2	-11.84	114.52	122.80
35	BB	1517	G	C6-C5-N7	-11.83	123.30	130.40
36	BC	155	C	O4'-C1'-N1	11.83	117.66	108.20
40	BG	25	G	C5'-C4'-C3'	-11.82	97.08	116.00
34	BA	504	A	N1-C6-N6	-11.81	111.51	118.60
35	BB	621	C	O4'-C1'-N1	11.81	117.65	108.20
35	BB	1327	U	C2-N3-C4	-11.81	119.91	127.00
38	BE	8	G	N9-C1'-C2'	-11.81	98.64	114.00
38	BE	25	U	N3-C2-O2	-11.81	113.93	122.20
85	AA	2061	C	P-O3'-C3'	11.81	133.87	119.70
85	AA	2244	G	N1-C6-O6	11.81	126.98	119.90
35	BB	713	U	P-O3'-C3'	11.80	133.87	119.70
35	BB	830	G	P-O3'-C3'	-11.80	105.54	119.70
85	AA	1909	C	O4'-C1'-N1	11.79	117.64	108.20
36	BC	13	U	C2-N1-C1'	-11.79	103.55	117.70
40	BG	64	C	P-O5'-C5'	11.79	139.77	120.90
85	AA	1035	C	C6-N1-C2	-11.79	115.58	120.30
34	BA	155	U	O4'-C1'-N1	11.79	117.63	108.20
85	AA	1457	C	P-O5'-C5'	11.79	139.76	120.90
85	AA	2119	C	C6-N1-C2	-11.79	115.58	120.30
35	BB	56	U	C6-N1-C2	-11.79	113.93	121.00
36	BC	157	U	C5'-C4'-C3'	11.78	134.84	116.00
35	BB	958	C	C5'-C4'-C3'	11.78	134.84	116.00
34	BA	687	G	N1-C6-O6	-11.77	112.84	119.90
38	BE	30	C	C6-N1-C1'	11.77	134.93	120.80
85	AA	1378	U	P-O3'-C3'	11.77	133.82	119.70
35	BB	3	C	P-O3'-C3'	11.77	133.82	119.70
85	AA	1471	G	C5-C6-O6	-11.77	121.54	128.60
85	AA	1455	C	C6-N1-C2	-11.76	115.59	120.30
24	AQ	63	ARG	NE-CZ-NH1	11.76	126.18	120.30
85	AA	753	U	P-O3'-C3'	11.76	133.81	119.70
35	BB	1177	U	O4'-C1'-N1	11.76	117.61	108.20
37	BD	48	G	P-O3'-C3'	11.76	133.81	119.70
35	BB	1026	G	P-O3'-C3'	-11.75	105.60	119.70
47	BN	52	PHE	CB-CG-CD2	-11.75	112.58	120.80
47	BN	206	ARG	NE-CZ-NH1	11.75	126.17	120.30
34	BA	210	G	C5'-C4'-C3'	11.74	134.79	116.00
39	BF	11	C	O4'-C1'-N1	11.74	117.59	108.20
34	BA	221	G	C4-N9-C1'	-11.74	111.24	126.50
34	BA	521	C	P-O3'-C3'	-11.74	105.62	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1736	A	C8-N9-C4	-11.74	101.11	105.80
85	AA	769	C	P-O3'-C3'	11.74	133.78	119.70
77	Br	32	ARG	NE-CZ-NH1	11.74	126.17	120.30
85	AA	1466	U	O4'-C1'-N1	-11.73	98.81	108.20
85	AA	1535	C	C2-N3-C4	-11.73	114.03	119.90
85	AA	1584	U	P-O3'-C3'	11.73	133.78	119.70
35	BB	148	C	C2-N1-C1'	11.73	131.70	118.80
85	AA	330	C	P-O3'-C3'	11.73	133.78	119.70
85	AA	1211	C	C2-N3-C4	-11.73	114.04	119.90
1	A0	83	TYR	CB-CG-CD2	-11.72	113.97	121.00
38	BE	30	C	C2-N3-C4	-11.72	114.04	119.90
85	AA	53	G	C8-N9-C1'	11.72	142.23	127.00
34	BA	959	G	C5-C6-O6	-11.72	121.57	128.60
35	BB	1511	U	C5'-C4'-C3'	11.72	134.75	116.00
34	BA	1820	G	N1-C6-O6	11.71	126.93	119.90
34	BA	109	A	N1-C6-N6	-11.71	111.57	118.60
36	BC	38	U	C1'-O4'-C4'	-11.71	100.53	109.90
85	AA	123	A	P-O3'-C3'	11.71	133.75	119.70
85	AA	653	A	P-O3'-C3'	11.71	133.75	119.70
85	AA	270	A	O4'-C1'-N9	11.71	117.56	108.20
34	BA	917	C	C6-N1-C2	-11.70	115.62	120.30
34	BA	538	G	P-O3'-C3'	11.70	133.74	119.70
34	BA	1440	C	O4'-C1'-N1	11.70	117.56	108.20
34	BA	1735	G	O4'-C1'-N9	11.70	117.56	108.20
34	BA	221	G	C8-N9-C1'	11.69	142.20	127.00
34	BA	250	G	P-O3'-C3'	-11.69	105.67	119.70
34	BA	1497	A	P-O3'-C3'	11.69	133.72	119.70
34	BA	897	U	P-O3'-C3'	11.69	133.72	119.70
38	BE	10	G	O4'-C1'-N9	11.69	117.55	108.20
41	BH	134	U	O4'-C1'-N1	11.69	117.55	108.20
85	AA	1458	G	C8-N9-C1'	11.69	142.19	127.00
34	BA	923	C	C6-N1-C2	-11.68	115.63	120.30
85	AA	133	G	C8-N9-C1'	11.68	142.19	127.00
34	BA	875	G	C4-N9-C1'	-11.68	111.31	126.50
35	BB	6	A	P-O5'-C5'	11.68	139.59	120.90
86	AB	7	A	N1-C6-N6	11.68	125.61	118.60
41	BH	135	U	P-O5'-C5'	11.67	139.58	120.90
85	AA	114	C	C5'-C4'-C3'	-11.67	97.32	116.00
85	AA	1578	G	O4'-C1'-N9	11.67	117.53	108.20
34	BA	593	G	P-O3'-C3'	11.66	133.70	119.70
34	BA	1708	A	O4'-C1'-N9	11.66	117.53	108.20
85	AA	730	G	C5-C6-O6	-11.66	121.60	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	57	A	C5-C6-N6	-11.66	114.37	123.70
39	BF	66	C	C6-N1-C2	-11.66	115.64	120.30
38	BE	20	C	P-O3'-C3'	-11.66	105.71	119.70
85	AA	1881	C	C6-N1-C2	-11.66	115.64	120.30
34	BA	383	G	C5'-C4'-C3'	-11.65	97.36	116.00
34	BA	1084	A	N1-C6-N6	-11.65	111.61	118.60
38	BE	1	U	N3-C2-O2	-11.65	114.04	122.20
34	BA	1294	C	C4'-C3'-C2'	-11.65	90.95	102.60
85	AA	626	G	C5-C6-O6	-11.65	121.61	128.60
85	AA	2060	G	P-O5'-C5'	11.64	139.53	120.90
34	BA	1085	G	O4'-C1'-N9	11.64	117.51	108.20
34	BA	286	C	O4'-C1'-N1	11.63	117.51	108.20
35	BB	1166	A	P-O3'-C3'	11.63	133.66	119.70
35	BB	1323	U	C2-N3-C4	-11.63	120.02	127.00
37	BD	73	U	P-O3'-C3'	11.63	133.66	119.70
85	AA	1226	A	O4'-C1'-N9	11.63	117.50	108.20
85	AA	1959	G	N1-C6-O6	11.63	126.88	119.90
34	BA	1725	U	C1'-O4'-C4'	-11.62	100.60	109.90
41	BH	39	G	P-O5'-C5'	11.62	139.50	120.90
85	AA	611	G	N1-C6-O6	11.62	126.88	119.90
39	BF	13	U	O4'-C1'-N1	11.62	117.50	108.20
34	BA	1809	G	C5-C6-O6	-11.62	121.63	128.60
82	Bw	41	PHE	CB-CG-CD1	-11.61	112.67	120.80
85	AA	778	C	C6-N1-C2	-11.61	115.65	120.30
34	BA	117	C	C1'-O4'-C4'	-11.61	100.61	109.90
85	AA	971	U	O4'-C1'-N1	11.61	117.49	108.20
85	AA	1051	A	P-O3'-C3'	11.61	133.63	119.70
44	BK	98	ARG	NE-CZ-NH1	11.61	126.10	120.30
85	AA	1260	G	O4'-C1'-N9	11.61	117.48	108.20
34	BA	631	G	C8-N9-C1'	11.60	142.08	127.00
34	BA	768	G	N1-C6-O6	11.60	126.86	119.90
34	BA	1509	U	C5'-C4'-C3'	11.60	134.56	116.00
36	BC	140	U	C6-N1-C2	-11.60	114.04	121.00
27	AT	83	TYR	CB-CG-CD2	-11.60	114.04	121.00
86	AB	14	A	N1-C6-N6	-11.60	111.64	118.60
34	BA	651	U	O4'-C1'-N1	11.59	117.47	108.20
34	BA	1592	U	C2-N3-C4	-11.59	120.04	127.00
40	BG	164	U	C2-N3-C4	-11.59	120.05	127.00
36	BC	140	U	P-O5'-C5'	11.59	139.44	120.90
35	BB	560	C	C4'-C3'-C2'	11.59	114.19	102.60
85	AA	790	A	P-O3'-C3'	-11.59	105.80	119.70
34	BA	1741	G	O4'-C1'-N9	11.58	117.46	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	10	G	O4'-C1'-C2'	-11.58	94.22	105.80
35	BB	1231	U	O4'-C1'-N1	11.58	117.46	108.20
38	BE	85	G	C8-N9-C1'	11.58	142.05	127.00
40	BG	22	G	O3'-P-O5'	-11.58	82.00	104.00
85	AA	1133	C	C5'-C4'-C3'	-11.57	97.48	116.00
59	BZ	37	ARG	NE-CZ-NH1	11.57	126.08	120.30
85	AA	633	C	O4'-C1'-N1	11.57	117.45	108.20
85	AA	588	G	C5-C6-O6	-11.56	121.66	128.60
85	AA	1441	G	C6-N1-C2	-11.56	118.16	125.10
85	AA	726	U	P-O5'-C5'	11.56	139.40	120.90
38	BE	168	C	C6-N1-C2	-11.56	115.67	120.30
85	AA	929	G	C5-C6-O6	-11.56	121.66	128.60
34	BA	523	A	N1-C6-N6	-11.56	111.66	118.60
39	BF	32	G	C5-C6-O6	-11.56	121.67	128.60
34	BA	1321	A	C5-C6-N6	-11.56	114.45	123.70
36	BC	146	U	P-O3'-C3'	11.56	133.57	119.70
85	AA	1458	G	C4-N9-C1'	-11.56	111.47	126.50
34	BA	487	A	C5-C6-N6	11.56	132.94	123.70
86	AB	3	C	C1'-O4'-C4'	-11.55	100.66	109.90
34	BA	507	U	C2-N3-C4	-11.55	120.07	127.00
85	AA	1158	U	C1'-O4'-C4'	-11.55	100.66	109.90
85	AA	2244	G	C5-C6-O6	-11.55	121.67	128.60
85	AA	368	C	C1'-O4'-C4'	-11.55	100.66	109.90
35	BB	1375	G	N1-C6-O6	-11.54	112.97	119.90
85	AA	83	U	P-O3'-C3'	11.55	133.56	119.70
39	BF	7	G	C4-N9-C1'	11.54	141.51	126.50
35	BB	964	G	C8-N9-C1'	11.53	141.99	127.00
35	BB	1038	G	C5-C6-O6	-11.53	121.68	128.60
41	BH	72	G	O5'-C5'-C4'	11.53	133.60	111.70
85	AA	1927	G	P-O3'-C3'	11.53	133.53	119.70
34	BA	1451	A	C4-C5-C6	-11.53	111.24	117.00
34	BA	1597	G	C8-N9-C4	11.53	111.01	106.40
85	AA	1985	C	C5'-C4'-C3'	-11.53	97.56	116.00
37	BD	41	G	C4-N9-C1'	-11.52	111.52	126.50
34	BA	282	A	OP1-P-OP2	-11.52	102.33	119.60
85	AA	264	A	N1-C6-N6	11.52	125.51	118.60
34	BA	740	A	O4'-C1'-N9	11.51	117.41	108.20
35	BB	1333	U	O4'-C1'-N1	11.51	117.41	108.20
37	BD	48	G	C6-N1-C2	-11.51	118.19	125.10
34	BA	993	C	C6-N1-C2	-11.51	115.70	120.30
35	BB	792	G	P-O3'-C3'	11.51	133.51	119.70
35	BB	362	A	N1-C6-N6	11.51	125.50	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	870	U	P-O5'-C5'	11.51	139.31	120.90
85	AA	1553	G	C5'-C4'-C3'	-11.50	97.60	116.00
34	BA	167	U	C2-N1-C1'	-11.50	103.90	117.70
38	BE	30	C	N3-C4-N4	-11.50	109.95	118.00
41	BH	84	A	N1-C2-N3	-11.50	123.55	129.30
34	BA	128	C	C2-N1-C1'	-11.50	106.15	118.80
39	BF	13	U	C2-N3-C4	-11.50	120.10	127.00
85	AA	1978	G	C8-N9-C4	11.49	111.00	106.40
34	BA	1606	A	O4'-C1'-N9	11.49	117.39	108.20
38	BE	129	G	C5'-C4'-C3'	-11.49	97.61	116.00
85	AA	1458	G	O4'-C1'-N9	11.49	117.39	108.20
35	BB	979	G	C5-C6-O6	-11.49	121.71	128.60
85	AA	743	C	O4'-C1'-N1	11.49	117.39	108.20
85	AA	518	A	P-O5'-C5'	11.48	139.28	120.90
34	BA	1846	G	C5-C6-N1	11.48	117.24	111.50
85	AA	207	G	C1'-O4'-C4'	-11.48	100.72	109.90
35	BB	635	A	P-O3'-C3'	-11.48	105.92	119.70
37	BD	74	A	N1-C6-N6	11.48	125.49	118.60
34	BA	331	G	O4'-C1'-N9	11.47	117.38	108.20
34	BA	1721	U	O4'-C1'-C2'	-11.47	94.33	105.80
35	BB	1510	G	C5'-C4'-C3'	-11.47	97.64	116.00
85	AA	262	G	P-O3'-C3'	11.47	133.47	119.70
35	BB	1533	U	C6-N1-C2	-11.47	114.12	121.00
37	BD	29	C	C2-N3-C4	-11.47	114.17	119.90
35	BB	384	A	O4'-C1'-N9	11.47	117.38	108.20
85	AA	35	U	C5'-C4'-C3'	-11.47	97.65	116.00
35	BB	526	A	N1-C6-N6	11.46	125.48	118.60
38	BE	203	C	C2-N3-C4	-11.46	114.17	119.90
85	AA	802	A	P-O5'-C5'	11.46	139.24	120.90
34	BA	414	A	P-O3'-C3'	-11.46	105.95	119.70
34	BA	597	C	C6-N1-C2	-11.46	115.72	120.30
41	BH	103	C	O4'-C1'-N1	11.46	117.36	108.20
44	BK	181	TYR	CB-CG-CD2	-11.46	114.13	121.00
34	BA	1454	G	N1-C6-O6	11.45	126.77	119.90
62	Bc	107	TYR	CB-CG-CD2	-11.45	114.13	121.00
34	BA	1299	G	C5-C6-N1	11.45	117.22	111.50
34	BA	1454	G	P-O5'-C5'	11.45	139.22	120.90
34	BA	1747	C	C6-N1-C1'	11.45	134.53	120.80
34	BA	1834	A	N1-C6-N6	-11.45	111.73	118.60
37	BD	83	A	P-O5'-C5'	11.44	139.21	120.90
34	BA	772	G	O4'-C1'-N9	11.44	117.35	108.20
35	BB	993	A	C8-N9-C4	11.44	110.38	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	1	G	N1-C6-O6	11.44	126.77	119.90
1	A0	83	TYR	CB-CG-CD1	11.44	127.86	121.00
85	AA	1801	U	P-O3'-C3'	11.44	133.43	119.70
34	BA	250	G	C5'-C4'-C3'	-11.44	97.70	116.00
34	BA	681	G	C4'-C3'-C2'	-11.44	91.16	102.60
85	AA	970	U	C5'-C4'-C3'	-11.44	97.70	116.00
5	A4	145	TRP	CB-CG-CD2	-11.43	111.74	126.60
34	BA	289	A	C6-C5-N7	-11.43	124.30	132.30
34	BA	684	G	O5'-C5'-C4'	11.43	133.42	111.70
34	BA	449	G	C8-N9-C1'	11.43	141.86	127.00
34	BA	1808	A	C5'-C4'-C3'	-11.43	97.72	116.00
38	BE	77	C	P-O3'-C3'	11.43	133.41	119.70
34	BA	629	G	O5'-P-OP1	-11.43	95.42	105.70
36	BC	134	G	P-O5'-C5'	11.43	139.18	120.90
53	BT	62	ARG	NE-CZ-NH1	11.42	126.01	120.30
34	BA	742	C	C5'-C4'-C3'	-11.42	97.73	116.00
85	AA	568	C	O4'-C1'-N1	11.42	117.33	108.20
34	BA	1609	U	C2-N1-C1'	-11.41	104.01	117.70
40	BG	2	U	C5'-C4'-C3'	-11.41	97.75	116.00
34	BA	1312	A	P-O3'-C3'	-11.41	106.01	119.70
34	BA	1430	C	C6-N1-C2	-11.41	115.74	120.30
35	BB	765	G	O3'-P-O5'	-11.41	82.33	104.00
85	AA	1844	A	N1-C6-N6	-11.41	111.76	118.60
34	BA	1240	G	C5-C6-O6	-11.40	121.76	128.60
40	BG	126	G	C5-C6-O6	-11.40	121.76	128.60
85	AA	520	A	O5'-C5'-C4'	11.40	133.36	111.70
85	AA	1933	G	O4'-C1'-N9	11.40	117.32	108.20
37	BD	75	G	N3-C2-N2	-11.39	111.92	119.90
34	BA	112	C	P-O3'-C3'	-11.39	106.03	119.70
38	BE	206	G	P-O3'-C3'	11.39	133.37	119.70
85	AA	1093	C	O4'-C1'-N1	11.39	117.31	108.20
39	BF	23	G	C5'-C4'-C3'	11.39	134.22	116.00
85	AA	1351	U	P-O3'-C3'	11.39	133.37	119.70
34	BA	608	G	P-O3'-C3'	-11.38	106.04	119.70
34	BA	219	U	O4'-C1'-N1	11.38	117.30	108.20
22	AO	89	TYR	CB-CG-CD1	-11.38	114.17	121.00
85	AA	927	A	C5-C6-N6	-11.38	114.60	123.70
85	AA	389	A	N1-C6-N6	-11.37	111.78	118.60
34	BA	15	G	C5-C6-O6	-11.37	121.78	128.60
35	BB	138	A	O4'-C1'-N9	11.37	117.29	108.20
34	BA	472	G	C8-N9-C1'	-11.36	112.23	127.00
35	BB	1226	G	C8-N9-C4	11.36	110.94	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	72	G	C5'-C4'-C3'	11.36	134.18	116.00
39	BF	34	C	P-O5'-C5'	11.36	139.08	120.90
34	BA	1131	G	P-O3'-C3'	11.36	133.33	119.70
20	AL	29	TYR	CB-CG-CD2	-11.35	114.19	121.00
34	BA	1225	A	C5-C6-N6	11.35	132.78	123.70
34	BA	89	G	C5-C6-O6	-11.35	121.79	128.60
85	AA	1994	G	C5'-C4'-C3'	11.35	134.16	116.00
34	BA	1694	C	C6-N1-C2	-11.35	115.76	120.30
34	BA	813	C	N3-C2-O2	-11.34	113.96	121.90
35	BB	1390	U	C2-N3-C4	-11.34	120.20	127.00
38	BE	106	C	O4'-C1'-N1	11.34	117.27	108.20
35	BB	1134	G	C5-C6-O6	-11.34	121.80	128.60
35	BB	1102	U	C1'-O4'-C4'	-11.33	100.84	109.90
85	AA	207	G	C4-N9-C1'	11.33	141.23	126.50
85	AA	264	A	C5-C6-N6	-11.33	114.64	123.70
34	BA	702	G	C4'-C3'-C2'	11.32	113.92	102.60
32	AY	33	ARG	NE-CZ-NH1	11.32	125.96	120.30
34	BA	383	G	C5-C6-O6	-11.32	121.81	128.60
35	BB	426	A	P-O3'-C3'	11.32	133.28	119.70
29	AV	45	ARG	NE-CZ-NH2	-11.31	114.64	120.30
85	AA	403	G	P-O3'-C3'	11.31	133.28	119.70
34	BA	547	C	P-O3'-C3'	11.31	133.27	119.70
35	BB	880	G	C8-N9-C4	-11.31	101.88	106.40
85	AA	682	C	O4'-C1'-N1	11.31	117.25	108.20
34	BA	488	C	C2-N1-C1'	11.30	131.24	118.80
38	BE	125	C	C6-N1-C2	-11.30	115.78	120.30
40	BG	118	U	C2-N1-C1'	-11.30	104.14	117.70
34	BA	136	A	C8-N9-C1'	11.30	148.04	127.70
38	BE	186	C	C1'-O4'-C4'	-11.30	100.86	109.90
34	BA	1284	G	N3-C2-N2	11.30	127.81	119.90
34	BA	756	A	N1-C6-N6	11.29	125.37	118.60
34	BA	1213	A	C8-N9-C4	11.29	110.31	105.80
34	BA	1215	U	C2-N1-C1'	-11.29	104.16	117.70
34	BA	548	G	P-O3'-C3'	11.28	133.24	119.70
41	BH	29	G	N1-C6-O6	-11.28	113.13	119.90
85	AA	1453	U	C2-N3-C4	-11.28	120.23	127.00
85	AA	92	G	P-O5'-C5'	11.28	138.95	120.90
35	BB	126	C	C6-N1-C2	-11.28	115.79	120.30
85	AA	773	G	N1-C6-O6	11.28	126.67	119.90
85	AA	1719	C	O4'-C1'-N1	11.28	117.22	108.20
17	AI	104	TYR	CB-CG-CD1	11.27	127.76	121.00
34	BA	383	G	C4-N9-C1'	-11.27	111.85	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	204	U	C2-N1-C1'	-11.27	104.18	117.70
35	BB	561	C	C6-N1-C1'	-11.26	107.29	120.80
85	AA	2215	C	O3'-P-O5'	11.26	125.40	104.00
5	A4	15	ARG	NE-CZ-NH1	11.26	125.93	120.30
34	BA	1069	U	C4'-C3'-C2'	11.26	113.86	102.60
34	BA	429	G	C4-C5-C6	-11.26	112.05	118.80
37	BD	104	C	C6-N1-C2	-11.26	115.80	120.30
35	BB	574	G	P-O3'-C3'	-11.26	106.19	119.70
37	BD	49	A	N1-C6-N6	-11.26	111.85	118.60
34	BA	763	U	C4'-C3'-C2'	-11.25	91.35	102.60
34	BA	1166	A	C4'-C3'-C2'	11.25	113.85	102.60
85	AA	2140	U	C6-N1-C2	-11.25	114.25	121.00
85	AA	120	C	C2-N1-C1'	11.24	131.17	118.80
38	BE	195	G	C5'-C4'-C3'	-11.24	98.01	116.00
41	BH	4	U	P-O5'-C5'	11.24	138.88	120.90
35	BB	836	U	C6-N1-C2	-11.24	114.26	121.00
85	AA	478	U	C2-N3-C4	-11.24	120.26	127.00
34	BA	1622	U	O4'-C1'-N1	11.24	117.19	108.20
34	BA	679	U	C5'-C4'-C3'	11.23	133.98	116.00
85	AA	1210	U	C2-N3-C4	-11.23	120.26	127.00
34	BA	481	A	C2-N3-C4	-11.23	104.99	110.60
41	BH	101	A	O5'-P-OP1	-11.22	95.60	105.70
34	BA	299	C	P-O3'-C3'	-11.22	106.23	119.70
85	AA	753	U	O4'-C1'-N1	11.22	117.18	108.20
86	AB	56	C	C5'-C4'-C3'	-11.22	98.05	116.00
41	BH	126	C	C2-N3-C4	-11.22	114.29	119.90
85	AA	505	U	C1'-O4'-C4'	-11.22	100.92	109.90
85	AA	77	C	O4'-C1'-N1	11.21	117.17	108.20
35	BB	378	C	C5'-C4'-C3'	-11.21	98.06	116.00
36	BC	157	U	O4'-C1'-N1	11.21	117.17	108.20
85	AA	1457	C	O5'-P-OP2	-11.21	95.61	105.70
85	AA	1931	C	C5'-C4'-C3'	11.21	133.94	116.00
85	AA	2234	C	O4'-C1'-N1	11.21	117.17	108.20
34	BA	1800	G	O4'-C1'-N9	11.21	117.17	108.20
37	BD	99	G	N1-C6-O6	11.21	126.62	119.90
34	BA	1454	G	C5-C6-O6	-11.21	121.88	128.60
38	BE	134	A	N1-C6-N6	-11.21	111.88	118.60
35	BB	912	C	P-O3'-C3'	11.20	133.14	119.70
38	BE	45	G	C4-N9-C1'	-11.20	111.94	126.50
85	AA	1730	C	P-O5'-C5'	11.20	138.82	120.90
36	BC	33	U	P-O5'-C5'	11.20	138.82	120.90
40	BG	168	A	C5-C6-N6	11.20	132.66	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	141	G	N3-C4-N9	-11.20	119.28	126.00
34	BA	868	C	O4'-C1'-N1	11.20	117.16	108.20
34	BA	1434	U	P-O3'-C3'	11.19	133.13	119.70
80	Bu	22	ARG	NE-CZ-NH1	11.19	125.90	120.30
41	BH	133	U	C5'-C4'-C3'	-11.19	98.09	116.00
38	BE	31	A	C5'-C4'-O4'	11.19	122.52	109.10
34	BA	1790	U	C6-N1-C2	-11.18	114.29	121.00
85	AA	24	U	P-O5'-C5'	11.18	138.79	120.90
85	AA	510	A	P-O5'-C5'	-11.18	103.01	120.90
34	BA	343	G	C5-C6-O6	-11.18	121.89	128.60
85	AA	836	A	P-O3'-C3'	11.18	133.11	119.70
34	BA	1725	U	P-O3'-C3'	11.17	133.11	119.70
36	BC	10	C	O4'-C4'-C3'	-11.17	92.83	104.00
85	AA	143	U	P-O3'-C3'	11.17	133.11	119.70
39	BF	17	U	P-O3'-C3'	11.17	133.10	119.70
35	BB	1541	G	P-O5'-C5'	11.17	138.77	120.90
35	BB	1226	G	C5'-C4'-C3'	11.16	133.86	116.00
35	BB	490	G	C5-C6-O6	-11.16	121.90	128.60
38	BE	112	G	C2-N3-C4	-11.16	106.32	111.90
85	AA	992	G	C4-N9-C1'	-11.16	112.00	126.50
37	BD	86	A	C5'-C4'-C3'	11.15	133.84	116.00
85	AA	1163	G	C5'-C4'-C3'	-11.15	98.15	116.00
34	BA	323	C	C3'-C2'-C1'	-11.15	92.58	101.50
38	BE	87	U	C5'-C4'-C3'	-11.15	98.17	116.00
34	BA	1828	A	P-O3'-C3'	-11.14	106.33	119.70
35	BB	797	C	N3-C4-N4	-11.14	110.20	118.00
85	AA	1283	C	C6-N1-C2	-11.14	115.84	120.30
34	BA	910	U	C2-N3-C4	-11.14	120.32	127.00
34	BA	23	A	N1-C6-N6	-11.14	111.92	118.60
85	AA	1731	G	C6-N1-C2	-11.14	118.42	125.10
34	BA	53	G	C4-N9-C1'	-11.13	112.02	126.50
34	BA	167	U	C6-N1-C1'	11.13	136.79	121.20
35	BB	1145	G	N1-C6-O6	11.13	126.58	119.90
34	BA	1313	U	O4'-C1'-N1	11.13	117.10	108.20
35	BB	115	A	N1-C6-N6	11.13	125.28	118.60
35	BB	1202	G	C5-C6-O6	-11.13	121.92	128.60
85	AA	180	A	N1-C6-N6	-11.12	111.93	118.60
34	BA	547	C	C5'-C4'-O4'	-11.12	95.76	109.10
35	BB	899	C	C6-N1-C1'	11.12	134.14	120.80
65	Bf	170	TYR	CB-CG-CD1	11.12	127.67	121.00
41	BH	126	C	C6-N1-C2	-11.12	115.85	120.30
40	BG	106	G	C1'-O4'-C4'	-11.11	101.01	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
65	Bf	339	TYR	CB-CG-CD2	-11.12	114.33	121.00
34	BA	135	G	P-O3'-C3'	-11.11	106.36	119.70
34	BA	1674	G	O4'-C1'-N9	11.11	117.09	108.20
35	BB	1202	G	P-O3'-C3'	-11.11	106.37	119.70
40	BG	84	U	O4'-C1'-N1	11.11	117.09	108.20
34	BA	690	G	C5'-C4'-C3'	-11.11	98.22	116.00
35	BB	521	U	P-O3'-C3'	11.11	133.03	119.70
34	BA	770	G	O4'-C1'-N9	11.11	117.09	108.20
34	BA	1500	G	C5'-C4'-C3'	11.11	133.78	116.00
85	AA	1361	A	C5-C6-N6	-11.11	114.81	123.70
35	BB	638	G	C5-C6-O6	-11.11	121.94	128.60
42	BI	18	HIS	C-N-CA	11.11	149.47	121.70
66	Bg	96	PHE	CB-CG-CD1	11.11	128.57	120.80
85	AA	901	C	C6-N1-C2	-11.11	115.86	120.30
85	AA	1464	G	C5-C6-N1	11.11	117.05	111.50
34	BA	548	G	O4'-C1'-N9	11.10	117.08	108.20
34	BA	201	A	C5-C6-N6	11.10	132.58	123.70
35	BB	13	A	P-O3'-C3'	-11.10	106.38	119.70
85	AA	2231	G	C5-C6-O6	-11.10	121.94	128.60
36	BC	156	A	P-O3'-C3'	11.10	133.01	119.70
85	AA	1598	A	O3'-P-O5'	-11.10	82.92	104.00
85	AA	707	U	P-O3'-C3'	-11.09	106.39	119.70
85	AA	1092	G	C8-N9-C4	-11.09	101.96	106.40
41	BH	91	G	C5-C6-O6	11.09	135.25	128.60
34	BA	1668	C	C2-N1-C1'	-11.08	106.61	118.80
85	AA	2211	G	C8-N9-C1'	11.08	141.41	127.00
36	BC	16	A	C8-N9-C4	11.08	110.23	105.80
43	BJ	59	ARG	NE-CZ-NH2	-11.08	114.76	120.30
35	BB	12	G	C5'-C4'-C3'	11.07	133.72	116.00
34	BA	131	A	N1-C6-N6	-11.07	111.96	118.60
85	AA	619	A	C1'-O4'-C4'	-11.06	101.05	109.90
34	BA	140	C	N3-C4-N4	11.06	125.74	118.00
34	BA	1673	G	N1-C6-O6	-11.06	113.26	119.90
34	BA	557	U	O4'-C4'-C3'	-11.06	92.94	104.00
35	BB	653	G	N1-C6-O6	11.06	126.54	119.90
35	BB	1426	G	N1-C6-O6	-11.06	113.26	119.90
85	AA	52	U	C5'-C4'-C3'	-11.06	98.31	116.00
34	BA	1535	G	C5-C6-O6	-11.06	121.97	128.60
7	A6	163	PHE	CB-CG-CD1	11.05	128.54	120.80
34	BA	1614	G	N1-C6-O6	11.05	126.53	119.90
85	AA	207	G	P-O5'-C5'	11.05	138.59	120.90
85	AA	1134	G	N1-C6-O6	11.05	126.53	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2196	G	C4-N9-C1'	-11.05	112.13	126.50
34	BA	678	C	C2-N3-C4	-11.05	114.38	119.90
38	BE	184	G	C5'-C4'-C3'	11.05	133.68	116.00
85	AA	53	G	C4-N9-C1'	-11.05	112.14	126.50
85	AA	578	U	C5'-C4'-C3'	-11.05	98.32	116.00
85	AA	556	C	O3'-P-O5'	-11.04	83.01	104.00
85	AA	817	G	P-O3'-C3'	11.05	132.96	119.70
34	BA	1486	U	C2-N3-C4	-11.04	120.38	127.00
39	BF	12	U	O4'-C1'-N1	11.04	117.03	108.20
35	BB	1138	A	P-O3'-C3'	-11.04	106.45	119.70
35	BB	1018	U	C5'-C4'-C3'	-11.04	98.34	116.00
35	BB	1460	G	C5-C6-O6	-11.04	121.98	128.60
85	AA	1977	G	C5-C6-O6	-11.04	121.98	128.60
85	AA	277	G	P-O5'-C5'	11.04	138.56	120.90
37	BD	99	G	C5-C6-O6	-11.03	121.98	128.60
40	BG	168	A	P-O3'-C3'	11.04	132.94	119.70
37	BD	87	G	C8-N9-C1'	11.03	141.34	127.00
85	AA	14	C	C6-N1-C2	-11.03	115.89	120.30
85	AA	1654	G	N1-C6-O6	11.03	126.52	119.90
34	BA	816	G	P-O3'-C3'	-11.03	106.47	119.70
34	BA	125	G	P-O3'-C3'	-11.02	106.47	119.70
34	BA	628	U	C1'-O4'-C4'	-11.02	101.08	109.90
34	BA	1719	G	N1-C6-O6	-11.02	113.28	119.90
35	BB	1466	A	P-O5'-C5'	11.02	138.54	120.90
34	BA	558	C	O5'-P-OP2	-11.02	95.78	105.70
85	AA	1562	U	O4'-C1'-N1	11.02	117.02	108.20
34	BA	481	A	N1-C2-N3	11.02	134.81	129.30
34	BA	1716	A	N1-C2-N3	-11.02	123.79	129.30
34	BA	1197	U	C2-N1-C1'	-11.01	104.48	117.70
35	BB	490	G	C5'-C4'-C3'	-11.01	98.39	116.00
35	BB	750	G	C5-C6-O6	-11.01	122.00	128.60
34	BA	1180	A	P-O3'-C3'	11.01	132.91	119.70
34	BA	105	U	C5'-C4'-C3'	-11.00	98.39	116.00
83	Bx	119	ARG	NE-CZ-NH1	11.00	125.80	120.30
85	AA	673	A	N1-C6-N6	-11.00	112.00	118.60
35	BB	1302	C	C2-N1-C1'	-11.00	106.70	118.80
35	BB	383	U	O4'-C1'-N1	11.00	117.00	108.20
34	BA	502	U	C2-N1-C1'	-10.99	104.51	117.70
39	BF	22	U	C2-N3-C4	-10.99	120.40	127.00
85	AA	466	A	C4-C5-N7	-10.99	105.20	110.70
85	AA	2194	U	C2-N3-C4	-10.99	120.40	127.00
35	BB	1328	C	P-O5'-C5'	10.98	138.47	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1458	A	C4'-C3'-C2'	10.98	113.58	102.60
37	BD	89	G	P-O3'-C3'	-10.98	106.53	119.70
34	BA	561	U	P-O3'-C3'	-10.98	106.53	119.70
35	BB	1045	G	C5'-C4'-C3'	10.98	133.56	116.00
34	BA	1194	G	C4-N9-C1'	-10.97	112.23	126.50
40	BG	31	G	C6-C5-N7	-10.97	123.81	130.40
34	BA	1056	C	C6-N1-C2	-10.97	115.91	120.30
35	BB	701	U	O4'-C1'-N1	10.97	116.98	108.20
35	BB	1529	G	C5-C6-O6	-10.97	122.02	128.60
38	BE	117	A	C4-N9-C1'	10.97	146.05	126.30
85	AA	619	A	C5'-C4'-O4'	10.97	122.26	109.10
35	BB	1143	A	C5'-C4'-C3'	-10.96	98.46	116.00
4	A3	24	ARG	NE-CZ-NH1	10.96	125.78	120.30
34	BA	306	G	N1-C6-O6	-10.95	113.33	119.90
85	AA	775	C	C2-N1-C1'	-10.95	106.75	118.80
85	AA	1698	A	P-O3'-C3'	10.95	132.84	119.70
34	BA	863	G	P-O3'-C3'	10.95	132.84	119.70
37	BD	65	G	C4-N9-C1'	-10.95	112.27	126.50
34	BA	896	U	C5'-C4'-C3'	-10.94	98.49	116.00
85	AA	2146	G	C4-N9-C1'	-10.95	112.27	126.50
34	BA	1595	G	P-O3'-C3'	10.94	132.83	119.70
35	BB	653	G	O5'-P-OP1	-10.94	95.85	105.70
85	AA	736	U	C6-N1-C2	-10.94	114.43	121.00
34	BA	249	A	O4'-C1'-N9	10.94	116.95	108.20
40	BG	131	U	C2-N3-C4	-10.94	120.44	127.00
34	BA	340	U	P-O5'-C5'	10.93	138.39	120.90
36	BC	34	U	C2-N3-C4	-10.93	120.44	127.00
38	BE	130	G	C4-N9-C1'	10.93	140.70	126.50
40	BG	61	A	C5-C6-N6	-10.93	114.96	123.70
85	AA	1421	U	O4'-C1'-N1	10.93	116.94	108.20
17	AI	130	TYR	CB-CG-CD2	-10.92	114.45	121.00
34	BA	140	C	C2-N1-C1'	-10.92	106.79	118.80
35	BB	653	G	C5'-C4'-C3'	10.91	133.46	116.00
85	AA	487	G	P-O5'-C5'	10.91	138.36	120.90
85	AA	2121	G	C8-N9-C1'	10.91	141.19	127.00
34	BA	322	U	N3-C4-O4	-10.91	111.76	119.40
35	BB	374	A	O4'-C1'-N9	10.91	116.93	108.20
85	AA	1110	A	N1-C6-N6	-10.91	112.06	118.60
34	BA	214	A	P-O3'-C3'	10.91	132.79	119.70
34	BA	1810	A	O4'-C1'-N9	10.91	116.93	108.20
35	BB	792	G	O4'-C1'-N9	10.91	116.93	108.20
39	BF	56	C	C2-N1-C1'	-10.91	106.80	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	13	A	O4'-C1'-N9	10.91	116.92	108.20
35	BB	1494	G	C4-N9-C1'	-10.90	112.32	126.50
38	BE	151	C	P-O5'-C5'	10.90	138.35	120.90
34	BA	718	U	O4'-C1'-N1	10.90	116.92	108.20
35	BB	1532	C	C2-N3-C4	-10.90	114.45	119.90
85	AA	67	C	C6-N1-C2	-10.90	115.94	120.30
85	AA	1484	G	C6-N1-C2	-10.90	118.56	125.10
38	BE	133	C	C4'-C3'-C2'	-10.89	91.71	102.60
35	BB	361	A	P-O3'-C3'	10.89	132.77	119.70
34	BA	599	U	O4'-C1'-N1	10.89	116.91	108.20
35	BB	1413	U	P-O5'-C5'	10.89	138.32	120.90
38	BE	28	C	C5'-C4'-O4'	10.88	122.16	109.10
85	AA	436	G	C5-C6-O6	-10.89	122.07	128.60
85	AA	1007	G	C8-N9-C1'	10.88	141.15	127.00
34	BA	1563	G	P-O3'-C3'	10.88	132.76	119.70
85	AA	838	G	C5-C6-O6	-10.88	122.07	128.60
37	BD	25	G	O4'-C1'-N9	10.88	116.91	108.20
34	BA	151	A	O4'-C1'-N9	10.88	116.90	108.20
34	BA	796	G	N1-C6-O6	10.88	126.43	119.90
34	BA	1299	G	C5-C6-O6	-10.88	122.07	128.60
85	AA	1538	C	O4'-C1'-N1	10.88	116.90	108.20
86	AB	68	C	C2-N1-C1'	-10.88	106.83	118.80
38	BE	9	C	O3'-P-O5'	10.88	124.67	104.00
85	AA	2084	U	P-O3'-C3'	-10.88	106.65	119.70
35	BB	454	U	C5'-C4'-C3'	-10.87	98.60	116.00
40	BG	24	A	C6-C5-N7	-10.87	124.69	132.30
85	AA	854	A	O5'-C5'-C4'	10.87	132.35	111.70
35	BB	1006	C	O4'-C1'-N1	10.87	116.89	108.20
34	BA	603	U	P-O3'-C3'	-10.87	106.66	119.70
36	BC	3	C	O4'-C1'-N1	10.87	116.89	108.20
34	BA	15	G	N1-C6-O6	10.86	126.42	119.90
34	BA	483	A	P-O5'-C5'	10.86	138.28	120.90
39	BF	5	U	C2-N3-C4	-10.86	120.48	127.00
85	AA	1814	U	C6-N1-C2	-10.86	114.48	121.00
39	BF	23	G	O4'-C4'-C3'	-10.86	93.14	104.00
34	BA	625	U	O4'-C1'-N1	10.86	116.89	108.20
38	BE	36	U	N1-C2-N3	10.86	121.41	114.90
34	BA	1442	A	C6-N1-C2	-10.86	112.09	118.60
35	BB	1100	C	C5'-C4'-C3'	-10.85	98.64	116.00
41	BH	72	G	C4'-C3'-C2'	-10.85	91.75	102.60
85	AA	648	G	C4-N9-C1'	-10.85	112.40	126.50
34	BA	1002	U	C1'-O4'-C4'	-10.85	101.22	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	72	C	C6-N1-C2	-10.84	115.96	120.30
40	BG	11	G	P-O5'-C5'	10.84	138.25	120.90
85	AA	967	C	P-O5'-C5'	10.84	138.25	120.90
36	BC	110	A	C1'-O4'-C4'	-10.84	101.23	109.90
38	BE	70	C	P-O3'-C3'	10.84	132.71	119.70
85	AA	802	A	N1-C6-N6	10.84	125.11	118.60
34	BA	382	G	C8-N9-C4	-10.84	102.06	106.40
34	BA	800	G	N3-C4-C5	-10.84	123.18	128.60
41	BH	30	C	P-O5'-C5'	10.84	138.24	120.90
35	BB	703	U	P-O5'-C5'	10.83	138.23	120.90
35	BB	802	G	N1-C6-O6	-10.83	113.40	119.90
85	AA	1111	A	P-O3'-C3'	10.83	132.70	119.70
85	AA	1701	G	C8-N9-C1'	10.83	141.08	127.00
34	BA	557	U	N1-C1'-C2'	-10.83	99.92	114.00
85	AA	687	G	C5-C6-N1	10.83	116.92	111.50
39	BF	15	U	O4'-C1'-N1	10.83	116.86	108.20
85	AA	424	A	C3'-C2'-C1'	-10.83	92.84	101.50
34	BA	398	G	C5'-C4'-C3'	10.82	133.32	116.00
34	BA	1670	A	N1-C6-N6	-10.82	112.11	118.60
35	BB	949	G	P-O3'-C3'	10.82	132.69	119.70
34	BA	616	G	O4'-C1'-N9	10.82	116.86	108.20
35	BB	1346	A	C6-N1-C2	-10.82	112.11	118.60
38	BE	111	C	C2-N1-C1'	10.82	130.70	118.80
38	BE	58	U	C6-N1-C2	-10.82	114.51	121.00
40	BG	182	G	N1-C6-O6	10.81	126.39	119.90
85	AA	469	G	N1-C2-N3	10.81	130.39	123.90
34	BA	214	A	N1-C6-N6	10.81	125.09	118.60
38	BE	85	G	C5'-C4'-C3'	10.81	133.29	116.00
38	BE	116	U	P-O3'-C3'	-10.81	106.73	119.70
41	BH	75	G	N1-C2-N3	-10.81	117.42	123.90
34	BA	166	G	C4-N9-C1'	-10.80	112.45	126.50
40	BG	31	G	C4-C5-C6	-10.81	112.32	118.80
40	BG	31	G	C5-C6-N1	10.80	116.90	111.50
85	AA	555	C	C6-N1-C2	-10.81	115.98	120.30
85	AA	589	A	P-O3'-C3'	10.80	132.66	119.70
34	BA	216	C	P-O3'-C3'	-10.80	106.74	119.70
35	BB	392	G	N1-C6-O6	-10.80	113.42	119.90
35	BB	557	C	C6-N1-C2	-10.80	115.98	120.30
35	BB	1306	G	C5-C6-O6	10.80	135.08	128.60
34	BA	91	C	C3'-C2'-C1'	-10.79	92.86	101.50
85	AA	1002	G	C5-C6-O6	-10.79	122.12	128.60
85	AA	1469	G	C8-N9-C4	-10.79	102.08	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	63	G	N1-C6-O6	10.79	126.37	119.90
85	AA	2056	C	C6-N1-C1'	10.79	133.75	120.80
34	BA	1792	U	P-O3'-C3'	10.79	132.64	119.70
85	AA	1004	G	C8-N9-C4	-10.79	102.08	106.40
38	BE	133	C	C6-N1-C2	-10.78	115.99	120.30
34	BA	1845	G	N3-C2-N2	-10.78	112.35	119.90
38	BE	14	C	C6-N1-C2	-10.78	115.99	120.30
34	BA	1835	A	C5'-C4'-C3'	-10.78	98.75	116.00
85	AA	355	G	P-O3'-C3'	10.78	132.64	119.70
34	BA	1729	G	O4'-C1'-N9	10.78	116.82	108.20
34	BA	764	G	C5'-C4'-C3'	10.78	133.24	116.00
35	BB	991	C	O4'-C1'-N1	10.78	116.82	108.20
34	BA	116	G	C1'-O4'-C4'	-10.78	101.28	109.90
34	BA	1112	U	C6-N1-C2	-10.78	114.53	121.00
35	BB	1401	G	C5-C6-O6	-10.78	122.13	128.60
38	BE	91	G	C8-N9-C1'	10.78	141.01	127.00
85	AA	2060	G	C5-C6-O6	-10.78	122.14	128.60
85	AA	2084	U	P-O5'-C5'	10.78	138.14	120.90
36	BC	87	C	C5'-C4'-C3'	-10.77	98.77	116.00
85	AA	1493	A	P-O3'-C3'	-10.77	106.78	119.70
85	AA	994	A	P-O3'-C3'	-10.77	106.78	119.70
34	BA	1325	G	C5-C6-O6	-10.77	122.14	128.60
85	AA	504	U	C1'-O4'-C4'	-10.77	101.29	109.90
35	BB	1145	G	C5-C6-O6	-10.77	122.14	128.60
34	BA	343	G	C6-N1-C2	-10.76	118.64	125.10
34	BA	631	G	C4-N9-C1'	-10.76	112.51	126.50
61	Bb	32	ARG	NE-CZ-NH1	10.76	125.68	120.30
35	BB	970	C	O4'-C1'-N1	10.76	116.81	108.20
36	BC	168	C	C5'-C4'-C3'	-10.76	98.79	116.00
85	AA	742	U	C6-N1-C2	-10.76	114.55	121.00
34	BA	593	G	C8-N9-C4	-10.75	102.10	106.40
35	BB	1030	U	C1'-O4'-C4'	-10.75	101.30	109.90
85	AA	860	C	C6-N1-C1'	10.75	133.70	120.80
85	AA	2099	C	C2-N3-C4	-10.75	114.53	119.90
85	AA	2231	G	P-O5'-C5'	10.75	138.10	120.90
35	BB	1517	G	C5-C6-O6	-10.75	122.15	128.60
35	BB	1470	G	C5-C6-O6	-10.75	122.15	128.60
85	AA	489	C	O4'-C1'-N1	10.75	116.80	108.20
34	BA	1797	A	P-O3'-C3'	10.74	132.59	119.70
37	BD	25	G	C4-N9-C1'	-10.74	112.53	126.50
85	AA	961	U	C5'-C4'-C3'	-10.74	98.81	116.00
85	AA	899	A	P-O3'-C3'	-10.74	106.81	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	322	U	C5-C4-O4	10.74	132.34	125.90
38	BE	97	G	C5-C6-O6	-10.74	122.16	128.60
85	AA	1491	G	P-O3'-C3'	10.74	132.59	119.70
34	BA	1829	A	P-O3'-C3'	-10.74	106.81	119.70
38	BE	101	C	P-O5'-C5'	10.74	138.08	120.90
34	BA	575	U	C4'-C3'-C2'	10.73	113.33	102.60
85	AA	133	G	C4-N9-C1'	-10.73	112.55	126.50
40	BG	171	A	O4'-C4'-C3'	-10.73	93.27	104.00
40	BG	75	C	C2-N3-C4	-10.73	114.54	119.90
38	BE	89	G	C5-C6-O6	10.73	135.04	128.60
34	BA	1613	G	P-O3'-C3'	-10.72	106.83	119.70
34	BA	1722	U	C1'-O4'-C4'	-10.72	101.32	109.90
35	BB	669	A	P-O5'-C5'	10.72	138.05	120.90
41	BH	113	G	N1-C6-O6	10.72	126.33	119.90
35	BB	1168	G	C8-N9-C1'	10.72	140.94	127.00
35	BB	1495	U	C2-N3-C4	-10.72	120.57	127.00
85	AA	972	G	C5'-C4'-C3'	10.72	133.15	116.00
85	AA	1872	G	C5-C6-O6	-10.72	122.17	128.60
34	BA	593	G	O4'-C1'-N9	10.71	116.77	108.20
34	BA	780	U	O4'-C4'-C3'	-10.72	93.28	104.00
36	BC	59	A	N1-C6-N6	-10.71	112.17	118.60
85	AA	1713	A	O4'-C1'-N9	10.71	116.77	108.20
34	BA	240	C	O4'-C1'-N1	10.71	116.77	108.20
59	BZ	48	ARG	NE-CZ-NH1	10.71	125.66	120.30
85	AA	1458	G	C8-N9-C4	-10.71	102.12	106.40
85	AA	1247	A	C5-C6-N6	-10.70	115.14	123.70
34	BA	6	C	P-O5'-C5'	10.70	138.02	120.90
34	BA	1313	U	C1'-O4'-C4'	-10.70	101.34	109.90
65	Bf	416	PHE	CB-CG-CD1	-10.70	113.31	120.80
38	BE	117	A	C5-N7-C8	-10.69	98.55	103.90
34	BA	660	C	C2-N1-C1'	-10.69	107.04	118.80
35	BB	145	G	P-O3'-C3'	10.69	132.53	119.70
34	BA	856	G	P-O5'-C5'	10.69	138.00	120.90
35	BB	389	G	C5-C6-O6	-10.69	122.19	128.60
35	BB	809	U	C2-N1-C1'	-10.69	104.87	117.70
47	BN	52	PHE	CB-CG-CD1	10.69	128.28	120.80
85	AA	157	G	C8-N9-C4	10.69	110.68	106.40
34	BA	1218	G	N1-C6-O6	-10.69	113.49	119.90
85	AA	863	C	C2-N3-C4	-10.69	114.56	119.90
35	BB	91	G	C6-N1-C2	-10.68	118.69	125.10
85	AA	211	C	C5'-C4'-C3'	10.68	133.09	116.00
34	BA	626	G	C8-N9-C4	-10.68	102.13	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	303	A	C5'-C4'-C3'	10.68	133.08	116.00
34	BA	1569	C	C6-N1-C2	-10.67	116.03	120.30
38	BE	115	U	O4'-C1'-N1	10.67	116.74	108.20
41	BH	91	G	N1-C6-O6	-10.67	113.50	119.90
85	AA	908	C	C5'-C4'-C3'	-10.67	98.92	116.00
85	AA	8	U	P-O3'-C3'	10.67	132.50	119.70
85	AA	450	A	C1'-O4'-C4'	-10.67	101.36	109.90
85	AA	2208	G	C8-N9-C4	-10.67	102.13	106.40
34	BA	1808	A	P-O3'-C3'	-10.67	106.90	119.70
34	BA	232	U	P-O5'-C5'	10.66	137.96	120.90
34	BA	587	U	P-O3'-C3'	10.66	132.49	119.70
34	BA	651	U	C3'-C2'-C1'	-10.66	92.97	101.50
40	BG	14	G	C5'-C4'-C3'	10.66	133.06	116.00
85	AA	925	G	C4-N9-C1'	-10.66	112.64	126.50
85	AA	2033	C	O4'-C1'-N1	10.66	116.73	108.20
34	BA	596	G	C5-C6-O6	-10.66	122.21	128.60
38	BE	127	G	C2-N3-C4	10.66	117.23	111.90
34	BA	1204	U	C5'-C4'-C3'	-10.66	98.95	116.00
35	BB	1294	C	O4'-C1'-N1	10.66	116.73	108.20
40	BG	9	G	C6-N1-C2	-10.66	118.71	125.10
34	BA	260	A	N1-C6-N6	-10.65	112.21	118.60
85	AA	877	G	C5-C6-N1	10.65	116.83	111.50
36	BC	129	C	C6-N1-C2	-10.65	116.04	120.30
35	BB	993	A	P-O3'-C3'	-10.65	106.92	119.70
85	AA	976	G	C4-N9-C1'	-10.65	112.66	126.50
85	AA	989	U	O4'-C1'-N1	10.65	116.72	108.20
34	BA	18	G	C6-N1-C2	-10.65	118.71	125.10
34	BA	743	A	C5-C6-N6	-10.65	115.18	123.70
34	BA	865	C	O4'-C1'-N1	10.65	116.72	108.20
38	BE	115	U	O3'-P-O5'	-10.65	83.77	104.00
34	BA	1182	U	P-O5'-C5'	10.64	137.93	120.90
85	AA	120	C	C6-N1-C1'	-10.64	108.03	120.80
85	AA	2023	U	O4'-C1'-N1	10.64	116.71	108.20
41	BH	129	G	O4'-C1'-N9	10.64	116.71	108.20
85	AA	2077	G	C5-C6-O6	-10.64	122.22	128.60
34	BA	1613	G	N1-C6-O6	10.64	126.28	119.90
34	BA	1599	A	C5'-C4'-C3'	-10.64	98.98	116.00
34	BA	1830	A	O4'-C1'-N9	10.64	116.71	108.20
35	BB	1250	A	N1-C6-N6	-10.64	112.22	118.60
35	BB	833	G	C8-N9-C1'	10.64	140.83	127.00
35	BB	989	C	C2-N1-C1'	10.63	130.50	118.80
38	BE	195	G	C5-C6-O6	-10.64	122.22	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	182	G	C5-C6-O6	-10.63	122.22	128.60
85	AA	691	U	C6-N1-C2	-10.64	114.62	121.00
85	AA	899	A	O4'-C1'-N9	10.63	116.70	108.20
41	BH	111	U	O4'-C1'-N1	10.63	116.70	108.20
85	AA	1464	G	C1'-O4'-C4'	-10.63	101.39	109.90
34	BA	573	U	C5'-C4'-C3'	10.63	133.00	116.00
40	BG	62	C	C6-N1-C2	-10.63	116.05	120.30
85	AA	286	C	O4'-C1'-N1	10.62	116.70	108.20
85	AA	442	G	C5-C6-O6	-10.62	122.22	128.60
34	BA	570	G	N9-C4-C5	-10.62	101.15	105.40
41	BH	6	U	C2-N1-C1'	-10.62	104.95	117.70
85	AA	337	C	O4'-C1'-N1	10.62	116.70	108.20
85	AA	455	G	C2-N3-C4	-10.62	106.59	111.90
34	BA	690	G	C4-N9-C1'	-10.62	112.70	126.50
34	BA	736	G	C5-C6-O6	-10.62	122.23	128.60
35	BB	975	G	C5'-C4'-C3'	10.62	132.99	116.00
34	BA	985	C	C4'-C3'-C2'	10.61	113.21	102.60
41	BH	20	A	C5'-C4'-C3'	-10.62	99.02	116.00
34	BA	524	G	O3'-P-O5'	10.61	124.17	104.00
34	BA	572	G	O3'-P-O5'	10.61	124.16	104.00
40	BG	9	G	C8-N9-C1'	10.61	140.79	127.00
41	BH	123	G	N9-C1'-C2'	-10.61	100.21	114.00
34	BA	25	C	O4'-C1'-N1	10.61	116.68	108.20
34	BA	1730	A	P-O5'-C5'	10.61	137.87	120.90
35	BB	692	G	C5-C6-O6	10.61	134.96	128.60
85	AA	325	C	P-O3'-C3'	10.61	132.43	119.70
85	AA	424	A	P-O3'-C3'	-10.61	106.97	119.70
34	BA	578	C	O4'-C1'-N1	10.60	116.68	108.20
35	BB	1212	C	C2-N3-C4	-10.60	114.60	119.90
34	BA	944	G	C4-N9-C1'	-10.60	112.72	126.50
39	BF	51	C	O3'-P-O5'	-10.60	83.86	104.00
35	BB	57	G	N9-C1'-C2'	-10.60	100.22	114.00
35	BB	1229	A	O3'-P-O5'	10.60	124.13	104.00
36	BC	169	G	C4-N9-C1'	-10.60	112.72	126.50
85	AA	1814	U	N3-C2-O2	-10.60	114.78	122.20
38	BE	136	G	P-O3'-C3'	-10.59	106.99	119.70
85	AA	1683	U	C2-N1-C1'	10.59	130.41	117.70
38	BE	117	A	C8-N9-C4	-10.59	101.56	105.80
40	BG	175	G	P-O5'-C5'	10.59	137.85	120.90
34	BA	248	G	C5-C6-O6	-10.59	122.25	128.60
34	BA	547	C	C3'-C2'-C1'	10.59	109.97	101.50
34	BA	1364	G	O4'-C1'-N9	10.59	116.67	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	54	U	C4'-C3'-C2'	-10.59	92.01	102.60
35	BB	60	A	C1'-O4'-C4'	-10.59	101.43	109.90
35	BB	1188	A	N1-C6-N6	-10.59	112.25	118.60
34	BA	870	C	O4'-C1'-N1	10.58	116.67	108.20
35	BB	1226	G	N1-C6-O6	10.58	126.25	119.90
85	AA	1979	A	O4'-C1'-N9	10.58	116.67	108.20
85	AA	2082	C	C6-N1-C2	-10.58	116.07	120.30
34	BA	882	G	C4-N9-C1'	-10.58	112.74	126.50
85	AA	668	A	P-O3'-C3'	10.58	132.40	119.70
34	BA	944	G	C5-C6-O6	-10.58	122.25	128.60
35	BB	780	U	C2-N3-C4	-10.58	120.65	127.00
35	BB	901	U	P-O5'-C5'	10.58	137.82	120.90
85	AA	687	G	C2'-C3'-O3'	10.58	132.77	109.50
36	BC	121	G	P-O3'-C3'	10.58	132.39	119.70
35	BB	1441	C	C6-N1-C1'	-10.57	108.11	120.80
85	AA	561	C	O4'-C1'-N1	10.57	116.66	108.20
85	AA	2075	C	P-O5'-C5'	10.57	137.82	120.90
27	AT	65	PHE	CB-CG-CD2	-10.57	113.40	120.80
35	BB	1520	C	O4'-C1'-N1	10.57	116.66	108.20
34	BA	140	C	C5-C4-N4	-10.57	112.80	120.20
85	AA	1921	G	O5'-C5'-C4'	10.57	131.78	111.70
35	BB	991	C	P-O3'-C3'	10.56	132.38	119.70
85	AA	1999	C	O4'-C1'-N1	10.56	116.65	108.20
34	BA	712	C	O4'-C1'-N1	10.56	116.65	108.20
85	AA	333	A	P-O3'-C3'	10.56	132.38	119.70
37	BD	47	U	P-O3'-C3'	10.56	132.37	119.70
85	AA	82	A	O3'-P-O5'	-10.56	83.94	104.00
34	BA	993	C	C2-N3-C4	-10.56	114.62	119.90
34	BA	1689	U	P-O3'-C3'	10.55	132.37	119.70
35	BB	1148	U	P-O3'-C3'	10.55	132.37	119.70
38	BE	55	C	C2-N1-C1'	10.56	130.41	118.80
35	BB	1535	G	P-O5'-C5'	-10.55	104.02	120.90
86	AB	67	C	C5'-C4'-C3'	-10.55	99.11	116.00
34	BA	605	G	C5'-C4'-O4'	10.55	121.76	109.10
34	BA	214	A	C5-C6-N6	-10.55	115.26	123.70
85	AA	510	A	P-O3'-C3'	10.55	132.36	119.70
34	BA	65	A	P-O5'-C5'	10.55	137.78	120.90
34	BA	136	A	C4-N9-C1'	-10.55	107.31	126.30
85	AA	386	G	C4-N9-C1'	10.55	140.21	126.50
35	BB	1063	C	C2-N1-C1'	-10.55	107.20	118.80
15	AG	121	ARG	NE-CZ-NH1	10.55	125.57	120.30
34	BA	326	A	P-O3'-C3'	-10.55	107.04	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	76	C	P-O3'-C3'	-10.55	107.05	119.70
37	BD	26	C	C3'-C2'-C1'	-10.55	93.06	101.50
85	AA	658	C	C6-N1-C2	-10.55	116.08	120.30
39	BF	7	G	C8-N9-C1'	-10.54	113.29	127.00
50	BQ	164	ARG	NE-CZ-NH2	-10.54	115.03	120.30
35	BB	17	U	P-O5'-C5'	10.54	137.77	120.90
23	AP	87	ARG	NE-CZ-NH1	10.54	125.57	120.30
34	BA	1070	G	N3-C2-N2	10.54	127.28	119.90
35	BB	2	C	O4'-C1'-N1	10.54	116.63	108.20
40	BG	4	A	N1-C6-N6	-10.54	112.28	118.60
41	BH	32	U	C2-N3-C4	-10.54	120.67	127.00
38	BE	135	A	O4'-C1'-N9	10.54	116.63	108.20
48	BO	103	ARG	NE-CZ-NH1	10.54	125.57	120.30
85	AA	740	A	O4'-C1'-N9	10.54	116.63	108.20
35	BB	882	U	P-O3'-C3'	10.53	132.34	119.70
35	BB	129	U	C2-N3-C4	-10.53	120.68	127.00
85	AA	1495	G	O5'-C5'-C4'	-10.53	91.69	111.70
35	BB	800	U	P-O5'-C5'	10.53	137.75	120.90
35	BB	868	C	C6-N1-C1'	10.53	133.43	120.80
85	AA	424	A	C4'-C3'-O3'	10.53	134.06	113.00
49	BP	11	ARG	NE-CZ-NH1	10.53	125.56	120.30
85	AA	1499	G	C4-N9-C1'	-10.53	112.82	126.50
34	BA	764	G	O4'-C1'-N9	10.52	116.62	108.20
85	AA	861	G	C2-N3-C4	-10.52	106.64	111.90
72	Bm	38	TYR	CB-CG-CD1	-10.52	114.69	121.00
34	BA	24	C	C6-N1-C2	-10.52	116.09	120.30
34	BA	670	U	O4'-C1'-N1	10.52	116.61	108.20
35	BB	796	C	O4'-C1'-N1	10.52	116.61	108.20
85	AA	1729	C	C6-N1-C2	-10.52	116.09	120.30
40	BG	24	A	P-O5'-C5'	-10.52	104.08	120.90
40	BG	105	A	P-O5'-C5'	10.52	137.73	120.90
54	BU	49	ARG	NE-CZ-NH1	10.52	125.56	120.30
86	AB	65	G	C5-C6-O6	-10.52	122.29	128.60
35	BB	816	U	N1-C2-N3	10.51	121.21	114.90
35	BB	756	C	O4'-C1'-N1	10.51	116.61	108.20
35	BB	1026	G	P-O5'-C5'	10.51	137.71	120.90
40	BG	85	C	C6-N1-C2	-10.51	116.10	120.30
31	AX	152	ARG	NE-CZ-NH1	10.50	125.55	120.30
35	BB	833	G	C1'-O4'-C4'	-10.50	101.50	109.90
85	AA	538	A	O4'-C1'-N9	10.50	116.60	108.20
35	BB	868	C	N3-C2-O2	-10.50	114.55	121.90
85	AA	986	U	C6-N1-C1'	10.50	135.90	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	23	G	C8-N9-C4	-10.50	102.20	106.40
42	BI	187	ARG	NE-CZ-NH1	10.50	125.55	120.30
65	Bf	170	TYR	CB-CG-CD2	-10.50	114.70	121.00
86	AB	39	U	O4'-C1'-N1	10.50	116.60	108.20
35	BB	691	A	N1-C6-N6	10.49	124.90	118.60
85	AA	357	C	O4'-C1'-N1	10.49	116.60	108.20
35	BB	472	C	C2-N1-C1'	-10.49	107.26	118.80
35	BB	261	C	O4'-C1'-N1	10.49	116.59	108.20
85	AA	203	C	O4'-C1'-N1	10.49	116.59	108.20
34	BA	604	G	O5'-P-OP2	-10.49	96.26	105.70
85	AA	964	C	C2-N1-C1'	10.48	130.33	118.80
34	BA	1786	C	C6-N1-C2	-10.48	116.11	120.30
85	AA	984	A	C5'-C4'-C3'	-10.48	99.23	116.00
85	AA	205	A	P-O3'-C3'	-10.48	107.12	119.70
85	AA	456	A	P-O3'-C3'	-10.48	107.12	119.70
85	AA	992	G	C8-N9-C1'	10.48	140.63	127.00
34	BA	519	G	O4'-C1'-N9	10.48	116.58	108.20
85	AA	1456	A	O4'-C1'-N9	10.48	116.58	108.20
85	AA	1095	C	O4'-C1'-N1	10.48	116.58	108.20
35	BB	815	G	C4-N9-C1'	-10.48	112.88	126.50
42	BI	159	PHE	CB-CG-CD2	-10.47	113.47	120.80
34	BA	223	U	C2-N3-C4	-10.47	120.72	127.00
35	BB	423	G	N1-C6-O6	-10.47	113.62	119.90
40	BG	14	G	C5-C6-O6	-10.47	122.32	128.60
85	AA	881	C	O4'-C1'-N1	10.47	116.58	108.20
85	AA	914	U	O4'-C1'-N1	10.46	116.57	108.20
41	BH	75	G	OP1-P-OP2	-10.46	103.91	119.60
34	BA	1711	G	C5'-C4'-C3'	-10.46	99.27	116.00
85	AA	721	C	C6-N1-C2	-10.46	116.12	120.30
35	BB	1509	G	C5'-C4'-C3'	-10.46	99.27	116.00
85	AA	1115	G	C6-N1-C2	-10.46	118.83	125.10
34	BA	1830	A	C5'-C4'-C3'	-10.46	99.27	116.00
85	AA	645	C	C6-N1-C2	-10.46	116.12	120.30
34	BA	1200	U	C2-N1-C1'	10.45	130.24	117.70
35	BB	1137	G	C5'-C4'-C3'	-10.45	99.28	116.00
38	BE	13	A	C4'-C3'-C2'	10.45	113.05	102.60
85	AA	715	G	O4'-C1'-N9	10.45	116.56	108.20
35	BB	76	C	O4'-C1'-N1	10.45	116.56	108.20
37	BD	47	U	C5'-C4'-C3'	10.45	132.71	116.00
41	BH	34	G	C5'-C4'-C3'	-10.45	99.29	116.00
85	AA	386	G	C8-N9-C1'	-10.45	113.42	127.00
85	AA	455	G	C5'-C4'-C3'	-10.44	99.29	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	804	G	C4'-C3'-C2'	-10.44	92.16	102.60
85	AA	1916	A	C2-N3-C4	-10.44	105.38	110.60
38	BE	71	A	P-O3'-C3'	10.44	132.22	119.70
35	BB	1312	U	P-O5'-C5'	10.44	137.60	120.90
34	BA	480	G	C8-N9-C4	10.44	110.57	106.40
5	A4	71	ARG	NE-CZ-NH1	10.43	125.52	120.30
34	BA	1641	G	C8-N9-C1'	10.43	140.56	127.00
85	AA	684	G	C6-N1-C2	-10.43	118.84	125.10
85	AA	1730	C	O3'-P-O5'	-10.43	84.18	104.00
41	BH	134	U	C2-N1-C1'	-10.43	105.19	117.70
85	AA	2082	C	C2-N1-C1'	10.43	130.27	118.80
34	BA	669	U	O4'-C1'-N1	10.43	116.54	108.20
35	BB	389	G	N1-C6-O6	10.43	126.16	119.90
35	BB	1357	C	C6-N1-C1'	-10.42	108.29	120.80
35	BB	1514	G	C8-N9-C1'	10.42	140.55	127.00
34	BA	1080	U	C5'-C4'-C3'	10.42	132.67	116.00
35	BB	867	C	O4'-C1'-N1	10.42	116.54	108.20
35	BB	1235	A	P-O5'-C5'	-10.42	104.23	120.90
41	BH	123	G	C5-C6-O6	-10.42	122.35	128.60
34	BA	242	U	C2-N3-C4	-10.42	120.75	127.00
34	BA	1568	A	N1-C6-N6	10.41	124.85	118.60
34	BA	255	G	C5'-C4'-C3'	-10.41	99.34	116.00
85	AA	801	U	P-O5'-C5'	10.41	137.56	120.90
7	A6	15	ARG	NE-CZ-NH1	10.41	125.50	120.30
34	BA	755	G	C5-C6-O6	-10.41	122.35	128.60
38	BE	191	U	P-O5'-C5'	-10.41	104.25	120.90
85	AA	68	A	C5'-C4'-C3'	-10.41	99.35	116.00
34	BA	521	C	O4'-C1'-N1	10.41	116.53	108.20
36	BC	123	G	N1-C6-O6	10.41	126.14	119.90
34	BA	141	G	N1-C6-O6	10.40	126.14	119.90
85	AA	1935	G	P-O5'-C5'	10.40	137.55	120.90
34	BA	668	G	P-O3'-C3'	-10.40	107.22	119.70
34	BA	1184	A	O4'-C1'-N9	10.40	116.52	108.20
34	BA	1735	G	N1-C6-O6	10.40	126.14	119.90
15	AG	94	ARG	NE-CZ-NH1	10.40	125.50	120.30
34	BA	1468	U	C5'-C4'-C3'	-10.40	99.36	116.00
35	BB	1151	A	C8-N9-C4	-10.40	101.64	105.80
34	BA	489	A	O4'-C1'-N9	10.40	116.52	108.20
84	By	94	TYR	CB-CG-CD1	-10.40	114.76	121.00
34	BA	837	U	O3'-P-O5'	-10.40	84.25	104.00
35	BB	1052	G	N1-C6-O6	10.39	126.14	119.90
35	BB	1167	C	C6-N1-C2	-10.39	116.14	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	90	A	O4'-C1'-N9	10.39	116.52	108.20
38	BE	130	G	C5'-C4'-C3'	10.39	132.63	116.00
35	BB	1410	G	C5'-C4'-C3'	-10.39	99.38	116.00
34	BA	81	C	C4'-C3'-C2'	10.39	112.99	102.60
35	BB	599	U	C5'-C4'-C3'	-10.39	99.38	116.00
35	BB	1480	G	N1-C6-O6	10.38	126.13	119.90
39	BF	16	C	C5'-C4'-C3'	10.38	132.61	116.00
85	AA	696	G	P-O3'-C3'	10.38	132.16	119.70
34	BA	804	G	C8-N9-C4	-10.38	102.25	106.40
34	BA	6	C	O5'-P-OP2	-10.38	96.36	105.70
34	BA	214	A	C3'-C2'-C1'	-10.38	93.19	101.50
35	BB	1302	C	C5-C4-N4	-10.38	112.93	120.20
34	BA	620	C	O4'-C1'-N1	10.38	116.50	108.20
41	BH	62	C	O4'-C1'-N1	10.38	116.50	108.20
85	AA	2241	C	O4'-C1'-N1	10.38	116.50	108.20
34	BA	1320	A	N1-C2-N3	-10.37	124.11	129.30
38	BE	132	U	C2-N1-C1'	-10.37	105.25	117.70
85	AA	731	U	C2-N3-C4	-10.37	120.78	127.00
34	BA	1688	G	C5-C6-O6	-10.37	122.38	128.60
35	BB	555	G	C5'-C4'-C3'	-10.37	99.41	116.00
85	AA	1310	G	N1-C6-O6	10.37	126.12	119.90
85	AA	1650	G	P-O3'-C3'	10.37	132.15	119.70
35	BB	6	A	C5'-C4'-C3'	-10.37	99.42	116.00
34	BA	574	U	O4'-C1'-N1	10.36	116.49	108.20
37	BD	92	G	O4'-C1'-N9	10.36	116.49	108.20
34	BA	111	U	C3'-C2'-C1'	-10.36	93.21	101.50
34	BA	692	U	N3-C2-O2	10.36	129.45	122.20
35	BB	1134	G	N1-C6-O6	10.36	126.11	119.90
34	BA	141	G	C5-C6-O6	-10.36	122.39	128.60
34	BA	194	G	N1-C6-O6	-10.35	113.69	119.90
34	BA	15	G	C8-N9-C4	10.35	110.54	106.40
34	BA	1732	A	C5'-C4'-O4'	10.35	121.52	109.10
85	AA	408	C	C6-N1-C2	-10.35	116.16	120.30
85	AA	963	U	C1'-O4'-C4'	-10.35	101.62	109.90
35	BB	619	A	P-O3'-C3'	-10.35	107.28	119.70
47	BN	120	ARG	NE-CZ-NH1	10.35	125.47	120.30
36	BC	25	C	C5-C4-N4	-10.35	112.96	120.20
85	AA	1506	U	C2-N1-C1'	-10.35	105.28	117.70
85	AA	1829	C	C2-N3-C4	-10.35	114.73	119.90
34	BA	172	A	N1-C6-N6	10.34	124.81	118.60
34	BA	72	U	O5'-P-OP1	10.34	123.11	110.70
34	BA	1474	G	C5'-C4'-C3'	-10.34	99.45	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	157	U	P-O3'-C3'	10.34	132.10	119.70
34	BA	526	C	P-O3'-C3'	10.34	132.11	119.70
34	BA	528	C	C5'-C4'-C3'	10.34	132.54	116.00
35	BB	115	A	C5-C6-N6	-10.34	115.43	123.70
85	AA	2200	A	C5'-C4'-C3'	-10.33	99.47	116.00
34	BA	875	G	N9-C4-C5	-10.33	101.27	105.40
85	AA	2018	U	C2-N3-C4	-10.33	120.80	127.00
34	BA	429	G	C5-C6-O6	-10.33	122.40	128.60
34	BA	618	G	C8-N9-C4	-10.33	102.27	106.40
85	AA	941	C	P-O3'-C3'	10.33	132.09	119.70
34	BA	1673	G	P-O3'-C3'	-10.32	107.31	119.70
35	BB	878	G	N9-C1'-C2'	-10.32	100.58	114.00
34	BA	1410	C	C5'-C4'-C3'	-10.32	99.49	116.00
35	BB	1062	G	C5-C6-O6	-10.32	122.41	128.60
34	BA	508	C	P-O3'-C3'	10.32	132.08	119.70
34	BA	1042	U	P-O3'-C3'	10.31	132.08	119.70
34	BA	1294	C	O4'-C1'-N1	10.31	116.45	108.20
85	AA	658	C	C5'-C4'-C3'	-10.31	99.50	116.00
35	BB	781	U	P-O5'-C5'	10.31	137.40	120.90
41	BH	98	U	O5'-P-OP1	10.31	123.07	110.70
34	BA	691	A	P-O3'-C3'	-10.30	107.33	119.70
35	BB	490	G	P-O5'-C5'	-10.30	104.42	120.90
85	AA	2220	U	P-O3'-C3'	-10.30	107.33	119.70
85	AA	730	G	C8-N9-C1'	10.30	140.39	127.00
34	BA	1416	C	C5'-C4'-C3'	10.30	132.48	116.00
85	AA	2014	G	C5-C6-O6	-10.30	122.42	128.60
85	AA	15	U	C2-N3-C4	-10.29	120.82	127.00
85	AA	313	A	C8-N9-C4	10.29	109.92	105.80
85	AA	816	A	O4'-C1'-N9	10.29	116.43	108.20
41	BH	133	U	C2-N3-C4	-10.29	120.83	127.00
85	AA	114	C	O4'-C1'-N1	10.29	116.43	108.20
34	BA	289	A	P-O3'-C3'	10.29	132.05	119.70
34	BA	801	U	P-O3'-C3'	-10.29	107.36	119.70
34	BA	923	C	C3'-C2'-C1'	-10.29	93.27	101.50
41	BH	119	U	C2-N1-C1'	-10.29	105.36	117.70
34	BA	787	A	P-O3'-C3'	10.28	132.04	119.70
34	BA	71	G	C8-N9-C4	-10.28	102.29	106.40
35	BB	530	C	P-O5'-C5'	10.28	137.35	120.90
35	BB	1465	U	P-O5'-C5'	-10.28	104.45	120.90
34	BA	939	C	C6-N1-C2	-10.28	116.19	120.30
34	BA	1455	C	C4'-C3'-C2'	10.28	112.88	102.60
40	BG	16	G	O4'-C1'-N9	10.28	116.42	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	149	G	O4'-C1'-N9	10.28	116.42	108.20
85	AA	1225	C	C2-N1-C1'	10.28	130.10	118.80
34	BA	103	G	C4-N9-C1'	-10.27	113.15	126.50
34	BA	480	G	P-O3'-C3'	-10.27	107.37	119.70
35	BB	1537	C	P-O5'-C5'	10.27	137.34	120.90
85	AA	327	G	O4'-C4'-C3'	-10.27	93.73	104.00
85	AA	1358	A	O4'-C1'-N9	10.27	116.42	108.20
86	AB	70	G	P-O3'-C3'	-10.27	107.38	119.70
34	BA	114	U	C5'-C4'-O4'	-10.27	96.78	109.10
34	BA	1744	C	O4'-C1'-N1	10.27	116.41	108.20
36	BC	6	G	C5-C6-O6	-10.27	122.44	128.60
34	BA	576	C	P-O3'-C3'	10.26	132.02	119.70
35	BB	1515	C	C6-N1-C2	-10.26	116.19	120.30
36	BC	124	A	C5'-C4'-C3'	10.26	132.42	116.00
85	AA	942	A	C1'-O4'-C4'	-10.26	101.69	109.90
85	AA	1909	C	P-O3'-C3'	10.26	132.02	119.70
85	AA	365	G	C4-N9-C1'	-10.26	113.16	126.50
42	BI	159	PHE	CB-CG-CD1	10.26	127.98	120.80
40	BG	157	A	N1-C6-N6	-10.26	112.44	118.60
85	AA	2011	C	O4'-C1'-N1	10.26	116.40	108.20
34	BA	768	G	C5-C6-O6	-10.25	122.45	128.60
85	AA	2066	C	C6-N1-C2	-10.25	116.20	120.30
34	BA	114	U	C2-N1-C1'	-10.25	105.40	117.70
84	By	113	PHE	CB-CG-CD2	-10.25	113.63	120.80
85	AA	2149	C	O4'-C1'-N1	10.25	116.40	108.20
85	AA	1125	G	C6-N1-C2	-10.25	118.95	125.10
85	AA	266	U	P-O3'-C3'	10.24	131.99	119.70
85	AA	370	A	O4'-C1'-N9	10.24	116.39	108.20
34	BA	1574	C	C5'-C4'-C3'	10.24	132.38	116.00
85	AA	788	G	C4'-C3'-C2'	10.24	112.84	102.60
38	BE	83	U	P-O3'-C3'	-10.24	107.42	119.70
85	AA	1799	C	C5-C4-N4	-10.24	113.03	120.20
34	BA	631	G	N3-C2-N2	-10.23	112.74	119.90
34	BA	1736	A	P-O3'-C3'	-10.23	107.42	119.70
35	BB	133	G	P-O3'-C3'	-10.23	107.42	119.70
38	BE	117	A	C5-C6-N6	-10.23	115.51	123.70
34	BA	115	U	C3'-C2'-C1'	-10.23	93.32	101.50
34	BA	1350	C	C6-N1-C2	-10.23	116.21	120.30
85	AA	87	C	C5'-C4'-C3'	-10.23	99.63	116.00
85	AA	1251	G	P-O3'-C3'	-10.23	107.42	119.70
34	BA	871	G	C5-N7-C8	10.23	109.41	104.30
34	BA	1193	A	N1-C6-N6	-10.23	112.46	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	495	G	C8-N9-C1'	10.23	140.29	127.00
38	BE	88	G	C5-C6-O6	-10.22	122.47	128.60
39	BF	32	G	N1-C6-O6	10.22	126.03	119.90
41	BH	50	A	P-O5'-C5'	10.22	137.26	120.90
22	AO	89	TYR	CB-CG-CD2	10.22	127.13	121.00
35	BB	589	U	C3'-C2'-C1'	-10.22	93.32	101.50
34	BA	753	G	C5-C6-O6	-10.22	122.47	128.60
37	BD	99	G	P-O5'-C5'	10.22	137.25	120.90
38	BE	23	G	C1'-O4'-C4'	-10.22	101.73	109.90
41	BH	41	A	C8-N9-C4	10.22	109.89	105.80
85	AA	1096	G	C8-N9-C1'	10.22	140.28	127.00
38	BE	188	C	C1'-O4'-C4'	-10.21	101.73	109.90
34	BA	247	U	C2-N3-C4	-10.21	120.87	127.00
34	BA	570	G	C8-N9-C4	10.21	110.48	106.40
34	BA	658	C	P-O5'-C5'	10.21	137.24	120.90
34	BA	1454	G	C8-N9-C4	10.21	110.48	106.40
35	BB	684	U	C6-N1-C2	-10.21	114.87	121.00
35	BB	1449	G	C5-C6-O6	-10.21	122.47	128.60
34	BA	291	C	O4'-C1'-N1	10.21	116.37	108.20
34	BA	1651	C	P-O3'-C3'	10.21	131.95	119.70
38	BE	160	C	P-O3'-C3'	-10.21	107.45	119.70
40	BG	35	G	P-O5'-C5'	10.21	137.23	120.90
85	AA	1469	G	C1'-O4'-C4'	10.21	118.06	109.90
36	BC	113	G	C5-C6-O6	-10.21	122.48	128.60
34	BA	582	U	C2-N3-C4	-10.20	120.88	127.00
38	BE	30	C	O4'-C1'-N1	10.20	116.36	108.20
85	AA	485	A	C5'-C4'-C3'	-10.21	99.67	116.00
34	BA	370	U	P-O3'-C3'	10.20	131.94	119.70
34	BA	991	U	O4'-C1'-N1	10.20	116.36	108.20
34	BA	1299	G	P-O3'-C3'	10.20	131.94	119.70
35	BB	88	U	C2-N3-C4	-10.20	120.88	127.00
35	BB	500	C	O4'-C1'-N1	10.20	116.36	108.20
35	BB	1347	C	P-O5'-C5'	10.20	137.22	120.90
86	AB	6	G	C8-N9-C1'	10.20	140.26	127.00
85	AA	2155	U	C6-N1-C2	-10.20	114.88	121.00
85	AA	778	C	C5'-C4'-C3'	10.19	132.31	116.00
85	AA	1132	A	P-O3'-C3'	-10.19	107.47	119.70
85	AA	2209	U	C3'-C2'-C1'	-10.19	93.35	101.50
35	BB	1347	C	C6-N1-C2	-10.19	116.22	120.30
34	BA	141	G	N1-C2-N2	10.18	125.37	116.20
34	BA	821	G	C5-C6-O6	-10.18	122.49	128.60
34	BA	440	A	N1-C6-N6	-10.18	112.49	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	765	U	C2-N1-C1'	10.18	129.92	117.70
34	BA	814	C	C6-N1-C1'	-10.18	108.58	120.80
85	AA	227	A	N1-C6-N6	-10.18	112.49	118.60
40	BG	33	G	C2-N3-C4	-10.18	106.81	111.90
34	BA	237	A	O4'-C1'-N9	10.18	116.34	108.20
34	BA	924	U	P-O5'-C5'	10.18	137.19	120.90
41	BH	23	G	C5-C6-O6	-10.18	122.49	128.60
85	AA	854	A	O4'-C1'-N9	10.18	116.34	108.20
34	BA	98	A	N1-C6-N6	10.18	124.71	118.60
34	BA	502	U	P-O5'-C5'	-10.18	104.62	120.90
36	BC	1	A	P-O3'-C3'	10.18	131.91	119.70
40	BG	174	G	C5'-C4'-C3'	10.18	132.28	116.00
85	AA	192	G	O4'-C1'-N9	10.18	116.34	108.20
85	AA	342	C	O4'-C1'-N1	10.18	116.34	108.20
85	AA	640	C	O4'-C1'-N1	10.18	116.34	108.20
85	AA	1209	U	C6-N1-C2	-10.17	114.90	121.00
34	BA	606	G	O3'-P-O5'	10.17	123.33	104.00
85	AA	523	U	P-O3'-C3'	10.17	131.91	119.70
85	AA	1468	G	C5'-C4'-C3'	10.17	132.28	116.00
34	BA	1519	G	N1-C6-O6	-10.17	113.80	119.90
34	BA	1145	U	C5'-C4'-C3'	-10.17	99.73	116.00
35	BB	834	U	C6-N1-C2	-10.17	114.90	121.00
70	Bk	120	ARG	NE-CZ-NH1	10.17	125.39	120.30
37	BD	62	A	P-O3'-C3'	10.17	131.90	119.70
38	BE	30	C	C6-N1-C2	-10.17	116.23	120.30
85	AA	571	G	O4'-C1'-N9	10.17	116.33	108.20
35	BB	560	C	C5'-C4'-C3'	10.16	132.26	116.00
34	BA	177	G	P-O3'-C3'	10.16	131.90	119.70
35	BB	1497	C	P-O5'-C5'	10.16	137.16	120.90
40	BG	24	A	C5-C6-N1	10.16	122.78	117.70
41	BH	126	C	C5'-C4'-C3'	-10.16	99.74	116.00
85	AA	588	G	N1-C6-O6	10.16	126.00	119.90
35	BB	566	A	N1-C6-N6	-10.16	112.50	118.60
41	BH	24	U	C3'-C2'-C1'	-10.16	93.37	101.50
85	AA	112	A	C5'-C4'-C3'	-10.16	99.74	116.00
85	AA	1463	A	C5'-C4'-C3'	-10.16	99.74	116.00
35	BB	1132	A	P-O5'-C5'	10.16	137.15	120.90
35	BB	1166	A	C8-N9-C4	10.16	109.86	105.80
36	BC	169	G	C8-N9-C1'	10.16	140.21	127.00
85	AA	5	U	O4'-C1'-N1	10.16	116.33	108.20
85	AA	268	A	O4'-C1'-N9	10.16	116.33	108.20
85	AA	1524	A	P-O5'-C5'	10.16	137.16	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1004	G	C8-N9-C1'	10.16	140.21	127.00
35	BB	773	G	N1-C6-O6	10.15	125.99	119.90
34	BA	110	C	N1-C1'-C2'	-10.15	100.80	114.00
85	AA	609	U	P-O5'-C5'	-10.15	104.66	120.90
34	BA	1188	U	P-O3'-C3'	10.15	131.88	119.70
85	AA	496	C	C6-N1-C2	-10.15	116.24	120.30
36	BC	7	U	C2-N1-C1'	10.15	129.88	117.70
36	BC	34	U	P-O5'-C5'	-10.15	104.66	120.90
41	BH	72	G	C4'-C3'-O3'	10.15	133.30	113.00
39	BF	22	U	N3-C2-O2	-10.15	115.10	122.20
34	BA	1199	U	O4'-C1'-N1	10.14	116.31	108.20
35	BB	797	C	C5-C4-N4	10.14	127.30	120.20
35	BB	1347	C	P-O3'-C3'	-10.14	107.53	119.70
1	A0	210	ARG	NE-CZ-NH1	-10.14	115.23	120.30
36	BC	17	U	C5'-C4'-C3'	-10.14	99.78	116.00
41	BH	77	G	C5-C6-O6	-10.14	122.52	128.60
85	AA	964	C	P-O3'-C3'	-10.14	107.54	119.70
34	BA	1626	U	C5'-C4'-O4'	-10.14	96.94	109.10
34	BA	1097	G	C5'-C4'-C3'	10.13	132.21	116.00
34	BA	1202	G	C5'-C4'-C3'	-10.13	99.78	116.00
85	AA	282	C	C6-N1-C2	-10.13	116.25	120.30
34	BA	1809	G	C4-N9-C1'	-10.13	113.33	126.50
41	BH	14	C	C6-N1-C1'	-10.13	108.64	120.80
69	Bj	62	HIS	CA-CB-CG	10.13	130.83	113.60
85	AA	909	C	P-O3'-C3'	10.13	131.86	119.70
7	A6	106	ARG	NE-CZ-NH2	-10.13	115.24	120.30
34	BA	175	G	C5'-C4'-O4'	-10.13	96.95	109.10
34	BA	702	G	N9-C1'-C2'	-10.13	100.83	114.00
35	BB	1249	G	P-O3'-C3'	10.13	131.85	119.70
34	BA	222	C	O4'-C1'-N1	10.12	116.30	108.20
37	BD	108	G	C8-N9-C1'	10.12	140.16	127.00
38	BE	89	G	C8-N9-C1'	10.12	140.16	127.00
38	BE	104	G	C4-N9-C1'	10.12	139.66	126.50
85	AA	2149	C	C5'-C4'-C3'	-10.12	99.80	116.00
34	BA	122	U	C6-N1-C2	-10.12	114.93	121.00
34	BA	294	C	C6-N1-C2	-10.12	116.25	120.30
35	BB	4	C	O4'-C1'-N1	10.12	116.30	108.20
69	Bj	89	ARG	NE-CZ-NH1	10.12	125.36	120.30
85	AA	940	G	P-O5'-C5'	-10.12	104.71	120.90
34	BA	539	C	O4'-C1'-N1	10.12	116.30	108.20
34	BA	967	C	P-O3'-C3'	-10.12	107.56	119.70
38	BE	136	G	N1-C6-O6	10.12	125.97	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1661	U	C6-N1-C2	-10.12	114.93	121.00
34	BA	1122	G	C5-C6-O6	-10.11	122.53	128.60
34	BA	1198	U	P-O3'-C3'	10.11	131.84	119.70
34	BA	1339	G	O4'-C1'-N9	10.11	116.29	108.20
34	BA	162	G	C6-C5-N7	-10.11	124.33	130.40
49	BP	67	ARG	NE-CZ-NH1	10.11	125.36	120.30
85	AA	542	G	P-O5'-C5'	10.11	137.08	120.90
37	BD	22	A	P-O3'-C3'	-10.11	107.57	119.70
85	AA	738	C	O4'-C1'-N1	10.11	116.29	108.20
3	A2	67	ARG	NE-CZ-NH1	10.10	125.35	120.30
85	AA	798	A	O4'-C1'-N9	10.10	116.28	108.20
85	AA	1252	A	C4'-C3'-C2'	-10.10	92.50	102.60
38	BE	50	G	P-O3'-C3'	-10.10	107.58	119.70
35	BB	24	C	P-O5'-C5'	10.10	137.05	120.90
35	BB	633	C	C6-N1-C2	-10.10	116.26	120.30
85	AA	65	A	O5'-P-OP1	-10.10	96.61	105.70
34	BA	179	U	O4'-C1'-N1	10.09	116.27	108.20
34	BA	223	U	N1-C2-N3	10.09	120.95	114.90
85	AA	2050	C	O4'-C1'-N1	10.09	116.27	108.20
34	BA	302	A	C5'-C4'-C3'	-10.09	99.86	116.00
34	BA	878	G	C5'-C4'-C3'	10.09	132.14	116.00
34	BA	1747	C	P-O5'-C5'	10.09	137.04	120.90
85	AA	893	G	C2-N3-C4	-10.09	106.86	111.90
85	AA	1575	G	C8-N9-C1'	10.09	140.11	127.00
34	BA	180	G	C5-C6-O6	-10.09	122.55	128.60
36	BC	108	A	O4'-C1'-N9	10.09	116.27	108.20
85	AA	382	G	C5'-C4'-C3'	10.09	132.13	116.00
85	AA	484	G	O4'-C1'-N9	10.08	116.27	108.20
85	AA	1310	G	C5-C6-O6	-10.08	122.55	128.60
85	AA	1930	U	P-O3'-C3'	10.08	131.80	119.70
35	BB	41	A	C5-C6-N6	-10.08	115.64	123.70
38	BE	178	G	C4-N9-C1'	-10.08	113.40	126.50
34	BA	546	U	C5'-C4'-C3'	-10.08	99.88	116.00
35	BB	1505	U	O4'-C1'-N1	10.07	116.26	108.20
34	BA	1782	C	O4'-C1'-N1	10.07	116.26	108.20
35	BB	363	A	N1-C6-N6	-10.07	112.56	118.60
35	BB	1345	A	P-O3'-C3'	10.07	131.78	119.70
41	BH	105	U	C2-N3-C4	-10.07	120.96	127.00
34	BA	584	A	P-O3'-C3'	10.07	131.78	119.70
35	BB	560	C	O4'-C1'-C2'	10.07	116.66	107.60
41	BH	73	A	OP1-P-OP2	-10.07	104.50	119.60
41	BH	128	G	N9-C4-C5	-10.06	101.37	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	259	C	C6-N1-C1'	10.06	132.88	120.80
34	BA	1477	C	N1-C1'-C2'	-10.06	100.92	114.00
34	BA	177	G	C4-N9-C1'	-10.06	113.42	126.50
34	BA	1613	G	C5-C6-O6	-10.06	122.56	128.60
39	BF	70	A	P-O5'-C5'	10.06	137.00	120.90
41	BH	10	U	C6-N1-C2	-10.06	114.96	121.00
45	BL	140	ARG	NE-CZ-NH1	10.06	125.33	120.30
85	AA	928	U	C2-N3-C4	-10.06	120.96	127.00
34	BA	594	G	C4-N9-C1'	10.06	139.57	126.50
34	BA	665	C	C2-N1-C1'	-10.06	107.74	118.80
34	BA	1846	G	C5'-C4'-C3'	-10.06	99.91	116.00
38	BE	111	C	O4'-C1'-N1	10.06	116.25	108.20
40	BG	29	U	C2-N3-C4	-10.06	120.97	127.00
85	AA	553	G	C8-N9-C4	-10.06	102.38	106.40
85	AA	755	G	O4'-C1'-N9	10.06	116.25	108.20
29	AV	68	MET	CG-SD-CE	-10.05	84.11	100.20
36	BC	107	C	P-O3'-C3'	-10.05	107.64	119.70
68	Bi	129	ARG	NE-CZ-NH1	10.05	125.33	120.30
34	BA	1326	U	P-O3'-C3'	10.05	131.76	119.70
40	BG	14	G	C4-N9-C1'	-10.05	113.43	126.50
85	AA	1854	U	C2-N3-C4	-10.05	120.97	127.00
34	BA	875	G	C5-C6-N1	10.05	116.52	111.50
35	BB	996	G	N9-C1'-C2'	-10.05	100.94	114.00
85	AA	2123	U	C6-N1-C1'	10.05	135.27	121.20
85	AA	96	C	C4'-C3'-C2'	10.05	112.65	102.60
85	AA	1106	A	O4'-C1'-N9	10.05	116.24	108.20
34	BA	258	C	C6-N1-C2	-10.04	116.28	120.30
38	BE	153	C	C6-N1-C2	-10.04	116.28	120.30
85	AA	1599	G	C5'-C4'-C3'	-10.04	99.93	116.00
82	Bw	82	ARG	NE-CZ-NH1	10.04	125.32	120.30
85	AA	1096	G	C5'-C4'-C3'	-10.04	99.94	116.00
85	AA	2198	G	C5'-C4'-C3'	-10.04	99.94	116.00
34	BA	1496	G	C5'-C4'-C3'	-10.04	99.94	116.00
85	AA	798	A	P-O3'-C3'	-10.04	107.66	119.70
85	AA	1490	A	C1'-O4'-C4'	-10.04	101.87	109.90
34	BA	653	U	C6-N1-C2	-10.04	114.98	121.00
35	BB	1187	G	C6-N1-C2	-10.04	119.08	125.10
39	BF	37	C	P-O3'-C3'	10.04	131.74	119.70
85	AA	2225	G	N1-C6-O6	10.04	125.92	119.90
34	BA	1033	G	N1-C6-O6	10.03	125.92	119.90
38	BE	113	C	P-O5'-C5'	10.03	136.95	120.90
85	AA	842	G	P-O3'-C3'	10.03	131.74	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1237	A	C1'-O4'-C4'	-10.03	101.88	109.90
86	AB	65	G	N1-C6-O6	10.03	125.92	119.90
35	BB	642	G	N1-C6-O6	-10.03	113.88	119.90
38	BE	208	G	O4'-C1'-N9	10.03	116.22	108.20
85	AA	597	A	C5'-C4'-C3'	-10.03	99.95	116.00
65	Bf	278	LYS	N-CA-CB	-10.03	92.55	110.60
85	AA	581	A	P-O3'-C3'	-10.03	107.67	119.70
85	AA	2157	G	C8-N9-C1'	10.02	140.03	127.00
85	AA	1090	A	O4'-C1'-N9	10.02	116.22	108.20
35	BB	871	C	O4'-C1'-N1	10.02	116.22	108.20
35	BB	1220	A	P-O3'-C3'	10.02	131.72	119.70
54	BU	92	ARG	NE-CZ-NH1	10.02	125.31	120.30
85	AA	2010	C	C5'-C4'-C3'	-10.02	99.97	116.00
38	BE	97	G	N1-C6-O6	10.02	125.91	119.90
48	BO	166	TRP	CB-CG-CD2	-10.02	113.58	126.60
35	BB	755	A	N1-C6-N6	-10.01	112.59	118.60
35	BB	1510	G	C6-N1-C2	-10.01	119.09	125.10
38	BE	117	A	N1-C6-N6	10.01	124.61	118.60
41	BH	23	G	P-O3'-C3'	10.01	131.72	119.70
85	AA	368	C	N3-C2-O2	-10.01	114.89	121.90
85	AA	1701	G	C4-N9-C1'	-10.01	113.48	126.50
85	AA	2002	A	C5'-C4'-C3'	10.01	132.02	116.00
34	BA	309	U	C2-N3-C4	-10.01	120.99	127.00
34	BA	1620	U	C5'-C4'-C3'	-10.01	99.98	116.00
85	AA	691	U	O3'-P-O5'	10.01	123.02	104.00
34	BA	1536	A	O4'-C1'-N9	10.01	116.20	108.20
34	BA	1580	U	P-O3'-C3'	-10.01	107.69	119.70
38	BE	180	G	P-O3'-C3'	10.01	131.71	119.70
85	AA	1268	C	C5'-C4'-C3'	-10.01	99.99	116.00
35	BB	993	A	C2-N3-C4	-10.01	105.60	110.60
38	BE	171	U	P-O3'-C3'	10.01	131.71	119.70
85	AA	1682	U	P-O3'-C3'	10.00	131.70	119.70
85	AA	1239	C	C5'-C4'-C3'	-10.00	100.00	116.00
85	AA	2055	G	C5-C6-O6	-10.00	122.60	128.60
85	AA	2058	C	C1'-O4'-C4'	-10.00	101.90	109.90
85	AA	1517	G	C5-C6-O6	-10.00	122.60	128.60
35	BB	797	C	C4'-C3'-C2'	-10.00	92.60	102.60
35	BB	1142	C	P-O3'-C3'	10.00	131.70	119.70
35	BB	1347	C	O4'-C1'-N1	10.00	116.20	108.20
34	BA	764	G	N3-C2-N2	10.00	126.90	119.90
35	BB	429	C	C6-N1-C2	-10.00	116.30	120.30
34	BA	1306	U	O4'-C1'-N1	10.00	116.20	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	469	C	C1'-O4'-C4'	-9.99	101.91	109.90
34	BA	1518	A	N9-C1'-C2'	-9.99	101.01	112.00
38	BE	129	G	C4'-C3'-C2'	-9.99	92.61	102.60
85	AA	473	C	O4'-C1'-N1	9.99	116.20	108.20
85	AA	662	U	N1-C1'-C2'	-9.99	101.01	112.00
34	BA	280	A	C8-N9-C4	-9.99	101.80	105.80
34	BA	741	A	C4'-C3'-C2'	-9.99	92.61	102.60
85	AA	1162	A	O4'-C1'-N9	9.99	116.19	108.20
85	AA	1819	U	P-O5'-C5'	9.99	136.88	120.90
34	BA	7	U	O3'-P-O5'	9.99	122.98	104.00
85	AA	386	G	O4'-C1'-N9	9.99	116.19	108.20
34	BA	1412	G	C5-C6-O6	9.99	134.59	128.60
34	BA	1846	G	C5-C6-O6	-9.99	122.61	128.60
35	BB	1513	U	C6-N1-C2	-9.99	115.01	121.00
85	AA	480	U	C2-N1-C1'	-9.99	105.72	117.70
85	AA	600	C	O4'-C1'-N1	9.99	116.19	108.20
33	AZ	97	ARG	NE-CZ-NH1	9.98	125.29	120.30
34	BA	517	A	C8-N9-C4	9.98	109.79	105.80
85	AA	988	C	O4'-C1'-N1	9.98	116.19	108.20
34	BA	261	A	O4'-C1'-N9	9.98	116.19	108.20
85	AA	295	U	P-O3'-C3'	9.98	131.68	119.70
85	AA	1000	U	C2-N1-C1'	-9.98	105.72	117.70
86	AB	3	C	C5'-C4'-O4'	9.98	121.08	109.10
34	BA	382	G	N3-C4-C5	-9.98	123.61	128.60
34	BA	1441	C	O4'-C1'-N1	9.98	116.18	108.20
85	AA	1882	U	P-O3'-C3'	9.98	131.68	119.70
34	BA	1657	A	C5'-C4'-C3'	9.98	131.96	116.00
78	Bs	35	ARG	NE-CZ-NH1	9.97	125.29	120.30
85	AA	251	A	P-O5'-C5'	9.97	136.86	120.90
85	AA	976	G	C8-N9-C1'	9.97	139.97	127.00
34	BA	248	G	P-O3'-C3'	-9.97	107.73	119.70
34	BA	575	U	P-O3'-C3'	9.97	131.66	119.70
34	BA	1469	G	C5-C6-O6	-9.97	122.62	128.60
36	BC	138	C	O4'-C1'-N1	9.97	116.18	108.20
39	BF	11	C	C2-N1-C1'	9.97	129.77	118.80
35	BB	540	G	C6-C5-N7	-9.97	124.42	130.40
34	BA	279	U	P-O5'-C5'	-9.96	104.96	120.90
34	BA	979	G	N1-C6-O6	9.96	125.88	119.90
34	BA	1321	A	C5'-C4'-C3'	-9.96	100.06	116.00
35	BB	25	A	C5-C6-N6	9.97	131.67	123.70
85	AA	289	G	C5-C6-O6	-9.97	122.62	128.60
34	BA	606	G	N3-C2-N2	9.96	126.87	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1711	G	P-O3'-C3'	-9.96	107.74	119.70
48	BO	136	ARG	NE-CZ-NH2	9.96	125.28	120.30
34	BA	1161	G	C5'-C4'-C3'	9.96	131.94	116.00
34	BA	1460	U	P-O5'-C5'	9.96	136.84	120.90
39	BF	9	C	O4'-C1'-N1	9.96	116.17	108.20
39	BF	39	C	O3'-P-O5'	-9.96	85.07	104.00
34	BA	1799	G	P-O3'-C3'	-9.96	107.75	119.70
38	BE	96	G	C2-N3-C4	-9.96	106.92	111.90
35	BB	1048	A	P-O3'-C3'	-9.96	107.75	119.70
35	BB	1459	U	C6-N1-C2	-9.96	115.03	121.00
36	BC	78	G	N1-C6-O6	-9.96	113.92	119.90
85	AA	4	C	C2-N3-C4	-9.96	114.92	119.90
85	AA	428	G	C5-C6-O6	-9.96	122.62	128.60
85	AA	833	U	O4'-C1'-N1	9.96	116.17	108.20
34	BA	1625	C	O4'-C1'-N1	9.96	116.17	108.20
85	AA	965	G	P-O5'-C5'	9.96	136.83	120.90
34	BA	1540	C	C2-N3-C4	-9.95	114.92	119.90
35	BB	39	C	C1'-O4'-C4'	-9.95	101.94	109.90
35	BB	1251	G	N1-C6-O6	-9.95	113.93	119.90
85	AA	1225	C	P-O5'-C5'	-9.95	104.98	120.90
38	BE	128	G	O3'-P-O5'	-9.95	85.09	104.00
85	AA	1674	G	C5-C6-O6	-9.95	122.63	128.60
36	BC	130	U	C2-N1-C1'	9.95	129.64	117.70
34	BA	1519	G	P-O3'-C3'	-9.95	107.77	119.70
35	BB	1470	G	N1-C6-O6	9.94	125.87	119.90
37	BD	93	G	C5-C6-O6	9.95	134.57	128.60
85	AA	371	C	C6-N1-C2	-9.95	116.32	120.30
85	AA	586	G	C5-C6-O6	-9.95	122.63	128.60
85	AA	895	C	C5'-C4'-O4'	9.95	121.03	109.10
85	AA	2213	A	P-O3'-C3'	-9.94	107.77	119.70
40	BG	10	U	P-O3'-C3'	-9.94	107.77	119.70
35	BB	687	C	C6-N1-C2	-9.94	116.32	120.30
35	BB	880	G	N7-C8-N9	9.94	118.07	113.10
38	BE	23	G	O4'-C1'-N9	9.94	116.15	108.20
85	AA	251	A	C8-N9-C4	-9.94	101.82	105.80
40	BG	98	A	C5'-C4'-O4'	9.94	121.03	109.10
85	AA	982	G	C6-N1-C2	-9.94	119.14	125.10
85	AA	1525	C	C6-N1-C2	-9.94	116.33	120.30
85	AA	2106	C	O4'-C1'-N1	9.94	116.15	108.20
85	AA	1756	C	C6-N1-C1'	9.94	132.72	120.80
34	BA	331	G	N1-C6-O6	-9.94	113.94	119.90
49	BP	20	ARG	NE-CZ-NH1	9.94	125.27	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
69	Bj	8	TYR	CB-CG-CD1	-9.94	115.04	121.00
85	AA	835	C	O4'-C1'-N1	9.94	116.15	108.20
85	AA	2002	A	C4'-C3'-C2'	9.93	112.53	102.60
34	BA	280	A	C2'-C3'-O3'	9.93	131.35	109.50
85	AA	917	A	C8-N9-C4	9.93	109.77	105.80
50	BQ	55	ARG	NE-CZ-NH2	-9.93	115.33	120.30
85	AA	2133	A	P-O5'-C5'	-9.93	105.02	120.90
34	BA	416	A	N1-C6-N6	-9.93	112.64	118.60
34	BA	919	A	N1-C6-N6	-9.93	112.64	118.60
35	BB	2	C	C2-N3-C4	-9.93	114.94	119.90
35	BB	379	U	C1'-O4'-C4'	-9.93	101.96	109.90
35	BB	1548	C	O4'-C1'-N1	9.93	116.14	108.20
85	AA	2237	G	C4'-C3'-C2'	9.93	112.53	102.60
34	BA	159	U	O4'-C1'-N1	9.92	116.14	108.20
35	BB	29	C	O4'-C1'-N1	9.92	116.14	108.20
35	BB	1069	C	O4'-C1'-N1	9.92	116.14	108.20
34	BA	20	A	O3'-P-O5'	-9.92	85.15	104.00
35	BB	991	C	C5'-C4'-C3'	9.92	131.87	116.00
34	BA	1477	C	P-O5'-C5'	-9.91	105.04	120.90
85	AA	1718	C	O4'-C1'-N1	9.91	116.13	108.20
34	BA	212	A	N1-C6-N6	-9.91	112.65	118.60
35	BB	41	A	P-O5'-C5'	-9.91	105.05	120.90
35	BB	295	U	P-O3'-C3'	9.91	131.59	119.70
85	AA	2123	U	N1-C2-N3	9.91	120.84	114.90
85	AA	605	A	C5-C6-N6	-9.91	115.78	123.70
85	AA	1139	G	O4'-C1'-N9	9.91	116.12	108.20
35	BB	823	G	C6-N1-C2	-9.90	119.16	125.10
40	BG	27	C	C5'-C4'-C3'	-9.90	100.15	116.00
34	BA	1776	G	P-O5'-C5'	9.90	136.74	120.90
35	BB	878	G	O4'-C4'-C3'	-9.90	94.10	104.00
35	BB	1334	C	O4'-C1'-N1	9.90	116.12	108.20
85	AA	100	A	C5'-C4'-C3'	-9.90	100.16	116.00
85	AA	526	G	C5-C6-O6	-9.90	122.66	128.60
34	BA	42	A	N1-C6-N6	-9.90	112.66	118.60
34	BA	751	A	P-O3'-C3'	9.90	131.58	119.70
34	BA	806	U	O4'-C1'-N1	9.90	116.12	108.20
34	BA	1175	G	O3'-P-O5'	-9.90	85.19	104.00
35	BB	1168	G	C4-N9-C1'	-9.90	113.63	126.50
34	BA	903	C	C2-N3-C4	-9.90	114.95	119.90
34	BA	1277	G	C8-N9-C1'	9.90	139.87	127.00
35	BB	638	G	C4'-C3'-C2'	9.90	112.50	102.60
38	BE	16	C	P-O3'-C3'	-9.90	107.83	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
83	Bx	234	ARG	NE-CZ-NH1	9.90	125.25	120.30
85	AA	273	C	C6-N1-C1'	9.90	132.68	120.80
34	BA	134	U	P-O3'-C3'	-9.89	107.83	119.70
34	BA	1591	G	C4-N9-C1'	-9.89	113.64	126.50
34	BA	194	G	P-O3'-C3'	-9.89	107.83	119.70
35	BB	618	U	C2-N3-C4	-9.89	121.06	127.00
35	BB	1289	G	C5-C6-O6	-9.89	122.67	128.60
38	BE	130	G	N3-C4-C5	-9.89	123.66	128.60
41	BH	128	G	P-O5'-C5'	-9.89	105.07	120.90
35	BB	726	A	O4'-C1'-N9	9.89	116.11	108.20
40	BG	16	G	C5'-C4'-C3'	9.89	131.82	116.00
34	BA	674	G	C5'-C4'-O4'	9.89	120.96	109.10
35	BB	545	C	C2-N1-C1'	9.89	129.67	118.80
38	BE	57	U	P-O3'-C3'	-9.89	107.84	119.70
85	AA	533	C	O4'-C1'-N1	9.89	116.11	108.20
17	AI	130	TYR	CB-CG-CD1	9.88	126.93	121.00
34	BA	585	G	C5-C6-O6	-9.88	122.67	128.60
34	BA	1192	A	O3'-P-O5'	-9.88	85.22	104.00
35	BB	609	G	C4-N9-C1'	-9.88	113.65	126.50
35	BB	993	A	C5'-C4'-C3'	-9.88	100.19	116.00
40	BG	33	G	N3-C2-N2	-9.88	112.98	119.90
35	BB	1475	U	C5'-C4'-C3'	-9.88	100.19	116.00
35	BB	10	C	C6-N1-C2	-9.88	116.35	120.30
40	BG	8	U	N3-C2-O2	-9.88	115.29	122.20
85	AA	72	C	C6-N1-C2	-9.88	116.35	120.30
34	BA	184	C	C5'-C4'-C3'	-9.88	100.20	116.00
34	BA	1197	U	C6-N1-C1'	9.88	135.03	121.20
35	BB	618	U	C4'-C3'-C2'	9.88	112.47	102.60
40	BG	16	G	C4'-C3'-C2'	9.88	112.48	102.60
85	AA	775	C	P-O5'-C5'	9.88	136.70	120.90
35	BB	851	U	O4'-C1'-N1	9.87	116.10	108.20
35	BB	1323	U	C4-C5-C6	-9.87	113.78	119.70
35	BB	1528	U	C2-N3-C4	-9.88	121.08	127.00
85	AA	1832	G	C5'-C4'-C3'	-9.88	100.20	116.00
85	AA	786	G	P-O3'-C3'	9.87	131.55	119.70
34	BA	1249	G	C5-C6-N1	9.87	116.44	111.50
34	BA	1413	G	C5-C6-O6	-9.87	122.68	128.60
35	BB	994	A	O4'-C1'-N9	9.87	116.10	108.20
40	BG	33	G	C5'-C4'-C3'	9.87	131.80	116.00
34	BA	1707	C	N3-C4-N4	-9.87	111.09	118.00
39	BF	17	U	C2-N3-C4	-9.87	121.08	127.00
85	AA	461	G	N1-C6-O6	9.87	125.82	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	AM	33	MET	CG-SD-CE	-9.87	84.42	100.20
34	BA	557	U	P-O5'-C5'	9.87	136.68	120.90
34	BA	1200	U	C1'-O4'-C4'	-9.87	102.01	109.90
41	BH	114	G	C5-C6-O6	-9.86	122.68	128.60
34	BA	1532	G	P-O5'-C5'	9.86	136.68	120.90
36	BC	26	U	N3-C2-O2	9.86	129.10	122.20
85	AA	73	A	P-O3'-C3'	9.86	131.53	119.70
34	BA	951	C	O4'-C1'-N1	9.86	116.09	108.20
35	BB	461	U	C2-N3-C4	-9.86	121.08	127.00
35	BB	1083	C	C2-N1-C1'	9.86	129.65	118.80
85	AA	1170	C	O4'-C1'-N1	9.86	116.09	108.20
4	A3	96	ARG	NE-CZ-NH1	9.86	125.23	120.30
34	BA	980	C	C2-N3-C4	-9.86	114.97	119.90
35	BB	1091	C	P-O3'-C3'	-9.86	107.87	119.70
85	AA	1106	A	P-O3'-C3'	9.86	131.53	119.70
35	BB	1401	G	N1-C6-O6	9.86	125.81	119.90
51	BR	155	GLU	N-CA-CB	9.86	128.34	110.60
67	Bh	167	TYR	CB-CG-CD1	9.86	126.91	121.00
85	AA	2123	U	C6-N1-C2	-9.86	115.09	121.00
34	BA	690	G	C8-N9-C1'	9.85	139.81	127.00
35	BB	1458	U	C1'-O4'-C4'	-9.85	102.02	109.90
85	AA	1006	C	C6-N1-C1'	9.85	132.62	120.80
85	AA	2209	U	C1'-O4'-C4'	-9.85	102.02	109.90
34	BA	80	U	C2-N1-C1'	-9.85	105.88	117.70
34	BA	1809	G	C6-N1-C2	-9.85	119.19	125.10
35	BB	609	G	C8-N9-C1'	9.85	139.80	127.00
34	BA	360	C	C6-N1-C2	-9.85	116.36	120.30
36	BC	10	C	C5'-C4'-C3'	9.85	131.75	116.00
34	BA	56	G	C5-C6-O6	-9.84	122.69	128.60
34	BA	1581	G	O3'-P-O5'	9.84	122.70	104.00
36	BC	156	A	C4'-C3'-C2'	9.84	112.44	102.60
38	BE	203	C	P-O3'-C3'	9.84	131.51	119.70
34	BA	758	G	C4'-C3'-C2'	-9.84	92.76	102.60
38	BE	130	G	P-O5'-C5'	-9.84	105.16	120.90
26	AS	131	TYR	CB-CG-CD1	9.84	126.90	121.00
37	BD	67	C	C3'-C2'-C1'	-9.84	93.63	101.50
85	AA	635	G	C4-N9-C1'	-9.84	113.71	126.50
39	BF	49	C	C6-N1-C2	-9.84	116.37	120.30
35	BB	41	A	C4-C5-C6	-9.83	112.08	117.00
85	AA	1485	G	P-O3'-C3'	9.83	131.50	119.70
34	BA	979	G	C5-C6-O6	-9.83	122.70	128.60
34	BA	1818	A	O5'-C5'-C4'	9.83	130.38	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1721	U	C4'-C3'-C2'	-9.83	92.77	102.60
41	BH	93	G	O4'-C1'-N9	9.83	116.06	108.20
85	AA	149	A	C5'-C4'-C3'	9.83	131.73	116.00
85	AA	1797	U	P-O3'-C3'	-9.83	107.91	119.70
34	BA	547	C	O5'-C5'-C4'	9.83	130.37	111.70
39	BF	35	C	C6-N1-C2	-9.83	116.37	120.30
34	BA	687	G	P-O5'-C5'	-9.82	105.18	120.90
85	AA	2061	C	O4'-C1'-N1	9.82	116.06	108.20
34	BA	1747	C	C2-N1-C1'	-9.82	108.00	118.80
38	BE	210	G	P-O5'-C5'	9.82	136.62	120.90
65	Bf	339	TYR	CB-CG-CD1	9.82	126.89	121.00
35	BB	1030	U	O4'-C1'-N1	9.82	116.06	108.20
37	BD	26	C	C5'-C4'-C3'	-9.82	100.29	116.00
47	BN	31	PHE	CB-CG-CD2	-9.82	113.92	120.80
85	AA	787	U	C2-N1-C1'	9.82	129.49	117.70
85	AA	853	G	P-O3'-C3'	-9.82	107.91	119.70
85	AA	886	A	P-O3'-C3'	-9.82	107.91	119.70
85	AA	1459	C	P-O5'-C5'	9.82	136.62	120.90
85	AA	2001	C	C2-N1-C1'	9.82	129.60	118.80
85	AA	736	U	C3'-C2'-C1'	-9.82	93.64	101.50
85	AA	856	G	O4'-C1'-N9	9.82	116.06	108.20
34	BA	501	U	C3'-C2'-C1'	9.82	109.35	101.50
35	BB	971	A	C3'-C2'-C1'	-9.82	93.65	101.50
36	BC	39	G	C5-C6-O6	-9.82	122.71	128.60
85	AA	1451	U	C2-N3-C4	-9.81	121.11	127.00
35	BB	91	G	C5-C6-O6	-9.81	122.71	128.60
34	BA	1658	G	C5-C6-O6	9.81	134.49	128.60
34	BA	1472	G	N1-C6-O6	-9.81	114.02	119.90
85	AA	965	G	C1'-O4'-C4'	-9.81	102.05	109.90
85	AA	1002	G	N1-C6-O6	9.81	125.79	119.90
34	BA	401	A	O5'-C5'-C4'	-9.80	93.07	111.70
34	BA	869	C	C2-N3-C4	-9.81	115.00	119.90
34	BA	165	C	C5'-C4'-O4'	9.80	120.86	109.10
34	BA	166	G	N9-C1'-C2'	-9.80	101.22	112.00
35	BB	1391	G	C5'-C4'-C3'	-9.80	100.31	116.00
59	BZ	71	TYR	CB-CG-CD2	-9.80	115.12	121.00
35	BB	667	G	P-O5'-C5'	9.80	136.58	120.90
85	AA	607	U	P-O5'-C5'	9.80	136.58	120.90
85	AA	2062	U	C6-N1-C2	-9.80	115.12	121.00
34	BA	214	A	C4-C5-N7	-9.80	105.80	110.70
37	BD	81	C	C1'-O4'-C4'	-9.80	102.06	109.90
34	BA	1539	A	O4'-C1'-N9	9.80	116.04	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	8	G	O4'-C1'-N9	9.80	116.04	108.20
34	BA	574	U	C6-N1-C2	-9.79	115.12	121.00
35	BB	134	G	C5'-C4'-C3'	9.79	131.67	116.00
38	BE	207	G	P-O3'-C3'	9.79	131.46	119.70
40	BG	59	G	C5-C6-O6	-9.80	122.72	128.60
85	AA	1896	G	C5-C6-O6	-9.79	122.72	128.60
85	AA	2074	G	O4'-C1'-N9	9.80	116.04	108.20
34	BA	214	A	C4-N9-C1'	-9.79	108.67	126.30
35	BB	831	C	C5'-C4'-O4'	9.79	120.85	109.10
34	BA	556	A	O5'-P-OP2	9.79	122.45	110.70
34	BA	931	G	N9-C1'-C2'	-9.79	101.23	112.00
35	BB	1026	G	C6-C5-N7	-9.79	124.53	130.40
35	BB	1224	C	O4'-C1'-N1	9.79	116.03	108.20
36	BC	145	G	N1-C6-O6	-9.79	114.02	119.90
85	AA	1074	U	O4'-C1'-N1	9.79	116.03	108.20
85	AA	1290	G	C5-C6-O6	-9.79	122.72	128.60
34	BA	367	G	C5-C6-O6	-9.79	122.73	128.60
34	BA	1538	G	C8-N9-C4	-9.79	102.48	106.40
85	AA	2208	G	C3'-C2'-C1'	-9.79	93.67	101.50
53	BT	97	ARG	NE-CZ-NH1	9.79	125.19	120.30
34	BA	237	A	P-O5'-C5'	-9.79	105.24	120.90
34	BA	631	G	C5-C6-O6	-9.79	122.73	128.60
36	BC	160	C	C2-N1-C1'	-9.79	108.04	118.80
39	BF	33	C	P-O5'-C5'	-9.79	105.25	120.90
34	BA	118	C	P-O5'-C5'	9.78	136.55	120.90
35	BB	836	U	P-O5'-C5'	-9.78	105.25	120.90
35	BB	1404	A	C5'-C4'-O4'	9.78	120.84	109.10
35	BB	22	A	C5-C6-N6	9.78	131.53	123.70
35	BB	1218	G	C8-N9-C1'	-9.78	114.28	127.00
41	BH	6	U	C2-N3-C4	-9.78	121.13	127.00
85	AA	84	C	C3'-C2'-C1'	-9.78	93.67	101.50
85	AA	744	C	N3-C4-C5	-9.78	117.99	121.90
34	BA	694	G	O4'-C1'-N9	9.78	116.02	108.20
40	BG	156	G	C5-C6-O6	-9.78	122.73	128.60
35	BB	448	G	C5'-C4'-C3'	-9.78	100.36	116.00
52	BS	151	PHE	CB-CG-CD1	9.78	127.64	120.80
85	AA	970	U	P-O5'-C5'	9.78	136.55	120.90
34	BA	113	G	P-O3'-C3'	-9.78	107.97	119.70
34	BA	596	G	C4-N9-C1'	9.78	139.21	126.50
85	AA	288	G	P-O5'-C5'	9.78	136.54	120.90
85	AA	2189	U	P-O3'-C3'	-9.78	107.97	119.70
34	BA	52	G	C5-C6-N1	9.77	116.39	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1122	G	N1-C6-O6	9.77	125.77	119.90
85	AA	2022	A	C5-C6-N6	9.77	131.52	123.70
35	BB	776	U	P-O5'-C5'	-9.77	105.27	120.90
35	BB	1488	G	C8-N9-C1'	9.77	139.70	127.00
42	BI	112	PHE	CB-CG-CD1	9.77	127.64	120.80
85	AA	2104	C	P-O3'-C3'	9.77	131.43	119.70
85	AA	727	U	C2-N3-C4	-9.77	121.14	127.00
38	BE	70	C	O4'-C1'-N1	9.77	116.01	108.20
85	AA	890	U	C2-N3-C4	-9.77	121.14	127.00
85	AA	2143	U	O4'-C1'-N1	9.77	116.02	108.20
34	BA	1472	G	C4-N9-C1'	-9.77	113.80	126.50
85	AA	1225	C	C6-N1-C1'	-9.77	109.08	120.80
85	AA	1268	C	O4'-C1'-N1	9.77	116.01	108.20
85	AA	2141	G	C5-C6-O6	-9.77	122.74	128.60
85	AA	963	U	C2-N3-C4	-9.76	121.14	127.00
85	AA	1133	C	C6-N1-C2	-9.76	116.39	120.30
34	BA	798	G	O4'-C4'-C3'	-9.76	94.24	104.00
85	AA	648	G	C8-N9-C1'	9.76	139.69	127.00
37	BD	13	A	C5'-C4'-C3'	-9.76	100.39	116.00
38	BE	195	G	N3-C4-C5	-9.76	123.72	128.60
34	BA	1412	G	O3'-P-O5'	-9.75	85.47	104.00
85	AA	1821	C	O4'-C1'-N1	9.75	116.00	108.20
85	AA	2211	G	C4-N9-C1'	-9.75	113.82	126.50
35	BB	56	U	N1-C2-N3	9.75	120.75	114.90
38	BE	36	U	C2-N3-C4	-9.75	121.15	127.00
38	BE	111	C	C6-N1-C1'	-9.75	109.10	120.80
85	AA	1246	G	P-O3'-C3'	9.75	131.40	119.70
34	BA	1296	U	C4-C5-C6	-9.75	113.85	119.70
35	BB	841	U	P-O3'-C3'	-9.75	108.00	119.70
85	AA	101	C	O4'-C1'-N1	9.75	116.00	108.20
85	AA	940	G	C5-C6-O6	-9.75	122.75	128.60
35	BB	700	C	P-O5'-C5'	9.74	136.49	120.90
85	AA	509	C	O3'-P-O5'	9.74	122.52	104.00
34	BA	816	G	C5-C6-O6	-9.74	122.75	128.60
1	A0	202	ARG	NE-CZ-NH2	-9.74	115.43	120.30
34	BA	162	G	C2-N3-C4	-9.74	107.03	111.90
35	BB	817	C	C5'-C4'-O4'	9.74	120.79	109.10
35	BB	842	G	C4-N9-C1'	-9.74	113.84	126.50
38	BE	36	U	C5'-C4'-C3'	9.74	131.59	116.00
40	BG	8	U	P-O3'-C3'	-9.74	108.01	119.70
40	BG	134	U	O4'-C1'-N1	9.74	115.99	108.20
41	BH	130	G	P-O5'-C5'	-9.74	105.31	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	765	U	P-O5'-C5'	9.74	136.48	120.90
85	AA	1283	C	C2-N3-C4	-9.74	115.03	119.90
35	BB	690	C	C5'-C4'-C3'	-9.74	100.42	116.00
34	BA	1609	U	C2-N3-C4	-9.74	121.16	127.00
34	BA	1637	G	C8-N9-C4	-9.74	102.51	106.40
39	BF	48	G	C5-C6-O6	-9.74	122.76	128.60
41	BH	31	A	C4'-C3'-C2'	9.74	112.34	102.60
41	BH	96	G	C5-C6-O6	-9.74	122.76	128.60
85	AA	357	C	C5'-C4'-C3'	-9.74	100.42	116.00
85	AA	533	C	N3-C4-N4	9.74	124.82	118.00
35	BB	798	A	P-O5'-C5'	-9.73	105.32	120.90
85	AA	688	C	C1'-O4'-C4'	-9.73	102.11	109.90
18	AJ	28	ARG	NE-CZ-NH1	9.73	125.17	120.30
34	BA	1259	C	C6-N1-C2	-9.73	116.41	120.30
35	BB	768	A	C2'-C3'-O3'	9.73	130.91	109.50
55	BV	92	TYR	CB-CG-CD2	-9.73	115.16	121.00
58	BY	60	ARG	NE-CZ-NH1	9.73	125.17	120.30
4	A3	85	PHE	CB-CG-CD2	-9.73	113.99	120.80
34	BA	1501	U	N3-C2-O2	-9.73	115.39	122.20
35	BB	4	C	C6-N1-C2	-9.73	116.41	120.30
35	BB	628	A	N1-C6-N6	-9.73	112.76	118.60
67	Bh	1	MET	CG-SD-CE	-9.73	84.63	100.20
85	AA	1134	G	C6-N1-C2	-9.73	119.26	125.10
34	BA	508	C	O4'-C1'-N1	9.73	115.98	108.20
63	Bd	62	ARG	NE-CZ-NH1	9.73	125.16	120.30
34	BA	679	U	C5-C4-O4	-9.73	120.06	125.90
38	BE	26	G	C5-C6-O6	-9.73	122.76	128.60
41	BH	92	A	O4'-C1'-N9	9.73	115.98	108.20
85	AA	2153	G	C4'-C3'-C2'	-9.73	92.87	102.60
34	BA	1175	G	P-O5'-C5'	9.72	136.46	120.90
34	BA	564	C	C2-N1-C1'	9.72	129.50	118.80
34	BA	399	G	C8-N9-C4	9.72	110.29	106.40
34	BA	543	A	P-O3'-C3'	-9.72	108.03	119.70
34	BA	743	A	C6-N1-C2	9.72	124.43	118.60
38	BE	26	G	C5'-C4'-O4'	-9.72	97.43	109.10
34	BA	542	A	P-O3'-C3'	-9.72	108.04	119.70
34	BA	1801	G	C4-N9-C1'	-9.72	113.86	126.50
35	BB	29	C	C1'-O4'-C4'	-9.72	102.12	109.90
35	BB	872	A	C1'-O4'-C4'	-9.72	102.12	109.90
35	BB	833	G	C5'-C4'-C3'	-9.72	100.45	116.00
37	BD	81	C	N3-C2-O2	-9.72	115.10	121.90
85	AA	863	C	N3-C2-O2	-9.72	115.10	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2141	G	C4-N9-C1'	-9.72	113.87	126.50
85	AA	2156	C	C6-N1-C2	-9.72	116.41	120.30
34	BA	1693	U	P-O3'-C3'	9.71	131.35	119.70
35	BB	1279	C	O4'-C1'-N1	9.71	115.97	108.20
85	AA	207	G	C8-N9-C4	-9.71	102.52	106.40
85	AA	1058	G	C5-C6-O6	-9.71	122.77	128.60
85	AA	1329	U	P-O3'-C3'	9.71	131.35	119.70
85	AA	1883	C	C5'-C4'-C3'	-9.71	100.46	116.00
35	BB	833	G	C4'-C3'-C2'	-9.71	92.89	102.60
34	BA	501	U	C6-N1-C2	-9.71	115.18	121.00
85	AA	865	G	C5-C6-O6	-9.71	122.78	128.60
34	BA	1221	A	N1-C6-N6	-9.71	112.78	118.60
35	BB	1543	C	C6-N1-C2	-9.71	116.42	120.30
38	BE	101	C	C6-N1-C2	-9.71	116.42	120.30
85	AA	364	C	O4'-C1'-N1	9.71	115.96	108.20
85	AA	743	C	O4'-C4'-C3'	-9.71	94.30	104.00
34	BA	12	G	C5'-C4'-C3'	9.70	131.53	116.00
34	BA	729	C	C4'-C3'-C2'	9.70	112.30	102.60
35	BB	1492	C	C6-N1-C2	-9.70	116.42	120.30
35	BB	1539	C	C2-N1-C1'	-9.70	108.13	118.80
37	BD	48	G	N7-C8-N9	-9.70	108.25	113.10
85	AA	326	C	C2-N1-C1'	9.70	129.47	118.80
36	BC	145	G	C5'-C4'-C3'	-9.70	100.48	116.00
60	Ba	90	ARG	NE-CZ-NH1	9.70	125.15	120.30
34	BA	643	U	O4'-C1'-N1	9.70	115.96	108.20
35	BB	112	G	C5-C6-O6	-9.70	122.78	128.60
38	BE	116	U	C2-N1-C1'	-9.70	106.06	117.70
38	BE	198	A	C4-C5-C6	-9.70	112.15	117.00
85	AA	585	G	C4-C5-C6	-9.70	112.98	118.80
85	AA	1231	G	P-O3'-C3'	9.70	131.34	119.70
85	AA	1368	G	C8-N9-C4	-9.70	102.52	106.40
23	AP	78	ASP	N-CA-CB	-9.70	93.15	110.60
40	BG	19	C	O4'-C1'-N1	9.70	115.96	108.20
34	BA	30	A	C4-N9-C1'	-9.70	108.85	126.30
85	AA	1247	A	N1-C6-N6	9.70	124.42	118.60
85	AA	1465	C	C1'-O4'-C4'	-9.70	102.14	109.90
34	BA	329	G	C4-N9-C1'	-9.69	113.90	126.50
39	BF	11	C	O5'-C5'-C4'	9.69	130.12	111.70
86	AB	44	G	C5-C6-O6	-9.69	122.78	128.60
85	AA	613	G	O4'-C1'-N9	9.69	115.95	108.20
85	AA	1921	G	O4'-C1'-N9	9.69	115.95	108.20
34	BA	480	G	N1-C6-O6	9.69	125.71	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	966	C	O4'-C1'-N1	9.69	115.95	108.20
36	BC	135	A	O4'-C1'-N9	9.69	115.95	108.20
34	BA	881	C	C2-N1-C1'	-9.69	108.14	118.80
34	BA	1120	U	P-O3'-C3'	-9.69	108.07	119.70
34	BA	1510	C	P-O5'-C5'	-9.69	105.40	120.90
66	Bg	81	TYR	CB-CG-CD2	-9.69	115.19	121.00
85	AA	382	G	C8-N9-C1'	9.69	139.60	127.00
85	AA	1637	C	O4'-C1'-N1	9.69	115.95	108.20
86	AB	6	G	C4-N9-C1'	-9.69	113.91	126.50
34	BA	1432	C	C6-N1-C2	-9.69	116.42	120.30
85	AA	242	G	P-O5'-C5'	-9.69	105.40	120.90
85	AA	20	G	C6-N1-C2	-9.69	119.29	125.10
85	AA	39	A	P-O3'-C3'	-9.69	108.08	119.70
85	AA	436	G	N1-C6-O6	9.69	125.71	119.90
85	AA	1459	C	O4'-C1'-N1	9.69	115.95	108.20
85	AA	2224	U	C5'-C4'-C3'	-9.69	100.50	116.00
34	BA	188	C	C6-N1-C2	-9.68	116.43	120.30
34	BA	424	U	O4'-C1'-N1	9.68	115.94	108.20
34	BA	1220	C	C2-N1-C1'	-9.68	108.15	118.80
38	BE	10	G	P-O3'-C3'	-9.68	108.08	119.70
40	BG	169	A	C8-N9-C4	9.68	109.67	105.80
34	BA	557	U	C3'-C2'-C1'	-9.68	93.76	101.50
34	BA	1060	C	O4'-C1'-N1	9.68	115.94	108.20
85	AA	753	U	C1'-O4'-C4'	-9.68	102.16	109.90
85	AA	1005	C	P-O5'-C5'	9.68	136.38	120.90
85	AA	2020	C	C5'-C4'-C3'	-9.68	100.52	116.00
34	BA	426	A	C5'-C4'-C3'	-9.67	100.52	116.00
34	BA	785	G	O4'-C1'-N9	9.67	115.94	108.20
35	BB	1060	U	C2-N3-C4	-9.67	121.20	127.00
34	BA	290	G	N9-C1'-C2'	-9.67	101.36	112.00
35	BB	811	C	P-O3'-C3'	-9.67	108.09	119.70
85	AA	618	A	P-O3'-C3'	-9.67	108.10	119.70
37	BD	79	G	C6-N1-C2	-9.67	119.30	125.10
34	BA	10	G	N1-C6-O6	-9.67	114.10	119.90
51	BR	126	ARG	NE-CZ-NH1	9.67	125.13	120.30
85	AA	138	C	O4'-C1'-N1	9.67	115.93	108.20
85	AA	1869	U	P-O3'-C3'	-9.67	108.10	119.70
35	BB	380	G	O4'-C1'-N9	9.66	115.93	108.20
85	AA	2131	C	P-O3'-C3'	-9.66	108.10	119.70
85	AA	2210	C	C6-N1-C1'	9.66	132.40	120.80
34	BA	2	A	O4'-C1'-N9	9.66	115.93	108.20
34	BA	662	U	O4'-C1'-N1	9.66	115.93	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	39	C	C6-N1-C2	-9.66	116.44	120.30
85	AA	1248	U	O4'-C1'-N1	9.66	115.93	108.20
34	BA	209	A	O4'-C1'-N9	9.66	115.93	108.20
34	BA	1041	U	C2-N3-C4	-9.66	121.20	127.00
35	BB	561	C	C2-N1-C1'	9.66	129.42	118.80
35	BB	768	A	C4'-C3'-C2'	-9.66	92.94	102.60
85	AA	639	C	C5'-C4'-C3'	9.66	131.46	116.00
85	AA	991	G	C6-C5-N7	-9.66	124.61	130.40
86	AB	26	A	O4'-C1'-N9	9.66	115.93	108.20
34	BA	1297	G	N1-C6-O6	9.66	125.69	119.90
85	AA	1538	C	P-O5'-C5'	9.66	136.35	120.90
85	AA	2123	U	C5'-C4'-C3'	9.66	131.45	116.00
34	BA	816	G	O4'-C1'-N9	9.66	115.92	108.20
35	BB	883	G	O4'-C1'-N9	9.66	115.92	108.20
62	Bc	130	ARG	NE-CZ-NH1	9.66	125.13	120.30
85	AA	894	A	C5'-C4'-O4'	9.66	120.69	109.10
34	BA	1310	C	P-O5'-C5'	9.65	136.35	120.90
41	BH	90	C	P-O5'-C5'	9.65	136.34	120.90
85	AA	1338	C	O3'-P-O5'	-9.65	85.66	104.00
85	AA	1492	U	N1-C2-N3	9.65	120.69	114.90
34	BA	266	G	N3-C2-N2	9.65	126.66	119.90
85	AA	66	U	P-O3'-C3'	9.65	131.28	119.70
85	AA	1467	U	P-O5'-C5'	9.65	136.34	120.90
34	BA	177	G	N9-C1'-C2'	-9.65	101.39	112.00
34	BA	996	U	C4'-C3'-C2'	9.65	112.25	102.60
35	BB	1114	A	C5'-C4'-C3'	-9.65	100.56	116.00
34	BA	129	U	O3'-P-O5'	-9.65	85.67	104.00
34	BA	1240	G	C5'-C4'-C3'	9.65	131.44	116.00
85	AA	557	G	C4-N9-C1'	-9.65	113.96	126.50
37	BD	74	A	C5-C6-N6	-9.65	115.98	123.70
85	AA	1110	A	C4'-C3'-C2'	-9.65	92.95	102.60
35	BB	1453	G	O4'-C1'-N9	9.64	115.92	108.20
85	AA	821	U	N1-C2-N3	9.64	120.69	114.90
85	AA	1007	G	C4-N9-C1'	-9.64	113.96	126.50
85	AA	1346	C	C6-N1-C2	-9.64	116.44	120.30
85	AA	315	U	P-O5'-C5'	9.64	136.33	120.90
35	BB	1385	C	P-O3'-C3'	9.64	131.27	119.70
38	BE	26	G	C5'-C4'-C3'	9.64	131.43	116.00
35	BB	1431	G	C4-N9-C1'	-9.64	113.97	126.50
85	AA	251	A	P-O3'-C3'	-9.64	108.13	119.70
85	AA	254	G	P-O5'-C5'	9.64	136.32	120.90
85	AA	1594	G	O4'-C1'-N9	9.64	115.91	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1162	A	C5'-C4'-O4'	9.64	120.67	109.10
85	AA	962	U	C2-N1-C1'	-9.64	106.14	117.70
34	BA	1440	C	C1'-O4'-C4'	-9.64	102.19	109.90
85	AA	847	G	C5'-C4'-C3'	-9.64	100.58	116.00
38	BE	127	G	C5-C6-O6	9.63	134.38	128.60
39	BF	9	C	C6-N1-C2	-9.63	116.45	120.30
85	AA	271	A	N1-C6-N6	9.63	124.38	118.60
85	AA	896	C	P-O3'-C3'	-9.63	108.14	119.70
85	AA	1240	A	N1-C6-N6	9.63	124.38	118.60
64	Be	70	ARG	NE-CZ-NH2	-9.63	115.48	120.30
34	BA	688	G	C5'-C4'-C3'	-9.63	100.59	116.00
34	BA	1257	U	P-O3'-C3'	9.63	131.25	119.70
85	AA	937	G	O5'-P-OP2	-9.63	97.03	105.70
85	AA	2099	C	C5'-C4'-C3'	-9.63	100.60	116.00
85	AA	2237	G	C5-C6-O6	9.63	134.38	128.60
34	BA	165	C	O4'-C1'-N1	9.62	115.90	108.20
34	BA	349	G	N1-C6-O6	-9.62	114.12	119.90
34	BA	683	C	C4'-C3'-O3'	9.63	132.25	113.00
35	BB	37	C	P-O3'-C3'	-9.63	108.15	119.70
34	BA	682	A	P-O3'-C3'	-9.62	108.15	119.70
35	BB	1346	A	C5-C6-N6	-9.62	116.00	123.70
38	BE	27	A	C5-C6-N6	-9.62	116.00	123.70
81	Bv	68	TYR	CB-CG-CD2	-9.62	115.22	121.00
34	BA	222	C	N3-C2-O2	-9.62	115.17	121.90
35	BB	3	C	C2-N1-C1'	9.62	129.38	118.80
35	BB	845	C	C6-N1-C1'	9.62	132.35	120.80
39	BF	14	C	C2-N3-C4	-9.62	115.09	119.90
35	BB	833	G	C4-N9-C1'	-9.62	113.99	126.50
34	BA	1489	U	O5'-P-OP2	-9.62	97.04	105.70
35	BB	1097	U	C5'-C4'-C3'	-9.62	100.61	116.00
85	AA	63	G	C8-N9-C1'	9.62	139.50	127.00
40	BG	148	C	P-O5'-C5'	-9.62	105.51	120.90
85	AA	789	A	O5'-C5'-C4'	9.62	129.97	111.70
85	AA	1464	G	C5'-C4'-C3'	-9.62	100.61	116.00
34	BA	788	C	P-O5'-C5'	-9.62	105.51	120.90
35	BB	631	G	N1-C6-O6	9.62	125.67	119.90
36	BC	147	G	C5-C6-O6	-9.61	122.83	128.60
34	BA	648	C	P-O5'-C5'	9.61	136.28	120.90
34	BA	761	U	P-O5'-C5'	-9.61	105.52	120.90
34	BA	1508	C	C6-N1-C2	-9.61	116.45	120.30
35	BB	630	A	O5'-C5'-C4'	-9.61	93.44	111.70
41	BH	76	G	OP1-P-OP2	-9.61	105.18	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	125	A	C2'-C3'-O3'	9.61	130.64	109.50
85	AA	1115	G	O4'-C1'-N9	9.61	115.89	108.20
36	BC	42	G	C4-N9-C1'	-9.61	114.01	126.50
85	AA	2090	C	C6-N1-C2	-9.61	116.46	120.30
35	BB	427	U	P-O5'-C5'	-9.61	105.53	120.90
35	BB	1259	A	C5'-C4'-C3'	9.61	131.37	116.00
35	BB	1336	G	C5-C6-O6	-9.61	122.83	128.60
40	BG	12	A	P-O3'-C3'	-9.61	108.17	119.70
38	BE	25	U	O3'-P-O5'	9.61	122.25	104.00
38	BE	173	G	C5-C6-O6	-9.61	122.84	128.60
65	Bf	72	ARG	NE-CZ-NH1	9.61	125.10	120.30
85	AA	457	G	C4-N9-C1'	-9.61	114.01	126.50
34	BA	572	G	N9-C4-C5	9.60	109.24	105.40
34	BA	890	G	C5'-C4'-C3'	9.60	131.36	116.00
34	BA	1500	G	N3-C4-C5	-9.60	123.80	128.60
71	Bl	95	TRP	C-N-CA	9.60	145.71	121.70
85	AA	2085	C	C2'-C3'-O3'	9.60	130.62	109.50
34	BA	880	G	N1-C6-O6	9.60	125.66	119.90
34	BA	667	U	P-O3'-C3'	9.60	131.22	119.70
34	BA	1410	C	C5'-C4'-O4'	9.60	120.62	109.10
35	BB	979	G	C5'-C4'-C3'	-9.60	100.64	116.00
85	AA	1541	G	C5-C6-N1	9.60	116.30	111.50
85	AA	1645	G	P-O5'-C5'	9.60	136.26	120.90
34	BA	1067	G	C4-N9-C1'	-9.60	114.02	126.50
37	BD	19	C	C5'-C4'-C3'	-9.60	100.64	116.00
84	By	22	ARG	NE-CZ-NH1	9.60	125.10	120.30
85	AA	806	G	P-O5'-C5'	9.60	136.25	120.90
85	AA	2126	U	P-O3'-C3'	-9.60	108.18	119.70
85	AA	1756	C	O4'-C1'-N1	9.59	115.88	108.20
85	AA	1177	G	P-O3'-C3'	9.59	131.21	119.70
85	AA	2202	G	C5'-C4'-C3'	9.59	131.35	116.00
85	AA	1879	U	C2-N3-C4	-9.59	121.25	127.00
3	A2	27	TYR	CB-CG-CD2	-9.59	115.25	121.00
34	BA	214	A	C1'-O4'-C4'	-9.59	102.23	109.90
34	BA	845	U	O4'-C1'-N1	9.59	115.87	108.20
85	AA	351	C	O4'-C1'-N1	9.59	115.87	108.20
85	AA	1009	G	P-O3'-C3'	9.59	131.21	119.70
70	Bk	44	ARG	NE-CZ-NH2	9.59	125.09	120.30
34	BA	1124	U	C5'-C4'-C3'	-9.58	100.67	116.00
85	AA	2134	U	C5'-C4'-C3'	-9.58	100.67	116.00
34	BA	558	C	OP1-P-OP2	-9.58	105.23	119.60
34	BA	572	G	N3-C4-C5	-9.58	123.81	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	768	A	O4'-C1'-C2'	-9.58	96.22	105.80
35	BB	1071	G	C8-N9-C4	-9.58	102.57	106.40
41	BH	94	G	O4'-C1'-N9	9.58	115.86	108.20
7	A6	18	PHE	CB-CG-CD2	-9.58	114.10	120.80
40	BG	10	U	C2-N1-C1'	-9.58	106.21	117.70
85	AA	207	G	C5'-C4'-O4'	9.58	120.59	109.10
85	AA	384	C	C2-N1-C1'	-9.57	108.27	118.80
34	BA	534	C	C6-N1-C2	-9.57	116.47	120.30
38	BE	175	U	C2-N3-C4	-9.57	121.26	127.00
39	BF	24	G	O4'-C1'-N9	9.57	115.86	108.20
41	BH	26	C	C2-N3-C4	-9.57	115.11	119.90
15	AG	128	TYR	CB-CG-CD1	-9.57	115.26	121.00
34	BA	1265	G	O4'-C1'-N9	9.57	115.86	108.20
35	BB	1352	C	C1'-O4'-C4'	-9.57	102.24	109.90
41	BH	72	G	C1'-O4'-C4'	-9.57	102.24	109.90
85	AA	569	A	C8-N9-C1'	9.57	144.93	127.70
85	AA	847	G	C5-C6-O6	-9.57	122.86	128.60
85	AA	1735	U	P-O3'-C3'	9.57	131.18	119.70
34	BA	944	G	C8-N9-C1'	9.57	139.44	127.00
34	BA	1704	G	C5'-C4'-C3'	-9.57	100.69	116.00
35	BB	1197	G	C5-C6-O6	-9.57	122.86	128.60
85	AA	307	G	P-O3'-C3'	-9.57	108.22	119.70
85	AA	849	A	O4'-C1'-N9	9.57	115.85	108.20
34	BA	486	G	C5-C6-O6	-9.56	122.86	128.60
34	BA	878	G	N1-C6-O6	9.56	125.64	119.90
34	BA	911	G	C4-N9-C1'	-9.56	114.07	126.50
41	BH	10	U	O4'-C1'-N1	9.56	115.85	108.20
34	BA	1651	C	O4'-C1'-N1	9.56	115.85	108.20
39	BF	8	C	C1'-O4'-C4'	-9.56	102.25	109.90
85	AA	2131	C	C5'-C4'-C3'	-9.56	100.70	116.00
36	BC	157	U	O5'-P-OP2	-9.56	97.10	105.70
40	BG	81	G	C4-N9-C1'	-9.56	114.08	126.50
35	BB	132	G	C4-N9-C1'	-9.56	114.08	126.50
85	AA	66	U	C2-N3-C4	-9.56	121.27	127.00
85	AA	314	C	C6-N1-C2	-9.56	116.48	120.30
35	BB	504	C	C5'-C4'-O4'	9.55	120.56	109.10
85	AA	723	U	P-O5'-C5'	9.55	136.18	120.90
34	BA	362	G	N1-C6-O6	-9.55	114.17	119.90
37	BD	84	U	N1-C2-N3	9.55	120.63	114.90
35	BB	1359	G	N1-C6-O6	9.55	125.63	119.90
34	BA	150	C	O4'-C1'-N1	9.55	115.84	108.20
34	BA	800	G	O4'-C1'-N9	-9.55	100.56	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	17	U	C2-N3-C4	-9.55	121.27	127.00
34	BA	99	G	C6-N1-C2	-9.54	119.37	125.10
35	BB	1129	C	P-O5'-C5'	9.54	136.17	120.90
34	BA	330	A	N1-C6-N6	9.54	124.33	118.60
34	BA	819	G	C5'-C4'-C3'	-9.54	100.73	116.00
34	BA	1809	G	C3'-C2'-C1'	-9.54	93.87	101.50
85	AA	1797	U	O4'-C1'-N1	9.54	115.83	108.20
35	BB	963	G	C8-N9-C1'	9.54	139.40	127.00
85	AA	100	A	N9-C1'-C2'	9.54	126.40	114.00
34	BA	548	G	C8-N9-C4	-9.54	102.58	106.40
35	BB	423	G	C5'-C4'-C3'	-9.54	100.74	116.00
35	BB	505	G	N1-C6-O6	-9.54	114.18	119.90
36	BC	128	U	O4'-C1'-N1	9.54	115.83	108.20
36	BC	38	U	O4'-C1'-N1	9.54	115.83	108.20
85	AA	1547	G	N1-C6-O6	-9.54	114.18	119.90
35	BB	1302	C	N3-C4-N4	9.53	124.67	118.00
85	AA	974	U	O4'-C1'-N1	9.53	115.83	108.20
34	BA	636	G	N1-C6-O6	-9.53	114.18	119.90
35	BB	1206	G	N3-C2-N2	9.53	126.57	119.90
85	AA	572	G	C4-N9-C1'	-9.53	114.11	126.50
85	AA	1281	G	C8-N9-C4	9.53	110.21	106.40
80	Bu	109	MET	CG-SD-CE	-9.53	84.95	100.20
35	BB	503	G	C5-C6-O6	-9.53	122.88	128.60
34	BA	862	C	C2-N3-C4	-9.53	115.14	119.90
34	BA	1793	G	O4'-C1'-N9	9.53	115.82	108.20
34	BA	1804	A	O4'-C1'-N9	9.53	115.82	108.20
35	BB	384	A	P-O3'-C3'	-9.53	108.27	119.70
35	BB	453	C	C5'-C4'-C3'	9.53	131.24	116.00
85	AA	1797	U	P-O5'-C5'	9.53	136.14	120.90
85	AA	1190	G	C8-N9-C4	9.52	110.21	106.40
34	BA	874	G	P-O5'-C5'	9.52	136.13	120.90
85	AA	242	G	O4'-C1'-N9	9.52	115.82	108.20
85	AA	1794	U	P-O5'-C5'	-9.52	105.67	120.90
85	AA	1921	G	C4-C5-N7	-9.52	106.99	110.80
34	BA	1662	U	C2-N3-C4	-9.52	121.29	127.00
35	BB	524	C	O4'-C1'-N1	9.52	115.81	108.20
85	AA	572	G	C8-N9-C1'	9.52	139.37	127.00
34	BA	1723	U	N1-C2-N3	9.52	120.61	114.90
35	BB	791	A	N1-C6-N6	-9.52	112.89	118.60
34	BA	102	G	N1-C6-O6	-9.52	114.19	119.90
34	BA	1023	G	C5-C6-O6	-9.52	122.89	128.60
34	BA	1215	U	P-O5'-C5'	9.52	136.13	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	164	G	C8-N9-C1'	9.52	139.37	127.00
35	BB	608	A	N1-C6-N6	-9.51	112.89	118.60
40	BG	11	G	C2-N3-C4	-9.51	107.14	111.90
40	BG	170	G	P-O3'-C3'	-9.51	108.28	119.70
80	Bu	108	ARG	NE-CZ-NH1	9.51	125.06	120.30
34	BA	360	C	O4'-C1'-N1	9.51	115.81	108.20
38	BE	110	U	P-O3'-C3'	-9.51	108.29	119.70
34	BA	882	G	C1'-O4'-C4'	-9.51	102.29	109.90
85	AA	465	A	P-O3'-C3'	9.51	131.11	119.70
34	BA	763	U	C6-N1-C2	-9.51	115.30	121.00
34	BA	1821	A	N1-C6-N6	-9.51	112.90	118.60
35	BB	1203	C	C5'-C4'-C3'	-9.51	100.79	116.00
36	BC	164	G	C4-N9-C1'	-9.51	114.14	126.50
85	AA	1235	G	P-O3'-C3'	-9.51	108.29	119.70
85	AA	2053	A	C8-N9-C4	-9.51	102.00	105.80
34	BA	1658	G	C5'-C4'-C3'	-9.50	100.80	116.00
36	BC	20	C	O5'-P-OP1	-9.50	97.15	105.70
34	BA	1477	C	C5'-C4'-O4'	9.50	120.50	109.10
35	BB	1133	C	P-O5'-C5'	-9.50	105.70	120.90
85	AA	1692	U	N3-C2-O2	-9.50	115.55	122.20
85	AA	1921	G	N9-C4-C5	9.50	109.20	105.40
86	AB	17	C	O4'-C1'-N1	9.50	115.80	108.20
34	BA	530	A	P-O3'-C3'	9.49	131.09	119.70
34	BA	777	C	C6-N1-C2	-9.49	116.50	120.30
35	BB	520	G	C5-C6-O6	-9.49	122.90	128.60
85	AA	1248	U	C2-N3-C4	-9.49	121.31	127.00
36	BC	31	A	C3'-C2'-C1'	-9.49	93.91	101.50
38	BE	46	G	C4-N9-C1'	-9.49	114.16	126.50
85	AA	1235	G	O4'-C1'-N9	9.49	115.79	108.20
35	BB	490	G	C5'-C4'-O4'	9.49	120.49	109.10
35	BB	1426	G	N3-C4-C5	-9.49	123.86	128.60
34	BA	21	C	O4'-C1'-N1	9.48	115.79	108.20
34	BA	571	G	C4-N9-C1'	9.48	138.83	126.50
34	BA	802	G	C4-N9-C1'	-9.48	114.17	126.50
85	AA	881	C	C6-N1-C1'	9.48	132.18	120.80
34	BA	554	A	O5'-P-OP2	-9.48	97.17	105.70
34	BA	253	U	C2-N1-C1'	-9.48	106.32	117.70
85	AA	898	A	O4'-C1'-N9	9.48	115.78	108.20
85	AA	1006	C	C2-N1-C1'	-9.48	108.37	118.80
85	AA	1471	G	C4-C5-N7	9.48	114.59	110.80
37	BD	83	A	O4'-C1'-N9	9.48	115.78	108.20
34	BA	1716	A	C5'-C4'-C3'	9.48	131.16	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	881	G	C8-N9-C1'	9.48	139.32	127.00
35	BB	1475	U	P-O3'-C3'	9.48	131.07	119.70
85	AA	535	G	N1-C6-O6	9.48	125.59	119.90
34	BA	43	U	C2-N1-C1'	-9.47	106.33	117.70
34	BA	564	C	O4'-C1'-N1	9.47	115.78	108.20
85	AA	209	C	C6-N1-C2	-9.47	116.51	120.30
85	AA	857	G	C8-N9-C1'	9.47	139.32	127.00
85	AA	1496	U	O3'-P-O5'	9.47	122.00	104.00
34	BA	220	U	P-O5'-C5'	9.47	136.06	120.90
35	BB	1138	A	O5'-C5'-C4'	-9.47	93.71	111.70
85	AA	434	U	P-O5'-C5'	-9.47	105.75	120.90
34	BA	1071	G	C5-C6-O6	-9.47	122.92	128.60
34	BA	486	G	C8-N9-C4	-9.47	102.61	106.40
40	BG	59	G	C5'-C4'-C3'	-9.47	100.85	116.00
34	BA	1138	C	C6-N1-C1'	9.46	132.16	120.80
35	BB	363	A	C1'-O4'-C4'	-9.46	102.33	109.90
85	AA	767	A	O4'-C1'-N9	9.46	115.77	108.20
34	BA	272	A	C5'-C4'-C3'	9.46	131.14	116.00
34	BA	658	C	O4'-C1'-N1	9.46	115.77	108.20
34	BA	761	U	C2-N1-C1'	-9.46	106.34	117.70
34	BA	1226	G	P-O3'-C3'	-9.46	108.35	119.70
35	BB	1323	U	C6-N1-C2	-9.46	115.32	121.00
34	BA	1506	C	C6-N1-C2	-9.46	116.52	120.30
34	BA	1724	G	C5-C6-N1	9.46	116.23	111.50
37	BD	83	A	C4'-C3'-C2'	-9.46	93.14	102.60
85	AA	494	G	C5'-C4'-C3'	-9.46	100.86	116.00
85	AA	907	G	C1'-O4'-C4'	-9.46	102.33	109.90
85	AA	1789	C	C5'-C4'-O4'	-9.46	97.75	109.10
85	AA	1878	C	P-O5'-C5'	9.46	136.04	120.90
34	BA	1733	G	P-O5'-C5'	9.46	136.03	120.90
85	AA	2179	C	O4'-C1'-N1	9.46	115.77	108.20
35	BB	1445	A	C5-C6-N6	9.46	131.26	123.70
86	AB	61	C	C2-N1-C1'	-9.46	108.40	118.80
38	BE	146	U	N3-C2-O2	-9.45	115.58	122.20
85	AA	1897	A	O4'-C1'-N9	9.46	115.76	108.20
5	A4	143	ARG	NE-CZ-NH1	9.45	125.03	120.30
34	BA	243	C	C6-N1-C1'	9.45	132.14	120.80
35	BB	545	C	C6-N1-C2	-9.45	116.52	120.30
85	AA	888	A	P-O3'-C3'	9.45	131.04	119.70
85	AA	1541	G	C6-N1-C2	-9.45	119.43	125.10
85	AA	1852	U	P-O3'-C3'	9.45	131.04	119.70
85	AA	2008	G	N9-C4-C5	9.45	109.18	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1868	G	O4'-C1'-N9	9.45	115.76	108.20
34	BA	906	A	O4'-C1'-N9	9.45	115.76	108.20
34	BA	933	U	C2-N1-C1'	-9.45	106.36	117.70
34	BA	2	A	C2'-C3'-O3'	9.45	130.28	109.50
34	BA	966	G	N1-C6-O6	-9.45	114.23	119.90
34	BA	1563	G	C8-N9-C4	-9.44	102.62	106.40
35	BB	871	C	C3'-C2'-C1'	-9.45	93.94	101.50
35	BB	977	G	C8-N9-C4	-9.45	102.62	106.40
35	BB	996	G	P-O5'-C5'	-9.44	105.79	120.90
35	BB	1482	A	P-O3'-C3'	9.44	131.03	119.70
38	BE	185	G	C8-N9-C1'	9.44	139.27	127.00
85	AA	889	G	N3-C2-N2	-9.44	113.29	119.90
34	BA	526	C	C4'-C3'-C2'	-9.44	93.16	102.60
34	BA	685	C	O4'-C4'-C3'	-9.44	94.56	104.00
35	BB	529	A	C5-C6-N6	-9.44	116.15	123.70
40	BG	9	G	P-O5'-C5'	-9.44	105.80	120.90
85	AA	63	G	C4-N9-C1'	-9.44	114.23	126.50
68	Bi	111	ARG	NE-CZ-NH1	9.44	125.02	120.30
85	AA	584	G	C4-N9-C1'	9.44	138.77	126.50
37	BD	71	G	P-O3'-C3'	9.43	131.02	119.70
41	BH	129	G	N1-C6-O6	-9.43	114.24	119.90
35	BB	813	C	O4'-C1'-N1	9.43	115.75	108.20
85	AA	937	G	C5-C6-O6	-9.43	122.94	128.60
7	A6	84	PHE	CB-CG-CD2	-9.43	114.20	120.80
34	BA	99	G	C4-N9-C1'	9.43	138.76	126.50
34	BA	1207	A	P-O3'-C3'	9.43	131.01	119.70
85	AA	2126	U	C5'-C4'-C3'	9.43	131.09	116.00
34	BA	943	G	C4-N9-C1'	-9.43	114.25	126.50
34	BA	1541	G	N1-C6-O6	9.43	125.56	119.90
85	AA	253	C	C6-N1-C2	-9.43	116.53	120.30
85	AA	817	G	N1-C6-O6	-9.43	114.25	119.90
34	BA	513	U	O4'-C1'-N1	9.42	115.74	108.20
34	BA	1260	G	C3'-C2'-C1'	-9.42	93.96	101.50
34	BA	1803	A	C8-N9-C4	9.42	109.57	105.80
35	BB	1093	C	C6-N1-C2	-9.42	116.53	120.30
35	BB	1480	G	C6-C5-N7	-9.42	124.75	130.40
85	AA	1913	G	C8-N9-C4	-9.42	102.63	106.40
34	BA	1194	G	C8-N9-C1'	9.42	139.24	127.00
34	BA	1060	C	P-O3'-C3'	9.42	131.00	119.70
85	AA	1010	U	C6-N1-C1'	9.42	134.38	121.20
2	A1	97	ARG	NE-CZ-NH1	9.41	125.01	120.30
34	BA	1041	U	P-O5'-C5'	9.41	135.96	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	270	A	P-O3'-C3'	-9.41	108.40	119.70
34	BA	1566	G	N1-C6-O6	9.41	125.55	119.90
34	BA	1653	G	C4-N9-C1'	-9.41	114.26	126.50
85	AA	2074	G	N1-C6-O6	9.41	125.55	119.90
34	BA	1631	U	C5'-C4'-C3'	9.41	131.06	116.00
35	BB	763	U	C2-N1-C1'	-9.41	106.40	117.70
35	BB	945	U	P-O3'-C3'	9.41	131.00	119.70
85	AA	2127	G	O4'-C1'-N9	9.41	115.73	108.20
15	AG	55	ARG	NE-CZ-NH2	-9.41	115.60	120.30
27	AT	120	ARG	NE-CZ-NH1	9.41	125.00	120.30
34	BA	611	A	C5'-C4'-C3'	-9.41	100.95	116.00
39	BF	22	U	N1-C2-N3	9.41	120.54	114.90
85	AA	817	G	C5-C6-O6	9.41	134.24	128.60
85	AA	1050	C	C6-N1-C2	-9.41	116.54	120.30
85	AA	1245	U	C5'-C4'-C3'	-9.41	100.95	116.00
34	BA	196	A	N9-C1'-C2'	-9.40	101.66	112.00
35	BB	1459	U	C2-N3-C4	-9.40	121.36	127.00
41	BH	6	U	N3-C2-O2	-9.40	115.62	122.20
34	BA	89	G	C4-N9-C1'	-9.40	114.28	126.50
35	BB	878	G	P-O3'-C3'	9.40	130.98	119.70
34	BA	111	U	C1'-O4'-C4'	-9.40	102.38	109.90
34	BA	297	A	C5'-C4'-O4'	9.40	120.38	109.10
85	AA	799	G	C5-C6-O6	-9.40	122.96	128.60
34	BA	1515	U	O4'-C1'-N1	9.40	115.72	108.20
35	BB	441	G	C4-N9-C1'	-9.40	114.28	126.50
35	BB	1193	G	C5'-C4'-C3'	9.40	131.04	116.00
85	AA	2128	G	C4-N9-C1'	-9.40	114.28	126.50
34	BA	260	A	O4'-C1'-N9	9.40	115.72	108.20
37	BD	95	G	N1-C6-O6	-9.40	114.26	119.90
8	A7	105	PHE	CB-CG-CD2	-9.39	114.22	120.80
38	BE	78	C	O4'-C1'-N1	9.39	115.72	108.20
45	BL	135	TYR	CB-CG-CD1	-9.39	115.36	121.00
85	AA	2196	G	C8-N9-C4	-9.39	102.64	106.40
34	BA	141	G	N9-C4-C5	9.39	109.16	105.40
34	BA	297	A	C2'-C3'-O3'	9.39	130.16	109.50
85	AA	1186	C	O4'-C1'-N1	9.39	115.72	108.20
34	BA	248	G	N1-C6-O6	9.39	125.53	119.90
34	BA	720	A	N1-C6-N6	9.39	124.23	118.60
35	BB	1426	G	N1-C2-N2	-9.39	107.75	116.20
38	BE	158	U	P-O3'-C3'	9.39	130.97	119.70
85	AA	1464	G	C3'-C2'-C1'	-9.39	93.99	101.50
85	AA	1661	U	C5'-C4'-C3'	-9.39	100.97	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
12	AD	17	PHE	CB-CG-CD2	-9.38	114.23	120.80
34	BA	1096	C	O4'-C1'-N1	9.38	115.71	108.20
35	BB	1204	C	P-O3'-C3'	9.38	130.96	119.70
40	BG	54	G	N3-C2-N2	-9.38	113.33	119.90
85	AA	365	G	C8-N9-C4	9.39	110.15	106.40
85	AA	1225	C	O4'-C1'-N1	9.38	115.71	108.20
34	BA	192	G	C5'-C4'-C3'	9.38	131.01	116.00
35	BB	1079	G	C4-N9-C1'	-9.38	114.30	126.50
85	AA	46	U	C2-N1-C1'	9.38	128.96	117.70
85	AA	424	A	C1'-O4'-C4'	-9.38	102.39	109.90
85	AA	1793	A	C8-N9-C1'	9.38	144.59	127.70
35	BB	1151	A	N1-C6-N6	-9.38	112.97	118.60
37	BD	29	C	C5'-C4'-C3'	-9.38	100.99	116.00
4	A3	157	ARG	NE-CZ-NH1	9.38	124.99	120.30
36	BC	58	G	C6-N1-C2	-9.38	119.47	125.10
34	BA	1265	G	C4-N9-C1'	9.38	138.69	126.50
35	BB	427	U	C1'-O4'-C4'	-9.38	102.40	109.90
40	BG	54	G	P-O3'-C3'	-9.38	108.45	119.70
85	AA	775	C	O4'-C1'-N1	9.38	115.70	108.20
34	BA	531	C	O4'-C1'-N1	9.38	115.70	108.20
34	BA	537	C	O4'-C1'-N1	9.38	115.70	108.20
34	BA	665	C	N3-C2-O2	-9.38	115.34	121.90
36	BC	50	C	O4'-C1'-N1	9.37	115.70	108.20
38	BE	142	A	N1-C6-N6	-9.38	112.97	118.60
85	AA	592	C	O4'-C1'-N1	9.38	115.70	108.20
85	AA	820	G	C4'-C3'-C2'	-9.37	93.23	102.60
85	AA	1088	C	O4'-C1'-N1	9.37	115.70	108.20
85	AA	2007	G	O4'-C1'-N9	9.37	115.70	108.20
12	AD	14	TYR	CB-CG-CD2	-9.37	115.38	121.00
85	AA	1645	G	C5-C6-O6	-9.37	122.98	128.60
85	AA	671	G	P-O5'-C5'	-9.37	105.91	120.90
85	AA	1806	C	C6-N1-C2	-9.37	116.55	120.30
85	AA	1558	U	C5'-C4'-C3'	9.37	130.99	116.00
85	AA	1763	G	C5-C6-O6	-9.37	122.98	128.60
34	BA	334	G	C5-C6-O6	9.37	134.22	128.60
34	BA	660	C	C6-N1-C1'	9.37	132.04	120.80
35	BB	799	A	C1'-O4'-C4'	9.37	117.39	109.90
35	BB	1201	G	C4'-C3'-O3'	9.37	131.73	113.00
37	BD	100	A	C5'-C4'-C3'	9.37	130.99	116.00
34	BA	1498	A	C5'-C4'-C3'	-9.36	101.02	116.00
85	AA	1676	G	C5'-C4'-C3'	-9.36	101.02	116.00
85	AA	1799	C	O5'-P-OP1	-9.36	97.27	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1845	G	P-O3'-C3'	-9.36	108.46	119.70
35	BB	853	U	P-O3'-C3'	9.36	130.94	119.70
36	BC	168	C	P-O3'-C3'	-9.36	108.47	119.70
38	BE	76	U	O4'-C1'-N1	9.36	115.69	108.20
38	BE	183	C	N3-C4-N4	9.36	124.55	118.00
85	AA	1117	G	N1-C6-O6	-9.36	114.28	119.90
85	AA	1636	C	O4'-C1'-N1	9.36	115.69	108.20
85	AA	618	A	O4'-C1'-N9	9.36	115.69	108.20
85	AA	807	A	C5'-C4'-C3'	-9.36	101.03	116.00
85	AA	1228	A	O4'-C1'-N9	9.36	115.69	108.20
85	AA	1647	G	N1-C6-O6	-9.36	114.28	119.90
85	AA	2056	C	C2-N1-C1'	-9.36	108.51	118.80
34	BA	1001	G	C5-C6-O6	-9.36	122.99	128.60
34	BA	1739	G	C5-C6-O6	-9.36	122.99	128.60
41	BH	113	G	C4-N9-C1'	-9.36	114.34	126.50
53	BT	107	ARG	NE-CZ-NH1	9.36	124.98	120.30
36	BC	74	U	C5-C6-N1	-9.35	118.02	122.70
38	BE	6	A	C6-N1-C2	-9.35	112.99	118.60
85	AA	277	G	C5'-C4'-C3'	-9.35	101.04	116.00
85	AA	392	G	C4-N9-C1'	-9.35	114.34	126.50
85	AA	1952	C	O4'-C1'-N1	9.35	115.68	108.20
85	AA	2046	G	C8-N9-C4	-9.35	102.66	106.40
34	BA	1426	A	C8-N9-C4	-9.35	102.06	105.80
34	BA	1847	G	C2-N3-C4	-9.35	107.22	111.90
39	BF	32	G	C5'-C4'-O4'	9.35	120.32	109.10
36	BC	126	G	O4'-C1'-N9	9.35	115.68	108.20
85	AA	253	C	C5'-C4'-C3'	-9.35	101.04	116.00
85	AA	604	C	O4'-C1'-N1	9.35	115.68	108.20
85	AA	1611	A	P-O5'-C5'	-9.35	105.94	120.90
35	BB	1467	A	O4'-C1'-N9	9.35	115.68	108.20
85	AA	326	C	C6-N1-C1'	-9.35	109.58	120.80
34	BA	146	G	C4'-C3'-C2'	-9.34	93.26	102.60
35	BB	522	A	C4'-C3'-C2'	9.34	111.94	102.60
85	AA	970	U	O4'-C1'-N1	9.34	115.67	108.20
34	BA	471	U	N3-C2-O2	9.34	128.74	122.20
34	BA	1579	G	N1-C6-O6	-9.34	114.30	119.90
40	BG	147	U	P-O5'-C5'	9.34	135.84	120.90
85	AA	684	G	C5-C6-N1	9.34	116.17	111.50
34	BA	1745	G	C8-N9-C1'	9.34	139.14	127.00
85	AA	977	U	C2-N3-C4	-9.34	121.40	127.00
34	BA	1052	G	N1-C6-O6	9.34	125.50	119.90
35	BB	544	C	C6-N1-C2	-9.34	116.57	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	120	C	O4'-C1'-N1	9.34	115.67	108.20
35	BB	1511	U	C5'-C4'-O4'	-9.34	97.90	109.10
85	AA	495	G	C4-N9-C1'	-9.34	114.36	126.50
85	AA	1299	A	O4'-C1'-N9	9.33	115.67	108.20
34	BA	108	A	P-O3'-C3'	9.33	130.90	119.70
34	BA	270	U	C2-N3-C4	-9.33	121.40	127.00
35	BB	881	G	C4-N9-C1'	-9.33	114.37	126.50
35	BB	1245	A	N1-C6-N6	-9.33	113.00	118.60
85	AA	1492	U	C2-N1-C1'	-9.33	106.50	117.70
34	BA	298	G	O4'-C1'-N9	9.33	115.66	108.20
34	BA	1190	A	O4'-C1'-N9	9.33	115.66	108.20
34	BA	1334	G	C5-C6-N1	9.33	116.16	111.50
34	BA	1579	G	O4'-C1'-N9	9.33	115.66	108.20
40	BG	2	U	C5'-C4'-O4'	9.33	120.29	109.10
85	AA	467	U	C2-N1-C1'	9.33	128.89	117.70
35	BB	262	C	O4'-C1'-N1	9.33	115.66	108.20
11	AC	216	ARG	NE-CZ-NH1	9.32	124.96	120.30
34	BA	839	U	O3'-P-O5'	-9.32	86.28	104.00
35	BB	1426	G	N3-C2-N2	9.32	126.43	119.90
37	BD	82	G	O4'-C1'-C2'	9.32	115.99	107.60
85	AA	1300	A	P-O3'-C3'	-9.32	108.51	119.70
85	AA	2018	U	O4'-C1'-N1	9.32	115.66	108.20
85	AA	2085	C	C2-N1-C1'	-9.32	108.54	118.80
34	BA	53	G	C4-C5-C6	-9.32	113.21	118.80
34	BA	701	G	C1'-O4'-C4'	-9.32	102.44	109.90
35	BB	1410	G	C5-C6-O6	-9.32	123.01	128.60
85	AA	240	A	O4'-C1'-N9	9.32	115.66	108.20
34	BA	450	G	C5-C6-O6	-9.32	123.01	128.60
39	BF	55	A	O4'-C1'-N9	9.32	115.66	108.20
85	AA	1717	G	P-O3'-C3'	9.32	130.88	119.70
85	AA	2092	A	P-O5'-C5'	9.32	135.81	120.90
34	BA	1668	C	P-O3'-C3'	-9.32	108.52	119.70
35	BB	432	C	C5'-C4'-C3'	9.32	130.91	116.00
36	BC	75	G	P-O3'-C3'	-9.32	108.52	119.70
35	BB	1083	C	C6-N1-C2	-9.32	116.57	120.30
38	BE	175	U	C6-N1-C2	-9.32	115.41	121.00
56	BW	37	TYR	CB-CG-CD2	-9.32	115.41	121.00
85	AA	1128	G	C4-N9-C1'	-9.32	114.39	126.50
29	AV	24	HIS	CA-CB-CG	9.31	129.43	113.60
34	BA	595	U	P-O3'-C3'	9.31	130.88	119.70
83	Bx	100	ARG	NE-CZ-NH1	9.31	124.96	120.30
85	AA	787	U	C5'-C4'-C3'	-9.31	101.10	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1919	G	C5-C6-O6	-9.31	123.01	128.60
34	BA	141	G	C5'-C4'-C3'	-9.31	101.10	116.00
35	BB	1218	G	C4-N9-C1'	9.31	138.60	126.50
85	AA	137	C	C6-N1-C2	-9.31	116.58	120.30
35	BB	545	C	P-O3'-C3'	9.31	130.87	119.70
83	Bx	48	GLY	N-CA-C	9.31	136.37	113.10
85	AA	450	A	C5'-C4'-C3'	-9.31	101.11	116.00
35	BB	1434	G	C5-C6-O6	-9.31	123.02	128.60
85	AA	388	G	C5-C6-O6	-9.31	123.02	128.60
85	AA	1962	U	C4'-C3'-C2'	9.31	111.91	102.60
35	BB	773	G	C5-C6-O6	-9.31	123.02	128.60
38	BE	30	C	C2-N1-C1'	-9.30	108.57	118.80
41	BH	41	A	C1'-O4'-C4'	-9.30	102.46	109.90
38	BE	195	G	N3-C2-N2	9.30	126.41	119.90
85	AA	32	U	C2-N3-C4	-9.30	121.42	127.00
85	AA	1090	A	O3'-P-O5'	-9.30	86.33	104.00
85	AA	1506	U	C2-N3-C4	-9.30	121.42	127.00
85	AA	2062	U	C2-N1-C1'	-9.30	106.54	117.70
40	BG	171	A	N9-C4-C5	9.30	109.52	105.80
41	BH	41	A	C3'-C2'-C1'	-9.30	94.06	101.50
85	AA	760	U	C3'-C2'-C1'	-9.30	94.06	101.50
85	AA	918	U	O4'-C1'-N1	9.30	115.64	108.20
85	AA	2225	G	C4-N9-C1'	-9.30	114.42	126.50
85	AA	159	G	C4-N9-C1'	-9.29	114.42	126.50
34	BA	190	U	O4'-C1'-N1	9.29	115.63	108.20
34	BA	357	A	N1-C6-N6	-9.29	113.02	118.60
34	BA	741	A	C1'-O4'-C4'	-9.29	102.47	109.90
34	BA	564	C	C6-N1-C1'	-9.29	109.65	120.80
35	BB	405	U	C4'-C3'-C2'	-9.29	93.31	102.60
38	BE	128	G	O5'-P-OP2	-9.29	97.34	105.70
38	BE	130	G	C8-N9-C4	-9.29	102.68	106.40
43	BJ	59	ARG	NE-CZ-NH1	9.29	124.95	120.30
85	AA	1594	G	N1-C6-O6	9.29	125.48	119.90
41	BH	58	C	C6-N1-C2	-9.29	116.58	120.30
42	BI	135	MET	N-CA-CB	-9.29	93.88	110.60
34	BA	236	A	C4'-C3'-C2'	9.29	111.89	102.60
35	BB	132	G	C5-C6-O6	-9.29	123.03	128.60
34	BA	480	G	C5-C6-O6	-9.29	123.03	128.60
34	BA	1210	A	P-O3'-C3'	-9.29	108.55	119.70
36	BC	121	G	C5-C6-O6	-9.29	123.03	128.60
85	AA	265	A	O4'-C1'-N9	9.29	115.63	108.20
34	BA	483	A	C1'-O4'-C4'	9.28	117.33	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	538	G	N1-C2-N2	-9.29	107.84	116.20
34	BA	642	U	O4'-C1'-N1	9.28	115.63	108.20
34	BA	666	C	C5'-C4'-C3'	-9.29	101.14	116.00
34	BA	349	G	C5-C6-O6	9.28	134.17	128.60
35	BB	43	G	P-O3'-C3'	-9.28	108.56	119.70
36	BC	137	C	O4'-C1'-N1	9.28	115.63	108.20
35	BB	605	C	C4'-C3'-C2'	9.28	111.88	102.60
22	AO	62	ARG	NE-CZ-NH1	9.28	124.94	120.30
34	BA	1206	C	O4'-C1'-N1	9.28	115.62	108.20
34	BA	1672	C	O4'-C1'-N1	9.28	115.62	108.20
85	AA	1240	A	C4'-C3'-C2'	9.28	111.88	102.60
13	AE	76	PHE	CB-CG-CD2	-9.28	114.31	120.80
34	BA	744	G	C3'-C2'-C1'	-9.28	94.08	101.50
34	BA	954	U	C2-N1-C1'	-9.28	106.57	117.70
85	AA	473	C	C6-N1-C2	-9.28	116.59	120.30
35	BB	439	G	C6-N1-C2	-9.28	119.53	125.10
53	BT	185	ARG	NE-CZ-NH1	9.27	124.94	120.30
34	BA	610	A	O4'-C1'-N9	9.27	115.62	108.20
34	BA	1789	A	C5'-C4'-C3'	-9.27	101.17	116.00
86	AB	21	A	O4'-C1'-N9	9.27	115.62	108.20
35	BB	82	G	C8-N9-C1'	9.27	139.05	127.00
35	BB	769	C	C5'-C4'-C3'	-9.27	101.17	116.00
35	BB	837	A	C8-N9-C4	-9.27	102.09	105.80
35	BB	1225	A	P-O3'-C3'	9.27	130.82	119.70
38	BE	117	A	O4'-C4'-C3'	9.27	113.52	106.10
49	BP	23	ARG	NE-CZ-NH2	-9.27	115.67	120.30
39	BF	39	C	C4'-C3'-C2'	9.27	111.87	102.60
35	BB	112	G	C6-N1-C2	-9.27	119.54	125.10
85	AA	353	G	C5-C6-O6	-9.27	123.04	128.60
85	AA	1086	U	C2-N1-C1'	-9.27	106.58	117.70
34	BA	112	C	C6-N1-C1'	-9.26	109.69	120.80
34	BA	516	U	P-O5'-C5'	9.26	135.72	120.90
34	BA	840	U	C1'-O4'-C4'	-9.26	102.49	109.90
34	BA	1711	G	C5-C6-O6	-9.26	123.04	128.60
35	BB	1331	U	O4'-C1'-N1	9.26	115.61	108.20
85	AA	866	U	O4'-C1'-N1	9.26	115.61	108.20
34	BA	229	C	P-O3'-C3'	9.26	130.81	119.70
34	BA	577	U	P-O3'-C3'	9.26	130.81	119.70
34	BA	1324	G	C5-C6-O6	-9.26	123.05	128.60
85	AA	452	A	O4'-C1'-N9	9.26	115.61	108.20
34	BA	237	A	C1'-O4'-C4'	-9.26	102.49	109.90
34	BA	889	U	N3-C2-O2	-9.26	115.72	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	755	G	N1-C6-O6	9.26	125.45	119.90
85	AA	84	C	P-O5'-C5'	9.26	135.71	120.90
85	AA	227	A	O4'-C1'-N9	9.26	115.60	108.20
22	AO	139	ARG	NE-CZ-NH1	9.25	124.93	120.30
34	BA	1516	G	C5'-C4'-C3'	-9.25	101.20	116.00
37	BD	118	C	P-O5'-C5'	9.25	135.70	120.90
85	AA	763	U	C1'-O4'-C4'	-9.25	102.50	109.90
34	BA	159	U	C5'-C4'-C3'	9.25	130.80	116.00
34	BA	329	G	C8-N9-C1'	9.25	139.03	127.00
40	BG	72	G	N1-C6-O6	-9.25	114.35	119.90
39	BF	45	G	P-O3'-C3'	-9.25	108.60	119.70
77	Br	313	TYR	CB-CG-CD2	-9.25	115.45	121.00
85	AA	126	U	O4'-C1'-N1	9.25	115.60	108.20
34	BA	12	G	P-O5'-C5'	9.25	135.70	120.90
34	BA	110	C	O4'-C1'-N1	9.25	115.60	108.20
34	BA	115	U	C1'-O4'-C4'	-9.25	102.50	109.90
34	BA	124	G	P-O3'-C3'	-9.25	108.60	119.70
34	BA	585	G	P-O3'-C3'	-9.25	108.60	119.70
34	BA	1475	G	C5'-C4'-C3'	9.25	130.80	116.00
34	BA	1707	C	C6-N1-C1'	9.25	131.90	120.80
35	BB	1000	U	O4'-C1'-N1	9.25	115.60	108.20
85	AA	382	G	C4-N9-C1'	-9.25	114.48	126.50
85	AA	476	C	C1'-O4'-C4'	-9.25	102.50	109.90
35	BB	789	G	P-O5'-C5'	-9.24	106.11	120.90
85	AA	608	A	O4'-C1'-N9	9.24	115.60	108.20
34	BA	1019	C	P-O3'-C3'	-9.24	108.61	119.70
37	BD	66	G	N3-C2-N2	9.24	126.37	119.90
38	BE	32	U	P-O5'-C5'	-9.24	106.11	120.90
85	AA	768	C	C6-N1-C1'	9.24	131.89	120.80
35	BB	1368	A	C8-N9-C4	9.24	109.50	105.80
36	BC	156	A	N1-C6-N6	9.24	124.14	118.60
85	AA	166	C	C6-N1-C2	-9.24	116.60	120.30
34	BA	540	G	C4-N9-C1'	-9.24	114.49	126.50
34	BA	1014	A	N1-C6-N6	-9.24	113.06	118.60
85	AA	1527	G	P-O3'-C3'	-9.24	108.61	119.70
35	BB	968	C	O4'-C1'-N1	9.24	115.59	108.20
85	AA	610	C	O4'-C1'-N1	9.24	115.59	108.20
85	AA	99	U	O3'-P-O5'	-9.23	86.45	104.00
85	AA	332	A	P-O3'-C3'	9.23	130.78	119.70
34	BA	175	G	O4'-C1'-N9	9.23	115.59	108.20
85	AA	486	G	C5-C6-O6	-9.23	123.06	128.60
85	AA	650	G	C4-N9-C1'	-9.23	114.49	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
13	AE	98	ARG	NE-CZ-NH2	-9.23	115.68	120.30
41	BH	121	A	C2-N3-C4	-9.23	105.98	110.60
85	AA	466	A	C5-C6-N1	-9.23	113.08	117.70
85	AA	923	A	O4'-C1'-N9	9.23	115.58	108.20
85	AA	78	A	C5'-C4'-C3'	-9.23	101.24	116.00
85	AA	400	G	C5-C6-O6	-9.23	123.06	128.60
85	AA	1469	G	C2-N3-C4	9.23	116.51	111.90
85	AA	2066	C	O4'-C1'-N1	9.23	115.58	108.20
34	BA	1419	A	C5-C6-N6	-9.23	116.32	123.70
35	BB	38	C	C6-N1-C2	-9.23	116.61	120.30
35	BB	1113	C	C3'-C2'-C1'	-9.23	94.12	101.50
85	AA	1549	G	C5-C6-O6	-9.23	123.06	128.60
35	BB	854	G	C8-N9-C1'	9.22	138.99	127.00
38	BE	68	U	C5'-C4'-C3'	9.22	130.76	116.00
13	AE	113	TYR	CA-CB-CG	-9.22	95.88	113.40
35	BB	672	C	O4'-C1'-N1	9.22	115.58	108.20
71	Bl	141	ARG	NE-CZ-NH2	-9.22	115.69	120.30
85	AA	16	G	C8-N9-C4	9.22	110.09	106.40
85	AA	1923	A	O4'-C1'-N9	9.22	115.58	108.20
34	BA	116	G	P-O5'-C5'	-9.22	106.15	120.90
34	BA	517	A	C1'-O4'-C4'	-9.22	102.52	109.90
34	BA	547	C	O4'-C1'-N1	9.22	115.58	108.20
34	BA	1526	C	C3'-C2'-C1'	-9.22	94.12	101.50
35	BB	1079	G	P-O3'-C3'	-9.22	108.63	119.70
85	AA	2181	G	C4-N9-C1'	-9.22	114.51	126.50
34	BA	650	C	O4'-C1'-N1	9.22	115.58	108.20
36	BC	130	U	O4'-C1'-N1	9.22	115.58	108.20
38	BE	117	A	C4-C5-C6	-9.22	112.39	117.00
39	BF	65	U	C2-N1-C1'	-9.22	106.64	117.70
80	Bu	30	TYR	CB-CG-CD2	-9.22	115.47	121.00
85	AA	624	A	P-O3'-C3'	9.22	130.76	119.70
85	AA	1106	A	C3'-C2'-C1'	-9.22	94.12	101.50
34	BA	464	U	O4'-C1'-N1	9.21	115.57	108.20
85	AA	1469	G	C5-C6-O6	-9.22	123.07	128.60
34	BA	1539	A	O5'-P-OP1	-9.21	97.41	105.70
35	BB	59	U	C5'-C4'-C3'	-9.21	101.26	116.00
85	AA	87	C	C6-N1-C2	-9.21	116.61	120.30
34	BA	82	A	P-O3'-C3'	-9.21	108.65	119.70
34	BA	1787	U	C5'-C4'-C3'	9.21	130.74	116.00
41	BH	4	U	P-O3'-C3'	-9.21	108.65	119.70
85	AA	2047	U	C2-N3-C4	-9.21	121.47	127.00
34	BA	1648	G	P-O5'-C5'	-9.21	106.16	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	715	G	O4'-C1'-N9	9.21	115.57	108.20
38	BE	139	U	C2-N1-C1'	-9.21	106.65	117.70
85	AA	293	A	P-O3'-C3'	9.21	130.75	119.70
85	AA	1912	U	P-O3'-C3'	9.21	130.75	119.70
85	AA	2064	A	N1-C6-N6	-9.21	113.07	118.60
85	AA	2127	G	C8-N9-C4	9.21	110.08	106.40
34	BA	692	U	O3'-P-O5'	9.21	121.50	104.00
34	BA	612	U	C5-C4-O4	-9.21	120.38	125.90
35	BB	1459	U	P-O5'-C5'	-9.21	106.17	120.90
85	AA	489	C	P-O3'-C3'	-9.21	108.65	119.70
85	AA	761	G	O4'-C1'-N9	9.21	115.57	108.20
85	AA	1159	C	O4'-C1'-N1	9.21	115.57	108.20
85	AA	1797	U	N1-C2-N3	9.21	120.42	114.90
34	BA	103	G	P-O3'-C3'	-9.21	108.65	119.70
34	BA	1317	U	P-O3'-C3'	-9.21	108.65	119.70
34	BA	1614	G	N9-C4-C5	-9.20	101.72	105.40
34	BA	1631	U	C2-N3-C4	-9.20	121.48	127.00
36	BC	94	C	C3'-C2'-C1'	-9.20	94.14	101.50
37	BD	95	G	N1-C2-N2	-9.21	107.92	116.20
47	BN	105	ARG	NE-CZ-NH1	9.20	124.90	120.30
85	AA	47	A	O4'-C1'-N9	9.21	115.56	108.20
85	AA	710	A	O4'-C1'-N9	9.21	115.56	108.20
85	AA	578	U	P-O5'-C5'	-9.20	106.17	120.90
34	BA	399	G	C5-N7-C8	-9.20	99.70	104.30
35	BB	466	A	N1-C6-N6	-9.20	113.08	118.60
35	BB	633	C	N3-C2-O2	-9.20	115.46	121.90
35	BB	1194	A	P-O3'-C3'	-9.20	108.66	119.70
37	BD	104	C	C3'-C2'-C1'	-9.20	94.14	101.50
35	BB	993	A	N9-C4-C5	-9.20	102.12	105.80
35	BB	1474	A	C4'-C3'-C2'	-9.20	93.40	102.60
36	BC	85	U	P-O3'-C3'	-9.20	108.66	119.70
37	BD	95	G	N3-C2-N2	9.20	126.34	119.90
85	AA	1008	C	O4'-C1'-N1	9.20	115.56	108.20
37	BD	78	C	C5-C4-N4	9.20	126.64	120.20
34	BA	7	U	N3-C2-O2	-9.20	115.76	122.20
34	BA	189	G	C5'-C4'-O4'	9.20	120.14	109.10
34	BA	994	G	N1-C6-O6	9.20	125.42	119.90
35	BB	969	C	O4'-C1'-N1	9.20	115.56	108.20
85	AA	696	G	N1-C6-O6	9.20	125.42	119.90
36	BC	32	U	P-O5'-C5'	9.20	135.61	120.90
85	AA	309	G	C5-C6-O6	-9.20	123.08	128.60
85	AA	2075	C	C5'-C4'-C3'	-9.19	101.29	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	164	C	C1'-O4'-C4'	-9.19	102.55	109.90
34	BA	642	U	C5-C4-O4	9.19	131.41	125.90
41	BH	88	C	O4'-C1'-N1	9.19	115.55	108.20
34	BA	557	U	C5'-C4'-O4'	9.19	120.13	109.10
34	BA	1524	G	C4-C5-C6	-9.19	113.29	118.80
34	BA	1568	A	P-O3'-C3'	9.19	130.73	119.70
38	BE	25	U	C6-N1-C1'	9.19	134.07	121.20
60	Ba	119	TYR	CB-CG-CD2	-9.19	115.49	121.00
34	BA	1225	A	P-O3'-C3'	-9.19	108.68	119.70
34	BA	1502	G	N9-C1'-C2'	-9.19	101.89	112.00
85	AA	368	C	C3'-C2'-C1'	-9.19	94.15	101.50
35	BB	1287	U	C5'-C4'-C3'	-9.18	101.31	116.00
85	AA	777	U	O4'-C1'-N1	9.18	115.55	108.20
34	BA	641	U	C5-C4-O4	9.18	131.41	125.90
15	AG	73	ARG	NE-CZ-NH1	9.18	124.89	120.30
38	BE	149	A	C5-N7-C8	-9.18	99.31	103.90
40	BG	128	U	C3'-C2'-C1'	-9.18	94.16	101.50
41	BH	45	G	C8-N9-C4	9.18	110.07	106.40
85	AA	869	A	C5'-C4'-C3'	-9.18	101.31	116.00
85	AA	923	A	C1'-O4'-C4'	-9.18	102.56	109.90
34	BA	241	U	P-O3'-C3'	-9.18	108.69	119.70
85	AA	2210	C	O4'-C1'-N1	9.18	115.54	108.20
34	BA	872	U	O4'-C1'-N1	9.18	115.54	108.20
34	BA	115	U	O4'-C1'-C2'	-9.17	96.63	105.80
41	BH	119	U	C6-N1-C1'	9.17	134.04	121.20
85	AA	639	C	C2-N1-C1'	9.17	128.89	118.80
85	AA	1190	G	C4-N9-C1'	-9.17	114.58	126.50
85	AA	1820	G	C5-C6-O6	-9.17	123.10	128.60
34	BA	3	G	C3'-C2'-C1'	-9.17	94.17	101.50
40	BG	171	A	O5'-C5'-C4'	-9.17	94.28	111.70
35	BB	4	C	C4'-C3'-C2'	-9.17	93.43	102.60
34	BA	11	U	O4'-C1'-N1	9.17	115.53	108.20
34	BA	1138	C	C2-N1-C1'	-9.17	108.72	118.80
35	BB	983	C	C1'-O4'-C4'	-9.17	102.57	109.90
35	BB	1326	U	P-O5'-C5'	9.17	135.56	120.90
35	BB	1503	U	O4'-C1'-N1	9.17	115.53	108.20
85	AA	1960	C	C2-N1-C1'	9.17	128.88	118.80
35	BB	1303	A	C5'-C4'-C3'	-9.16	101.34	116.00
35	BB	1361	A	C5'-C4'-C3'	-9.16	101.34	116.00
85	AA	987	C	C6-N1-C2	-9.16	116.63	120.30
86	AB	71	G	C5-C6-O6	-9.16	123.10	128.60
35	BB	1475	U	P-O5'-C5'	-9.16	106.25	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1535	G	C5'-C4'-C3'	-9.16	101.35	116.00
35	BB	90	G	C4-N9-C1'	-9.16	114.60	126.50
38	BE	11	A	P-O5'-C5'	-9.16	106.25	120.90
41	BH	66	G	C8-N9-C1'	9.16	138.90	127.00
85	AA	282	C	C5'-C4'-C3'	-9.16	101.35	116.00
85	AA	1128	G	C8-N9-C1'	9.16	138.90	127.00
34	BA	228	A	O4'-C1'-N9	9.15	115.52	108.20
34	BA	588	C	C5'-C4'-C3'	-9.15	101.35	116.00
35	BB	80	C	C6-N1-C2	-9.15	116.64	120.30
35	BB	706	G	C6-N1-C2	-9.15	119.61	125.10
35	BB	1203	C	C5'-C4'-O4'	9.15	120.08	109.10
35	BB	1511	U	C4'-C3'-C2'	-9.15	93.44	102.60
37	BD	80	G	C4-N9-C1'	-9.15	114.60	126.50
85	AA	519	A	P-O3'-C3'	9.15	130.69	119.70
85	AA	1829	C	C6-N1-C2	-9.15	116.64	120.30
85	AA	2051	G	O4'-C1'-N9	9.15	115.52	108.20
85	AA	2056	C	O4'-C1'-N1	9.15	115.52	108.20
34	BA	590	U	P-O3'-C3'	-9.15	108.72	119.70
34	BA	1447	C	O4'-C1'-N1	9.15	115.52	108.20
35	BB	87	G	N1-C6-O6	9.15	125.39	119.90
85	AA	361	U	O4'-C1'-N1	9.15	115.52	108.20
35	BB	789	G	O4'-C1'-N9	9.15	115.52	108.20
69	Bj	9	ARG	NE-CZ-NH1	9.15	124.88	120.30
34	BA	838	U	O4'-C1'-N1	9.15	115.52	108.20
34	BA	1609	U	N3-C2-O2	-9.15	115.80	122.20
85	AA	424	A	O4'-C1'-N9	9.15	115.52	108.20
85	AA	1958	C	C6-N1-C2	-9.15	116.64	120.30
35	BB	831	C	N1-C1'-C2'	-9.14	101.94	112.00
34	BA	678	C	O4'-C1'-N1	9.14	115.51	108.20
38	BE	26	G	N1-C6-O6	9.14	125.39	119.90
41	BH	130	G	N1-C6-O6	-9.14	114.41	119.90
85	AA	573	U	O4'-C1'-N1	9.14	115.52	108.20
85	AA	1904	C	C2-N3-C4	9.14	124.47	119.90
18	AJ	97	ARG	NE-CZ-NH1	9.14	124.87	120.30
35	BB	620	G	C4-N9-C1'	-9.14	114.62	126.50
85	AA	2199	G	O4'-C1'-N9	9.14	115.51	108.20
85	AA	201	U	P-O3'-C3'	-9.14	108.73	119.70
34	BA	555	C	O4'-C1'-N1	9.14	115.51	108.20
34	BA	1329	U	C3'-C2'-C1'	-9.14	94.19	101.50
35	BB	1356	G	N1-C2-N2	-9.14	107.98	116.20
34	BA	1168	C	P-O3'-C3'	9.13	130.66	119.70
34	BA	1540	C	C2-N1-C1'	9.14	128.85	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1515	C	C2-N1-C1'	-9.13	108.75	118.80
38	BE	130	G	C3'-C2'-C1'	-9.13	94.19	101.50
85	AA	117	C	C1'-O4'-C4'	-9.13	102.59	109.90
85	AA	363	A	C3'-C2'-C1'	-9.13	94.19	101.50
34	BA	1225	A	O4'-C1'-N9	9.13	115.51	108.20
35	BB	607	G	C4-N9-C1'	-9.13	114.63	126.50
41	BH	123	G	N1-C6-O6	9.13	125.38	119.90
85	AA	386	G	P-O3'-C3'	-9.13	108.74	119.70
34	BA	778	U	P-O5'-C5'	-9.13	106.29	120.90
35	BB	1035	C	C5'-C4'-C3'	-9.13	101.39	116.00
77	Br	158	ARG	NE-CZ-NH1	9.13	124.87	120.30
85	AA	303	A	O4'-C1'-N9	9.13	115.50	108.20
85	AA	501	A	C8-N9-C4	-9.13	102.15	105.80
85	AA	1755	U	C5'-C4'-C3'	9.13	130.61	116.00
40	BG	169	A	O4'-C1'-C2'	-9.13	96.67	105.80
13	AE	97	ARG	NE-CZ-NH1	9.13	124.86	120.30
35	BB	726	A	C5'-C4'-C3'	-9.13	101.40	116.00
39	BF	4	A	P-O3'-C3'	-9.13	108.75	119.70
48	BO	100	ARG	NE-CZ-NH1	9.13	124.86	120.30
34	BA	766	A	O4'-C1'-N9	9.12	115.50	108.20
34	BA	1485	U	N3-C4-O4	-9.12	113.01	119.40
35	BB	579	A	N1-C6-N6	9.12	124.08	118.60
85	AA	374	C	O5'-C5'-C4'	-9.12	94.36	111.70
85	AA	975	G	P-O5'-C5'	-9.12	106.30	120.90
34	BA	1597	G	C6-N1-C2	-9.12	119.63	125.10
34	BA	660	C	C6-N1-C2	-9.12	116.65	120.30
34	BA	851	C	N3-C2-O2	-9.12	115.52	121.90
35	BB	145	G	C4-N9-C1'	-9.12	114.64	126.50
37	BD	98	G	C5'-C4'-C3'	-9.12	101.41	116.00
35	BB	1484	A	O4'-C1'-N9	9.12	115.50	108.20
34	BA	89	G	C6-N1-C2	-9.12	119.63	125.10
34	BA	437	G	C4-N9-C1'	9.12	138.35	126.50
37	BD	71	G	C1'-O4'-C4'	-9.12	102.61	109.90
34	BA	1189	A	P-O5'-C5'	9.11	135.48	120.90
85	AA	1826	U	C5'-C4'-C3'	-9.11	101.42	116.00
34	BA	1277	G	C4-N9-C1'	-9.11	114.65	126.50
34	BA	403	A	O4'-C1'-N9	9.11	115.49	108.20
34	BA	525	A	O5'-P-OP2	-9.11	97.50	105.70
34	BA	1240	G	C4'-C3'-C2'	9.11	111.71	102.60
34	BA	1299	G	C6-N1-C2	-9.11	119.63	125.10
35	BB	1126	A	O4'-C1'-N9	9.11	115.49	108.20
37	BD	49	A	C5-C6-N6	9.11	130.99	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	815	G	N3-C4-C5	-9.11	124.04	128.60
85	AA	938	A	N1-C6-N6	9.11	124.07	118.60
34	BA	227	C	C5'-C4'-C3'	-9.11	101.43	116.00
35	BB	12	G	C4'-C3'-C2'	9.11	111.71	102.60
85	AA	70	U	C5'-C4'-C3'	-9.11	101.42	116.00
85	AA	2186	U	C5'-C4'-C3'	-9.11	101.42	116.00
85	AA	859	G	C3'-C2'-C1'	-9.11	94.21	101.50
34	BA	878	G	C4'-C3'-C2'	9.11	111.70	102.60
34	BA	1088	G	C8-N9-C4	9.11	110.04	106.40
35	BB	3	C	N3-C4-N4	9.11	124.37	118.00
41	BH	101	A	O4'-C1'-N9	9.11	115.48	108.20
69	Bj	9	ARG	NE-CZ-NH2	-9.11	115.75	120.30
40	BG	167	C	P-O3'-C3'	-9.10	108.78	119.70
34	BA	781	U	P-O3'-C3'	-9.10	108.78	119.70
34	BA	1300	G	C5-C6-O6	-9.10	123.14	128.60
35	BB	483	C	O4'-C1'-N1	9.10	115.48	108.20
35	BB	622	G	C5'-C4'-C3'	9.10	130.57	116.00
38	BE	90	G	C8-N9-C1'	9.10	138.83	127.00
40	BG	34	A	C8-N9-C4	9.10	109.44	105.80
85	AA	466	A	C6-N1-C2	9.10	124.06	118.60
35	BB	709	G	O4'-C1'-C2'	9.10	115.79	107.60
35	BB	1441	C	C3'-C2'-C1'	-9.10	94.22	101.50
85	AA	115	U	P-O3'-C3'	9.10	130.62	119.70
38	BE	157	C	O4'-C1'-N1	9.10	115.48	108.20
85	AA	788	G	O3'-P-O5'	-9.10	86.71	104.00
85	AA	1885	A	O4'-C1'-N9	9.10	115.48	108.20
34	BA	226	A	P-O3'-C3'	9.10	130.61	119.70
34	BA	1247	G	C5'-C4'-C3'	-9.10	101.45	116.00
35	BB	750	G	N1-C6-O6	9.10	125.36	119.90
38	BE	25	U	C5-C4-O4	-9.10	120.44	125.90
34	BA	316	G	P-O3'-C3'	9.09	130.61	119.70
34	BA	644	C	O4'-C1'-N1	9.09	115.47	108.20
34	BA	813	C	C6-N1-C2	-9.09	116.66	120.30
35	BB	1222	A	P-O3'-C3'	-9.09	108.79	119.70
69	Bj	114	VAL	O-C-N	-9.09	108.15	122.70
85	AA	1487	G	C4-N9-C1'	-9.09	114.68	126.50
34	BA	966	G	P-O3'-C3'	9.09	130.61	119.70
85	AA	330	C	C6-N1-C1'	-9.09	109.89	120.80
85	AA	1894	G	C5-C6-N1	9.09	116.05	111.50
85	AA	189	G	C8-N9-C1'	9.09	138.81	127.00
85	AA	374	C	C1'-O4'-C4'	-9.09	102.63	109.90
85	AA	1292	A	P-O5'-C5'	-9.09	106.36	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	641	C	O4'-C1'-N1	9.08	115.47	108.20
85	AA	699	U	C3'-C2'-C1'	-9.08	94.23	101.50
69	Bj	77	ARG	NE-CZ-NH1	9.08	124.84	120.30
85	AA	1786	G	N1-C6-O6	9.08	125.35	119.90
34	BA	1202	G	C4'-C3'-C2'	-9.08	93.52	102.60
40	BG	176	G	C6-N1-C2	-9.08	119.65	125.10
34	BA	230	A	C5'-C4'-O4'	-9.08	98.21	109.10
85	AA	774	C	C6-N1-C2	-9.08	116.67	120.30
34	BA	1616	A	O5'-P-OP2	-9.08	97.53	105.70
34	BA	1724	G	C3'-C2'-C1'	-9.08	94.24	101.50
85	AA	99	U	C5-C4-O4	-9.08	120.45	125.90
85	AA	541	A	C1'-O4'-C4'	-9.08	102.64	109.90
85	AA	1575	G	C4-N9-C1'	-9.07	114.70	126.50
85	AA	901	C	C2-N1-C1'	9.07	128.78	118.80
35	BB	1176	G	C1'-O4'-C4'	-9.07	102.64	109.90
34	BA	91	C	P-O5'-C5'	9.07	135.41	120.90
85	AA	2215	C	C5'-C4'-O4'	-9.07	98.22	109.10
85	AA	378	A	C4-C5-C6	-9.07	112.47	117.00
34	BA	10	G	C8-N9-C1'	9.07	138.79	127.00
38	BE	117	A	C5'-C4'-C3'	-9.07	101.49	116.00
40	BG	81	G	C8-N9-C1'	9.07	138.79	127.00
34	BA	469	C	C5'-C4'-C3'	-9.07	101.49	116.00
85	AA	143	U	C2-N3-C4	-9.06	121.56	127.00
35	BB	83	G	O4'-C1'-N9	9.06	115.45	108.20
35	BB	1232	A	C3'-C2'-C1'	-9.06	94.25	101.50
85	AA	930	G	P-O3'-C3'	9.06	130.58	119.70
34	BA	90	G	O5'-P-OP1	-9.06	97.55	105.70
34	BA	114	U	O4'-C1'-N1	9.06	115.45	108.20
34	BA	1539	A	N9-C1'-C2'	-9.06	102.03	112.00
37	BD	36	C	O4'-C1'-N1	9.06	115.45	108.20
41	BH	12	U	P-O3'-C3'	-9.06	108.83	119.70
85	AA	252	G	C5'-C4'-C3'	9.06	130.50	116.00
85	AA	1035	C	C5'-C4'-C3'	-9.06	101.50	116.00
85	AA	1289	U	C2-N1-C1'	-9.06	106.83	117.70
85	AA	1660	U	C4'-C3'-C2'	9.06	111.66	102.60
86	AB	7	A	P-O5'-C5'	9.06	135.40	120.90
34	BA	515	U	C4'-C3'-C2'	9.06	111.66	102.60
34	BA	869	C	OP1-P-OP2	9.06	133.19	119.60
85	AA	434	U	P-O3'-C3'	9.06	130.57	119.70
34	BA	658	C	P-O3'-C3'	9.06	130.57	119.70
34	BA	971	G	C6-N1-C2	-9.05	119.67	125.10
35	BB	91	G	N1-C6-O6	9.06	125.33	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1419	A	N1-C6-N6	9.05	124.03	118.60
34	BA	1518	A	C2-N3-C4	-9.05	106.07	110.60
36	BC	89	U	O4'-C1'-N1	9.05	115.44	108.20
37	BD	25	G	N9-C1'-C2'	-9.05	102.04	112.00
38	BE	203	C	N3-C4-N4	9.05	124.34	118.00
85	AA	930	G	N1-C6-O6	9.05	125.33	119.90
85	AA	1278	C	P-O5'-C5'	-9.05	106.41	120.90
85	AA	1877	G	C8-N9-C4	-9.05	102.78	106.40
34	BA	371	U	P-O3'-C3'	-9.05	108.84	119.70
40	BG	48	U	O4'-C1'-N1	9.05	115.44	108.20
40	BG	59	G	N1-C6-O6	9.05	125.33	119.90
34	BA	163	G	O4'-C1'-N9	9.05	115.44	108.20
34	BA	1658	G	O5'-P-OP2	-9.05	97.56	105.70
37	BD	10	C	P-O3'-C3'	9.05	130.56	119.70
85	AA	1096	G	C8-N9-C4	-9.05	102.78	106.40
34	BA	1616	A	P-O5'-C5'	-9.05	106.43	120.90
38	BE	32	U	C2'-C3'-O3'	9.04	129.40	109.50
40	BG	111	C	C6-N1-C2	-9.04	116.68	120.30
82	Bw	151	TYR	CB-CG-CD2	-9.05	115.57	121.00
85	AA	1236	G	O4'-C1'-N9	9.05	115.44	108.20
85	AA	2167	A	O4'-C1'-N9	9.04	115.44	108.20
34	BA	155	U	O4'-C4'-C3'	-9.04	94.96	104.00
34	BA	1170	A	N1-C6-N6	-9.04	113.17	118.60
34	BA	1192	A	C5'-C4'-C3'	-9.04	101.53	116.00
34	BA	1194	G	O4'-C1'-N9	9.04	115.43	108.20
34	BA	1724	G	C4-C5-C6	-9.04	113.38	118.80
37	BD	97	U	C2-N3-C4	-9.04	121.58	127.00
38	BE	117	A	O5'-C5'-C4'	-9.04	94.52	111.70
85	AA	201	U	P-O5'-C5'	9.04	135.37	120.90
85	AA	986	U	C2-N1-C1'	-9.04	106.85	117.70
34	BA	1365	U	P-O3'-C3'	9.04	130.55	119.70
34	BA	1322	A	C3'-C2'-C1'	-9.04	94.27	101.50
34	BA	1781	A	P-O3'-C3'	-9.04	108.85	119.70
36	BC	134	G	O4'-C1'-N9	9.04	115.43	108.20
85	AA	131	C	O4'-C1'-N1	9.04	115.43	108.20
35	BB	82	G	C4-N9-C1'	-9.04	114.75	126.50
37	BD	36	C	C3'-C2'-C1'	-9.04	94.27	101.50
85	AA	109	G	N1-C6-O6	-9.04	114.48	119.90
85	AA	1873	U	C2-N3-C4	-9.04	121.58	127.00
85	AA	233	C	O4'-C1'-N1	9.04	115.43	108.20
85	AA	905	C	O4'-C1'-N1	9.04	115.43	108.20
35	BB	1375	G	C5-C6-O6	9.04	134.02	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	95	U	C2-N1-C1'	-9.03	106.86	117.70
41	BH	53	C	O4'-C1'-N1	9.04	115.43	108.20
85	AA	1106	A	O3'-P-O5'	-9.04	86.83	104.00
50	BQ	164	ARG	NE-CZ-NH1	9.03	124.82	120.30
85	AA	1373	U	C2-N3-C4	-9.04	121.58	127.00
85	AA	1520	A	N1-C6-N6	9.04	124.02	118.60
34	BA	739	A	C8-N9-C4	-9.03	102.19	105.80
34	BA	875	G	P-O3'-C3'	9.03	130.54	119.70
35	BB	1491	G	C4-N9-C1'	-9.03	114.76	126.50
36	BC	18	G	C2-N3-C4	-9.03	107.38	111.90
39	BF	32	G	C4'-C3'-C2'	9.03	111.63	102.60
41	BH	29	G	N9-C1'-C2'	-9.03	102.06	112.00
41	BH	127	A	N1-C6-N6	-9.03	113.18	118.60
85	AA	240	A	P-O3'-C3'	-9.03	108.86	119.70
34	BA	437	G	C8-N9-C1'	-9.03	115.26	127.00
35	BB	382	U	C6-N1-C2	-9.03	115.58	121.00
85	AA	473	C	C1'-O4'-C4'	-9.03	102.68	109.90
34	BA	943	G	C8-N9-C1'	9.03	138.74	127.00
34	BA	1705	C	O3'-P-O5'	9.03	121.15	104.00
35	BB	1132	A	N1-C6-N6	-9.03	113.18	118.60
35	BB	1334	C	C4'-C3'-C2'	9.03	111.63	102.60
35	BB	1546	C	C6-N1-C2	-9.03	116.69	120.30
41	BH	77	G	N1-C6-O6	9.03	125.32	119.90
85	AA	245	A	P-O3'-C3'	9.03	130.53	119.70
85	AA	2089	G	C5'-C4'-C3'	9.03	130.44	116.00
85	AA	1799	C	N3-C4-N4	9.03	124.32	118.00
34	BA	27	G	C5'-C4'-C3'	-9.02	101.56	116.00
34	BA	678	C	C4'-C3'-C2'	-9.02	93.58	102.60
38	BE	69	C	C6-N1-C2	-9.02	116.69	120.30
85	AA	2095	U	C5'-C4'-C3'	9.02	130.44	116.00
34	BA	431	A	C5-C6-N6	-9.02	116.48	123.70
34	BA	1233	U	C2-N3-C4	-9.02	121.59	127.00
85	AA	729	U	O4'-C1'-N1	9.02	115.42	108.20
37	BD	63	C	P-O3'-C3'	-9.02	108.88	119.70
85	AA	1431	U	O4'-C1'-N1	9.02	115.42	108.20
34	BA	221	G	O4'-C1'-N9	9.02	115.42	108.20
35	BB	417	A	N1-C6-N6	9.02	124.01	118.60
83	Bx	46	ASP	C-N-CA	9.02	144.25	121.70
34	BA	547	C	OP1-P-OP2	-9.02	106.08	119.60
34	BA	744	G	C1'-O4'-C4'	-9.02	102.69	109.90
40	BG	33	G	N9-C1'-C2'	-9.02	102.08	112.00
85	AA	551	C	C4'-C3'-C2'	-9.02	93.58	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1464	C	C6-N1-C2	-9.02	116.69	120.30
52	BS	165	ARG	NE-CZ-NH1	9.02	124.81	120.30
34	BA	1262	A	C8-N9-C4	9.01	109.41	105.80
34	BA	749	G	C5-C6-O6	-9.01	123.19	128.60
35	BB	995	C	O5'-C5'-C4'	9.01	128.83	111.70
38	BE	132	U	P-O3'-C3'	-9.01	108.89	119.70
85	AA	383	C	P-O5'-C5'	9.01	135.32	120.90
85	AA	114	C	C6-N1-C2	-9.01	116.69	120.30
86	AB	3	C	C2-N1-C1'	9.01	128.71	118.80
35	BB	714	U	C6-N1-C2	-9.01	115.59	121.00
35	BB	840	C	P-O5'-C5'	9.01	135.32	120.90
85	AA	785	C	P-O5'-C5'	9.01	135.31	120.90
85	AA	984	A	C3'-C2'-C1'	-9.01	94.29	101.50
85	AA	1098	C	C1'-O4'-C4'	-9.01	102.69	109.90
34	BA	174	A	O3'-P-O5'	-9.01	86.89	104.00
35	BB	1474	A	C5'-C4'-O4'	-9.01	98.29	109.10
85	AA	716	G	O4'-C1'-N9	9.01	115.41	108.20
85	AA	765	U	O5'-P-OP1	-9.01	97.59	105.70
85	AA	1589	G	C8-N9-C1'	9.01	138.71	127.00
34	BA	514	U	O4'-C1'-N1	9.01	115.40	108.20
34	BA	745	A	C5-C6-N6	9.01	130.90	123.70
34	BA	1720	U	O4'-C1'-N1	9.01	115.40	108.20
85	AA	2172	A	C8-N9-C4	-9.01	102.20	105.80
34	BA	247	U	C6-N1-C2	-9.00	115.60	121.00
34	BA	681	G	O4'-C4'-C3'	9.00	113.30	106.10
85	AA	822	U	C2-N3-C4	-9.00	121.60	127.00
38	BE	186	C	O4'-C1'-N1	9.00	115.40	108.20
34	BA	814	C	O4'-C1'-N1	9.00	115.40	108.20
35	BB	362	A	C5-C6-N6	-9.00	116.50	123.70
85	AA	1191	G	O5'-P-OP2	-9.00	97.60	105.70
85	AA	2049	U	P-O3'-C3'	9.00	130.50	119.70
34	BA	212	A	P-O5'-C5'	9.00	135.29	120.90
34	BA	1223	C	O4'-C1'-N1	9.00	115.40	108.20
37	BD	75	G	C4-N9-C1'	-9.00	114.80	126.50
38	BE	21	C	O4'-C1'-N1	8.99	115.40	108.20
34	BA	1087	A	C8-N9-C4	8.99	109.40	105.80
35	BB	111	C	O4'-C1'-N1	8.99	115.39	108.20
40	BG	66	C	C4'-C3'-C2'	8.99	111.59	102.60
85	AA	1523	G	C5-C6-O6	-8.99	123.20	128.60
35	BB	1421	C	N3-C2-O2	-8.99	115.61	121.90
85	AA	25	C	O5'-P-OP2	-8.99	97.61	105.70
40	BG	102	G	N3-C4-C5	-8.99	124.11	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	559	C	P-O5'-C5'	8.99	135.28	120.90
35	BB	28	G	P-O5'-C5'	8.99	135.28	120.90
85	AA	2164	G	C5-C6-O6	-8.99	123.21	128.60
34	BA	605	G	N9-C4-C5	8.99	109.00	105.40
85	AA	1441	G	C5'-C4'-C3'	-8.99	101.62	116.00
34	BA	1491	U	O5'-C5'-C4'	8.99	128.77	111.70
34	BA	1496	G	C1'-O4'-C4'	-8.99	102.71	109.90
85	AA	390	U	O4'-C1'-N1	8.99	115.39	108.20
85	AA	479	C	O4'-C1'-N1	8.99	115.39	108.20
85	AA	575	G	O4'-C1'-C2'	8.99	115.69	107.60
85	AA	1301	C	O4'-C1'-N1	8.99	115.39	108.20
34	BA	1442	A	N3-C4-C5	-8.98	120.51	126.80
35	BB	845	C	O4'-C1'-N1	8.98	115.39	108.20
35	BB	1253	U	C2-N3-C4	-8.98	121.61	127.00
67	Bh	167	TYR	CB-CG-CD2	-8.98	115.61	121.00
5	A4	170	ARG	NE-CZ-NH1	8.98	124.79	120.30
34	BA	842	U	C5'-C4'-C3'	8.98	130.37	116.00
35	BB	850	U	C5'-C4'-C3'	8.98	130.37	116.00
85	AA	2084	U	C5'-C4'-C3'	-8.98	101.63	116.00
34	BA	213	A	C5'-C4'-C3'	8.98	130.37	116.00
34	BA	1739	G	C8-N9-C1'	8.98	138.67	127.00
35	BB	415	A	C5'-C4'-C3'	-8.98	101.63	116.00
35	BB	1506	C	O4'-C1'-N1	8.98	115.38	108.20
86	AB	19	G	N3-C2-N2	-8.98	113.61	119.90
34	BA	1223	C	N1-C2-O2	8.98	124.29	118.90
35	BB	1416	A	N1-C6-N6	-8.98	113.21	118.60
41	BH	120	C	C5'-C4'-C3'	8.98	130.36	116.00
62	Bc	70	ASN	N-CA-C	-8.97	86.77	111.00
85	AA	688	C	C5'-C4'-C3'	-8.97	101.64	116.00
34	BA	102	G	P-O3'-C3'	-8.97	108.93	119.70
34	BA	950	C	C6-N1-C2	-8.97	116.71	120.30
36	BC	5	U	C5'-C4'-C3'	-8.97	101.64	116.00
34	BA	1466	U	P-O3'-C3'	-8.97	108.94	119.70
38	BE	204	U	C3'-C2'-C1'	-8.97	94.32	101.50
66	Bg	49	ARG	NE-CZ-NH1	8.97	124.78	120.30
85	AA	875	C	O4'-C1'-N1	8.97	115.38	108.20
85	AA	1731	G	C5'-C4'-C3'	-8.97	101.65	116.00
34	BA	1011	G	C4-N9-C1'	-8.97	114.84	126.50
34	BA	1500	G	C5-C6-N1	8.97	115.98	111.50
40	BG	180	C	C5'-C4'-C3'	-8.97	101.65	116.00
34	BA	1525	G	O4'-C1'-N9	8.96	115.37	108.20
38	BE	201	A	C8-N9-C4	8.97	109.39	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	139	G	P-O5'-C5'	-8.96	106.56	120.90
36	BC	33	U	C5'-C4'-O4'	8.96	119.86	109.10
85	AA	575	G	C5-C6-O6	-8.96	123.22	128.60
34	BA	890	G	N3-C2-N2	8.96	126.17	119.90
35	BB	1507	U	O5'-P-OP2	-8.96	97.63	105.70
38	BE	198	A	C5-C6-N6	-8.96	116.53	123.70
85	AA	10	G	C5'-C4'-C3'	-8.96	101.66	116.00
7	A6	157	PHE	CB-CG-CD1	-8.96	114.53	120.80
85	AA	211	C	P-O3'-C3'	8.96	130.46	119.70
85	AA	2183	U	C5'-C4'-C3'	-8.96	101.66	116.00
38	BE	139	U	C1'-O4'-C4'	-8.96	102.73	109.90
53	BT	74	ARG	NE-CZ-NH2	8.96	124.78	120.30
34	BA	1817	G	C2-N3-C4	-8.96	107.42	111.90
61	Bb	4	ARG	NE-CZ-NH1	8.96	124.78	120.30
85	AA	1492	U	C2-N3-C4	-8.96	121.63	127.00
34	BA	386	A	P-O5'-C5'	8.95	135.23	120.90
35	BB	1187	G	C8-N9-C4	-8.95	102.82	106.40
35	BB	1293	C	P-O3'-C3'	-8.96	108.95	119.70
35	BB	1491	G	C8-N9-C1'	8.96	138.64	127.00
85	AA	1499	G	C3'-C2'-C1'	-8.96	94.34	101.50
85	AA	1599	G	C5-C6-O6	-8.96	123.23	128.60
85	AA	2123	U	C2-N1-C1'	-8.96	106.95	117.70
34	BA	121	A	C3'-C2'-C1'	-8.95	94.34	101.50
34	BA	538	G	N3-C2-N2	8.95	126.17	119.90
34	BA	1828	A	O4'-C1'-N9	8.95	115.36	108.20
38	BE	14	C	O4'-C1'-N1	8.95	115.36	108.20
86	AB	16	U	C5'-C4'-C3'	8.95	130.32	116.00
35	BB	380	G	C4-N9-C1'	-8.95	114.86	126.50
41	BH	123	G	C8-N9-C4	8.95	109.98	106.40
44	BK	154	ARG	NE-CZ-NH1	8.95	124.78	120.30
85	AA	516	G	C5-C6-O6	-8.95	123.23	128.60
34	BA	1519	G	C5-C6-O6	8.95	133.97	128.60
85	AA	736	U	O4'-C1'-N1	8.95	115.36	108.20
35	BB	807	U	P-O3'-C3'	-8.95	108.96	119.70
85	AA	179	G	C6-N1-C2	-8.95	119.73	125.10
36	BC	123	G	O3'-P-O5'	8.94	120.99	104.00
34	BA	229	C	C6-N1-C2	-8.94	116.72	120.30
34	BA	238	C	P-O5'-C5'	8.94	135.21	120.90
35	BB	1518	U	P-O3'-C3'	8.94	130.43	119.70
37	BD	91	U	O3'-P-O5'	-8.94	87.01	104.00
85	AA	613	G	C5-C6-O6	-8.94	123.23	128.60
85	AA	1100	U	O4'-C1'-N1	8.94	115.35	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1786	G	C5-C6-O6	-8.94	123.23	128.60
37	BD	51	G	C5'-C4'-C3'	-8.94	101.70	116.00
40	BG	110	U	C5'-C4'-C3'	8.94	130.30	116.00
35	BB	1546	C	O4'-C1'-N1	8.94	115.35	108.20
38	BE	8	G	N3-C4-C5	-8.94	124.13	128.60
85	AA	383	C	P-O3'-C3'	-8.94	108.97	119.70
85	AA	822	U	C2-N1-C1'	-8.94	106.97	117.70
35	BB	660	G	C5-C6-O6	-8.94	123.24	128.60
35	BB	963	G	C4-N9-C1'	-8.94	114.88	126.50
34	BA	283	U	O4'-C1'-N1	8.94	115.35	108.20
34	BA	820	C	P-O3'-C3'	-8.94	108.98	119.70
38	BE	102	U	C2-N1-C1'	8.94	128.42	117.70
52	BS	120	TYR	CB-CG-CD2	-8.94	115.64	121.00
52	BS	151	PHE	CB-CG-CD2	-8.94	114.55	120.80
85	AA	687	G	P-O3'-C3'	8.94	130.42	119.70
85	AA	1125	G	N9-C4-C5	-8.94	101.83	105.40
85	AA	2168	C	C5'-C4'-C3'	-8.94	101.70	116.00
34	BA	1735	G	C4-C5-C6	-8.93	113.44	118.80
35	BB	878	G	C1'-O4'-C4'	8.93	117.05	109.90
40	BG	7	U	P-O3'-C3'	-8.93	108.98	119.70
85	AA	165	C	O4'-C1'-N1	8.93	115.35	108.20
85	AA	310	U	N3-C2-O2	-8.93	115.95	122.20
85	AA	991	G	N3-C2-N2	-8.93	113.65	119.90
37	BD	78	C	N3-C4-N4	-8.93	111.75	118.00
85	AA	824	C	O4'-C1'-N1	8.93	115.34	108.20
34	BA	1697	U	C5-C6-N1	8.93	127.16	122.70
35	BB	1079	G	C8-N9-C1'	8.93	138.61	127.00
35	BB	1489	A	C1'-O4'-C4'	-8.93	102.76	109.90
38	BE	20	C	C6-N1-C1'	-8.93	110.08	120.80
85	AA	887	A	C1'-O4'-C4'	-8.93	102.75	109.90
85	AA	262	G	C5'-C4'-C3'	8.93	130.29	116.00
85	AA	1595	G	C5'-C4'-C3'	-8.93	101.71	116.00
85	AA	773	G	C5'-C4'-C3'	-8.93	101.72	116.00
85	AA	779	G	P-O3'-C3'	-8.93	108.99	119.70
35	BB	1205	A	N1-C6-N6	8.93	123.96	118.60
39	BF	8	C	O4'-C1'-N1	8.93	115.34	108.20
85	AA	1468	G	C5'-C4'-O4'	-8.93	98.39	109.10
85	AA	2036	A	P-O5'-C5'	8.93	135.18	120.90
34	BA	926	A	O5'-C5'-C4'	-8.93	94.74	111.70
85	AA	1813	C	O4'-C1'-N1	8.93	115.34	108.20
37	BD	63	C	O4'-C1'-N1	8.92	115.34	108.20
38	BE	89	G	C4-N9-C1'	-8.92	114.90	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1109	G	P-O3'-C3'	-8.92	108.99	119.70
34	BA	821	G	O4'-C1'-N9	8.92	115.34	108.20
85	AA	1124	G	C8-N9-C1'	8.92	138.60	127.00
85	AA	2227	A	C8-N9-C4	8.92	109.37	105.80
35	BB	1354	C	O4'-C1'-N1	8.92	115.33	108.20
35	BB	1544	A	C8-N9-C1'	8.92	143.76	127.70
85	AA	552	C	C6-N1-C2	-8.92	116.73	120.30
85	AA	1731	G	C4'-C3'-C2'	-8.92	93.68	102.60
34	BA	380	A	N1-C6-N6	-8.92	113.25	118.60
35	BB	1202	G	N3-C4-N9	8.92	131.35	126.00
85	AA	210	G	P-O3'-C3'	-8.92	109.00	119.70
38	BE	41	C	C5'-C4'-C3'	8.92	130.27	116.00
34	BA	406	G	C5-C6-O6	-8.91	123.25	128.60
34	BA	942	G	C5'-C4'-C3'	-8.91	101.74	116.00
54	BU	84	THR	N-CA-CB	8.91	127.24	110.30
85	AA	2210	C	C6-N1-C2	-8.91	116.73	120.30
35	BB	542	A	N1-C6-N6	8.91	123.95	118.60
41	BH	114	G	N1-C6-O6	8.91	125.25	119.90
85	AA	646	C	O4'-C1'-N1	8.91	115.33	108.20
85	AA	815	G	N3-C4-N9	8.91	131.35	126.00
85	AA	1058	G	N1-C6-O6	8.91	125.25	119.90
2	A1	95	ARG	NE-CZ-NH1	8.91	124.76	120.30
34	BA	747	G	C8-N9-C4	-8.91	102.84	106.40
36	BC	8	C	P-O3'-C3'	8.91	130.39	119.70
4	A3	64	MET	CG-SD-CE	-8.91	85.95	100.20
17	AI	17	ARG	NE-CZ-NH1	8.91	124.75	120.30
34	BA	763	U	C6-N1-C1'	-8.91	108.73	121.20
34	BA	1493	U	C5'-C4'-C3'	8.91	130.25	116.00
36	BC	38	U	C3'-C2'-C1'	-8.91	94.37	101.50
85	AA	1669	G	C4-N9-C1'	-8.91	114.92	126.50
39	BF	1	C	O4'-C1'-N1	8.91	115.33	108.20
34	BA	510	U	P-O3'-C3'	8.90	130.39	119.70
34	BA	585	G	P-O5'-C5'	-8.90	106.65	120.90
38	BE	72	C	O4'-C1'-N1	8.90	115.32	108.20
41	BH	33	G	C4-N9-C1'	-8.90	114.92	126.50
85	AA	786	G	O4'-C1'-N9	8.90	115.32	108.20
85	AA	1595	G	N1-C6-O6	8.90	125.24	119.90
85	AA	1832	G	C1'-O4'-C4'	-8.90	102.78	109.90
85	AA	2206	A	P-O3'-C3'	-8.90	109.01	119.70
39	BF	21	C	O4'-C1'-N1	8.90	115.32	108.20
85	AA	1139	G	C4'-C3'-C2'	-8.90	93.70	102.60
34	BA	524	G	P-O5'-C5'	8.90	135.14	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	971	A	C5'-C4'-O4'	-8.90	98.42	109.10
35	BB	1477	C	O4'-C1'-N1	8.90	115.32	108.20
40	BG	14	G	C8-N9-C1'	8.90	138.56	127.00
41	BH	72	G	C4-N9-C1'	-8.90	114.93	126.50
34	BA	608	G	C1'-O4'-C4'	8.89	117.02	109.90
85	AA	2194	U	P-O3'-C3'	8.89	130.37	119.70
34	BA	594	G	O5'-P-OP2	8.89	121.37	110.70
34	BA	961	C	C1'-O4'-C4'	-8.89	102.78	109.90
35	BB	148	C	C6-N1-C1'	-8.89	110.13	120.80
35	BB	691	A	C5-C6-N6	-8.89	116.58	123.70
58	BY	51	ARG	NE-CZ-NH1	8.89	124.75	120.30
15	AG	131	ARG	NE-CZ-NH1	8.89	124.75	120.30
34	BA	98	A	C5-C6-N6	-8.89	116.59	123.70
40	BG	118	U	N3-C2-O2	-8.89	115.98	122.20
41	BH	41	A	O4'-C1'-N9	8.89	115.31	108.20
52	BS	7	ARG	NE-CZ-NH1	8.89	124.75	120.30
85	AA	495	G	C5'-C4'-O4'	-8.89	98.43	109.10
34	BA	238	C	C5-C6-N1	8.89	125.44	121.00
34	BA	1675	C	O5'-C5'-C4'	-8.89	94.81	111.70
41	BH	39	G	N3-C4-C5	-8.89	124.16	128.60
85	AA	1617	G	P-O3'-C3'	8.89	130.37	119.70
34	BA	300	C	C4'-C3'-C2'	-8.89	93.71	102.60
34	BA	470	C	OP2-P-O3'	8.89	124.76	105.20
34	BA	690	G	C2'-C3'-O3'	8.89	129.06	109.50
34	BA	875	G	O5'-C5'-C4'	-8.89	94.81	111.70
34	BA	1071	G	C4-N9-C1'	-8.89	114.95	126.50
34	BA	1258	G	C5-C6-O6	-8.89	123.27	128.60
34	BA	1801	G	C8-N9-C1'	8.89	138.56	127.00
34	BA	1506	C	C4'-C3'-C2'	8.89	111.49	102.60
40	BG	179	C	C4'-C3'-C2'	8.89	111.49	102.60
85	AA	989	U	O5'-P-OP1	-8.89	97.70	105.70
34	BA	1211	G	C6-N1-C2	-8.89	119.77	125.10
38	BE	133	C	C5'-C4'-C3'	-8.89	101.78	116.00
34	BA	1736	A	O4'-C1'-N9	8.88	115.31	108.20
35	BB	480	C	O4'-C1'-N1	8.88	115.31	108.20
38	BE	178	G	P-O3'-C3'	-8.88	109.04	119.70
85	AA	82	A	C5'-C4'-O4'	8.88	119.76	109.10
85	AA	1698	A	P-O5'-C5'	-8.88	106.69	120.90
34	BA	124	G	C4-N9-C1'	-8.88	114.96	126.50
34	BA	63	A	P-O5'-C5'	-8.88	106.70	120.90
34	BA	1195	G	C4-N9-C1'	-8.88	114.96	126.50
37	BD	79	G	C5'-C4'-C3'	8.88	130.20	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	6	A	C8-N9-C4	8.88	109.35	105.80
62	Bc	107	TYR	CB-CG-CD1	8.88	126.33	121.00
38	BE	104	G	C8-N9-C4	-8.88	102.85	106.40
85	AA	400	G	C6-N1-C2	-8.88	119.77	125.10
85	AA	1683	U	C6-N1-C1'	-8.88	108.77	121.20
34	BA	232	U	O4'-C1'-N1	8.88	115.30	108.20
34	BA	622	G	C5-C6-O6	-8.88	123.28	128.60
38	BE	123	A	C2-N3-C4	-8.88	106.16	110.60
85	AA	886	A	C3'-C2'-C1'	-8.88	94.40	101.50
34	BA	395	G	N9-C1'-C2'	-8.87	102.24	112.00
35	BB	1496	C	C6-N1-C2	-8.87	116.75	120.30
85	AA	2132	A	P-O5'-C5'	-8.87	106.70	120.90
34	BA	1411	C	P-O3'-C3'	-8.87	109.06	119.70
35	BB	739	C	O4'-C1'-N1	8.87	115.30	108.20
48	BO	55	ARG	NE-CZ-NH1	8.87	124.74	120.30
85	AA	1184	A	C6-N1-C2	-8.87	113.28	118.60
36	BC	157	U	P-O5'-C5'	-8.87	106.71	120.90
62	Bc	76	MET	CG-SD-CE	-8.87	86.00	100.20
85	AA	1872	G	N1-C6-O6	8.87	125.22	119.90
85	AA	1003	G	C8-N9-C4	-8.87	102.85	106.40
85	AA	2048	C	C5'-C4'-C3'	8.87	130.20	116.00
34	BA	1299	G	O4'-C1'-N9	8.87	115.30	108.20
34	BA	1380	G	O4'-C1'-N9	8.87	115.30	108.20
38	BE	203	C	C6-N1-C1'	-8.87	110.16	120.80
34	BA	508	C	C2-N1-C1'	8.87	128.55	118.80
34	BA	578	C	C3'-C2'-C1'	8.87	108.59	101.50
34	BA	692	U	N1-C2-O2	-8.87	116.59	122.80
52	BS	42	ARG	NE-CZ-NH1	8.87	124.73	120.30
65	Bf	456	ARG	NE-CZ-NH2	-8.87	115.87	120.30
85	AA	1516	A	C4'-C3'-C2'	8.87	111.47	102.60
34	BA	980	C	P-O5'-C5'	-8.87	106.72	120.90
34	BA	1485	U	C2-N1-C1'	-8.87	107.06	117.70
35	BB	80	C	P-O3'-C3'	8.87	130.34	119.70
34	BA	398	G	N1-C6-O6	8.86	125.22	119.90
35	BB	818	U	C6-N1-C1'	8.86	133.61	121.20
34	BA	4	A	C5'-C4'-C3'	8.86	130.18	116.00
34	BA	75	U	O4'-C1'-N1	8.86	115.29	108.20
34	BA	540	G	C8-N9-C1'	8.86	138.52	127.00
34	BA	1707	C	C2-N3-C4	-8.86	115.47	119.90
40	BG	140	G	C5'-C4'-C3'	-8.86	101.82	116.00
85	AA	845	A	O4'-C1'-N9	8.86	115.29	108.20
34	BA	730	C	C6-N1-C2	-8.86	116.76	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	958	G	C4'-C3'-C2'	-8.86	93.74	102.60
35	BB	1235	A	P-O3'-C3'	8.86	130.33	119.70
40	BG	94	G	C8-N9-C4	8.86	109.94	106.40
85	AA	2157	G	C4-N9-C1'	-8.86	114.98	126.50
16	AH	43	ARG	NE-CZ-NH1	8.86	124.73	120.30
34	BA	568	G	C5'-C4'-C3'	-8.86	101.83	116.00
34	BA	1731	A	C5'-C4'-C3'	8.86	130.17	116.00
35	BB	638	G	C8-N9-C4	8.86	109.94	106.40
85	AA	919	U	P-O3'-C3'	-8.86	109.07	119.70
85	AA	1180	C	C4'-C3'-C2'	8.86	111.45	102.60
85	AA	1426	G	C5'-C4'-C3'	-8.86	101.83	116.00
34	BA	917	C	C5'-C4'-C3'	-8.85	101.84	116.00
34	BA	994	G	C5-C6-O6	-8.85	123.29	128.60
35	BB	815	G	O4'-C1'-N9	8.85	115.28	108.20
35	BB	1362	G	C5-C6-O6	-8.85	123.29	128.60
85	AA	982	G	N3-C2-N2	-8.85	113.70	119.90
85	AA	2105	G	C5-C6-O6	-8.85	123.29	128.60
34	BA	402	G	O4'-C1'-N9	8.85	115.28	108.20
36	BC	160	C	N3-C2-O2	-8.85	115.70	121.90
38	BE	13	A	C5'-C4'-O4'	8.85	119.72	109.10
85	AA	1793	A	C4-N9-C1'	-8.85	110.37	126.30
35	BB	1295	A	P-O3'-C3'	-8.85	109.08	119.70
41	BH	74	G	OP1-P-OP2	-8.85	106.33	119.60
45	BL	160	ARG	NE-CZ-NH1	8.85	124.72	120.30
85	AA	38	C	C1'-O4'-C4'	-8.85	102.82	109.90
34	BA	472	G	C8-N9-C4	-8.85	102.86	106.40
35	BB	1507	U	C1'-O4'-C4'	-8.85	102.82	109.90
37	BD	106	G	C6-N1-C2	-8.85	119.79	125.10
60	Ba	87	ASP	CB-CG-OD1	8.85	126.26	118.30
85	AA	467	U	O4'-C1'-N1	8.85	115.28	108.20
40	BG	173	C	O4'-C1'-N1	8.85	115.28	108.20
34	BA	10	G	O4'-C1'-C2'	8.84	115.56	107.60
34	BA	655	U	C2-N1-C1'	-8.84	107.09	117.70
38	BE	64	A	C5-C6-N6	-8.84	116.62	123.70
85	AA	1899	A	N1-C6-N6	8.84	123.91	118.60
34	BA	550	U	C5-C4-O4	8.84	131.20	125.90
34	BA	866	C	N3-C4-C5	-8.84	118.36	121.90
35	BB	411	A	P-O3'-C3'	-8.84	109.09	119.70
35	BB	708	C	P-O5'-C5'	-8.84	106.76	120.90
35	BB	1453	G	C6-C5-N7	-8.84	125.09	130.40
85	AA	2215	C	O4'-C1'-N1	8.84	115.27	108.20
34	BA	17	A	C4'-C3'-C2'	8.84	111.44	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	574	U	N3-C2-O2	-8.84	116.01	122.20
85	AA	1165	C	O4'-C1'-N1	8.84	115.27	108.20
85	AA	1927	G	C4-N9-C1'	-8.84	115.01	126.50
34	BA	1013	A	O4'-C1'-N9	8.84	115.27	108.20
35	BB	610	U	O3'-P-O5'	8.84	120.79	104.00
85	AA	2096	G	C5-C6-O6	-8.84	123.30	128.60
34	BA	1538	G	C6-N1-C2	-8.84	119.80	125.10
85	AA	539	A	C5-C6-N6	-8.84	116.63	123.70
85	AA	557	G	P-O5'-C5'	8.84	135.04	120.90
34	BA	527	C	P-O3'-C3'	-8.83	109.10	119.70
35	BB	800	U	C6-N1-C2	-8.83	115.70	121.00
35	BB	1546	C	C5'-C4'-C3'	-8.83	101.87	116.00
85	AA	305	A	P-O5'-C5'	8.83	135.03	120.90
85	AA	740	A	C5-C6-N6	8.83	130.77	123.70
34	BA	112	C	N3-C2-O2	-8.83	115.72	121.90
34	BA	660	C	N3-C2-O2	-8.83	115.72	121.90
35	BB	35	G	C6-N1-C2	-8.83	119.80	125.10
38	BE	112	G	O5'-P-OP1	8.83	121.30	110.70
85	AA	1191	G	C4-N9-C1'	-8.83	115.02	126.50
34	BA	1568	A	C5-C6-N6	-8.83	116.64	123.70
35	BB	1506	C	P-O5'-C5'	8.83	135.02	120.90
85	AA	857	G	C4-N9-C1'	-8.83	115.03	126.50
34	BA	1294	C	P-O3'-C3'	8.82	130.29	119.70
35	BB	1004	A	C8-N9-C4	-8.82	102.27	105.80
35	BB	1157	G	C5-C6-O6	-8.82	123.31	128.60
38	BE	92	C	C6-N1-C2	-8.82	116.77	120.30
39	BF	67	A	N1-C6-N6	8.82	123.89	118.60
85	AA	1286	C	P-O3'-C3'	8.82	130.29	119.70
34	BA	1093	G	O4'-C1'-N9	8.82	115.26	108.20
38	BE	13	A	C8-N9-C4	-8.82	102.27	105.80
20	AL	80	ARG	NE-CZ-NH2	-8.82	115.89	120.30
85	AA	457	G	C8-N9-C1'	8.82	138.47	127.00
85	AA	755	G	N7-C8-N9	8.82	117.51	113.10
36	BC	4	G	C4-N9-C1'	-8.82	115.03	126.50
85	AA	921	C	C2-N1-C1'	8.82	128.50	118.80
85	AA	1499	G	C8-N9-C1'	8.82	138.47	127.00
34	BA	1647	G	O4'-C1'-N9	8.82	115.25	108.20
35	BB	815	G	N9-C1'-C2'	-8.82	102.30	112.00
35	BB	989	C	C6-N1-C1'	-8.82	110.22	120.80
38	BE	30	C	C5'-C4'-C3'	8.82	130.11	116.00
38	BE	177	U	P-O5'-C5'	8.82	135.01	120.90
51	BR	42	ARG	NE-CZ-NH1	8.82	124.71	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1658	G	C5-C6-O6	-8.82	123.31	128.60
4	A3	91	TYR	CB-CG-CD1	8.81	126.29	121.00
34	BA	151	A	O5'-C5'-C4'	8.81	128.45	111.70
41	BH	113	G	C8-N9-C1'	8.81	138.46	127.00
85	AA	163	C	C2-N1-C1'	8.81	128.50	118.80
85	AA	1108	U	C6-N1-C2	-8.81	115.71	121.00
85	AA	1260	G	C4-N9-C1'	-8.81	115.04	126.50
85	AA	1553	G	C5-C6-O6	-8.81	123.31	128.60
85	AA	2152	C	O4'-C1'-N1	8.81	115.25	108.20
34	BA	548	G	OP1-P-OP2	-8.81	106.38	119.60
34	BA	1445	U	O4'-C1'-N1	8.81	115.25	108.20
34	BA	289	A	C4-N9-C1'	-8.81	110.44	126.30
38	BE	19	G	O5'-P-OP2	-8.81	97.77	105.70
85	AA	744	C	C2-N3-C4	8.81	124.30	119.90
85	AA	1645	G	C3'-C2'-C1'	-8.81	94.45	101.50
85	AA	1543	C	C5'-C4'-C3'	8.81	130.09	116.00
85	AA	2039	G	P-O3'-C3'	-8.81	109.13	119.70
39	BF	46	G	C4-N9-C1'	-8.81	115.05	126.50
40	BG	12	A	O4'-C1'-N9	8.81	115.25	108.20
34	BA	527	C	C1'-O4'-C4'	-8.80	102.86	109.90
85	AA	331	G	C8-N9-C4	-8.80	102.88	106.40
85	AA	790	A	C1'-O4'-C4'	8.81	116.94	109.90
34	BA	1224	A	C4'-C3'-C2'	8.80	111.40	102.60
35	BB	807	U	C2-N1-C1'	-8.80	107.14	117.70
37	BD	95	G	N7-C8-N9	8.80	117.50	113.10
65	Bf	398	ARG	NE-CZ-NH1	8.80	124.70	120.30
85	AA	1056	C	O4'-C1'-N1	8.80	115.24	108.20
34	BA	114	U	C5'-C4'-C3'	8.80	130.08	116.00
34	BA	1722	U	N3-C2-O2	-8.80	116.04	122.20
34	BA	1776	G	C4-N9-C1'	-8.80	115.06	126.50
40	BG	148	C	C1'-O4'-C4'	-8.80	102.86	109.90
85	AA	821	U	C2-N3-C4	-8.80	121.72	127.00
85	AA	1540	A	C5'-C4'-C3'	-8.80	101.92	116.00
85	AA	815	G	N3-C2-N2	8.80	126.06	119.90
34	BA	857	C	C5'-C4'-C3'	8.80	130.07	116.00
35	BB	803	U	C1'-O4'-C4'	-8.80	102.86	109.90
37	BD	84	U	N3-C2-O2	-8.80	116.04	122.20
85	AA	90	A	C1'-O4'-C4'	-8.80	102.86	109.90
35	BB	546	A	C6-N1-C2	-8.80	113.32	118.60
34	BA	435	U	C2-N3-C4	-8.79	121.72	127.00
38	BE	184	G	P-O5'-C5'	8.79	134.97	120.90
60	Ba	83	ARG	NE-CZ-NH1	8.79	124.70	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	273	C	N3-C2-O2	-8.79	115.74	121.90
85	AA	1818	C	C6-N1-C2	-8.79	116.78	120.30
4	A3	91	TYR	CB-CG-CD2	-8.79	115.72	121.00
23	AP	217	ARG	NE-CZ-NH2	8.79	124.70	120.30
34	BA	990	G	C5'-C4'-C3'	-8.79	101.93	116.00
34	BA	1722	U	C2-N1-C1'	-8.79	107.15	117.70
35	BB	643	G	N1-C6-O6	-8.79	114.62	119.90
35	BB	747	A	P-O3'-C3'	8.79	130.25	119.70
85	AA	707	U	C5'-C4'-C3'	-8.79	101.93	116.00
85	AA	1934	A	C5-C6-N6	-8.79	116.67	123.70
35	BB	1458	U	C5'-C4'-C3'	-8.79	101.93	116.00
38	BE	132	U	C6-N1-C1'	8.79	133.51	121.20
15	AG	91	LEU	CB-CA-C	8.79	126.90	110.20
34	BA	557	U	OP2-P-O3'	8.79	124.54	105.20
34	BA	1054	U	P-O3'-C3'	8.79	130.25	119.70
85	AA	48	G	C1'-O4'-C4'	-8.79	102.87	109.90
34	BA	1142	C	C5'-C4'-C3'	-8.79	101.94	116.00
85	AA	786	G	C5-C6-O6	-8.79	123.33	128.60
35	BB	607	G	C8-N9-C1'	8.79	138.42	127.00
35	BB	887	G	C8-N9-C4	-8.79	102.89	106.40
85	AA	1260	G	C8-N9-C1'	8.79	138.42	127.00
35	BB	265	C	O4'-C1'-N1	8.79	115.23	108.20
35	BB	576	A	O4'-C1'-N9	8.79	115.23	108.20
35	BB	1376	G	C5-C6-O6	-8.79	123.33	128.60
85	AA	613	G	N1-C6-O6	8.79	125.17	119.90
34	BA	610	A	P-O3'-C3'	-8.78	109.16	119.70
35	BB	1519	U	P-O3'-C3'	-8.78	109.16	119.70
13	AE	98	ARG	NE-CZ-NH1	8.78	124.69	120.30
34	BA	495	A	C1'-O4'-C4'	-8.78	102.88	109.90
34	BA	1126	U	C6-N1-C2	-8.78	115.73	121.00
35	BB	549	U	C2-N3-C4	-8.78	121.73	127.00
35	BB	768	A	P-O5'-C5'	8.78	134.95	120.90
85	AA	1105	G	O4'-C1'-N9	8.78	115.23	108.20
36	BC	127	C	O4'-C1'-N1	8.78	115.22	108.20
52	BS	47	MET	CG-SD-CE	-8.78	86.15	100.20
85	AA	301	U	O4'-C1'-N1	8.78	115.22	108.20
34	BA	1211	G	O5'-P-OP1	8.78	121.23	110.70
85	AA	831	C	O4'-C1'-N1	8.78	115.22	108.20
37	BD	98	G	P-O3'-C3'	8.78	130.23	119.70
34	BA	881	C	C6-N1-C1'	8.78	131.33	120.80
36	BC	137	C	C6-N1-C2	-8.78	116.79	120.30
34	BA	55	G	C6-N1-C2	-8.78	119.83	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	167	U	C6-N1-C2	-8.78	115.73	121.00
85	AA	414	C	C2-N1-C1'	-8.78	109.15	118.80
85	AA	1451	U	O4'-C1'-N1	8.78	115.22	108.20
35	BB	987	U	C2-N1-C1'	8.77	128.23	117.70
36	BC	76	C	O4'-C1'-N1	8.77	115.22	108.20
40	BG	118	U	C6-N1-C1'	8.77	133.48	121.20
34	BA	875	G	C5-C6-O6	-8.77	123.34	128.60
85	AA	850	U	P-O3'-C3'	-8.77	109.17	119.70
34	BA	1262	A	C4'-C3'-C2'	-8.77	93.83	102.60
85	AA	1874	G	O4'-C1'-N9	8.77	115.22	108.20
34	BA	1166	A	O4'-C1'-C2'	8.77	115.49	107.60
85	AA	644	A	P-O3'-C3'	8.77	130.22	119.70
85	AA	655	U	P-O3'-C3'	-8.77	109.18	119.70
85	AA	974	U	C4'-C3'-C2'	8.77	111.37	102.60
85	AA	1977	G	N1-C6-O6	8.77	125.16	119.90
34	BA	323	C	O4'-C4'-C3'	-8.77	95.23	104.00
50	BQ	176	ARG	NE-CZ-NH1	8.77	124.68	120.30
85	AA	1289	U	O4'-C1'-N1	8.77	115.21	108.20
85	AA	1973	G	C4-N9-C1'	-8.77	115.10	126.50
35	BB	1464	G	C8-N9-C1'	8.77	138.39	127.00
36	BC	8	C	N3-C4-C5	-8.77	118.39	121.90
85	AA	272	C	C1'-O4'-C4'	-8.77	102.89	109.90
85	AA	708	G	C5-C6-N1	8.77	115.88	111.50
34	BA	654	C	O4'-C1'-N1	8.76	115.21	108.20
34	BA	1668	C	C6-N1-C1'	8.76	131.31	120.80
34	BA	1570	C	C2-N3-C4	-8.76	115.52	119.90
34	BA	1711	G	C3'-C2'-C1'	-8.76	94.49	101.50
44	BK	3	ARG	NE-CZ-NH2	-8.76	115.92	120.30
85	AA	1004	G	P-O3'-C3'	-8.76	109.19	119.70
35	BB	771	U	O3'-P-O5'	-8.76	87.36	104.00
35	BB	782	A	N1-C6-N6	8.76	123.86	118.60
85	AA	475	A	C4-N9-C1'	-8.76	110.53	126.30
85	AA	1597	C	O4'-C1'-N1	8.76	115.21	108.20
85	AA	1920	A	P-O3'-C3'	8.76	130.21	119.70
34	BA	871	G	OP1-P-OP2	8.76	132.74	119.60
35	BB	439	G	C5'-C4'-C3'	-8.76	101.99	116.00
85	AA	851	G	C5-C6-O6	-8.76	123.34	128.60
41	BH	129	G	C5-C6-O6	8.76	133.85	128.60
85	AA	1363	U	C5'-C4'-C3'	-8.76	101.99	116.00
85	AA	635	G	C8-N9-C1'	8.76	138.38	127.00
35	BB	705	C	P-O3'-C3'	-8.75	109.19	119.70
35	BB	1545	U	O4'-C1'-N1	8.75	115.20	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	61	G	C8-N9-C1'	8.75	138.38	127.00
34	BA	672	G	C4'-C3'-C2'	-8.75	93.85	102.60
35	BB	1303	A	C4'-C3'-C2'	8.75	111.35	102.60
65	Bf	376	ARG	NE-CZ-NH2	8.75	124.68	120.30
85	AA	794	A	N1-C6-N6	8.75	123.85	118.60
85	AA	942	A	P-O5'-C5'	-8.75	106.90	120.90
85	AA	1840	C	P-O5'-C5'	8.75	134.90	120.90
4	A3	210	ARG	NE-CZ-NH1	8.75	124.67	120.30
38	BE	36	U	C2-N1-C1'	-8.75	107.20	117.70
40	BG	76	C	C5-C4-N4	-8.75	114.08	120.20
85	AA	847	G	C5'-C4'-O4'	8.75	119.60	109.10
85	AA	1198	U	N3-C2-O2	-8.75	116.08	122.20
85	AA	1539	A	N1-C6-N6	8.75	123.85	118.60
34	BA	98	A	C8-N9-C4	8.75	109.30	105.80
34	BA	137	C	C4'-C3'-C2'	-8.75	93.85	102.60
36	BC	16	A	C4-C5-C6	-8.75	112.63	117.00
85	AA	670	C	P-O3'-C3'	-8.75	109.20	119.70
85	AA	105	A	C1'-O4'-C4'	-8.75	102.90	109.90
85	AA	1268	C	N1-C1'-C2'	-8.75	102.38	112.00
34	BA	1257	U	P-O5'-C5'	-8.74	106.91	120.90
35	BB	494	C	O4'-C1'-N1	8.74	115.19	108.20
85	AA	180	A	C5-C6-N6	8.74	130.70	123.70
85	AA	958	C	O4'-C1'-N1	8.74	115.20	108.20
85	AA	2029	G	C5'-C4'-C3'	-8.74	102.01	116.00
35	BB	590	G	N3-C2-N2	8.74	126.02	119.90
35	BB	996	G	N3-C2-N2	8.74	126.02	119.90
36	BC	62	A	C2-N3-C4	-8.74	106.23	110.60
34	BA	591	G	O5'-C5'-C4'	-8.74	95.09	111.70
40	BG	112	C	C6-N1-C2	-8.74	116.80	120.30
85	AA	861	G	N1-C6-O6	8.74	125.14	119.90
34	BA	1745	G	C4-N9-C1'	-8.74	115.14	126.50
85	AA	1157	U	C2-N3-C4	-8.74	121.76	127.00
85	AA	1650	G	C5'-C4'-C3'	-8.74	102.02	116.00
85	AA	2249	U	C1'-O4'-C4'	-8.74	102.91	109.90
85	AA	873	U	C2-N3-C4	-8.74	121.76	127.00
15	AG	114	ARG	NE-CZ-NH1	8.74	124.67	120.30
34	BA	799	A	O4'-C1'-N9	8.74	115.19	108.20
34	BA	828	A	C5-C6-N6	8.74	130.69	123.70
35	BB	637	G	C5'-C4'-C3'	-8.74	102.02	116.00
36	BC	7	U	C6-N1-C1'	-8.74	108.97	121.20
85	AA	784	C	C6-N1-C2	-8.74	116.81	120.30
85	AA	814	G	N1-C6-O6	8.74	125.14	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1594	G	C5-C6-O6	-8.74	123.36	128.60
85	AA	2002	A	C8-N9-C4	-8.74	102.31	105.80
85	AA	1622	G	P-O3'-C3'	8.73	130.18	119.70
34	BA	299	C	C5'-C4'-C3'	-8.73	102.03	116.00
34	BA	638	U	O4'-C1'-N1	8.73	115.19	108.20
85	AA	967	C	O4'-C1'-N1	8.73	115.19	108.20
40	BG	105	A	C5-C6-N6	8.73	130.69	123.70
34	BA	1448	G	C5-C6-O6	-8.73	123.36	128.60
41	BH	1	U	P-O3'-C3'	-8.73	109.22	119.70
85	AA	292	C	O4'-C1'-N1	8.73	115.19	108.20
34	BA	1073	G	C6-N1-C2	-8.73	119.86	125.10
35	BB	329	U	O4'-C1'-N1	8.73	115.18	108.20
35	BB	1027	U	C5'-C4'-O4'	8.73	119.58	109.10
85	AA	2189	U	O4'-C1'-N1	8.73	115.19	108.20
35	BB	83	G	C5-C6-O6	-8.73	123.36	128.60
41	BH	36	C	C2-N1-C1'	-8.73	109.20	118.80
85	AA	104	C	C6-N1-C2	-8.73	116.81	120.30
85	AA	1720	C	P-O5'-C5'	-8.73	106.93	120.90
34	BA	356	C	O4'-C1'-N1	8.73	115.18	108.20
85	AA	82	A	O4'-C1'-N9	8.73	115.18	108.20
85	AA	459	C	O4'-C1'-N1	8.73	115.18	108.20
19	AK	88	ARG	NE-CZ-NH1	8.72	124.66	120.30
37	BD	78	C	C5'-C4'-C3'	-8.72	102.04	116.00
38	BE	90	G	C4'-C3'-C2'	8.72	111.33	102.60
38	BE	10	G	C4'-C3'-C2'	-8.72	93.88	102.60
72	Bm	63	ARG	NE-CZ-NH1	8.72	124.66	120.30
34	BA	1454	G	C4'-C3'-C2'	-8.72	93.88	102.60
34	BA	1832	A	C5'-C4'-C3'	-8.72	102.05	116.00
41	BH	9	C	C2-N1-C1'	8.72	128.39	118.80
85	AA	2124	G	P-O3'-C3'	8.72	130.16	119.70
34	BA	1477	C	C4'-C3'-C2'	-8.72	93.88	102.60
35	BB	15	C	C6-N1-C2	-8.72	116.81	120.30
85	AA	1927	G	C8-N9-C1'	8.72	138.34	127.00
85	AA	2089	G	C4-N9-C1'	-8.72	115.17	126.50
35	BB	1483	A	C8-N9-C4	-8.72	102.31	105.80
34	BA	659	U	P-O3'-C3'	8.71	130.16	119.70
85	AA	1163	G	C1'-O4'-C4'	-8.72	102.93	109.90
40	BG	14	G	C5-C6-N1	8.71	115.86	111.50
85	AA	3	U	C2-N1-C1'	8.71	128.16	117.70
85	AA	1260	G	P-O3'-C3'	-8.71	109.24	119.70
85	AA	1900	C	C5'-C4'-C3'	-8.71	102.06	116.00
34	BA	144	C	C2-N3-C4	-8.71	115.54	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	196	A	C4'-C3'-C2'	8.71	111.31	102.60
34	BA	977	G	C5'-C4'-C3'	-8.71	102.06	116.00
34	BA	1344	G	C1'-O4'-C4'	-8.71	102.93	109.90
34	BA	1657	A	N1-C6-N6	-8.71	113.37	118.60
35	BB	682	U	C2-N3-C4	-8.71	121.77	127.00
35	BB	1453	G	N9-C4-C5	-8.71	101.92	105.40
37	BD	95	G	C5-C6-N1	8.71	115.86	111.50
35	BB	804	U	N1-C2-N3	8.71	120.13	114.90
38	BE	185	G	C4-N9-C1'	-8.71	115.18	126.50
47	BN	44	ARG	NE-CZ-NH1	8.71	124.66	120.30
80	Bu	33	ARG	NE-CZ-NH1	8.71	124.66	120.30
41	BH	119	U	C4'-C3'-C2'	-8.71	93.89	102.60
85	AA	840	A	P-O3'-C3'	-8.71	109.25	119.70
35	BB	1167	C	O4'-C4'-C3'	8.71	113.07	106.10
35	BB	1419	G	C5-C6-O6	-8.71	123.38	128.60
86	AB	49	C	O4'-C1'-N1	8.71	115.17	108.20
34	BA	480	G	C5'-C4'-C3'	-8.71	102.07	116.00
34	BA	1650	G	C3'-C2'-C1'	-8.71	94.53	101.50
37	BD	83	A	C1'-O4'-C4'	-8.71	102.94	109.90
85	AA	1486	G	O4'-C1'-N9	8.71	115.17	108.20
35	BB	1356	G	C5-C6-O6	-8.71	123.38	128.60
38	BE	198	A	N1-C2-N3	-8.71	124.95	129.30
34	BA	1184	A	P-O3'-C3'	8.70	130.14	119.70
34	BA	316	G	C4-N9-C1'	-8.70	115.19	126.50
34	BA	657	C	C6-N1-C2	-8.70	116.82	120.30
35	BB	641	C	C6-N1-C2	-8.70	116.82	120.30
35	BB	839	G	N1-C6-O6	-8.70	114.68	119.90
38	BE	8	G	N3-C4-N9	8.70	131.22	126.00
63	Bd	61	GLU	OE1-CD-OE2	8.70	133.74	123.30
34	BA	1790	U	C2-N3-C4	-8.70	121.78	127.00
62	Bc	13	ARG	N-CA-CB	8.70	126.26	110.60
85	AA	243	A	O5'-P-OP2	8.70	121.14	110.70
34	BA	1478	G	P-O3'-C3'	8.70	130.14	119.70
34	BA	1493	U	O3'-P-O5'	8.70	120.53	104.00
40	BG	35	G	P-O3'-C3'	-8.70	109.26	119.70
40	BG	164	U	C6-N1-C2	-8.70	115.78	121.00
34	BA	508	C	C5'-C4'-O4'	8.70	119.54	109.10
34	BA	564	C	C4-C5-C6	8.70	121.75	117.40
35	BB	842	G	C8-N9-C1'	8.70	138.31	127.00
85	AA	1663	U	O4'-C1'-N1	8.70	115.16	108.20
35	BB	653	G	C4-N9-C1'	-8.70	115.20	126.50
85	AA	977	U	C5'-C4'-C3'	-8.70	102.09	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1086	U	O4'-C1'-N1	8.70	115.16	108.20
85	AA	1242	A	C5'-C4'-C3'	-8.70	102.08	116.00
85	AA	2020	C	C2-N1-C1'	8.70	128.37	118.80
40	BG	6	A	C5'-C4'-C3'	8.69	129.91	116.00
85	AA	860	C	O4'-C1'-N1	8.70	115.16	108.20
85	AA	1226	A	O4'-C4'-C3'	-8.69	95.31	104.00
85	AA	1912	U	O4'-C1'-N1	8.69	115.16	108.20
34	BA	538	G	O4'-C1'-N9	8.69	115.16	108.20
34	BA	637	G	C5-C6-O6	8.69	133.81	128.60
34	BA	1286	C	C2-N3-C4	8.69	124.25	119.90
85	AA	1496	U	C2-N3-C4	-8.69	121.79	127.00
34	BA	513	U	C6-N1-C2	-8.69	115.79	121.00
34	BA	1202	G	C1'-O4'-C4'	-8.69	102.95	109.90
65	Bf	167	ARG	NE-CZ-NH1	8.69	124.64	120.30
35	BB	836	U	N3-C4-O4	8.69	125.48	119.40
85	AA	2150	G	O4'-C1'-N9	8.69	115.15	108.20
34	BA	347	A	P-O3'-C3'	8.69	130.12	119.70
34	BA	1693	U	O5'-P-OP2	-8.69	97.88	105.70
34	BA	1743	U	O4'-C1'-N1	8.69	115.15	108.20
35	BB	816	U	C6-N1-C1'	8.69	133.36	121.20
35	BB	1424	G	C5'-C4'-C3'	-8.69	102.10	116.00
39	BF	64	U	C2-N3-C4	-8.69	121.79	127.00
41	BH	95	C	O4'-C1'-N1	8.69	115.15	108.20
85	AA	385	A	C3'-C2'-C1'	-8.69	94.55	101.50
85	AA	1116	G	C5'-C4'-C3'	-8.69	102.10	116.00
34	BA	232	U	P-O3'-C3'	-8.68	109.28	119.70
34	BA	429	G	N9-C4-C5	-8.68	101.93	105.40
85	AA	442	G	C5-C6-N1	8.68	115.84	111.50
85	AA	1812	C	C2-N3-C4	-8.68	115.56	119.90
35	BB	852	G	O5'-P-OP1	-8.68	97.89	105.70
34	BA	94	G	C5'-C4'-C3'	-8.68	102.11	116.00
34	BA	807	U	C6-N1-C2	-8.68	115.79	121.00
34	BA	1735	G	P-O3'-C3'	8.68	130.12	119.70
35	BB	656	A	P-O3'-C3'	-8.68	109.28	119.70
35	BB	1391	G	C5'-C4'-O4'	8.68	119.52	109.10
49	BP	120	ARG	NE-CZ-NH2	-8.68	115.96	120.30
34	BA	259	C	C2-N1-C1'	-8.68	109.26	118.80
34	BA	575	U	C6-N1-C2	-8.68	115.79	121.00
34	BA	1176	C	O5'-C5'-C4'	-8.68	95.21	111.70
34	BA	1658	G	P-O3'-C3'	-8.68	109.29	119.70
85	AA	1900	C	C2-N3-C4	-8.68	115.56	119.90
85	AA	2057	G	C5'-C4'-C3'	-8.68	102.12	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1716	A	C1'-O4'-C4'	-8.68	102.96	109.90
85	AA	1832	G	C3'-C2'-C1'	-8.68	94.56	101.50
85	AA	1956	C	P-O5'-C5'	-8.67	107.02	120.90
34	BA	606	G	C5'-C4'-C3'	-8.67	102.12	116.00
35	BB	1146	C	O4'-C1'-N1	8.67	115.14	108.20
39	BF	5	U	C2-N1-C1'	8.67	128.11	117.70
39	BF	11	C	C6-N1-C2	-8.67	116.83	120.30
85	AA	313	A	N1-C2-N3	-8.67	124.97	129.30
85	AA	632	U	P-O3'-C3'	8.67	130.10	119.70
85	AA	796	U	C5'-C4'-C3'	8.67	129.87	116.00
85	AA	709	A	O4'-C1'-N9	8.67	115.14	108.20
85	AA	1661	U	P-O3'-C3'	8.67	130.10	119.70
85	AA	1984	A	N1-C6-N6	8.67	123.80	118.60
2	A1	127	TYR	CA-CB-CG	-8.67	96.93	113.40
34	BA	1072	U	C5-C6-N1	-8.67	118.37	122.70
34	BA	27	G	P-O3'-C3'	-8.67	109.30	119.70
34	BA	678	C	C5-C6-N1	8.67	125.33	121.00
85	AA	357	C	C6-N1-C2	-8.67	116.83	120.30
85	AA	1482	C	P-O3'-C3'	-8.67	109.30	119.70
35	BB	1066	G	C5-C6-O6	-8.67	123.40	128.60
77	Br	312	ARG	NE-CZ-NH1	8.66	124.63	120.30
34	BA	606	G	N1-C2-N2	-8.66	108.40	116.20
35	BB	14	C	P-O3'-C3'	-8.66	109.30	119.70
35	BB	1018	U	C1'-O4'-C4'	-8.66	102.97	109.90
38	BE	127	G	C5'-C4'-C3'	8.66	129.86	116.00
85	AA	671	G	C5'-C4'-C3'	-8.66	102.14	116.00
34	BA	165	C	O4'-C1'-C2'	-8.66	97.14	105.80
34	BA	506	U	O4'-C1'-N1	8.66	115.13	108.20
34	BA	938	C	C4'-C3'-C2'	8.66	111.26	102.60
34	BA	1265	G	C8-N9-C1'	-8.66	115.74	127.00
35	BB	1460	G	N1-C6-O6	8.66	125.10	119.90
36	BC	113	G	N1-C6-O6	8.66	125.10	119.90
85	AA	1441	G	C5-C6-O6	-8.66	123.40	128.60
15	AG	55	ARG	NE-CZ-NH1	8.66	124.63	120.30
34	BA	186	G	P-O3'-C3'	-8.66	109.31	119.70
34	BA	447	U	C2-N1-C1'	-8.66	107.31	117.70
34	BA	1137	U	C5'-C4'-C3'	-8.66	102.15	116.00
35	BB	118	A	N1-C6-N6	8.66	123.80	118.60
38	BE	75	C	P-O5'-C5'	8.66	134.75	120.90
85	AA	414	C	N3-C2-O2	-8.66	115.84	121.90
85	AA	495	G	O4'-C1'-N9	8.66	115.13	108.20
42	BI	19	THR	N-CA-C	8.66	134.37	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	806	G	O4'-C1'-N9	8.66	115.12	108.20
34	BA	465	A	O4'-C1'-N9	8.65	115.12	108.20
34	BA	877	U	O4'-C1'-N1	8.65	115.12	108.20
85	AA	482	C	P-O3'-C3'	-8.65	109.31	119.70
34	BA	1563	G	C2'-C3'-O3'	8.65	128.54	109.50
34	BA	1568	A	O4'-C1'-N9	8.65	115.12	108.20
35	BB	600	C	O4'-C1'-N1	8.65	115.12	108.20
35	BB	1485	G	O4'-C1'-N9	8.65	115.12	108.20
85	AA	2200	A	P-O3'-C3'	-8.65	109.31	119.70
34	BA	223	U	N3-C2-O2	-8.65	116.14	122.20
34	BA	576	C	P-O5'-C5'	-8.65	107.06	120.90
34	BA	1157	A	P-O3'-C3'	-8.65	109.32	119.70
37	BD	80	G	C5-C6-O6	-8.65	123.41	128.60
34	BA	293	A	O4'-C1'-N9	8.65	115.12	108.20
35	BB	1200	A	O4'-C4'-C3'	-8.65	95.35	104.00
36	BC	51	A	O4'-C1'-N9	8.65	115.12	108.20
37	BD	70	C	C6-N1-C1'	8.65	131.18	120.80
40	BG	145	C	C6-N1-C2	-8.65	116.84	120.30
34	BA	143	A	O5'-C5'-C4'	-8.65	95.27	111.70
35	BB	1226	G	O4'-C1'-C2'	8.65	115.38	107.60
36	BC	156	A	C5-C6-N6	-8.65	116.78	123.70
85	AA	732	G	P-O5'-C5'	-8.65	107.06	120.90
34	BA	348	U	C5'-C4'-C3'	-8.65	102.17	116.00
35	BB	1547	U	C2-N3-C4	-8.65	121.81	127.00
36	BC	7	U	N1-C2-N3	-8.65	109.71	114.90
85	AA	1763	G	P-O5'-C5'	8.65	134.74	120.90
34	BA	1458	A	C5-C6-N6	-8.64	116.78	123.70
34	BA	1730	A	O4'-C4'-C3'	-8.64	95.36	104.00
35	BB	367	C	O4'-C1'-N1	8.64	115.12	108.20
36	BC	123	G	C6-N1-C2	-8.64	119.91	125.10
71	BI	138	ARG	NE-CZ-NH1	8.64	124.62	120.30
85	AA	2062	U	P-O5'-C5'	-8.64	107.07	120.90
85	AA	68	A	C1'-O4'-C4'	-8.64	102.99	109.90
85	AA	904	U	C5'-C4'-C3'	-8.64	102.17	116.00
34	BA	1202	G	C4-N9-C1'	-8.64	115.27	126.50
34	BA	1454	G	N9-C4-C5	-8.64	101.94	105.40
35	BB	1530	U	C2-N1-C1'	-8.64	107.33	117.70
38	BE	204	U	N3-C2-O2	-8.64	116.15	122.20
40	BG	21	C	O4'-C1'-C2'	-8.64	97.16	105.80
34	BA	1537	G	C5'-C4'-C3'	-8.64	102.18	116.00
34	BA	1822	U	P-O3'-C3'	-8.64	109.33	119.70
35	BB	516	G	P-O3'-C3'	8.64	130.07	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	528	G	C4-N9-C1'	-8.64	115.27	126.50
85	AA	239	G	P-O3'-C3'	-8.64	109.34	119.70
85	AA	1375	U	P-O3'-C3'	8.64	130.06	119.70
34	BA	387	A	P-O5'-C5'	8.64	134.72	120.90
35	BB	396	C	O4'-C1'-N1	8.63	115.11	108.20
35	BB	1181	A	C8-N9-C4	8.63	109.25	105.80
37	BD	35	C	O4'-C1'-N1	8.63	115.11	108.20
85	AA	802	A	O5'-P-OP2	-8.63	97.93	105.70
85	AA	1275	A	N1-C6-N6	-8.64	113.42	118.60
34	BA	479	U	C2-N3-C4	-8.63	121.82	127.00
34	BA	743	A	C5-C6-N1	-8.63	113.38	117.70
34	BA	1586	U	O4'-C1'-N1	8.63	115.10	108.20
34	BA	1702	G	P-O3'-C3'	-8.63	109.34	119.70
37	BD	79	G	C5-C6-O6	-8.63	123.42	128.60
38	BE	111	C	C5'-C4'-C3'	-8.63	102.19	116.00
83	Bx	49	ILE	C-N-CA	8.63	140.43	122.30
85	AA	931	G	C5-C6-O6	-8.63	123.42	128.60
67	Bh	150	ARG	NE-CZ-NH1	8.63	124.61	120.30
85	AA	1099	U	C6-N1-C2	-8.63	115.82	121.00
35	BB	571	C	O4'-C1'-N1	8.63	115.10	108.20
40	BG	166	C	N3-C4-N4	-8.63	111.96	118.00
34	BA	818	G	C5'-C4'-C3'	8.63	129.80	116.00
85	AA	690	G	N3-C2-N2	8.63	125.94	119.90
35	BB	1224	C	P-O3'-C3'	-8.62	109.35	119.70
38	BE	126	G	C4'-C3'-C2'	-8.62	93.98	102.60
85	AA	435	A	C5'-C4'-O4'	8.62	119.45	109.10
34	BA	9	A	C5'-C4'-C3'	-8.62	102.21	116.00
34	BA	244	A	C5'-C4'-C3'	-8.62	102.21	116.00
34	BA	853	A	P-O5'-C5'	8.62	134.69	120.90
34	BA	1618	A	C1'-O4'-C4'	-8.62	103.00	109.90
34	BA	1659	G	C5'-C4'-C3'	-8.62	102.20	116.00
85	AA	877	G	C6-N1-C2	-8.62	119.93	125.10
5	A4	188	PHE	CB-CG-CD1	-8.62	114.77	120.80
34	BA	740	A	C5-C6-N1	8.62	122.01	117.70
35	BB	1065	G	C5-C6-O6	-8.62	123.43	128.60
49	BP	17	ARG	NE-CZ-NH2	-8.62	115.99	120.30
60	Ba	60	SER	N-CA-CB	8.62	123.43	110.50
12	AD	17	PHE	CB-CG-CD1	8.62	126.83	120.80
34	BA	637	G	N1-C6-O6	-8.62	114.73	119.90
34	BA	673	U	P-O3'-C3'	8.62	130.04	119.70
34	BA	788	C	O4'-C1'-N1	8.62	115.09	108.20
34	BA	1011	G	C5-C6-O6	-8.62	123.43	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	895	U	C6-N1-C2	-8.62	115.83	121.00
41	BH	63	G	C4-N9-C1'	8.62	137.70	126.50
85	AA	1991	C	P-O5'-C5'	8.62	134.69	120.90
34	BA	11	U	C2-N3-C4	-8.62	121.83	127.00
34	BA	168	U	N3-C2-O2	-8.61	116.17	122.20
34	BA	1487	U	C2-N3-C4	-8.61	121.83	127.00
35	BB	323	C	O4'-C1'-N1	8.62	115.09	108.20
35	BB	1087	A	O3'-P-O5'	8.61	120.37	104.00
40	BG	10	U	C1'-O4'-C4'	-8.62	103.01	109.90
85	AA	1282	A	C4-N9-C1'	-8.62	110.79	126.30
85	AA	243	A	O5'-C5'-C4'	8.61	128.07	111.70
34	BA	180	G	P-O3'-C3'	-8.61	109.36	119.70
34	BA	1211	G	N7-C8-N9	8.61	117.41	113.10
34	BA	596	G	P-O3'-C3'	8.61	130.03	119.70
35	BB	96	A	C5-C6-N6	-8.61	116.81	123.70
35	BB	587	A	N9-C1'-C2'	-8.61	102.53	112.00
41	BH	58	C	C6-N1-C1'	8.61	131.13	120.80
85	AA	1128	G	N1-C6-O6	-8.61	114.73	119.90
85	AA	2161	C	C6-N1-C2	-8.61	116.86	120.30
85	AA	1174	G	C5-C6-O6	-8.61	123.43	128.60
34	BA	866	C	N3-C4-N4	8.61	124.03	118.00
34	BA	1556	A	C8-N9-C4	8.61	109.24	105.80
35	BB	26	C	C2-N3-C4	-8.61	115.60	119.90
35	BB	899	C	O4'-C1'-N1	8.61	115.09	108.20
40	BG	2	U	C3'-C2'-C1'	-8.61	94.61	101.50
41	BH	34	G	C8-N9-C4	-8.61	102.96	106.40
85	AA	1266	C	O4'-C1'-N1	8.61	115.09	108.20
85	AA	2215	C	C6-N1-C1'	8.61	131.13	120.80
34	BA	640	U	O4'-C1'-N1	8.61	115.09	108.20
34	BA	502	U	C1'-O4'-C4'	-8.61	103.02	109.90
34	BA	784	C	C4'-C3'-C2'	8.61	111.21	102.60
38	BE	129	G	O4'-C1'-C2'	-8.61	97.19	105.80
85	AA	1833	C	C6-N1-C2	-8.61	116.86	120.30
34	BA	134	U	C2-N3-C4	-8.60	121.84	127.00
34	BA	494	A	C8-N9-C4	8.60	109.24	105.80
35	BB	39	C	C3'-C2'-C1'	-8.60	94.62	101.50
36	BC	17	U	N3-C2-O2	8.60	128.22	122.20
40	BG	31	G	C6-N1-C2	-8.60	119.94	125.10
34	BA	3	G	C1'-O4'-C4'	-8.60	103.02	109.90
34	BA	417	A	C8-N9-C4	8.60	109.24	105.80
35	BB	652	G	C5'-C4'-C3'	-8.60	102.23	116.00
35	BB	1453	G	C4-C5-C6	-8.60	113.64	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	19	A	C4'-C3'-C2'	8.60	111.20	102.60
84	By	47	ASN	CB-CA-C	8.60	127.60	110.40
85	AA	1117	G	C4-N9-C1'	-8.60	115.32	126.50
34	BA	373	G	C4-N9-C1'	-8.60	115.32	126.50
34	BA	594	G	C8-N9-C1'	-8.60	115.82	127.00
34	BA	1433	U	O4'-C1'-C2'	8.60	115.34	107.60
35	BB	1441	C	C2-N1-C1'	8.60	128.26	118.80
36	BC	147	G	P-O5'-C5'	8.60	134.66	120.90
85	AA	1117	G	C8-N9-C1'	8.60	138.18	127.00
38	BE	11	A	N1-C6-N6	-8.60	113.44	118.60
35	BB	41	A	C5'-C4'-C3'	-8.60	102.25	116.00
35	BB	433	C	P-O3'-C3'	8.60	130.01	119.70
35	BB	762	C	C5-C4-N4	-8.60	114.18	120.20
85	AA	168	A	C8-N9-C4	-8.60	102.36	105.80
85	AA	787	U	C6-N1-C1'	-8.60	109.17	121.20
85	AA	1869	U	P-O5'-C5'	8.60	134.65	120.90
34	BA	837	U	P-O3'-C3'	8.59	130.01	119.70
35	BB	1061	G	N1-C6-O6	-8.59	114.75	119.90
40	BG	24	A	C2-N3-C4	-8.59	106.30	110.60
85	AA	719	C	O4'-C1'-N1	8.59	115.08	108.20
85	AA	858	G	C8-N9-C1'	8.59	138.17	127.00
50	BQ	221	ARG	NE-CZ-NH1	8.59	124.60	120.30
34	BA	527	C	C5'-C4'-C3'	-8.59	102.25	116.00
34	BA	991	U	N3-C2-O2	-8.59	116.19	122.20
85	AA	1190	G	C5-C6-O6	-8.59	123.45	128.60
11	AC	222	ARG	NE-CZ-NH2	-8.59	116.00	120.30
34	BA	489	A	P-O5'-C5'	8.59	134.64	120.90
40	BG	71	C	C2-N3-C4	-8.59	115.61	119.90
47	BN	187	TYR	CA-CB-CG	8.59	129.72	113.40
51	BR	23	ARG	NE-CZ-NH1	8.59	124.59	120.30
54	BU	20	ARG	NE-CZ-NH1	8.59	124.59	120.30
85	AA	1098	C	P-O5'-C5'	-8.59	107.16	120.90
85	AA	2033	C	P-O3'-C3'	-8.59	109.39	119.70
85	AA	2139	G	C5-C6-O6	-8.59	123.45	128.60
34	BA	384	U	C3'-C2'-C1'	-8.59	94.63	101.50
36	BC	114	C	C5'-C4'-C3'	-8.59	102.26	116.00
85	AA	1990	U	O4'-C1'-N1	8.59	115.07	108.20
34	BA	1290	A	O4'-C1'-N9	8.59	115.07	108.20
38	BE	127	G	C8-N9-C1'	8.59	138.16	127.00
39	BF	5	U	N3-C2-O2	-8.59	116.19	122.20
40	BG	110	U	P-O3'-C3'	-8.59	109.40	119.70
85	AA	983	A	N1-C6-N6	8.59	123.75	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	245	U	C2-N3-C4	-8.58	121.85	127.00
34	BA	864	G	C2-N3-C4	8.58	116.19	111.90
36	BC	167	U	P-O3'-C3'	8.58	130.00	119.70
60	Ba	35	ARG	NE-CZ-NH1	8.58	124.59	120.30
34	BA	206	C	O4'-C1'-N1	8.58	115.06	108.20
34	BA	748	C	C5'-C4'-C3'	-8.58	102.27	116.00
85	AA	210	G	C5-C6-N1	8.58	115.79	111.50
85	AA	1458	G	N3-C4-C5	-8.58	124.31	128.60
34	BA	895	U	P-O3'-C3'	8.58	129.99	119.70
34	BA	1783	C	O4'-C1'-N1	8.58	115.06	108.20
35	BB	96	A	N1-C6-N6	8.58	123.75	118.60
34	BA	53	G	C8-N9-C4	8.58	109.83	106.40
34	BA	208	A	O4'-C1'-N9	8.58	115.06	108.20
35	BB	964	G	O4'-C1'-N9	8.58	115.06	108.20
85	AA	252	G	C6-N1-C2	-8.58	119.95	125.10
85	AA	656	U	O5'-P-OP2	-8.58	97.98	105.70
35	BB	1457	A	P-O3'-C3'	-8.57	109.41	119.70
85	AA	964	C	C5'-C4'-O4'	8.57	119.39	109.10
85	AA	42	G	C5-C6-O6	-8.57	123.46	128.60
85	AA	1358	A	P-O3'-C3'	8.57	129.99	119.70
85	AA	1674	G	C4'-C3'-C2'	8.57	111.17	102.60
20	AL	81	ARG	NE-CZ-NH2	-8.57	116.01	120.30
34	BA	197	A	P-O3'-C3'	-8.57	109.42	119.70
34	BA	739	A	C2'-C3'-O3'	8.57	128.36	109.50
35	BB	62	C	O3'-P-O5'	8.57	120.29	104.00
37	BD	52	U	C5'-C4'-C3'	-8.57	102.28	116.00
34	BA	819	G	N9-C1'-C2'	-8.57	102.57	112.00
35	BB	957	A	O4'-C1'-N9	8.57	115.06	108.20
85	AA	1389	G	P-O3'-C3'	8.57	129.99	119.70
37	BD	102	C	C1'-O4'-C4'	-8.57	103.05	109.90
40	BG	112	C	C5'-C4'-C3'	-8.57	102.29	116.00
85	AA	183	C	O4'-C1'-N1	8.57	115.06	108.20
85	AA	687	G	C6-N1-C2	-8.57	119.96	125.10
85	AA	743	C	O3'-P-O5'	8.57	120.28	104.00
85	AA	1371	C	C5'-C4'-C3'	8.57	129.71	116.00
34	BA	1649	A	C4-C5-C6	-8.57	112.72	117.00
35	BB	1488	G	C4-N9-C1'	-8.57	115.36	126.50
34	BA	373	G	P-O5'-C5'	-8.57	107.19	120.90
34	BA	550	U	O5'-P-OP2	-8.57	97.99	105.70
35	BB	65	A	C5'-C4'-C3'	8.57	129.71	116.00
77	Br	181	ARG	NE-CZ-NH1	8.57	124.58	120.30
85	AA	1290	G	N1-C6-O6	8.57	125.04	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	34	G	P-O3'-C3'	-8.56	109.42	119.70
34	BA	920	U	O4'-C1'-N1	8.56	115.05	108.20
35	BB	125	G	C3'-C2'-C1'	-8.56	94.65	101.50
35	BB	812	G	C8-N9-C1'	8.56	138.13	127.00
35	BB	1397	G	N1-C6-O6	-8.56	114.76	119.90
35	BB	34	G	C5'-C4'-C3'	-8.56	102.31	116.00
35	BB	788	U	C2-N1-C1'	-8.56	107.43	117.70
35	BB	1204	C	O4'-C1'-N1	8.56	115.05	108.20
41	BH	105	U	C6-N1-C2	-8.56	115.86	121.00
44	BK	3	ARG	NE-CZ-NH1	8.56	124.58	120.30
85	AA	79	G	C4-N9-C1'	-8.56	115.38	126.50
85	AA	771	A	C5'-C4'-C3'	-8.56	102.31	116.00
34	BA	662	U	P-O3'-C3'	-8.56	109.43	119.70
35	BB	1416	A	C4'-C3'-C2'	8.56	111.16	102.60
85	AA	532	G	C4-N9-C1'	-8.56	115.38	126.50
85	AA	1175	A	O4'-C1'-N9	8.55	115.04	108.20
35	BB	845	C	C2-N1-C1'	-8.55	109.39	118.80
62	Bc	127	ARG	CD-NE-CZ	-8.55	111.63	123.60
85	AA	2064	A	C5-C6-N6	8.55	130.54	123.70
85	AA	2212	U	C5'-C4'-C3'	-8.55	102.31	116.00
35	BB	106	A	N1-C6-N6	8.55	123.73	118.60
41	BH	73	A	N1-C2-N3	-8.55	125.02	129.30
2	A1	186	ARG	NE-CZ-NH1	8.55	124.58	120.30
34	BA	609	G	N9-C1'-C2'	-8.55	102.59	112.00
35	BB	1308	G	C5-C6-O6	-8.55	123.47	128.60
35	BB	1485	G	C8-N9-C1'	8.55	138.12	127.00
41	BH	38	G	C4'-C3'-C2'	8.55	111.15	102.60
85	AA	581	A	C5'-C4'-C3'	-8.55	102.32	116.00
35	BB	709	G	C4'-C3'-C2'	8.55	111.15	102.60
36	BC	108	A	C1'-O4'-C4'	-8.55	103.06	109.90
85	AA	80	G	C4-N9-C1'	-8.55	115.39	126.50
85	AA	2112	G	P-O3'-C3'	-8.55	109.44	119.70
35	BB	1391	G	C5-C6-O6	-8.55	123.47	128.60
41	BH	101	A	C2-N3-C4	8.55	114.87	110.60
6	A5	7	ARG	NE-CZ-NH1	8.54	124.57	120.30
35	BB	778	A	C1'-O4'-C4'	-8.54	103.06	109.90
35	BB	818	U	N3-C2-O2	-8.54	116.22	122.20
35	BB	1487	G	C5-C6-O6	-8.54	123.47	128.60
53	BT	181	ARG	NE-CZ-NH1	8.54	124.57	120.30
85	AA	970	U	C4'-C3'-C2'	-8.54	94.06	102.60
34	BA	871	G	C2-N3-C4	-8.54	107.63	111.90
35	BB	782	A	C5-C6-N6	-8.54	116.86	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	657	A	O5'-P-OP1	-8.54	98.01	105.70
85	AA	375	C	C6-N1-C2	-8.54	116.88	120.30
85	AA	1576	G	P-O3'-C3'	8.54	129.95	119.70
85	AA	1863	A	N1-C6-N6	-8.54	113.47	118.60
34	BA	610	A	C5-C6-N6	-8.54	116.87	123.70
35	BB	1497	C	O4'-C1'-N1	8.54	115.03	108.20
37	BD	67	C	C6-N1-C2	-8.54	116.88	120.30
40	BG	146	C	P-O3'-C3'	8.54	129.95	119.70
85	AA	865	G	N1-C6-O6	8.54	125.02	119.90
85	AA	1137	C	P-O3'-C3'	8.54	129.94	119.70
85	AA	2108	C	O4'-C1'-N1	8.54	115.03	108.20
85	AA	2242	U	P-O3'-C3'	8.54	129.95	119.70
27	AT	83	TYR	CB-CG-CD1	8.54	126.12	121.00
34	BA	239	C	C2-N3-C4	-8.54	115.63	119.90
34	BA	1202	G	N3-C2-N2	8.54	125.88	119.90
34	BA	1451	A	P-O3'-C3'	-8.54	109.46	119.70
35	BB	128	C	C2-N1-C1'	8.54	128.19	118.80
35	BB	1357	C	C2-N1-C1'	8.53	128.19	118.80
41	BH	66	G	C4-N9-C1'	-8.54	115.40	126.50
41	BH	100	A	C2-N3-C4	8.53	114.87	110.60
85	AA	1	G	P-O3'-C3'	-8.54	109.46	119.70
85	AA	765	U	O4'-C1'-N1	8.54	115.03	108.20
85	AA	1441	G	N3-C2-N2	-8.53	113.93	119.90
85	AA	1974	C	C2-N1-C1'	8.54	128.19	118.80
1	A0	30	TYR	CB-CG-CD1	-8.53	115.88	121.00
14	AF	97	ARG	NE-CZ-NH1	8.53	124.57	120.30
34	BA	71	G	P-O3'-C3'	8.53	129.94	119.70
34	BA	310	C	P-O5'-C5'	8.53	134.55	120.90
34	BA	1222	C	C2-N3-C4	-8.53	115.63	119.90
34	BA	1811	A	P-O5'-C5'	8.53	134.55	120.90
35	BB	333	C	O4'-C1'-N1	8.53	115.03	108.20
36	BC	13	U	P-O3'-C3'	8.53	129.94	119.70
37	BD	20	C	P-O3'-C3'	8.53	129.94	119.70
37	BD	59	G	C5'-C4'-C3'	-8.53	102.35	116.00
38	BE	168	C	C6-N1-C1'	8.53	131.04	120.80
85	AA	741	G	N3-C2-N2	8.53	125.87	119.90
85	AA	1864	G	C5-C6-O6	-8.53	123.48	128.60
37	BD	107	G	C8-N9-C1'	8.53	138.09	127.00
31	AX	193	MET	CG-SD-CE	-8.53	86.56	100.20
77	Br	120	ARG	NE-CZ-NH2	-8.53	116.04	120.30
85	AA	516	G	P-O5'-C5'	-8.53	107.25	120.90
85	AA	1573	A	P-O5'-C5'	8.53	134.54	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1872	G	P-O3'-C3'	-8.53	109.47	119.70
85	AA	2070	C	C5'-C4'-C3'	-8.53	102.36	116.00
2	A1	143	ARG	NE-CZ-NH1	8.53	124.56	120.30
34	BA	14	G	C5-C6-O6	-8.53	123.48	128.60
57	BX	95	MET	CG-SD-CE	-8.53	86.56	100.20
85	AA	247	G	C5-C6-O6	-8.53	123.48	128.60
34	BA	401	A	C5-C6-N6	8.52	130.52	123.70
34	BA	1468	U	N3-C2-O2	-8.52	116.23	122.20
35	BB	444	U	C5'-C4'-C3'	8.52	129.64	116.00
35	BB	959	C	C6-N1-C2	-8.52	116.89	120.30
85	AA	1731	G	O3'-P-O5'	-8.52	87.81	104.00
85	AA	1830	U	C5'-C4'-C3'	-8.52	102.36	116.00
34	BA	921	G	P-O3'-C3'	-8.52	109.47	119.70
77	Br	43	MET	CG-SD-CE	-8.52	86.56	100.20
85	AA	1625	C	O4'-C1'-N1	8.52	115.02	108.20
34	BA	469	C	O4'-C1'-N1	8.52	115.01	108.20
35	BB	1227	G	O4'-C1'-N9	8.52	115.01	108.20
35	BB	1356	G	C6-N1-C2	-8.52	119.99	125.10
36	BC	121	G	O3'-P-O5'	-8.52	87.82	104.00
85	AA	339	A	C4-N9-C1'	-8.52	110.97	126.30
85	AA	1122	U	C2-N1-C1'	-8.52	107.48	117.70
34	BA	410	G	O3'-P-O5'	-8.51	87.82	104.00
34	BA	911	G	N1-C6-O6	-8.51	114.79	119.90
34	BA	1284	G	N1-C6-O6	-8.51	114.79	119.90
34	BA	1454	G	C6-C5-N7	-8.51	125.29	130.40
35	BB	512	C	C6-N1-C2	-8.51	116.89	120.30
35	BB	1284	U	N3-C2-O2	-8.51	116.24	122.20
36	BC	88	A	C1'-O4'-C4'	-8.51	103.09	109.90
85	AA	942	A	O5'-C5'-C4'	8.51	127.87	111.70
85	AA	1505	G	C6-N1-C2	-8.51	119.99	125.10
85	AA	1644	G	P-O5'-C5'	8.51	134.52	120.90
85	AA	15	U	N3-C2-O2	-8.51	116.24	122.20
85	AA	877	G	O4'-C1'-N9	8.51	115.01	108.20
34	BA	232	U	O4'-C4'-C3'	-8.51	95.49	104.00
36	BC	140	U	C2-N3-C4	-8.51	121.89	127.00
85	AA	28	A	C5'-C4'-C3'	8.51	129.61	116.00
41	BH	50	A	C8-N9-C4	-8.51	102.40	105.80
85	AA	567	G	O3'-P-O5'	-8.51	87.84	104.00
85	AA	819	G	N9-C1'-C2'	-8.51	102.64	112.00
85	AA	925	G	C8-N9-C1'	8.51	138.06	127.00
85	AA	1229	G	C3'-C2'-C1'	-8.51	94.69	101.50
34	BA	626	G	O4'-C1'-N9	8.51	115.00	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	49	C	C2-N1-C1'	8.51	128.16	118.80
85	AA	1350	A	P-O3'-C3'	8.51	129.91	119.70
6	A5	31	ARG	NE-CZ-NH2	-8.50	116.05	120.30
34	BA	431	A	N1-C6-N6	8.50	123.70	118.60
34	BA	867	C	N3-C4-C5	8.50	125.30	121.90
34	BA	944	G	P-O3'-C3'	8.50	129.90	119.70
35	BB	1141	A	C5'-C4'-O4'	8.50	119.30	109.10
85	AA	89	C	O4'-C1'-N1	8.50	115.00	108.20
85	AA	1345	C	P-O3'-C3'	8.50	129.90	119.70
85	AA	2054	G	C5'-C4'-C3'	-8.50	102.40	116.00
34	BA	228	A	C8-N9-C4	-8.50	102.40	105.80
34	BA	243	C	O4'-C1'-N1	8.50	115.00	108.20
34	BA	1202	G	P-O3'-C3'	-8.50	109.50	119.70
34	BA	1307	U	C6-N1-C2	-8.50	115.90	121.00
35	BB	1243	A	C8-N9-C4	8.50	109.20	105.80
35	BB	1475	U	C4'-C3'-C2'	-8.50	94.10	102.60
34	BA	254	U	N1-C2-O2	8.50	128.75	122.80
34	BA	548	G	C5'-C4'-O4'	-8.50	98.90	109.10
34	BA	605	G	N3-C4-N9	8.50	131.10	126.00
34	BA	684	G	O3'-P-O5'	8.50	120.14	104.00
34	BA	1283	U	C3'-C2'-C1'	-8.50	94.70	101.50
34	BA	1673	G	N3-C2-N2	8.50	125.85	119.90
34	BA	1844	U	O4'-C1'-N1	8.50	115.00	108.20
35	BB	1291	G	N3-C2-N2	8.50	125.85	119.90
58	BY	11	PHE	CB-CG-CD1	8.50	126.75	120.80
38	BE	175	U	C4'-C3'-O3'	8.50	129.99	113.00
41	BH	54	U	N3-C2-O2	-8.50	116.25	122.20
34	BA	129	U	C5'-C4'-C3'	8.49	129.59	116.00
34	BA	1081	U	P-O5'-C5'	8.49	134.49	120.90
35	BB	1479	C	O4'-C1'-N1	8.49	114.99	108.20
65	Bf	165	PHE	CB-CG-CD2	-8.49	114.86	120.80
85	AA	392	G	O4'-C1'-C2'	8.49	115.24	107.60
85	AA	736	U	C2-N1-C1'	-8.49	107.51	117.70
85	AA	2129	U	O4'-C1'-N1	8.49	114.99	108.20
34	BA	80	U	P-O3'-C3'	8.49	129.89	119.70
34	BA	1250	C	P-O3'-C3'	8.49	129.89	119.70
40	BG	16	G	N3-C4-C5	-8.49	124.36	128.60
85	AA	412	G	C5'-C4'-C3'	-8.49	102.42	116.00
20	AL	29	TYR	CA-CB-CG	8.49	129.53	113.40
34	BA	202	A	O5'-C5'-C4'	-8.49	95.58	111.70
34	BA	952	G	C4-N9-C1'	-8.49	115.47	126.50
34	BA	1301	G	C5-C6-O6	-8.49	123.51	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	137	A	P-O3'-C3'	8.49	129.88	119.70
78	Bs	52	ARG	NE-CZ-NH1	8.49	124.54	120.30
83	Bx	51	ARG	N-CA-C	8.49	133.92	111.00
85	AA	122	A	C5'-C4'-C3'	8.49	129.58	116.00
85	AA	1581	C	P-O3'-C3'	-8.49	109.52	119.70
85	AA	1974	C	C6-N1-C2	-8.49	116.91	120.30
34	BA	251	U	O3'-P-O5'	-8.48	87.88	104.00
34	BA	1610	A	P-O3'-C3'	-8.48	109.52	119.70
35	BB	1486	C	C6-N1-C2	-8.48	116.91	120.30
86	AB	55	U	O4'-C1'-N1	8.48	114.99	108.20
34	BA	174	A	C6-C5-N7	-8.48	126.36	132.30
34	BA	481	A	N3-C4-N9	-8.48	120.61	127.40
34	BA	1119	A	C5'-C4'-C3'	-8.48	102.43	116.00
40	BG	148	C	O4'-C4'-C3'	8.48	112.89	106.10
85	AA	511	A	P-O3'-C3'	-8.48	109.52	119.70
85	AA	1788	U	O4'-C1'-N1	8.48	114.99	108.20
34	BA	874	G	O5'-P-OP1	8.48	120.88	110.70
35	BB	687	C	P-O5'-C5'	8.48	134.47	120.90
35	BB	847	U	C1'-O4'-C4'	8.48	116.69	109.90
41	BH	74	G	O5'-P-OP1	8.48	120.88	110.70
85	AA	579	U	P-O5'-C5'	-8.48	107.33	120.90
27	AT	67	PHE	CB-CG-CD2	8.48	126.73	120.80
21	AM	133	ARG	NE-CZ-NH1	8.48	124.54	120.30
34	BA	269	G	C5'-C4'-O4'	8.48	119.27	109.10
34	BA	815	C	O4'-C1'-N1	8.48	114.98	108.20
34	BA	894	G	N9-C4-C5	-8.48	102.01	105.40
34	BA	1707	C	C5-C4-N4	8.48	126.13	120.20
35	BB	1130	U	C2-N1-C1'	-8.48	107.53	117.70
85	AA	767	A	P-O3'-C3'	8.48	129.87	119.70
34	BA	1354	G	C5-C6-O6	-8.47	123.52	128.60
35	BB	41	A	C6-N1-C2	-8.47	113.52	118.60
35	BB	381	C	P-O3'-C3'	8.47	129.87	119.70
35	BB	1547	U	O4'-C1'-N1	8.47	114.98	108.20
38	BE	10	G	C5'-C4'-C3'	-8.47	102.44	116.00
85	AA	1897	A	C6-N1-C2	-8.47	113.52	118.60
85	AA	2188	C	C4'-C3'-C2'	8.47	111.07	102.60
35	BB	847	U	P-O5'-C5'	8.47	134.46	120.90
34	BA	1721	U	O4'-C4'-C3'	-8.47	95.53	104.00
35	BB	524	C	C2-N1-C1'	8.47	128.12	118.80
36	BC	95	A	P-O3'-C3'	-8.47	109.53	119.70
37	BD	92	G	C5-C6-N1	8.47	115.74	111.50
37	BD	106	G	C5'-C4'-C3'	-8.47	102.45	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	9	C	C1'-O4'-C4'	-8.47	103.12	109.90
85	AA	889	G	C8-N9-C4	-8.47	103.01	106.40
85	AA	1116	G	C5-C6-O6	8.47	133.68	128.60
36	BC	124	A	C4-N9-C1'	8.47	141.54	126.30
85	AA	1491	G	O4'-C1'-N9	8.47	114.97	108.20
35	BB	1027	U	C6-N1-C2	-8.47	115.92	121.00
38	BE	141	A	P-O3'-C3'	-8.47	109.54	119.70
40	BG	104	A	C8-N9-C4	-8.47	102.41	105.80
85	AA	1255	C	C3'-C2'-C1'	-8.47	94.72	101.50
39	BF	60	C	P-O5'-C5'	8.47	134.45	120.90
41	BH	125	U	P-O3'-C3'	-8.47	109.54	119.70
34	BA	250	G	C5'-C4'-O4'	8.47	119.26	109.10
34	BA	596	G	N1-C6-O6	8.46	124.98	119.90
85	AA	234	G	N1-C6-O6	8.47	124.98	119.90
34	BA	1571	C	P-O3'-C3'	8.46	129.86	119.70
35	BB	999	G	N9-C1'-C2'	-8.46	102.69	112.00
37	BD	20	C	C6-N1-C2	-8.46	116.91	120.30
34	BA	558	C	O5'-P-OP1	-8.46	98.08	105.70
34	BA	892	C	N3-C2-O2	-8.46	115.98	121.90
35	BB	585	U	C2-N3-C4	-8.46	121.92	127.00
85	AA	2060	G	C4-N9-C1'	-8.46	115.50	126.50
34	BA	110	C	C6-N1-C2	-8.46	116.92	120.30
34	BA	185	A	N1-C6-N6	8.46	123.68	118.60
35	BB	1025	A	P-O3'-C3'	8.46	129.85	119.70
36	BC	39	G	P-O5'-C5'	8.46	134.44	120.90
65	Bf	463	ARG	NE-CZ-NH2	-8.46	116.07	120.30
85	AA	251	A	C2'-C3'-O3'	8.46	128.11	109.50
85	AA	347	U	P-O5'-C5'	-8.46	107.36	120.90
34	BA	621	G	C8-N9-C1'	8.46	138.00	127.00
38	BE	129	G	C5'-C4'-O4'	8.46	119.25	109.10
85	AA	1056	C	C2-N1-C1'	8.46	128.10	118.80
35	BB	65	A	O4'-C1'-N9	8.46	114.97	108.20
50	BQ	221	ARG	NE-CZ-NH2	-8.46	116.07	120.30
58	BY	21	TYR	CB-CG-CD2	-8.46	115.93	121.00
85	AA	495	G	C1'-O4'-C4'	-8.46	103.13	109.90
85	AA	737	G	C4-N9-C1'	8.46	137.49	126.50
34	BA	736	G	C1'-O4'-C4'	-8.46	103.14	109.90
34	BA	892	C	N3-C4-N4	-8.46	112.08	118.00
35	BB	792	G	C5-C6-O6	-8.46	123.53	128.60
35	BB	1431	G	N1-C6-O6	-8.45	114.83	119.90
35	BB	1482	A	C4'-C3'-C2'	-8.46	94.14	102.60
40	BG	118	U	C5-C6-N1	-8.45	118.47	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	270	U	C1'-O4'-C4'	-8.45	103.14	109.90
34	BA	331	G	C5'-C6-O6	8.45	133.67	128.60
34	BA	486	G	O3'-P-O5'	8.45	120.06	104.00
34	BA	1088	G	C5'-C6-O6	-8.45	123.53	128.60
34	BA	1116	G	P-O3'-C3'	8.45	129.84	119.70
41	BH	14	C	C2-N1-C1'	8.45	128.10	118.80
85	AA	1133	C	C5'-C4'-O4'	8.45	119.24	109.10
34	BA	1488	C	C5'-C4'-C3'	8.45	129.52	116.00
85	AA	1997	G	C5'-C4'-C3'	8.45	129.52	116.00
34	BA	888	G	C1'-O4'-C4'	-8.45	103.14	109.90
34	BA	1661	U	O4'-C1'-C2'	-8.45	97.35	105.80
34	BA	1835	A	O4'-C1'-N9	8.45	114.96	108.20
85	AA	900	G	P-O5'-C5'	8.45	134.42	120.90
85	AA	1996	A	C6-N1-C2	8.45	123.67	118.60
34	BA	468	A	O3'-P-O5'	-8.45	87.95	104.00
34	BA	502	U	C2-N3-C4	-8.45	121.93	127.00
34	BA	782	C	C2-N3-C4	-8.45	115.68	119.90
34	BA	992	A	P-O3'-C3'	-8.45	109.56	119.70
37	BD	72	U	P-O3'-C3'	8.45	129.84	119.70
85	AA	423	G	P-O3'-C3'	-8.45	109.56	119.70
85	AA	569	A	C4-N9-C1'	-8.45	111.10	126.30
35	BB	847	U	O4'-C4'-C3'	-8.44	95.56	104.00
35	BB	1141	A	O5'-C5'-C4'	8.45	127.75	111.70
38	BE	67	A	P-O3'-C3'	-8.44	109.57	119.70
85	AA	1719	C	C5'-C4'-O4'	8.44	119.23	109.10
86	AB	19	G	C5'-C4'-C3'	-8.44	102.49	116.00
34	BA	881	C	N3-C2-O2	-8.44	115.99	121.90
39	BF	37	C	C6-N1-C2	-8.44	116.92	120.30
58	BY	11	PHE	CB-CG-CD2	-8.44	114.89	120.80
85	AA	65	A	O4'-C1'-N9	8.44	114.95	108.20
34	BA	1592	U	P-O3'-C3'	-8.44	109.58	119.70
39	BF	46	G	C8-N9-C1'	8.44	137.97	127.00
85	AA	839	C	O4'-C1'-N1	8.44	114.95	108.20
85	AA	1162	A	C5'-C4'-O4'	8.44	119.22	109.10
85	AA	1715	C	C5'-C4'-C3'	-8.44	102.50	116.00
85	AA	1720	C	C6-N1-C2	-8.44	116.93	120.30
85	AA	2251	U	C2-N1-C1'	8.44	127.82	117.70
34	BA	1311	G	C5'-C4'-C3'	8.43	129.49	116.00
39	BF	54	U	C6-N1-C2	-8.43	115.94	121.00
85	AA	2008	G	C2-N3-C4	8.43	116.12	111.90
52	BS	89	TYR	CB-CG-CD1	8.43	126.06	121.00
35	BB	743	C	C6-N1-C2	-8.43	116.93	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1388	A	C5-C6-N1	8.43	121.92	117.70
85	AA	961	U	O4'-C1'-N1	8.43	114.94	108.20
85	AA	1670	U	C2-N3-C4	8.43	132.06	127.00
34	BA	1819	U	P-O5'-C5'	-8.43	107.42	120.90
40	BG	37	G	C8-N9-C1'	8.43	137.96	127.00
85	AA	1002	G	O4'-C1'-N9	8.43	114.94	108.20
41	BH	28	U	C6-N1-C2	-8.43	115.94	121.00
34	BA	253	U	O4'-C1'-N1	8.43	114.94	108.20
36	BC	25	C	N3-C4-N4	8.43	123.90	118.00
41	BH	57	A	O3'-P-O5'	-8.43	87.99	104.00
41	BH	120	C	P-O3'-C3'	8.43	129.81	119.70
85	AA	209	C	O4'-C1'-N1	8.43	114.94	108.20
85	AA	489	C	C6-N1-C2	-8.43	116.93	120.30
34	BA	50	G	C8-N9-C4	8.42	109.77	106.40
34	BA	843	G	C5'-C4'-C3'	-8.42	102.52	116.00
34	BA	1480	C	O4'-C1'-N1	8.42	114.94	108.20
35	BB	1361	A	C6-N1-C2	-8.42	113.55	118.60
83	Bx	198	ARG	NE-CZ-NH2	-8.42	116.09	120.30
85	AA	1284	A	P-O3'-C3'	-8.42	109.59	119.70
34	BA	117	C	P-O3'-C3'	-8.42	109.60	119.70
34	BA	814	C	P-O3'-C3'	-8.42	109.60	119.70
37	BD	80	G	P-O3'-C3'	-8.42	109.60	119.70
38	BE	24	G	P-O5'-C5'	-8.42	107.43	120.90
48	BO	115	ARG	NE-CZ-NH1	8.42	124.51	120.30
85	AA	124	A	C8-N9-C4	-8.42	102.43	105.80
85	AA	327	G	N9-C1'-C2'	-8.42	102.74	112.00
85	AA	967	C	P-O3'-C3'	-8.42	109.60	119.70
85	AA	1237	A	C5'-C4'-O4'	8.42	119.20	109.10
34	BA	1135	U	P-O3'-C3'	-8.42	109.60	119.70
34	BA	1335	A	O5'-C5'-C4'	-8.42	95.71	111.70
35	BB	1098	G	P-O3'-C3'	8.42	129.80	119.70
38	BE	199	A	O4'-C1'-N9	8.42	114.93	108.20
41	BH	75	G	P-O3'-C3'	8.42	129.80	119.70
53	BT	80	ARG	NE-CZ-NH1	8.42	124.51	120.30
85	AA	1282	A	C8-N9-C1'	8.42	142.85	127.70
85	AA	2062	U	C4'-C3'-C2'	8.42	111.02	102.60
4	A3	220	ARG	NE-CZ-NH1	8.41	124.51	120.30
34	BA	1078	U	C5'-C4'-C3'	-8.41	102.53	116.00
35	BB	617	C	O4'-C1'-N1	8.41	114.93	108.20
77	Br	160	TYR	CB-CG-CD2	-8.41	115.95	121.00
80	Bu	204	ARG	NE-CZ-NH1	8.41	124.51	120.30
85	AA	86	G	C5-C6-O6	-8.41	123.55	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	641	A	N1-C6-N6	-8.41	113.55	118.60
85	AA	1667	C	C6-N1-C2	-8.41	116.94	120.30
13	AE	113	TYR	CB-CG-CD2	-8.41	115.95	121.00
34	BA	336	A	C5-C6-N6	-8.41	116.97	123.70
34	BA	754	G	C6-N1-C2	-8.41	120.05	125.10
34	BA	780	U	C6-N1-C2	-8.41	115.95	121.00
34	BA	1260	G	P-O3'-C3'	-8.41	109.61	119.70
35	BB	1231	U	O4'-C4'-C3'	-8.41	95.59	104.00
37	BD	29	C	C1'-O4'-C4'	-8.41	103.17	109.90
38	BE	155	C	C2-N1-C1'	8.41	128.05	118.80
85	AA	749	C	P-O3'-C3'	8.41	129.79	119.70
41	BH	28	U	C5-C4-O4	-8.41	120.86	125.90
41	BH	123	G	C4-N9-C1'	-8.41	115.57	126.50
34	BA	1057	C	O4'-C1'-N1	8.41	114.93	108.20
39	BF	65	U	C5-C6-N1	-8.41	118.50	122.70
85	AA	987	C	C2-N3-C4	8.41	124.10	119.90
85	AA	1720	C	C5'-C4'-C3'	-8.41	102.55	116.00
34	BA	1192	A	O4'-C1'-N9	8.40	114.92	108.20
34	BA	684	G	C4-N9-C1'	8.40	137.42	126.50
35	BB	306	U	O4'-C1'-N1	8.40	114.92	108.20
35	BB	524	C	C6-N1-C1'	-8.40	110.71	120.80
85	AA	2001	C	C6-N1-C1'	-8.40	110.71	120.80
35	BB	855	G	C5-C6-O6	-8.40	123.56	128.60
37	BD	71	G	C8-N9-C1'	8.40	137.93	127.00
38	BE	100	U	C3'-C2'-C1'	-8.40	94.78	101.50
38	BE	108	U	C3'-C2'-C1'	-8.40	94.78	101.50
51	BR	86	LYS	N-CA-CB	-8.40	95.47	110.60
85	AA	163	C	C6-N1-C2	-8.40	116.94	120.30
85	AA	783	C	C6-N1-C2	-8.40	116.94	120.30
34	BA	71	G	C6-N1-C2	-8.40	120.06	125.10
34	BA	298	G	P-O3'-C3'	8.40	129.78	119.70
39	BF	3	A	P-O3'-C3'	-8.40	109.62	119.70
7	A6	39	ARG	NE-CZ-NH2	-8.40	116.10	120.30
35	BB	637	G	C8-N9-C4	-8.40	103.04	106.40
40	BG	4	A	C8-N9-C4	-8.40	102.44	105.80
85	AA	1285	C	P-O3'-C3'	-8.40	109.62	119.70
85	AA	1936	C	O4'-C1'-N1	8.40	114.92	108.20
35	BB	678	U	P-O3'-C3'	-8.40	109.62	119.70
34	BA	1690	U	O4'-C1'-N1	8.40	114.92	108.20
35	BB	775	U	C2-N3-C4	-8.40	121.96	127.00
40	BG	164	U	C5'-C4'-C3'	-8.40	102.56	116.00
47	BN	7	ALA	C-N-CA	8.40	142.69	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1483	A	C5'-C4'-C3'	-8.40	102.57	116.00
40	BG	24	A	C5'-C4'-C3'	-8.40	102.56	116.00
40	BG	76	C	N3-C4-N4	8.40	123.88	118.00
85	AA	744	C	P-O3'-C3'	-8.40	109.62	119.70
85	AA	1865	C	O4'-C1'-N1	8.40	114.92	108.20
40	BG	19	C	C6-N1-C2	-8.39	116.94	120.30
34	BA	1495	A	O4'-C1'-N9	8.39	114.92	108.20
35	BB	1462	G	N1-C6-O6	-8.39	114.86	119.90
36	BC	28	C	N1-C2-O2	-8.39	113.86	118.90
77	Br	120	ARG	NE-CZ-NH1	8.39	124.50	120.30
85	AA	773	G	C6-N1-C2	-8.39	120.06	125.10
85	AA	1655	G	P-O3'-C3'	-8.39	109.63	119.70
34	BA	723	C	C6-N1-C1'	-8.39	110.73	120.80
34	BA	1616	A	O5'-C5'-C4'	8.39	127.64	111.70
41	BH	10	U	P-O5'-C5'	-8.39	107.47	120.90
85	AA	2083	G	C5-C6-N1	8.39	115.70	111.50
85	AA	537	G	P-O3'-C3'	-8.39	109.63	119.70
85	AA	1078	A	O4'-C1'-N9	8.39	114.91	108.20
34	BA	1172	C	C4'-C3'-C2'	8.39	110.99	102.60
34	BA	1792	U	N1-C2-N3	8.39	119.93	114.90
39	BF	33	C	O5'-C5'-C4'	-8.39	95.76	111.70
85	AA	520	A	O4'-C1'-N9	8.39	114.91	108.20
85	AA	1107	A	C8-N9-C4	-8.39	102.44	105.80
34	BA	1565	U	O5'-C5'-C4'	8.39	127.63	111.70
35	BB	802	G	N3-C4-C5	-8.38	124.41	128.60
36	BC	107	C	O4'-C1'-N1	8.39	114.91	108.20
38	BE	91	G	N1-C6-O6	8.38	124.93	119.90
85	AA	1024	G	P-O3'-C3'	-8.39	109.64	119.70
40	BG	20	U	N3-C2-O2	-8.38	116.33	122.20
85	AA	3	U	O4'-C1'-N1	8.38	114.91	108.20
85	AA	586	G	C4-N9-C1'	-8.38	115.60	126.50
34	BA	745	A	P-O3'-C3'	-8.38	109.64	119.70
34	BA	1175	G	O4'-C1'-N9	8.38	114.91	108.20
35	BB	1495	U	C2-N1-C1'	-8.38	107.64	117.70
38	BE	25	U	N1-C2-N3	8.38	119.93	114.90
85	AA	622	G	C4-N9-C1'	-8.38	115.60	126.50
34	BA	76	U	C2-N3-C4	-8.38	121.97	127.00
35	BB	808	U	C6-N1-C1'	8.38	132.93	121.20
41	BH	134	U	P-O3'-C3'	8.38	129.76	119.70
35	BB	1222	A	N1-C6-N6	8.38	123.63	118.60
39	BF	69	A	P-O3'-C3'	-8.38	109.64	119.70
34	BA	575	U	C6-N1-C1'	-8.38	109.47	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	282	C	P-O5'-C5'	-8.38	107.50	120.90
85	AA	1647	G	O4'-C1'-N9	8.38	114.90	108.20
85	AA	1955	U	C1'-O4'-C4'	-8.38	103.20	109.90
34	BA	251	U	C5'-C4'-O4'	8.38	119.15	109.10
85	AA	790	A	C8-N9-C4	-8.38	102.45	105.80
85	AA	1732	G	C5-C6-O6	-8.38	123.57	128.60
34	BA	152	C	C6-N1-C2	-8.37	116.95	120.30
34	BA	732	A	N1-C6-N6	8.38	123.62	118.60
34	BA	1202	G	O4'-C1'-N9	8.38	114.90	108.20
85	AA	604	C	P-O5'-C5'	8.38	134.30	120.90
85	AA	2189	U	C5'-C4'-C3'	-8.38	102.60	116.00
34	BA	1650	G	N1-C6-O6	-8.37	114.88	119.90
35	BB	832	C	O4'-C1'-N1	8.37	114.90	108.20
36	BC	16	A	C5-C6-N1	8.38	121.89	117.70
41	BH	105	U	N3-C2-O2	-8.37	116.34	122.20
34	BA	1195	G	C8-N9-C1'	8.37	137.88	127.00
85	AA	2066	C	C3'-C2'-C1'	-8.37	94.80	101.50
35	BB	872	A	P-O5'-C5'	-8.37	107.51	120.90
37	BD	48	G	P-O5'-C5'	-8.37	107.51	120.90
38	BE	148	C	O5'-C5'-C4'	-8.37	95.79	111.70
85	AA	385	A	C1'-O4'-C4'	-8.37	103.20	109.90
85	AA	944	C	P-O3'-C3'	-8.37	109.66	119.70
35	BB	1511	U	P-O5'-C5'	-8.37	107.51	120.90
85	AA	488	G	P-O5'-C5'	8.37	134.29	120.90
85	AA	1584	U	O4'-C1'-N1	8.37	114.90	108.20
34	BA	228	A	C6-N1-C2	-8.37	113.58	118.60
85	AA	584	G	C5-C6-O6	-8.37	123.58	128.60
34	BA	384	U	C4-C5-C6	-8.37	114.68	119.70
35	BB	1230	A	C5-C6-N6	-8.37	117.01	123.70
38	BE	8	G	C6-N1-C2	-8.37	120.08	125.10
85	AA	1586	C	O4'-C1'-N1	8.37	114.89	108.20
85	AA	1991	C	N3-C4-C5	-8.37	118.55	121.90
34	BA	35	U	C2-N3-C4	-8.36	121.98	127.00
34	BA	486	G	C6-N1-C2	-8.36	120.08	125.10
34	BA	1071	G	C8-N9-C1'	8.36	137.87	127.00
35	BB	266	C	O4'-C1'-N1	8.36	114.89	108.20
35	BB	497	C	O4'-C1'-N1	8.36	114.89	108.20
35	BB	1115	G	C8-N9-C1'	8.37	137.87	127.00
54	BU	86	ARG	NE-CZ-NH1	8.37	124.48	120.30
35	BB	127	U	P-O5'-C5'	8.36	134.28	120.90
34	BA	596	G	C8-N9-C1'	-8.36	116.13	127.00
37	BD	17	G	C5-C6-O6	-8.36	123.58	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	BN	7	ALA	CB-CA-C	8.36	122.64	110.10
85	AA	196	U	P-O3'-C3'	8.36	129.73	119.70
34	BA	1661	U	P-O5'-C5'	-8.36	107.52	120.90
34	BA	937	G	N3-C2-N2	8.36	125.75	119.90
35	BB	840	C	C4'-C3'-C2'	8.36	110.96	102.60
35	BB	1221	G	N1-C6-O6	-8.36	114.88	119.90
85	AA	21	U	C2-N3-C4	-8.36	121.98	127.00
85	AA	818	C	N3-C4-C5	-8.36	118.56	121.90
85	AA	414	C	C3'-C2'-C1'	-8.36	94.81	101.50
85	AA	552	C	C1'-O4'-C4'	-8.36	103.21	109.90
85	AA	686	U	C4'-C3'-C2'	-8.36	94.24	102.60
34	BA	482	C	O3'-P-O5'	8.36	119.87	104.00
34	BA	727	G	N1-C6-O6	-8.36	114.89	119.90
35	BB	51	U	C5'-C4'-O4'	8.36	119.13	109.10
35	BB	145	G	C8-N9-C1'	8.36	137.86	127.00
35	BB	366	G	C1'-O4'-C4'	-8.36	103.22	109.90
41	BH	109	G	C4-N9-C1'	-8.36	115.64	126.50
85	AA	1542	A	N1-C6-N6	-8.36	113.59	118.60
85	AA	55	A	P-O5'-C5'	-8.36	107.53	120.90
85	AA	62	A	P-O5'-C5'	-8.36	107.53	120.90
85	AA	1865	C	C6-N1-C2	-8.36	116.96	120.30
85	AA	2128	G	C8-N9-C1'	8.36	137.86	127.00
85	AA	2238	C	O4'-C1'-C2'	8.36	115.12	107.60
34	BA	1507	C	C6-N1-C2	-8.35	116.96	120.30
34	BA	147	U	C4-C5-C6	-8.35	114.69	119.70
34	BA	154	A	N1-C6-N6	8.35	123.61	118.60
85	AA	684	G	C5-C6-O6	-8.35	123.59	128.60
85	AA	820	G	C5-N7-C8	8.35	108.48	104.30
34	BA	488	C	C6-N1-C1'	-8.35	110.78	120.80
34	BA	888	G	C4-N9-C1'	-8.35	115.64	126.50
34	BA	1792	U	O4'-C1'-N1	8.35	114.88	108.20
37	BD	68	C	C3'-C2'-C1'	-8.35	94.82	101.50
80	Bu	30	TYR	CB-CG-CD1	8.35	126.01	121.00
85	AA	987	C	N3-C4-C5	-8.35	118.56	121.90
85	AA	1281	G	C5-C6-O6	-8.35	123.59	128.60
85	AA	1603	G	C5-C6-O6	-8.35	123.59	128.60
34	BA	732	A	C5-C6-N6	-8.35	117.02	123.70
38	BE	83	U	C4'-C3'-C2'	-8.35	94.25	102.60
35	BB	1498	G	C8-N9-C4	8.35	109.74	106.40
38	BE	67	A	C5'-C4'-C3'	-8.35	102.64	116.00
85	AA	767	A	O5'-C5'-C4'	8.35	127.56	111.70
85	AA	1427	A	O4'-C1'-N9	8.35	114.88	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2249	U	P-O3'-C3'	8.35	129.72	119.70
22	AO	60	ARG	NE-CZ-NH1	8.35	124.47	120.30
34	BA	754	G	O4'-C1'-N9	8.35	114.88	108.20
41	BH	3	U	C4'-C3'-C2'	8.35	110.95	102.60
41	BH	41	A	C4-N9-C1'	-8.35	111.28	126.30
85	AA	607	U	O4'-C1'-N1	8.35	114.88	108.20
34	BA	500	C	C2-N1-C1'	-8.34	109.62	118.80
35	BB	1534	U	C2-N1-C1'	-8.34	107.69	117.70
38	BE	97	G	C3'-C2'-C1'	-8.34	94.83	101.50
39	BF	22	U	C6-N1-C2	-8.34	115.99	121.00
40	BG	80	G	N1-C6-O6	-8.34	114.89	119.90
41	BH	51	C	O4'-C1'-N1	8.34	114.87	108.20
85	AA	1155	A	C5-C6-N6	-8.34	117.03	123.70
85	AA	2076	C	C5'-C4'-C3'	8.34	129.35	116.00
37	BD	82	G	N1-C6-O6	-8.34	114.90	119.90
26	AS	119	ARG	N-CA-CB	-8.34	95.59	110.60
34	BA	870	C	OP1-P-OP2	8.34	132.11	119.60
85	AA	1536	C	C6-N1-C2	-8.34	116.96	120.30
85	AA	2164	G	P-O3'-C3'	-8.34	109.69	119.70
35	BB	716	G	C8-N9-C4	-8.34	103.06	106.40
35	BB	1004	A	C1'-O4'-C4'	-8.34	103.23	109.90
85	AA	817	G	O3'-P-O5'	-8.34	88.15	104.00
85	AA	1476	C	C4'-C3'-C2'	8.34	110.94	102.60
34	BA	1069	U	O4'-C1'-N1	8.34	114.87	108.20
34	BA	1079	C	N3-C4-C5	-8.34	118.56	121.90
35	BB	706	G	C5-C6-N1	8.34	115.67	111.50
35	BB	788	U	C5'-C4'-C3'	8.34	129.34	116.00
41	BH	80	C	O4'-C1'-N1	8.34	114.87	108.20
85	AA	1492	U	N3-C2-O2	-8.34	116.36	122.20
34	BA	1406	U	P-O3'-C3'	-8.34	109.70	119.70
34	BA	1502	G	P-O3'-C3'	-8.34	109.70	119.70
35	BB	557	C	C6-N1-C1'	8.34	130.80	120.80
85	AA	981	A	O4'-C1'-N9	8.34	114.87	108.20
85	AA	1454	U	O4'-C1'-N1	8.34	114.87	108.20
34	BA	134	U	C6-N1-C1'	8.33	132.87	121.20
34	BA	1730	A	O4'-C1'-N9	8.33	114.87	108.20
35	BB	1039	A	P-O3'-C3'	8.33	129.70	119.70
63	Bd	61	GLU	CG-CD-OE1	-8.33	101.63	118.30
68	Bi	20	ARG	NE-CZ-NH1	8.33	124.47	120.30
5	A4	71	ARG	NE-CZ-NH2	-8.33	116.14	120.30
35	BB	1036	G	C4-N9-C1'	-8.33	115.67	126.50
34	BA	53	G	C5-C6-N1	8.33	115.66	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	315	U	O3'-P-O5'	-8.33	88.17	104.00
34	BA	681	G	O4'-C1'-N9	-8.33	101.54	108.20
34	BA	1652	G	N9-C1'-C2'	-8.33	102.84	112.00
35	BB	391	G	C4-N9-C1'	-8.33	115.67	126.50
85	AA	448	G	P-O3'-C3'	-8.33	109.70	119.70
85	AA	1014	U	O4'-C1'-N1	8.33	114.86	108.20
34	BA	174	A	C5'-C4'-C3'	-8.33	102.68	116.00
34	BA	773	A	P-O3'-C3'	-8.33	109.71	119.70
38	BE	37	C	C2-N1-C1'	8.33	127.96	118.80
34	BA	1846	G	P-O5'-C5'	-8.33	107.58	120.90
40	BG	5	G	O4'-C1'-N9	8.33	114.86	108.20
60	Ba	115	PHE	CB-CG-CD2	-8.33	114.97	120.80
85	AA	464	A	C5'-C4'-C3'	-8.33	102.68	116.00
85	AA	485	A	P-O3'-C3'	-8.33	109.71	119.70
85	AA	2186	U	C2-N3-C4	-8.33	122.00	127.00
34	BA	1054	U	C2-N3-C4	-8.32	122.01	127.00
35	BB	796	C	P-O3'-C3'	-8.32	109.71	119.70
35	BB	973	G	C8-N9-C4	-8.32	103.07	106.40
85	AA	532	G	C8-N9-C1'	8.32	137.82	127.00
85	AA	1521	U	C5'-C4'-C3'	-8.32	102.68	116.00
38	BE	42	C	O4'-C1'-N1	8.32	114.86	108.20
41	BH	79	A	N1-C6-N6	8.32	123.59	118.60
53	BT	181	ARG	NE-CZ-NH2	-8.32	116.14	120.30
35	BB	1328	C	C5'-C4'-C3'	8.32	129.31	116.00
35	BB	999	G	P-O3'-C3'	-8.32	109.72	119.70
40	BG	167	C	C6-N1-C2	-8.32	116.97	120.30
67	Bh	38	TRP	CB-CA-C	-8.32	93.76	110.40
85	AA	415	G	C4-N9-C1'	-8.32	115.68	126.50
85	AA	752	C	C6-N1-C2	-8.32	116.97	120.30
85	AA	1101	C	O4'-C1'-N1	8.32	114.86	108.20
34	BA	797	A	O4'-C1'-N9	8.32	114.85	108.20
35	BB	472	C	P-O3'-C3'	-8.32	109.72	119.70
35	BB	1132	A	O4'-C1'-N9	8.32	114.85	108.20
38	BE	123	A	C8-N9-C4	8.32	109.13	105.80
38	BE	184	G	C8-N9-C1'	8.32	137.81	127.00
85	AA	444	U	P-O5'-C5'	8.32	134.21	120.90
85	AA	568	C	C6-N1-C2	-8.32	116.97	120.30
34	BA	330	A	C8-N9-C4	8.31	109.12	105.80
36	BC	27	U	N3-C2-O2	8.31	128.02	122.20
38	BE	198	A	N9-C4-C5	-8.31	102.47	105.80
85	AA	1140	G	O4'-C1'-N9	8.31	114.85	108.20
85	AA	927	A	C5'-C4'-C3'	-8.31	102.70	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1545	U	P-O3'-C3'	8.31	129.68	119.70
4	A3	73	ARG	NE-CZ-NH1	8.31	124.46	120.30
35	BB	504	C	N3-C4-C5	-8.31	118.58	121.90
35	BB	690	C	P-O3'-C3'	-8.31	109.72	119.70
35	BB	979	G	O4'-C1'-N9	8.31	114.85	108.20
85	AA	788	G	P-O5'-C5'	8.31	134.20	120.90
85	AA	1159	C	C5'-C4'-O4'	-8.31	99.13	109.10
35	BB	717	A	C5'-C4'-C3'	-8.31	102.70	116.00
35	BB	736	G	C5-C6-O6	-8.31	123.61	128.60
67	Bh	36	ARG	NE-CZ-NH1	8.31	124.45	120.30
69	Bj	71	HIS	CA-CB-CG	8.31	127.73	113.60
85	AA	927	A	C5-C6-N1	8.31	121.86	117.70
85	AA	1955	U	C4'-C3'-C2'	-8.31	94.29	102.60
34	BA	526	C	C1'-O4'-C4'	-8.31	103.25	109.90
34	BA	1776	G	C8-N9-C1'	8.31	137.80	127.00
85	AA	366	A	P-O3'-C3'	-8.31	109.73	119.70
34	BA	280	A	O5'-C5'-C4'	-8.30	95.92	111.70
34	BA	623	U	P-O5'-C5'	-8.30	107.61	120.90
34	BA	774	A	N1-C6-N6	-8.30	113.62	118.60
34	BA	1197	U	C2'-C3'-O3'	8.30	127.77	109.50
34	BA	1478	G	C8-N9-C4	-8.30	103.08	106.40
35	BB	1054	G	N3-C2-N2	8.30	125.71	119.90
35	BB	1228	A	O4'-C1'-N9	8.30	114.84	108.20
85	AA	1479	U	P-O3'-C3'	-8.31	109.73	119.70
35	BB	1474	A	C5'-C4'-C3'	8.30	129.29	116.00
37	BD	57	C	C5'-C4'-C3'	-8.30	102.72	116.00
37	BD	108	G	C4-N9-C1'	-8.30	115.71	126.50
38	BE	141	A	C1'-O4'-C4'	-8.30	103.26	109.90
85	AA	1112	G	C5-C6-O6	-8.30	123.62	128.60
85	AA	1668	G	C8-N9-C1'	8.30	137.80	127.00
34	BA	166	G	O4'-C1'-N9	8.30	114.84	108.20
38	BE	57	U	P-O5'-C5'	8.30	134.18	120.90
85	AA	781	G	P-O3'-C3'	-8.30	109.74	119.70
85	AA	1640	G	C5-C6-O6	-8.30	123.62	128.60
12	AD	54	ARG	NE-CZ-NH1	8.30	124.45	120.30
34	BA	777	C	O4'-C1'-N1	8.30	114.84	108.20
35	BB	902	C	P-O5'-C5'	-8.30	107.62	120.90
35	BB	1194	A	N1-C6-N6	8.30	123.58	118.60
34	BA	1626	U	C1'-O4'-C4'	8.30	116.54	109.90
44	BK	21	ARG	NE-CZ-NH1	8.30	124.45	120.30
85	AA	675	A	O4'-C1'-N9	8.30	114.84	108.20
85	AA	1656	C	O4'-C1'-N1	8.30	114.84	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	100	A	C5'-C4'-C3'	8.29	129.27	116.00
34	BA	1442	A	C5-C6-N1	8.30	121.85	117.70
38	BE	174	U	C2-N3-C4	-8.30	122.02	127.00
34	BA	113	G	O4'-C1'-N9	8.29	114.83	108.20
35	BB	1124	G	N1-C6-O6	8.29	124.88	119.90
85	AA	189	G	C4-N9-C1'	-8.29	115.72	126.50
85	AA	1013	C	O4'-C1'-N1	8.29	114.83	108.20
34	BA	294	C	C2-N3-C4	-8.29	115.75	119.90
35	BB	762	C	N3-C4-N4	8.29	123.81	118.00
35	BB	1108	G	C5-C6-N1	8.29	115.65	111.50
38	BE	84	U	O4'-C1'-N1	8.29	114.83	108.20
35	BB	365	U	C4'-C3'-C2'	8.29	110.89	102.60
38	BE	88	G	P-O3'-C3'	8.29	129.65	119.70
85	AA	960	G	N7-C8-N9	8.29	117.25	113.10
37	BD	3	G	C4-N9-C1'	-8.29	115.72	126.50
85	AA	636	G	C5'-C4'-C3'	-8.29	102.74	116.00
86	AB	6	G	O4'-C1'-N9	8.29	114.83	108.20
34	BA	201	A	O4'-C1'-N9	8.29	114.83	108.20
37	BD	71	G	C4-N9-C1'	-8.29	115.72	126.50
38	BE	15	A	O3'-P-O5'	8.29	119.75	104.00
85	AA	474	C	N3-C2-O2	-8.29	116.10	121.90
85	AA	1253	G	C3'-C2'-C1'	-8.29	94.87	101.50
85	AA	1663	U	C2-N3-C4	-8.29	122.03	127.00
85	AA	1828	C	C5'-C4'-C3'	-8.29	102.74	116.00
85	AA	747	U	O4'-C1'-N1	8.29	114.83	108.20
85	AA	936	C	P-O3'-C3'	8.29	129.65	119.70
85	AA	1420	U	P-O3'-C3'	8.29	129.65	119.70
85	AA	1779	C	O4'-C1'-N1	8.29	114.83	108.20
34	BA	1489	U	P-O3'-C3'	-8.29	109.76	119.70
34	BA	1502	G	C5-C6-N1	8.29	115.64	111.50
2	A1	88	GLU	N-CA-CB	8.28	125.51	110.60
34	BA	502	U	C5'-C4'-C3'	-8.28	102.75	116.00
35	BB	852	G	C4'-C3'-C2'	-8.28	94.32	102.60
37	BD	108	G	C5'-C4'-C3'	-8.29	102.74	116.00
77	Br	250	TRP	CB-CG-CD2	-8.28	115.83	126.60
85	AA	370	A	O4'-C4'-C3'	-8.28	95.72	104.00
85	AA	1207	C	C5'-C4'-C3'	-8.29	102.74	116.00
34	BA	393	G	O4'-C1'-N9	8.28	114.83	108.20
34	BA	614	A	P-O3'-C3'	-8.28	109.76	119.70
34	BA	348	U	C1'-O4'-C4'	-8.28	103.28	109.90
34	BA	642	U	N3-C4-O4	-8.28	113.60	119.40
40	BG	176	G	C4-N9-C1'	-8.28	115.73	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	744	G	C8-N9-C4	-8.28	103.09	106.40
35	BB	379	U	C3'-C2'-C1'	-8.28	94.88	101.50
35	BB	1514	G	C4-N9-C1'	-8.28	115.73	126.50
41	BH	96	G	N1-C6-O6	8.28	124.87	119.90
85	AA	83	U	C6-N1-C2	-8.28	116.03	121.00
85	AA	207	G	C8-N9-C1'	-8.28	116.23	127.00
85	AA	407	G	P-O5'-C5'	8.28	134.15	120.90
85	AA	1094	G	O4'-C1'-N9	8.28	114.83	108.20
34	BA	1084	A	C5-C6-N6	8.28	130.32	123.70
40	BG	95	U	N3-C2-O2	-8.28	116.40	122.20
85	AA	921	C	C6-N1-C1'	-8.28	110.86	120.80
38	BE	129	G	P-O5'-C5'	8.28	134.14	120.90
85	AA	771	A	C2-N3-C4	8.28	114.74	110.60
85	AA	820	G	C1'-O4'-C4'	-8.28	103.28	109.90
34	BA	204	U	P-O5'-C5'	-8.28	107.66	120.90
34	BA	587	U	C5'-C4'-O4'	8.28	119.03	109.10
34	BA	1494	G	C5'-C4'-C3'	-8.28	102.76	116.00
35	BB	839	G	O3'-P-O5'	8.28	119.72	104.00
35	BB	1529	G	N1-C6-O6	8.28	124.86	119.90
38	BE	173	G	N1-C6-O6	8.28	124.87	119.90
74	Bo	8	MET	CG-SD-CE	-8.28	86.96	100.20
2	A1	100	TYR	CB-CG-CD1	-8.27	116.04	121.00
16	AH	124	ASP	N-CA-CB	8.27	125.49	110.60
34	BA	322	U	C2-N1-C1'	-8.27	107.77	117.70
34	BA	882	G	C8-N9-C4	-8.27	103.09	106.40
35	BB	1478	G	P-O3'-C3'	-8.27	109.77	119.70
38	BE	121	G	N1-C6-O6	8.27	124.86	119.90
38	BE	175	U	C2-N1-C1'	-8.27	107.77	117.70
34	BA	189	G	C6-N1-C2	-8.27	120.14	125.10
39	BF	23	G	O4'-C1'-N9	8.27	114.82	108.20
41	BH	44	A	N1-C6-N6	8.27	123.56	118.60
34	BA	501	U	C5'-C4'-C3'	8.27	129.23	116.00
34	BA	782	C	C5-C4-N4	-8.27	114.41	120.20
34	BA	840	U	C5'-C4'-C3'	-8.27	102.77	116.00
34	BA	1482	A	N1-C6-N6	-8.27	113.64	118.60
37	BD	85	C	C2-N3-C4	-8.27	115.76	119.90
41	BH	118	U	C4'-C3'-C2'	8.27	110.87	102.60
85	AA	1125	G	C8-N9-C4	8.27	109.71	106.40
86	AB	8	U	C1'-O4'-C4'	-8.27	103.28	109.90
35	BB	388	C	C5'-C4'-C3'	-8.27	102.77	116.00
35	BB	1506	C	P-O3'-C3'	8.27	129.62	119.70
85	AA	1199	C	P-O5'-C5'	-8.27	107.67	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1586	C	C3'-C2'-C1'	-8.27	94.89	101.50
35	BB	66	G	P-O3'-C3'	-8.27	109.78	119.70
85	AA	659	A	C1'-O4'-C4'	-8.27	103.29	109.90
35	BB	863	U	O4'-C1'-N1	8.27	114.81	108.20
35	BB	1102	U	C2-N3-C4	-8.27	122.04	127.00
36	BC	47	C	C6-N1-C2	-8.27	116.99	120.30
85	AA	201	U	C6-N1-C2	-8.27	116.04	121.00
85	AA	578	U	O4'-C1'-N1	8.27	114.81	108.20
85	AA	909	C	C2-N1-C1'	8.27	127.89	118.80
85	AA	2197	A	C8-N9-C4	-8.27	102.49	105.80
85	AA	86	G	C5'-C4'-O4'	-8.26	99.18	109.10
35	BB	871	C	C6-N1-C1'	8.26	130.71	120.80
85	AA	147	G	C5-C6-O6	-8.26	123.64	128.60
85	AA	557	G	O4'-C1'-N9	8.26	114.81	108.20
85	AA	818	C	O4'-C1'-N1	8.26	114.81	108.20
34	BA	624	G	C8-N9-C4	-8.26	103.10	106.40
41	BH	128	G	C6-N1-C2	-8.26	120.14	125.10
85	AA	2014	G	P-O5'-C5'	-8.26	107.68	120.90
38	BE	1	U	C6-N1-C2	-8.26	116.05	121.00
61	Bb	19	TYR	CB-CG-CD2	-8.26	116.05	121.00
61	Bb	42	ARG	NE-CZ-NH2	-8.26	116.17	120.30
85	AA	771	A	P-O5'-C5'	8.26	134.12	120.90
85	AA	1010	U	C2-N1-C1'	-8.26	107.79	117.70
34	BA	301	U	C2-N1-C1'	-8.26	107.79	117.70
34	BA	1027	C	C6-N1-C2	-8.26	117.00	120.30
34	BA	1260	G	C1'-O4'-C4'	-8.26	103.29	109.90
38	BE	191	U	C5'-C4'-C3'	-8.26	102.79	116.00
85	AA	493	A	P-O5'-C5'	-8.26	107.69	120.90
86	AB	7	A	C5-C6-N6	-8.26	117.09	123.70
34	BA	631	G	N1-C6-O6	8.26	124.85	119.90
34	BA	1354	G	C1'-O4'-C4'	-8.26	103.30	109.90
34	BA	1562	G	N9-C4-C5	-8.26	102.10	105.40
35	BB	1212	C	N3-C4-C5	-8.26	118.60	121.90
37	BD	25	G	C8-N9-C1'	8.26	137.73	127.00
35	BB	534	C	O4'-C1'-N1	8.25	114.80	108.20
41	BH	41	A	O4'-C1'-C2'	8.25	115.03	107.60
34	BA	201	A	N9-C4-C5	8.25	109.10	105.80
34	BA	882	G	C5-C6-N1	8.25	115.63	111.50
41	BH	100	A	OP1-P-OP2	-8.25	107.22	119.60
85	AA	509	C	C5'-C4'-C3'	-8.25	102.80	116.00
85	AA	540	A	C1'-O4'-C4'	-8.25	103.30	109.90
36	BC	7	U	C4'-C3'-C2'	-8.25	94.35	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	121	G	P-O3'-C3'	-8.25	109.80	119.70
33	AZ	89	MET	CG-SD-CE	-8.25	87.00	100.20
34	BA	895	U	O4'-C4'-C3'	-8.25	95.75	104.00
85	AA	788	G	C5'-C4'-C3'	8.25	129.20	116.00
85	AA	863	C	P-O5'-C5'	-8.25	107.70	120.90
85	AA	2082	C	C6-N1-C1'	-8.25	110.90	120.80
34	BA	118	C	N3-C4-N4	-8.25	112.23	118.00
34	BA	123	C	O4'-C1'-N1	8.25	114.80	108.20
34	BA	1483	U	P-O3'-C3'	-8.25	109.80	119.70
35	BB	64	U	C2-N3-C4	-8.25	122.05	127.00
34	BA	952	G	C8-N9-C1'	8.25	137.72	127.00
34	BA	1443	U	C4'-C3'-C2'	-8.25	94.35	102.60
82	Bw	33	PHE	CB-CG-CD1	8.25	126.57	120.80
34	BA	206	C	C6-N1-C2	-8.24	117.00	120.30
34	BA	239	C	C1'-O4'-C4'	-8.24	103.31	109.90
35	BB	506	G	N1-C6-O6	-8.24	114.95	119.90
40	BG	44	G	C3'-C2'-C1'	-8.24	94.90	101.50
85	AA	426	C	C1'-O4'-C4'	-8.24	103.31	109.90
35	BB	799	A	C5-C6-N6	-8.24	117.11	123.70
85	AA	857	G	P-O3'-C3'	-8.24	109.81	119.70
85	AA	1973	G	C5'-C4'-C3'	-8.24	102.81	116.00
34	BA	861	C	O4'-C1'-N1	8.24	114.79	108.20
34	BA	1413	G	O5'-P-OP1	8.24	120.59	110.70
35	BB	363	A	C5'-C4'-O4'	8.24	118.99	109.10
35	BB	866	A	C5'-C4'-C3'	-8.24	102.82	116.00
35	BB	1024	G	N7-C8-N9	8.24	117.22	113.10
36	BC	82	C	C5-C4-N4	-8.24	114.43	120.20
37	BD	54	A	N1-C6-N6	8.24	123.54	118.60
85	AA	982	G	C5'-C4'-C3'	-8.24	102.82	116.00
19	AK	115	TYR	CB-CG-CD2	-8.24	116.06	121.00
34	BA	526	C	N1-C1'-C2'	-8.24	102.94	112.00
34	BA	1832	A	P-O3'-C3'	8.24	129.59	119.70
34	BA	513	U	N1-C2-N3	8.24	119.84	114.90
34	BA	1089	U	P-O3'-C3'	-8.24	109.82	119.70
34	BA	1149	C	N3-C4-N4	-8.24	112.23	118.00
35	BB	2	C	C4-C5-C6	-8.24	113.28	117.40
47	BN	31	PHE	CB-CG-CD1	8.24	126.57	120.80
85	AA	899	A	O4'-C4'-C3'	-8.24	95.76	104.00
85	AA	1919	G	N1-C6-O6	8.24	124.84	119.90
34	BA	1495	A	C5'-C4'-O4'	8.23	118.98	109.10
34	BA	336	A	N1-C6-N6	8.23	123.54	118.60
34	BA	588	C	P-O5'-C5'	8.23	134.07	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	84	U	C1'-O4'-C4'	-8.23	103.31	109.90
34	BA	398	G	N3-C2-N2	-8.23	114.14	119.90
34	BA	935	A	P-O3'-C3'	-8.23	109.82	119.70
34	BA	1615	A	O3'-P-O5'	8.23	119.64	104.00
36	BC	23	G	C1'-O4'-C4'	-8.23	103.31	109.90
85	AA	1098	C	O5'-C5'-C4'	-8.23	96.06	111.70
35	BB	1260	A	N1-C6-N6	-8.23	113.66	118.60
85	AA	270	A	C8-N9-C1'	8.23	142.51	127.70
85	AA	1500	C	O4'-C1'-N1	8.23	114.78	108.20
34	BA	833	U	O4'-C1'-N1	8.23	114.78	108.20
34	BA	1100	A	C8-N9-C4	8.23	109.09	105.80
35	BB	52	G	N9-C1'-C2'	-8.23	102.95	112.00
35	BB	1016	C	P-O3'-C3'	-8.23	109.83	119.70
35	BB	1226	G	P-O3'-C3'	8.23	129.57	119.70
85	AA	28	A	N1-C6-N6	-8.23	113.66	118.60
34	BA	400	A	C5'-C4'-C3'	-8.23	102.84	116.00
34	BA	557	U	O3'-P-O5'	8.23	119.63	104.00
34	BA	617	G	N1-C6-O6	-8.23	114.96	119.90
34	BA	690	G	C5-C6-O6	-8.23	123.66	128.60
34	BA	1085	G	P-O3'-C3'	8.23	129.57	119.70
35	BB	1389	C	O4'-C1'-N1	8.23	114.78	108.20
85	AA	363	A	C1'-O4'-C4'	-8.23	103.32	109.90
85	AA	805	A	P-O3'-C3'	8.23	129.57	119.70
85	AA	1589	G	C4-N9-C1'	-8.23	115.81	126.50
35	BB	736	G	O3'-P-O5'	-8.22	88.37	104.00
85	AA	495	G	C3'-C2'-C1'	-8.22	94.92	101.50
34	BA	311	C	P-O3'-C3'	8.22	129.57	119.70
34	BA	1411	C	O4'-C1'-N1	8.22	114.78	108.20
34	BA	1451	A	P-O5'-C5'	-8.22	107.74	120.90
35	BB	1108	G	N1-C6-O6	-8.22	114.97	119.90
44	BK	139	ARG	NE-CZ-NH1	8.22	124.41	120.30
85	AA	487	G	C1'-O4'-C4'	-8.22	103.32	109.90
85	AA	765	U	C5'-C4'-O4'	8.22	118.97	109.10
34	BA	1077	G	C5-C6-O6	-8.22	123.67	128.60
4	A3	203	ARG	NE-CZ-NH1	8.22	124.41	120.30
35	BB	474	G	C8-N9-C4	8.22	109.69	106.40
35	BB	1071	G	C8-N9-C1'	8.22	137.69	127.00
37	BD	95	G	C8-N9-C1'	8.22	137.69	127.00
40	BG	95	U	N1-C2-O2	8.22	128.56	122.80
85	AA	198	U	O4'-C1'-N1	8.22	114.78	108.20
85	AA	210	G	P-O5'-C5'	8.22	134.05	120.90
85	AA	417	U	C4'-C3'-C2'	-8.22	94.38	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	916	A	P-O5'-C5'	-8.22	107.75	120.90
85	AA	1363	U	C6-N1-C2	-8.22	116.07	121.00
85	AA	1695	G	C8-N9-C4	8.22	109.69	106.40
86	AB	8	U	C5'-C4'-C3'	8.22	129.15	116.00
35	BB	870	C	P-O3'-C3'	-8.22	109.84	119.70
35	BB	964	G	C5-C6-O6	-8.22	123.67	128.60
85	AA	155	U	O4'-C1'-N1	8.22	114.77	108.20
85	AA	352	G	C8-N9-C4	8.22	109.69	106.40
85	AA	1690	A	C8-N9-C4	8.22	109.09	105.80
34	BA	210	G	C4'-C3'-C2'	8.21	110.81	102.60
37	BD	78	C	C2-N1-C1'	-8.22	109.76	118.80
85	AA	8	U	C4'-C3'-C2'	8.22	110.82	102.60
34	BA	449	G	O4'-C1'-N9	8.21	114.77	108.20
85	AA	1439	A	P-O3'-C3'	-8.21	109.84	119.70
34	BA	61	G	C4-N9-C1'	-8.21	115.82	126.50
34	BA	258	C	C1'-O4'-C4'	-8.21	103.33	109.90
34	BA	295	G	C6-N1-C2	-8.21	120.17	125.10
34	BA	793	A	N1-C6-N6	8.21	123.53	118.60
35	BB	1178	A	C5'-C4'-O4'	8.21	118.95	109.10
85	AA	1720	C	C2'-C3'-O3'	8.21	127.57	109.50
34	BA	1321	A	C1'-O4'-C4'	-8.21	103.33	109.90
36	BC	165	U	P-O3'-C3'	-8.21	109.85	119.70
39	BF	47	C	O4'-C1'-N1	8.21	114.77	108.20
52	BS	89	TYR	CA-CB-CG	8.21	129.00	113.40
85	AA	194	U	C2-N1-C1'	-8.21	107.84	117.70
85	AA	494	G	C1'-O4'-C4'	-8.21	103.33	109.90
85	AA	1458	G	N9-C4-C5	8.21	108.69	105.40
85	AA	674	U	C6-N1-C1'	8.21	132.69	121.20
85	AA	722	G	C8-N9-C1'	8.21	137.67	127.00
34	BA	300	C	N3-C2-O2	-8.21	116.15	121.90
34	BA	763	U	C5-C4-O4	-8.21	120.97	125.90
35	BB	983	C	O4'-C1'-N1	8.21	114.77	108.20
85	AA	1116	G	C5-C6-N1	8.21	115.60	111.50
34	BA	763	U	P-O3'-C3'	8.21	129.54	119.70
34	BA	875	G	C6-N1-C2	-8.21	120.18	125.10
34	BA	1675	C	C2-N1-C1'	-8.21	109.77	118.80
34	BA	1688	G	P-O3'-C3'	8.21	129.55	119.70
36	BC	139	A	C5'-C4'-C3'	8.21	129.13	116.00
41	BH	74	G	N1-C2-N3	-8.21	118.98	123.90
43	BJ	212	ARG	NE-CZ-NH1	8.21	124.40	120.30
85	AA	373	G	N9-C1'-C2'	-8.21	102.97	112.00
85	AA	394	C	O4'-C1'-N1	8.20	114.76	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	600	C	C5'-C4'-C3'	8.20	129.13	116.00
85	AA	1450	U	O4'-C1'-N1	8.21	114.76	108.20
34	BA	280	A	N1-C6-N6	8.20	123.52	118.60
34	BA	680	C	O5'-P-OP2	8.20	120.54	110.70
34	BA	1200	U	C6-N1-C1'	-8.20	109.72	121.20
36	BC	30	U	P-O3'-C3'	8.20	129.54	119.70
40	BG	164	U	C4-C5-C6	-8.20	114.78	119.70
34	BA	1205	A	C4-C5-C6	-8.20	112.90	117.00
76	Bq	25	TYR	CB-CG-CD2	-8.20	116.08	121.00
85	AA	813	G	P-O3'-C3'	-8.20	109.86	119.70
85	AA	1607	A	C8-N9-C4	-8.20	102.52	105.80
34	BA	239	C	C3'-C2'-C1'	-8.20	94.94	101.50
34	BA	944	G	N1-C6-O6	8.20	124.82	119.90
37	BD	29	C	C2-N1-C1'	-8.20	109.78	118.80
41	BH	129	G	P-O3'-C3'	-8.20	109.86	119.70
85	AA	390	U	C1'-O4'-C4'	-8.20	103.34	109.90
34	BA	3	G	N1-C6-O6	8.20	124.82	119.90
34	BA	144	C	O4'-C1'-N1	8.20	114.76	108.20
34	BA	296	G	C5'-C4'-O4'	-8.20	99.26	109.10
34	BA	422	C	O4'-C1'-N1	8.20	114.76	108.20
39	BF	22	U	P-O5'-C5'	8.20	134.02	120.90
85	AA	944	C	N3-C4-N4	8.20	123.74	118.00
85	AA	2017	U	N1-C1'-C2'	-8.20	102.98	112.00
4	A3	196	ARG	NE-CZ-NH1	8.20	124.40	120.30
34	BA	1736	A	N7-C8-N9	8.20	117.90	113.80
38	BE	181	U	P-O3'-C3'	8.20	129.54	119.70
85	AA	651	G	C5-C6-O6	-8.20	123.68	128.60
34	BA	130	U	O4'-C1'-N1	8.19	114.75	108.20
34	BA	732	A	C8-N9-C4	8.19	109.08	105.80
34	BA	1216	G	O4'-C1'-N9	8.20	114.76	108.20
35	BB	620	G	C8-N9-C1'	8.20	137.65	127.00
38	BE	15	A	C1'-O4'-C4'	-8.20	103.34	109.90
85	AA	307	G	C5'-C4'-C3'	8.20	129.11	116.00
35	BB	1063	C	C2'-C3'-O3'	8.19	127.53	109.50
38	BE	70	C	C6-N1-C1'	8.19	130.63	120.80
35	BB	406	A	O4'-C1'-C2'	-8.19	97.61	105.80
35	BB	1303	A	N1-C6-N6	8.19	123.52	118.60
85	AA	337	C	P-O3'-C3'	-8.19	109.87	119.70
85	AA	875	C	C2-N3-C4	-8.19	115.80	119.90
85	AA	1863	A	C3'-C2'-C1'	-8.19	94.95	101.50
35	BB	1131	C	C6-N1-C2	-8.19	117.02	120.30
85	AA	172	A	P-O3'-C3'	-8.19	109.87	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	927	A	N1-C6-N6	8.19	123.51	118.60
34	BA	711	C	O4'-C1'-N1	8.19	114.75	108.20
34	BA	771	A	C1'-O4'-C4'	-8.19	103.35	109.90
34	BA	1541	G	C6-C5-N7	-8.19	125.49	130.40
38	BE	131	C	P-O3'-C3'	8.19	129.53	119.70
56	BW	88	ARG	NE-CZ-NH1	8.19	124.39	120.30
85	AA	44	C	O4'-C1'-N1	8.19	114.75	108.20
85	AA	580	C	C6-N1-C2	-8.19	117.02	120.30
34	BA	1234	U	P-O5'-C5'	-8.19	107.80	120.90
35	BB	732	G	C5'-C4'-C3'	-8.19	102.90	116.00
38	BE	96	G	C6-N1-C2	-8.19	120.19	125.10
85	AA	556	C	C4'-C3'-C2'	8.19	110.79	102.60
85	AA	639	C	C6-N1-C1'	-8.19	110.97	120.80
85	AA	751	C	O4'-C1'-N1	8.19	114.75	108.20
85	AA	1254	A	C3'-C2'-C1'	-8.19	94.95	101.50
34	BA	1254	C	O4'-C1'-N1	8.19	114.75	108.20
34	BA	1439	C	C2-N1-C1'	-8.19	109.80	118.80
40	BG	166	C	C6-N1-C2	-8.19	117.03	120.30
35	BB	858	U	P-O3'-C3'	-8.19	109.88	119.70
38	BE	162	U	C2-N3-C4	-8.19	122.09	127.00
85	AA	139	G	O4'-C1'-N9	8.19	114.75	108.20
85	AA	246	C	O4'-C1'-N1	8.19	114.75	108.20
85	AA	1247	A	O4'-C1'-N9	8.19	114.75	108.20
85	AA	1458	G	C2'-C3'-O3'	8.19	127.51	109.50
85	AA	1832	G	C8-N9-C1'	8.19	137.64	127.00
34	BA	372	U	C6-N1-C1'	8.18	132.66	121.20
34	BA	1637	G	N3-C4-C5	-8.18	124.51	128.60
35	BB	42	A	C5'-C4'-C3'	8.18	129.09	116.00
35	BB	658	G	P-O5'-C5'	8.18	133.99	120.90
38	BE	90	G	C5-C6-O6	-8.18	123.69	128.60
37	BD	29	C	C3'-C2'-C1'	-8.18	94.95	101.50
39	BF	24	G	P-O5'-C5'	-8.18	107.81	120.90
85	AA	417	U	P-O5'-C5'	-8.18	107.81	120.90
85	AA	1236	G	C8-N9-C1'	8.18	137.64	127.00
85	AA	1667	C	C5'-C4'-C3'	-8.18	102.91	116.00
34	BA	1620	U	C2'-C3'-O3'	8.18	127.50	109.50
34	BA	1699	A	C5-C6-N6	8.18	130.25	123.70
35	BB	1442	C	O4'-C1'-N1	8.18	114.75	108.20
38	BE	59	U	N1-C2-O2	8.18	128.53	122.80
34	BA	1284	G	N1-C2-N2	-8.18	108.84	116.20
85	AA	494	G	C5'-C4'-O4'	8.18	118.92	109.10
82	Bw	68	ARG	NE-CZ-NH1	8.18	124.39	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1468	G	O4'-C1'-N9	8.18	114.74	108.20
34	BA	343	G	P-O3'-C3'	-8.18	109.89	119.70
34	BA	593	G	N9-C1'-C2'	8.18	124.63	114.00
34	BA	605	G	P-O3'-C3'	-8.18	109.89	119.70
34	BA	1677	C	P-O5'-C5'	-8.18	107.82	120.90
35	BB	1228	A	P-O5'-C5'	8.18	133.98	120.90
35	BB	1285	U	O4'-C1'-N1	8.18	114.74	108.20
38	BE	133	C	C2-N3-C4	-8.18	115.81	119.90
85	AA	1280	U	P-O3'-C3'	8.18	129.51	119.70
85	AA	1566	A	O4'-C1'-N9	8.18	114.74	108.20
85	AA	2130	G	C5-C6-O6	-8.18	123.69	128.60
19	AK	99	TYR	CB-CG-CD2	-8.17	116.10	121.00
34	BA	73	G	C4-N9-C1'	-8.17	115.87	126.50
34	BA	89	G	C8-N9-C4	8.17	109.67	106.40
34	BA	232	U	C5'-C4'-C3'	8.17	129.08	116.00
34	BA	805	A	P-O5'-C5'	8.17	133.98	120.90
34	BA	535	G	N3-C4-C5	-8.17	124.51	128.60
34	BA	804	G	P-O5'-C5'	-8.17	107.83	120.90
34	BA	1591	G	N9-C1'-C2'	-8.17	103.01	112.00
35	BB	168	U	P-O5'-C5'	8.17	133.98	120.90
40	BG	96	C	P-O5'-C5'	-8.17	107.83	120.90
41	BH	127	A	P-O3'-C3'	8.17	129.51	119.70
85	AA	210	G	O4'-C1'-N9	8.17	114.74	108.20
34	BA	678	C	N1-C2-N3	8.17	124.92	119.20
34	BA	903	C	P-O3'-C3'	-8.17	109.89	119.70
34	BA	1160	U	C5'-C4'-C3'	-8.17	102.93	116.00
35	BB	1445	A	O3'-P-O5'	8.17	119.52	104.00
34	BA	1265	G	C5-C6-O6	-8.17	123.70	128.60
40	BG	9	G	C5-C6-N1	8.17	115.58	111.50
41	BH	20	A	C6-N1-C2	-8.17	113.70	118.60
85	AA	1881	C	P-O3'-C3'	8.17	129.50	119.70
85	AA	2153	G	N1-C6-O6	8.17	124.80	119.90
34	BA	1003	A	O4'-C1'-C2'	8.17	114.95	107.60
35	BB	799	A	C4'-C3'-C2'	8.17	110.77	102.60
35	BB	1260	A	P-O5'-C5'	-8.17	107.83	120.90
38	BE	133	C	O4'-C1'-N1	8.17	114.73	108.20
85	AA	570	U	O4'-C1'-N1	8.17	114.73	108.20
83	Bx	263	ARG	NE-CZ-NH1	8.16	124.38	120.30
85	AA	389	A	P-O5'-C5'	-8.16	107.84	120.90
85	AA	1015	U	C6-N1-C2	-8.16	116.10	121.00
85	AA	2119	C	C3'-C2'-C1'	-8.16	94.97	101.50
34	BA	1510	C	O4'-C1'-N1	8.16	114.73	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1199	A	C3'-C2'-C1'	-8.16	94.97	101.50
85	AA	1110	A	O4'-C1'-N9	8.16	114.73	108.20
85	AA	1556	G	O4'-C1'-N9	8.16	114.73	108.20
34	BA	960	C	O4'-C1'-N1	8.16	114.73	108.20
35	BB	1376	G	C5'-C4'-C3'	-8.16	102.95	116.00
36	BC	143	C	C6-N1-C2	-8.16	117.04	120.30
85	AA	506	G	P-O3'-C3'	-8.16	109.91	119.70
85	AA	2201	A	C8-N9-C4	8.16	109.06	105.80
34	BA	876	C	C5'-C4'-C3'	8.16	129.05	116.00
35	BB	905	C	C6-N1-C2	-8.16	117.04	120.30
85	AA	713	G	C3'-C2'-C1'	-8.16	94.97	101.50
34	BA	462	C	O4'-C1'-N1	8.16	114.72	108.20
34	BA	1120	U	O4'-C1'-N1	8.16	114.72	108.20
34	BA	1505	G	C6-C5-N7	-8.16	125.51	130.40
35	BB	757	C	O4'-C1'-N1	8.16	114.72	108.20
35	BB	823	G	P-O3'-C3'	-8.16	109.91	119.70
41	BH	34	G	N1-C6-O6	-8.16	115.01	119.90
51	BR	52	LEU	CB-CA-C	8.16	125.70	110.20
85	AA	146	U	O4'-C1'-N1	8.16	114.72	108.20
85	AA	205	A	C3'-C2'-C1'	-8.16	94.97	101.50
34	BA	376	U	C1'-O4'-C4'	-8.15	103.38	109.90
34	BA	926	A	C4-N9-C1'	-8.15	111.62	126.30
34	BA	1203	G	P-O5'-C5'	-8.15	107.86	120.90
34	BA	1493	U	O5'-C5'-C4'	8.15	127.19	111.70
85	AA	415	G	C5-C6-O6	-8.15	123.71	128.60
85	AA	2075	C	C2-N1-C1'	-8.15	109.83	118.80
34	BA	1638	U	C6-N1-C1'	8.15	132.61	121.20
38	BE	130	G	C8-N9-C1'	-8.15	116.40	127.00
66	Bg	94	ARG	NE-CZ-NH1	8.15	124.38	120.30
35	BB	839	G	C8-N9-C1'	8.15	137.60	127.00
40	BG	16	G	N9-C1'-C2'	-8.15	103.03	112.00
85	AA	1612	C	O4'-C1'-N1	8.15	114.72	108.20
35	BB	14	C	C2-N3-C4	-8.15	115.83	119.90
41	BH	21	G	C6-N1-C2	-8.15	120.21	125.10
85	AA	268	A	C3'-C2'-C1'	-8.15	94.98	101.50
85	AA	1980	A	O4'-C1'-N9	8.15	114.72	108.20
4	A3	49	TYR	CB-CG-CD1	-8.15	116.11	121.00
34	BA	882	G	C2-N3-C4	8.15	115.97	111.90
35	BB	653	G	C8-N9-C1'	8.15	137.59	127.00
37	BD	91	U	C6-N1-C2	-8.15	116.11	121.00
38	BE	34	C	P-O5'-C5'	-8.15	107.86	120.90
85	AA	422	G	C8-N9-C1'	8.15	137.59	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1049	G	N1-C6-O6	8.15	124.79	119.90
34	BA	1475	G	C8-N9-C4	8.15	109.66	106.40
35	BB	455	G	P-O3'-C3'	8.15	129.48	119.70
39	BF	31	U	O3'-P-O5'	-8.15	88.52	104.00
85	AA	1	G	O4'-C1'-N9	8.15	114.72	108.20
85	AA	392	G	N1-C6-O6	-8.15	115.01	119.90
6	A5	22	ARG	NE-CZ-NH1	8.14	124.37	120.30
34	BA	1533	G	C5'-C4'-C3'	8.14	129.03	116.00
34	BA	764	G	O4'-C4'-C3'	-8.14	95.86	104.00
85	AA	339	A	C8-N9-C1'	8.14	142.36	127.70
85	AA	2095	U	C2-N3-C4	-8.14	122.11	127.00
85	AA	1713	A	C1'-O4'-C4'	-8.14	103.39	109.90
85	AA	620	U	C1'-O4'-C4'	-8.14	103.39	109.90
25	AR	17	TYR	CB-CG-CD2	-8.14	116.12	121.00
34	BA	1086	A	O4'-C1'-N9	8.14	114.71	108.20
35	BB	539	G	N9-C1'-C2'	-8.14	103.05	112.00
85	AA	2181	G	C8-N9-C1'	8.14	137.58	127.00
85	AA	2242	U	O5'-P-OP2	-8.14	98.37	105.70
34	BA	557	U	O5'-P-OP2	8.14	120.47	110.70
34	BA	1103	G	C8-N9-C4	8.14	109.66	106.40
85	AA	932	A	O5'-C5'-C4'	-8.14	96.24	111.70
34	BA	1483	U	C4'-C3'-C2'	8.14	110.74	102.60
35	BB	701	U	C2-N1-C1'	8.14	127.47	117.70
35	BB	1029	U	C2'-C3'-O3'	8.14	127.40	109.50
58	BY	60	ARG	NE-CZ-NH2	-8.14	116.23	120.30
38	BE	32	U	P-O3'-C3'	8.14	129.46	119.70
38	BE	123	A	N1-C2-N3	-8.14	125.23	129.30
41	BH	46	C	C6-N1-C1'	8.14	130.56	120.80
85	AA	476	C	C6-N1-C2	-8.14	117.05	120.30
85	AA	1397	U	O4'-C1'-N1	8.14	114.71	108.20
85	AA	1490	A	P-O3'-C3'	8.14	129.47	119.70
85	AA	1368	G	P-O3'-C3'	-8.14	109.94	119.70
34	BA	816	G	N9-C1'-C2'	-8.14	103.05	112.00
35	BB	642	G	C4-N9-C1'	8.13	137.08	126.50
35	BB	846	A	C1'-O4'-C4'	-8.14	103.39	109.90
35	BB	1220	A	C5-C6-N6	-8.13	117.19	123.70
85	AA	274	A	P-O3'-C3'	-8.13	109.94	119.70
85	AA	492	C	C6-N1-C2	-8.13	117.05	120.30
85	AA	494	G	C3'-C2'-C1'	-8.13	94.99	101.50
85	AA	736	U	C6-N1-C1'	8.13	132.59	121.20
85	AA	975	G	N1-C6-O6	-8.13	115.02	119.90
40	BG	15	G	N9-C1'-C2'	-8.13	103.06	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1549	U	C2-N1-C1'	-8.13	107.94	117.70
35	BB	1122	C	O4'-C1'-N1	8.13	114.70	108.20
35	BB	1266	A	O4'-C1'-N9	8.13	114.70	108.20
45	BL	130	TYR	CB-CG-CD2	-8.13	116.12	121.00
85	AA	1247	A	P-O5'-C5'	-8.13	107.89	120.90
85	AA	1700	C	P-O5'-C5'	8.13	133.91	120.90
85	AA	2191	C	O4'-C1'-N1	8.13	114.70	108.20
34	BA	101	G	N3-C2-N2	8.13	125.59	119.90
34	BA	661	C	C4'-C3'-C2'	8.13	110.73	102.60
34	BA	691	A	O4'-C1'-N9	8.13	114.70	108.20
34	BA	1282	G	C3'-C2'-C1'	-8.13	95.00	101.50
34	BA	1662	U	C5-C4-O4	8.13	130.78	125.90
39	BF	39	C	C5'-C4'-O4'	8.13	118.86	109.10
85	AA	790	A	O5'-C5'-C4'	8.13	127.14	111.70
34	BA	1321	A	P-O3'-C3'	-8.13	109.95	119.70
35	BB	380	G	C5'-C4'-C3'	-8.13	103.00	116.00
38	BE	132	U	N3-C2-O2	-8.13	116.51	122.20
85	AA	532	G	O4'-C1'-N9	8.13	114.70	108.20
85	AA	643	C	C6-N1-C2	-8.13	117.05	120.30
85	AA	1975	G	C8-N9-C1'	8.13	137.56	127.00
34	BA	1441	C	C5'-C4'-C3'	8.13	129.00	116.00
35	BB	427	U	C2-N3-C4	-8.12	122.12	127.00
85	AA	203	C	C6-N1-C1'	-8.13	111.05	120.80
85	AA	1121	U	C1'-O4'-C4'	-8.12	103.40	109.90
85	AA	1810	C	O4'-C1'-N1	8.12	114.70	108.20
85	AA	1836	U	O4'-C1'-N1	8.12	114.70	108.20
85	AA	1970	A	O4'-C1'-N9	8.13	114.70	108.20
34	BA	110	C	C5'-C4'-C3'	-8.12	103.01	116.00
35	BB	21	C	O4'-C1'-N1	8.12	114.70	108.20
35	BB	1315	C	C2-N3-C4	-8.12	115.84	119.90
36	BC	147	G	C8-N9-C4	-8.12	103.15	106.40
38	BE	13	A	C3'-C2'-C1'	-8.12	95.00	101.50
85	AA	779	G	C5-C6-O6	-8.12	123.73	128.60
85	AA	1515	A	C4'-C3'-C2'	-8.12	94.48	102.60
86	AB	14	A	C5'-C4'-C3'	8.12	128.99	116.00
35	BB	144	G	P-O5'-C5'	8.12	133.89	120.90
35	BB	817	C	C5'-C4'-C3'	-8.12	103.01	116.00
35	BB	1515	C	N3-C2-O2	-8.12	116.22	121.90
85	AA	647	C	P-O3'-C3'	8.12	129.44	119.70
85	AA	994	A	C5'-C4'-C3'	-8.12	103.01	116.00
34	BA	252	A	N9-C1'-C2'	-8.12	103.07	112.00
34	BA	1316	G	P-O5'-C5'	-8.12	107.91	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
62	Bc	16	SER	N-CA-CB	8.12	122.68	110.50
71	Bl	95	TRP	CA-C-N	8.12	135.06	117.20
85	AA	106	G	N3-C2-N2	8.12	125.58	119.90
34	BA	77	C	O4'-C1'-N1	8.12	114.69	108.20
34	BA	634	U	O5'-P-OP2	-8.12	98.40	105.70
34	BA	1486	U	N3-C2-O2	-8.12	116.52	122.20
85	AA	370	A	P-O3'-C3'	-8.12	109.96	119.70
35	BB	2	C	C6-N1-C2	-8.11	117.05	120.30
36	BC	25	C	C6-N1-C2	-8.11	117.05	120.30
38	BE	90	G	C2'-C3'-O3'	8.12	127.35	109.50
85	AA	1535	C	C5'-C4'-C3'	-8.12	103.01	116.00
85	AA	2088	U	P-O3'-C3'	-8.12	109.96	119.70
85	AA	2153	G	O4'-C1'-N9	8.12	114.69	108.20
34	BA	214	A	N7-C8-N9	8.11	117.86	113.80
38	BE	32	U	C5'-C4'-C3'	-8.11	103.02	116.00
41	BH	129	G	C4-N9-C1'	-8.11	115.95	126.50
57	BX	139	ASP	CB-CG-OD1	8.11	125.60	118.30
35	BB	25	A	O4'-C1'-N9	8.11	114.69	108.20
35	BB	1234	G	C5-C6-N1	8.11	115.56	111.50
62	Bc	78	LYS	CD-CE-NZ	8.11	130.35	111.70
85	AA	380	C	O4'-C1'-N1	8.11	114.69	108.20
85	AA	1239	C	C6-N1-C2	-8.11	117.06	120.30
85	AA	2119	C	N3-C2-O2	-8.11	116.22	121.90
85	AA	1456	A	N9-C1'-C2'	-8.11	103.08	112.00
34	BA	1379	G	C8-N9-C4	-8.11	103.16	106.40
34	BA	102	G	O4'-C1'-N9	8.11	114.69	108.20
34	BA	1320	A	P-O5'-C5'	8.11	133.87	120.90
35	BB	973	G	C5'-C4'-C3'	-8.11	103.03	116.00
35	BB	1020	U	C2-N3-C4	-8.11	122.13	127.00
35	BB	1301	U	P-O3'-C3'	8.11	129.43	119.70
35	BB	1458	U	P-O5'-C5'	-8.11	107.93	120.90
41	BH	18	C	O4'-C1'-N1	8.11	114.69	108.20
85	AA	893	G	C4-C5-C6	-8.11	113.93	118.80
85	AA	1231	G	C8-N9-C1'	8.11	137.54	127.00
40	BG	133	C	C2-N1-C1'	8.11	127.72	118.80
34	BA	146	G	C6-N1-C2	-8.11	120.24	125.10
34	BA	843	G	N3-C2-N2	8.11	125.57	119.90
34	BA	1058	C	P-O3'-C3'	8.11	129.43	119.70
35	BB	1141	A	O4'-C4'-C3'	-8.11	95.89	104.00
35	BB	1270	C	O4'-C1'-N1	8.11	114.69	108.20
36	BC	94	C	O4'-C1'-N1	8.11	114.69	108.20
85	AA	358	U	C5'-C4'-C3'	8.11	128.97	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	534	A	N1-C6-N6	-8.11	113.74	118.60
38	BE	195	G	N3-C4-N9	8.11	130.86	126.00
85	AA	230	U	O4'-C1'-N1	8.11	114.68	108.20
85	AA	610	C	C4'-C3'-C2'	-8.11	94.49	102.60
85	AA	862	U	N3-C2-O2	-8.11	116.53	122.20
85	AA	1665	G	C3'-C2'-C1'	-8.11	95.02	101.50
34	BA	1261	G	C1'-O4'-C4'	-8.10	103.42	109.90
34	BA	998	U	C5'-C4'-C3'	8.10	128.96	116.00
34	BA	1281	U	C1'-O4'-C4'	-8.10	103.42	109.90
35	BB	449	C	N3-C2-O2	-8.10	116.23	121.90
35	BB	858	U	C5'-C4'-C3'	-8.10	103.03	116.00
85	AA	646	C	P-O3'-C3'	-8.10	109.97	119.70
35	BB	1534	U	C5'-C4'-C3'	8.10	128.96	116.00
85	AA	874	A	C8-N9-C4	8.10	109.04	105.80
36	BC	86	U	C3'-C2'-C1'	-8.10	95.02	101.50
40	BG	58	G	C5'-C4'-C3'	-8.10	103.04	116.00
85	AA	755	G	N3-C2-N2	8.10	125.57	119.90
85	AA	1560	A	C5'-C4'-O4'	8.10	118.82	109.10
34	BA	8	G	N9-C1'-C2'	-8.10	103.09	112.00
34	BA	210	G	P-O3'-C3'	8.10	129.42	119.70
35	BB	62	C	C6-N1-C2	-8.10	117.06	120.30
38	BE	177	U	P-O3'-C3'	-8.10	109.98	119.70
56	BW	73	ARG	CD-NE-CZ	8.10	134.94	123.60
65	Bf	194	ARG	NE-CZ-NH1	8.10	124.35	120.30
85	AA	375	C	C4'-C3'-C2'	8.10	110.70	102.60
85	AA	414	C	C5-C4-N4	8.10	125.87	120.20
35	BB	1141	A	O4'-C1'-N9	8.10	114.68	108.20
41	BH	47	G	C5-C6-N1	8.10	115.55	111.50
85	AA	164	G	O5'-P-OP2	8.10	120.42	110.70
85	AA	1896	G	N1-C6-O6	8.10	124.76	119.90
34	BA	680	C	C3'-C2'-C1'	-8.10	95.02	101.50
85	AA	1520	A	C5-C6-N6	-8.10	117.22	123.70
34	BA	1103	G	C4-N9-C1'	-8.09	115.98	126.50
35	BB	42	A	C4'-C3'-C2'	8.09	110.69	102.60
35	BB	632	U	P-O3'-C3'	-8.09	109.99	119.70
35	BB	1432	U	P-O5'-C5'	-8.09	107.95	120.90
52	BS	157	ARG	C-N-CA	8.09	141.94	121.70
38	BE	96	G	C4-C5-C6	-8.09	113.94	118.80
85	AA	699	U	C5'-C4'-C3'	-8.09	103.05	116.00
34	BA	134	U	C6-N1-C2	-8.09	116.15	121.00
34	BA	221	G	C5'-C4'-C3'	-8.09	103.05	116.00
34	BA	1450	G	C5-C6-O6	-8.09	123.75	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	256	G	N1-C6-O6	8.09	124.75	119.90
35	BB	399	A	N1-C6-N6	8.09	123.45	118.60
36	BC	158	U	O4'-C1'-N1	8.09	114.67	108.20
79	Bt	23	PHE	CB-CG-CD2	-8.09	115.14	120.80
85	AA	893	G	P-O3'-C3'	-8.09	109.99	119.70
85	AA	2157	G	O4'-C1'-N9	8.09	114.67	108.20
34	BA	308	C	C3'-C2'-C1'	-8.09	95.03	101.50
34	BA	361	C	C6-N1-C2	-8.09	117.06	120.30
85	AA	1560	A	O4'-C1'-N9	8.09	114.67	108.20
34	BA	217	C	C6-N1-C1'	-8.09	111.10	120.80
34	BA	238	C	P-O3'-C3'	-8.09	110.00	119.70
34	BA	616	G	C8-N9-C1'	8.09	137.51	127.00
34	BA	894	G	C2-N3-C4	-8.09	107.86	111.90
34	BA	967	C	C1'-O4'-C4'	-8.09	103.43	109.90
35	BB	1207	C	P-O3'-C3'	-8.09	110.00	119.70
85	AA	443	A	C8-N9-C4	8.09	109.03	105.80
85	AA	1890	C	O4'-C1'-N1	8.09	114.67	108.20
85	AA	1992	A	C5'-C4'-C3'	8.09	128.94	116.00
34	BA	1229	G	C5-C6-O6	-8.09	123.75	128.60
38	BE	16	C	O4'-C4'-C3'	-8.09	95.92	104.00
85	AA	1459	C	N3-C2-O2	-8.09	116.24	121.90
35	BB	1212	C	O4'-C1'-N1	8.08	114.67	108.20
85	AA	428	G	N1-C6-O6	8.08	124.75	119.90
85	AA	1506	U	N3-C2-O2	-8.08	116.54	122.20
85	AA	2048	C	C6-N1-C2	-8.08	117.07	120.30
20	AL	124	VAL	C-N-CA	8.08	141.91	121.70
34	BA	1173	C	C1'-O4'-C4'	-8.08	103.44	109.90
34	BA	1732	A	N3-C4-N9	8.08	133.87	127.40
35	BB	1512	C	P-O5'-C5'	8.08	133.83	120.90
37	BD	84	U	C5'-C4'-C3'	-8.08	103.07	116.00
38	BE	104	G	C4'-C3'-C2'	-8.08	94.52	102.60
40	BG	64	C	C2-N3-C4	-8.08	115.86	119.90
47	BN	214	GLU	N-CA-CB	-8.08	96.05	110.60
85	AA	126	U	C3'-C2'-C1'	-8.08	95.03	101.50
34	BA	1503	U	O4'-C1'-N1	8.08	114.66	108.20
34	BA	1617	U	P-O3'-C3'	-8.08	110.00	119.70
85	AA	1982	C	C6-N1-C2	-8.08	117.07	120.30
34	BA	315	U	O5'-C5'-C4'	8.08	127.05	111.70
34	BA	447	U	C6-N1-C1'	8.08	132.51	121.20
34	BA	867	C	N3-C4-N4	-8.08	112.34	118.00
35	BB	161	G	O4'-C1'-N9	8.08	114.66	108.20
35	BB	1288	G	C5'-C4'-C3'	8.08	128.93	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2225	G	O4'-C1'-N9	8.08	114.67	108.20
85	AA	1884	A	O4'-C1'-N9	8.08	114.66	108.20
34	BA	709	C	C6-N1-C2	-8.08	117.07	120.30
35	BB	786	A	N1-C6-N6	-8.08	113.75	118.60
85	AA	1235	G	C5'-C4'-C3'	-8.08	103.08	116.00
35	BB	1431	G	C8-N9-C1'	8.08	137.50	127.00
36	BC	52	A	C3'-C2'-C1'	-8.08	95.04	101.50
85	AA	1682	U	O3'-P-O5'	-8.08	88.65	104.00
35	BB	139	G	C4-N9-C1'	-8.07	116.00	126.50
35	BB	679	G	N9-C1'-C2'	-8.07	103.12	112.00
35	BB	832	C	C6-N1-C2	-8.07	117.07	120.30
35	BB	1475	U	C1'-O4'-C4'	-8.07	103.44	109.90
34	BA	62	A	C5'-C4'-C3'	-8.07	103.08	116.00
34	BA	103	G	C3'-C2'-C1'	-8.07	95.04	101.50
34	BA	1156	U	C2-N1-C1'	-8.07	108.01	117.70
36	BC	55	U	C5'-C4'-C3'	-8.07	103.08	116.00
85	AA	1411	C	O4'-C1'-N1	8.07	114.66	108.20
85	AA	1517	G	C5'-C4'-C3'	-8.07	103.08	116.00
39	BF	30	C	P-O3'-C3'	-8.07	110.01	119.70
85	AA	1645	G	C4-N9-C1'	-8.07	116.00	126.50
86	AB	63	G	C1'-O4'-C4'	-8.07	103.44	109.90
35	BB	708	C	N3-C4-C5	-8.07	118.67	121.90
34	BA	166	G	C8-N9-C1'	8.07	137.49	127.00
34	BA	1467	U	C3'-C2'-C1'	-8.07	95.05	101.50
35	BB	1421	C	O4'-C1'-N1	8.07	114.66	108.20
40	BG	46	G	C5-C6-O6	-8.07	123.76	128.60
81	Bv	68	TYR	CB-CG-CD1	8.07	125.84	121.00
85	AA	463	G	C5-C6-O6	-8.07	123.76	128.60
85	AA	743	C	C2-N1-C1'	-8.07	109.92	118.80
34	BA	762	A	C5'-C4'-C3'	-8.07	103.09	116.00
34	BA	1294	C	N1-C1'-C2'	8.07	124.49	114.00
34	BA	1738	G	C8-N9-C1'	8.07	137.49	127.00
35	BB	652	G	C4'-C3'-C2'	-8.07	94.53	102.60
37	BD	48	G	N9-C4-C5	-8.07	102.17	105.40
35	BB	1538	G	C4-N9-C1'	-8.07	116.01	126.50
37	BD	94	C	P-O5'-C5'	-8.07	107.99	120.90
85	AA	549	A	C4-N9-C1'	-8.07	111.78	126.30
85	AA	735	G	C8-N9-C4	-8.07	103.17	106.40
85	AA	2241	C	C1'-O4'-C4'	-8.06	103.45	109.90
34	BA	470	C	P-O3'-C3'	8.06	129.38	119.70
34	BA	748	C	C1'-O4'-C4'	-8.06	103.45	109.90
85	AA	698	G	C1'-O4'-C4'	-8.06	103.45	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	780	U	O4'-C1'-N1	8.06	114.65	108.20
37	BD	64	A	N1-C6-N6	8.06	123.44	118.60
38	BE	127	G	C4'-C3'-C2'	8.06	110.66	102.60
85	AA	456	A	C8-N9-C4	8.06	109.03	105.80
85	AA	841	U	P-O3'-C3'	-8.06	110.02	119.70
85	AA	1787	G	P-O3'-C3'	8.06	129.38	119.70
34	BA	53	G	C8-N9-C1'	8.06	137.48	127.00
37	BD	31	U	O4'-C1'-N1	8.06	114.65	108.20
82	Bw	218	TRP	CB-CG-CD2	-8.06	116.12	126.60
85	AA	919	U	C2-N1-C1'	8.06	127.37	117.70
85	AA	484	G	P-O5'-C5'	8.06	133.80	120.90
85	AA	2014	G	N3-C2-N2	-8.06	114.26	119.90
34	BA	819	G	P-O3'-C3'	-8.06	110.03	119.70
35	BB	1212	C	C4'-C3'-C2'	8.06	110.66	102.60
38	BE	168	C	O4'-C1'-N1	8.06	114.65	108.20
85	AA	855	G	P-O3'-C3'	-8.06	110.03	119.70
85	AA	989	U	O5'-C5'-C4'	-8.06	96.39	111.70
85	AA	775	C	C3'-C2'-C1'	-8.06	95.05	101.50
85	AA	1281	G	C1'-O4'-C4'	-8.06	103.45	109.90
85	AA	1471	G	C5-C6-N1	8.06	115.53	111.50
85	AA	1839	G	C5'-C4'-C3'	-8.06	103.11	116.00
85	AA	2075	C	C6-N1-C1'	8.06	130.47	120.80
34	BA	847	U	P-O5'-C5'	8.05	133.79	120.90
34	BA	1798	G	P-O5'-C5'	8.05	133.79	120.90
35	BB	706	G	C5'-C4'-C3'	8.05	128.89	116.00
35	BB	1141	A	P-O5'-C5'	-8.05	108.01	120.90
35	BB	1426	G	C5-C6-N1	8.05	115.53	111.50
41	BH	81	U	O4'-C1'-N1	8.05	114.64	108.20
70	Bk	94	ARG	NE-CZ-NH1	8.05	124.33	120.30
85	AA	650	G	C8-N9-C1'	8.05	137.47	127.00
85	AA	1229	G	N3-C4-C5	-8.06	124.57	128.60
38	BE	7	U	N3-C2-O2	-8.05	116.56	122.20
34	BA	1415	C	O4'-C1'-N1	8.05	114.64	108.20
34	BA	1448	G	N1-C6-O6	8.05	124.73	119.90
77	Br	139	ARG	NE-CZ-NH2	-8.05	116.28	120.30
85	AA	768	C	C2-N1-C1'	-8.05	109.94	118.80
34	BA	110	C	C1'-O4'-C4'	-8.05	103.46	109.90
34	BA	314	A	C1'-O4'-C4'	-8.05	103.46	109.90
35	BB	37	C	C3'-C2'-C1'	-8.05	95.06	101.50
35	BB	109	U	C5'-C4'-C3'	-8.05	103.12	116.00
35	BB	534	C	C6-N1-C2	-8.05	117.08	120.30
35	BB	1016	C	O4'-C1'-N1	8.05	114.64	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1453	G	C2-N3-C4	-8.05	107.88	111.90
35	BB	1495	U	C1'-O4'-C4'	-8.05	103.46	109.90
44	BK	185	ARG	NE-CZ-NH1	8.05	124.32	120.30
85	AA	848	C	O4'-C1'-N1	8.05	114.64	108.20
85	AA	1560	A	C1'-O4'-C4'	-8.05	103.46	109.90
85	AA	1909	C	C6-N1-C2	-8.05	117.08	120.30
35	BB	1230	A	P-O5'-C5'	-8.05	108.03	120.90
40	BG	9	G	C1'-O4'-C4'	-8.05	103.46	109.90
34	BA	389	U	O4'-C1'-N1	8.04	114.64	108.20
41	BH	109	G	N1-C6-O6	-8.04	115.07	119.90
53	BT	132	PHE	CB-CG-CD1	8.05	126.43	120.80
75	Bp	27	HIS	CA-CB-CG	-8.04	99.92	113.60
85	AA	1544	G	N1-C6-O6	-8.04	115.07	119.90
85	AA	1845	G	C4-N9-C1'	-8.05	116.04	126.50
86	AB	19	G	P-O5'-C5'	-8.04	108.03	120.90
6	A5	25	MET	CG-SD-CE	-8.04	87.33	100.20
34	BA	1672	C	C6-N1-C2	-8.04	117.08	120.30
38	BE	135	A	C1'-O4'-C4'	-8.04	103.47	109.90
85	AA	1012	C	P-O3'-C3'	8.04	129.35	119.70
73	Bn	56	ARG	NE-CZ-NH1	8.04	124.32	120.30
84	By	44	PHE	CB-CG-CD1	8.04	126.43	120.80
85	AA	1988	A	N1-C6-N6	-8.04	113.78	118.60
34	BA	65	A	C5'-C4'-C3'	8.04	128.86	116.00
34	BA	290	G	C4-N9-C1'	-8.04	116.05	126.50
34	BA	1190	A	O3'-P-O5'	-8.04	88.72	104.00
35	BB	1480	G	O4'-C1'-N9	8.04	114.63	108.20
38	BE	75	C	C6-N1-C1'	8.04	130.45	120.80
38	BE	113	C	C6-N1-C2	-8.04	117.08	120.30
38	BE	176	G	C5'-C4'-C3'	8.04	128.86	116.00
39	BF	37	C	O4'-C1'-N1	8.04	114.63	108.20
34	BA	525	A	C4-N9-C1'	-8.04	111.83	126.30
34	BA	566	G	N9-C1'-C2'	-8.04	103.16	112.00
34	BA	1699	A	N9-C1'-C2'	-8.04	103.16	112.00
35	BB	816	U	P-O5'-C5'	-8.04	108.04	120.90
37	BD	35	C	P-O3'-C3'	-8.04	110.05	119.70
85	AA	1557	U	P-O3'-C3'	8.04	129.35	119.70
35	BB	1014	U	N3-C2-O2	-8.04	116.57	122.20
37	BD	77	A	N9-C1'-C2'	-8.04	103.16	112.00
40	BG	132	U	C2-N1-C1'	-8.04	108.06	117.70
34	BA	292	C	O4'-C1'-N1	8.04	114.63	108.20
34	BA	398	G	C5-C6-O6	-8.04	123.78	128.60
34	BA	1244	G	C5-C6-O6	-8.04	123.78	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1	U	C5'-C4'-C3'	8.04	128.86	116.00
35	BB	1032	U	C1'-O4'-C4'	-8.04	103.47	109.90
36	BC	5	U	C2-N3-C4	-8.04	122.18	127.00
85	AA	918	U	C1'-O4'-C4'	-8.04	103.47	109.90
36	BC	53	A	C8-N9-C4	8.03	109.01	105.80
41	BH	58	C	O4'-C1'-N1	8.03	114.63	108.20
85	AA	866	U	P-O3'-C3'	-8.04	110.06	119.70
85	AA	889	G	O5'-C5'-C4'	-8.03	96.44	111.70
85	AA	964	C	C6-N1-C1'	-8.04	111.16	120.80
85	AA	2169	C	O4'-C1'-N1	8.03	114.63	108.20
34	BA	696	A	P-O5'-C5'	-8.03	108.05	120.90
35	BB	522	A	C3'-C2'-C1'	-8.03	95.07	101.50
45	BL	179	PHE	CB-CG-CD2	-8.03	115.18	120.80
34	BA	254	U	O4'-C1'-N1	8.03	114.62	108.20
35	BB	1039	A	C5'-C4'-C3'	-8.03	103.15	116.00
36	BC	141	C	O4'-C1'-N1	8.03	114.62	108.20
41	BH	134	U	O4'-C4'-C3'	-8.03	95.97	104.00
85	AA	285	C	O5'-C5'-C4'	8.03	126.96	111.70
27	AT	65	PHE	CB-CG-CD1	8.03	126.42	120.80
34	BA	1080	U	C4'-C3'-C2'	-8.03	94.57	102.60
35	BB	589	U	C1'-O4'-C4'	-8.03	103.48	109.90
35	BB	1003	G	C3'-C2'-C1'	-8.03	95.08	101.50
36	BC	156	A	O4'-C1'-N9	8.03	114.62	108.20
38	BE	134	A	C5'-C4'-O4'	-8.03	99.46	109.10
85	AA	1727	U	P-O3'-C3'	-8.03	110.06	119.70
59	BZ	10	ARG	NE-CZ-NH1	8.03	124.31	120.30
85	AA	514	U	C2-N3-C4	-8.03	122.18	127.00
34	BA	502	U	N3-C2-O2	-8.03	116.58	122.20
85	AA	14	C	C2-N3-C4	-8.03	115.89	119.90
85	AA	565	G	C5'-C4'-C3'	-8.03	103.16	116.00
85	AA	774	C	C2-N3-C4	-8.03	115.89	119.90
34	BA	636	G	C5-C6-O6	8.03	133.41	128.60
34	BA	736	G	N1-C6-O6	8.03	124.72	119.90
34	BA	771	A	O4'-C1'-N9	8.03	114.62	108.20
34	BA	948	C	O4'-C1'-N1	8.03	114.62	108.20
34	BA	1460	U	O4'-C1'-N1	8.03	114.62	108.20
35	BB	503	G	N1-C6-O6	8.03	124.72	119.90
35	BB	556	U	C3'-C2'-C1'	-8.03	95.08	101.50
85	AA	1034	U	O4'-C1'-N1	8.03	114.62	108.20
85	AA	2187	G	P-O5'-C5'	-8.03	108.06	120.90
35	BB	1026	G	C2-N3-C4	-8.03	107.89	111.90
85	AA	335	G	C3'-C2'-C1'	-8.03	95.08	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	472	G	C6-C5-N7	-8.02	125.59	130.40
37	BD	30	A	N1-C6-N6	8.02	123.42	118.60
85	AA	1185	G	C5'-C4'-C3'	8.02	128.84	116.00
34	BA	1165	A	C5'-C4'-C3'	-8.02	103.17	116.00
34	BA	1482	A	C5'-C4'-C3'	8.02	128.84	116.00
34	BA	1503	U	N1-C1'-C2'	-8.02	103.18	112.00
34	BA	1528	U	P-O5'-C5'	8.02	133.74	120.90
34	BA	1561	C	P-O5'-C5'	8.02	133.74	120.90
47	BN	72	MET	CG-SD-CE	-8.02	87.36	100.20
36	BC	15	G	O4'-C1'-N9	8.02	114.62	108.20
45	BL	36	ARG	NE-CZ-NH1	8.02	124.31	120.30
70	Bk	121	ARG	NE-CZ-NH1	8.02	124.31	120.30
85	AA	726	U	P-O3'-C3'	8.02	129.32	119.70
85	AA	1502	A	N1-C6-N6	8.02	123.41	118.60
85	AA	2084	U	O4'-C1'-N1	8.02	114.62	108.20
34	BA	289	A	C2-N3-C4	-8.02	106.59	110.60
34	BA	610	A	C6-N1-C2	-8.02	113.79	118.60
34	BA	1281	U	C3'-C2'-C1'	-8.02	95.08	101.50
34	BA	1499	A	O4'-C1'-N9	8.02	114.61	108.20
35	BB	1487	G	N3-C2-N2	8.02	125.51	119.90
40	BG	16	G	C5-C6-O6	8.02	133.41	128.60
40	BG	163	G	P-O5'-C5'	-8.02	108.07	120.90
65	Bf	348	GLU	C-N-CD	-8.02	102.96	120.60
79	Bt	38	ARG	NE-CZ-NH2	-8.02	116.29	120.30
85	AA	1623	U	P-O5'-C5'	-8.02	108.07	120.90
85	AA	1757	C	C6-N1-C2	-8.02	117.09	120.30
5	A4	68	TYR	CB-CG-CD1	-8.02	116.19	121.00
34	BA	482	C	C4'-C3'-C2'	8.02	110.62	102.60
34	BA	560	U	OP1-P-OP2	-8.02	107.58	119.60
34	BA	621	G	O3'-P-O5'	8.02	119.23	104.00
34	BA	873	G	O5'-P-OP2	8.02	120.32	110.70
34	BA	23	A	C1'-O4'-C4'	-8.02	103.49	109.90
34	BA	198	U	P-O3'-C3'	-8.02	110.08	119.70
34	BA	765	U	O5'-C5'-C4'	8.02	126.93	111.70
35	BB	812	G	C4-N9-C1'	-8.02	116.08	126.50
40	BG	133	C	C6-N1-C1'	-8.02	111.18	120.80
85	AA	406	U	P-O5'-C5'	-8.02	108.08	120.90
85	AA	1941	C	O4'-C1'-N1	8.02	114.61	108.20
85	AA	2125	A	C2'-C3'-O3'	8.02	127.14	109.50
34	BA	926	A	N1-C2-N3	-8.01	125.29	129.30
34	BA	1735	G	C8-N9-C4	8.01	109.61	106.40
35	BB	30	A	O4'-C1'-N9	8.01	114.61	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1042	U	C5'-C4'-C3'	8.01	128.82	116.00
39	BF	36	G	C5'-C4'-C3'	-8.01	103.18	116.00
40	BG	50	G	C6-N1-C2	-8.01	120.29	125.10
40	BG	77	U	O4'-C1'-N1	8.01	114.61	108.20
82	Bw	204	TYR	CB-CG-CD2	-8.01	116.19	121.00
85	AA	92	G	C5-C6-O6	-8.01	123.79	128.60
85	AA	1644	G	N7-C8-N9	8.01	117.11	113.10
86	AB	2	C	O4'-C1'-N1	8.01	114.61	108.20
4	A3	85	PHE	CB-CG-CD1	8.01	126.41	120.80
85	AA	738	C	O4'-C1'-C2'	-8.01	97.79	105.80
85	AA	1553	G	O4'-C1'-N9	8.01	114.61	108.20
34	BA	14	G	C2-N3-C4	-8.01	107.89	111.90
34	BA	262	A	P-O3'-C3'	8.01	129.31	119.70
85	AA	1346	C	P-O5'-C5'	8.01	133.72	120.90
34	BA	896	U	N3-C4-C5	-8.01	109.79	114.60
34	BA	1069	U	N3-C2-O2	-8.01	116.59	122.20
34	BA	1500	G	N3-C2-N2	8.01	125.51	119.90
35	BB	642	G	O4'-C1'-N9	8.01	114.61	108.20
41	BH	115	A	P-O5'-C5'	8.01	133.71	120.90
85	AA	1622	G	C5'-C4'-C3'	-8.01	103.19	116.00
85	AA	1701	G	N1-C2-N2	8.01	123.41	116.20
34	BA	628	U	O4'-C1'-N1	8.01	114.61	108.20
34	BA	1149	C	N3-C2-O2	-8.01	116.29	121.90
34	BA	1309	U	P-O3'-C3'	8.01	129.31	119.70
34	BA	1782	C	N1-C1'-C2'	-8.01	103.19	112.00
35	BB	13	A	C3'-C2'-C1'	-8.01	95.09	101.50
40	BG	33	G	C5-C6-O6	-8.01	123.80	128.60
85	AA	77	C	C2-N1-C1'	8.01	127.61	118.80
37	BD	78	C	N1-C2-O2	8.01	123.70	118.90
85	AA	156	G	O4'-C1'-N9	8.01	114.61	108.20
85	AA	168	A	O4'-C1'-N9	8.01	114.61	108.20
85	AA	730	G	C4-N9-C1'	-8.01	116.09	126.50
85	AA	1457	C	C4'-C3'-C2'	-8.01	94.59	102.60
85	AA	1531	G	C5-C6-O6	-8.01	123.80	128.60
38	BE	131	C	O4'-C1'-N1	8.01	114.60	108.20
85	AA	75	U	O4'-C1'-N1	8.01	114.60	108.20
85	AA	1721	A	O4'-C1'-N9	8.01	114.60	108.20
85	AA	1810	C	C5'-C4'-O4'	8.01	118.71	109.10
34	BA	138	C	N3-C2-O2	-8.00	116.30	121.90
34	BA	213	A	N1-C6-N6	-8.00	113.80	118.60
34	BA	516	U	C4'-C3'-C2'	8.00	110.60	102.60
34	BA	850	C	O4'-C1'-N1	8.00	114.60	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1540	C	C6-N1-C2	-8.00	117.10	120.30
40	BG	60	A	O4'-C1'-N9	8.00	114.60	108.20
85	AA	4	C	C1'-O4'-C4'	-8.00	103.50	109.90
34	BA	4	A	C4'-C3'-C2'	8.00	110.60	102.60
34	BA	767	U	P-O3'-C3'	-8.00	110.10	119.70
56	BW	73	ARG	NE-CZ-NH1	8.00	124.30	120.30
77	Br	288	THR	N-CA-CB	-8.00	95.10	110.30
85	AA	182	C	C2-N1-C1'	8.00	127.60	118.80
85	AA	321	C	C6-N1-C2	-8.00	117.10	120.30
85	AA	1657	C	O4'-C1'-N1	8.00	114.60	108.20
34	BA	1637	G	O4'-C1'-N9	8.00	114.60	108.20
38	BE	95	G	C5-C6-O6	8.00	133.40	128.60
40	BG	148	C	C5'-C4'-C3'	-8.00	103.20	116.00
41	BH	120	C	O4'-C1'-N1	8.00	114.60	108.20
85	AA	485	A	C8-N9-C4	8.00	109.00	105.80
37	BD	70	C	C2-N1-C1'	-8.00	110.00	118.80
38	BE	2	G	C3'-C2'-C1'	-8.00	95.10	101.50
38	BE	64	A	N1-C6-N6	8.00	123.40	118.60
38	BE	141	A	N1-C6-N6	8.00	123.40	118.60
49	BP	48	ARG	NE-CZ-NH1	8.00	124.30	120.30
85	AA	926	C	C2-N3-C4	-8.00	115.90	119.90
85	AA	1048	C	O4'-C1'-N1	8.00	114.60	108.20
85	AA	1537	A	C6-N1-C2	-8.00	113.80	118.60
35	BB	866	A	C4'-C3'-C2'	-8.00	94.61	102.60
35	BB	885	U	O4'-C1'-N1	8.00	114.60	108.20
16	AH	34	PHE	CB-CG-CD2	-7.99	115.20	120.80
34	BA	596	G	N1-C2-N3	-7.99	119.10	123.90
34	BA	1379	G	C2-N3-C4	7.99	115.90	111.90
35	BB	781	U	C3'-C2'-C1'	-7.99	95.11	101.50
85	AA	46	U	C6-N1-C1'	-7.99	110.01	121.20
85	AA	478	U	C6-N1-C2	-7.99	116.20	121.00
85	AA	2008	G	C4-N9-C1'	-7.99	116.11	126.50
49	BP	174	ARG	NE-CZ-NH1	7.99	124.30	120.30
34	BA	103	G	C8-N9-C1'	7.99	137.39	127.00
34	BA	1707	C	C2-N1-C1'	-7.99	110.01	118.80
41	BH	122	U	O4'-C1'-N1	7.99	114.59	108.20
85	AA	1808	G	C4-N9-C1'	7.99	136.89	126.50
34	BA	214	A	N3-C4-C5	-7.99	121.21	126.80
34	BA	481	A	C5'-C4'-C3'	-7.99	103.22	116.00
85	AA	427	G	N3-C2-N2	7.99	125.49	119.90
85	AA	1022	G	C2'-C3'-O3'	7.99	127.08	109.50
85	AA	1926	A	C5'-C4'-C3'	-7.99	103.22	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2130	G	C8-N9-C1'	7.99	137.38	127.00
34	BA	1798	G	C8-N9-C1'	-7.99	116.62	127.00
35	BB	902	C	O4'-C1'-N1	7.99	114.59	108.20
85	AA	94	C	O4'-C1'-N1	7.99	114.59	108.20
85	AA	1432	C	C2-N1-C1'	7.99	127.59	118.80
7	A6	22	ARG	NE-CZ-NH1	7.99	124.29	120.30
34	BA	99	G	C8-N9-C1'	-7.99	116.62	127.00
34	BA	401	A	C8-N9-C4	7.99	108.99	105.80
34	BA	551	U	O4'-C1'-N1	7.99	114.59	108.20
34	BA	957	A	C5'-C4'-O4'	7.99	118.68	109.10
36	BC	137	C	P-O3'-C3'	-7.99	110.12	119.70
38	BE	90	G	C4-N9-C1'	-7.99	116.12	126.50
38	BE	174	U	P-O3'-C3'	7.99	129.28	119.70
85	AA	1426	G	P-O5'-C5'	7.99	133.68	120.90
34	BA	892	C	C5-C4-N4	7.98	125.79	120.20
37	BD	82	G	C8-N9-C1'	7.98	137.38	127.00
38	BE	33	C	P-O3'-C3'	-7.98	110.12	119.70
41	BH	133	U	P-O5'-C5'	-7.98	108.13	120.90
35	BB	791	A	O4'-C1'-N9	7.98	114.59	108.20
36	BC	158	U	O4'-C4'-C3'	-7.98	96.02	104.00
34	BA	1732	A	C4-C5-C6	7.98	120.99	117.00
35	BB	564	U	P-O5'-C5'	7.98	133.67	120.90
85	AA	1293	U	O4'-C1'-N1	7.98	114.58	108.20
35	BB	375	G	P-O3'-C3'	-7.98	110.13	119.70
65	Bf	61	ARG	NE-CZ-NH1	7.98	124.29	120.30
85	AA	66	U	O4'-C1'-N1	7.98	114.58	108.20
86	AB	71	G	C2'-C3'-O3'	7.98	127.05	109.50
34	BA	320	G	P-O3'-C3'	-7.98	110.13	119.70
35	BB	463	C	C6-N1-C2	-7.98	117.11	120.30
36	BC	151	G	C6-N1-C2	-7.98	120.31	125.10
85	AA	1105	G	C4-N9-C1'	7.98	136.87	126.50
27	AT	17	PHE	CB-CG-CD1	-7.98	115.22	120.80
34	BA	122	U	P-O5'-C5'	-7.98	108.14	120.90
34	BA	317	U	O4'-C1'-N1	7.98	114.58	108.20
34	BA	1019	C	O4'-C1'-N1	7.98	114.58	108.20
52	BS	137	ARG	NE-CZ-NH2	-7.98	116.31	120.30
34	BA	1268	C	P-O3'-C3'	-7.97	110.13	119.70
34	BA	1815	G	C5-C6-N1	7.97	115.49	111.50
35	BB	16	G	C4-N9-C1'	-7.97	116.13	126.50
38	BE	199	A	C6-N1-C2	-7.97	113.81	118.60
40	BG	59	G	C1'-O4'-C4'	-7.97	103.52	109.90
41	BH	85	C	C6-N1-C2	-7.97	117.11	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
49	BP	93	ARG	NE-CZ-NH1	7.97	124.29	120.30
74	Bo	49	ARG	NE-CZ-NH1	7.97	124.29	120.30
85	AA	1655	G	C4'-C3'-C2'	7.97	110.57	102.60
34	BA	231	U	C5'-C4'-C3'	7.97	128.75	116.00
34	BA	405	C	C5'-C4'-C3'	7.97	128.76	116.00
35	BB	2	C	N3-C4-C5	-7.97	118.71	121.90
35	BB	1129	C	C5'-C4'-C3'	-7.97	103.24	116.00
35	BB	1490	G	N9-C1'-C2'	-7.97	103.23	112.00
36	BC	129	C	C6-N1-C1'	7.97	130.37	120.80
37	BD	112	U	C2-N1-C1'	-7.97	108.13	117.70
38	BE	127	G	O4'-C1'-N9	7.97	114.58	108.20
85	AA	133	G	O4'-C1'-N9	7.97	114.58	108.20
85	AA	655	U	C2-N3-C4	-7.97	122.22	127.00
85	AA	1683	U	C2-N3-C4	-7.97	122.22	127.00
34	BA	14	G	C8-N9-C4	7.97	109.59	106.40
34	BA	167	U	N3-C2-O2	-7.97	116.62	122.20
34	BA	296	G	C6-C5-N7	-7.97	125.62	130.40
65	Bf	400	MET	CG-SD-CE	-7.97	87.45	100.20
34	BA	1708	A	C2-N3-C4	7.97	114.58	110.60
34	BA	1711	G	N3-C2-N2	7.97	125.48	119.90
35	BB	822	G	N3-C4-N9	7.97	130.78	126.00
36	BC	16	A	P-O5'-C5'	-7.97	108.15	120.90
85	AA	483	G	P-O5'-C5'	7.97	133.65	120.90
85	AA	991	G	C6-N1-C2	-7.97	120.32	125.10
85	AA	1183	C	C2-N1-C1'	7.97	127.57	118.80
34	BA	1510	C	C1'-O4'-C4'	-7.97	103.53	109.90
35	BB	604	C	C3'-C2'-C1'	-7.97	95.13	101.50
34	BA	8	G	O5'-P-OP2	-7.97	98.53	105.70
35	BB	1509	G	P-O3'-C3'	-7.97	110.14	119.70
85	AA	421	G	P-O3'-C3'	-7.97	110.14	119.70
85	AA	1633	A	P-O3'-C3'	7.97	129.26	119.70
85	AA	1950	G	O4'-C1'-N9	7.97	114.57	108.20
85	AA	2058	C	C2-N1-C1'	-7.97	110.04	118.80
85	AA	2168	C	O4'-C1'-N1	7.97	114.57	108.20
34	BA	1816	G	C5-C6-N1	7.96	115.48	111.50
35	BB	438	G	C4-N9-C1'	-7.96	116.14	126.50
35	BB	878	G	C4-N9-C1'	-7.96	116.15	126.50
35	BB	1268	C	O4'-C1'-N1	7.96	114.57	108.20
37	BD	36	C	C5'-C4'-C3'	-7.96	103.26	116.00
39	BF	57	C	P-O3'-C3'	-7.96	110.14	119.70
85	AA	16	G	C5-C6-N1	7.96	115.48	111.50
85	AA	708	G	C5-C6-O6	-7.96	123.82	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	545	U	C5'-C4'-C3'	-7.96	103.26	116.00
34	BA	615	A	O3'-P-O5'	-7.96	88.87	104.00
40	BG	105	A	C5-C6-N1	7.96	121.68	117.70
85	AA	2143	U	C5'-C4'-C3'	-7.96	103.26	116.00
34	BA	1809	G	P-O3'-C3'	7.96	129.25	119.70
35	BB	1424	G	C4-N9-C1'	-7.96	116.15	126.50
85	AA	2231	G	N1-C6-O6	7.96	124.68	119.90
35	BB	798	A	O4'-C1'-N9	7.96	114.57	108.20
37	BD	92	G	C8-N9-C4	-7.96	103.22	106.40
49	BP	122	TYR	CB-CG-CD1	-7.96	116.22	121.00
85	AA	352	G	C1'-O4'-C4'	-7.96	103.53	109.90
34	BA	1724	G	O4'-C1'-N9	7.96	114.57	108.20
85	AA	917	A	C1'-O4'-C4'	-7.96	103.53	109.90
85	AA	924	A	O5'-P-OP2	-7.96	98.54	105.70
85	AA	928	U	O4'-C1'-N1	7.96	114.57	108.20
85	AA	1720	C	C5'-C4'-O4'	7.96	118.65	109.10
85	AA	2182	A	N1-C6-N6	7.96	123.38	118.60
34	BA	1666	U	C5'-C4'-C3'	-7.96	103.27	116.00
34	BA	1768	G	O4'-C1'-N9	7.96	114.56	108.20
35	BB	839	G	C1'-O4'-C4'	-7.96	103.53	109.90
37	BD	48	G	N1-C6-O6	7.96	124.67	119.90
57	BX	61	ARG	NE-CZ-NH1	7.96	124.28	120.30
79	Bt	103	ASP	N-CA-CB	-7.96	96.28	110.60
85	AA	887	A	C8-N9-C4	7.96	108.98	105.80
85	AA	925	G	O4'-C1'-N9	7.96	114.56	108.20
85	AA	1529	A	P-O3'-C3'	7.96	129.25	119.70
35	BB	648	G	P-O3'-C3'	-7.96	110.15	119.70
34	BA	1297	G	C4-N9-C1'	7.95	136.84	126.50
34	BA	1671	A	N1-C6-N6	-7.95	113.83	118.60
35	BB	16	G	C8-N9-C1'	7.95	137.34	127.00
35	BB	1035	C	C6-N1-C2	-7.95	117.12	120.30
38	BE	28	C	P-O3'-C3'	-7.95	110.16	119.70
41	BH	2	U	O4'-C1'-N1	7.95	114.56	108.20
59	BZ	80	ASP	N-CA-CB	-7.95	96.28	110.60
85	AA	111	A	N1-C6-N6	-7.95	113.83	118.60
85	AA	704	A	C4-C5-C6	-7.95	113.02	117.00
85	AA	2056	C	C3'-C2'-C1'	-7.95	95.14	101.50
86	AB	16	U	C2-N3-C4	7.95	131.77	127.00
41	BH	133	U	N3-C2-O2	-7.95	116.63	122.20
85	AA	295	U	O4'-C1'-N1	7.95	114.56	108.20
85	AA	1082	U	O4'-C1'-N1	7.95	114.56	108.20
85	AA	1701	G	C6-N1-C2	-7.95	120.33	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2058	C	N3-C2-O2	-7.95	116.33	121.90
24	AQ	48	ARG	NE-CZ-NH1	7.95	124.28	120.30
36	BC	101	U	O4'-C1'-N1	7.95	114.56	108.20
34	BA	431	A	P-O5'-C5'	-7.95	108.18	120.90
34	BA	758	G	O4'-C1'-N9	7.95	114.56	108.20
34	BA	904	G	N9-C1'-C2'	-7.95	103.26	112.00
35	BB	878	G	O4'-C1'-N9	7.95	114.56	108.20
35	BB	1025	A	O5'-C5'-C4'	7.95	126.80	111.70
37	BD	2	G	C4-N9-C1'	-7.95	116.17	126.50
38	BE	208	G	N9-C1'-C2'	-7.95	103.26	112.00
39	BF	29	U	O4'-C1'-N1	7.95	114.56	108.20
41	BH	40	C	P-O3'-C3'	-7.95	110.16	119.70
85	AA	57	G	C5'-C4'-C3'	-7.95	103.28	116.00
34	BA	23	A	C5-C6-N1	7.95	121.67	117.70
34	BA	400	A	N1-C6-N6	-7.95	113.83	118.60
35	BB	560	C	C2-N1-C1'	-7.95	110.06	118.80
35	BB	93	A	P-O5'-C5'	7.95	133.61	120.90
35	BB	278	U	O4'-C1'-N1	7.95	114.56	108.20
38	BE	204	U	C6-N1-C2	-7.95	116.23	121.00
85	AA	1954	C	C6-N1-C2	-7.95	117.12	120.30
11	AC	138	ARG	NE-CZ-NH1	7.94	124.27	120.30
34	BA	815	C	C4'-C3'-C2'	7.94	110.54	102.60
35	BB	1416	A	P-O5'-C5'	7.94	133.61	120.90
37	BD	85	C	C6-N1-C2	-7.94	117.12	120.30
34	BA	750	C	O4'-C1'-N1	7.94	114.55	108.20
35	BB	488	G	N7-C8-N9	-7.94	109.13	113.10
36	BC	43	A	P-O3'-C3'	-7.94	110.17	119.70
38	BE	88	G	N1-C6-O6	7.94	124.67	119.90
85	AA	188	G	C8-N9-C1'	7.94	137.32	127.00
85	AA	1986	G	P-O3'-C3'	-7.94	110.17	119.70
34	BA	1730	A	C5'-C4'-C3'	7.94	128.70	116.00
35	BB	373	C	O4'-C1'-N1	7.94	114.55	108.20
35	BB	454	U	C2-N1-C1'	-7.94	108.17	117.70
35	BB	1015	U	N3-C2-O2	-7.94	116.64	122.20
35	BB	1026	G	N3-C4-N9	-7.94	121.23	126.00
35	BB	1196	A	C4'-C3'-C2'	7.94	110.54	102.60
36	BC	35	C	C5'-C4'-C3'	7.94	128.70	116.00
38	BE	96	G	C3'-C2'-C1'	-7.94	95.15	101.50
38	BE	178	G	N9-C1'-C2'	-7.94	103.27	112.00
39	BF	52	A	C8-N9-C4	-7.94	102.62	105.80
85	AA	282	C	C1'-O4'-C4'	-7.94	103.55	109.90
85	AA	1509	A	C8-N9-C4	-7.94	102.62	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2218	G	N1-C6-O6	7.94	124.67	119.90
85	AA	86	G	C5'-C4'-C3'	7.94	128.70	116.00
85	AA	1831	U	C5'-C4'-C3'	-7.94	103.30	116.00
34	BA	247	U	C2-N1-C1'	7.94	127.23	117.70
34	BA	1182	U	P-O3'-C3'	-7.94	110.17	119.70
34	BA	1411	C	C2-N3-C4	-7.94	115.93	119.90
35	BB	610	U	C2-N1-C1'	-7.94	108.18	117.70
36	BC	133	C	P-O5'-C5'	7.94	133.60	120.90
38	BE	63	C	C4'-C3'-C2'	7.94	110.54	102.60
39	BF	35	C	O5'-P-OP1	-7.94	98.56	105.70
85	AA	309	G	N1-C6-O6	7.94	124.66	119.90
85	AA	375	C	N3-C2-O2	-7.94	116.34	121.90
85	AA	2153	G	C5'-C4'-C3'	-7.94	103.30	116.00
35	BB	1206	G	C5'-C4'-C3'	-7.94	103.30	116.00
40	BG	78	C	C5'-C4'-O4'	7.94	118.62	109.10
65	Bf	377	ASN	N-CA-CB	7.94	124.89	110.60
85	AA	820	G	C4-N9-C1'	-7.94	116.18	126.50
34	BA	567	U	C3'-C2'-C1'	-7.93	95.15	101.50
34	BA	585	G	N1-C6-O6	7.93	124.66	119.90
34	BA	684	G	N3-C2-N2	7.93	125.45	119.90
34	BA	1735	G	P-O5'-C5'	-7.93	108.21	120.90
35	BB	872	A	C3'-C2'-C1'	-7.93	95.15	101.50
85	AA	553	G	C5'-C4'-C3'	-7.93	103.31	116.00
85	AA	1204	A	C8-N9-C4	-7.93	102.63	105.80
34	BA	135	G	C5'-C4'-C3'	-7.93	103.31	116.00
34	BA	571	G	C8-N9-C1'	-7.93	116.69	127.00
35	BB	622	G	P-O3'-C3'	-7.93	110.18	119.70
39	BF	32	G	C2'-C3'-O3'	7.93	126.95	109.50
85	AA	1254	A	P-O3'-C3'	-7.93	110.18	119.70
85	AA	2069	A	C1'-O4'-C4'	-7.93	103.55	109.90
85	AA	2130	G	C4-N9-C1'	-7.93	116.19	126.50
34	BA	1408	C	O4'-C1'-N1	7.93	114.55	108.20
37	BD	51	G	O5'-C5'-C4'	-7.93	96.63	111.70
85	AA	842	G	P-O5'-C5'	-7.93	108.21	120.90
34	BA	1669	C	P-O3'-C3'	-7.93	110.18	119.70
35	BB	1506	C	C5'-C4'-C3'	7.93	128.69	116.00
85	AA	128	U	C2-N1-C1'	7.93	127.22	117.70
85	AA	714	U	P-O3'-C3'	7.93	129.22	119.70
85	AA	924	A	O3'-P-O5'	7.93	119.07	104.00
35	BB	814	A	C8-N9-C1'	7.93	141.97	127.70
34	BA	935	A	N1-C6-N6	7.93	123.36	118.60
34	BA	1694	C	C3'-C2'-C1'	-7.93	95.16	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	788	U	C6-N1-C1'	7.93	132.30	121.20
35	BB	869	G	C4-N9-C1'	-7.93	116.20	126.50
82	Bw	140	ASN	CB-CA-C	-7.93	94.55	110.40
85	AA	179	G	C4'-C3'-C2'	-7.93	94.67	102.60
85	AA	2142	A	C8-N9-C4	7.93	108.97	105.80
34	BA	1665	G	C8-N9-C1'	7.92	137.30	127.00
35	BB	378	C	C1'-O4'-C4'	-7.92	103.56	109.90
35	BB	438	G	C8-N9-C1'	7.92	137.30	127.00
35	BB	979	G	N1-C6-O6	7.92	124.65	119.90
37	BD	79	G	C4-N9-C1'	-7.92	116.20	126.50
39	BF	67	A	C5-C6-N6	-7.92	117.36	123.70
85	AA	1588	A	P-O5'-C5'	7.92	133.58	120.90
35	BB	57	G	C4-N9-C1'	-7.92	116.20	126.50
41	BH	132	C	O3'-P-O5'	-7.92	88.95	104.00
77	Br	298	ARG	NE-CZ-NH1	7.92	124.26	120.30
34	BA	501	U	P-O3'-C3'	7.92	129.21	119.70
34	BA	608	G	C5'-C4'-C3'	7.92	128.67	116.00
39	BF	53	G	P-O3'-C3'	7.92	129.21	119.70
85	AA	1176	C	C6-N1-C1'	7.92	130.31	120.80
34	BA	242	U	O4'-C1'-N1	7.92	114.54	108.20
35	BB	645	C	C6-N1-C2	-7.92	117.13	120.30
35	BB	1195	A	O4'-C1'-N9	7.92	114.54	108.20
34	BA	325	A	P-O3'-C3'	7.92	129.20	119.70
34	BA	875	G	C2'-C3'-O3'	7.92	126.92	109.50
34	BA	1746	G	C8-N9-C1'	7.92	137.29	127.00
35	BB	26	C	P-O3'-C3'	-7.92	110.20	119.70
35	BB	540	G	C5-C6-O6	-7.92	123.85	128.60
35	BB	1523	U	P-O3'-C3'	-7.92	110.20	119.70
37	BD	87	G	C6-N1-C2	-7.92	120.35	125.10
34	BA	94	G	N1-C6-O6	-7.92	115.15	119.90
35	BB	472	C	N3-C2-O2	-7.92	116.36	121.90
39	BF	23	G	O3'-P-O5'	7.92	119.04	104.00
34	BA	1440	C	P-O5'-C5'	-7.92	108.24	120.90
35	BB	580	A	P-O5'-C5'	7.92	133.56	120.90
38	BE	104	G	N3-C4-C5	-7.92	124.64	128.60
85	AA	351	C	O3'-P-O5'	7.92	119.04	104.00
21	AM	114	ARG	NE-CZ-NH2	-7.91	116.34	120.30
34	BA	115	U	N1-C1'-C2'	7.91	124.29	114.00
35	BB	1362	G	C5'-C4'-C3'	7.91	128.66	116.00
37	BD	73	U	C4'-C3'-C2'	7.91	110.51	102.60
77	Br	349	ARG	NE-CZ-NH1	7.91	124.26	120.30
85	AA	16	G	C4-N9-C1'	-7.91	116.21	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	17	C	O4'-C1'-N1	7.91	114.53	108.20
85	AA	926	C	P-O5'-C5'	-7.91	108.24	120.90
85	AA	1178	A	P-O3'-C3'	7.91	129.20	119.70
35	BB	1372	G	C5'-C4'-C3'	-7.91	103.34	116.00
85	AA	1252	A	C1'-O4'-C4'	-7.91	103.57	109.90
35	BB	384	A	P-O5'-C5'	-7.91	108.24	120.90
62	Bc	119	ARG	NE-CZ-NH1	7.91	124.25	120.30
85	AA	753	U	C5'-C4'-O4'	7.91	118.59	109.10
3	A2	149	PHE	CB-CG-CD1	7.91	126.34	120.80
34	BA	574	U	O4'-C4'-C3'	-7.91	96.09	104.00
34	BA	1223	C	C4'-C3'-C2'	7.91	110.51	102.60
35	BB	1130	U	N3-C2-O2	-7.91	116.67	122.20
36	BC	23	G	N3-C2-N2	7.91	125.44	119.90
40	BG	71	C	C5-C4-N4	-7.91	114.66	120.20
85	AA	40	A	N1-C6-N6	7.91	123.34	118.60
34	BA	796	G	N7-C8-N9	7.91	117.05	113.10
34	BA	889	U	C2-N3-C4	-7.91	122.26	127.00
34	BA	563	A	O4'-C1'-N9	7.91	114.52	108.20
34	BA	684	G	O5'-P-OP1	-7.91	98.58	105.70
35	BB	532	C	C6-N1-C2	-7.91	117.14	120.30
35	BB	1168	G	C6-N1-C2	-7.91	120.36	125.10
35	BB	1313	C	C6-N1-C2	-7.91	117.14	120.30
5	A4	91	PHE	CB-CG-CD1	-7.90	115.27	120.80
85	AA	413	G	O4'-C1'-N9	7.90	114.52	108.20
85	AA	2248	A	N1-C6-N6	-7.90	113.86	118.60
34	BA	377	G	P-O5'-C5'	-7.90	108.26	120.90
65	Bf	143	TYR	N-CA-CB	7.90	124.83	110.60
85	AA	821	U	N3-C2-O2	-7.90	116.67	122.20
7	A6	25	ARG	NE-CZ-NH2	-7.90	116.35	120.30
34	BA	1558	C	O4'-C1'-N1	7.90	114.52	108.20
35	BB	1127	A	P-O3'-C3'	-7.90	110.22	119.70
35	BB	1510	G	C4'-C3'-C2'	-7.90	94.70	102.60
36	BC	23	G	C5-C6-N1	7.90	115.45	111.50
36	BC	89	U	P-O5'-C5'	-7.90	108.26	120.90
36	BC	124	A	C8-N9-C4	-7.90	102.64	105.80
38	BE	70	C	C5'-C4'-C3'	-7.90	103.36	116.00
19	AK	88	ARG	NE-CZ-NH2	-7.90	116.35	120.30
34	BA	1555	G	C4-N9-C1'	-7.90	116.23	126.50
35	BB	1084	A	C4'-C3'-C2'	-7.90	94.70	102.60
35	BB	1542	C	O4'-C1'-N1	7.90	114.52	108.20
85	AA	1119	A	C1'-O4'-C4'	-7.90	103.58	109.90
34	BA	86	A	P-O5'-C5'	7.90	133.54	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	166	G	C8-N9-C4	7.90	109.56	106.40
35	BB	802	G	P-O3'-C3'	-7.90	110.22	119.70
35	BB	1100	C	P-O3'-C3'	-7.90	110.22	119.70
85	AA	272	C	O4'-C1'-N1	7.90	114.52	108.20
85	AA	983	A	C5'-C4'-C3'	-7.90	103.36	116.00
34	BA	559	C	O4'-C1'-N1	7.90	114.52	108.20
34	BA	557	U	OP1-P-OP2	-7.89	107.76	119.60
35	BB	528	G	O4'-C1'-N9	7.89	114.52	108.20
35	BB	1099	U	O4'-C1'-N1	7.89	114.52	108.20
39	BF	8	C	C6-N1-C2	-7.89	117.14	120.30
85	AA	1754	G	P-O5'-C5'	7.89	133.53	120.90
34	BA	211	C	N3-C2-O2	-7.89	116.38	121.90
34	BA	249	A	C8-N9-C4	-7.89	102.64	105.80
34	BA	363	G	N1-C6-O6	7.89	124.64	119.90
35	BB	711	C	C2-N1-C1'	-7.89	110.12	118.80
40	BG	94	G	P-O5'-C5'	7.89	133.53	120.90
41	BH	47	G	C6-N1-C2	-7.89	120.36	125.10
85	AA	643	C	C6-N1-C1'	7.89	130.27	120.80
85	AA	1512	U	C2-N3-C4	-7.89	122.26	127.00
34	BA	1437	G	C5-C6-N1	7.89	115.45	111.50
35	BB	488	G	C8-N9-C4	7.89	109.56	106.40
35	BB	604	C	C6-N1-C2	-7.89	117.14	120.30
5	A4	10	LEU	N-CA-CB	-7.89	94.62	110.40
34	BA	1594	G	N1-C6-O6	7.89	124.63	119.90
35	BB	738	G	C5-C6-O6	-7.89	123.87	128.60
35	BB	870	C	C5'-C4'-C3'	-7.89	103.38	116.00
37	BD	97	U	C2-N1-C1'	-7.89	108.23	117.70
71	Bl	114	ARG	NE-CZ-NH1	7.89	124.24	120.30
85	AA	124	A	P-O3'-C3'	7.89	129.17	119.70
85	AA	1455	C	O4'-C1'-N1	7.89	114.51	108.20
85	AA	1500	C	C6-N1-C2	-7.89	117.14	120.30
34	BA	55	G	C5-C6-N1	7.89	115.44	111.50
34	BA	127	U	C2-N1-C1'	-7.89	108.23	117.70
34	BA	1565	U	P-O3'-C3'	-7.89	110.23	119.70
34	BA	1802	C	C5'-C4'-C3'	7.89	128.62	116.00
34	BA	1515	U	C5'-C4'-O4'	7.89	118.56	109.10
85	AA	533	C	C5-C4-N4	-7.89	114.68	120.20
85	AA	825	U	O4'-C1'-N1	7.89	114.51	108.20
34	BA	504	A	C5-C6-N6	7.88	130.01	123.70
34	BA	1814	U	O4'-C1'-N1	7.88	114.51	108.20
35	BB	367	C	C5'-C4'-C3'	-7.88	103.38	116.00
36	BC	54	G	C6-N1-C2	-7.88	120.37	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	46	G	C8-N9-C1'	7.88	137.25	127.00
40	BG	8	U	C2-N1-C1'	-7.88	108.24	117.70
85	AA	1226	A	C3'-C2'-C1'	-7.88	95.19	101.50
85	AA	1670	U	P-O3'-C3'	-7.88	110.24	119.70
34	BA	1397	C	C6-N1-C2	-7.88	117.15	120.30
35	BB	964	G	N9-C1'-C2'	-7.88	103.33	112.00
39	BF	4	A	C8-N9-C4	7.88	108.95	105.80
85	AA	1190	G	P-O5'-C5'	-7.88	108.29	120.90
4	A3	162	ARG	NE-CZ-NH1	7.88	124.24	120.30
5	A4	136	TYR	CB-CG-CD1	-7.88	116.27	121.00
34	BA	519	G	C1'-O4'-C4'	-7.88	103.59	109.90
34	BA	661	C	C5'-C4'-C3'	7.88	128.61	116.00
34	BA	1601	C	C4'-C3'-C2'	7.88	110.48	102.60
35	BB	852	G	C5'-C4'-C3'	-7.88	103.39	116.00
35	BB	1191	G	C4-N9-C1'	-7.88	116.25	126.50
38	BE	134	A	C4-C5-C6	-7.88	113.06	117.00
40	BG	126	G	C5-C6-N1	7.88	115.44	111.50
56	BW	88	ARG	NE-CZ-NH2	-7.88	116.36	120.30
85	AA	2204	A	C5'-C4'-C3'	-7.88	103.39	116.00
34	BA	1707	C	P-O3'-C3'	-7.88	110.24	119.70
35	BB	814	A	C4-N9-C1'	-7.88	112.11	126.30
34	BA	243	C	C2-N1-C1'	-7.88	110.13	118.80
34	BA	677	U	C5'-C4'-C3'	-7.88	103.39	116.00
35	BB	1524	G	C4-N9-C1'	-7.88	116.26	126.50
38	BE	173	G	O3'-P-O5'	7.88	118.97	104.00
40	BG	87	G	O3'-P-O5'	-7.88	89.03	104.00
85	AA	737	G	C8-N9-C1'	-7.88	116.76	127.00
85	AA	1525	C	C2-N1-C1'	-7.88	110.13	118.80
85	AA	1690	A	P-O3'-C3'	7.88	129.16	119.70
85	AA	2123	U	N3-C2-O2	-7.88	116.69	122.20
34	BA	559	C	C5-C6-N1	7.88	124.94	121.00
34	BA	895	U	O4'-C1'-N1	7.88	114.50	108.20
34	BA	1186	U	O4'-C1'-N1	7.88	114.50	108.20
34	BA	1343	A	N1-C6-N6	-7.88	113.87	118.60
35	BB	1413	U	O4'-C1'-N1	7.88	114.50	108.20
41	BH	48	G	C8-N9-C4	-7.88	103.25	106.40
41	BH	86	G	C5-C6-O6	-7.88	123.87	128.60
85	AA	330	C	C5-C6-N1	7.88	124.94	121.00
85	AA	959	C	P-O5'-C5'	-7.88	108.30	120.90
85	AA	1275	A	C5'-C4'-O4'	7.88	118.55	109.10
85	AA	2217	A	P-O5'-C5'	-7.88	108.30	120.90
34	BA	836	U	O4'-C1'-N1	7.88	114.50	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1369	C	O4'-C1'-N1	7.88	114.50	108.20
34	BA	485	C	C5'-C4'-C3'	7.87	128.60	116.00
34	BA	1469	G	C5'-C4'-C3'	7.87	128.60	116.00
34	BA	1567	G	O4'-C1'-N9	7.87	114.50	108.20
34	BA	1639	U	C2-N1-C1'	-7.87	108.25	117.70
35	BB	418	G	C1'-O4'-C4'	-7.87	103.60	109.90
36	BC	39	G	N1-C6-O6	7.87	124.62	119.90
39	BF	46	G	C3'-C2'-C1'	-7.87	95.20	101.50
40	BG	52	A	C5'-C4'-C3'	-7.87	103.40	116.00
41	BH	39	G	C5-C6-N1	7.87	115.44	111.50
85	AA	974	U	O5'-C5'-C4'	7.87	126.66	111.70
34	BA	493	G	P-O5'-C5'	-7.87	108.31	120.90
37	BD	97	U	C4'-C3'-C2'	7.87	110.47	102.60
85	AA	281	C	O4'-C1'-N1	7.87	114.50	108.20
35	BB	92	C	O4'-C1'-N1	7.87	114.50	108.20
34	BA	372	U	C3'-C2'-C1'	-7.87	95.20	101.50
35	BB	493	U	O4'-C1'-N1	7.87	114.49	108.20
81	Bv	42	ARG	NE-CZ-NH2	-7.87	116.36	120.30
85	AA	353	G	C4-N9-C1'	-7.87	116.27	126.50
37	BD	52	U	P-O3'-C3'	-7.87	110.26	119.70
38	BE	194	A	P-O5'-C5'	7.87	133.49	120.90
40	BG	152	G	N3-C2-N2	7.87	125.41	119.90
57	BX	55	ARG	NE-CZ-NH1	7.87	124.23	120.30
34	BA	1577	U	C2-N1-C1'	-7.87	108.26	117.70
35	BB	599	U	C2-N3-C4	-7.87	122.28	127.00
40	BG	125	C	O4'-C1'-N1	7.87	114.49	108.20
41	BH	86	G	C2-N3-C4	7.87	115.83	111.90
85	AA	507	C	P-O3'-C3'	-7.87	110.26	119.70
85	AA	1291	A	C3'-C2'-C1'	-7.87	95.21	101.50
34	BA	283	U	C6-N1-C2	-7.86	116.28	121.00
34	BA	1508	C	O3'-P-O5'	7.86	118.94	104.00
35	BB	590	G	O4'-C1'-N9	7.86	114.49	108.20
35	BB	704	G	O5'-C5'-C4'	-7.86	96.76	111.70
35	BB	1441	C	C2-N3-C4	-7.86	115.97	119.90
38	BE	70	C	C2-N1-C1'	-7.86	110.15	118.80
85	AA	768	C	O4'-C1'-N1	7.86	114.49	108.20
85	AA	792	A	P-O3'-C3'	7.86	129.14	119.70
15	AG	20	ARG	NE-CZ-NH1	7.86	124.23	120.30
35	BB	1201	G	C1'-O4'-C4'	-7.86	103.61	109.90
85	AA	818	C	C6-N1-C2	-7.86	117.16	120.30
85	AA	316	C	C6-N1-C2	-7.86	117.16	120.30
85	AA	321	C	O4'-C1'-N1	7.86	114.49	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	116	C	N3-C2-O2	-7.86	116.40	121.90
39	BF	49	C	C1'-O4'-C4'	-7.86	103.61	109.90
85	AA	1000	U	C6-N1-C2	-7.86	116.28	121.00
85	AA	1252	A	P-O5'-C5'	-7.86	108.33	120.90
85	AA	1509	A	C5'-C4'-C3'	7.86	128.57	116.00
35	BB	805	G	C5-C6-N1	-7.86	107.57	111.50
35	BB	1314	G	C1'-O4'-C4'	-7.86	103.61	109.90
37	BD	65	G	C8-N9-C1'	7.86	137.22	127.00
38	BE	20	C	O4'-C4'-C3'	-7.86	96.14	104.00
38	BE	132	U	C6-N1-C2	-7.86	116.29	121.00
38	BE	147	G	N1-C6-O6	7.86	124.61	119.90
34	BA	525	A	P-O3'-C3'	7.86	129.13	119.70
34	BA	1307	U	O4'-C1'-N1	7.86	114.48	108.20
34	BA	1767	G	O4'-C1'-N9	7.86	114.48	108.20
57	BX	55	ARG	NE-CZ-NH2	-7.86	116.37	120.30
85	AA	766	G	P-O5'-C5'	7.86	133.47	120.90
85	AA	1548	A	P-O3'-C3'	7.86	129.13	119.70
34	BA	1379	G	O3'-P-O5'	-7.85	89.08	104.00
38	BE	204	U	P-O5'-C5'	-7.85	108.33	120.90
40	BG	115	C	O4'-C1'-N1	7.85	114.48	108.20
41	BH	26	C	O5'-P-OP2	7.85	120.12	110.70
34	BA	932	G	C4'-C3'-C2'	-7.85	94.75	102.60
34	BA	1591	G	N7-C8-N9	-7.85	109.17	113.10
37	BD	106	G	C5-C6-N1	7.85	115.43	111.50
85	AA	266	U	C6-N1-C1'	-7.85	110.21	121.20
85	AA	469	G	N1-C2-N2	-7.85	109.13	116.20
85	AA	686	U	C5'-C4'-C3'	-7.85	103.44	116.00
85	AA	737	G	C5'-C4'-C3'	-7.85	103.44	116.00
85	AA	1720	C	O4'-C4'-C3'	7.85	112.38	106.10
38	BE	166	G	C5-C6-O6	-7.85	123.89	128.60
85	AA	551	C	C1'-O4'-C4'	-7.85	103.62	109.90
18	AJ	111	MET	CG-SD-CE	-7.85	87.64	100.20
19	AK	129	ARG	NE-CZ-NH1	7.85	124.22	120.30
34	BA	166	G	OP2-P-O3'	7.85	122.47	105.20
34	BA	1722	U	O4'-C1'-N1	7.85	114.48	108.20
35	BB	704	G	N3-C2-N2	7.85	125.39	119.90
35	BB	1426	G	C5'-C4'-C3'	-7.85	103.44	116.00
40	BG	16	G	C2-N3-C4	7.85	115.83	111.90
85	AA	527	A	P-O5'-C5'	7.85	133.46	120.90
85	AA	690	G	N1-C6-O6	-7.85	115.19	119.90
85	AA	1536	C	P-O3'-C3'	-7.85	110.28	119.70
34	BA	306	G	C1'-O4'-C4'	-7.85	103.62	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	829	U	C3'-C2'-C1'	-7.85	95.22	101.50
71	Bl	37	ARG	NE-CZ-NH1	7.85	124.22	120.30
85	AA	636	G	C5-C6-O6	-7.85	123.89	128.60
34	BA	13	U	C2'-C3'-O3'	7.85	126.76	109.50
85	AA	802	A	C5'-C4'-C3'	7.85	128.55	116.00
85	AA	1162	A	C1'-O4'-C4'	-7.85	103.62	109.90
85	AA	2171	A	O3'-P-O5'	-7.85	89.09	104.00
34	BA	86	A	C5-C6-N6	-7.84	117.42	123.70
35	BB	805	G	O5'-C5'-C4'	-7.84	96.80	111.70
85	AA	422	G	C4-N9-C1'	-7.84	116.30	126.50
85	AA	1457	C	O4'-C1'-N1	7.84	114.48	108.20
85	AA	1687	U	C1'-O4'-C4'	-7.84	103.62	109.90
85	AA	21	U	C4'-C3'-C2'	7.84	110.44	102.60
85	AA	302	C	O5'-C5'-C4'	7.84	126.60	111.70
85	AA	365	G	N1-C6-O6	-7.84	115.19	119.90
85	AA	1827	U	P-O3'-C3'	-7.84	110.29	119.70
34	BA	367	G	C4-N9-C1'	7.84	136.69	126.50
34	BA	772	G	C8-N9-C4	-7.84	103.26	106.40
34	BA	1746	G	C4-N9-C1'	-7.84	116.31	126.50
35	BB	751	A	O4'-C1'-N9	7.84	114.47	108.20
85	AA	24	U	O5'-C5'-C4'	-7.84	96.80	111.70
85	AA	822	U	C3'-C2'-C1'	-7.84	95.23	101.50
85	AA	980	U	C5'-C4'-C3'	-7.84	103.45	116.00
85	AA	1449	C	O4'-C1'-N1	7.84	114.47	108.20
34	BA	568	G	C1'-O4'-C4'	-7.84	103.63	109.90
34	BA	832	C	O4'-C1'-N1	7.84	114.47	108.20
34	BA	851	C	O4'-C1'-N1	7.84	114.47	108.20
34	BA	1482	A	C8-N9-C4	7.84	108.94	105.80
34	BA	1829	A	C4'-C3'-C2'	-7.84	94.76	102.60
38	BE	154	A	P-O3'-C3'	-7.84	110.29	119.70
85	AA	145	C	P-O5'-C5'	7.84	133.44	120.90
85	AA	469	G	C5-N7-C8	7.84	108.22	104.30
85	AA	611	G	P-O5'-C5'	-7.84	108.36	120.90
85	AA	1146	C	C5'-C4'-C3'	-7.84	103.46	116.00
85	AA	1799	C	C5'-C4'-C3'	-7.84	103.46	116.00
85	AA	1976	G	C8-N9-C1'	7.84	137.19	127.00
35	BB	505	G	N3-C4-C5	-7.84	124.68	128.60
85	AA	115	U	C5'-C4'-C3'	7.84	128.54	116.00
85	AA	1172	A	N1-C6-N6	7.84	123.30	118.60
34	BA	573	U	C4'-C3'-C2'	-7.84	94.76	102.60
34	BA	1792	U	N3-C2-O2	-7.84	116.71	122.20
35	BB	1536	G	C5-C6-O6	-7.84	123.90	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	35	U	C6-N1-C2	-7.84	116.30	121.00
85	AA	113	U	C3'-C2'-C1'	-7.84	95.23	101.50
85	AA	1271	U	O4'-C1'-N1	7.84	114.47	108.20
85	AA	1536	C	O5'-C5'-C4'	7.84	126.59	111.70
34	BA	72	U	P-O5'-C5'	7.83	133.44	120.90
34	BA	379	C	C6-N1-C2	-7.83	117.17	120.30
34	BA	1110	A	N1-C6-N6	-7.83	113.90	118.60
34	BA	157	U	C5'-C4'-C3'	7.83	128.53	116.00
34	BA	945	A	P-O3'-C3'	7.83	129.10	119.70
34	BA	1288	U	P-O5'-C5'	-7.83	108.37	120.90
34	BA	1395	C	O4'-C1'-N1	7.83	114.47	108.20
35	BB	385	C	O4'-C1'-N1	7.83	114.47	108.20
40	BG	81	G	C5-C6-O6	-7.83	123.90	128.60
60	Ba	93	ARG	NE-CZ-NH1	7.83	124.22	120.30
85	AA	289	G	O4'-C1'-N9	7.83	114.47	108.20
85	AA	427	G	P-O5'-C5'	7.83	133.43	120.90
85	AA	750	A	C4-N9-C1'	-7.83	112.20	126.30
85	AA	1204	A	C5-C6-N6	7.83	129.97	123.70
85	AA	1468	G	P-O5'-C5'	7.83	133.43	120.90
85	AA	1495	G	C5-C6-N1	7.83	115.42	111.50
85	AA	2248	A	O4'-C1'-N9	7.83	114.47	108.20
16	AH	9	TYR	CB-CG-CD2	-7.83	116.30	121.00
34	BA	437	G	C5-C6-O6	-7.83	123.90	128.60
35	BB	68	G	O3'-P-O5'	7.83	118.88	104.00
35	BB	971	A	N1-C6-N6	7.83	123.30	118.60
38	BE	117	A	P-O3'-C3'	-7.83	110.30	119.70
85	AA	1140	G	P-O5'-C5'	-7.83	108.37	120.90
85	AA	2173	A	P-O5'-C5'	7.83	133.43	120.90
34	BA	241	U	C5'-C4'-C3'	-7.83	103.47	116.00
34	BA	937	G	N1-C2-N2	-7.83	109.15	116.20
35	BB	1137	G	C5'-C4'-O4'	7.83	118.50	109.10
34	BA	1505	G	C2-N3-C4	-7.83	107.98	111.90
35	BB	272	C	O4'-C1'-N1	7.83	114.46	108.20
35	BB	533	U	C1'-O4'-C4'	-7.83	103.64	109.90
85	AA	383	C	O4'-C1'-N1	7.83	114.46	108.20
85	AA	1702	G	O5'-C5'-C4'	-7.83	96.83	111.70
85	AA	1864	G	C5'-C4'-C3'	-7.83	103.47	116.00
34	BA	1725	U	C2'-C3'-O3'	7.83	126.72	109.50
35	BB	871	C	C2-N1-C1'	-7.83	110.19	118.80
35	BB	956	G	P-O5'-C5'	7.83	133.42	120.90
34	BA	497	U	N3-C2-O2	-7.83	116.72	122.20
34	BA	632	U	O4'-C1'-N1	7.83	114.46	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	763	U	O4'-C1'-C2'	-7.83	97.97	105.80
38	BE	43	A	P-O3'-C3'	-7.83	110.31	119.70
85	AA	64	A	C4'-C3'-C2'	-7.83	94.78	102.60
85	AA	964	C	O3'-P-O5'	-7.83	89.13	104.00
85	AA	1594	G	P-O3'-C3'	-7.83	110.31	119.70
1	A0	177	ARG	NE-CZ-NH1	7.82	124.21	120.30
34	BA	316	G	C5-C6-O6	-7.82	123.91	128.60
35	BB	963	G	C3'-C2'-C1'	-7.82	95.24	101.50
40	BG	112	C	C2-N1-C1'	-7.82	110.19	118.80
85	AA	330	C	N3-C4-C5	-7.82	118.77	121.90
85	AA	1848	G	C6-C5-N7	-7.82	125.71	130.40
86	AB	48	C	N1-C2-O2	7.82	123.59	118.90
34	BA	487	A	O4'-C1'-C2'	-7.82	97.98	105.80
34	BA	641	U	O4'-C1'-N1	7.82	114.46	108.20
85	AA	2052	U	P-O3'-C3'	-7.82	110.31	119.70
34	BA	6	C	O4'-C1'-N1	7.82	114.46	108.20
34	BA	127	U	C2'-C3'-O3'	7.82	126.71	109.50
34	BA	740	A	O4'-C4'-C3'	-7.82	96.18	104.00
34	BA	1247	G	O4'-C1'-N9	7.82	114.46	108.20
35	BB	1461	C	C3'-C2'-C1'	-7.82	95.24	101.50
35	BB	1500	U	C5'-C4'-C3'	7.82	128.51	116.00
40	BG	55	A	C8-N9-C4	7.82	108.93	105.80
84	By	128	VAL	N-CA-C	-7.82	89.88	111.00
85	AA	118	C	O4'-C1'-N1	7.82	114.46	108.20
85	AA	237	G	O4'-C1'-N9	7.82	114.46	108.20
86	AB	62	C	O4'-C1'-N1	7.82	114.46	108.20
35	BB	572	G	N9-C1'-C2'	-7.82	103.40	112.00
34	BA	883	C	C2-N1-C1'	7.82	127.40	118.80
34	BA	1468	U	C2-N1-C1'	-7.82	108.32	117.70
35	BB	838	G	C8-N9-C4	-7.82	103.27	106.40
35	BB	1507	U	P-O3'-C3'	7.82	129.08	119.70
38	BE	105	A	N1-C6-N6	7.82	123.29	118.60
85	AA	611	G	C8-N9-C1'	7.82	137.16	127.00
85	AA	831	C	C4'-C3'-C2'	-7.82	94.78	102.60
85	AA	881	C	C2-N1-C1'	-7.82	110.20	118.80
34	BA	222	C	N1-C2-N3	7.82	124.67	119.20
34	BA	556	A	C2-N3-C4	-7.82	106.69	110.60
34	BA	782	C	N1-C2-O2	-7.82	114.21	118.90
36	BC	88	A	C8-N9-C4	-7.82	102.67	105.80
85	AA	251	A	N1-C6-N6	-7.82	113.91	118.60
34	BA	966	G	N9-C1'-C2'	-7.81	103.41	112.00
34	BA	531	C	O4'-C4'-C3'	-7.81	96.19	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	611	A	O4'-C1'-N9	7.81	114.45	108.20
38	BE	6	A	C5-C6-N6	-7.81	117.45	123.70
34	BA	582	U	N1-C2-O2	7.81	128.27	122.80
34	BA	864	G	OP1-P-OP2	-7.81	107.88	119.60
34	BA	1705	C	C5'-C4'-C3'	7.81	128.50	116.00
35	BB	431	U	N3-C2-O2	-7.81	116.73	122.20
34	BA	1219	G	C5'-C4'-C3'	-7.81	103.50	116.00
35	BB	1284	U	O4'-C1'-N1	7.81	114.45	108.20
39	BF	12	U	O5'-P-OP1	-7.81	98.67	105.70
85	AA	903	G	O4'-C1'-N9	7.81	114.45	108.20
85	AA	2024	U	C1'-O4'-C4'	-7.81	103.65	109.90
34	BA	698	U	C3'-C2'-C1'	-7.81	95.25	101.50
35	BB	587	A	N1-C6-N6	7.81	123.28	118.60
35	BB	1484	A	C8-N9-C4	-7.81	102.68	105.80
38	BE	182	U	O4'-C1'-N1	7.81	114.45	108.20
40	BG	103	C	C6-N1-C2	-7.81	117.18	120.30
85	AA	575	G	C4-N9-C1'	-7.81	116.35	126.50
85	AA	967	C	C6-N1-C1'	7.81	130.17	120.80
85	AA	1179	A	C4'-C3'-C2'	-7.81	94.79	102.60
38	BE	83	U	O5'-P-OP2	7.81	120.07	110.70
5	A4	155	MET	CB-CA-C	7.80	126.01	110.40
19	AK	78	SER	N-CA-CB	7.80	122.21	110.50
34	BA	763	U	C3'-C2'-C1'	7.80	107.74	101.50
34	BA	1206	C	C2-N1-C1'	7.80	127.39	118.80
34	BA	1324	G	C5-C6-N1	7.80	115.40	111.50
36	BC	113	G	P-O3'-C3'	-7.80	110.33	119.70
39	BF	51	C	C5'-C4'-C3'	-7.80	103.51	116.00
74	Bo	56	ARG	N-CA-CB	7.80	124.65	110.60
85	AA	1459	C	C2-N3-C4	-7.80	116.00	119.90
35	BB	141	G	C5-C6-O6	-7.80	123.92	128.60
38	BE	18	U	O3'-P-O5'	7.80	118.83	104.00
45	BL	140	ARG	NE-CZ-NH2	-7.80	116.40	120.30
34	BA	281	C	O5'-C5'-C4'	-7.80	96.88	111.70
34	BA	685	C	C6-N1-C1'	-7.80	111.44	120.80
34	BA	1083	A	N1-C2-N3	-7.80	125.40	129.30
34	BA	1739	G	N1-C6-O6	7.80	124.58	119.90
35	BB	412	A	O4'-C1'-N9	7.80	114.44	108.20
85	AA	162	A	C5'-C4'-C3'	-7.80	103.52	116.00
85	AA	492	C	C5'-C4'-C3'	-7.80	103.52	116.00
85	AA	1213	U	C2-N3-C4	-7.80	122.32	127.00
85	AA	1719	C	O5'-C5'-C4'	7.80	126.52	111.70
34	BA	782	C	P-O3'-C3'	-7.80	110.34	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	950	C	C5'-C4'-C3'	-7.80	103.52	116.00
34	BA	1164	C	O4'-C1'-N1	7.80	114.44	108.20
38	BE	63	C	C5'-C4'-C3'	7.80	128.48	116.00
40	BG	149	U	C1'-O4'-C4'	-7.80	103.66	109.90
85	AA	80	G	P-O5'-C5'	-7.80	108.42	120.90
85	AA	265	A	C3'-C2'-C1'	-7.80	95.26	101.50
85	AA	488	G	C5-C6-N1	7.80	115.40	111.50
85	AA	2194	U	C2-N1-C1'	-7.80	108.34	117.70
34	BA	81	C	O4'-C1'-C2'	7.80	114.62	107.60
34	BA	817	U	C5'-C4'-O4'	7.80	118.46	109.10
34	BA	980	C	C5'-C4'-C3'	-7.80	103.52	116.00
85	AA	534	A	C4-N9-C1'	-7.80	112.26	126.30
34	BA	1450	G	N1-C6-O6	7.80	124.58	119.90
34	BA	1611	A	C8-N9-C4	-7.80	102.68	105.80
45	BL	56	ARG	NE-CZ-NH2	-7.80	116.40	120.30
8	A7	5	TYR	CB-CG-CD2	-7.79	116.32	121.00
84	By	178	TYR	CB-CG-CD1	7.79	125.68	121.00
85	AA	158	C	C6-N1-C1'	7.79	130.15	120.80
85	AA	1009	G	C8-N9-C1'	7.79	137.13	127.00
34	BA	97	A	N1-C6-N6	-7.79	113.92	118.60
35	BB	52	G	C4-N9-C1'	-7.79	116.37	126.50
35	BB	132	G	C8-N9-C1'	7.79	137.13	127.00
35	BB	1399	A	C8-N9-C4	7.79	108.92	105.80
35	BB	1544	A	C4-N9-C1'	-7.79	112.27	126.30
85	AA	569	A	C5-C6-N6	7.79	129.93	123.70
85	AA	1823	G	C4'-C3'-C2'	7.79	110.39	102.60
35	BB	38	C	C5'-C4'-C3'	-7.79	103.53	116.00
35	BB	603	U	C5'-C4'-O4'	7.79	118.45	109.10
35	BB	888	U	O4'-C1'-N1	7.79	114.43	108.20
77	Br	77	ARG	NE-CZ-NH1	7.79	124.20	120.30
85	AA	187	C	C1'-O4'-C4'	-7.79	103.67	109.90
7	A6	170	ARG	NE-CZ-NH1	7.79	124.19	120.30
34	BA	528	C	C5-C6-N1	7.79	124.89	121.00
34	BA	1300	G	C5'-C4'-C3'	-7.79	103.54	116.00
35	BB	2	C	P-O3'-C3'	-7.79	110.35	119.70
53	BT	110	ARG	NE-CZ-NH1	7.79	124.19	120.30
84	By	111	ARG	NE-CZ-NH1	7.79	124.19	120.30
85	AA	304	G	O3'-P-O5'	7.79	118.80	104.00
85	AA	857	G	C3'-C2'-C1'	-7.79	95.27	101.50
35	BB	1167	C	C2'-C3'-O3'	7.79	126.63	109.50
15	AG	128	TYR	CB-CG-CD2	7.79	125.67	121.00
34	BA	1300	G	C1'-O4'-C4'	-7.79	103.67	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1809	G	O3'-P-O5'	7.79	118.79	104.00
35	BB	1168	G	P-O3'-C3'	-7.79	110.36	119.70
40	BG	146	C	C1'-O4'-C4'	-7.79	103.67	109.90
53	BT	65	GLN	N-CA-CB	-7.79	96.59	110.60
67	Bh	148	ARG	CD-NE-CZ	-7.79	112.70	123.60
35	BB	806	U	C2-N3-C4	-7.78	122.33	127.00
85	AA	1432	C	N3-C4-N4	7.78	123.45	118.00
85	AA	1837	U	O4'-C1'-N1	7.78	114.43	108.20
38	BE	184	G	C8-N9-C4	-7.78	103.29	106.40
38	BE	195	G	N1-C2-N2	-7.78	109.20	116.20
85	AA	824	C	P-O3'-C3'	7.78	129.04	119.70
85	AA	1213	U	C2-N1-C1'	7.78	127.04	117.70
34	BA	124	G	C8-N9-C1'	7.78	137.12	127.00
34	BA	565	U	C5'-C4'-C3'	-7.78	103.55	116.00
34	BA	772	G	N7-C8-N9	7.78	116.99	113.10
34	BA	1022	C	O4'-C1'-N1	7.78	114.42	108.20
34	BA	1109	G	N9-C1'-C2'	-7.78	103.44	112.00
34	BA	1261	G	P-O3'-C3'	-7.78	110.36	119.70
35	BB	865	C	P-O3'-C3'	-7.78	110.36	119.70
35	BB	1232	A	C1'-O4'-C4'	-7.78	103.67	109.90
36	BC	23	G	C5-C6-O6	7.78	133.27	128.60
38	BE	32	U	O4'-C1'-N1	-7.78	101.98	108.20
34	BA	104	A	C3'-C2'-C1'	-7.78	95.28	101.50
35	BB	1477	C	C6-N1-C2	-7.78	117.19	120.30
40	BG	12	A	C5'-C4'-O4'	7.78	118.44	109.10
85	AA	584	G	N1-C6-O6	7.78	124.57	119.90
85	AA	1595	G	C5-C6-O6	-7.78	123.93	128.60
34	BA	601	A	C5'-C4'-C3'	7.78	128.44	116.00
34	BA	743	A	C6-C5-N7	-7.78	126.86	132.30
34	BA	779	U	O4'-C1'-N1	7.78	114.42	108.20
35	BB	600	C	N1-C1'-C2'	-7.78	103.44	112.00
37	BD	69	U	N3-C2-O2	-7.78	116.76	122.20
85	AA	203	C	C2-N1-C1'	7.78	127.36	118.80
85	AA	342	C	P-O3'-C3'	-7.78	110.37	119.70
85	AA	371	C	P-O3'-C3'	-7.78	110.37	119.70
85	AA	909	C	C6-N1-C1'	-7.78	111.47	120.80
34	BA	227	C	O4'-C1'-N1	7.78	114.42	108.20
34	BA	279	U	C2-N3-C4	-7.78	122.33	127.00
34	BA	550	U	N3-C4-O4	-7.78	113.96	119.40
35	BB	1211	C	P-O5'-C5'	7.78	133.34	120.90
85	AA	1528	A	P-O3'-C3'	7.78	129.03	119.70
85	AA	2246	U	O5'-P-OP1	-7.78	98.70	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1363	A	C1'-O4'-C4'	-7.77	103.68	109.90
34	BA	1816	G	P-O3'-C3'	7.77	129.03	119.70
35	BB	652	G	O4'-C1'-N9	7.77	114.42	108.20
35	BB	1024	G	N3-C4-C5	-7.77	124.71	128.60
34	BA	1202	G	C8-N9-C1'	7.77	137.10	127.00
35	BB	993	A	P-O5'-C5'	7.77	133.34	120.90
36	BC	11	G	N3-C2-N2	7.77	125.34	119.90
38	BE	32	U	C2-N3-C4	-7.77	122.34	127.00
85	AA	385	A	N1-C6-N6	7.77	123.26	118.60
85	AA	386	G	N3-C2-N2	7.77	125.34	119.90
85	AA	1009	G	O4'-C1'-N9	7.77	114.42	108.20
86	AB	16	U	O5'-C5'-C4'	7.77	126.47	111.70
34	BA	1485	U	O4'-C1'-N1	7.77	114.42	108.20
35	BB	57	G	C8-N9-C1'	7.77	137.10	127.00
35	BB	1053	G	C2-N3-C4	-7.77	108.01	111.90
85	AA	1291	A	O4'-C4'-C3'	-7.77	96.23	104.00
35	BB	677	U	C4'-C3'-C2'	7.77	110.37	102.60
40	BG	169	A	C4-C5-C6	-7.77	113.11	117.00
41	BH	60	A	O4'-C1'-N9	7.77	114.42	108.20
64	Be	133	TYR	CB-CG-CD2	-7.77	116.34	121.00
85	AA	1487	G	N9-C1'-C2'	-7.77	103.45	112.00
34	BA	1305	A	C5-C6-N6	7.77	129.91	123.70
35	BB	141	G	N1-C6-O6	7.77	124.56	119.90
41	BH	30	C	C5'-C4'-O4'	7.77	118.42	109.10
85	AA	915	G	C5-C6-O6	-7.77	123.94	128.60
85	AA	659	A	C3'-C2'-C1'	-7.77	95.29	101.50
85	AA	854	A	O4'-C4'-C3'	-7.77	96.23	104.00
34	BA	756	A	C5'-C4'-C3'	7.76	128.42	116.00
35	BB	534	C	C3'-C2'-C1'	-7.76	95.29	101.50
35	BB	1208	G	O4'-C1'-C2'	-7.76	98.04	105.80
41	BH	120	C	O5'-P-OP1	-7.76	98.71	105.70
85	AA	789	A	C5'-C4'-O4'	7.76	118.42	109.10
85	AA	1265	C	O4'-C1'-N1	7.76	114.41	108.20
85	AA	2209	U	C2-N1-C1'	-7.76	108.38	117.70
16	AH	114	ARG	NE-CZ-NH2	-7.76	116.42	120.30
35	BB	1063	C	N3-C2-O2	-7.76	116.47	121.90
35	BB	1531	G	O4'-C1'-N9	7.76	114.41	108.20
38	BE	36	U	C6-N1-C2	-7.76	116.34	121.00
85	AA	649	C	P-O5'-C5'	-7.76	108.48	120.90
85	AA	858	G	C4-N9-C1'	-7.76	116.41	126.50
10	A9	140	ARG	NE-CZ-NH1	7.76	124.18	120.30
34	BA	246	G	C2'-C3'-O3'	7.76	126.58	109.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1800	G	C1'-O4'-C4'	-7.76	103.69	109.90
35	BB	285	C	C6-N1-C2	-7.76	117.19	120.30
35	BB	708	C	C2-N3-C4	7.76	123.78	119.90
35	BB	775	U	O4'-C1'-N1	7.76	114.41	108.20
35	BB	901	U	C5'-C4'-C3'	-7.76	103.58	116.00
35	BB	1475	U	C2-N3-C4	-7.76	122.34	127.00
38	BE	62	C	C2-N1-C1'	-7.76	110.26	118.80
85	AA	587	G	C4-N9-C1'	-7.76	116.41	126.50
85	AA	594	C	C2-N1-C1'	-7.76	110.26	118.80
85	AA	962	U	P-O3'-C3'	-7.76	110.39	119.70
85	AA	1432	C	C5'-C4'-C3'	7.76	128.42	116.00
85	AA	1570	A	C1'-O4'-C4'	-7.76	103.69	109.90
34	BA	499	C	C2-N1-C1'	-7.76	110.26	118.80
34	BA	889	U	C2-N1-C1'	-7.76	108.39	117.70
34	BA	1119	A	O5'-C5'-C4'	-7.76	96.96	111.70
34	BA	1473	A	C1'-O4'-C4'	-7.76	103.69	109.90
34	BA	1656	A	C5-C6-N1	7.76	121.58	117.70
38	BE	25	U	O5'-P-OP1	7.76	120.01	110.70
41	BH	7	C	P-O5'-C5'	-7.76	108.48	120.90
85	AA	45	U	P-O5'-C5'	-7.76	108.49	120.90
85	AA	2092	A	O4'-C1'-N9	7.76	114.41	108.20
34	BA	1000	G	P-O3'-C3'	-7.76	110.39	119.70
34	BA	1814	U	C6-N1-C2	-7.76	116.34	121.00
85	AA	966	G	C5'-C4'-C3'	-7.76	103.59	116.00
34	BA	755	G	C6-N1-C2	-7.76	120.45	125.10
34	BA	896	U	P-O3'-C3'	7.76	129.01	119.70
34	BA	1411	C	C5'-C4'-C3'	-7.76	103.59	116.00
34	BA	1721	U	P-O5'-C5'	-7.76	108.49	120.90
38	BE	1	U	C3'-C2'-C1'	-7.76	95.29	101.50
40	BG	8	U	O4'-C1'-N1	7.76	114.41	108.20
84	By	51	ARG	NE-CZ-NH2	-7.76	116.42	120.30
34	BA	532	C	C5'-C4'-O4'	7.75	118.41	109.10
35	BB	268	G	C5-C6-O6	-7.75	123.95	128.60
35	BB	380	G	C8-N9-C1'	7.75	137.08	127.00
40	BG	35	G	C5'-C4'-C3'	-7.75	103.59	116.00
85	AA	157	G	C1'-O4'-C4'	-7.75	103.70	109.90
85	AA	158	C	C1'-O4'-C4'	-7.75	103.70	109.90
85	AA	1102	C	C6-N1-C2	-7.75	117.20	120.30
34	BA	389	U	P-O3'-C3'	-7.75	110.40	119.70
34	BA	1572	G	N1-C6-O6	-7.75	115.25	119.90
35	BB	827	U	P-O5'-C5'	7.75	133.30	120.90
38	BE	82	C	O3'-P-O5'	-7.75	89.27	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
59	BZ	51	ASP	N-CA-CB	-7.75	96.64	110.60
85	AA	1701	G	N9-C4-C5	7.75	108.50	105.40
85	AA	1974	C	O4'-C1'-N1	7.75	114.40	108.20
34	BA	530	A	C2'-C3'-O3'	7.75	126.56	109.50
34	BA	543	A	C5'-C4'-C3'	-7.75	103.60	116.00
35	BB	771	U	P-O3'-C3'	7.75	129.00	119.70
85	AA	449	G	C8-N9-C4	7.75	109.50	106.40
34	BA	382	G	C5-C6-O6	7.75	133.25	128.60
34	BA	1002	U	C3'-C2'-C1'	-7.75	95.30	101.50
85	AA	1541	G	O5'-C5'-C4'	-7.75	96.97	111.70
34	BA	145	U	C5'-C4'-C3'	-7.75	103.60	116.00
34	BA	1326	U	C2-N1-C1'	7.75	127.00	117.70
34	BA	1506	C	C5'-C4'-C3'	7.75	128.40	116.00
34	BA	1518	A	N9-C4-C5	-7.75	102.70	105.80
85	AA	447	C	O4'-C1'-N1	7.75	114.40	108.20
34	BA	1119	A	C5-C6-N6	-7.75	117.50	123.70
36	BC	90	U	O4'-C1'-N1	7.75	114.40	108.20
38	BE	104	G	C8-N9-C1'	-7.75	116.93	127.00
85	AA	723	U	O4'-C1'-N1	7.75	114.40	108.20
85	AA	860	C	N1-C2-O2	7.75	123.55	118.90
85	AA	917	A	C2-N3-C4	-7.75	106.73	110.60
85	AA	1845	G	C8-N9-C1'	7.75	137.07	127.00
85	AA	1865	C	C2-N3-C4	-7.75	116.03	119.90
34	BA	766	A	C5'-C4'-O4'	7.75	118.39	109.10
35	BB	621	C	O3'-P-O5'	7.75	118.72	104.00
39	BF	48	G	C1'-O4'-C4'	-7.75	103.70	109.90
85	AA	463	G	O3'-P-O5'	-7.75	89.28	104.00
85	AA	1734	A	C5'-C4'-C3'	7.75	128.39	116.00
34	BA	149	G	C1'-O4'-C4'	-7.74	103.70	109.90
34	BA	505	U	P-O5'-C5'	-7.74	108.51	120.90
34	BA	761	U	C6-N1-C1'	7.74	132.04	121.20
34	BA	1083	A	P-O3'-C3'	7.74	128.99	119.70
35	BB	2	C	C4'-C3'-C2'	-7.74	94.86	102.60
35	BB	1432	U	C2-N1-C1'	-7.74	108.41	117.70
35	BB	1532	C	C5'-C4'-C3'	7.74	128.39	116.00
39	BF	62	U	P-O5'-C5'	7.74	133.29	120.90
40	BG	71	C	C6-N1-C2	-7.74	117.20	120.30
41	BH	9	C	P-O3'-C3'	-7.74	110.41	119.70
41	BH	71	C	O4'-C1'-N1	7.74	114.39	108.20
83	Bx	263	ARG	NE-CZ-NH2	-7.74	116.43	120.30
85	AA	65	A	C8-N9-C1'	-7.74	113.76	127.70
85	AA	1278	C	O4'-C1'-N1	7.74	114.39	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	797	C	O5'-C5'-C4'	7.74	126.41	111.70
35	BB	1115	G	C4-N9-C1'	-7.74	116.44	126.50
85	AA	2122	A	C2'-C3'-O3'	7.74	126.53	109.50
35	BB	1386	C	C6-N1-C2	-7.74	117.20	120.30
85	AA	254	G	C5-C6-O6	-7.74	123.95	128.60
85	AA	518	A	C5'-C4'-O4'	7.74	118.39	109.10
85	AA	566	U	O4'-C1'-N1	7.74	114.39	108.20
85	AA	643	C	C2-N1-C1'	-7.74	110.29	118.80
85	AA	1415	G	P-O3'-C3'	7.74	128.99	119.70
34	BA	182	U	C2-N3-C4	-7.74	122.36	127.00
34	BA	688	G	C5-C6-O6	-7.74	123.96	128.60
35	BB	681	G	P-O3'-C3'	-7.74	110.41	119.70
35	BB	824	C	O4'-C4'-C3'	-7.74	96.26	104.00
35	BB	888	U	P-O3'-C3'	-7.74	110.41	119.70
35	BB	905	C	O4'-C1'-N1	7.74	114.39	108.20
35	BB	1399	A	N1-C6-N6	-7.74	113.96	118.60
85	AA	1730	C	C3'-C2'-C1'	-7.74	95.31	101.50
34	BA	288	U	N1-C2-N3	7.74	119.54	114.90
85	AA	803	C	C5'-C4'-C3'	-7.74	103.62	116.00
34	BA	540	G	C5-C6-O6	-7.74	123.96	128.60
34	BA	883	C	C6-N1-C2	-7.74	117.20	120.30
34	BA	1029	C	O4'-C1'-N1	7.74	114.39	108.20
34	BA	1510	C	N3-C4-N4	-7.74	112.58	118.00
35	BB	378	C	C3'-C2'-C1'	-7.74	95.31	101.50
35	BB	834	U	C5'-C4'-O4'	7.74	118.38	109.10
35	BB	1450	G	C5-C6-O6	-7.74	123.96	128.60
38	BE	107	U	C2-N3-C4	-7.74	122.36	127.00
85	AA	968	U	C5'-C4'-C3'	7.74	128.38	116.00
85	AA	2014	G	C6-N1-C2	-7.74	120.46	125.10
34	BA	532	C	C5'-C4'-C3'	-7.73	103.62	116.00
47	BN	41	ARG	NE-CZ-NH1	7.73	124.17	120.30
85	AA	748	C	O4'-C1'-N1	7.73	114.39	108.20
85	AA	1832	G	C4-N9-C1'	-7.73	116.45	126.50
34	BA	1418	G	C5-C6-O6	-7.73	123.96	128.60
34	BA	1495	A	C1'-O4'-C4'	-7.73	103.71	109.90
35	BB	429	C	C1'-O4'-C4'	-7.73	103.71	109.90
35	BB	484	G	C5-C6-O6	-7.73	123.96	128.60
35	BB	1517	G	N1-C6-O6	7.73	124.54	119.90
36	BC	116	C	C1'-O4'-C4'	-7.73	103.71	109.90
40	BG	25	G	P-O3'-C3'	-7.73	110.42	119.70
34	BA	89	G	N3-C2-N2	-7.73	114.49	119.90
34	BA	815	C	P-O3'-C3'	-7.73	110.42	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1011	G	N3-C2-N2	-7.73	114.49	119.90
34	BA	1822	U	C6-N1-C1'	-7.73	110.38	121.20
35	BB	567	G	C5'-C4'-C3'	-7.73	103.63	116.00
35	BB	1020	U	N3-C2-O2	-7.73	116.79	122.20
56	BW	12	ARG	NE-CZ-NH1	7.73	124.17	120.30
85	AA	1809	G	C5-C6-O6	-7.73	123.96	128.60
34	BA	271	C	C6-N1-C2	-7.73	117.21	120.30
34	BA	766	A	N1-C6-N6	-7.73	113.96	118.60
34	BA	1485	U	C5-C4-O4	7.73	130.54	125.90
34	BA	1636	C	P-O5'-C5'	-7.73	108.53	120.90
34	BA	52	G	P-O3'-C3'	-7.73	110.43	119.70
34	BA	807	U	C6-N1-C1'	-7.73	110.38	121.20
34	BA	1419	A	P-O5'-C5'	-7.73	108.53	120.90
35	BB	1025	A	C3'-C2'-C1'	7.73	107.68	101.50
55	BV	92	TYR	CB-CG-CD1	7.73	125.64	121.00
85	AA	327	G	C3'-C2'-C1'	-7.73	95.32	101.50
85	AA	508	C	O5'-P-OP2	-7.73	98.75	105.70
34	BA	582	U	N1-C2-N3	-7.73	110.26	114.90
37	BD	30	A	C5-C6-N6	-7.73	117.52	123.70
85	AA	744	C	N1-C2-O2	7.73	123.54	118.90
29	AV	59	SER	N-CA-CB	7.72	122.09	110.50
35	BB	1453	G	P-O3'-C3'	-7.72	110.43	119.70
40	BG	14	G	C6-N1-C2	-7.72	120.47	125.10
41	BH	26	C	C3'-C2'-C1'	7.72	107.68	101.50
85	AA	2196	G	N9-C4-C5	7.72	108.49	105.40
34	BA	606	G	N3-C4-C5	-7.72	124.74	128.60
34	BA	937	G	C1'-O4'-C4'	-7.72	103.72	109.90
34	BA	1113	A	C5'-C4'-C3'	7.72	128.36	116.00
34	BA	1503	U	O4'-C4'-C3'	-7.72	96.28	104.00
38	BE	55	C	C6-N1-C2	-7.72	117.21	120.30
38	BE	183	C	O5'-C5'-C4'	7.72	126.37	111.70
41	BH	35	G	C5'-C4'-C3'	-7.72	103.64	116.00
60	Ba	64	ARG	CD-NE-CZ	-7.72	112.79	123.60
85	AA	1270	C	C2-N1-C1'	-7.72	110.31	118.80
29	AV	62	TYR	CB-CG-CD2	-7.72	116.37	121.00
34	BA	1567	G	C4-N9-C1'	-7.72	116.46	126.50
35	BB	1463	A	C8-N9-C4	-7.72	102.71	105.80
34	BA	7	U	C6-N1-C2	-7.72	116.37	121.00
34	BA	288	U	C1'-O4'-C4'	-7.72	103.72	109.90
34	BA	602	G	N3-C2-N2	-7.72	114.50	119.90
34	BA	889	U	C4'-C3'-C2'	7.72	110.32	102.60
34	BA	1040	G	N3-C2-N2	7.72	125.30	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1665	G	C4-N9-C1'	-7.72	116.46	126.50
40	BG	148	C	C6-N1-C2	-7.72	117.21	120.30
85	AA	311	U	C4'-C3'-C2'	7.72	110.32	102.60
34	BA	1530	G	C4-N9-C1'	-7.72	116.47	126.50
34	BA	1653	G	C8-N9-C1'	7.72	137.03	127.00
68	Bi	44	ARG	NE-CZ-NH1	7.72	124.16	120.30
26	AS	67	ARG	NE-CZ-NH1	7.72	124.16	120.30
34	BA	1047	U	P-O3'-C3'	-7.72	110.44	119.70
38	BE	127	G	O4'-C1'-C2'	7.72	114.55	107.60
38	BE	169	C	O4'-C1'-N1	7.72	114.37	108.20
41	BH	37	U	P-O3'-C3'	7.72	128.96	119.70
85	AA	331	G	C4'-C3'-C2'	-7.72	94.88	102.60
85	AA	945	A	O4'-C1'-N9	7.72	114.37	108.20
85	AA	1456	A	C8-N9-C4	7.72	108.89	105.80
35	BB	391	G	N3-C2-N2	7.71	125.30	119.90
35	BB	663	G	C5-C6-N1	7.71	115.36	111.50
77	Br	258	LEU	N-CA-CB	-7.71	94.97	110.40
85	AA	527	A	O4'-C1'-N9	7.71	114.37	108.20
85	AA	1756	C	C2-N1-C1'	-7.71	110.31	118.80
4	A3	7	TYR	CB-CG-CD1	7.71	125.63	121.00
35	BB	839	G	C5-C6-O6	7.71	133.23	128.60
38	BE	135	A	C4'-C3'-C2'	-7.71	94.89	102.60
40	BG	111	C	N3-C2-O2	-7.71	116.50	121.90
41	BH	100	A	C6-C5-N7	-7.71	126.90	132.30
85	AA	680	U	P-O3'-C3'	7.71	128.96	119.70
34	BA	1284	G	C5-C6-O6	7.71	133.23	128.60
34	BA	1305	A	C4-N9-C1'	7.71	140.18	126.30
34	BA	1356	C	O4'-C1'-N1	7.71	114.37	108.20
35	BB	537	A	P-O5'-C5'	-7.71	108.56	120.90
71	Bl	37	ARG	NE-CZ-NH2	-7.71	116.44	120.30
85	AA	150	U	P-O5'-C5'	7.71	133.24	120.90
85	AA	959	C	O4'-C1'-N1	7.71	114.37	108.20
85	AA	1322	C	O4'-C1'-N1	7.71	114.37	108.20
85	AA	1609	U	O4'-C1'-N1	7.71	114.37	108.20
85	AA	1825	A	C4'-C3'-C2'	7.71	110.31	102.60
85	AA	2198	G	P-O3'-C3'	-7.71	110.45	119.70
34	BA	1421	A	P-O3'-C3'	-7.71	110.45	119.70
34	BA	1644	A	C5'-C4'-C3'	-7.71	103.66	116.00
35	BB	1031	G	C1'-O4'-C4'	-7.71	103.73	109.90
35	BB	1294	C	N1-C1'-C2'	-7.71	103.52	112.00
85	AA	188	G	C3'-C2'-C1'	-7.71	95.33	101.50
85	AA	1083	C	O4'-C1'-N1	7.71	114.37	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1994	G	C8-N9-C1'	7.71	137.02	127.00
34	BA	63	A	O4'-C1'-C2'	-7.71	98.09	105.80
34	BA	686	U	O4'-C4'-C3'	-7.71	96.29	104.00
35	BB	1155	U	N3-C2-O2	-7.71	116.81	122.20
35	BB	1222	A	O5'-P-OP1	-7.71	98.76	105.70
45	BL	179	PHE	CB-CG-CD1	7.71	126.20	120.80
70	Bk	45	LEU	N-CA-C	-7.71	90.19	111.00
85	AA	218	U	O4'-C1'-N1	7.71	114.37	108.20
85	AA	252	G	O4'-C1'-N9	-7.71	102.03	108.20
85	AA	858	G	C3'-C2'-C1'	-7.71	95.33	101.50
34	BA	1297	G	P-O5'-C5'	7.71	133.23	120.90
35	BB	1186	A	C4-C5-C6	-7.71	113.15	117.00
35	BB	1423	U	C1'-O4'-C4'	-7.71	103.73	109.90
40	BG	168	A	C4-N9-C1'	-7.71	112.43	126.30
34	BA	82	A	C3'-C2'-C1'	-7.71	95.34	101.50
12	AD	14	TYR	CB-CG-CD1	7.70	125.62	121.00
34	BA	1013	A	C5'-C4'-C3'	-7.70	103.67	116.00
34	BA	1179	U	P-O3'-C3'	7.70	128.94	119.70
34	BA	1501	U	O4'-C1'-N1	7.70	114.36	108.20
85	AA	562	C	O4'-C1'-N1	7.70	114.36	108.20
85	AA	2225	G	C8-N9-C1'	7.70	137.01	127.00
34	BA	798	G	N9-C4-C5	-7.70	102.32	105.40
85	AA	800	A	P-O3'-C3'	7.70	128.94	119.70
2	A1	63	ARG	NE-CZ-NH1	7.70	124.15	120.30
34	BA	312	U	C5'-C4'-C3'	7.70	128.32	116.00
85	AA	581	A	C5'-C4'-O4'	7.70	118.34	109.10
85	AA	962	U	C6-N1-C1'	7.70	131.98	121.20
34	BA	794	G	N9-C1'-C2'	-7.70	103.53	112.00
35	BB	1003	G	C8-N9-C1'	7.70	137.01	127.00
35	BB	1394	A	C6-N1-C2	-7.70	113.98	118.60
36	BC	99	U	C2-N3-C4	-7.70	122.38	127.00
85	AA	325	C	O4'-C1'-N1	7.70	114.36	108.20
35	BB	1251	G	N3-C4-C5	-7.70	124.75	128.60
35	BB	1394	A	C5-C6-N1	7.70	121.55	117.70
38	BE	25	U	N3-C4-C5	7.70	119.22	114.60
52	BS	9	TYR	CB-CG-CD2	-7.70	116.38	121.00
27	AT	38	CYS	CA-CB-SG	7.70	127.85	114.00
35	BB	724	G	O4'-C1'-N9	7.70	114.36	108.20
35	BB	1108	G	C5'-C4'-C3'	7.70	128.31	116.00
35	BB	1519	U	O4'-C1'-N1	7.70	114.36	108.20
84	By	54	THR	N-CA-CB	7.70	124.92	110.30
84	By	132	ARG	NE-CZ-NH1	7.70	124.15	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	585	G	N9-C4-C5	-7.70	102.32	105.40
3	A2	43	GLN	N-CA-C	7.69	131.77	111.00
34	BA	91	C	C2-N1-C1'	-7.69	110.34	118.80
34	BA	1126	U	C5-C6-N1	7.69	126.55	122.70
34	BA	1619	U	C2'-C3'-O3'	7.69	126.43	109.50
35	BB	1201	G	C5'-C4'-C3'	7.69	128.31	116.00
34	BA	410	G	O4'-C1'-N9	7.69	114.35	108.20
85	AA	300	C	O4'-C1'-N1	7.69	114.35	108.20
85	AA	666	A	N1-C6-N6	-7.69	113.98	118.60
85	AA	1034	U	C3'-C2'-C1'	7.69	107.65	101.50
85	AA	1243	G	C5'-C4'-C3'	7.69	128.31	116.00
85	AA	1371	C	C6-N1-C1'	7.69	130.03	120.80
85	AA	1711	C	O4'-C1'-C2'	7.69	114.52	107.60
34	BA	107	C	O4'-C1'-N1	7.69	114.35	108.20
34	BA	481	A	N3-C4-C5	7.69	132.18	126.80
34	BA	802	G	C8-N9-C1'	7.69	137.00	127.00
35	BB	1133	C	C6-N1-C1'	7.69	130.03	120.80
38	BE	67	A	O3'-P-O5'	7.69	118.61	104.00
38	BE	185	G	C5'-C4'-C3'	-7.69	103.69	116.00
85	AA	557	G	C6-N1-C2	-7.69	120.48	125.10
85	AA	626	G	N1-C6-O6	7.69	124.51	119.90
85	AA	887	A	P-O3'-C3'	-7.69	110.47	119.70
85	AA	1109	G	O4'-C1'-N9	7.69	114.35	108.20
85	AA	1474	U	O4'-C1'-N1	7.69	114.35	108.20
85	AA	2141	G	O4'-C1'-C2'	7.69	114.52	107.60
34	BA	452	A	N1-C6-N6	7.69	123.21	118.60
34	BA	1003	A	O5'-P-OP2	-7.69	98.78	105.70
34	BA	1428	G	P-O5'-C5'	-7.69	108.60	120.90
35	BB	1461	C	C5-C4-N4	-7.69	114.82	120.20
36	BC	88	A	P-O3'-C3'	-7.69	110.47	119.70
85	AA	1494	C	O5'-P-OP2	7.69	119.93	110.70
34	BA	188	C	O4'-C1'-N1	7.69	114.35	108.20
34	BA	813	C	C5'-C4'-O4'	-7.69	99.88	109.10
34	BA	1537	G	C1'-O4'-C4'	-7.69	103.75	109.90
35	BB	1016	C	C6-N1-C2	-7.69	117.22	120.30
39	BF	23	G	C5'-C4'-O4'	7.69	118.33	109.10
71	Bl	62	ARG	NE-CZ-NH1	7.69	124.14	120.30
85	AA	272	C	C6-N1-C2	-7.69	117.22	120.30
85	AA	1294	U	N1-C2-O2	-7.69	117.42	122.80
34	BA	1494	G	C5-C6-N1	7.69	115.34	111.50
34	BA	1505	G	C8-N9-C4	-7.69	103.33	106.40
34	BA	1540	C	O4'-C1'-N1	7.69	114.35	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1112	U	C6-N1-C2	-7.69	116.39	121.00
35	BB	1475	U	O4'-C1'-N1	7.69	114.35	108.20
85	AA	743	C	C5'-C4'-C3'	7.69	128.30	116.00
85	AA	2070	C	C6-N1-C2	-7.69	117.23	120.30
34	BA	386	A	C4'-C3'-C2'	7.68	110.28	102.60
35	BB	743	C	O4'-C1'-N1	7.68	114.35	108.20
35	BB	800	U	O5'-C5'-C4'	7.68	126.30	111.70
85	AA	268	A	P-O3'-C3'	-7.68	110.48	119.70
85	AA	1294	U	C4'-C3'-C2'	-7.68	94.92	102.60
35	BB	313	C	O4'-C1'-N1	7.68	114.35	108.20
35	BB	887	G	O3'-P-O5'	-7.68	89.40	104.00
36	BC	23	G	C4'-C3'-C2'	-7.68	94.92	102.60
37	BD	4	U	C5'-C4'-C3'	-7.68	103.71	116.00
85	AA	989	U	C5'-C4'-C3'	7.68	128.29	116.00
85	AA	1460	G	C5-C6-O6	-7.68	123.99	128.60
85	AA	1983	C	P-O3'-C3'	7.68	128.92	119.70
85	AA	2018	U	C5-C4-O4	-7.68	121.29	125.90
34	BA	930	A	C5-C6-N6	7.68	129.84	123.70
35	BB	118	A	C1'-O4'-C4'	-7.68	103.76	109.90
40	BG	118	U	C2-N3-C4	-7.68	122.39	127.00
85	AA	432	A	C8-N9-C1'	7.68	141.53	127.70
34	BA	183	G	O4'-C1'-N9	7.68	114.34	108.20
34	BA	470	C	O4'-C1'-N1	7.68	114.34	108.20
34	BA	530	A	C4-C5-C6	-7.68	113.16	117.00
34	BA	836	U	P-O5'-C5'	-7.68	108.61	120.90
35	BB	1358	A	N1-C6-N6	-7.68	113.99	118.60
38	BE	20	C	C2-N1-C1'	7.68	127.25	118.80
38	BE	142	A	C5'-C4'-O4'	7.68	118.31	109.10
47	BN	203	ARG	NE-CZ-NH1	7.68	124.14	120.30
85	AA	391	G	N1-C6-O6	-7.68	115.29	119.90
85	AA	1722	G	P-O3'-C3'	7.68	128.91	119.70
85	AA	633	C	C1'-O4'-C4'	-7.68	103.76	109.90
34	BA	468	A	O4'-C1'-N9	7.68	114.34	108.20
34	BA	827	A	C1'-O4'-C4'	-7.68	103.76	109.90
34	BA	1472	G	C4-C5-C6	-7.68	114.19	118.80
76	Bq	4	PHE	CB-CG-CD1	7.68	126.17	120.80
34	BA	534	C	O4'-C1'-N1	7.67	114.34	108.20
85	AA	1484	G	N1-C6-O6	-7.67	115.30	119.90
34	BA	1094	U	C5'-C4'-C3'	-7.67	103.72	116.00
85	AA	269	G	C5-C6-O6	-7.67	124.00	128.60
85	AA	598	C	O4'-C1'-N1	7.67	114.34	108.20
34	BA	871	G	O5'-P-OP2	-7.67	98.80	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	987	C	O4'-C1'-N1	7.67	114.34	108.20
35	BB	831	C	P-O3'-C3'	-7.67	110.49	119.70
35	BB	1168	G	C5'-C4'-O4'	7.67	118.31	109.10
37	BD	2	G	C8-N9-C1'	7.67	136.97	127.00
85	AA	251	A	C4-N9-C1'	7.67	140.11	126.30
34	BA	1283	U	O4'-C1'-N1	7.67	114.34	108.20
34	BA	1408	C	P-O5'-C5'	-7.67	108.63	120.90
40	BG	16	G	C5-C6-N1	7.67	115.33	111.50
40	BG	140	G	C4-N9-C1'	-7.67	116.53	126.50
85	AA	288	G	C5-C6-O6	-7.67	124.00	128.60
16	AH	114	ARG	NE-CZ-NH1	7.67	124.14	120.30
35	BB	551	C	O4'-C1'-N1	7.67	114.33	108.20
85	AA	852	C	O4'-C1'-N1	7.67	114.33	108.20
85	AA	1953	G	O4'-C1'-N9	7.67	114.33	108.20
34	BA	208	A	N1-C6-N6	-7.67	114.00	118.60
34	BA	878	G	O4'-C1'-C2'	7.67	114.50	107.60
34	BA	1556	A	P-O3'-C3'	7.67	128.90	119.70
34	BA	1707	C	N3-C2-O2	-7.67	116.53	121.90
34	BA	1788	U	C6-N1-C2	-7.67	116.40	121.00
35	BB	714	U	C2-N3-C4	-7.67	122.40	127.00
38	BE	168	C	C2-N1-C1'	-7.67	110.37	118.80
40	BG	169	A	C1'-O4'-C4'	-7.67	103.77	109.90
85	AA	421	G	N1-C6-O6	-7.67	115.30	119.90
85	AA	939	A	O3'-P-O5'	7.67	118.57	104.00
85	AA	2156	C	P-O5'-C5'	7.67	133.16	120.90
35	BB	404	A	C4-C5-C6	-7.67	113.17	117.00
85	AA	1106	A	O4'-C4'-C3'	-7.67	96.33	104.00
85	AA	1465	C	C4'-C3'-C2'	-7.67	94.94	102.60
31	AX	50	ARG	NE-CZ-NH1	7.66	124.13	120.30
34	BA	1090	A	C5'-C4'-C3'	7.66	128.26	116.00
34	BA	1160	U	P-O3'-C3'	-7.66	110.50	119.70
41	BH	71	C	O3'-P-O5'	7.66	118.56	104.00
50	BQ	192	ARG	CD-NE-CZ	-7.66	112.87	123.60
85	AA	2207	A	O4'-C1'-N9	7.66	114.33	108.20
34	BA	110	C	C3'-C2'-C1'	-7.66	95.37	101.50
34	BA	168	U	O4'-C1'-N1	7.66	114.33	108.20
41	BH	10	U	C2-N1-C1'	-7.66	108.51	117.70
60	Ba	28	THR	N-CA-CB	7.66	124.85	110.30
85	AA	714	U	C4'-C3'-C2'	-7.66	94.94	102.60
85	AA	1553	G	C4'-C3'-C2'	-7.66	94.94	102.60
34	BA	871	G	N7-C8-N9	-7.66	109.27	113.10
34	BA	956	G	O4'-C1'-N9	7.66	114.33	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	36	C	P-O5'-C5'	7.66	133.15	120.90
85	AA	1096	G	C4-N9-C1'	-7.66	116.55	126.50
34	BA	343	G	C5-C6-N1	7.66	115.33	111.50
34	BA	741	A	P-O5'-C5'	-7.66	108.65	120.90
34	BA	1036	G	C5-C6-O6	-7.66	124.01	128.60
34	BA	1107	A	N1-C2-N3	7.66	133.13	129.30
34	BA	1113	A	C5-C6-N6	-7.66	117.57	123.70
35	BB	879	G	C8-N9-C4	-7.66	103.34	106.40
38	BE	8	G	O4'-C4'-C3'	-7.66	96.34	104.00
39	BF	6	C	P-O3'-C3'	7.66	128.89	119.70
34	BA	683	C	P-O5'-C5'	7.66	133.15	120.90
35	BB	1102	U	C3'-C2'-C1'	-7.66	95.38	101.50
36	BC	96	A	N1-C6-N6	-7.66	114.01	118.60
38	BE	109	C	O5'-C5'-C4'	7.66	126.24	111.70
40	BG	8	U	N1-C2-O2	7.66	128.16	122.80
41	BH	4	U	C2-N3-C4	-7.66	122.41	127.00
41	BH	27	A	N1-C6-N6	-7.66	114.01	118.60
66	Bg	49	ARG	NE-CZ-NH2	-7.66	116.47	120.30
85	AA	520	A	C5'-C4'-C3'	7.66	128.25	116.00
85	AA	534	A	C8-N9-C1'	7.66	141.48	127.70
85	AA	1541	G	P-O3'-C3'	-7.66	110.51	119.70
7	A6	157	PHE	CB-CG-CD2	-7.65	115.44	120.80
34	BA	168	U	C6-N1-C1'	7.65	131.91	121.20
36	BC	136	G	O4'-C1'-N9	7.65	114.32	108.20
41	BH	39	G	C5'-C4'-C3'	7.65	128.25	116.00
85	AA	1539	A	C4'-C3'-C2'	-7.65	94.95	102.60
85	AA	1793	A	P-O3'-C3'	-7.65	110.52	119.70
85	AA	2055	G	O4'-C1'-N9	7.65	114.32	108.20
8	A7	105	PHE	CB-CG-CD1	7.65	126.16	120.80
37	BD	51	G	P-O3'-C3'	7.65	128.88	119.70
40	BG	102	G	N3-C2-N2	7.65	125.26	119.90
85	AA	1426	G	P-O3'-C3'	-7.65	110.52	119.70
85	AA	2121	G	N3-C4-C5	-7.65	124.77	128.60
34	BA	333	A	N1-C6-N6	7.65	123.19	118.60
34	BA	535	G	C5-C6-O6	7.65	133.19	128.60
34	BA	537	C	P-O3'-C3'	-7.65	110.52	119.70
34	BA	1295	U	P-O5'-C5'	7.65	133.14	120.90
34	BA	1735	G	C4-N9-C1'	-7.65	116.56	126.50
35	BB	787	A	C4-N9-C1'	-7.65	112.53	126.30
36	BC	35	C	P-O3'-C3'	-7.65	110.52	119.70
38	BE	139	U	C5-C6-N1	-7.65	118.88	122.70
85	AA	479	C	P-O5'-C5'	7.65	133.14	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	486	G	O4'-C1'-N9	7.65	114.32	108.20
85	AA	1368	G	C5'-C4'-C3'	-7.65	103.76	116.00
85	AA	1514	A	N1-C6-N6	7.65	123.19	118.60
85	AA	1795	C	C3'-C2'-C1'	-7.65	95.38	101.50
34	BA	56	G	P-O5'-C5'	7.65	133.14	120.90
34	BA	467	A	C5-C6-N6	-7.65	117.58	123.70
34	BA	783	U	C6-N1-C2	-7.65	116.41	121.00
34	BA	1166	A	C4-N9-C1'	-7.65	112.53	126.30
38	BE	16	C	N3-C4-N4	7.65	123.35	118.00
34	BA	1665	G	C5'-C4'-C3'	-7.65	103.77	116.00
34	BA	1741	G	C3'-C2'-C1'	-7.65	95.38	101.50
34	BA	1846	G	C1'-O4'-C4'	-7.65	103.78	109.90
35	BB	41	A	N1-C6-N6	7.65	123.19	118.60
35	BB	1323	U	C1'-O4'-C4'	-7.65	103.78	109.90
35	BB	1532	C	P-O5'-C5'	-7.65	108.66	120.90
85	AA	1176	C	C2-N3-C4	-7.65	116.08	119.90
85	AA	1811	C	C6-N1-C2	-7.65	117.24	120.30
34	BA	334	G	C1'-O4'-C4'	-7.65	103.78	109.90
34	BA	1246	G	C3'-C2'-C1'	-7.65	95.38	101.50
82	Bw	105	ARG	NE-CZ-NH1	7.65	124.12	120.30
85	AA	1246	G	C5-C6-N1	7.65	115.32	111.50
85	AA	1939	C	O4'-C1'-N1	7.65	114.32	108.20
34	BA	1487	U	C5'-C4'-C3'	7.64	128.23	116.00
71	Bl	141	ARG	CG-CD-NE	-7.64	95.75	111.80
85	AA	1002	G	C8-N9-C1'	7.64	136.94	127.00
85	AA	1215	A	C3'-C2'-C1'	-7.64	95.39	101.50
34	BA	245	U	O4'-C1'-N1	7.64	114.31	108.20
34	BA	1272	U	C2-N1-C1'	7.64	126.87	117.70
34	BA	1296	U	P-O3'-C3'	7.64	128.87	119.70
35	BB	1487	G	O4'-C1'-N9	7.64	114.31	108.20
38	BE	5	A	P-O3'-C3'	7.64	128.87	119.70
85	AA	686	U	C6-N1-C2	-7.64	116.42	121.00
85	AA	1787	G	O4'-C1'-N9	7.64	114.31	108.20
85	AA	572	G	O4'-C1'-N9	7.64	114.31	108.20
85	AA	584	G	C8-N9-C1'	-7.64	117.07	127.00
85	AA	1503	G	C4-N9-C1'	-7.64	116.57	126.50
34	BA	403	A	C5'-C4'-O4'	7.64	118.27	109.10
34	BA	894	G	C8-N9-C4	7.64	109.46	106.40
35	BB	1182	A	C8-N9-C4	7.64	108.86	105.80
35	BB	1471	A	P-O5'-C5'	7.64	133.12	120.90
35	BB	1473	U	O4'-C1'-N1	7.64	114.31	108.20
41	BH	133	U	N1-C2-N3	7.64	119.48	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
83	Bx	73	ARG	NE-CZ-NH1	7.64	124.12	120.30
85	AA	1647	G	C5'-C4'-C3'	-7.64	103.78	116.00
85	AA	1659	C	O4'-C1'-N1	7.64	114.31	108.20
85	AA	2209	U	C5'-C4'-C3'	-7.64	103.78	116.00
34	BA	1304	C	C5-C4-N4	7.64	125.55	120.20
35	BB	1484	A	C5'-C4'-C3'	-7.64	103.78	116.00
37	BD	85	C	P-O3'-C3'	-7.64	110.53	119.70
38	BE	172	U	C4'-C3'-C2'	7.64	110.24	102.60
85	AA	2050	C	C4'-C3'-C2'	-7.64	94.96	102.60
34	BA	400	A	N9-C1'-C2'	-7.64	103.60	112.00
34	BA	843	G	N3-C4-C5	-7.64	124.78	128.60
34	BA	1668	C	C5'-C4'-C3'	-7.64	103.78	116.00
37	BD	65	G	N9-C1'-C2'	-7.64	103.60	112.00
85	AA	681	G	C4'-C3'-C2'	7.64	110.24	102.60
85	AA	961	U	C5'-C4'-O4'	7.64	118.26	109.10
34	BA	532	C	P-O3'-C3'	-7.63	110.54	119.70
34	BA	683	C	C4'-C3'-C2'	-7.63	94.97	102.60
34	BA	743	A	N3-C4-N9	7.63	133.51	127.40
35	BB	367	C	C6-N1-C2	-7.63	117.25	120.30
40	BG	178	G	C4-N9-C1'	-7.63	116.58	126.50
85	AA	20	G	C5-C6-O6	-7.63	124.02	128.60
34	BA	681	G	N9-C4-C5	-7.63	102.35	105.40
34	BA	1398	C	C6-N1-C2	-7.63	117.25	120.30
34	BA	1741	G	P-O3'-C3'	-7.63	110.54	119.70
35	BB	1238	A	C5'-C4'-C3'	-7.63	103.79	116.00
35	BB	1298	C	N3-C4-N4	-7.63	112.66	118.00
37	BD	92	G	C4'-C3'-C2'	-7.63	94.97	102.60
34	BA	614	A	P-O5'-C5'	7.63	133.11	120.90
34	BA	1562	G	P-O5'-C5'	-7.63	108.69	120.90
35	BB	152	G	N1-C6-O6	7.63	124.48	119.90
35	BB	779	C	C4'-C3'-C2'	7.63	110.23	102.60
41	BH	2	U	O4'-C4'-C3'	-7.63	96.37	104.00
34	BA	478	G	N3-C2-N2	7.63	125.24	119.90
38	BE	201	A	P-O3'-C3'	-7.63	110.54	119.70
77	Br	275	PHE	N-CA-CB	-7.63	96.87	110.60
85	AA	1830	U	P-O3'-C3'	-7.63	110.54	119.70
34	BA	372	U	N3-C2-O2	-7.63	116.86	122.20
34	BA	572	G	C4-C5-N7	-7.63	107.75	110.80
34	BA	597	C	C2-N1-C1'	7.63	127.19	118.80
34	BA	1846	G	O5'-P-OP2	7.63	119.85	110.70
38	BE	148	C	C5'-C4'-C3'	-7.63	103.80	116.00
85	AA	57	G	N1-C2-N2	-7.63	109.33	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	197	C	O4'-C1'-N1	7.63	114.30	108.20
34	BA	487	A	O4'-C1'-N9	7.63	114.30	108.20
34	BA	863	G	N1-C6-O6	-7.63	115.33	119.90
34	BA	1474	G	N9-C1'-C2'	-7.63	103.61	112.00
35	BB	825	U	P-O5'-C5'	-7.63	108.70	120.90
85	AA	270	A	C4-N9-C1'	-7.63	112.57	126.30
85	AA	637	U	N1-C2-N3	7.63	119.48	114.90
34	BA	740	A	N1-C2-N3	7.62	133.11	129.30
35	BB	589	U	C6-N1-C2	-7.62	116.42	121.00
35	BB	901	U	C1'-O4'-C4'	-7.62	103.80	109.90
85	AA	1938	G	C8-N9-C1'	7.62	136.91	127.00
34	BA	121	A	C8-N9-C4	7.62	108.85	105.80
34	BA	178	C	C5'-C4'-O4'	-7.62	99.95	109.10
34	BA	1272	U	P-O3'-C3'	-7.62	110.55	119.70
34	BA	1649	A	C8-N9-C4	7.62	108.85	105.80
35	BB	434	A	C8-N9-C4	7.62	108.85	105.80
35	BB	1524	G	C8-N9-C1'	7.62	136.91	127.00
38	BE	94	U	C5'-C4'-C3'	7.62	128.20	116.00
52	BS	45	ARG	NE-CZ-NH1	7.62	124.11	120.30
85	AA	252	G	C2-N3-C4	-7.62	108.09	111.90
85	AA	1055	U	P-O5'-C5'	7.62	133.10	120.90
85	AA	1283	C	N1-C2-O2	7.62	123.47	118.90
85	AA	2166	G	O4'-C1'-N9	7.62	114.30	108.20
5	A4	191	MET	CG-SD-CE	-7.62	88.00	100.20
34	BA	1551	G	C5'-C4'-C3'	7.62	128.19	116.00
34	BA	1628	A	P-O3'-C3'	7.62	128.85	119.70
34	BA	1695	G	C3'-C2'-C1'	-7.62	95.40	101.50
35	BB	868	C	C5-C6-N1	7.62	124.81	121.00
37	BD	77	A	C8-N9-C4	7.62	108.85	105.80
40	BG	140	G	C5-C6-O6	-7.62	124.03	128.60
41	BH	29	G	C6-N1-C2	-7.62	120.53	125.10
85	AA	1822	G	C4-N9-C1'	-7.62	116.59	126.50
38	BE	109	C	O4'-C4'-C3'	-7.62	96.38	104.00
82	Bw	210	PHE	N-CA-CB	-7.62	96.88	110.60
34	BA	613	A	C8-N9-C4	7.62	108.85	105.80
34	BA	1175	G	C5'-C4'-C3'	-7.62	103.81	116.00
34	BA	1748	G	C5-C6-O6	-7.62	124.03	128.60
38	BE	27	A	C3'-C2'-C1'	-7.62	95.41	101.50
38	BE	66	A	N1-C6-N6	7.62	123.17	118.60
85	AA	944	C	C5-C4-N4	-7.62	114.87	120.20
85	AA	2179	C	N3-C2-O2	-7.62	116.57	121.90
6	A5	42	ARG	N-CA-C	-7.62	90.43	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	894	G	C5'-C4'-C3'	7.62	128.19	116.00
85	AA	2209	U	C6-N1-C1'	7.62	131.86	121.20
34	BA	212	A	C4-N9-C1'	-7.62	112.59	126.30
34	BA	1738	G	O4'-C1'-N9	7.62	114.29	108.20
35	BB	449	C	O4'-C1'-N1	7.62	114.29	108.20
35	BB	827	U	P-O3'-C3'	-7.62	110.56	119.70
35	BB	1140	C	O4'-C1'-N1	7.62	114.29	108.20
34	BA	78	U	C2-N3-C4	-7.61	122.43	127.00
34	BA	127	U	C6-N1-C1'	7.61	131.86	121.20
34	BA	482	C	C2'-C3'-O3'	7.61	126.25	109.50
34	BA	575	U	N1-C2-N3	-7.61	110.33	114.90
34	BA	1677	C	O4'-C1'-N1	7.61	114.29	108.20
35	BB	51	U	C3'-C2'-C1'	-7.61	95.41	101.50
35	BB	493	U	C5'-C4'-C3'	-7.61	103.82	116.00
85	AA	82	A	P-O5'-C5'	-7.61	108.72	120.90
85	AA	1173	A	N1-C6-N6	7.61	123.17	118.60
34	BA	86	A	O5'-P-OP2	-7.61	98.85	105.70
34	BA	959	G	N1-C6-O6	7.61	124.47	119.90
36	BC	62	A	C8-N9-C4	7.61	108.84	105.80
85	AA	84	C	P-O3'-C3'	-7.61	110.57	119.70
85	AA	413	G	C8-N9-C1'	7.61	136.90	127.00
85	AA	982	G	C5-C6-O6	-7.61	124.03	128.60
34	BA	470	C	O3'-P-O5'	-7.61	89.54	104.00
34	BA	1445	U	C2-N1-C1'	7.61	126.83	117.70
38	BE	1	U	C4'-C3'-C2'	-7.61	94.99	102.60
52	BS	120	TYR	N-CA-CB	-7.61	96.90	110.60
85	AA	927	A	O4'-C1'-N9	7.61	114.29	108.20
34	BA	1513	G	N3-C4-C5	-7.61	124.80	128.60
85	AA	9	U	P-O5'-C5'	-7.61	108.72	120.90
85	AA	327	G	C5'-C4'-C3'	7.61	128.18	116.00
85	AA	1432	C	C5-C4-N4	-7.61	114.87	120.20
85	AA	1644	G	N1-C6-O6	7.61	124.47	119.90
85	AA	2155	U	N3-C2-O2	-7.61	116.87	122.20
34	BA	220	U	P-O3'-C3'	-7.61	110.57	119.70
34	BA	1510	C	N3-C2-O2	-7.61	116.57	121.90
34	BA	1794	A	N9-C1'-C2'	-7.61	103.63	112.00
65	Bf	349	PRO	CA-N-CD	-7.61	100.85	111.50
85	AA	186	U	C3'-C2'-C1'	-7.61	95.41	101.50
35	BB	1334	C	O4'-C4'-C3'	-7.61	96.39	104.00
85	AA	525	C	C6-N1-C2	-7.61	117.26	120.30
85	AA	1463	A	C6-N1-C2	-7.61	114.04	118.60
85	AA	1864	G	C3'-C2'-C1'	-7.60	95.42	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	629	G	O4'-C1'-N9	7.60	114.28	108.20
34	BA	1355	G	O4'-C1'-N9	7.60	114.28	108.20
34	BA	1580	U	C5-C4-O4	-7.60	121.34	125.90
34	BA	1581	G	C5-C6-N1	7.60	115.30	111.50
35	BB	857	G	C1'-O4'-C4'	-7.60	103.82	109.90
35	BB	1197	G	N1-C6-O6	7.60	124.46	119.90
35	BB	1309	A	P-O3'-C3'	-7.60	110.58	119.70
38	BE	87	U	C1'-O4'-C4'	-7.60	103.82	109.90
39	BF	20	U	C2-N3-C4	-7.60	122.44	127.00
40	BG	175	G	C5'-C4'-C3'	7.60	128.16	116.00
85	AA	700	U	C2-N3-C4	-7.60	122.44	127.00
85	AA	2241	C	N3-C4-N4	-7.60	112.68	118.00
34	BA	1210	A	C1'-O4'-C4'	-7.60	103.82	109.90
39	BF	2	G	C6-C5-N7	-7.60	125.84	130.40
85	AA	1928	A	C4'-C3'-C2'	7.60	110.20	102.60
26	AS	18	ARG	NE-CZ-NH2	7.60	124.10	120.30
34	BA	329	G	C5-C6-N1	7.60	115.30	111.50
35	BB	1500	U	C2-N1-C1'	-7.60	108.58	117.70
40	BG	9	G	N9-C1'-C2'	-7.60	103.64	112.00
40	BG	130	G	C8-N9-C4	7.60	109.44	106.40
85	AA	28	A	O4'-C1'-N9	7.60	114.28	108.20
85	AA	275	A	O4'-C1'-N9	7.60	114.28	108.20
85	AA	1955	U	P-O5'-C5'	7.60	133.06	120.90
85	AA	1960	C	O4'-C1'-N1	7.60	114.28	108.20
86	AB	68	C	N3-C2-O2	-7.60	116.58	121.90
34	BA	543	A	N9-C1'-C2'	-7.60	103.64	112.00
34	BA	843	G	O4'-C1'-N9	7.60	114.28	108.20
35	BB	736	G	O4'-C1'-N9	7.60	114.28	108.20
35	BB	879	G	C5'-C4'-C3'	7.60	128.16	116.00
35	BB	1475	U	C5-C6-N1	-7.60	118.90	122.70
38	BE	54	U	C2-N3-C4	-7.60	122.44	127.00
85	AA	882	C	C2-N1-C1'	-7.60	110.44	118.80
34	BA	1005	C	O4'-C1'-N1	7.60	114.28	108.20
35	BB	438	G	C1'-O4'-C4'	-7.60	103.82	109.90
35	BB	1164	U	N1-C2-N3	7.60	119.46	114.90
41	BH	79	A	OP1-P-OP2	-7.60	108.21	119.60
85	AA	2010	C	C4'-C3'-C2'	-7.60	95.00	102.60
34	BA	470	C	O5'-C5'-C4'	-7.59	97.27	111.70
34	BA	519	G	P-O3'-C3'	-7.59	110.59	119.70
34	BA	1149	C	O4'-C1'-N1	7.59	114.28	108.20
34	BA	1722	U	C6-N1-C2	-7.59	116.44	121.00
37	BD	106	G	C8-N9-C1'	7.59	136.87	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	94	G	C4'-C3'-C2'	7.59	110.19	102.60
63	Bd	8	THR	C-N-CA	7.59	140.69	121.70
85	AA	810	C	O4'-C1'-N1	7.59	114.28	108.20
85	AA	1102	C	O4'-C1'-N1	7.59	114.28	108.20
85	AA	1426	G	C4'-C3'-C2'	-7.59	95.01	102.60
85	AA	1438	C	P-O3'-C3'	-7.59	110.59	119.70
34	BA	612	U	C2-N3-C4	-7.59	122.44	127.00
35	BB	512	C	C5'-C4'-C3'	-7.59	103.85	116.00
39	BF	34	C	O4'-C1'-N1	7.59	114.27	108.20
85	AA	2186	U	O4'-C1'-N1	7.59	114.27	108.20
20	AL	36	MET	CG-SD-CE	-7.59	88.05	100.20
34	BA	140	C	O4'-C1'-N1	7.59	114.27	108.20
34	BA	270	U	C5'-C4'-O4'	7.59	118.21	109.10
34	BA	621	G	C4-N9-C1'	-7.59	116.63	126.50
34	BA	1599	A	P-O5'-C5'	7.59	133.05	120.90
85	AA	708	G	C4-C5-C6	-7.59	114.25	118.80
85	AA	802	A	C5-C6-N6	-7.59	117.63	123.70
85	AA	1287	C	C5'-C4'-O4'	7.59	118.21	109.10
85	AA	1666	U	O4'-C1'-N1	7.59	114.27	108.20
85	AA	1835	U	C1'-O4'-C4'	-7.59	103.83	109.90
85	AA	2121	G	C2'-C3'-O3'	7.59	126.20	109.50
35	BB	119	G	N9-C1'-C2'	-7.59	103.65	112.00
35	BB	1469	A	C8-N9-C4	7.59	108.84	105.80
40	BG	144	G	O4'-C1'-N9	7.59	114.27	108.20
51	BR	50	GLN	N-CA-C	7.59	131.49	111.00
85	AA	296	A	N1-C6-N6	-7.59	114.05	118.60
85	AA	423	G	C1'-O4'-C4'	-7.59	103.83	109.90
85	AA	2217	A	N1-C6-N6	7.59	123.15	118.60
34	BA	1116	G	P-O5'-C5'	7.59	133.04	120.90
35	BB	1206	G	P-O5'-C5'	7.59	133.04	120.90
85	AA	450	A	C3'-C2'-C1'	-7.59	95.43	101.50
34	BA	141	G	C2'-C3'-O3'	7.59	126.19	109.50
34	BA	655	U	N3-C2-O2	-7.59	116.89	122.20
34	BA	1540	C	P-O3'-C3'	-7.59	110.60	119.70
34	BA	1613	G	N9-C1'-C2'	-7.59	103.66	112.00
35	BB	710	A	O4'-C1'-N9	7.59	114.27	108.20
35	BB	1150	A	C5-C6-N6	7.59	129.77	123.70
40	BG	149	U	O4'-C1'-C2'	7.59	114.43	107.60
85	AA	1603	G	P-O5'-C5'	-7.59	108.76	120.90
34	BA	48	C	N1-C1'-C2'	-7.58	103.66	112.00
34	BA	1171	C	C1'-O4'-C4'	-7.58	103.83	109.90
35	BB	429	C	O4'-C1'-N1	7.58	114.27	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	BU	39	TYR	CB-CG-CD2	-7.58	116.45	121.00
85	AA	1982	C	O4'-C1'-N1	7.58	114.27	108.20
14	AF	41	ARG	NE-CZ-NH1	7.58	124.09	120.30
34	BA	19	G	C4-N9-C1'	-7.58	116.64	126.50
34	BA	492	G	N9-C1'-C2'	-7.58	103.66	112.00
34	BA	1688	G	N1-C6-O6	7.58	124.45	119.90
35	BB	40	C	N3-C2-O2	-7.58	116.59	121.90
35	BB	368	C	O4'-C1'-N1	7.58	114.27	108.20
38	BE	125	C	C3'-C2'-C1'	-7.58	95.43	101.50
85	AA	179	G	P-O5'-C5'	7.58	133.03	120.90
85	AA	289	G	N1-C6-O6	7.58	124.45	119.90
85	AA	1188	A	P-O3'-C3'	-7.58	110.60	119.70
85	AA	1212	C	O4'-C1'-N1	7.58	114.27	108.20
34	BA	145	U	C2-N3-C4	-7.58	122.45	127.00
34	BA	177	G	C8-N9-C4	7.58	109.43	106.40
35	BB	843	G	P-O3'-C3'	-7.58	110.60	119.70
35	BB	1259	A	O5'-C5'-C4'	7.58	126.11	111.70
37	BD	119	U	P-O5'-C5'	-7.58	108.77	120.90
40	BG	159	A	C6-N1-C2	-7.58	114.05	118.60
41	BH	40	C	O4'-C1'-N1	7.58	114.27	108.20
85	AA	1757	C	O4'-C1'-N1	7.58	114.27	108.20
85	AA	1902	C	O4'-C1'-N1	7.58	114.27	108.20
85	AA	2010	C	O4'-C1'-N1	7.58	114.27	108.20
34	BA	899	G	P-O5'-C5'	7.58	133.03	120.90
34	BA	966	G	C5-C6-O6	7.58	133.15	128.60
35	BB	1314	G	C5'-C4'-C3'	-7.58	103.87	116.00
61	Bb	42	ARG	NE-CZ-NH1	7.58	124.09	120.30
65	Bf	280	HIS	CB-CA-C	-7.58	95.24	110.40
85	AA	385	A	C5'-C4'-O4'	7.58	118.20	109.10
85	AA	2103	C	O4'-C1'-N1	7.58	114.26	108.20
34	BA	490	A	N9-C1'-C2'	-7.58	103.66	112.00
34	BA	875	G	N7-C8-N9	-7.58	109.31	113.10
34	BA	877	U	C4'-C3'-C2'	7.58	110.18	102.60
34	BA	1612	C	P-O3'-C3'	-7.58	110.61	119.70
85	AA	1146	C	C6-N1-C2	-7.58	117.27	120.30
85	AA	1497	U	C5'-C4'-C3'	7.58	128.13	116.00
34	BA	158	U	O4'-C1'-N1	7.58	114.26	108.20
34	BA	587	U	C2'-C3'-O3'	7.58	126.17	109.50
35	BB	263	C	O4'-C1'-N1	7.58	114.26	108.20
84	By	173	PHE	CB-CG-CD2	-7.58	115.50	120.80
34	BA	829	U	O4'-C1'-N1	7.58	114.26	108.20
34	BA	1646	U	N3-C2-O2	-7.58	116.90	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1655	G	C5'-C4'-C3'	7.58	128.12	116.00
35	BB	1489	A	C2'-C3'-O3'	7.58	126.16	109.50
36	BC	12	A	C5-C6-N1	7.58	121.49	117.70
36	BC	90	U	O4'-C4'-C3'	-7.58	96.42	104.00
41	BH	72	G	C5-C6-O6	-7.58	124.06	128.60
85	AA	717	G	C5'-C4'-C3'	-7.58	103.88	116.00
85	AA	1465	C	N3-C2-O2	-7.58	116.60	121.90
85	AA	1670	U	C2-N1-C1'	-7.58	108.61	117.70
34	BA	852	C	N3-C4-N4	-7.57	112.70	118.00
34	BA	1676	A	O4'-C4'-C3'	-7.57	96.43	104.00
35	BB	1161	G	P-O3'-C3'	7.57	128.79	119.70
35	BB	653	G	P-O5'-C5'	7.57	133.01	120.90
39	BF	16	C	C2-N1-C1'	7.57	127.13	118.80
85	AA	1436	A	N1-C6-N6	-7.57	114.06	118.60
34	BA	471	U	C4-C5-C6	-7.57	115.16	119.70
34	BA	785	G	C4'-C3'-C2'	-7.57	95.03	102.60
34	BA	835	U	P-O5'-C5'	-7.57	108.79	120.90
34	BA	1272	U	C6-N1-C1'	-7.57	110.60	121.20
85	AA	345	U	O4'-C1'-N1	7.57	114.26	108.20
85	AA	392	G	C8-N9-C1'	7.57	136.84	127.00
34	BA	1445	U	C1'-O4'-C4'	-7.57	103.84	109.90
37	BD	48	G	C1'-O4'-C4'	-7.57	103.84	109.90
85	AA	469	G	N9-C1'-C2'	-7.57	103.67	112.00
34	BA	197	A	O4'-C1'-N9	7.57	114.25	108.20
34	BA	279	U	C3'-C2'-C1'	-7.57	95.45	101.50
34	BA	496	G	C4-N9-C1'	-7.57	116.66	126.50
85	AA	80	G	C8-N9-C1'	7.57	136.84	127.00
85	AA	1475	A	P-O3'-C3'	7.57	128.78	119.70
85	AA	1997	G	O3'-P-O5'	7.57	118.38	104.00
85	AA	2238	C	C2-N3-C4	-7.57	116.12	119.90
31	AX	155	TYR	CB-CG-CD2	-7.57	116.46	121.00
33	AZ	60	ARG	NE-CZ-NH2	-7.57	116.52	120.30
34	BA	395	G	C1'-O4'-C4'	-7.57	103.85	109.90
34	BA	782	C	N3-C4-N4	7.57	123.30	118.00
35	BB	1442	C	C4'-C3'-C2'	7.57	110.17	102.60
70	Bk	53	LYS	CB-CA-C	7.57	125.53	110.40
85	AA	76	G	O4'-C1'-N9	7.57	114.25	108.20
85	AA	272	C	C5'-C4'-C3'	-7.57	103.89	116.00
85	AA	1518	A	C8-N9-C4	7.57	108.83	105.80
85	AA	2187	G	P-O3'-C3'	-7.57	110.62	119.70
85	AA	2200	A	C8-N9-C4	7.57	108.83	105.80
35	BB	684	U	C6-N1-C1'	7.56	131.79	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	178	C	C2-N1-C1'	-7.56	110.48	118.80
34	BA	367	G	P-O3'-C3'	-7.56	110.62	119.70
34	BA	513	U	N3-C4-O4	-7.56	114.11	119.40
34	BA	1175	G	P-O3'-C3'	-7.56	110.62	119.70
35	BB	392	G	C5-C6-N1	7.56	115.28	111.50
35	BB	1022	C	O4'-C1'-N1	7.56	114.25	108.20
37	BD	81	C	C3'-C2'-C1'	-7.56	95.45	101.50
85	AA	37	U	P-O3'-C3'	-7.56	110.62	119.70
85	AA	1176	C	P-O3'-C3'	-7.56	110.62	119.70
86	AB	43	C	O4'-C1'-N1	7.56	114.25	108.20
34	BA	1552	C	C6-N1-C2	-7.56	117.28	120.30
85	AA	2094	U	P-O5'-C5'	-7.56	108.80	120.90
34	BA	694	G	N3-C2-N2	7.56	125.19	119.90
34	BA	718	U	O5'-P-OP1	-7.56	98.90	105.70
34	BA	1471	U	C2-N1-C1'	7.56	126.77	117.70
34	BA	1591	G	C8-N9-C4	7.56	109.42	106.40
85	AA	57	G	N3-C2-N2	7.56	125.19	119.90
85	AA	1092	G	C8-N9-C1'	7.56	136.83	127.00
85	AA	1626	U	P-O3'-C3'	-7.56	110.63	119.70
34	BA	180	G	C6-N1-C2	-7.56	120.56	125.10
34	BA	194	G	C5-C6-O6	7.56	133.13	128.60
34	BA	684	G	P-O3'-C3'	7.56	128.77	119.70
34	BA	720	A	P-O3'-C3'	-7.56	110.63	119.70
37	BD	91	U	N1-C2-N3	7.56	119.43	114.90
40	BG	110	U	C4'-C3'-C2'	7.56	110.16	102.60
52	BS	165	ARG	NE-CZ-NH2	-7.56	116.52	120.30
85	AA	966	G	C6-N1-C2	-7.56	120.57	125.10
35	BB	836	U	C2-N3-C4	-7.56	122.47	127.00
41	BH	47	G	C5-C6-O6	7.56	133.13	128.60
59	BZ	22	HIS	CA-CB-CG	7.56	126.44	113.60
85	AA	820	G	N3-C4-C5	-7.56	124.82	128.60
85	AA	856	G	C5'-C4'-C3'	7.56	128.09	116.00
34	BA	1	C	O3'-P-O5'	-7.55	89.65	104.00
34	BA	141	G	P-O5'-C5'	-7.55	108.81	120.90
34	BA	1226	G	C8-N9-C1'	7.55	136.82	127.00
35	BB	1367	U	N3-C2-O2	-7.55	116.91	122.20
41	BH	105	U	N3-C4-C5	7.55	119.13	114.60
53	BT	100	ARG	CB-CA-C	7.55	125.51	110.40
85	AA	1892	G	C4-N9-C1'	-7.55	116.68	126.50
35	BB	638	G	N1-C6-O6	7.55	124.43	119.90
35	BB	987	U	C6-N1-C1'	-7.55	110.63	121.20
35	BB	1071	G	C5'-C4'-C3'	-7.55	103.92	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1127	A	N1-C6-N6	-7.55	114.07	118.60
34	BA	420	A	C8-N9-C4	7.55	108.82	105.80
34	BA	1232	C	N3-C2-O2	-7.55	116.61	121.90
34	BA	1547	G	C5'-C4'-C3'	-7.55	103.92	116.00
34	BA	1635	A	O5'-P-OP1	-7.55	98.90	105.70
37	BD	93	G	C4-N9-C1'	-7.55	116.68	126.50
85	AA	2217	A	P-O3'-C3'	-7.55	110.64	119.70
34	BA	22	C	P-O3'-C3'	-7.55	110.64	119.70
34	BA	638	U	C5-C4-O4	7.55	130.43	125.90
34	BA	1530	G	C8-N9-C1'	7.55	136.81	127.00
35	BB	1161	G	N3-C2-N2	-7.55	114.62	119.90
35	BB	1512	C	C5-C6-N1	7.55	124.77	121.00
40	BG	23	C	C6-N1-C2	-7.55	117.28	120.30
41	BH	35	G	N7-C8-N9	7.55	116.88	113.10
49	BP	114	ARG	NE-CZ-NH2	-7.55	116.53	120.30
85	AA	157	G	C4'-C3'-C2'	-7.55	95.05	102.60
85	AA	160	A	O4'-C1'-N9	7.55	114.24	108.20
85	AA	198	U	C2-N3-C4	-7.55	122.47	127.00
85	AA	1564	U	C5'-C4'-C3'	7.55	128.08	116.00
34	BA	349	G	P-O5'-C5'	-7.55	108.82	120.90
34	BA	777	C	O3'-P-O5'	7.55	118.34	104.00
34	BA	1059	U	P-O5'-C5'	-7.55	108.82	120.90
34	BA	1397	C	O4'-C1'-N1	7.55	114.24	108.20
38	BE	126	G	C8-N9-C1'	7.55	136.81	127.00
85	AA	121	C	C5'-C4'-O4'	-7.55	100.04	109.10
34	BA	1176	C	C5'-C4'-C3'	-7.55	103.93	116.00
35	BB	847	U	P-O3'-C3'	-7.55	110.64	119.70
35	BB	1314	G	N3-C2-N2	7.55	125.18	119.90
85	AA	741	G	N1-C2-N2	-7.55	109.41	116.20
85	AA	787	U	C5'-C4'-O4'	7.55	118.16	109.10
41	BH	39	G	C6-N1-C2	-7.54	120.57	125.10
85	AA	995	G	N9-C1'-C2'	-7.54	103.70	112.00
34	BA	296	G	N9-C1'-C2'	7.54	123.81	114.00
34	BA	357	A	P-O5'-C5'	7.54	132.97	120.90
34	BA	514	U	C5'-C4'-O4'	7.54	118.15	109.10
34	BA	1023	G	P-O3'-C3'	-7.54	110.65	119.70
38	BE	65	U	O4'-C1'-N1	7.54	114.23	108.20
85	AA	628	C	O4'-C1'-N1	7.54	114.23	108.20
85	AA	867	G	C4'-C3'-C2'	-7.54	95.06	102.60
34	BA	1642	A	P-O5'-C5'	-7.54	108.83	120.90
34	BA	1699	A	C5'-C4'-C3'	7.54	128.07	116.00
35	BB	374	A	C5-C6-N1	7.54	121.47	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1137	G	P-O3'-C3'	-7.54	110.65	119.70
36	BC	65	G	C5'-C4'-C3'	-7.54	103.93	116.00
39	BF	57	C	C1'-O4'-C4'	-7.54	103.87	109.90
40	BG	46	G	C4-N9-C1'	-7.54	116.69	126.50
41	BH	48	G	C6-N1-C2	-7.54	120.58	125.10
85	AA	129	U	P-O3'-C3'	-7.54	110.65	119.70
34	BA	1846	G	C3'-C2'-C1'	-7.54	95.47	101.50
34	BA	1046	G	C5'-C4'-C3'	7.54	128.06	116.00
35	BB	897	C	C1'-O4'-C4'	-7.54	103.87	109.90
35	BB	1231	U	O5'-C5'-C4'	7.54	126.02	111.70
38	BE	37	C	O4'-C1'-N1	7.54	114.23	108.20
85	AA	1580	A	P-O3'-C3'	7.54	128.75	119.70
86	AB	20	U	O4'-C1'-N1	7.54	114.23	108.20
34	BA	436	U	C5'-C4'-C3'	-7.54	103.94	116.00
34	BA	1491	U	P-O3'-C3'	-7.54	110.66	119.70
34	BA	1514	A	N1-C6-N6	-7.54	114.08	118.60
35	BB	830	G	P-O5'-C5'	-7.54	108.84	120.90
35	BB	831	C	C1'-O4'-C4'	-7.54	103.87	109.90
35	BB	1425	A	P-O3'-C3'	-7.54	110.66	119.70
35	BB	1480	G	N3-C4-N9	-7.54	121.48	126.00
85	AA	713	G	N9-C1'-C2'	-7.54	103.71	112.00
34	BA	1067	G	C8-N9-C1'	7.53	136.79	127.00
34	BA	1489	U	C4'-C3'-C2'	-7.53	95.07	102.60
34	BA	1524	G	C5-C6-N1	7.53	115.27	111.50
35	BB	499	A	N1-C6-N6	-7.53	114.08	118.60
35	BB	652	G	P-O5'-C5'	7.53	132.95	120.90
35	BB	1000	U	P-O3'-C3'	-7.53	110.66	119.70
35	BB	1159	U	C2-N1-C1'	-7.53	108.66	117.70
35	BB	1169	A	P-O3'-C3'	-7.53	110.66	119.70
85	AA	1235	G	N3-C4-C5	-7.53	124.83	128.60
85	AA	1355	U	P-O5'-C5'	7.53	132.96	120.90
85	AA	1531	G	C4-N9-C1'	-7.53	116.71	126.50
34	BA	631	G	N1-C2-N2	7.53	122.98	116.20
34	BA	1845	G	N3-C4-N9	-7.53	121.48	126.00
35	BB	423	G	O4'-C1'-N9	7.53	114.22	108.20
35	BB	1404	A	P-O3'-C3'	-7.53	110.66	119.70
85	AA	4	C	C5'-C4'-O4'	7.53	118.14	109.10
85	AA	1217	U	O4'-C1'-N1	7.53	114.23	108.20
85	AA	1273	C	O4'-C1'-N1	7.53	114.22	108.20
85	AA	1637	C	C6-N1-C2	-7.53	117.29	120.30
34	BA	1723	U	N3-C2-O2	-7.53	116.93	122.20
35	BB	977	G	O4'-C1'-N9	7.53	114.22	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	80	G	C8-N9-C1'	7.53	136.79	127.00
64	Be	80	GLU	N-CA-C	7.53	131.33	111.00
85	AA	1006	C	N3-C2-O2	-7.53	116.63	121.90
40	BG	4	A	O4'-C1'-N9	7.53	114.22	108.20
80	Bu	15	ARG	NE-CZ-NH1	7.53	124.06	120.30
34	BA	416	A	C5-C6-N6	7.53	129.72	123.70
34	BA	1842	U	C5'-C4'-C3'	-7.53	103.96	116.00
35	BB	5	A	C5-C6-N1	7.53	121.46	117.70
36	BC	4	G	C8-N9-C1'	7.53	136.79	127.00
36	BC	123	G	C8-N9-C1'	7.53	136.79	127.00
36	BC	146	U	O4'-C1'-N1	7.53	114.22	108.20
40	BG	67	A	C5'-C4'-C3'	-7.53	103.95	116.00
41	BH	70	U	O4'-C1'-N1	7.53	114.22	108.20
49	BP	106	THR	N-CA-CB	7.53	124.60	110.30
34	BA	53	G	N7-C8-N9	-7.53	109.34	113.10
34	BA	182	U	N3-C2-O2	-7.53	116.93	122.20
34	BA	266	G	N1-C2-N2	-7.53	109.43	116.20
35	BB	775	U	N3-C2-O2	-7.53	116.93	122.20
37	BD	80	G	C1'-O4'-C4'	-7.53	103.88	109.90
40	BG	88	G	C5-C6-N1	7.53	115.26	111.50
41	BH	92	A	N1-C6-N6	7.53	123.11	118.60
85	AA	140	C	O4'-C1'-N1	7.53	114.22	108.20
85	AA	749	C	O4'-C1'-N1	7.53	114.22	108.20
35	BB	1004	A	C3'-C2'-C1'	-7.52	95.48	101.50
38	BE	56	U	C4'-C3'-C2'	-7.52	95.08	102.60
23	AP	220	PHE	CB-CG-CD2	-7.52	115.53	120.80
34	BA	1502	G	N3-C2-N2	7.52	125.17	119.90
35	BB	431	U	C2'-C3'-O3'	7.52	126.05	109.50
35	BB	657	A	C5-C6-N6	7.52	129.72	123.70
35	BB	889	U	O4'-C1'-N1	7.52	114.22	108.20
37	BD	46	G	C5'-C4'-C3'	-7.52	103.96	116.00
40	BG	128	U	C5'-C4'-C3'	7.52	128.04	116.00
85	AA	5	U	O5'-P-OP2	-7.52	98.93	105.70
85	AA	577	U	C5'-C4'-C3'	-7.52	103.97	116.00
85	AA	1464	G	C5-C6-O6	-7.52	124.09	128.60
34	BA	211	C	C2-N1-C1'	-7.52	110.53	118.80
35	BB	793	A	C5'-C4'-C3'	-7.52	103.97	116.00
35	BB	1461	C	C5'-C4'-C3'	7.52	128.03	116.00
64	Be	190	ARG	NE-CZ-NH1	7.52	124.06	120.30
85	AA	177	A	C8-N9-C4	7.52	108.81	105.80
85	AA	1812	C	C2-N1-C1'	-7.52	110.53	118.80
34	BA	566	G	C3'-C2'-C1'	-7.52	95.48	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	678	C	N3-C2-O2	-7.52	116.64	121.90
34	BA	1073	G	C5-C6-O6	-7.52	124.09	128.60
35	BB	381	C	O4'-C1'-N1	7.52	114.22	108.20
41	BH	97	C	O5'-P-OP2	7.52	119.72	110.70
80	Bu	186	PHE	CB-CG-CD1	-7.52	115.54	120.80
85	AA	131	C	C5'-C4'-C3'	-7.52	103.97	116.00
85	AA	432	A	C4-N9-C1'	-7.52	112.76	126.30
85	AA	466	A	N1-C6-N6	7.52	123.11	118.60
4	A3	31	TYR	CB-CG-CD2	-7.52	116.49	121.00
35	BB	1281	G	N9-C1'-C2'	-7.52	103.73	112.00
85	AA	238	C	O4'-C1'-N1	7.52	114.22	108.20
85	AA	241	U	C5'-C4'-C3'	-7.52	103.97	116.00
85	AA	369	A	C8-N9-C4	7.52	108.81	105.80
85	AA	1734	A	P-O3'-C3'	7.52	128.72	119.70
35	BB	299	U	O4'-C1'-N1	7.52	114.21	108.20
35	BB	886	G	C5-C6-O6	-7.52	124.09	128.60
40	BG	141	A	N1-C6-N6	7.52	123.11	118.60
34	BA	55	G	C5-C6-O6	-7.51	124.09	128.60
34	BA	1742	G	C5'-C4'-C3'	7.51	128.02	116.00
35	BB	424	U	C4'-C3'-C2'	-7.51	95.08	102.60
35	BB	1482	A	O3'-P-O5'	-7.51	89.72	104.00
72	Bm	38	TYR	CB-CG-CD2	-7.51	116.49	121.00
85	AA	579	U	O4'-C1'-N1	7.51	114.21	108.20
85	AA	769	C	P-O5'-C5'	7.51	132.92	120.90
85	AA	1731	G	P-O5'-C5'	7.51	132.92	120.90
85	AA	1991	C	C2-N1-C1'	7.51	127.07	118.80
34	BA	1240	G	N1-C6-O6	7.51	124.41	119.90
35	BB	654	C	P-O5'-C5'	-7.51	108.88	120.90
85	AA	18	C	O3'-P-O5'	7.51	118.27	104.00
15	AG	105	ASN	C-N-CA	7.51	140.48	121.70
35	BB	576	A	N1-C6-N6	-7.51	114.09	118.60
40	BG	180	C	C5'-C4'-O4'	7.51	118.11	109.10
56	BW	73	ARG	CA-C-N	-7.51	100.67	117.20
61	Bb	9	ARG	NE-CZ-NH1	7.51	124.06	120.30
85	AA	304	G	P-O3'-C3'	7.51	128.71	119.70
85	AA	1923	A	P-O3'-C3'	7.51	128.71	119.70
34	BA	742	C	N3-C4-C5	-7.51	118.90	121.90
34	BA	1357	C	O4'-C1'-N1	7.51	114.21	108.20
34	BA	1792	U	C2-N3-C4	-7.51	122.49	127.00
35	BB	592	G	O4'-C1'-N9	7.51	114.21	108.20
35	BB	1491	G	O4'-C1'-N9	7.51	114.21	108.20
36	BC	78	G	C5-C6-N1	7.51	115.25	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	88	A	C8-N9-C1'	7.51	141.22	127.70
37	BD	69	U	C2-N1-C1'	-7.51	108.69	117.70
85	AA	15	U	C2-N1-C1'	-7.51	108.69	117.70
85	AA	1020	C	P-O5'-C5'	-7.51	108.89	120.90
85	AA	1496	U	O5'-P-OP2	-7.51	98.94	105.70
85	AA	1719	C	C1'-O4'-C4'	-7.51	103.89	109.90
38	BE	16	C	O4'-C1'-N1	7.51	114.21	108.20
85	AA	1449	C	C4'-C3'-C2'	-7.51	95.09	102.60
3	A2	132	ARG	NE-CZ-NH1	7.51	124.05	120.30
34	BA	2	A	C4-N9-C1'	-7.51	112.79	126.30
34	BA	590	U	C4-C5-C6	-7.51	115.20	119.70
34	BA	748	C	C6-N1-C1'	7.51	129.81	120.80
34	BA	1705	C	O4'-C1'-N1	7.51	114.20	108.20
35	BB	1084	A	C8-N9-C4	7.51	108.80	105.80
35	BB	1387	C	P-O3'-C3'	-7.51	110.69	119.70
37	BD	3	G	C8-N9-C1'	7.51	136.76	127.00
85	AA	302	C	O3'-P-O5'	7.51	118.26	104.00
85	AA	1115	G	C5-C6-N1	7.51	115.25	111.50
85	AA	1516	A	N9-C1'-C2'	-7.51	103.74	112.00
39	BF	45	G	C8-N9-C4	-7.50	103.40	106.40
77	Br	61	HIS	CA-CB-CG	-7.50	100.84	113.60
85	AA	1126	G	N3-C2-N2	7.50	125.15	119.90
85	AA	1947	A	P-O5'-C5'	7.50	132.91	120.90
34	BA	687	G	N9-C1'-C2'	-7.50	103.75	112.00
35	BB	40	C	N1-C2-O2	7.50	123.40	118.90
35	BB	981	A	C4-N9-C1'	7.50	139.81	126.30
35	BB	1458	U	C4'-C3'-C2'	-7.50	95.10	102.60
38	BE	176	G	N3-C2-N2	-7.50	114.65	119.90
38	BE	200	A	N9-C1'-C2'	-7.50	103.75	112.00
41	BH	52	G	P-O3'-C3'	-7.50	110.70	119.70
85	AA	569	A	C3'-C2'-C1'	-7.50	95.50	101.50
85	AA	1163	G	C4-N9-C1'	-7.50	116.75	126.50
85	AA	1587	C	C6-N1-C2	-7.50	117.30	120.30
34	BA	756	A	C5-C6-N6	-7.50	117.70	123.70
34	BA	779	U	C5'-C4'-C3'	7.50	128.00	116.00
35	BB	1187	G	N9-C4-C5	7.50	108.40	105.40
35	BB	1461	C	O4'-C1'-N1	7.50	114.20	108.20
48	BO	208	MET	CG-SD-CE	-7.50	88.20	100.20
85	AA	936	C	C2'-C3'-O3'	7.50	126.00	109.50
34	BA	387	A	N1-C6-N6	-7.50	114.10	118.60
34	BA	1720	U	C5-C6-N1	-7.50	118.95	122.70
35	BB	77	A	C4-N9-C1'	-7.50	112.80	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	806	U	C5-C4-O4	-7.50	121.40	125.90
37	BD	1	G	O4'-C1'-N9	7.50	114.20	108.20
85	AA	1154	A	C5'-C4'-O4'	7.50	118.10	109.10
35	BB	814	A	O4'-C1'-N9	7.50	114.20	108.20
35	BB	1200	A	C6-N1-C2	-7.50	114.10	118.60
35	BB	1219	A	P-O3'-C3'	7.50	128.70	119.70
35	BB	1285	U	C2-N3-C4	-7.50	122.50	127.00
34	BA	398	G	N1-C2-N2	7.50	122.95	116.20
34	BA	974	G	C5'-C4'-O4'	7.50	118.10	109.10
35	BB	1206	G	N1-C2-N2	-7.50	109.45	116.20
36	BC	140	U	O5'-P-OP2	-7.50	98.95	105.70
40	BG	57	A	C8-N9-C4	-7.50	102.80	105.80
85	AA	1436	A	P-O5'-C5'	7.50	132.89	120.90
34	BA	968	G	C5'-C4'-C3'	7.50	127.99	116.00
35	BB	51	U	C5'-C4'-C3'	-7.50	104.01	116.00
85	AA	92	G	N1-C6-O6	7.50	124.40	119.90
85	AA	1287	C	C2-N3-C4	-7.50	116.15	119.90
19	AK	128	ARG	NE-CZ-NH2	-7.49	116.55	120.30
34	BA	211	C	C3'-C2'-C1'	-7.49	95.51	101.50
34	BA	1011	G	O4'-C1'-C2'	7.49	114.34	107.60
34	BA	1210	A	C5'-C4'-O4'	7.49	118.09	109.10
35	BB	535	U	P-O3'-C3'	-7.49	110.71	119.70
35	BB	995	C	O4'-C1'-C2'	7.49	114.34	107.60
61	Bb	9	ARG	NE-CZ-NH2	-7.49	116.55	120.30
85	AA	860	C	N1-C1'-C2'	-7.49	103.76	112.00
34	BA	66	C	P-O3'-C3'	-7.49	110.71	119.70
34	BA	1275	G	C4-N9-C1'	-7.49	116.76	126.50
34	BA	1604	A	P-O3'-C3'	7.49	128.69	119.70
34	BA	1704	G	C4-N9-C1'	-7.49	116.76	126.50
40	BG	27	C	C5-C4-N4	7.49	125.44	120.20
41	BH	109	G	C2-N3-C4	7.49	115.65	111.90
85	AA	328	U	P-O3'-C3'	-7.49	110.71	119.70
34	BA	73	G	C8-N9-C1'	7.49	136.74	127.00
34	BA	1027	C	N3-C2-O2	-7.49	116.66	121.90
35	BB	666	A	N1-C6-N6	-7.49	114.11	118.60
35	BB	731	U	P-O5'-C5'	7.49	132.89	120.90
35	BB	831	C	O4'-C1'-N1	7.49	114.19	108.20
35	BB	844	G	C8-N9-C1'	7.49	136.74	127.00
35	BB	1089	A	O4'-C1'-C2'	7.49	114.34	107.60
38	BE	25	U	P-O3'-C3'	7.49	128.69	119.70
85	AA	533	C	C5'-C4'-C3'	-7.49	104.02	116.00
85	AA	1427	A	N1-C6-N6	-7.49	114.11	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	368	U	P-O3'-C3'	-7.49	110.71	119.70
34	BA	545	U	O4'-C1'-N1	7.49	114.19	108.20
35	BB	976	U	N3-C2-O2	-7.49	116.96	122.20
36	BC	62	A	C5-N7-C8	-7.49	100.16	103.90
40	BG	60	A	C3'-C2'-C1'	-7.49	95.51	101.50
50	BQ	70	TYR	CB-CG-CD2	-7.49	116.51	121.00
85	AA	66	U	C5-C6-N1	-7.49	118.96	122.70
85	AA	659	A	N1-C6-N6	-7.49	114.11	118.60
85	AA	660	G	C4'-C3'-C2'	-7.49	95.11	102.60
85	AA	820	G	C8-N9-C1'	7.49	136.73	127.00
85	AA	1095	C	C6-N1-C2	-7.49	117.31	120.30
85	AA	1461	A	P-O3'-C3'	7.49	128.69	119.70
85	AA	1763	G	N1-C6-O6	7.49	124.39	119.90
34	BA	1189	A	C4'-C3'-C2'	-7.49	95.11	102.60
37	BD	113	G	P-O5'-C5'	7.49	132.88	120.90
38	BE	155	C	C1'-O4'-C4'	-7.49	103.91	109.90
85	AA	1089	G	O4'-C1'-N9	7.49	114.19	108.20
34	BA	744	G	N3-C2-N2	7.49	125.14	119.90
34	BA	1320	A	C5-C6-N6	-7.49	117.71	123.70
35	BB	334	G	O4'-C1'-N9	7.49	114.19	108.20
35	BB	413	A	C5'-C4'-C3'	-7.49	104.02	116.00
35	BB	423	G	P-O5'-C5'	7.49	132.88	120.90
35	BB	587	A	C5-C6-N6	-7.49	117.71	123.70
36	BC	49	G	N1-C6-O6	-7.49	115.41	119.90
85	AA	490	A	C6-N1-C2	-7.49	114.11	118.60
85	AA	1528	A	C1'-O4'-C4'	-7.49	103.91	109.90
85	AA	2114	U	C3'-C2'-C1'	-7.49	95.51	101.50
34	BA	105	U	C2-N3-C4	-7.48	122.51	127.00
36	BC	105	C	C5'-C4'-C3'	-7.48	104.03	116.00
84	By	178	TYR	CB-CG-CD2	-7.48	116.51	121.00
85	AA	204	U	C5'-C4'-C3'	7.48	127.97	116.00
34	BA	592	G	C5'-C4'-O4'	-7.48	100.12	109.10
34	BA	1036	G	O4'-C1'-N9	7.48	114.19	108.20
34	BA	1662	U	N3-C4-O4	-7.48	114.16	119.40
85	AA	730	G	O4'-C1'-N9	7.48	114.19	108.20
85	AA	789	A	C5'-C4'-C3'	7.48	127.97	116.00
85	AA	938	A	O4'-C1'-N9	7.48	114.19	108.20
85	AA	2092	A	O5'-C5'-C4'	-7.48	97.48	111.70
5	A4	119	ARG	NE-CZ-NH1	7.48	124.04	120.30
34	BA	316	G	O4'-C1'-N9	7.48	114.18	108.20
35	BB	145	G	C5-C6-O6	-7.48	124.11	128.60
65	Bf	438	TYR	CA-CB-CG	-7.48	99.19	113.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	590	U	P-O3'-C3'	7.48	128.68	119.70
35	BB	765	G	C4-N9-C1'	-7.48	116.78	126.50
38	BE	43	A	N1-C6-N6	-7.48	114.11	118.60
34	BA	700	G	C8-N9-C1'	7.48	136.72	127.00
34	BA	960	C	C1'-O4'-C4'	-7.48	103.92	109.90
34	BA	1485	U	C5-C6-N1	-7.48	118.96	122.70
40	BG	33	G	O4'-C1'-C2'	7.48	114.33	107.60
85	AA	214	C	O4'-C1'-N1	7.48	114.18	108.20
85	AA	241	U	N3-C2-O2	-7.48	116.97	122.20
85	AA	388	G	O3'-P-O5'	7.48	118.20	104.00
85	AA	478	U	C1'-O4'-C4'	-7.48	103.92	109.90
34	BA	113	G	N9-C1'-C2'	-7.48	103.78	112.00
34	BA	541	C	C6-N1-C2	-7.48	117.31	120.30
34	BA	1638	U	C2-N1-C1'	-7.48	108.73	117.70
35	BB	295	U	O4'-C1'-N1	7.48	114.18	108.20
35	BB	579	A	C5-C6-N6	-7.48	117.72	123.70
35	BB	1462	G	C5-C6-O6	7.48	133.09	128.60
52	BS	7	ARG	NE-CZ-NH2	-7.48	116.56	120.30
85	AA	46	U	P-O3'-C3'	7.48	128.67	119.70
34	BA	640	U	N3-C2-O2	-7.47	116.97	122.20
34	BA	1491	U	C5'-C4'-C3'	-7.47	104.04	116.00
34	BA	1799	G	N1-C6-O6	-7.47	115.42	119.90
40	BG	102	G	N1-C2-N2	-7.47	109.47	116.20
63	Bd	28	TYR	CB-CG-CD1	-7.47	116.52	121.00
85	AA	2142	A	C1'-O4'-C4'	-7.47	103.92	109.90
34	BA	917	C	C5-C4-N4	7.47	125.43	120.20
34	BA	1278	A	C1'-O4'-C4'	-7.47	103.92	109.90
34	BA	1478	G	C5-C6-O6	-7.47	124.12	128.60
35	BB	493	U	C1'-O4'-C4'	-7.47	103.92	109.90
81	Bv	75	ARG	CB-CA-C	-7.47	95.45	110.40
85	AA	2141	G	C6-C5-N7	-7.47	125.92	130.40
34	BA	10	G	C5-C6-O6	7.47	133.08	128.60
34	BA	1717	C	O4'-C1'-N1	7.47	114.18	108.20
40	BG	176	G	C8-N9-C1'	7.47	136.71	127.00
47	BN	14	ARG	NE-CZ-NH1	-7.47	116.56	120.30
34	BA	342	U	C2-N3-C4	-7.47	122.52	127.00
34	BA	1619	U	C4'-C3'-O3'	-7.47	93.71	109.40
35	BB	586	U	C2-N1-C1'	-7.47	108.74	117.70
35	BB	1537	C	C3'-C2'-C1'	-7.47	95.52	101.50
40	BG	64	C	P-O3'-C3'	-7.47	110.74	119.70
40	BG	97	G	N1-C6-O6	7.47	124.38	119.90
83	Bx	47	PHE	N-CA-C	7.47	131.17	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	604	C	C4'-C3'-C2'	-7.47	95.13	102.60
85	AA	1252	A	C5'-C4'-C3'	-7.47	104.05	116.00
85	AA	2007	G	O4'-C1'-C2'	7.47	114.32	107.60
34	BA	674	G	C5-C6-O6	-7.47	124.12	128.60
34	BA	884	G	N1-C6-O6	7.47	124.38	119.90
34	BA	1087	A	C4-C5-C6	-7.47	113.27	117.00
41	BH	8	C	P-O3'-C3'	7.47	128.66	119.70
34	BA	800	G	N1-C6-O6	7.47	124.38	119.90
34	BA	1420	A	O4'-C1'-N9	7.47	114.17	108.20
34	BA	1646	U	C2-N3-C4	-7.47	122.52	127.00
35	BB	1032	U	O4'-C1'-C2'	-7.47	98.33	105.80
35	BB	1065	G	O4'-C1'-N9	7.47	114.17	108.20
35	BB	1274	G	O4'-C1'-C2'	-7.47	98.33	105.80
35	BB	136	A	C1'-O4'-C4'	-7.46	103.93	109.90
35	BB	534	C	C1'-O4'-C4'	-7.46	103.93	109.90
35	BB	678	U	C2-N3-C4	-7.46	122.52	127.00
36	BC	28	C	C6-N1-C1'	-7.46	111.84	120.80
41	BH	19	G	C4'-C3'-C2'	7.46	110.06	102.60
85	AA	1814	U	N1-C2-N3	7.46	119.38	114.90
27	AT	72	GLY	C-N-CA	7.46	137.97	122.30
34	BA	616	G	C4-N9-C1'	-7.46	116.80	126.50
39	BF	25	G	C5-C6-O6	-7.46	124.12	128.60
85	AA	1840	C	C6-N1-C2	-7.46	117.31	120.30
13	AE	83	ARG	NE-CZ-NH1	-7.46	116.57	120.30
34	BA	112	C	C2-N1-C1'	-7.46	110.59	118.80
34	BA	310	C	O4'-C1'-N1	7.46	114.17	108.20
34	BA	483	A	C2'-C3'-O3'	7.46	125.91	109.50
34	BA	820	C	N1-C1'-C2'	-7.46	103.79	112.00
34	BA	1686	G	C5-C6-N1	7.46	115.23	111.50
35	BB	551	C	C6-N1-C2	-7.46	117.32	120.30
35	BB	765	G	C5-C6-O6	-7.46	124.12	128.60
37	BD	69	U	N3-C4-C5	7.46	119.08	114.60
38	BE	112	G	O4'-C1'-N9	7.46	114.17	108.20
40	BG	156	G	N1-C6-O6	7.46	124.38	119.90
85	AA	132	G	P-O3'-C3'	-7.46	110.75	119.70
35	BB	680	A	C4'-C3'-C2'	7.46	110.06	102.60
35	BB	797	C	O4'-C4'-C3'	-7.46	96.54	104.00
35	BB	1144	A	C5-C6-N1	7.46	121.43	117.70
49	BP	163	ARG	NE-CZ-NH1	7.46	124.03	120.30
85	AA	867	G	C5'-C4'-C3'	-7.46	104.06	116.00
85	AA	2153	G	C5-C6-O6	-7.46	124.12	128.60
35	BB	829	C	N3-C2-O2	-7.46	116.68	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	995	G	C8-N9-C4	7.46	109.38	106.40
86	AB	17	C	C5'-C4'-C3'	-7.46	104.06	116.00
34	BA	862	C	P-O5'-C5'	-7.46	108.97	120.90
34	BA	1210	A	P-O5'-C5'	7.46	132.83	120.90
35	BB	764	C	O4'-C1'-N1	7.46	114.17	108.20
35	BB	1166	A	C1'-O4'-C4'	-7.46	103.94	109.90
35	BB	1254	G	C4-N9-C1'	-7.46	116.81	126.50
85	AA	767	A	C1'-O4'-C4'	-7.46	103.94	109.90
85	AA	1930	U	O4'-C1'-N1	7.46	114.17	108.20
34	BA	191	G	N3-C2-N2	7.46	125.12	119.90
36	BC	27	U	N1-C2-N3	-7.46	110.43	114.90
37	BD	95	G	N3-C4-N9	7.46	130.47	126.00
39	BF	35	C	C5-C6-N1	7.46	124.73	121.00
24	AQ	33	MET	CG-SD-CE	-7.45	88.27	100.20
34	BA	582	U	C6-N1-C2	-7.45	116.53	121.00
35	BB	834	U	C5'-C4'-C3'	-7.45	104.07	116.00
37	BD	24	U	C3'-C2'-C1'	-7.45	95.54	101.50
38	BE	90	G	C5'-C4'-C3'	7.45	127.92	116.00
38	BE	111	C	C5'-C4'-O4'	7.45	118.04	109.10
38	BE	155	C	C5'-C4'-O4'	7.45	118.04	109.10
50	BQ	85	ARG	NE-CZ-NH2	-7.45	116.57	120.30
85	AA	2014	G	N1-C6-O6	7.45	124.37	119.90
23	AP	154	TRP	CB-CG-CD2	-7.45	116.91	126.60
34	BA	519	G	P-O5'-C5'	7.45	132.82	120.90
34	BA	322	U	C5'-C4'-C3'	-7.45	104.08	116.00
35	BB	75	A	P-O3'-C3'	-7.45	110.76	119.70
85	AA	383	C	N3-C4-N4	-7.45	112.78	118.00
85	AA	600	C	C6-N1-C2	-7.45	117.32	120.30
34	BA	579	U	O4'-C1'-N1	-7.45	102.24	108.20
35	BB	77	A	C8-N9-C4	7.45	108.78	105.80
35	BB	1037	A	P-O3'-C3'	7.45	128.64	119.70
35	BB	1397	G	P-O3'-C3'	7.45	128.64	119.70
35	BB	997	G	N1-C6-O6	7.45	124.37	119.90
85	AA	478	U	P-O5'-C5'	-7.45	108.98	120.90
85	AA	700	U	C5'-C4'-C3'	-7.45	104.08	116.00
85	AA	749	C	C5-C4-N4	7.45	125.41	120.20
14	AF	97	ARG	NE-CZ-NH2	-7.45	116.58	120.30
34	BA	1620	U	C4'-C3'-C2'	-7.45	95.16	102.60
35	BB	317	C	C6-N1-C2	-7.45	117.32	120.30
35	BB	1368	A	C1'-O4'-C4'	-7.45	103.94	109.90
35	BB	1458	U	C5'-C4'-O4'	7.45	118.03	109.10
38	BE	66	A	C5'-C4'-C3'	7.45	127.91	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	993	G	C5-C6-O6	-7.45	124.13	128.60
34	BA	1294	C	C5-C4-N4	7.44	125.41	120.20
35	BB	701	U	C6-N1-C1'	-7.44	110.78	121.20
35	BB	1483	A	C2'-C3'-O3'	7.44	125.88	109.50
85	AA	570	U	C6-N1-C1'	7.44	131.62	121.20
34	BA	71	G	C5-C6-O6	-7.44	124.13	128.60
34	BA	1429	A	P-O3'-C3'	-7.44	110.77	119.70
35	BB	1486	C	O4'-C1'-N1	7.44	114.15	108.20
40	BG	21	C	C1'-O4'-C4'	-7.44	103.95	109.90
85	AA	1132	A	O4'-C1'-N9	7.44	114.15	108.20
85	AA	2037	A	N1-C6-N6	7.44	123.07	118.60
34	BA	640	U	N1-C2-O2	7.44	128.01	122.80
34	BA	929	A	N1-C6-N6	7.44	123.06	118.60
34	BA	1205	A	P-O3'-C3'	7.44	128.63	119.70
34	BA	1434	U	C4'-C3'-C2'	-7.44	95.16	102.60
34	BA	1482	A	P-O3'-C3'	-7.44	110.77	119.70
35	BB	1132	A	O5'-C5'-C4'	-7.44	97.56	111.70
35	BB	1319	U	O5'-C5'-C4'	7.44	125.84	111.70
85	AA	995	G	C4-N9-C1'	-7.44	116.83	126.50
85	AA	2090	C	O4'-C1'-N1	7.44	114.15	108.20
19	AK	129	ARG	NE-CZ-NH2	-7.44	116.58	120.30
35	BB	1186	A	N9-C4-C5	-7.44	102.83	105.80
85	AA	47	A	C6-N1-C2	-7.44	114.14	118.60
85	AA	453	G	P-O3'-C3'	-7.44	110.77	119.70
85	AA	2011	C	N1-C1'-C2'	-7.44	103.82	112.00
34	BA	375	C	C1'-O4'-C4'	-7.44	103.95	109.90
40	BG	170	G	C1'-O4'-C4'	-7.44	103.95	109.90
85	AA	596	A	C5-C6-N6	-7.44	117.75	123.70
34	BA	572	G	C3'-C2'-C1'	7.44	107.45	101.50
68	Bi	14	ARG	NE-CZ-NH1	7.44	124.02	120.30
85	AA	57	G	C4'-C3'-C2'	-7.44	95.16	102.60
85	AA	1281	G	C4-C5-C6	-7.44	114.34	118.80
85	AA	2039	G	C4-N9-C1'	-7.44	116.83	126.50
35	BB	558	U	O4'-C1'-N1	7.43	114.15	108.20
35	BB	1438	U	P-O3'-C3'	-7.43	110.78	119.70
36	BC	163	A	C4-N9-C1'	-7.43	112.92	126.30
40	BG	68	U	O4'-C1'-N1	7.43	114.15	108.20
40	BG	179	C	P-O3'-C3'	-7.43	110.78	119.70
85	AA	315	U	O4'-C1'-N1	7.43	114.15	108.20
85	AA	360	C	O4'-C1'-N1	7.43	114.15	108.20
85	AA	675	A	N1-C6-N6	-7.43	114.14	118.60
85	AA	1706	A	C3'-C2'-C1'	-7.43	95.55	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	121	A	N1-C6-N6	7.43	123.06	118.60
34	BA	1612	C	C6-N1-C2	-7.43	117.33	120.30
35	BB	16	G	P-O3'-C3'	7.43	128.62	119.70
35	BB	128	C	P-O3'-C3'	-7.43	110.78	119.70
85	AA	773	G	C1'-O4'-C4'	-7.43	103.95	109.90
85	AA	891	G	C4-N9-C1'	7.43	136.16	126.50
85	AA	2114	U	O4'-C1'-N1	7.43	114.15	108.20
34	BA	7	U	P-O3'-C3'	-7.43	110.78	119.70
85	AA	645	C	C4'-C3'-C2'	7.43	110.03	102.60
85	AA	878	U	C2-N3-C4	-7.43	122.54	127.00
85	AA	979	U	P-O3'-C3'	7.43	128.62	119.70
85	AA	1149	A	C5'-C4'-C3'	-7.43	104.11	116.00
85	AA	1535	C	C2-N1-C1'	7.43	126.97	118.80
34	BA	86	A	N1-C6-N6	7.43	123.06	118.60
34	BA	436	U	C2-N1-C1'	-7.43	108.78	117.70
34	BA	729	C	N3-C2-O2	-7.43	116.70	121.90
34	BA	1177	C	C1'-O4'-C4'	-7.43	103.96	109.90
85	AA	455	G	C4-N9-C1'	-7.43	116.84	126.50
85	AA	740	A	C3'-C2'-C1'	-7.43	95.56	101.50
85	AA	910	G	C4-N9-C1'	-7.43	116.84	126.50
85	AA	2185	U	P-O3'-C3'	-7.43	110.78	119.70
40	BG	39	A	C8-N9-C4	7.43	108.77	105.80
85	AA	1364	U	P-O3'-C3'	-7.43	110.79	119.70
85	AA	1829	C	C3'-C2'-C1'	-7.43	95.56	101.50
34	BA	18	G	C4-N9-C1'	-7.43	116.84	126.50
34	BA	1329	U	N3-C2-O2	-7.43	117.00	122.20
34	BA	1640	G	C6-N1-C2	-7.43	120.64	125.10
34	BA	1789	A	C5'-C4'-O4'	7.43	118.01	109.10
35	BB	127	U	O4'-C1'-N1	7.43	114.14	108.20
35	BB	1185	G	C5-C6-N1	7.43	115.21	111.50
35	BB	1512	C	C2-N1-C1'	-7.43	110.63	118.80
37	BD	55	A	N1-C6-N6	-7.43	114.14	118.60
39	BF	50	C	C3'-C2'-C1'	-7.43	95.56	101.50
40	BG	17	A	P-O5'-C5'	-7.43	109.02	120.90
40	BG	45	G	C5'-C4'-C3'	7.43	127.88	116.00
85	AA	35	U	C3'-C2'-C1'	-7.43	95.56	101.50
85	AA	520	A	O5'-P-OP2	7.43	119.61	110.70
85	AA	556	C	P-O3'-C3'	-7.43	110.79	119.70
85	AA	762	U	O5'-P-OP2	7.43	119.61	110.70
85	AA	942	A	O4'-C1'-N9	7.43	114.14	108.20
85	AA	1001	G	N9-C1'-C2'	-7.43	103.83	112.00
85	AA	1644	G	C4-N9-C1'	7.43	136.16	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A2	149	PHE	CB-CG-CD2	-7.42	115.60	120.80
34	BA	166	G	P-O3'-C3'	-7.42	110.79	119.70
85	AA	1231	G	C4-N9-C1'	-7.42	116.85	126.50
85	AA	1676	G	C2-N3-C4	-7.42	108.19	111.90
34	BA	606	G	N3-C4-N9	7.42	130.45	126.00
34	BA	1517	U	N1-C1'-C2'	-7.42	103.83	112.00
35	BB	732	G	C1'-O4'-C4'	-7.42	103.96	109.90
35	BB	879	G	O4'-C4'-C3'	-7.42	96.58	104.00
26	AS	107	ARG	NE-CZ-NH1	7.42	124.01	120.30
34	BA	1164	C	P-O3'-C3'	-7.42	110.79	119.70
34	BA	1629	A	O3'-P-O5'	7.42	118.10	104.00
41	BH	102	C	O4'-C1'-N1	7.42	114.14	108.20
80	Bu	22	ARG	NE-CZ-NH2	-7.42	116.59	120.30
85	AA	36	U	N3-C2-O2	-7.42	117.00	122.20
85	AA	1889	U	C5'-C4'-C3'	-7.42	104.12	116.00
34	BA	1260	G	C4-N9-C1'	-7.42	116.85	126.50
35	BB	1224	C	C5'-C4'-C3'	-7.42	104.13	116.00
40	BG	25	G	C4-N9-C1'	-7.42	116.86	126.50
34	BA	892	C	C6-N1-C2	-7.42	117.33	120.30
34	BA	1211	G	C5-N7-C8	-7.42	100.59	104.30
35	BB	8	U	C5-C4-O4	-7.42	121.45	125.90
37	BD	83	A	OP1-P-O3'	7.42	121.52	105.20
61	Bb	19	TYR	CB-CG-CD1	7.42	125.45	121.00
82	Bw	188	TYR	CB-CG-CD1	-7.42	116.55	121.00
85	AA	1605	G	P-O3'-C3'	7.42	128.60	119.70
34	BA	91	C	C5'-C4'-O4'	7.42	118.00	109.10
34	BA	679	U	C2-N1-C1'	-7.42	108.80	117.70
34	BA	831	U	C6-N1-C2	-7.42	116.55	121.00
34	BA	1249	G	N3-C2-N2	7.42	125.09	119.90
35	BB	986	C	C6-N1-C1'	-7.42	111.90	120.80
35	BB	1512	C	C6-N1-C1'	7.42	129.70	120.80
40	BG	138	C	O4'-C1'-N1	7.42	114.13	108.20
64	Be	163	ARG	NE-CZ-NH1	7.42	124.01	120.30
85	AA	735	G	C5-C6-O6	-7.42	124.15	128.60
85	AA	740	A	C8-N9-C4	-7.42	102.83	105.80
85	AA	2100	A	C5'-C4'-C3'	7.42	127.87	116.00
35	BB	452	A	C5'-C4'-O4'	-7.42	100.20	109.10
35	BB	844	G	P-O3'-C3'	-7.42	110.80	119.70
85	AA	928	U	C5-C4-O4	-7.42	121.45	125.90
85	AA	1808	G	C8-N9-C1'	-7.42	117.36	127.00
34	BA	544	U	N1-C1'-C2'	-7.41	103.84	112.00
35	BB	80	C	N3-C2-O2	-7.41	116.71	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1454	G	C4'-C3'-C2'	-7.41	95.19	102.60
35	BB	1523	U	C3'-C2'-C1'	-7.41	95.57	101.50
85	AA	352	G	N9-C1'-C2'	-7.41	103.85	112.00
34	BA	1503	U	O3'-P-O5'	-7.41	89.92	104.00
35	BB	997	G	N9-C1'-C2'	-7.41	103.85	112.00
39	BF	72	A	O4'-C1'-N9	7.41	114.13	108.20
85	AA	390	U	C5'-C4'-C3'	-7.41	104.14	116.00
85	AA	690	G	C5-C6-N1	7.41	115.21	111.50
34	BA	373	G	C8-N9-C1'	7.41	136.63	127.00
34	BA	543	A	C6-N1-C2	-7.41	114.15	118.60
35	BB	866	A	C1'-O4'-C4'	-7.41	103.97	109.90
35	BB	1476	C	N1-C1'-C2'	-7.41	103.85	112.00
85	AA	2234	C	C6-N1-C2	-7.41	117.34	120.30
29	AV	91	ARG	NE-CZ-NH1	7.41	124.00	120.30
34	BA	926	A	C2'-C3'-O3'	7.41	125.80	109.50
34	BA	1711	G	O5'-C5'-C4'	-7.41	97.62	111.70
85	AA	2111	C	C6-N1-C2	-7.41	117.34	120.30
85	AA	2202	G	N9-C1'-C2'	-7.41	103.85	112.00
35	BB	1201	G	P-O3'-C3'	-7.41	110.81	119.70
37	BD	83	A	C3'-C2'-C1'	-7.41	95.57	101.50
34	BA	81	C	O5'-C5'-C4'	-7.41	97.63	111.70
35	BB	90	G	C8-N9-C4	7.41	109.36	106.40
85	AA	120	C	N3-C2-O2	-7.41	116.72	121.90
85	AA	1644	G	C5-C6-O6	-7.41	124.16	128.60
85	AA	2119	C	N1-C2-N3	7.41	124.38	119.20
86	AB	14	A	C5-C6-N6	7.41	129.62	123.70
36	BC	14	G	C4-N9-C1'	-7.40	116.88	126.50
85	AA	1245	U	C2-N3-C4	-7.40	122.56	127.00
85	AA	1732	G	O4'-C1'-N9	7.40	114.12	108.20
8	A7	245	PHE	CB-CG-CD2	-7.40	115.62	120.80
34	BA	301	U	C2-N3-C4	-7.40	122.56	127.00
34	BA	451	A	P-O3'-C3'	7.40	128.58	119.70
34	BA	1179	U	C5'-C4'-C3'	-7.40	104.16	116.00
34	BA	1834	A	C5'-C4'-C3'	-7.40	104.16	116.00
35	BB	92	C	C6-N1-C2	-7.40	117.34	120.30
35	BB	729	G	C5'-C4'-C3'	7.40	127.84	116.00
40	BG	38	A	C8-N9-C4	7.40	108.76	105.80
85	AA	103	U	C5'-C4'-C3'	-7.40	104.16	116.00
85	AA	812	C	O4'-C1'-N1	7.40	114.12	108.20
34	BA	1484	A	C6-N1-C2	-7.40	114.16	118.60
40	BG	141	A	C5-C6-N6	-7.40	117.78	123.70
85	AA	274	A	C1'-O4'-C4'	-7.40	103.98	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1929	G	P-O5'-C5'	-7.40	109.06	120.90
34	BA	238	C	N3-C4-C5	-7.40	118.94	121.90
34	BA	748	C	C2-N1-C1'	-7.40	110.66	118.80
34	BA	1664	C	P-O3'-C3'	-7.40	110.82	119.70
35	BB	599	U	P-O3'-C3'	-7.40	110.82	119.70
35	BB	1014	U	C6-N1-C1'	7.40	131.56	121.20
36	BC	38	U	O4'-C1'-C2'	-7.40	98.40	105.80
38	BE	83	U	C5'-C4'-O4'	7.40	117.98	109.10
85	AA	1454	U	P-O3'-C3'	-7.40	110.82	119.70
85	AA	1696	U	O4'-C1'-N1	7.40	114.12	108.20
35	BB	102	G	P-O3'-C3'	-7.40	110.83	119.70
19	AK	144	PHE	CB-CG-CD1	7.39	125.98	120.80
34	BA	362	G	P-O3'-C3'	-7.39	110.83	119.70
34	BA	1017	C	O4'-C1'-N1	7.39	114.11	108.20
34	BA	1096	C	C1'-O4'-C4'	-7.39	103.98	109.90
34	BA	1542	A	C5-C6-N6	-7.39	117.78	123.70
35	BB	460	C	C5'-C4'-C3'	7.39	127.83	116.00
35	BB	773	G	C4-N9-C1'	-7.39	116.89	126.50
52	BS	89	TYR	CB-CG-CD2	-7.39	116.56	121.00
82	Bw	220	PHE	CB-CG-CD2	-7.39	115.62	120.80
86	AB	5	G	C8-N9-C1'	7.39	136.61	127.00
34	BA	79	C	C6-N1-C2	-7.39	117.34	120.30
34	BA	579	U	C2'-C3'-O3'	7.39	125.76	109.50
34	BA	857	C	C2-N1-C1'	-7.39	110.67	118.80
34	BA	1102	A	C5-C6-N1	7.39	121.40	117.70
34	BA	1610	A	P-O5'-C5'	-7.39	109.07	120.90
35	BB	7	C	C5-C4-N4	7.39	125.37	120.20
35	BB	854	G	C4-N9-C1'	-7.39	116.89	126.50
37	BD	86	A	C8-N9-C1'	7.39	141.01	127.70
37	BD	92	G	C8-N9-C1'	7.39	136.61	127.00
39	BF	65	U	P-O3'-C3'	7.39	128.57	119.70
85	AA	95	U	C3'-C2'-C1'	-7.39	95.58	101.50
85	AA	289	G	C3'-C2'-C1'	-7.39	95.59	101.50
85	AA	487	G	N1-C6-O6	-7.39	115.46	119.90
85	AA	2136	C	C2-N1-C1'	-7.39	110.67	118.80
34	BA	294	C	O5'-C5'-C4'	-7.39	97.66	111.70
34	BA	868	C	C2-N3-C4	-7.39	116.20	119.90
34	BA	1352	G	C8-N9-C1'	7.39	136.61	127.00
35	BB	1007	U	O4'-C1'-N1	7.39	114.11	108.20
38	BE	8	G	C5-C6-N1	7.39	115.20	111.50
85	AA	493	A	C4-N9-C1'	-7.39	113.00	126.30
85	AA	967	C	C5'-C4'-C3'	-7.39	104.17	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	508	C	C6-N1-C1'	-7.39	111.93	120.80
34	BA	513	U	C1'-O4'-C4'	-7.39	103.99	109.90
34	BA	761	U	N1-C1'-C2'	-7.39	103.87	112.00
34	BA	1686	G	C5-C6-O6	-7.39	124.17	128.60
35	BB	61	A	O3'-P-O5'	-7.39	89.96	104.00
35	BB	291	C	O4'-C1'-N1	7.39	114.11	108.20
35	BB	1212	C	C5-C4-N4	7.39	125.37	120.20
36	BC	158	U	C5'-C4'-O4'	7.39	117.97	109.10
38	BE	96	G	N3-C4-C5	-7.39	124.91	128.60
40	BG	73	U	C2-N3-C4	-7.39	122.57	127.00
85	AA	585	G	C6-C5-N7	-7.39	125.97	130.40
85	AA	1370	G	C8-N9-C1'	7.39	136.61	127.00
3	A2	131	ARG	NE-CZ-NH1	7.39	123.99	120.30
34	BA	1162	U	P-O3'-C3'	-7.39	110.83	119.70
40	BG	169	A	C4-N9-C1'	-7.39	113.00	126.30
85	AA	400	G	O5'-P-OP2	7.39	119.57	110.70
85	AA	866	U	C4'-C3'-C2'	7.39	109.99	102.60
85	AA	1651	C	O4'-C1'-N1	7.39	114.11	108.20
11	AC	124	PHE	CB-CG-CD2	-7.39	115.63	120.80
34	BA	122	U	P-O3'-C3'	7.39	128.56	119.70
34	BA	782	C	O4'-C1'-N1	7.39	114.11	108.20
34	BA	1794	A	P-O3'-C3'	-7.39	110.84	119.70
36	BC	46	G	N3-C2-N2	7.39	125.07	119.90
40	BG	168	A	C4-C5-C6	-7.39	113.31	117.00
85	AA	367	A	N1-C6-N6	-7.39	114.17	118.60
34	BA	674	G	C8-N9-C4	7.38	109.35	106.40
34	BA	1615	A	C4'-C3'-C2'	-7.38	95.22	102.60
34	BA	1846	G	C5'-C4'-O4'	7.38	117.96	109.10
38	BE	127	G	C5-C6-N1	7.38	115.19	111.50
38	BE	131	C	C6-N1-C2	-7.38	117.35	120.30
40	BG	59	G	C4'-C3'-C2'	-7.38	95.22	102.60
85	AA	834	U	C1'-O4'-C4'	-7.38	103.99	109.90
85	AA	1856	G	C4-N9-C1'	-7.38	116.90	126.50
85	AA	1924	C	P-O5'-C5'	7.38	132.72	120.90
20	AL	100	ARG	NE-CZ-NH1	7.38	123.99	120.30
34	BA	505	U	O4'-C1'-N1	7.38	114.11	108.20
35	BB	876	G	C4-N9-C1'	-7.38	116.90	126.50
36	BC	26	U	C6-N1-C2	-7.38	116.57	121.00
40	BG	122	G	C4-N9-C1'	-7.38	116.90	126.50
34	BA	437	G	N1-C6-O6	7.38	124.33	119.90
34	BA	550	U	O4'-C1'-N1	7.38	114.10	108.20
34	BA	602	G	C2'-C3'-O3'	7.38	125.74	109.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	645	U	C5-C4-O4	-7.38	121.47	125.90
34	BA	1288	U	P-O3'-C3'	-7.38	110.84	119.70
35	BB	544	C	N3-C2-O2	-7.38	116.73	121.90
37	BD	15	U	P-O3'-C3'	-7.38	110.84	119.70
37	BD	111	U	C5'-C4'-C3'	-7.38	104.19	116.00
40	BG	15	G	C8-N9-C1'	7.38	136.60	127.00
85	AA	780	U	P-O5'-C5'	7.38	132.71	120.90
85	AA	1250	A	P-O3'-C3'	-7.38	110.84	119.70
34	BA	134	U	N1-C2-N3	7.38	119.33	114.90
34	BA	764	G	N1-C2-N2	-7.38	109.56	116.20
35	BB	1155	U	P-O3'-C3'	-7.38	110.84	119.70
85	AA	937	G	C8-N9-C1'	7.38	136.59	127.00
34	BA	888	G	C8-N9-C1'	7.38	136.59	127.00
34	BA	1295	U	C6-N1-C1'	7.38	131.53	121.20
68	Bi	58	TYR	CA-CB-CG	-7.38	99.38	113.40
85	AA	1153	G	C3'-C2'-C1'	-7.38	95.60	101.50
85	AA	1523	G	O4'-C1'-N9	7.38	114.10	108.20
85	AA	2200	A	N9-C1'-C2'	-7.38	103.88	112.00
34	BA	684	G	C4'-C3'-C2'	-7.38	95.22	102.60
35	BB	756	C	P-O3'-C3'	-7.38	110.85	119.70
36	BC	27	U	P-O5'-C5'	7.38	132.70	120.90
58	BY	2	ARG	NE-CZ-NH2	-7.38	116.61	120.30
85	AA	1810	C	P-O3'-C3'	-7.38	110.85	119.70
14	AF	55	ARG	NE-CZ-NH1	7.38	123.99	120.30
85	AA	1387	C	O4'-C1'-N1	7.38	114.10	108.20
34	BA	666	C	O4'-C1'-N1	7.37	114.10	108.20
34	BA	669	U	N1-C1'-C2'	-7.37	103.89	112.00
34	BA	996	U	N3-C2-O2	-7.37	117.04	122.20
34	BA	1318	G	P-O3'-C3'	-7.37	110.85	119.70
35	BB	399	A	P-O3'-C3'	-7.37	110.85	119.70
35	BB	701	U	C5'-C4'-O4'	7.37	117.95	109.10
35	BB	901	U	O5'-C5'-C4'	-7.37	97.69	111.70
38	BE	124	G	C2-N3-C4	-7.37	108.21	111.90
85	AA	364	C	C6-N1-C2	7.37	123.25	120.30
85	AA	1259	U	O4'-C1'-N1	7.37	114.10	108.20
85	AA	2244	G	C8-N9-C1'	-7.37	117.42	127.00
35	BB	440	U	C1'-O4'-C4'	-7.37	104.00	109.90
35	BB	590	G	N1-C6-O6	-7.37	115.48	119.90
39	BF	50	C	C2-N3-C4	-7.37	116.22	119.90
85	AA	744	C	C5'-C4'-C3'	-7.37	104.21	116.00
85	AA	2123	U	O4'-C1'-N1	7.37	114.10	108.20
35	BB	1475	U	O3'-P-O5'	7.37	118.00	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
53	BT	63	TRP	CA-CB-CG	7.37	127.70	113.70
85	AA	970	U	C2-N1-C1'	7.37	126.54	117.70
4	A3	38	ASP	CA-CB-CG	-7.37	97.19	113.40
34	BA	1220	C	N3-C2-O2	-7.37	116.74	121.90
34	BA	1535	G	C1'-O4'-C4'	-7.37	104.00	109.90
35	BB	993	A	O3'-P-O5'	-7.37	90.00	104.00
38	BE	146	U	N1-C2-N3	7.37	119.32	114.90
40	BG	171	A	O4'-C1'-C2'	-7.37	98.43	105.80
34	BA	563	A	P-O3'-C3'	-7.37	110.86	119.70
34	BA	719	G	C5'-C4'-C3'	-7.37	104.21	116.00
35	BB	877	A	O4'-C4'-C3'	-7.37	96.63	104.00
37	BD	87	G	O4'-C1'-N9	7.37	114.09	108.20
41	BH	109	G	C8-N9-C1'	7.37	136.58	127.00
85	AA	1670	U	C6-N1-C1'	7.37	131.51	121.20
34	BA	198	U	C2-N3-C4	-7.37	122.58	127.00
34	BA	223	U	P-O3'-C3'	7.37	128.54	119.70
34	BA	961	C	P-O5'-C5'	-7.37	109.11	120.90
34	BA	1454	G	C2-N3-C4	-7.37	108.22	111.90
35	BB	789	G	C1'-O4'-C4'	-7.37	104.01	109.90
35	BB	1414	A	N1-C6-N6	-7.37	114.18	118.60
36	BC	106	G	C1'-O4'-C4'	-7.37	104.01	109.90
38	BE	75	C	P-O3'-C3'	-7.37	110.86	119.70
40	BG	138	C	C4'-C3'-C2'	-7.37	95.23	102.60
40	BG	157	A	C5'-C4'-O4'	7.37	117.94	109.10
85	AA	820	G	N1-C6-O6	-7.37	115.48	119.90
85	AA	1367	C	O5'-C5'-C4'	-7.37	97.71	111.70
85	AA	2188	C	C5'-C4'-C3'	-7.37	104.22	116.00
34	BA	1821	A	C4'-C3'-C2'	-7.36	95.24	102.60
35	BB	319	C	O4'-C1'-N1	7.36	114.09	108.20
35	BB	629	C	P-O5'-C5'	7.36	132.68	120.90
35	BB	799	A	P-O5'-C5'	-7.36	109.12	120.90
67	Bh	36	ARG	NE-CZ-NH2	-7.36	116.62	120.30
85	AA	370	A	C5'-C4'-O4'	-7.36	100.26	109.10
85	AA	838	G	O4'-C1'-N9	7.36	114.09	108.20
85	AA	1922	A	O4'-C1'-C2'	-7.36	98.44	105.80
85	AA	559	G	C5'-C4'-C3'	-7.36	104.22	116.00
85	AA	1581	C	O4'-C1'-N1	7.36	114.09	108.20
34	BA	1011	G	P-O3'-C3'	-7.36	110.87	119.70
35	BB	1101	C	O4'-C1'-N1	7.36	114.09	108.20
38	BE	210	G	O4'-C1'-N9	7.36	114.09	108.20
85	AA	465	A	C2'-C3'-O3'	7.36	125.69	109.50
85	AA	529	G	O3'-P-O5'	-7.36	90.02	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1248	U	C4'-C3'-C2'	7.36	109.96	102.60
34	BA	110	C	O3'-P-O5'	7.36	117.98	104.00
34	BA	513	U	C5-C6-N1	-7.36	119.02	122.70
34	BA	655	U	P-O5'-C5'	-7.36	109.13	120.90
34	BA	1814	U	N3-C2-O2	-7.36	117.05	122.20
35	BB	694	C	N1-C2-O2	7.36	123.32	118.90
35	BB	1032	U	C4'-C3'-C2'	-7.36	95.24	102.60
38	BE	94	U	C2-N3-C4	-7.36	122.58	127.00
70	Bk	95	ARG	NE-CZ-NH1	7.36	123.98	120.30
85	AA	548	G	N9-C1'-C2'	-7.36	103.91	112.00
85	AA	1168	C	C4'-C3'-C2'	7.36	109.96	102.60
85	AA	1867	G	C5-C6-O6	-7.36	124.18	128.60
85	AA	2185	U	O4'-C1'-N1	7.36	114.09	108.20
34	BA	87	G	O4'-C1'-N9	7.36	114.09	108.20
34	BA	346	A	O4'-C1'-N9	7.36	114.09	108.20
34	BA	758	G	C5-N7-C8	-7.36	100.62	104.30
34	BA	1341	A	C5'-C4'-O4'	7.36	117.93	109.10
35	BB	37	C	C1'-O4'-C4'	-7.36	104.01	109.90
35	BB	880	G	C5-C6-O6	-7.36	124.19	128.60
41	BH	11	C	C6-N1-C2	-7.36	117.36	120.30
41	BH	118	U	O3'-P-O5'	7.36	117.98	104.00
85	AA	2018	U	N1-C2-O2	-7.36	117.65	122.80
34	BA	1472	G	C8-N9-C1'	7.36	136.56	127.00
35	BB	1331	U	C5'-C4'-C3'	-7.36	104.23	116.00
36	BC	32	U	O3'-P-O5'	7.36	117.97	104.00
36	BC	153	C	P-O5'-C5'	7.36	132.67	120.90
38	BE	8	G	C5'-C4'-O4'	7.36	117.92	109.10
39	BF	13	U	N1-C1'-C2'	-7.36	103.91	112.00
40	BG	164	U	P-O3'-C3'	-7.36	110.87	119.70
48	BO	112	ARG	NE-CZ-NH1	7.36	123.98	120.30
85	AA	23	G	C8-N9-C1'	7.36	136.56	127.00
85	AA	1274	A	O3'-P-O5'	7.36	117.97	104.00
85	AA	472	A	C8-N9-C1'	7.35	140.94	127.70
34	BA	172	A	C5-C6-N6	-7.35	117.82	123.70
34	BA	1011	G	C8-N9-C4	7.35	109.34	106.40
36	BC	154	A	P-O5'-C5'	-7.35	109.14	120.90
38	BE	8	G	C4-N9-C1'	7.35	136.06	126.50
61	Bb	3	THR	CA-CB-CG2	-7.35	102.11	112.40
35	BB	18	A	P-O3'-C3'	-7.35	110.88	119.70
35	BB	569	G	C5-C6-O6	7.35	133.01	128.60
35	BB	825	U	O4'-C1'-N1	7.35	114.08	108.20
35	BB	1543	C	O4'-C1'-N1	7.35	114.08	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	224	C	P-O3'-C3'	7.35	128.52	119.70
85	AA	392	G	C5-C6-O6	7.35	133.01	128.60
85	AA	944	C	O4'-C1'-N1	7.35	114.08	108.20
85	AA	1158	U	C6-N1-C2	-7.35	116.59	121.00
86	AB	52	G	C1'-O4'-C4'	-7.35	104.02	109.90
19	AK	76	ARG	NE-CZ-NH1	7.35	123.97	120.30
34	BA	1492	G	O4'-C1'-N9	7.35	114.08	108.20
35	BB	920	C	O4'-C1'-N1	7.35	114.08	108.20
35	BB	1218	G	O4'-C4'-C3'	-7.35	96.65	104.00
38	BE	183	C	C2-N3-C4	-7.35	116.23	119.90
40	BG	20	U	C6-N1-C2	-7.35	116.59	121.00
42	BI	59	ARG	NE-CZ-NH1	7.35	123.97	120.30
85	AA	288	G	N1-C6-O6	7.35	124.31	119.90
85	AA	875	C	C5'-C4'-C3'	-7.35	104.24	116.00
85	AA	1278	C	N1-C1'-C2'	-7.35	103.92	112.00
85	AA	1820	G	C4-N9-C1'	-7.35	116.95	126.50
34	BA	481	A	C5-N7-C8	-7.35	100.23	103.90
34	BA	974	G	N9-C1'-C2'	-7.35	103.92	112.00
34	BA	1595	G	C5-C6-O6	-7.35	124.19	128.60
85	AA	1284	A	N1-C6-N6	-7.35	114.19	118.60
85	AA	1809	G	N1-C6-O6	7.35	124.31	119.90
34	BA	700	G	C1'-O4'-C4'	-7.34	104.03	109.90
34	BA	973	U	P-O5'-C5'	-7.34	109.15	120.90
34	BA	1412	G	N3-C2-N2	7.34	125.04	119.90
38	BE	13	A	C1'-O4'-C4'	-7.34	104.03	109.90
85	AA	130	G	C4'-C3'-C2'	-7.34	95.26	102.60
85	AA	731	U	O4'-C1'-N1	7.34	114.08	108.20
85	AA	1617	G	O4'-C1'-N9	7.34	114.08	108.20
34	BA	790	G	C5-C6-O6	-7.34	124.19	128.60
70	Bk	121	ARG	NE-CZ-NH2	-7.34	116.63	120.30
85	AA	64	A	O3'-P-O5'	7.34	117.95	104.00
85	AA	1905	A	C5'-C4'-C3'	-7.34	104.25	116.00
34	BA	986	G	C6-N1-C2	-7.34	120.70	125.10
85	AA	975	G	C1'-O4'-C4'	-7.34	104.03	109.90
85	AA	1217	U	C6-N1-C1'	7.34	131.48	121.20
34	BA	111	U	O4'-C1'-C2'	7.34	114.21	107.60
34	BA	1297	G	C8-N9-C1'	-7.34	117.46	127.00
34	BA	1297	G	N3-C4-N9	7.34	130.40	126.00
34	BA	1341	A	C3'-C2'-C1'	-7.34	95.63	101.50
34	BA	1518	A	C8-N9-C4	7.34	108.74	105.80
35	BB	119	G	C4'-C3'-C2'	7.34	109.94	102.60
38	BE	21	C	P-O5'-C5'	-7.34	109.16	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	104	G	N3-C4-N9	7.34	130.40	126.00
53	BT	170	ARG	NE-CZ-NH1	7.34	123.97	120.30
80	Bu	248	MET	CG-SD-CE	-7.34	88.46	100.20
85	AA	800	A	O4'-C1'-N9	7.34	114.07	108.20
85	AA	860	C	N3-C4-N4	-7.34	112.86	118.00
85	AA	966	G	C8-N9-C4	-7.34	103.47	106.40
85	AA	1181	U	O4'-C1'-N1	7.34	114.07	108.20
85	AA	2176	U	C4'-C3'-C2'	7.34	109.94	102.60
85	AA	2193	A	C5-N7-C8	-7.34	100.23	103.90
34	BA	1738	G	C8-N9-C4	-7.34	103.47	106.40
41	BH	129	G	P-O5'-C5'	7.34	132.64	120.90
85	AA	940	G	N1-C6-O6	7.34	124.30	119.90
85	AA	1679	U	O4'-C1'-N1	7.34	114.07	108.20
34	BA	867	C	O4'-C1'-N1	7.34	114.07	108.20
34	BA	894	G	N1-C2-N2	-7.34	109.60	116.20
35	BB	717	A	P-O5'-C5'	-7.34	109.16	120.90
35	BB	779	C	C2-N1-C1'	-7.34	110.73	118.80
35	BB	1538	G	C8-N9-C1'	7.34	136.54	127.00
38	BE	86	C	O4'-C1'-N1	7.34	114.07	108.20
39	BF	19	A	O4'-C1'-N9	7.34	114.07	108.20
40	BG	33	G	C8-N9-C4	7.34	109.33	106.40
40	BG	71	C	P-O3'-C3'	-7.34	110.90	119.70
40	BG	109	C	C2-N1-C1'	-7.34	110.73	118.80
85	AA	588	G	C4-N9-C1'	-7.33	116.97	126.50
85	AA	1966	C	P-O5'-C5'	7.33	132.63	120.90
1	A0	202	ARG	NE-CZ-NH1	7.33	123.97	120.30
32	AY	47	LYS	CA-CB-CG	7.33	129.53	113.40
34	BA	941	G	N3-C2-N2	7.33	125.03	119.90
34	BA	1526	C	C5'-C4'-C3'	-7.33	104.27	116.00
34	BA	1566	G	P-O3'-C3'	-7.33	110.90	119.70
35	BB	1278	A	C8-N9-C4	7.33	108.73	105.80
37	BD	32	A	C4-N9-C1'	-7.33	113.10	126.30
53	BT	133	ARG	NE-CZ-NH1	7.33	123.97	120.30
85	AA	833	U	C6-N1-C1'	-7.33	110.93	121.20
85	AA	1098	C	C6-N1-C2	-7.33	117.37	120.30
85	AA	2141	G	N1-C6-O6	7.33	124.30	119.90
21	AM	40	ARG	NE-CZ-NH1	7.33	123.97	120.30
34	BA	860	G	C5'-C4'-C3'	-7.33	104.27	116.00
34	BA	1739	G	C4-N9-C1'	-7.33	116.97	126.50
35	BB	408	U	P-O3'-C3'	-7.33	110.90	119.70
35	BB	1546	C	C3'-C2'-C1'	-7.33	95.64	101.50
58	BY	37	ARG	N-CA-CB	7.33	123.80	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	66	U	O3'-P-O5'	-7.33	90.07	104.00
34	BA	3	G	C5-C6-O6	-7.33	124.20	128.60
35	BB	1243	A	N1-C6-N6	-7.33	114.20	118.60
40	BG	14	G	C4'-C3'-C2'	-7.33	95.27	102.60
40	BG	78	C	C5'-C4'-C3'	-7.33	104.27	116.00
85	AA	1896	G	C2'-C3'-O3'	7.33	125.63	109.50
34	BA	320	G	C6-N1-C2	-7.33	120.70	125.10
35	BB	878	G	O3'-P-O5'	7.33	117.92	104.00
35	BB	1215	U	C2-N3-C4	-7.33	122.60	127.00
34	BA	30	A	C8-N9-C1'	7.33	140.89	127.70
34	BA	201	A	C4'-C3'-C2'	7.33	109.93	102.60
34	BA	1282	G	C6-N1-C2	-7.33	120.70	125.10
34	BA	1506	C	O5'-C5'-C4'	7.33	125.62	111.70
37	BD	14	C	C4'-C3'-C2'	7.33	109.93	102.60
37	BD	79	G	C8-N9-C1'	7.33	136.53	127.00
34	BA	372	U	P-O3'-C3'	-7.33	110.91	119.70
34	BA	1003	A	C8-N9-C4	7.33	108.73	105.80
34	BA	1385	U	P-O5'-C5'	7.33	132.62	120.90
34	BA	1450	G	C8-N9-C1'	7.33	136.52	127.00
38	BE	89	G	P-O3'-C3'	7.33	128.49	119.70
67	Bh	60	ALA	CB-CA-C	-7.33	99.11	110.10
85	AA	676	U	P-O3'-C3'	-7.33	110.91	119.70
85	AA	1525	C	N3-C2-O2	-7.33	116.77	121.90
34	BA	435	U	N1-C2-N3	7.32	119.30	114.90
34	BA	1607	U	N3-C2-O2	-7.32	117.07	122.20
35	BB	384	A	C8-N9-C4	-7.32	102.87	105.80
38	BE	163	A	N1-C2-N3	-7.32	125.64	129.30
85	AA	9	U	C1'-O4'-C4'	-7.32	104.04	109.90
85	AA	730	G	C8-N9-C4	-7.32	103.47	106.40
35	BB	428	G	P-O3'-C3'	-7.32	110.91	119.70
36	BC	130	U	P-O3'-C3'	-7.32	110.91	119.70
38	BE	203	C	C5-C4-N4	-7.32	115.08	120.20
85	AA	293	A	C3'-C2'-C1'	-7.32	95.64	101.50
85	AA	450	A	O4'-C1'-C2'	7.32	114.19	107.60
13	AE	106	TYR	CB-CG-CD2	-7.32	116.61	121.00
20	AL	82	MET	CG-SD-CE	-7.32	88.49	100.20
34	BA	809	U	C2-N1-C1'	-7.32	108.92	117.70
34	BA	1478	G	N1-C6-O6	7.32	124.29	119.90
35	BB	823	G	C4'-C3'-C2'	-7.32	95.28	102.60
35	BB	1407	U	P-O3'-C3'	7.32	128.48	119.70
35	BB	1546	C	P-O3'-C3'	-7.32	110.92	119.70
36	BC	71	A	N9-C1'-C2'	-7.32	103.95	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	30	C	O5'-C5'-C4'	-7.32	97.79	111.70
40	BG	31	G	C2-N3-C4	-7.32	108.24	111.90
85	AA	132	G	C5'-C4'-C3'	-7.32	104.29	116.00
35	BB	1220	A	O3'-P-O5'	-7.32	90.09	104.00
38	BE	68	U	N3-C2-O2	-7.32	117.08	122.20
85	AA	265	A	O4'-C1'-C2'	-7.32	98.48	105.80
85	AA	527	A	C6-N1-C2	-7.32	114.21	118.60
85	AA	716	G	C8-N9-C1'	7.32	136.51	127.00
85	AA	1863	A	P-O3'-C3'	-7.32	110.92	119.70
85	AA	2196	G	P-O3'-C3'	7.32	128.48	119.70
34	BA	814	C	C2-N1-C1'	7.32	126.85	118.80
35	BB	356	C	O4'-C1'-N1	7.32	114.05	108.20
35	BB	528	G	C8-N9-C1'	7.32	136.51	127.00
57	BX	82	PHE	CB-CG-CD2	-7.32	115.68	120.80
62	Bc	22	ARG	NE-CZ-NH1	7.32	123.96	120.30
13	AE	71	ASP	CB-CG-OD1	7.32	124.88	118.30
15	AG	127	ARG	NE-CZ-NH1	7.32	123.96	120.30
17	AI	147	ARG	NE-CZ-NH1	7.32	123.96	120.30
34	BA	274	C	O4'-C1'-N1	7.32	114.05	108.20
34	BA	554	A	N1-C6-N6	7.32	122.99	118.60
34	BA	653	U	C5-C6-N1	7.32	126.36	122.70
34	BA	997	U	N3-C2-O2	-7.32	117.08	122.20
34	BA	1097	G	O3'-P-O5'	-7.32	90.10	104.00
34	BA	1300	G	C4-N9-C1'	-7.32	116.99	126.50
35	BB	411	A	C8-N9-C4	7.32	108.73	105.80
35	BB	1384	A	C3'-C2'-C1'	-7.32	95.65	101.50
41	BH	47	G	N9-C1'-C2'	-7.32	103.95	112.00
42	BI	147	ARG	NE-CZ-NH2	-7.32	116.64	120.30
47	BN	206	ARG	CB-CA-C	7.32	125.03	110.40
56	BW	47	ARG	NE-CZ-NH2	-7.32	116.64	120.30
85	AA	556	C	C2'-C3'-O3'	7.32	125.59	109.50
85	AA	984	A	C1'-O4'-C4'	-7.32	104.05	109.90
85	AA	1221	G	N9-C1'-C2'	-7.32	103.95	112.00
85	AA	1991	C	N3-C4-N4	7.32	123.12	118.00
23	AP	229	ARG	NE-CZ-NH1	7.31	123.96	120.30
85	AA	537	G	C4-N9-C1'	-7.31	116.99	126.50
85	AA	742	U	O4'-C1'-N1	7.31	114.05	108.20
85	AA	821	U	C5-C6-N1	7.31	126.36	122.70
85	AA	1448	A	C6-N1-C2	-7.31	114.21	118.60
85	AA	1976	G	C4-N9-C1'	-7.31	116.99	126.50
34	BA	34	U	C1'-O4'-C4'	-7.31	104.05	109.90
35	BB	344	U	O4'-C1'-N1	7.31	114.05	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	38	G	C5-C6-O6	-7.31	124.21	128.60
85	AA	1867	G	P-O3'-C3'	-7.31	110.93	119.70
85	AA	1994	G	C4-N9-C1'	-7.31	116.99	126.50
34	BA	224	G	P-O3'-C3'	7.31	128.47	119.70
40	BG	37	G	C4-N9-C1'	-7.31	117.00	126.50
23	AP	259	MET	CG-SD-CE	-7.31	88.50	100.20
34	BA	1058	C	C4'-C3'-C2'	7.31	109.91	102.60
34	BA	1094	U	C2-N1-C1'	-7.31	108.93	117.70
34	BA	1807	G	N1-C2-N2	-7.31	109.62	116.20
35	BB	479	U	C2-N1-C1'	-7.31	108.93	117.70
35	BB	788	U	O4'-C1'-C2'	7.31	114.18	107.60
35	BB	995	C	C5'-C4'-C3'	-7.31	104.31	116.00
36	BC	34	U	C1'-O4'-C4'	-7.31	104.05	109.90
36	BC	125	A	N9-C1'-C2'	-7.31	103.96	112.00
62	Bc	82	MET	CG-SD-CE	-7.31	88.50	100.20
85	AA	233	C	P-O3'-C3'	-7.31	110.93	119.70
34	BA	2	A	C8-N9-C4	-7.31	102.88	105.80
34	BA	1296	U	OP1-P-OP2	-7.31	108.64	119.60
35	BB	359	A	O4'-C1'-N9	7.31	114.05	108.20
35	BB	832	C	C3'-C2'-C1'	-7.31	95.65	101.50
85	AA	1286	C	C5-C4-N4	-7.31	115.08	120.20
85	AA	2179	C	N1-C1'-C2'	-7.31	103.96	112.00
34	BA	1415	C	P-O5'-C5'	7.31	132.59	120.90
35	BB	1303	A	C5-C6-N6	-7.31	117.86	123.70
38	BE	13	A	N1-C6-N6	-7.31	114.22	118.60
38	BE	21	C	C2'-C3'-O3'	7.31	125.57	109.50
39	BF	23	G	C3'-C2'-C1'	-7.31	95.66	101.50
85	AA	146	U	N1-C2-N3	-7.31	110.52	114.90
85	AA	441	C	C5'-C4'-C3'	-7.31	104.31	116.00
85	AA	1447	U	C4'-C3'-C2'	-7.31	95.29	102.60
85	AA	1480	C	P-O3'-C3'	-7.31	110.93	119.70
34	BA	1587	C	O4'-C1'-N1	7.30	114.04	108.20
38	BE	66	A	C3'-C2'-C1'	-7.30	95.66	101.50
38	BE	190	U	C4'-C3'-C2'	7.30	109.91	102.60
40	BG	162	A	P-O5'-C5'	-7.30	109.21	120.90
85	AA	71	G	O4'-C1'-N9	7.30	114.04	108.20
85	AA	458	C	O4'-C1'-N1	7.30	114.04	108.20
85	AA	2017	U	O3'-P-O5'	-7.30	90.12	104.00
35	BB	943	U	O4'-C1'-N1	7.30	114.04	108.20
85	AA	1225	C	C6-N1-C2	-7.30	117.38	120.30
85	AA	1671	G	C4-N9-C1'	-7.30	117.01	126.50
85	AA	2167	A	C3'-C2'-C1'	-7.30	95.66	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	813	C	P-O3'-C3'	7.30	128.46	119.70
34	BA	1177	C	C5'-C4'-C3'	-7.30	104.32	116.00
34	BA	1298	U	O4'-C1'-N1	7.30	114.04	108.20
34	BA	1467	U	C1'-O4'-C4'	-7.30	104.06	109.90
38	BE	8	G	C8-N9-C1'	-7.30	117.51	127.00
85	AA	475	A	C8-N9-C1'	7.30	140.84	127.70
85	AA	537	G	C3'-C2'-C1'	-7.30	95.66	101.50
85	AA	867	G	C5-C6-O6	-7.30	124.22	128.60
85	AA	1124	G	C3'-C2'-C1'	-7.30	95.66	101.50
85	AA	2111	C	O4'-C1'-N1	7.30	114.04	108.20
1	A0	215	ARG	NE-CZ-NH1	7.30	123.95	120.30
34	BA	288	U	C3'-C2'-C1'	7.30	107.34	101.50
35	BB	746	A	O4'-C1'-N9	7.30	114.04	108.20
35	BB	1542	C	P-O5'-C5'	7.30	132.58	120.90
36	BC	50	C	C6-N1-C2	-7.30	117.38	120.30
37	BD	14	C	O5'-C5'-C4'	-7.30	97.83	111.70
35	BB	876	G	C8-N9-C1'	7.30	136.49	127.00
35	BB	1390	U	P-O5'-C5'	-7.30	109.22	120.90
37	BD	84	U	O4'-C1'-N1	7.30	114.04	108.20
85	AA	278	C	C4'-C3'-C2'	-7.30	95.30	102.60
85	AA	1485	G	N1-C6-O6	7.30	124.28	119.90
85	AA	2007	G	C4-N9-C1'	-7.30	117.01	126.50
13	AE	58	PHE	CB-CG-CD2	-7.30	115.69	120.80
34	BA	26	C	C5'-C4'-C3'	-7.30	104.33	116.00
34	BA	536	C	C6-N1-C2	-7.30	117.38	120.30
34	BA	1510	C	C5-C4-N4	7.30	125.31	120.20
35	BB	679	G	N3-C2-N2	7.30	125.01	119.90
35	BB	1410	G	O4'-C1'-N9	7.30	114.04	108.20
85	AA	17	C	C1'-O4'-C4'	-7.30	104.06	109.90
86	AB	44	G	N1-C6-O6	7.30	124.28	119.90
34	BA	192	G	C5-C6-O6	-7.29	124.22	128.60
34	BA	699	G	P-O3'-C3'	-7.29	110.95	119.70
35	BB	1341	U	O4'-C1'-N1	7.29	114.04	108.20
36	BC	134	G	C8-N9-C4	-7.29	103.48	106.40
85	AA	97	A	N9-C1'-C2'	-7.29	103.98	112.00
85	AA	746	G	O4'-C1'-N9	7.29	114.04	108.20
85	AA	868	A	C8-N9-C4	-7.29	102.88	105.80
34	BA	896	U	C3'-C2'-C1'	7.29	107.33	101.50
35	BB	1510	G	P-O3'-C3'	-7.29	110.95	119.70
40	BG	169	A	N9-C4-C5	-7.29	102.88	105.80
80	Bu	218	ARG	NE-CZ-NH2	-7.29	116.65	120.30
85	AA	1671	G	P-O3'-C3'	-7.29	110.95	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A2	140	MET	CA-CB-CG	7.29	125.70	113.30
4	A3	157	ARG	NE-CZ-NH2	-7.29	116.66	120.30
28	AU	105	ARG	NE-CZ-NH1	7.29	123.95	120.30
34	BA	665	C	P-O3'-C3'	-7.29	110.95	119.70
34	BA	1674	G	C8-N9-C1'	-7.29	117.52	127.00
35	BB	663	G	C6-N1-C2	-7.29	120.72	125.10
40	BG	95	U	P-O3'-C3'	-7.29	110.95	119.70
85	AA	90	A	O4'-C1'-C2'	-7.29	98.51	105.80
85	AA	188	G	O4'-C1'-N9	7.29	114.03	108.20
85	AA	260	A	N1-C6-N6	-7.29	114.23	118.60
85	AA	1084	A	N1-C6-N6	-7.29	114.22	118.60
85	AA	1281	G	N1-C6-O6	7.29	124.28	119.90
85	AA	1757	C	C6-N1-C1'	7.29	129.55	120.80
34	BA	362	G	C5-C6-N1	7.29	115.14	111.50
34	BA	424	U	C5'-C4'-O4'	7.29	117.85	109.10
34	BA	484	A	N1-C2-N3	-7.29	125.66	129.30
35	BB	1001	G	N3-C4-C5	-7.29	124.96	128.60
35	BB	1395	G	P-O3'-C3'	-7.29	110.95	119.70
36	BC	106	G	P-O3'-C3'	-7.29	110.95	119.70
41	BH	122	U	C1'-O4'-C4'	-7.29	104.07	109.90
48	BO	179	ARG	CD-NE-CZ	-7.29	113.40	123.60
85	AA	1277	C	C5-C4-N4	-7.29	115.10	120.20
85	AA	1362	A	C5'-C4'-O4'	-7.29	100.35	109.10
85	AA	1579	A	C1'-O4'-C4'	-7.29	104.07	109.90
24	AQ	24	ARG	NE-CZ-NH1	7.29	123.94	120.30
34	BA	1785	G	C4-N9-C1'	-7.29	117.03	126.50
35	BB	879	G	N3-C2-N2	7.29	125.00	119.90
38	BE	16	C	O5'-C5'-C4'	7.29	125.55	111.70
38	BE	32	U	C6-N1-C2	-7.29	116.63	121.00
80	Bu	190	ASN	CB-CA-C	-7.29	95.83	110.40
85	AA	260	A	C4'-C3'-C2'	-7.29	95.31	102.60
85	AA	1721	A	P-O5'-C5'	-7.29	109.24	120.90
86	AB	19	G	C5-C6-O6	-7.29	124.23	128.60
34	BA	460	G	C4-N9-C1'	-7.29	117.03	126.50
34	BA	1669	C	C2-N3-C4	-7.29	116.26	119.90
34	BA	1706	A	N1-C6-N6	-7.29	114.23	118.60
35	BB	837	A	C5-N7-C8	-7.29	100.26	103.90
35	BB	1513	U	C3'-C2'-C1'	-7.29	95.67	101.50
37	BD	80	G	C8-N9-C1'	7.29	136.47	127.00
38	BE	50	G	N3-C2-N2	7.29	125.00	119.90
65	Bf	134	PRO	N-CA-CB	7.29	112.04	103.30
85	AA	714	U	C6-N1-C2	-7.29	116.63	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1700	C	O4'-C1'-N1	7.29	114.03	108.20
85	AA	2155	U	N1-C2-N3	7.29	119.27	114.90
34	BA	22	C	C5'-C4'-C3'	-7.28	104.35	116.00
34	BA	237	A	C8-N9-C4	7.28	108.71	105.80
34	BA	519	G	C6-C5-N7	-7.28	126.03	130.40
34	BA	605	G	N1-C2-N3	7.28	128.27	123.90
34	BA	837	U	C3'-C2'-C1'	-7.28	95.67	101.50
35	BB	446	U	C5'-C4'-C3'	7.28	127.66	116.00
85	AA	147	G	C5'-C4'-C3'	-7.28	104.35	116.00
85	AA	865	G	C5'-C4'-C3'	-7.28	104.35	116.00
85	AA	879	G	C1'-O4'-C4'	-7.28	104.07	109.90
85	AA	978	U	C1'-O4'-C4'	-7.28	104.07	109.90
34	BA	1017	C	O5'-C5'-C4'	-7.28	97.86	111.70
34	BA	1697	U	N1-C2-O2	-7.28	117.70	122.80
85	AA	858	G	P-O3'-C3'	-7.28	110.96	119.70
85	AA	2102	A	N1-C6-N6	7.28	122.97	118.60
85	AA	2111	C	C6-N1-C1'	7.28	129.54	120.80
34	BA	218	G	C5-C6-O6	-7.28	124.23	128.60
34	BA	257	G	C8-N9-C4	7.28	109.31	106.40
34	BA	397	A	C1'-O4'-C4'	-7.28	104.08	109.90
34	BA	1216	G	P-O3'-C3'	7.28	128.44	119.70
35	BB	419	G	C1'-O4'-C4'	-7.28	104.08	109.90
35	BB	1463	A	C5'-C4'-C3'	7.28	127.65	116.00
38	BE	117	A	C3'-C2'-C1'	7.28	107.32	101.50
38	BE	142	A	C1'-O4'-C4'	-7.28	104.08	109.90
40	BG	117	C	C6-N1-C2	-7.28	117.39	120.30
85	AA	54	C	O4'-C1'-N1	7.28	114.02	108.20
85	AA	471	U	C2-N1-C1'	-7.28	108.96	117.70
85	AA	1878	C	O4'-C1'-N1	7.28	114.02	108.20
21	AM	131	ARG	NE-CZ-NH1	-7.28	116.66	120.30
34	BA	42	A	C5'-C4'-C3'	-7.28	104.35	116.00
85	AA	509	C	C4'-C3'-C2'	7.28	109.88	102.60
85	AA	1296	G	O4'-C1'-N9	7.28	114.02	108.20
85	AA	1540	A	P-O5'-C5'	7.28	132.55	120.90
85	AA	1608	U	C5'-C4'-C3'	-7.28	104.35	116.00
26	AS	136	GLN	N-CA-CB	7.28	123.70	110.60
34	BA	546	U	O3'-P-O5'	7.28	117.83	104.00
34	BA	1529	G	C5-C6-O6	7.28	132.97	128.60
35	BB	1017	U	P-O3'-C3'	-7.28	110.97	119.70
39	BF	3	A	P-O5'-C5'	7.28	132.54	120.90
64	Be	83	PHE	C-N-CA	7.28	139.89	121.70
85	AA	972	G	C5-C6-O6	-7.28	124.23	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1145	U	P-O3'-C3'	-7.28	110.97	119.70
85	AA	1206	A	C3'-C2'-C1'	-7.28	95.68	101.50
85	AA	1241	A	C4'-C3'-C2'	-7.28	95.32	102.60
85	AA	1676	G	C8-N9-C1'	7.28	136.46	127.00
34	BA	1828	A	O5'-P-OP2	-7.28	99.15	105.70
35	BB	87	G	C5-C6-O6	-7.28	124.23	128.60
41	BH	97	C	O4'-C1'-N1	7.28	114.02	108.20
85	AA	100	A	C4-C5-C6	-7.28	113.36	117.00
85	AA	415	G	C8-N9-C1'	7.28	136.46	127.00
85	AA	484	G	C8-N9-C4	7.28	109.31	106.40
85	AA	1167	G	N1-C6-O6	-7.28	115.53	119.90
85	AA	1366	A	C8-N9-C1'	7.28	140.79	127.70
85	AA	1487	G	C8-N9-C4	7.28	109.31	106.40
34	BA	941	G	N9-C1'-C2'	-7.27	104.00	112.00
34	BA	1497	A	C6-C5-N7	-7.27	127.21	132.30
35	BB	1179	C	C5'-C4'-C3'	7.27	127.64	116.00
38	BE	184	G	C5-C6-O6	7.27	132.96	128.60
85	AA	252	G	N3-C2-N2	7.27	124.99	119.90
85	AA	991	G	C4-C5-N7	-7.27	107.89	110.80
85	AA	1528	A	N1-C6-N6	-7.27	114.24	118.60
85	AA	1541	G	C5-C6-O6	-7.27	124.24	128.60
34	BA	18	G	C2-N3-C4	7.27	115.54	111.90
34	BA	590	U	C5'-C4'-C3'	-7.27	104.36	116.00
35	BB	333	C	P-O5'-C5'	7.27	132.54	120.90
35	BB	636	G	C5-C6-O6	-7.27	124.24	128.60
35	BB	653	G	C8-N9-C4	-7.27	103.49	106.40
35	BB	1083	C	C6-N1-C1'	-7.27	112.07	120.80
35	BB	1126	A	C1'-O4'-C4'	-7.27	104.08	109.90
36	BC	69	U	C2-N1-C1'	-7.27	108.97	117.70
40	BG	17	A	P-O3'-C3'	-7.27	110.97	119.70
41	BH	135	U	C5'-C4'-C3'	-7.27	104.36	116.00
85	AA	124	A	N1-C6-N6	-7.27	114.24	118.60
85	AA	1622	G	C2'-C3'-O3'	7.27	125.50	109.50
85	AA	2002	A	C4-N9-C1'	7.27	139.39	126.30
34	BA	961	C	C6-N1-C1'	-7.27	112.08	120.80
35	BB	532	C	C1'-O4'-C4'	-7.27	104.08	109.90
85	AA	39	A	C1'-O4'-C4'	-7.27	104.08	109.90
85	AA	2199	G	C8-N9-C1'	7.27	136.45	127.00
34	BA	482	C	P-O5'-C5'	-7.27	109.27	120.90
35	BB	823	G	C5-C6-N1	7.27	115.13	111.50
35	BB	1165	A	C5-C6-N1	7.27	121.33	117.70
56	BW	87	TRP	N-CA-CB	7.27	123.69	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
67	Bh	150	ARG	NE-CZ-NH2	-7.27	116.67	120.30
83	Bx	76	ARG	NE-CZ-NH1	7.27	123.93	120.30
34	BA	500	C	C3'-C2'-C1'	-7.27	95.69	101.50
34	BA	588	C	C6-N1-C2	-7.27	117.39	120.30
34	BA	1113	A	N1-C6-N6	7.27	122.96	118.60
34	BA	1737	A	C1'-O4'-C4'	-7.27	104.09	109.90
34	BA	1781	A	N1-C6-N6	7.27	122.96	118.60
37	BD	66	G	N1-C2-N2	-7.27	109.66	116.20
39	BF	72	A	P-O3'-C3'	7.27	128.42	119.70
85	AA	50	C	C6-N1-C2	-7.27	117.39	120.30
85	AA	111	A	C5'-C4'-O4'	7.27	117.82	109.10
85	AA	658	C	C3'-C2'-C1'	-7.27	95.69	101.50
85	AA	1576	G	C1'-O4'-C4'	-7.27	104.09	109.90
34	BA	1181	G	C4'-C3'-C2'	7.27	109.87	102.60
37	BD	37	G	O4'-C1'-N9	7.27	114.01	108.20
85	AA	570	U	N1-C2-N3	7.27	119.26	114.90
85	AA	1057	G	O4'-C1'-N9	7.27	114.01	108.20
34	BA	663	U	O4'-C1'-N1	7.26	114.01	108.20
34	BA	1487	U	C5-C4-O4	7.26	130.26	125.90
35	BB	1404	A	C5'-C4'-C3'	-7.26	104.38	116.00
36	BC	44	A	P-O5'-C5'	-7.26	109.28	120.90
38	BE	121	G	C5-C6-O6	-7.26	124.24	128.60
40	BG	157	A	C5-C6-N6	7.26	129.51	123.70
85	AA	822	U	C1'-O4'-C4'	-7.26	104.09	109.90
85	AA	2155	U	C6-N1-C1'	7.26	131.37	121.20
34	BA	655	U	C6-N1-C1'	7.26	131.37	121.20
35	BB	753	A	O4'-C1'-N9	7.26	114.01	108.20
85	AA	303	A	O4'-C4'-C3'	-7.26	96.74	104.00
85	AA	505	U	O5'-C5'-C4'	-7.26	97.90	111.70
85	AA	2138	G	O4'-C1'-N9	7.26	114.01	108.20
34	BA	495	A	C3'-C2'-C1'	-7.26	95.69	101.50
34	BA	1630	A	N9-C1'-C2'	-7.26	104.01	112.00
34	BA	1781	A	C8-N9-C4	-7.26	102.90	105.80
82	Bw	8	LEU	N-CA-CB	7.26	124.92	110.40
85	AA	70	U	C2-N1-C1'	-7.26	108.99	117.70
85	AA	587	G	C5'-C4'-C3'	7.26	127.62	116.00
85	AA	832	U	C5'-C4'-O4'	-7.26	100.39	109.10
85	AA	1124	G	C4-N9-C1'	-7.26	117.06	126.50
34	BA	373	G	C1'-O4'-C4'	-7.26	104.09	109.90
34	BA	576	C	C6-N1-C1'	-7.26	112.09	120.80
34	BA	894	G	C4'-C3'-C2'	7.26	109.86	102.60
34	BA	1087	A	C5-C6-N6	-7.26	117.89	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1500	G	C2-N3-C4	7.26	115.53	111.90
35	BB	391	G	N1-C6-O6	-7.26	115.54	119.90
35	BB	614	U	C6-N1-C2	-7.26	116.64	121.00
35	BB	855	G	N1-C6-O6	7.26	124.26	119.90
36	BC	29	C	P-O3'-C3'	7.26	128.41	119.70
36	BC	58	G	C5-C6-N1	7.26	115.13	111.50
37	BD	35	C	C1'-O4'-C4'	-7.26	104.09	109.90
43	BJ	51	ARG	NE-CZ-NH1	7.26	123.93	120.30
51	BR	62	ARG	NE-CZ-NH1	7.26	123.93	120.30
52	BS	14	ARG	NE-CZ-NH1	7.26	123.93	120.30
85	AA	111	A	O4'-C1'-N9	7.26	114.01	108.20
85	AA	130	G	O4'-C1'-N9	7.26	114.01	108.20
85	AA	1112	G	N1-C6-O6	7.26	124.26	119.90
85	AA	1287	C	P-O5'-C5'	-7.26	109.28	120.90
85	AA	1483	A	C1'-O4'-C4'	-7.26	104.09	109.90
86	AB	52	G	C5'-C4'-C3'	-7.26	104.39	116.00
85	AA	137	C	P-O3'-C3'	-7.26	110.99	119.70
85	AA	745	C	O4'-C1'-N1	7.26	114.01	108.20
34	BA	1197	U	C5-C4-O4	7.26	130.25	125.90
35	BB	128	C	C6-N1-C2	-7.26	117.40	120.30
35	BB	986	C	C2-N1-C1'	7.26	126.78	118.80
35	BB	1217	C	C5'-C4'-O4'	7.26	117.81	109.10
5	A4	155	MET	N-CA-CB	-7.25	97.54	110.60
35	BB	652	G	C6-N1-C2	-7.25	120.75	125.10
85	AA	844	C	C6-N1-C2	-7.25	117.40	120.30
34	BA	631	G	P-O3'-C3'	-7.25	111.00	119.70
34	BA	1482	A	C4'-C3'-C2'	7.25	109.85	102.60
35	BB	364	U	C2-N3-C4	-7.25	122.65	127.00
35	BB	834	U	O5'-P-OP2	7.25	119.40	110.70
37	BD	32	A	N9-C1'-C2'	-7.25	104.02	112.00
85	AA	1155	A	N1-C6-N6	7.25	122.95	118.60
85	AA	1520	A	P-O5'-C5'	-7.25	109.29	120.90
34	BA	487	A	C1'-O4'-C4'	-7.25	104.10	109.90
34	BA	766	A	C5-C6-N6	7.25	129.50	123.70
34	BA	1586	U	C2-N3-C4	-7.25	122.65	127.00
35	BB	375	G	N1-C6-O6	-7.25	115.55	119.90
35	BB	838	G	N3-C4-C5	-7.25	124.97	128.60
41	BH	90	C	OP1-P-OP2	-7.25	108.72	119.60
70	Bk	52	ARG	NE-CZ-NH1	7.25	123.93	120.30
85	AA	159	G	C8-N9-C1'	7.25	136.43	127.00
85	AA	2163	G	N1-C6-O6	-7.25	115.55	119.90
35	BB	1464	G	P-O3'-C3'	-7.25	111.00	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
74	Bo	25	ARG	NE-CZ-NH1	7.25	123.92	120.30
85	AA	1788	U	P-O3'-C3'	-7.25	111.00	119.70
34	BA	155	U	C6-N1-C2	-7.25	116.65	121.00
34	BA	301	U	N1-C1'-C2'	-7.25	104.03	112.00
34	BA	637	G	O4'-C1'-N9	7.25	114.00	108.20
35	BB	1200	A	C5-C6-N1	7.25	121.33	117.70
38	BE	34	C	O4'-C1'-N1	7.25	114.00	108.20
85	AA	173	A	C5'-C4'-C3'	-7.25	104.40	116.00
85	AA	1671	G	C8-N9-C1'	7.25	136.42	127.00
25	AR	26	ASN	CA-C-N	-7.25	101.26	117.20
34	BA	1341	A	C1'-O4'-C4'	-7.25	104.10	109.90
34	BA	1496	G	C2'-C3'-O3'	7.25	125.44	109.50
35	BB	26	C	C5'-C4'-C3'	-7.25	104.41	116.00
35	BB	899	C	C5-C6-N1	7.25	124.62	121.00
35	BB	1072	C	C6-N1-C2	7.25	123.20	120.30
53	BT	60	ARG	NE-CZ-NH1	7.25	123.92	120.30
85	AA	1174	G	C6-C5-N7	-7.25	126.05	130.40
85	AA	1300	A	O4'-C1'-N9	7.25	114.00	108.20
85	AA	1958	C	O4'-C1'-N1	7.25	114.00	108.20
34	BA	71	G	N3-C4-C5	-7.25	124.98	128.60
34	BA	1494	G	C6-N1-C2	-7.25	120.75	125.10
35	BB	63	A	O4'-C1'-N9	7.25	114.00	108.20
37	BD	26	C	O4'-C1'-N1	7.25	114.00	108.20
37	BD	98	G	C4-N9-C1'	-7.25	117.08	126.50
41	BH	10	U	C5-C4-O4	7.25	130.25	125.90
41	BH	35	G	C8-N9-C1'	7.25	136.42	127.00
82	Bw	229	MET	CG-SD-CE	-7.25	88.61	100.20
85	AA	486	G	C6-N1-C2	-7.25	120.75	125.10
5	A4	143	ARG	NE-CZ-NH2	-7.24	116.68	120.30
18	AJ	64	ASN	CA-CB-CG	-7.24	97.47	113.40
35	BB	133	G	N9-C1'-C2'	-7.24	104.03	112.00
35	BB	942	G	O4'-C1'-N9	7.24	113.99	108.20
35	BB	1046	C	O4'-C1'-N1	7.24	114.00	108.20
38	BE	126	G	C6-N1-C2	-7.24	120.75	125.10
85	AA	1713	A	C5-C6-N6	-7.24	117.91	123.70
85	AA	2229	G	N1-C6-O6	-7.24	115.55	119.90
38	BE	182	U	C5'-C4'-C3'	-7.24	104.41	116.00
40	BG	80	G	P-O3'-C3'	-7.24	111.01	119.70
85	AA	1924	C	O4'-C1'-N1	7.24	113.99	108.20
34	BA	67	A	C5'-C4'-C3'	-7.24	104.42	116.00
35	BB	1287	U	P-O3'-C3'	-7.24	111.01	119.70
37	BD	51	G	C1'-O4'-C4'	-7.24	104.11	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1099	U	P-O3'-C3'	-7.24	111.01	119.70
34	BA	1615	A	N9-C1'-C2'	-7.24	104.04	112.00
34	BA	1834	A	C5-C6-N1	7.24	121.32	117.70
35	BB	3	C	C1'-O4'-C4'	-7.24	104.11	109.90
35	BB	49	A	P-O3'-C3'	7.24	128.38	119.70
35	BB	1396	G	N3-C2-N2	7.24	124.97	119.90
35	BB	1422	G	N3-C4-N9	-7.24	121.66	126.00
35	BB	1489	A	O4'-C1'-N9	7.24	113.99	108.20
38	BE	129	G	C1'-O4'-C4'	-7.24	104.11	109.90
56	BW	3	LYS	C-N-CA	7.24	139.80	121.70
85	AA	150	U	N3-C2-O2	-7.24	117.13	122.20
85	AA	2218	G	C5-C6-O6	-7.24	124.26	128.60
34	BA	236	A	C6-N1-C2	-7.24	114.26	118.60
34	BA	991	U	C2-N3-C4	-7.24	122.66	127.00
34	BA	1194	G	O4'-C1'-C2'	7.24	114.11	107.60
34	BA	1531	G	C5-C6-N1	7.24	115.12	111.50
85	AA	999	A	O3'-P-O5'	7.24	117.75	104.00
85	AA	1114	A	C2-N3-C4	-7.24	106.98	110.60
35	BB	119	G	O4'-C1'-C2'	7.24	114.11	107.60
35	BB	256	G	C5-C6-O6	-7.24	124.26	128.60
35	BB	378	C	C4'-C3'-C2'	-7.24	95.36	102.60
35	BB	835	C	N3-C4-C5	-7.24	119.01	121.90
37	BD	119	U	C1'-O4'-C4'	-7.24	104.11	109.90
40	BG	54	G	C4-N9-C1'	-7.24	117.09	126.50
85	AA	187	C	C2-N3-C4	-7.24	116.28	119.90
85	AA	2121	G	C4-N9-C1'	-7.24	117.09	126.50
82	Bw	16	MET	CG-SD-CE	-7.23	88.63	100.20
85	AA	70	U	P-O5'-C5'	-7.23	109.33	120.90
85	AA	100	A	N3-C4-N9	-7.23	121.61	127.40
85	AA	170	C	C2'-C3'-O3'	7.23	125.42	109.50
34	BA	247	U	O4'-C1'-N1	7.23	113.99	108.20
34	BA	1176	C	C5-C4-N4	7.23	125.26	120.20
62	Bc	137	HIS	CA-CB-CG	7.23	125.90	113.60
85	AA	726	U	C2-N1-C1'	7.23	126.38	117.70
85	AA	1196	C	P-O3'-C3'	-7.23	111.02	119.70
85	AA	1525	C	C2'-C3'-O3'	7.23	125.41	109.50
34	BA	1644	A	C8-N9-C4	7.23	108.69	105.80
35	BB	104	G	C8-N9-C4	7.23	109.29	106.40
35	BB	623	A	C4-N9-C1'	-7.23	113.29	126.30
35	BB	797	C	OP2-P-O3'	7.23	121.11	105.20
35	BB	1124	G	P-O5'-C5'	7.23	132.47	120.90
40	BG	24	A	C4-N9-C1'	-7.23	113.28	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	359	A	C5'-C4'-C3'	7.23	127.57	116.00
85	AA	597	A	P-O5'-C5'	-7.23	109.33	120.90
85	AA	1276	A	P-O3'-C3'	-7.23	111.03	119.70
85	AA	1669	G	C8-N9-C1'	7.23	136.40	127.00
85	AA	2129	U	P-O3'-C3'	-7.23	111.02	119.70
36	BC	97	U	P-O3'-C3'	-7.23	111.03	119.70
40	BG	105	A	C4'-C3'-C2'	7.23	109.83	102.60
85	AA	354	C	C3'-C2'-C1'	-7.23	95.72	101.50
34	BA	975	A	C1'-O4'-C4'	-7.23	104.12	109.90
34	BA	1310	C	O4'-C1'-N1	7.23	113.98	108.20
35	BB	825	U	C3'-C2'-C1'	-7.23	95.72	101.50
35	BB	1246	C	C1'-O4'-C4'	-7.23	104.12	109.90
37	BD	70	C	P-O3'-C3'	-7.23	111.03	119.70
37	BD	106	G	C4-N9-C1'	-7.23	117.11	126.50
41	BH	36	C	C5'-C4'-C3'	-7.23	104.44	116.00
56	BW	47	ARG	NE-CZ-NH1	7.23	123.91	120.30
57	BX	59	THR	C-N-CA	7.23	139.77	121.70
80	Bu	141	ASP	C-N-CA	7.23	139.77	121.70
85	AA	1170	C	C5-C4-N4	7.23	125.26	120.20
85	AA	1668	G	C4-N9-C1'	-7.23	117.10	126.50
85	AA	1731	G	C8-N9-C4	-7.23	103.51	106.40
34	BA	113	G	C5'-C4'-O4'	-7.23	100.43	109.10
38	BE	127	G	N3-C4-C5	-7.23	124.99	128.60
40	BG	140	G	P-O5'-C5'	7.23	132.46	120.90
85	AA	823	C	P-O3'-C3'	-7.23	111.03	119.70
85	AA	1975	G	C8-N9-C4	-7.23	103.51	106.40
34	BA	1118	C	O4'-C1'-N1	7.22	113.98	108.20
34	BA	1577	U	N3-C2-O2	-7.22	117.14	122.20
35	BB	556	U	C1'-O4'-C4'	-7.22	104.12	109.90
35	BB	700	C	C4-C5-C6	-7.22	113.79	117.40
38	BE	7	U	C2-N3-C4	-7.22	122.67	127.00
57	BX	82	PHE	CA-CB-CG	7.22	131.24	113.90
85	AA	39	A	N9-C1'-C2'	-7.22	104.05	112.00
85	AA	57	G	P-O5'-C5'	-7.22	109.34	120.90
85	AA	211	C	O3'-P-O5'	7.22	117.73	104.00
85	AA	1978	G	N9-C4-C5	-7.22	102.51	105.40
34	BA	210	G	C8-N9-C1'	7.22	136.39	127.00
85	AA	794	A	C1'-O4'-C4'	-7.22	104.12	109.90
85	AA	2215	C	C2-N1-C1'	-7.22	110.86	118.80
34	BA	471	U	C2-N3-C4	-7.22	122.67	127.00
34	BA	1519	G	C1'-O4'-C4'	-7.22	104.12	109.90
36	BC	125	A	O4'-C1'-N9	7.22	113.98	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
77	Br	155	ASP	CB-CG-OD1	7.22	124.80	118.30
34	BA	72	U	O5'-P-OP2	-7.22	99.20	105.70
34	BA	257	G	C1'-O4'-C4'	-7.22	104.12	109.90
34	BA	1745	G	O4'-C1'-N9	7.22	113.97	108.20
37	BD	73	U	C5-C6-N1	-7.22	119.09	122.70
38	BE	142	A	P-O3'-C3'	7.22	128.36	119.70
65	Bf	173	TRP	CB-CG-CD2	-7.22	117.21	126.60
85	AA	448	G	O4'-C1'-N9	7.22	113.98	108.20
85	AA	877	G	N1-C6-O6	-7.22	115.57	119.90
85	AA	2210	C	C2-N1-C1'	-7.22	110.86	118.80
1	A0	117	ARG	NE-CZ-NH1	7.22	123.91	120.30
35	BB	1134	G	P-O5'-C5'	7.22	132.45	120.90
85	AA	701	C	O4'-C1'-N1	7.22	113.97	108.20
85	AA	1923	A	O4'-C1'-C2'	-7.22	98.58	105.80
30	AW	6	SER	C-N-CA	7.22	139.74	121.70
34	BA	1263	A	N1-C6-N6	-7.22	114.27	118.60
34	BA	1286	C	N3-C4-N4	-7.22	112.95	118.00
35	BB	1001	G	O3'-P-O5'	7.22	117.71	104.00
41	BH	14	C	O4'-C1'-N1	7.22	113.97	108.20
41	BH	47	G	N3-C4-C5	-7.22	124.99	128.60
85	AA	519	A	C8-N9-C4	7.22	108.69	105.80
85	AA	1176	C	C5'-C4'-C3'	7.22	127.55	116.00
85	AA	1935	G	C5-C6-O6	-7.22	124.27	128.60
10	A9	99	ARG	NE-CZ-NH2	-7.21	116.69	120.30
35	BB	22	A	P-O3'-C3'	7.21	128.36	119.70
35	BB	751	A	N1-C6-N6	-7.21	114.27	118.60
35	BB	816	U	N3-C4-C5	7.21	118.93	114.60
35	BB	904	C	C6-N1-C2	-7.21	117.41	120.30
85	AA	1998	A	N1-C6-N6	7.21	122.93	118.60
35	BB	1045	G	O5'-C5'-C4'	7.21	125.40	111.70
85	AA	354	C	O4'-C1'-N1	7.21	113.97	108.20
85	AA	478	U	C4-C5-C6	-7.21	115.37	119.70
85	AA	496	C	C2-N1-C1'	-7.21	110.87	118.80
85	AA	1100	U	C6-N1-C2	-7.21	116.67	121.00
34	BA	678	C	C5-C4-N4	-7.21	115.15	120.20
34	BA	1102	A	C6-N1-C2	-7.21	114.27	118.60
40	BG	133	C	N1-C1'-C2'	-7.21	104.07	112.00
41	BH	67	G	C5'-C4'-C3'	-7.21	104.46	116.00
84	By	111	ARG	NE-CZ-NH2	-7.21	116.69	120.30
85	AA	992	G	P-O3'-C3'	-7.21	111.05	119.70
85	AA	1822	G	C8-N9-C1'	7.21	136.38	127.00
85	AA	1883	C	O4'-C1'-N1	7.21	113.97	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2139	G	O4'-C1'-N9	7.21	113.97	108.20
34	BA	1499	A	C8-N9-C1'	7.21	140.68	127.70
35	BB	338	C	C6-N1-C2	-7.21	117.42	120.30
36	BC	31	A	C8-N9-C4	7.21	108.68	105.80
34	BA	1692	U	N3-C2-O2	-7.21	117.15	122.20
35	BB	56	U	P-O5'-C5'	7.21	132.43	120.90
35	BB	649	A	P-O3'-C3'	-7.21	111.05	119.70
35	BB	1043	C	O4'-C1'-N1	7.21	113.97	108.20
38	BE	163	A	P-O3'-C3'	-7.21	111.05	119.70
38	BE	183	C	C5-C4-N4	-7.21	115.15	120.20
74	Bo	13	ARG	NE-CZ-NH1	7.21	123.90	120.30
84	By	74	HIS	CA-CB-CG	-7.21	101.35	113.60
85	AA	586	G	P-O5'-C5'	7.21	132.44	120.90
85	AA	1120	G	P-O3'-C3'	-7.21	111.05	119.70
85	AA	1996	A	C5-C6-N1	-7.21	114.10	117.70
5	A4	67	LEU	N-CA-CB	-7.21	95.98	110.40
34	BA	288	U	C6-N1-C2	-7.21	116.68	121.00
36	BC	15	G	P-O3'-C3'	7.21	128.35	119.70
38	BE	21	C	N3-C2-O2	-7.21	116.86	121.90
85	AA	906	U	O4'-C1'-N1	7.21	113.97	108.20
85	AA	1501	A	C5'-C4'-C3'	-7.21	104.47	116.00
85	AA	1534	A	P-O3'-C3'	-7.21	111.05	119.70
85	AA	1955	U	N1-C2-N3	7.21	119.22	114.90
34	BA	798	G	C3'-C2'-C1'	-7.21	95.74	101.50
38	BE	43	A	C5'-C4'-C3'	7.21	127.53	116.00
38	BE	180	G	C5'-C4'-C3'	-7.21	104.47	116.00
34	BA	140	C	C5'-C4'-C3'	-7.20	104.47	116.00
35	BB	45	A	C2-N3-C4	-7.20	107.00	110.60
35	BB	703	U	C5'-C4'-C3'	-7.20	104.47	116.00
35	BB	874	G	O4'-C1'-N9	7.20	113.96	108.20
40	BG	36	G	C3'-C2'-C1'	-7.20	95.74	101.50
85	AA	715	G	O4'-C4'-C3'	-7.20	96.80	104.00
85	AA	747	U	C1'-O4'-C4'	-7.20	104.14	109.90
34	BA	1766	G	O4'-C1'-N9	7.20	113.96	108.20
53	BT	165	LYS	CB-CA-C	-7.20	96.00	110.40
78	Bs	47	MET	CG-SD-CE	-7.20	88.68	100.20
34	BA	1742	G	C5-C6-O6	-7.20	124.28	128.60
35	BB	829	C	C5'-C4'-C3'	7.20	127.52	116.00
40	BG	84	U	P-O3'-C3'	-7.20	111.06	119.70
85	AA	21	U	C2'-C3'-O3'	7.20	125.34	109.50
85	AA	761	G	C4-N9-C1'	7.20	135.86	126.50
85	AA	1921	G	O4'-C4'-C3'	-7.20	96.80	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
7	A6	39	ARG	NE-CZ-NH1	7.20	123.90	120.30
23	AP	78	ASP	CB-CA-C	7.20	124.80	110.40
34	BA	384	U	C6-N1-C2	-7.20	116.68	121.00
34	BA	432	A	C5'-C4'-C3'	-7.20	104.48	116.00
35	BB	77	A	C5'-C4'-C3'	7.20	127.52	116.00
35	BB	1346	A	N1-C6-N6	7.20	122.92	118.60
35	BB	1465	U	P-O3'-C3'	7.20	128.34	119.70
40	BG	67	A	C1'-O4'-C4'	-7.20	104.14	109.90
67	Bh	122	THR	N-CA-C	-7.20	91.56	111.00
85	AA	189	G	N1-C6-O6	-7.20	115.58	119.90
85	AA	341	C	O4'-C1'-N1	7.20	113.96	108.20
85	AA	415	G	N1-C6-O6	7.20	124.22	119.90
85	AA	2160	U	P-O3'-C3'	7.20	128.34	119.70
34	BA	1707	C	N1-C2-N3	7.20	124.24	119.20
34	BA	252	A	C5'-C4'-C3'	-7.20	104.49	116.00
34	BA	1207	A	O4'-C1'-N9	7.20	113.96	108.20
35	BB	1522	G	C3'-C2'-C1'	-7.20	95.74	101.50
38	BE	162	U	P-O5'-C5'	7.20	132.41	120.90
40	BG	32	U	C5'-C4'-C3'	-7.20	104.49	116.00
40	BG	178	G	C8-N9-C1'	7.20	136.35	127.00
85	AA	908	C	P-O3'-C3'	-7.20	111.06	119.70
86	AB	25	C	O4'-C1'-N1	7.20	113.96	108.20
34	BA	126	G	C5-C6-O6	-7.19	124.28	128.60
34	BA	1101	A	C1'-O4'-C4'	-7.19	104.14	109.90
35	BB	798	A	C6-C5-N7	-7.19	127.26	132.30
35	BB	1306	G	O3'-P-O5'	-7.19	90.33	104.00
35	BB	1315	C	P-O5'-C5'	7.19	132.41	120.90
51	BR	133	HIS	CA-CB-CG	-7.19	101.37	113.60
82	Bw	103	ARG	NE-CZ-NH2	-7.19	116.70	120.30
2	A1	146	TYR	CB-CG-CD1	-7.19	116.68	121.00
34	BA	210	G	C4-N9-C1'	-7.19	117.15	126.50
34	BA	1663	U	N3-C2-O2	-7.19	117.17	122.20
35	BB	390	G	O4'-C1'-N9	7.19	113.95	108.20
35	BB	764	C	N3-C2-O2	-7.19	116.86	121.90
35	BB	802	G	N9-C1'-C2'	-7.19	104.09	112.00
35	BB	839	G	C5'-C4'-C3'	7.19	127.51	116.00
37	BD	50	A	C5'-C4'-C3'	7.19	127.51	116.00
41	BH	4	U	C6-N1-C1'	7.19	131.27	121.20
41	BH	35	G	C8-N9-C4	-7.19	103.52	106.40
41	BH	54	U	O4'-C1'-N1	7.19	113.95	108.20
80	Bu	134	THR	N-CA-CB	7.19	123.97	110.30
85	AA	41	G	C5'-C4'-C3'	-7.19	104.49	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	400	G	C4-N9-C1'	-7.19	117.15	126.50
85	AA	814	G	C6-N1-C2	-7.19	120.78	125.10
85	AA	1989	A	C4-C5-C6	-7.19	113.40	117.00
23	AP	136	ARG	NE-CZ-NH1	7.19	123.89	120.30
34	BA	161	U	N3-C4-C5	-7.19	110.28	114.60
34	BA	201	A	C5'-C4'-C3'	-7.19	104.49	116.00
34	BA	365	A	C4'-C3'-C2'	7.19	109.79	102.60
34	BA	967	C	C3'-C2'-C1'	-7.19	95.75	101.50
34	BA	1476	G	C8-N9-C1'	7.19	136.35	127.00
34	BA	1485	U	N3-C2-O2	-7.19	117.17	122.20
36	BC	107	C	C1'-O4'-C4'	-7.19	104.15	109.90
40	BG	179	C	O4'-C1'-N1	7.19	113.95	108.20
85	AA	99	U	C2-N3-C4	-7.19	122.69	127.00
85	AA	145	C	O4'-C1'-N1	7.19	113.95	108.20
85	AA	815	G	N9-C4-C5	-7.19	102.52	105.40
85	AA	969	U	C6-N1-C2	-7.19	116.69	121.00
85	AA	1831	U	O4'-C1'-N1	7.19	113.95	108.20
85	AA	2071	U	C2-N1-C1'	-7.19	109.07	117.70
85	AA	2126	U	O4'-C1'-N1	7.19	113.95	108.20
34	BA	1070	G	P-O3'-C3'	-7.19	111.07	119.70
34	BA	1488	C	N1-C2-N3	-7.19	114.17	119.20
35	BB	1199	A	C5-C6-N6	7.19	129.45	123.70
85	AA	161	A	C5'-C4'-C3'	-7.19	104.50	116.00
85	AA	2113	U	C1'-O4'-C4'	-7.19	104.15	109.90
34	BA	110	C	C5-C6-N1	-7.19	117.41	121.00
34	BA	151	A	P-O5'-C5'	-7.19	109.40	120.90
34	BA	689	C	C5'-C4'-C3'	7.19	127.50	116.00
34	BA	1178	U	C5'-C4'-C3'	-7.19	104.50	116.00
34	BA	1231	C	O4'-C1'-N1	7.19	113.95	108.20
38	BE	6	A	C5'-C4'-C3'	-7.19	104.50	116.00
85	AA	614	U	C4'-C3'-C2'	-7.19	95.41	102.60
86	AB	13	C	C4'-C3'-C2'	-7.19	95.41	102.60
22	AO	84	ARG	NE-CZ-NH1	7.19	123.89	120.30
34	BA	1441	C	C6-N1-C2	-7.19	117.43	120.30
34	BA	1445	U	C6-N1-C1'	-7.19	111.14	121.20
85	AA	751	C	C4'-C3'-C2'	-7.19	95.41	102.60
11	AC	45	ARG	NE-CZ-NH1	7.18	123.89	120.30
34	BA	5	C	O4'-C1'-N1	7.18	113.95	108.20
34	BA	222	C	C2-N3-C4	-7.18	116.31	119.90
34	BA	502	U	C6-N1-C1'	7.18	131.26	121.20
34	BA	556	A	O4'-C1'-N9	7.18	113.95	108.20
34	BA	803	U	C2-N1-C1'	-7.18	109.08	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	816	G	N1-C6-O6	7.18	124.21	119.90
34	BA	1538	G	C5'-C4'-C3'	7.18	127.50	116.00
35	BB	137	A	C4-C5-C6	-7.18	113.41	117.00
35	BB	1458	U	C2-N3-C4	-7.18	122.69	127.00
39	BF	65	U	C6-N1-C2	-7.18	116.69	121.00
85	AA	580	C	P-O3'-C3'	-7.18	111.08	119.70
85	AA	1876	U	C2-N1-C1'	-7.18	109.08	117.70
34	BA	436	U	C6-N1-C1'	7.18	131.25	121.20
34	BA	512	U	C2'-C3'-O3'	7.18	125.30	109.50
34	BA	536	C	C6-N1-C1'	7.18	129.42	120.80
34	BA	602	G	N1-C6-O6	7.18	124.21	119.90
34	BA	984	U	C2-N3-C4	-7.18	122.69	127.00
34	BA	1729	G	P-O3'-C3'	-7.18	111.08	119.70
35	BB	449	C	C2-N3-C4	-7.18	116.31	119.90
38	BE	149	A	N7-C8-N9	7.18	117.39	113.80
85	AA	1814	U	C2-N3-C4	-7.18	122.69	127.00
85	AA	1988	A	P-O3'-C3'	7.18	128.32	119.70
85	AA	2180	C	C5'-C4'-C3'	7.18	127.49	116.00
34	BA	251	U	C5'-C4'-C3'	-7.18	104.51	116.00
35	BB	804	U	P-O5'-C5'	-7.18	109.41	120.90
85	AA	771	A	C8-N9-C4	-7.18	102.93	105.80
34	BA	755	G	C3'-C2'-C1'	-7.18	95.76	101.50
34	BA	795	G	C5-C6-O6	-7.18	124.29	128.60
35	BB	580	A	C5'-C4'-C3'	7.18	127.49	116.00
35	BB	620	G	P-O3'-C3'	-7.18	111.08	119.70
42	BI	187	ARG	NE-CZ-NH2	-7.18	116.71	120.30
85	AA	642	G	C4-N9-C1'	-7.18	117.17	126.50
85	AA	681	G	O5'-C5'-C4'	7.18	125.34	111.70
24	AQ	76	THR	N-CA-C	7.18	130.38	111.00
34	BA	1579	G	O4'-C1'-C2'	-7.18	98.62	105.80
35	BB	673	C	O4'-C1'-N1	7.18	113.94	108.20
65	Bf	318	TYR	CB-CG-CD2	-7.18	116.69	121.00
77	Br	301	LYS	N-CA-CB	-7.18	97.68	110.60
34	BA	758	G	P-O5'-C5'	-7.18	109.42	120.90
34	BA	1053	U	P-O3'-C3'	7.18	128.31	119.70
34	BA	1402	C	O4'-C1'-N1	7.18	113.94	108.20
34	BA	1510	C	N1-C2-O2	7.18	123.20	118.90
34	BA	1826	C	C2-N3-C4	-7.18	116.31	119.90
36	BC	41	A	C5'-C4'-C3'	7.18	127.48	116.00
37	BD	119	U	O4'-C1'-N1	7.18	113.94	108.20
70	Bk	94	ARG	NE-CZ-NH2	-7.18	116.71	120.30
79	Bt	10	MET	CG-SD-CE	-7.18	88.72	100.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	477	U	P-O3'-C3'	7.18	128.31	119.70
85	AA	1603	G	N1-C6-O6	7.18	124.21	119.90
34	BA	805	A	C4'-C3'-C2'	-7.17	95.42	102.60
34	BA	873	G	O4'-C1'-N9	7.17	113.94	108.20
35	BB	701	U	C6-N1-C2	-7.17	116.70	121.00
35	BB	1043	C	O5'-P-OP1	7.17	119.31	110.70
39	BF	38	C	C5-C6-N1	7.17	124.59	121.00
85	AA	406	U	O4'-C1'-N1	7.17	113.94	108.20
85	AA	595	A	O4'-C1'-N9	7.17	113.94	108.20
85	AA	1301	C	N3-C2-O2	-7.17	116.88	121.90
85	AA	2231	G	O3'-P-O5'	-7.17	90.37	104.00
35	BB	577	U	P-O5'-C5'	7.17	132.38	120.90
35	BB	1326	U	O4'-C1'-N1	7.17	113.94	108.20
39	BF	70	A	P-O3'-C3'	-7.17	111.09	119.70
41	BH	10	U	N3-C2-O2	-7.17	117.18	122.20
85	AA	163	C	N3-C4-N4	7.17	123.02	118.00
85	AA	269	G	N1-C6-O6	7.17	124.20	119.90
15	AG	121	ARG	NE-CZ-NH2	-7.17	116.71	120.30
34	BA	1172	C	O4'-C1'-C2'	7.17	114.05	107.60
35	BB	877	A	O4'-C1'-N9	7.17	113.94	108.20
38	BE	39	U	O4'-C1'-N1	7.17	113.94	108.20
38	BE	67	A	C1'-O4'-C4'	-7.17	104.16	109.90
41	BH	122	U	C6-N1-C1'	-7.17	111.16	121.20
61	Bb	76	ASP	CB-CG-OD2	7.17	124.75	118.30
85	AA	477	U	C2-N3-C4	-7.17	122.70	127.00
85	AA	1393	C	O4'-C1'-N1	7.17	113.94	108.20
85	AA	1647	G	C4'-C3'-C2'	-7.17	95.43	102.60
34	BA	377	G	N3-C2-N2	7.17	124.92	119.90
85	AA	861	G	C5'-C4'-C3'	-7.17	104.53	116.00
4	A3	224	ARG	NE-CZ-NH1	7.17	123.88	120.30
34	BA	696	A	O4'-C1'-N9	7.17	113.94	108.20
35	BB	483	C	N1-C1'-C2'	-7.17	104.11	112.00
35	BB	1208	G	O4'-C1'-N9	7.17	113.94	108.20
40	BG	102	G	C5-C6-N1	7.17	115.08	111.50
85	AA	980	U	O4'-C1'-N1	7.17	113.93	108.20
34	BA	1709	A	C3'-C2'-C1'	-7.17	95.77	101.50
35	BB	1089	A	C8-N9-C4	7.17	108.67	105.80
35	BB	1209	A	N1-C6-N6	7.17	122.90	118.60
35	BB	1425	A	C8-N9-C4	7.17	108.67	105.80
38	BE	196	C	N3-C4-N4	-7.17	112.98	118.00
39	BF	12	U	C6-N1-C1'	-7.17	111.17	121.20
40	BG	137	G	C5-C6-N1	7.17	115.08	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	BN	9	PRO	N-CA-CB	-7.17	94.70	103.30
81	Bv	105	ARG	NE-CZ-NH1	7.17	123.88	120.30
85	AA	767	A	P-O5'-C5'	7.17	132.37	120.90
85	AA	975	G	C5'-C4'-C3'	-7.17	104.53	116.00
34	BA	1458	A	N1-C6-N6	7.17	122.90	118.60
35	BB	1161	G	N9-C1'-C2'	-7.17	104.12	112.00
85	AA	291	G	N9-C1'-C2'	-7.17	104.12	112.00
85	AA	391	G	O4'-C1'-N9	7.17	113.93	108.20
85	AA	483	G	C5'-C4'-O4'	7.17	117.70	109.10
85	AA	1892	G	C5'-C4'-C3'	-7.17	104.54	116.00
34	BA	19	G	N1-C6-O6	-7.16	115.60	119.90
34	BA	996	U	P-O5'-C5'	7.16	132.36	120.90
35	BB	830	G	C3'-C2'-C1'	-7.16	95.77	101.50
35	BB	1348	C	C1'-O4'-C4'	-7.16	104.17	109.90
40	BG	33	G	N1-C2-N3	-7.16	119.60	123.90
85	AA	919	U	C6-N1-C1'	-7.16	111.17	121.20
85	AA	2249	U	C5'-C4'-C3'	-7.16	104.54	116.00
35	BB	1435	G	C5-C6-O6	-7.16	124.30	128.60
85	AA	13	U	P-O3'-C3'	-7.16	111.11	119.70
85	AA	496	C	C6-N1-C1'	7.16	129.39	120.80
85	AA	1192	C	P-O5'-C5'	-7.16	109.44	120.90
34	BA	112	C	C2-N3-C4	-7.16	116.32	119.90
34	BA	512	U	C6-N1-C2	-7.16	116.70	121.00
34	BA	722	A	P-O3'-C3'	7.16	128.29	119.70
34	BA	1280	A	C6-N1-C2	-7.16	114.30	118.60
34	BA	1426	A	C5'-C4'-C3'	-7.16	104.54	116.00
35	BB	750	G	C4-N9-C1'	-7.16	117.19	126.50
35	BB	1291	G	C4-N9-C1'	-7.16	117.19	126.50
35	BB	1376	G	O5'-C5'-C4'	-7.16	98.09	111.70
35	BB	1514	G	C5'-C4'-C3'	-7.16	104.54	116.00
38	BE	109	C	O4'-C1'-N1	7.16	113.93	108.20
42	BI	13	ARG	NE-CZ-NH2	-7.16	116.72	120.30
53	BT	38	ARG	CA-CB-CG	7.16	129.15	113.40
65	Bf	170	TYR	N-CA-C	-7.16	91.67	111.00
85	AA	696	G	N3-C2-N2	-7.16	114.89	119.90
85	AA	1735	U	C3'-C2'-C1'	-7.16	95.77	101.50
85	AA	2154	C	O3'-P-O5'	-7.16	90.40	104.00
21	AM	143	ARG	NE-CZ-NH1	7.16	123.88	120.30
34	BA	62	A	N9-C1'-C2'	-7.16	104.13	112.00
34	BA	605	G	C4-C5-C6	7.16	123.09	118.80
34	BA	684	G	C5'-C4'-C3'	7.16	127.45	116.00
34	BA	867	C	N1-C2-O2	7.16	123.19	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1226	G	C4-N9-C1'	-7.16	117.19	126.50
35	BB	59	U	C4'-C3'-C2'	-7.16	95.44	102.60
35	BB	877	A	C3'-C2'-C1'	-7.16	95.77	101.50
35	BB	910	C	C6-N1-C2	-7.16	117.44	120.30
35	BB	1118	G	C1'-O4'-C4'	-7.16	104.17	109.90
35	BB	1261	U	C5'-C4'-C3'	7.16	127.45	116.00
35	BB	1483	A	N7-C8-N9	7.16	117.38	113.80
77	Br	333	GLN	N-CA-CB	-7.16	97.72	110.60
85	AA	1486	G	P-O3'-C3'	-7.16	111.11	119.70
85	AA	1599	G	P-O3'-C3'	-7.16	111.11	119.70
86	AB	9	A	P-O3'-C3'	-7.16	111.11	119.70
34	BA	82	A	C8-N9-C4	7.16	108.66	105.80
34	BA	290	G	O5'-P-OP2	-7.16	99.26	105.70
41	BH	66	G	C3'-C2'-C1'	-7.16	95.78	101.50
85	AA	1105	G	N1-C2-N2	-7.16	109.76	116.20
34	BA	568	G	C8-N9-C1'	7.16	136.30	127.00
34	BA	988	U	C2-N1-C1'	-7.16	109.11	117.70
35	BB	271	C	C6-N1-C2	-7.16	117.44	120.30
38	BE	54	U	O4'-C1'-N1	7.16	113.92	108.20
53	BT	63	TRP	N-CA-CB	7.16	123.48	110.60
73	Bn	72	ARG	CD-NE-CZ	-7.16	113.58	123.60
85	AA	655	U	C3'-C2'-C1'	-7.16	95.78	101.50
85	AA	1032	U	C1'-O4'-C4'	-7.16	104.17	109.90
85	AA	1328	U	O4'-C1'-N1	7.16	113.92	108.20
35	BB	899	C	C2-N1-C1'	-7.15	110.93	118.80
35	BB	1337	C	C2-N3-C4	-7.15	116.32	119.90
41	BH	13	C	P-O3'-C3'	-7.15	111.12	119.70
85	AA	1896	G	C8-N9-C4	-7.15	103.54	106.40
34	BA	26	C	C2-N3-C4	-7.15	116.32	119.90
34	BA	223	U	C6-N1-C1'	7.15	131.21	121.20
34	BA	486	G	C5'-C4'-C3'	7.15	127.45	116.00
34	BA	1593	U	N3-C2-O2	-7.15	117.19	122.20
35	BB	29	C	C3'-C2'-C1'	-7.15	95.78	101.50
35	BB	1227	G	O3'-P-O5'	7.15	117.59	104.00
40	BG	5	G	P-O3'-C3'	-7.15	111.12	119.70
40	BG	13	A	C3'-C2'-C1'	-7.15	95.78	101.50
85	AA	107	A	C5'-C4'-O4'	7.15	117.68	109.10
85	AA	405	C	P-O5'-C5'	7.15	132.34	120.90
85	AA	1249	U	O4'-C1'-N1	7.15	113.92	108.20
85	AA	2202	G	C6-N1-C2	-7.15	120.81	125.10
8	A7	46	ASN	N-CA-C	-7.15	91.69	111.00
25	AR	50	ILE	CA-C-N	-7.15	101.47	117.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	689	C	C5-C4-N4	-7.15	115.19	120.20
34	BA	1293	A	C5-C6-N1	7.15	121.28	117.70
34	BA	1454	G	N3-C4-C5	7.15	132.18	128.60
35	BB	88	U	C1'-O4'-C4'	-7.15	104.18	109.90
35	BB	1117	G	C3'-C2'-C1'	-7.15	95.78	101.50
37	BD	93	G	O4'-C1'-N9	7.15	113.92	108.20
40	BG	14	G	N9-C1'-C2'	-7.15	104.14	112.00
35	BB	1108	G	N3-C4-C5	-7.15	125.03	128.60
35	BB	1400	C	P-O5'-C5'	7.15	132.34	120.90
85	AA	588	G	C8-N9-C1'	7.15	136.29	127.00
85	AA	929	G	C2'-C3'-O3'	7.15	125.23	109.50
34	BA	34	U	N3-C2-O2	-7.15	117.20	122.20
34	BA	1026	C	N3-C2-O2	-7.15	116.90	121.90
34	BA	1837	U	N3-C2-O2	-7.15	117.20	122.20
35	BB	288	C	O4'-C1'-N1	7.15	113.92	108.20
35	BB	541	U	C5-C4-O4	-7.15	121.61	125.90
35	BB	1508	G	O4'-C1'-N9	7.15	113.92	108.20
37	BD	31	U	C5'-C4'-C3'	-7.15	104.56	116.00
38	BE	153	C	C6-N1-C1'	7.15	129.38	120.80
51	BR	56	ARG	NE-CZ-NH2	-7.15	116.73	120.30
65	Bf	432	LYS	N-CA-CB	-7.15	97.73	110.60
31	AX	152	ARG	NE-CZ-NH2	-7.15	116.73	120.30
34	BA	821	G	N1-C6-O6	7.15	124.19	119.90
35	BB	103	C	O4'-C1'-N1	7.15	113.92	108.20
85	AA	730	G	N3-C2-N2	-7.15	114.90	119.90
85	AA	1954	C	P-O3'-C3'	-7.15	111.12	119.70
34	BA	1550	G	C4-N9-C1'	-7.14	117.21	126.50
34	BA	1618	A	P-O3'-C3'	7.14	128.27	119.70
57	BX	91	THR	CA-CB-CG2	-7.14	102.40	112.40
67	Bh	64	ARG	N-CA-CB	7.14	123.46	110.60
85	AA	466	A	N3-C4-C5	7.14	131.80	126.80
85	AA	2213	A	C5'-C4'-C3'	7.14	127.43	116.00
34	BA	1231	C	C6-N1-C2	-7.14	117.44	120.30
34	BA	1277	G	C1'-O4'-C4'	-7.14	104.19	109.90
34	BA	1332	U	O4'-C1'-N1	7.14	113.91	108.20
35	BB	524	C	C1'-O4'-C4'	-7.14	104.19	109.90
35	BB	577	U	C5'-C4'-C3'	-7.14	104.57	116.00
85	AA	365	G	P-O3'-C3'	-7.14	111.13	119.70
86	AB	40	C	O4'-C1'-N1	7.14	113.92	108.20
34	BA	315	U	O5'-P-OP2	7.14	119.27	110.70
34	BA	1502	G	O4'-C1'-N9	7.14	113.91	108.20
35	BB	519	A	N1-C6-N6	7.14	122.89	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A2	174	TYR	CB-CG-CD2	-7.14	116.72	121.00
34	BA	622	G	C5'-C4'-C3'	-7.14	104.58	116.00
34	BA	868	C	N3-C4-C5	7.14	124.76	121.90
34	BA	1501	U	P-O3'-C3'	-7.14	111.13	119.70
34	BA	1598	U	C4'-C3'-C2'	-7.14	95.46	102.60
34	BA	1635	A	C1'-O4'-C4'	-7.14	104.19	109.90
34	BA	1654	G	O4'-C1'-C2'	7.14	114.03	107.60
35	BB	879	G	N1-C2-N2	-7.14	109.77	116.20
38	BE	58	U	C2-N3-C4	-7.14	122.72	127.00
72	Bm	30	ARG	NE-CZ-NH1	7.14	123.87	120.30
80	Bu	263	ARG	NE-CZ-NH1	7.14	123.87	120.30
85	AA	100	A	O4'-C4'-C3'	7.14	111.81	106.10
85	AA	487	G	O5'-C5'-C4'	-7.14	98.13	111.70
85	AA	491	G	O4'-C1'-N9	7.14	113.91	108.20
85	AA	2199	G	C5-C6-O6	-7.14	124.32	128.60
34	BA	295	G	C8-N9-C1'	-7.14	117.72	127.00
35	BB	513	G	O5'-C5'-C4'	-7.14	98.14	111.70
35	BB	608	A	P-O3'-C3'	-7.14	111.14	119.70
85	AA	158	C	C2-N1-C1'	-7.14	110.95	118.80
6	A5	160	ARG	CB-CA-C	7.14	124.67	110.40
34	BA	1166	A	C3'-C2'-C1'	-7.14	95.79	101.50
35	BB	544	C	C6-N1-C1'	7.14	129.37	120.80
35	BB	983	C	C3'-C2'-C1'	-7.14	95.79	101.50
36	BC	8	C	O4'-C1'-N1	7.14	113.91	108.20
36	BC	131	C	C5'-C4'-O4'	7.14	117.67	109.10
37	BD	64	A	C5-C6-N6	-7.14	117.99	123.70
38	BE	29	C	O4'-C1'-N1	7.14	113.91	108.20
59	BZ	71	TYR	CB-CG-CD1	7.14	125.28	121.00
85	AA	234	G	C5-C6-O6	-7.14	124.32	128.60
85	AA	1278	C	P-O3'-C3'	-7.14	111.14	119.70
35	BB	1473	U	C3'-C2'-C1'	-7.13	95.79	101.50
37	BD	108	G	C5'-C4'-O4'	7.13	117.66	109.10
34	BA	554	A	O4'-C1'-N9	7.13	113.91	108.20
34	BA	1429	A	C5'-C4'-C3'	-7.13	104.59	116.00
35	BB	505	G	N1-C2-N2	-7.13	109.78	116.20
63	Bd	29	MET	N-CA-C	-7.13	91.74	111.00
85	AA	510	A	O3'-P-O5'	-7.13	90.45	104.00
85	AA	1714	G	P-O5'-C5'	-7.13	109.49	120.90
85	AA	1928	A	C8-N9-C4	7.13	108.65	105.80
34	BA	130	U	P-O5'-C5'	7.13	132.31	120.90
34	BA	228	A	C4'-C3'-C2'	-7.13	95.47	102.60
34	BA	413	A	C3'-C2'-C1'	-7.13	95.80	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	7	C	N3-C4-N4	-7.13	113.01	118.00
35	BB	822	G	C5'-C4'-C3'	7.13	127.41	116.00
35	BB	1244	U	N1-C1'-C2'	-7.13	104.16	112.00
35	BB	1453	G	N3-C4-N9	-7.13	121.72	126.00
36	BC	70	C	O4'-C1'-C2'	7.13	114.02	107.60
85	AA	548	G	C8-N9-C1'	7.13	136.27	127.00
34	BA	381	A	C4'-C3'-C2'	7.13	109.73	102.60
34	BA	611	A	C2-N3-C4	-7.13	107.03	110.60
35	BB	8	U	C2-N3-C4	-7.13	122.72	127.00
40	BG	181	C	O4'-C4'-C3'	-7.13	96.87	104.00
85	AA	244	G	P-O3'-C3'	-7.13	111.14	119.70
34	BA	76	U	C5-C6-N1	-7.13	119.14	122.70
34	BA	228	A	C5'-C4'-C3'	-7.13	104.59	116.00
34	BA	841	G	P-O3'-C3'	-7.13	111.15	119.70
34	BA	991	U	C4'-C3'-C2'	7.13	109.73	102.60
34	BA	1254	C	P-O3'-C3'	-7.13	111.14	119.70
34	BA	1535	G	C6-N1-C2	-7.13	120.82	125.10
35	BB	844	G	C3'-C2'-C1'	-7.13	95.80	101.50
36	BC	121	G	C1'-O4'-C4'	-7.13	104.20	109.90
38	BE	135	A	OP1-P-OP2	-7.13	108.91	119.60
42	BI	101	ARG	NE-CZ-NH2	-7.13	116.73	120.30
85	AA	106	G	O4'-C1'-N9	7.13	113.90	108.20
85	AA	527	A	C4'-C3'-C2'	-7.13	95.47	102.60
85	AA	594	C	C3'-C2'-C1'	-7.13	95.80	101.50
85	AA	680	U	C4'-C3'-C2'	-7.13	95.47	102.60
85	AA	1126	G	N1-C6-O6	-7.13	115.62	119.90
34	BA	163	G	N1-C6-O6	7.13	124.18	119.90
34	BA	186	G	C5'-C4'-C3'	-7.13	104.60	116.00
34	BA	1355	G	N1-C6-O6	7.13	124.18	119.90
34	BA	1747	C	C6-N1-C2	-7.13	117.45	120.30
35	BB	384	A	C8-N9-C1'	-7.13	114.87	127.70
35	BB	773	G	C8-N9-C1'	7.13	136.26	127.00
36	BC	37	U	C3'-C2'-C1'	-7.13	95.80	101.50
36	BC	169	G	C2'-C3'-O3'	7.13	125.18	109.50
38	BE	18	U	C2'-C3'-O3'	7.13	125.18	109.50
40	BG	46	G	C8-N9-C1'	7.13	136.26	127.00
85	AA	692	U	C2-N1-C1'	-7.13	109.15	117.70
85	AA	1012	C	O4'-C1'-N1	7.13	113.90	108.20
85	AA	1176	C	C6-N1-C2	-7.13	117.45	120.30
85	AA	1420	U	O4'-C1'-N1	7.13	113.90	108.20
85	AA	1448	A	C5-C6-N1	7.13	121.26	117.70
85	AA	1495	G	C3'-C2'-C1'	-7.13	95.80	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	181	G	C5-C6-O6	-7.12	124.33	128.60
35	BB	736	G	C6-N1-C2	-7.12	120.83	125.10
37	BD	118	C	C2-N1-C1'	-7.12	110.96	118.80
85	AA	819	G	C8-N9-C4	-7.12	103.55	106.40
85	AA	1089	G	P-O3'-C3'	-7.12	111.15	119.70
85	AA	1472	G	C4-N9-C1'	-7.12	117.24	126.50
85	AA	1978	G	P-O3'-C3'	7.12	128.25	119.70
85	AA	2103	C	OP1-P-OP2	-7.12	108.91	119.60
85	AA	2244	G	C4-N9-C1'	7.12	135.76	126.50
34	BA	803	U	C4'-C3'-C2'	-7.12	95.48	102.60
34	BA	1489	U	N1-C1'-C2'	-7.12	104.16	112.00
34	BA	395	G	N1-C6-O6	-7.12	115.63	119.90
34	BA	615	A	O4'-C1'-N9	7.12	113.90	108.20
35	BB	855	G	O4'-C1'-N9	7.12	113.90	108.20
38	BE	116	U	C5'-C4'-C3'	-7.12	104.60	116.00
53	BT	109	TYR	CB-CG-CD2	-7.12	116.73	121.00
66	Bg	79	HIS	CA-CB-CG	-7.12	101.49	113.60
85	AA	372	U	N1-C1'-C2'	-7.12	104.17	112.00
85	AA	793	C	O4'-C1'-N1	7.12	113.90	108.20
85	AA	2199	G	N1-C2-N2	-7.12	109.79	116.20
1	A0	26	ARG	NE-CZ-NH2	-7.12	116.74	120.30
34	BA	1121	U	O4'-C1'-N1	7.12	113.90	108.20
34	BA	1144	A	N1-C6-N6	-7.12	114.33	118.60
34	BA	1337	A	C4-N9-C1'	-7.12	113.48	126.30
34	BA	1542	A	N1-C6-N6	7.12	122.87	118.60
34	BA	1751	C	C6-N1-C2	-7.12	117.45	120.30
35	BB	459	U	P-O3'-C3'	-7.12	111.16	119.70
85	AA	656	U	C2-N1-C1'	7.12	126.24	117.70
16	AH	77	ARG	NE-CZ-NH1	7.12	123.86	120.30
34	BA	1063	G	N1-C6-O6	-7.12	115.63	119.90
34	BA	1145	U	C2-N1-C1'	-7.12	109.16	117.70
34	BA	1432	C	P-O3'-C3'	-7.12	111.16	119.70
34	BA	1659	G	O5'-P-OP1	-7.12	99.29	105.70
34	BA	1698	C	N1-C1'-C2'	-7.12	104.17	112.00
34	BA	1782	C	P-O3'-C3'	-7.12	111.16	119.70
40	BG	112	C	C4'-C3'-C2'	7.12	109.72	102.60
85	AA	173	A	C8-N9-C4	-7.12	102.95	105.80
85	AA	549	A	C8-N9-C1'	7.12	140.51	127.70
85	AA	1715	C	C6-N1-C2	-7.12	117.45	120.30
11	AC	153	MET	CG-SD-CE	-7.12	88.81	100.20
34	BA	1337	A	C8-N9-C4	7.12	108.65	105.80
47	BN	14	ARG	NE-CZ-NH2	7.12	123.86	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	531	C	C6-N1-C2	-7.12	117.45	120.30
34	BA	668	G	C8-N9-C4	-7.12	103.55	106.40
34	BA	720	A	C5-C6-N6	-7.12	118.01	123.70
39	BF	27	G	P-O3'-C3'	7.12	128.24	119.70
40	BG	151	A	P-O3'-C3'	-7.12	111.16	119.70
41	BH	135	U	C6-N1-C2	-7.12	116.73	121.00
85	AA	39	A	C5'-C4'-O4'	7.12	117.64	109.10
85	AA	205	A	N1-C6-N6	-7.12	114.33	118.60
85	AA	667	A	C8-N9-C4	7.12	108.65	105.80
85	AA	1191	G	C8-N9-C1'	7.12	136.25	127.00
34	BA	465	A	C5'-C4'-O4'	7.11	117.64	109.10
34	BA	579	U	C2-N3-C4	-7.11	122.73	127.00
35	BB	370	A	C4'-C3'-C2'	7.11	109.71	102.60
35	BB	387	G	N7-C8-N9	7.11	116.66	113.10
35	BB	451	A	C8-N9-C4	7.11	108.65	105.80
35	BB	1330	A	P-O5'-C5'	-7.11	109.52	120.90
35	BB	1369	A	O4'-C1'-N9	7.11	113.89	108.20
36	BC	49	G	N1-C2-N2	-7.11	109.80	116.20
38	BE	116	U	C6-N1-C1'	7.11	131.16	121.20
41	BH	20	A	C4-N9-C1'	-7.11	113.50	126.30
77	Br	92	GLY	C-N-CA	7.11	139.48	121.70
85	AA	727	U	O4'-C1'-N1	7.11	113.89	108.20
85	AA	917	A	C5'-C4'-C3'	-7.11	104.62	116.00
85	AA	2121	G	O4'-C1'-N9	7.11	113.89	108.20
34	BA	1193	A	C6-N1-C2	-7.11	114.33	118.60
35	BB	653	G	O4'-C4'-C3'	-7.11	96.89	104.00
38	BE	196	C	O4'-C1'-C2'	7.11	114.00	107.60
68	Bi	99	HIS	C-N-CA	7.11	139.48	121.70
73	Bn	68	LYS	N-CA-CB	-7.11	97.80	110.60
85	AA	355	G	C5-C6-O6	-7.11	124.33	128.60
85	AA	2241	C	C5-C4-N4	7.11	125.18	120.20
34	BA	589	A	C1'-O4'-C4'	-7.11	104.21	109.90
34	BA	829	U	C5'-C4'-C3'	-7.11	104.62	116.00
34	BA	1075	U	N3-C2-O2	-7.11	117.22	122.20
34	BA	1267	A	C3'-C2'-C1'	7.11	107.19	101.50
34	BA	1311	G	N3-C2-N2	7.11	124.88	119.90
34	BA	1521	C	C3'-C2'-C1'	-7.11	95.81	101.50
34	BA	1594	G	C5-C6-O6	-7.11	124.33	128.60
34	BA	1676	A	C4'-C3'-C2'	7.11	109.71	102.60
35	BB	677	U	C5'-C4'-O4'	7.11	117.63	109.10
35	BB	999	G	C5-C6-N1	7.11	115.06	111.50
36	BC	160	C	C6-N1-C1'	7.11	129.33	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	128	U	C1'-O4'-C4'	-7.11	104.21	109.90
56	BW	44	TYR	CB-CG-CD1	7.11	125.27	121.00
85	AA	597	A	C4'-C3'-C2'	-7.11	95.49	102.60
34	BA	175	G	C8-N9-C1'	7.11	136.24	127.00
35	BB	152	G	C5-C6-O6	-7.11	124.33	128.60
36	BC	130	U	C6-N1-C1'	-7.11	111.25	121.20
34	BA	944	G	O4'-C1'-C2'	7.11	114.00	107.60
34	BA	1464	C	C2-N1-C1'	-7.11	110.98	118.80
35	BB	58	G	C8-N9-C4	7.11	109.24	106.40
35	BB	689	C	N3-C2-O2	-7.11	116.92	121.90
36	BC	123	G	N9-C4-C5	7.11	108.24	105.40
39	BF	34	C	N3-C2-O2	-7.11	116.92	121.90
40	BG	21	C	OP2-P-O3'	7.11	120.84	105.20
85	AA	2130	G	O4'-C1'-N9	7.11	113.89	108.20
34	BA	7	U	C3'-C2'-C1'	-7.11	95.82	101.50
34	BA	214	A	C4'-C3'-C2'	-7.11	95.49	102.60
34	BA	681	G	C8-N9-C4	7.11	109.24	106.40
34	BA	1632	G	C5'-C4'-O4'	7.11	117.63	109.10
34	BA	1718	C	P-O3'-C3'	-7.11	111.17	119.70
44	BK	7	ARG	NE-CZ-NH1	7.11	123.85	120.30
55	BV	90	PHE	CB-CG-CD2	-7.11	115.83	120.80
58	BY	64	ARG	N-CA-CB	7.11	123.39	110.60
85	AA	65	A	P-O3'-C3'	7.11	128.23	119.70
85	AA	117	C	C5'-C4'-C3'	-7.11	104.63	116.00
85	AA	153	C	C1'-O4'-C4'	-7.11	104.22	109.90
85	AA	492	C	N3-C4-N4	-7.11	113.03	118.00
85	AA	898	A	C4'-C3'-C2'	-7.11	95.49	102.60
85	AA	1553	G	N1-C6-O6	7.11	124.16	119.90
85	AA	1822	G	C3'-C2'-C1'	-7.11	95.82	101.50
85	AA	2164	G	N1-C6-O6	7.11	124.16	119.90
34	BA	50	G	N1-C6-O6	-7.10	115.64	119.90
34	BA	596	G	C4-C5-N7	7.10	113.64	110.80
34	BA	1205	A	C5-C6-N6	7.10	129.38	123.70
35	BB	555	G	C5'-C4'-O4'	7.10	117.62	109.10
35	BB	800	U	P-O3'-C3'	7.10	128.22	119.70
35	BB	1539	C	O4'-C1'-N1	7.10	113.88	108.20
40	BG	14	G	C5'-C4'-O4'	-7.10	100.58	109.10
85	AA	1633	A	N1-C6-N6	7.10	122.86	118.60
34	BA	259	C	C3'-C2'-C1'	-7.10	95.82	101.50
34	BA	1039	G	C4-N9-C1'	-7.10	117.27	126.50
34	BA	1223	C	C2-N3-C4	7.10	123.45	119.90
34	BA	1450	G	C4-N9-C1'	-7.10	117.27	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1676	A	C8-N9-C4	7.10	108.64	105.80
35	BB	362	A	O4'-C1'-N9	7.10	113.88	108.20
35	BB	978	C	O4'-C1'-N1	7.10	113.88	108.20
85	AA	913	U	P-O3'-C3'	-7.10	111.18	119.70
85	AA	1116	G	N3-C2-N2	7.10	124.87	119.90
85	AA	545	A	O4'-C1'-N9	7.10	113.88	108.20
85	AA	1254	A	C8-N9-C4	7.10	108.64	105.80
34	BA	47	U	C2-N1-C1'	-7.10	109.18	117.70
34	BA	485	C	P-O3'-C3'	7.10	128.22	119.70
34	BA	603	U	C5-C4-O4	-7.10	121.64	125.90
34	BA	657	C	O5'-C5'-C4'	7.10	125.19	111.70
35	BB	566	A	C6-N1-C2	-7.10	114.34	118.60
35	BB	1203	C	C4'-C3'-C2'	-7.10	95.50	102.60
85	AA	1922	A	O4'-C1'-N9	-7.10	102.52	108.20
85	AA	2176	U	P-O5'-C5'	7.10	132.26	120.90
34	BA	3	G	P-O5'-C5'	-7.10	109.54	120.90
34	BA	94	G	C5-C6-O6	7.10	132.86	128.60
34	BA	163	G	C5-C6-O6	-7.10	124.34	128.60
34	BA	400	A	C4'-C3'-C2'	7.10	109.70	102.60
34	BA	1114	G	N3-C2-N2	7.10	124.87	119.90
34	BA	1535	G	N9-C1'-C2'	-7.10	104.19	112.00
34	BA	1844	U	P-O3'-C3'	-7.10	111.18	119.70
35	BB	3	C	C5-C6-N1	7.10	124.55	121.00
35	BB	810	G	N1-C6-O6	-7.10	115.64	119.90
35	BB	842	G	C5'-C4'-C3'	7.10	127.35	116.00
36	BC	147	G	C1'-O4'-C4'	-7.10	104.22	109.90
40	BG	117	C	C1'-O4'-C4'	-7.10	104.22	109.90
85	AA	98	U	O4'-C1'-N1	7.10	113.88	108.20
85	AA	1847	U	P-O3'-C3'	7.10	128.22	119.70
34	BA	326	A	C5'-C4'-O4'	7.10	117.61	109.10
35	BB	463	C	C1'-O4'-C4'	-7.10	104.22	109.90
35	BB	877	A	C2-N3-C4	7.10	114.15	110.60
35	BB	1187	G	N1-C2-N3	7.10	128.16	123.90
35	BB	1454	G	P-O5'-C5'	7.10	132.25	120.90
36	BC	50	C	P-O3'-C3'	-7.10	111.19	119.70
34	BA	884	G	C5-C6-O6	-7.09	124.34	128.60
34	BA	1630	A	O3'-P-O5'	7.09	117.48	104.00
35	BB	677	U	C5'-C4'-C3'	-7.09	104.65	116.00
35	BB	798	A	P-O3'-C3'	7.09	128.21	119.70
37	BD	29	C	O4'-C1'-N1	7.09	113.88	108.20
38	BE	113	C	C4'-C3'-C2'	-7.09	95.51	102.60
38	BE	119	U	C2-N1-C1'	-7.09	109.19	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	413	G	C3'-C2'-C1'	-7.09	95.82	101.50
85	AA	1727	U	O3'-P-O5'	7.09	117.48	104.00
85	AA	1891	U	C2-N1-C1'	-7.09	109.19	117.70
35	BB	952	U	O4'-C1'-N1	7.09	113.87	108.20
85	AA	1314	C	P-O3'-C3'	7.09	128.21	119.70
85	AA	1728	G	C6-C5-N7	-7.09	126.14	130.40
34	BA	314	A	O4'-C1'-N9	7.09	113.87	108.20
34	BA	644	C	N3-C4-N4	7.09	122.97	118.00
35	BB	366	G	C5'-C4'-C3'	-7.09	104.66	116.00
35	BB	784	C	O5'-C5'-C4'	-7.09	98.23	111.70
37	BD	107	G	C4-N9-C1'	-7.09	117.28	126.50
85	AA	362	G	C5-C6-O6	-7.09	124.34	128.60
85	AA	2056	C	O3'-P-O5'	-7.09	90.53	104.00
34	BA	37	A	O4'-C1'-N9	7.09	113.87	108.20
35	BB	1040	C	O4'-C1'-N1	7.09	113.87	108.20
35	BB	1452	U	C4'-C3'-C2'	7.09	109.69	102.60
37	BD	98	G	O3'-P-O5'	-7.09	90.53	104.00
47	BN	40	ARG	NE-CZ-NH1	7.09	123.84	120.30
85	AA	75	U	C6-N1-C2	-7.09	116.75	121.00
85	AA	1097	G	C8-N9-C1'	7.09	136.22	127.00
85	AA	1373	U	C3'-C2'-C1'	-7.09	95.83	101.50
6	A5	164	TYR	CB-CG-CD2	-7.09	116.75	121.00
34	BA	556	A	C5-C6-N6	-7.09	118.03	123.70
52	BS	156	ARG	NE-CZ-NH1	7.09	123.84	120.30
85	AA	2078	A	P-O5'-C5'	7.09	132.24	120.90
31	AX	189	ARG	NE-CZ-NH1	7.09	123.84	120.30
34	BA	800	G	P-O3'-C3'	-7.09	111.20	119.70
34	BA	810	A	N9-C1'-C2'	-7.09	104.20	112.00
35	BB	1170	U	P-O3'-C3'	7.09	128.20	119.70
38	BE	159	A	O4'-C1'-N9	7.09	113.87	108.20
73	Bn	24	ARG	NE-CZ-NH1	7.09	123.84	120.30
84	By	6	HIS	C-N-CA	7.09	139.42	121.70
85	AA	436	G	C4-N9-C1'	-7.09	117.29	126.50
85	AA	1210	U	N1-C2-O2	7.09	127.76	122.80
85	AA	1957	C	O4'-C1'-N1	7.09	113.87	108.20
86	AB	13	C	C2-N3-C4	7.09	123.44	119.90
35	BB	441	G	C3'-C2'-C1'	-7.08	95.83	101.50
35	BB	1248	A	P-O3'-C3'	-7.08	111.20	119.70
85	AA	1279	A	P-O5'-C5'	7.08	132.24	120.90
85	AA	1760	C	O4'-C1'-N1	7.08	113.87	108.20
34	BA	1696	G	C1'-O4'-C4'	-7.08	104.23	109.90
35	BB	1076	U	C2-N3-C4	-7.08	122.75	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1191	G	C5-C6-O6	-7.08	124.35	128.60
35	BB	1292	G	N1-C6-O6	-7.08	115.65	119.90
40	BG	102	G	C8-N9-C4	-7.08	103.57	106.40
85	AA	429	G	C5-C6-O6	-7.08	124.35	128.60
85	AA	431	G	C4-N9-C1'	-7.08	117.29	126.50
85	AA	767	A	N9-C1'-C2'	-7.08	104.21	112.00
85	AA	1046	C	O4'-C1'-N1	7.08	113.87	108.20
85	AA	1119	A	C3'-C2'-C1'	-7.08	95.83	101.50
85	AA	1840	C	C2-N1-C1'	-7.08	111.01	118.80
85	AA	1978	G	C4-C5-C6	-7.08	114.55	118.80
34	BA	73	G	O4'-C1'-N9	7.08	113.87	108.20
34	BA	1233	U	C4'-C3'-C2'	7.08	109.68	102.60
34	BA	1614	G	N3-C2-N2	7.08	124.86	119.90
35	BB	616	U	C5'-C4'-O4'	7.08	117.60	109.10
35	BB	1273	G	C5-C6-O6	-7.08	124.35	128.60
38	BE	2	G	C5-C6-O6	7.08	132.85	128.60
66	Bg	63	ARG	NE-CZ-NH1	7.08	123.84	120.30
85	AA	266	U	C2-N3-C4	-7.08	122.75	127.00
85	AA	467	U	N3-C2-O2	-7.08	117.24	122.20
85	AA	635	G	O4'-C1'-N9	7.08	113.87	108.20
85	AA	811	A	N1-C6-N6	-7.08	114.35	118.60
85	AA	1006	C	C3'-C2'-C1'	-7.08	95.83	101.50
35	BB	1445	A	C8-N9-C1'	7.08	140.44	127.70
34	BA	1545	C	N3-C2-O2	-7.08	116.94	121.90
40	BG	64	C	O5'-C5'-C4'	-7.08	98.25	111.70
51	BR	63	TYR	CB-CG-CD1	-7.08	116.75	121.00
85	AA	23	G	C4-N9-C1'	-7.08	117.30	126.50
34	BA	519	G	C4-N9-C1'	7.08	135.70	126.50
34	BA	959	G	O3'-P-O5'	-7.08	90.55	104.00
35	BB	508	U	O4'-C1'-N1	7.08	113.86	108.20
35	BB	699	U	C2-N3-C4	-7.08	122.75	127.00
38	BE	126	G	C4-N9-C1'	-7.08	117.30	126.50
40	BG	148	C	C4'-C3'-C2'	-7.08	95.52	102.60
80	Bu	158	ARG	NE-CZ-NH1	7.08	123.84	120.30
85	AA	82	A	O4'-C4'-C3'	-7.08	96.92	104.00
85	AA	289	G	C1'-O4'-C4'	-7.08	104.24	109.90
85	AA	2008	G	N3-C4-N9	-7.08	121.75	126.00
1	A0	102	PHE	N-CA-CB	-7.08	97.86	110.60
34	BA	109	A	C5-C6-N1	7.08	121.24	117.70
34	BA	1358	A	O4'-C1'-N9	7.08	113.86	108.20
34	BA	1532	G	N3-C2-N2	7.08	124.85	119.90
35	BB	621	C	C6-N1-C2	-7.08	117.47	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	624	A	N9-C1'-C2'	-7.08	104.22	112.00
35	BB	1464	G	C4-N9-C1'	-7.08	117.30	126.50
36	BC	168	C	N3-C4-C5	-7.08	119.07	121.90
38	BE	21	C	N1-C2-O2	7.08	123.15	118.90
39	BF	31	U	C1'-O4'-C4'	-7.08	104.24	109.90
85	AA	863	C	C4'-C3'-C2'	-7.08	95.53	102.60
85	AA	2008	G	C8-N9-C1'	7.08	136.20	127.00
34	BA	332	U	P-O3'-C3'	-7.07	111.21	119.70
34	BA	831	U	O4'-C1'-N1	7.07	113.86	108.20
34	BA	875	G	C1'-O4'-C4'	-7.07	104.24	109.90
34	BA	1390	C	C5'-C4'-C3'	-7.07	104.68	116.00
40	BG	162	A	C5-C6-N6	-7.07	118.04	123.70
41	BH	31	A	C5'-C4'-O4'	-7.07	100.61	109.10
68	Bi	63	ARG	NE-CZ-NH1	7.07	123.84	120.30
72	Bm	95	ARG	NE-CZ-NH1	7.07	123.84	120.30
85	AA	54	C	C6-N1-C2	-7.07	117.47	120.30
85	AA	122	A	P-O3'-C3'	-7.07	111.21	119.70
85	AA	927	A	P-O5'-C5'	-7.07	109.58	120.90
85	AA	1148	G	N9-C1'-C2'	-7.07	104.22	112.00
34	BA	1586	U	C1'-O4'-C4'	-7.07	104.24	109.90
39	BF	10	A	C8-N9-C4	-7.07	102.97	105.80
85	AA	1664	G	C5-C6-O6	-7.07	124.36	128.60
35	BB	1061	G	C5-C6-O6	7.07	132.84	128.60
35	BB	1319	U	O4'-C1'-N1	7.07	113.86	108.20
53	BT	19	ARG	NE-CZ-NH2	-7.07	116.77	120.30
85	AA	340	G	P-O3'-C3'	7.07	128.18	119.70
85	AA	759	G	P-O3'-C3'	7.07	128.19	119.70
13	AE	26	ARG	NE-CZ-NH1	7.07	123.83	120.30
34	BA	553	A	C6-N1-C2	-7.07	114.36	118.60
35	BB	732	G	C5'-C4'-O4'	7.07	117.58	109.10
85	AA	1492	U	P-O3'-C3'	-7.07	111.22	119.70
85	AA	1544	G	C4-N9-C1'	-7.07	117.31	126.50
34	BA	1119	A	C5-C6-N1	7.07	121.23	117.70
35	BB	475	A	C1'-O4'-C4'	-7.07	104.25	109.90
35	BB	767	A	C5'-C4'-C3'	-7.07	104.69	116.00
39	BF	23	G	P-O3'-C3'	-7.07	111.22	119.70
40	BG	47	G	C5-C6-O6	-7.07	124.36	128.60
68	Bi	62	ARG	NE-CZ-NH1	7.07	123.83	120.30
85	AA	1731	G	C2-N3-C4	7.07	115.43	111.90
6	A5	197	ARG	NE-CZ-NH2	-7.07	116.77	120.30
34	BA	1617	U	O3'-P-O5'	-7.07	90.58	104.00
35	BB	628	A	C5'-C4'-C3'	-7.07	104.69	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1230	A	C4-N9-C1'	-7.07	113.58	126.30
35	BB	1276	U	O4'-C1'-N1	7.07	113.85	108.20
40	BG	130	G	C5-C6-O6	-7.07	124.36	128.60
84	By	8	GLN	N-CA-CB	7.07	123.32	110.60
85	AA	246	C	C6-N1-C2	-7.07	117.47	120.30
85	AA	1721	A	C4'-C3'-C2'	-7.07	95.53	102.60
85	AA	2151	U	N3-C2-O2	-7.07	117.25	122.20
34	BA	168	U	C2-N3-C4	-7.06	122.76	127.00
34	BA	1608	C	O5'-C5'-C4'	-7.06	98.28	111.70
41	BH	93	G	C2-N3-C4	-7.06	108.37	111.90
85	AA	1460	G	C5-C6-N1	7.06	115.03	111.50
34	BA	401	A	O5'-P-OP1	-7.06	99.34	105.70
34	BA	776	U	P-O3'-C3'	-7.06	111.22	119.70
35	BB	121	A	C5-C6-N1	7.06	121.23	117.70
35	BB	1510	G	C5-C6-N1	7.06	115.03	111.50
36	BC	59	A	O4'-C1'-N9	7.06	113.85	108.20
38	BE	89	G	N3-C2-N2	7.06	124.84	119.90
39	BF	14	C	C3'-C2'-C1'	-7.06	95.85	101.50
41	BH	3	U	P-O3'-C3'	7.06	128.18	119.70
85	AA	515	C	C6-N1-C2	-7.06	117.47	120.30
85	AA	808	A	C8-N9-C4	-7.06	102.97	105.80
85	AA	856	G	N9-C1'-C2'	-7.06	104.23	112.00
85	AA	865	G	N9-C1'-C2'	-7.06	104.23	112.00
85	AA	1055	U	C5'-C4'-O4'	7.06	117.58	109.10
2	A1	179	MET	CG-SD-CE	-7.06	88.90	100.20
35	BB	1194	A	N9-C1'-C2'	-7.06	104.23	112.00
85	AA	802	A	P-O3'-C3'	7.06	128.17	119.70
85	AA	2199	G	C4-N9-C1'	-7.06	117.32	126.50
34	BA	52	G	O5'-P-OP2	7.06	119.17	110.70
34	BA	1196	C	C2-N3-C4	7.06	123.43	119.90
35	BB	448	G	C5'-C4'-O4'	7.06	117.57	109.10
35	BB	762	C	C6-N1-C2	-7.06	117.48	120.30
35	BB	873	C	O4'-C1'-N1	7.06	113.85	108.20
85	AA	257	U	P-O3'-C3'	-7.06	111.23	119.70
85	AA	445	U	O4'-C1'-N1	7.06	113.85	108.20
85	AA	457	G	C3'-C2'-C1'	-7.06	95.85	101.50
85	AA	747	U	C3'-C2'-C1'	-7.06	95.85	101.50
85	AA	913	U	C5'-C4'-C3'	-7.06	104.70	116.00
85	AA	1465	C	C6-N1-C1'	7.06	129.27	120.80
34	BA	219	U	C5'-C4'-O4'	7.06	117.57	109.10
34	BA	295	G	O3'-P-O5'	7.06	117.41	104.00
34	BA	811	C	P-O5'-C5'	-7.06	109.61	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1333	G	C8-N9-C4	7.06	109.22	106.40
35	BB	549	U	C5-C4-O4	-7.06	121.67	125.90
85	AA	187	C	C5'-C4'-C3'	-7.06	104.71	116.00
85	AA	438	G	C5'-C4'-C3'	-7.06	104.71	116.00
85	AA	2127	G	C1'-O4'-C4'	-7.06	104.25	109.90
7	A6	46	MET	CG-SD-CE	-7.06	88.91	100.20
34	BA	882	G	C3'-C2'-C1'	-7.06	95.86	101.50
34	BA	1542	A	P-O3'-C3'	-7.06	111.23	119.70
35	BB	490	G	P-O3'-C3'	-7.06	111.23	119.70
85	AA	1002	G	C4-N9-C1'	-7.06	117.33	126.50
34	BA	543	A	C3'-C2'-C1'	-7.05	95.86	101.50
34	BA	748	C	O4'-C1'-C2'	7.05	113.95	107.60
34	BA	1819	U	P-O3'-C3'	7.05	128.17	119.70
35	BB	777	C	C6-N1-C2	-7.05	117.48	120.30
85	AA	1721	A	C8-N9-C1'	7.05	140.40	127.70
35	BB	1374	U	C4'-C3'-C2'	7.05	109.65	102.60
38	BE	1	U	N1-C2-O2	7.05	127.74	122.80
60	Ba	90	ARG	NE-CZ-NH2	-7.05	116.77	120.30
62	Bc	9	TRP	CB-CG-CD1	-7.05	117.83	127.00
85	AA	136	U	C5'-C4'-C3'	7.05	127.28	116.00
85	AA	708	G	C4-N9-C1'	-7.05	117.33	126.50
34	BA	25	C	P-O3'-C3'	-7.05	111.24	119.70
35	BB	733	G	C5-C6-O6	-7.05	124.37	128.60
35	BB	1469	A	C5-C6-N6	-7.05	118.06	123.70
59	BZ	40	TYR	CB-CG-CD2	-7.05	116.77	121.00
85	AA	475	A	O4'-C1'-C2'	7.05	113.95	107.60
85	AA	1456	A	C4'-C3'-C2'	-7.05	95.55	102.60
85	AA	1542	A	C6-N1-C2	-7.05	114.37	118.60
34	BA	1259	C	C3'-C2'-C1'	-7.05	95.86	101.50
35	BB	618	U	O4'-C1'-C2'	7.05	113.94	107.60
35	BB	1512	C	C2'-C3'-O3'	7.05	125.01	109.50
39	BF	12	U	C2-N1-C1'	7.05	126.16	117.70
40	BG	58	G	O5'-C5'-C4'	-7.05	98.31	111.70
65	Bf	356	TYR	CB-CG-CD1	-7.05	116.77	121.00
85	AA	69	C	O4'-C1'-N1	7.05	113.84	108.20
85	AA	472	A	N1-C6-N6	-7.05	114.37	118.60
85	AA	880	A	O4'-C1'-N9	7.05	113.84	108.20
85	AA	937	G	C4-N9-C1'	-7.05	117.33	126.50
85	AA	1955	U	C2-N3-C4	-7.05	122.77	127.00
86	AB	63	G	O4'-C1'-N9	7.05	113.84	108.20
34	BA	325	A	O3'-P-O5'	7.05	117.39	104.00
34	BA	1440	C	C5'-C4'-O4'	7.05	117.56	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	395	U	O4'-C1'-N1	7.05	113.84	108.20
38	BE	110	U	C1'-O4'-C4'	-7.05	104.26	109.90
42	BI	147	ARG	NE-CZ-NH1	7.05	123.82	120.30
72	Bm	44	LYS	N-CA-CB	-7.05	97.92	110.60
85	AA	699	U	C1'-O4'-C4'	-7.05	104.26	109.90
85	AA	1005	C	O4'-C1'-N1	7.05	113.84	108.20
85	AA	1018	G	O4'-C1'-N9	7.05	113.84	108.20
34	BA	777	C	C5'-C4'-C3'	-7.04	104.73	116.00
34	BA	1549	U	C2-N3-C4	-7.04	122.77	127.00
35	BB	1001	G	C8-N9-C4	-7.04	103.58	106.40
39	BF	6	C	P-O5'-C5'	7.04	132.17	120.90
64	Be	154	GLN	C-N-CA	7.04	139.31	121.70
85	AA	1921	G	C6-N1-C2	-7.04	120.87	125.10
27	AT	118	ARG	CD-NE-CZ	-7.04	113.74	123.60
34	BA	90	G	N1-C6-O6	7.04	124.13	119.90
34	BA	852	C	C5-C4-N4	7.04	125.13	120.20
35	BB	1094	A	C1'-O4'-C4'	-7.04	104.27	109.90
36	BC	142	C	O4'-C1'-N1	7.04	113.83	108.20
85	AA	841	U	C2-N3-C4	-7.04	122.77	127.00
85	AA	1103	A	N1-C6-N6	7.04	122.83	118.60
34	BA	1294	C	C3'-C2'-C1'	-7.04	95.87	101.50
34	BA	1736	A	N1-C6-N6	7.04	122.83	118.60
35	BB	118	A	N9-C1'-C2'	-7.04	104.25	112.00
35	BB	654	C	C2-N3-C4	-7.04	116.38	119.90
38	BE	179	A	N1-C6-N6	-7.04	114.38	118.60
72	Bm	64	ARG	NE-CZ-NH1	7.04	123.82	120.30
77	Br	359	ARG	NE-CZ-NH2	-7.04	116.78	120.30
85	AA	196	U	C2-N1-C1'	-7.04	109.25	117.70
85	AA	336	C	C5-C4-N4	-7.04	115.27	120.20
85	AA	836	A	C6-N1-C2	-7.04	114.38	118.60
85	AA	879	G	C5'-C4'-C3'	-7.04	104.73	116.00
85	AA	1222	A	C5-C6-N1	7.04	121.22	117.70
34	BA	1120	U	O4'-C4'-C3'	-7.04	96.96	104.00
34	BA	1721	U	P-O3'-C3'	-7.04	111.25	119.70
35	BB	701	U	C5'-C4'-C3'	-7.04	104.74	116.00
36	BC	62	A	C4-C5-C6	-7.04	113.48	117.00
40	BG	69	G	C5-C6-O6	-7.04	124.38	128.60
85	AA	2017	U	O4'-C1'-N1	7.04	113.83	108.20
34	BA	108	A	P-O5'-C5'	-7.04	109.64	120.90
34	BA	1725	U	O4'-C1'-N1	7.04	113.83	108.20
34	BA	1791	C	C6-N1-C2	-7.04	117.48	120.30
35	BB	639	A	O4'-C1'-C2'	-7.04	98.76	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	715	G	C4'-C3'-C2'	-7.04	95.56	102.60
35	BB	822	G	C6-N1-C2	-7.04	120.88	125.10
35	BB	1173	C	P-O5'-C5'	-7.04	109.64	120.90
37	BD	106	G	N1-C6-O6	-7.04	115.68	119.90
41	BH	102	C	N3-C2-O2	-7.04	116.97	121.90
72	Bm	37	ARG	NE-CZ-NH1	7.04	123.82	120.30
77	Br	324	ARG	NE-CZ-NH1	7.04	123.82	120.30
85	AA	414	C	N3-C4-N4	-7.04	113.07	118.00
85	AA	764	U	C2-N1-C1'	-7.04	109.25	117.70
34	BA	874	G	P-O3'-C3'	-7.04	111.25	119.70
34	BA	1328	U	C1'-O4'-C4'	-7.04	104.27	109.90
35	BB	895	U	C2-N3-C4	-7.04	122.78	127.00
35	BB	1230	A	O3'-P-O5'	7.04	117.37	104.00
85	AA	606	A	O4'-C1'-N9	7.04	113.83	108.20
85	AA	930	G	C6-N1-C2	-7.04	120.88	125.10
85	AA	1094	G	C8-N9-C1'	7.04	136.15	127.00
85	AA	1578	G	O5'-C5'-C4'	7.04	125.07	111.70
85	AA	1966	C	C4'-C3'-C2'	-7.04	95.56	102.60
34	BA	930	A	O4'-C1'-N9	7.04	113.83	108.20
35	BB	848	A	C4'-C3'-C2'	-7.04	95.56	102.60
35	BB	1515	C	O4'-C1'-N1	7.04	113.83	108.20
85	AA	424	A	O4'-C4'-C3'	-7.04	96.96	104.00
85	AA	963	U	C5'-C4'-O4'	7.04	117.54	109.10
34	BA	131	A	C5-C6-N6	7.03	129.33	123.70
34	BA	375	C	C6-N1-C2	-7.03	117.49	120.30
34	BA	537	C	C6-N1-C2	-7.03	117.49	120.30
34	BA	852	C	P-O3'-C3'	7.03	128.14	119.70
34	BA	1666	U	O4'-C1'-N1	7.03	113.83	108.20
38	BE	114	G	P-O3'-C3'	-7.03	111.26	119.70
41	BH	116	A	O4'-C1'-N9	7.03	113.83	108.20
42	BI	112	PHE	CB-CG-CD2	-7.03	115.88	120.80
56	BW	19	LEU	N-CA-C	-7.03	92.01	111.00
75	Bp	40	ARG	NE-CZ-NH1	7.03	123.82	120.30
85	AA	991	G	N9-C4-C5	-7.03	102.59	105.40
85	AA	1778	C	O4'-C1'-N1	7.03	113.83	108.20
34	BA	20	A	N9-C1'-C2'	-7.03	104.27	112.00
34	BA	1281	U	P-O3'-C3'	-7.03	111.26	119.70
34	BA	1428	G	N3-C2-N2	7.03	124.82	119.90
34	BA	1500	G	N3-C4-N9	7.03	130.22	126.00
85	AA	650	G	C3'-C2'-C1'	-7.03	95.88	101.50
85	AA	790	A	C5-N7-C8	-7.03	100.38	103.90
34	BA	5	C	P-O5'-C5'	7.03	132.15	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	813	C	P-O5'-C5'	7.03	132.15	120.90
34	BA	1037	C	C3'-C2'-C1'	-7.03	95.88	101.50
34	BA	1258	G	N1-C6-O6	7.03	124.12	119.90
35	BB	839	G	C4-N9-C1'	-7.03	117.36	126.50
36	BC	60	U	C1'-O4'-C4'	-7.03	104.28	109.90
37	BD	75	G	C6-N1-C2	-7.03	120.88	125.10
38	BE	165	U	N1-C2-N3	7.03	119.12	114.90
40	BG	35	G	C4-N9-C1'	-7.03	117.36	126.50
80	Bu	56	THR	N-CA-CB	7.03	123.66	110.30
85	AA	79	G	O4'-C1'-N9	7.03	113.83	108.20
85	AA	714	U	C2-N3-C4	-7.03	122.78	127.00
85	AA	1595	G	C2'-C3'-O3'	7.03	124.97	109.50
2	A1	48	ARG	NE-CZ-NH1	7.03	123.81	120.30
34	BA	433	G	P-O3'-C3'	7.03	128.13	119.70
36	BC	166	G	P-O5'-C5'	-7.03	109.65	120.90
85	AA	365	G	C8-N9-C1'	7.03	136.14	127.00
85	AA	940	G	O5'-C5'-C4'	7.03	125.06	111.70
34	BA	42	A	C1'-O4'-C4'	-7.03	104.28	109.90
34	BA	190	U	C1'-O4'-C4'	-7.03	104.28	109.90
34	BA	1485	U	O5'-C5'-C4'	-7.03	98.35	111.70
34	BA	1770	U	O4'-C1'-N1	7.03	113.82	108.20
35	BB	498	G	N1-C6-O6	7.03	124.12	119.90
38	BE	146	U	C5-C4-O4	7.03	130.12	125.90
80	Bu	13	TYR	CB-CG-CD1	-7.03	116.78	121.00
85	AA	155	U	C5'-C4'-C3'	-7.03	104.75	116.00
85	AA	179	G	C8-N9-C4	-7.03	103.59	106.40
85	AA	431	G	C8-N9-C1'	7.03	136.13	127.00
85	AA	540	A	P-O3'-C3'	7.03	128.13	119.70
85	AA	659	A	C5-C6-N6	7.03	129.32	123.70
85	AA	897	A	C3'-C2'-C1'	-7.03	95.88	101.50
85	AA	1866	A	P-O5'-C5'	-7.03	109.66	120.90
85	AA	2085	C	P-O3'-C3'	7.03	128.13	119.70
34	BA	521	C	C1'-O4'-C4'	-7.03	104.28	109.90
34	BA	1783	C	C3'-C2'-C1'	-7.03	95.88	101.50
35	BB	1427	A	C6-N1-C2	-7.03	114.38	118.60
38	BE	174	U	N3-C2-O2	-7.03	117.28	122.20
39	BF	32	G	O4'-C1'-C2'	7.03	113.92	107.60
40	BG	22	G	C5-C6-O6	-7.03	124.38	128.60
40	BG	51	U	O4'-C1'-N1	7.03	113.82	108.20
47	BN	94	TYR	CB-CG-CD2	-7.03	116.78	121.00
85	AA	183	C	P-O3'-C3'	7.03	128.13	119.70
85	AA	733	C	O4'-C1'-N1	7.03	113.82	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1520	A	O3'-P-O5'	-7.03	90.65	104.00
86	AB	19	G	C6-N1-C2	-7.03	120.89	125.10
8	A7	203	THR	N-CA-CB	7.02	123.65	110.30
34	BA	953	G	C4'-C3'-C2'	-7.02	95.58	102.60
34	BA	1542	A	C5'-C4'-C3'	7.02	127.24	116.00
35	BB	1322	A	P-O5'-C5'	-7.02	109.66	120.90
35	BB	1527	A	P-O5'-C5'	7.02	132.14	120.90
38	BE	205	G	C4-N9-C1'	-7.02	117.37	126.50
85	AA	587	G	P-O5'-C5'	-7.02	109.66	120.90
34	BA	251	U	O4'-C1'-N1	7.02	113.82	108.20
34	BA	665	C	O4'-C1'-N1	7.02	113.82	108.20
34	BA	1704	G	C1'-O4'-C4'	-7.02	104.28	109.90
37	BD	3	G	P-O5'-C5'	-7.02	109.66	120.90
38	BE	106	C	P-O5'-C5'	-7.02	109.66	120.90
85	AA	711	C	P-O5'-C5'	-7.02	109.67	120.90
34	BA	1461	A	C5'-C4'-C3'	-7.02	104.77	116.00
34	BA	1591	G	C5-C6-N1	7.02	115.01	111.50
36	BC	45	C	P-O3'-C3'	-7.02	111.28	119.70
53	BT	52	ARG	NE-CZ-NH1	7.02	123.81	120.30
85	AA	730	G	C5-C6-N1	7.02	115.01	111.50
85	AA	1099	U	O4'-C1'-N1	7.02	113.82	108.20
85	AA	1683	U	P-O3'-C3'	7.02	128.12	119.70
34	BA	357	A	O3'-P-O5'	7.02	117.34	104.00
34	BA	574	U	C5'-C4'-C3'	7.02	127.23	116.00
35	BB	433	C	C5'-C4'-C3'	7.02	127.23	116.00
38	BE	55	C	C6-N1-C1'	-7.02	112.38	120.80
38	BE	131	C	P-O5'-C5'	7.02	132.13	120.90
52	BS	3	ARG	NE-CZ-NH2	7.02	123.81	120.30
85	AA	775	C	P-O3'-C3'	7.02	128.12	119.70
85	AA	1031	G	P-O5'-C5'	7.02	132.13	120.90
85	AA	1622	G	N3-C2-N2	7.02	124.81	119.90
85	AA	1982	C	P-O5'-C5'	-7.02	109.67	120.90
34	BA	294	C	O4'-C1'-N1	7.02	113.81	108.20
34	BA	438	A	N1-C6-N6	-7.02	114.39	118.60
34	BA	579	U	OP1-P-OP2	-7.02	109.08	119.60
34	BA	799	A	C5-N7-C8	-7.02	100.39	103.90
34	BA	1090	A	O4'-C1'-N9	7.02	113.81	108.20
35	BB	1492	C	N3-C2-O2	-7.02	116.99	121.90
35	BB	1525	G	O4'-C1'-N9	7.02	113.81	108.20
37	BD	104	C	N1-C1'-C2'	-7.02	104.28	112.00
41	BH	75	G	C5-N7-C8	-7.02	100.79	104.30
43	BJ	177	ARG	NE-CZ-NH2	-7.02	116.79	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
67	Bh	104	ARG	NE-CZ-NH1	7.02	123.81	120.30
85	AA	744	C	N1-C2-N3	-7.02	114.29	119.20
85	AA	788	G	C8-N9-C1'	7.02	136.12	127.00
85	AA	1891	U	C6-N1-C1'	7.02	131.03	121.20
85	AA	1991	C	N1-C1'-C2'	7.02	123.12	114.00
34	BA	1151	A	O4'-C1'-N9	7.02	113.81	108.20
35	BB	1445	A	C4-N9-C1'	-7.02	113.67	126.30
85	AA	412	G	C3'-C2'-C1'	-7.02	95.89	101.50
85	AA	1506	U	C2'-C3'-O3'	7.02	124.94	109.50
85	AA	2036	A	C4-N9-C1'	7.02	138.93	126.30
19	AK	115	TYR	CA-CB-CG	-7.01	100.07	113.40
34	BA	260	A	O4'-C1'-C2'	7.01	113.91	107.60
34	BA	699	G	C5'-C4'-C3'	-7.01	104.78	116.00
34	BA	1087	A	C5-C6-N1	7.01	121.21	117.70
34	BA	1152	A	N9-C4-C5	-7.01	102.99	105.80
34	BA	1193	A	O4'-C1'-N9	7.01	113.81	108.20
35	BB	610	U	C6-N1-C1'	7.01	131.02	121.20
35	BB	1512	C	C3'-C2'-C1'	-7.01	95.89	101.50
37	BD	73	U	O4'-C1'-N1	-7.01	102.59	108.20
37	BD	86	A	N1-C6-N6	-7.01	114.39	118.60
64	Be	221	HIS	CA-CB-CG	-7.01	101.68	113.60
85	AA	674	U	N3-C4-O4	-7.01	114.49	119.40
85	AA	1197	U	C5'-C4'-O4'	7.01	117.52	109.10
85	AA	1551	G	P-O3'-C3'	7.01	128.12	119.70
85	AA	1649	U	C2-N1-C1'	-7.01	109.28	117.70
85	AA	2104	C	C2'-C3'-O3'	7.01	124.93	109.50
34	BA	520	G	P-O5'-C5'	7.01	132.12	120.90
35	BB	395	U	C3'-C2'-C1'	-7.01	95.89	101.50
35	BB	833	G	C3'-C2'-C1'	-7.01	95.89	101.50
35	BB	1141	A	C1'-O4'-C4'	-7.01	104.29	109.90
35	BB	1203	C	O3'-P-O5'	-7.01	90.68	104.00
35	BB	1507	U	C5'-C4'-C3'	-7.01	104.78	116.00
85	AA	510	A	C5'-C4'-O4'	7.01	117.51	109.10
85	AA	1220	A	C8-N9-C4	-7.01	103.00	105.80
34	BA	578	C	N1-C1'-C2'	-7.01	104.29	112.00
34	BA	839	U	C3'-C2'-C1'	-7.01	95.89	101.50
34	BA	882	G	O4'-C1'-C2'	7.01	113.91	107.60
34	BA	1425	G	P-O3'-C3'	7.01	128.11	119.70
34	BA	1787	U	C5'-C4'-O4'	-7.01	100.69	109.10
34	BA	1796	A	C8-N9-C4	7.01	108.60	105.80
35	BB	632	U	O4'-C1'-N1	7.01	113.81	108.20
35	BB	809	U	N1-C1'-C2'	-7.01	104.29	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1138	A	C5'-C4'-C3'	-7.01	104.79	116.00
38	BE	52	U	P-O3'-C3'	-7.01	111.29	119.70
38	BE	87	U	C5'-C4'-O4'	7.01	117.51	109.10
41	BH	20	A	C5-C6-N1	7.01	121.20	117.70
48	BO	148	TYR	CB-CG-CD2	-7.01	116.79	121.00
79	Bt	45	ARG	NE-CZ-NH1	7.01	123.81	120.30
85	AA	360	C	C5'-C4'-C3'	7.01	127.22	116.00
85	AA	1062	U	C5'-C4'-C3'	-7.01	104.78	116.00
34	BA	743	A	N1-C2-N3	7.01	132.80	129.30
34	BA	1378	A	P-O3'-C3'	-7.01	111.29	119.70
35	BB	496	C	P-O3'-C3'	-7.01	111.29	119.70
35	BB	799	A	C2'-C3'-O3'	7.01	124.92	109.50
37	BD	40	C	P-O5'-C5'	-7.01	109.69	120.90
38	BE	68	U	C2-N1-C1'	-7.01	109.29	117.70
77	Br	308	LYS	CA-C-N	-7.01	101.78	117.20
85	AA	575	G	C8-N9-C1'	7.01	136.11	127.00
85	AA	772	C	P-O3'-C3'	-7.01	111.29	119.70
85	AA	876	U	O4'-C1'-N1	7.01	113.81	108.20
85	AA	1501	A	P-O3'-C3'	-7.01	111.29	119.70
85	AA	1757	C	C2-N1-C1'	-7.01	111.09	118.80
85	AA	2056	C	C1'-O4'-C4'	-7.01	104.29	109.90
34	BA	288	U	P-O5'-C5'	-7.01	109.69	120.90
34	BA	804	G	P-O3'-C3'	7.01	128.11	119.70
34	BA	1086	A	P-O3'-C3'	-7.01	111.29	119.70
34	BA	1598	U	O3'-P-O5'	-7.01	90.69	104.00
38	BE	20	C	P-O5'-C5'	-7.01	109.69	120.90
38	BE	64	A	O4'-C1'-N9	7.01	113.81	108.20
73	Bn	46	ARG	NE-CZ-NH1	7.01	123.80	120.30
85	AA	207	G	P-O3'-C3'	-7.01	111.29	119.70
85	AA	377	U	C1'-O4'-C4'	-7.01	104.30	109.90
85	AA	1213	U	C6-N1-C1'	-7.01	111.39	121.20
85	AA	1554	C	O4'-C1'-N1	7.01	113.80	108.20
34	BA	596	G	N3-C4-C5	-7.00	125.10	128.60
34	BA	907	A	N1-C6-N6	-7.00	114.40	118.60
34	BA	911	G	N9-C1'-C2'	-7.00	104.29	112.00
35	BB	1476	C	O4'-C1'-N1	7.00	113.80	108.20
85	AA	475	A	N9-C1'-C2'	-7.00	104.30	112.00
34	BA	6	C	N1-C2-O2	7.00	123.10	118.90
34	BA	243	C	P-O3'-C3'	-7.00	111.30	119.70
34	BA	696	A	C5'-C4'-C3'	7.00	127.21	116.00
34	BA	1704	G	P-O3'-C3'	-7.00	111.30	119.70
35	BB	533	U	C3'-C2'-C1'	-7.00	95.90	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	706	G	C5-C6-O6	-7.00	124.40	128.60
35	BB	977	G	C4-N9-C1'	7.00	135.60	126.50
85	AA	790	A	N7-C8-N9	7.00	117.30	113.80
85	AA	1247	A	C8-N9-C4	7.00	108.60	105.80
85	AA	1535	C	C1'-O4'-C4'	-7.00	104.30	109.90
34	BA	584	A	O4'-C4'-C3'	-7.00	97.00	104.00
34	BA	609	G	P-O5'-C5'	7.00	132.10	120.90
34	BA	853	A	P-O3'-C3'	-7.00	111.30	119.70
35	BB	801	G	O4'-C1'-N9	7.00	113.80	108.20
35	BB	1186	A	C8-N9-C4	7.00	108.60	105.80
38	BE	4	A	P-O5'-C5'	-7.00	109.70	120.90
38	BE	51	C	P-O3'-C3'	-7.00	111.30	119.70
38	BE	146	U	N3-C4-O4	-7.00	114.50	119.40
48	BO	77	ARG	NE-CZ-NH1	7.00	123.80	120.30
52	BS	9	TYR	CB-CG-CD1	7.00	125.20	121.00
54	BU	65	TRP	CB-CG-CD2	-7.00	117.50	126.60
73	Bn	16	HIS	N-CA-C	-7.00	92.10	111.00
85	AA	1122	U	C2-N3-C4	-7.00	122.80	127.00
34	BA	862	C	N1-C1'-C2'	-7.00	104.30	112.00
36	BC	23	G	N1-C2-N2	-7.00	109.90	116.20
41	BH	135	U	O4'-C1'-N1	7.00	113.80	108.20
34	BA	1434	U	O5'-C5'-C4'	-7.00	98.40	111.70
35	BB	710	A	P-O5'-C5'	-7.00	109.70	120.90
35	BB	1132	A	C4-N9-C1'	-7.00	113.70	126.30
36	BC	85	U	O4'-C1'-N1	7.00	113.80	108.20
53	BT	64	ARG	NE-CZ-NH1	7.00	123.80	120.30
73	Bn	58	ARG	NE-CZ-NH1	7.00	123.80	120.30
85	AA	195	C	P-O3'-C3'	-7.00	111.30	119.70
85	AA	279	C	C5-C6-N1	7.00	124.50	121.00
85	AA	663	C	C5'-C4'-C3'	-7.00	104.80	116.00
85	AA	2240	G	C8-N9-C4	7.00	109.20	106.40
86	AB	8	U	C4'-C3'-C2'	7.00	109.60	102.60
34	BA	1524	G	N9-C4-C5	-7.00	102.60	105.40
34	BA	1599	A	C5'-C4'-O4'	7.00	117.50	109.10
35	BB	148	C	N3-C2-O2	-7.00	117.00	121.90
35	BB	1322	A	C3'-C2'-C1'	-7.00	95.90	101.50
38	BE	130	G	N3-C2-N2	7.00	124.80	119.90
39	BF	25	G	P-O5'-C5'	-7.00	109.71	120.90
41	BH	26	C	C6-N1-C2	-7.00	117.50	120.30
42	BI	13	ARG	NE-CZ-NH1	7.00	123.80	120.30
50	BQ	37	ARG	NE-CZ-NH1	7.00	123.80	120.30
85	AA	193	C	O4'-C1'-N1	7.00	113.80	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	339	A	C3'-C2'-C1'	-7.00	95.90	101.50
34	BA	98	A	O4'-C1'-C2'	7.00	113.90	107.60
34	BA	714	G	C5-C6-O6	-7.00	124.40	128.60
34	BA	1815	G	N1-C6-O6	-7.00	115.70	119.90
35	BB	5	A	C4-C5-C6	-7.00	113.50	117.00
35	BB	434	A	C6-N1-C2	-7.00	114.40	118.60
38	BE	85	G	C5-C6-O6	-7.00	124.40	128.60
84	By	113	PHE	CB-CG-CD1	7.00	125.70	120.80
34	BA	281	C	O4'-C1'-N1	6.99	113.79	108.20
34	BA	817	U	C2-N3-C4	-6.99	122.80	127.00
35	BB	763	U	O4'-C1'-N1	6.99	113.80	108.20
85	AA	1189	A	C4'-C3'-C2'	-6.99	95.61	102.60
85	AA	1356	U	C6-N1-C1'	-6.99	111.41	121.20
34	BA	301	U	O4'-C1'-N1	6.99	113.79	108.20
34	BA	366	G	C5'-C4'-O4'	6.99	117.49	109.10
34	BA	640	U	C2-N1-C1'	6.99	126.09	117.70
85	AA	1892	G	C5-C6-O6	-6.99	124.41	128.60
85	AA	2243	G	O4'-C1'-N9	6.99	113.79	108.20
4	A3	9	ARG	NE-CZ-NH1	6.99	123.80	120.30
34	BA	138	C	O5'-C5'-C4'	-6.99	98.42	111.70
34	BA	1779	U	C2-N1-C1'	-6.99	109.31	117.70
34	BA	1794	A	O4'-C1'-N9	6.99	113.79	108.20
35	BB	1306	G	N3-C2-N2	6.99	124.79	119.90
38	BE	101	C	N1-C1'-C2'	-6.99	104.31	112.00
38	BE	167	U	C2-N3-C4	-6.99	122.81	127.00
39	BF	41	U	C5'-C4'-C3'	-6.99	104.81	116.00
41	BH	112	U	N3-C2-O2	-6.99	117.31	122.20
85	AA	251	A	C5-C6-N6	6.99	129.29	123.70
85	AA	1217	U	C2-N1-C1'	-6.99	109.31	117.70
85	AA	2019	G	C5'-C4'-C3'	-6.99	104.81	116.00
85	AA	2064	A	P-O3'-C3'	-6.99	111.31	119.70
86	AB	3	C	C6-N1-C1'	-6.99	112.41	120.80
34	BA	1107	A	P-O3'-C3'	-6.99	111.31	119.70
35	BB	800	U	C2-N3-C4	-6.99	122.81	127.00
38	BE	61	A	N1-C6-N6	-6.99	114.41	118.60
41	BH	74	G	C5-N7-C8	-6.99	100.81	104.30
65	Bf	406	ARG	NE-CZ-NH2	-6.99	116.81	120.30
85	AA	210	G	C5'-C4'-C3'	-6.99	104.82	116.00
85	AA	1660	U	P-O3'-C3'	6.99	128.09	119.70
34	BA	664	C	C1'-O4'-C4'	-6.99	104.31	109.90
34	BA	1173	C	C6-N1-C1'	6.99	129.19	120.80
36	BC	106	G	C5'-C4'-C3'	-6.99	104.82	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	542	G	C3'-C2'-C1'	-6.99	95.91	101.50
34	BA	422	C	P-O3'-C3'	-6.99	111.32	119.70
34	BA	1215	U	N3-C2-O2	-6.99	117.31	122.20
34	BA	1458	A	C5'-C4'-C3'	6.99	127.18	116.00
35	BB	16	G	C5'-C4'-C3'	6.99	127.18	116.00
38	BE	32	U	C4'-C3'-C2'	-6.99	95.61	102.60
38	BE	98	C	O4'-C1'-N1	6.99	113.79	108.20
82	Bw	211	ARG	NE-CZ-NH1	6.99	123.79	120.30
85	AA	182	C	C6-N1-C1'	-6.99	112.42	120.80
85	AA	1048	C	C6-N1-C1'	6.99	129.18	120.80
85	AA	1345	C	O4'-C1'-N1	6.99	113.79	108.20
85	AA	1528	A	O5'-C5'-C4'	-6.99	98.43	111.70
85	AA	2109	G	P-O3'-C3'	-6.99	111.32	119.70
34	BA	1404	A	C2-N3-C4	6.98	114.09	110.60
35	BB	1480	G	C1'-O4'-C4'	-6.98	104.31	109.90
85	AA	1236	G	C4-N9-C1'	-6.98	117.42	126.50
34	BA	209	A	C8-N9-C1'	6.98	140.27	127.70
34	BA	610	A	N9-C1'-C2'	-6.98	104.32	112.00
34	BA	692	U	C2'-C3'-O3'	6.98	124.87	113.70
34	BA	961	C	C3'-C2'-C1'	-6.98	95.91	101.50
34	BA	1275	G	C8-N9-C1'	6.98	136.08	127.00
35	BB	1170	U	C3'-C2'-C1'	-6.98	95.91	101.50
35	BB	1369	A	P-O3'-C3'	6.98	128.08	119.70
36	BC	42	G	P-O3'-C3'	-6.98	111.32	119.70
41	BH	46	C	P-O3'-C3'	-6.98	111.32	119.70
85	AA	48	G	N3-C2-N2	6.98	124.79	119.90
85	AA	586	G	P-O3'-C3'	-6.98	111.32	119.70
85	AA	674	U	N1-C2-N3	6.98	119.09	114.90
85	AA	1518	A	C1'-O4'-C4'	-6.98	104.31	109.90
85	AA	1803	U	O4'-C1'-N1	6.98	113.79	108.20
15	AG	18	TYR	CB-CG-CD2	-6.98	116.81	121.00
34	BA	801	U	C6-N1-C2	-6.98	116.81	121.00
35	BB	1218	G	N9-C1'-C2'	-6.98	104.32	112.00
36	BC	78	G	C1'-O4'-C4'	-6.98	104.32	109.90
85	AA	124	A	O3'-P-O5'	-6.98	90.74	104.00
85	AA	287	G	C8-N9-C4	-6.98	103.61	106.40
85	AA	1465	C	O3'-P-O5'	6.98	117.26	104.00
85	AA	1971	G	C5'-C4'-C3'	-6.98	104.83	116.00
35	BB	967	G	C3'-C2'-C1'	-6.98	95.92	101.50
65	Bf	212	ARG	NE-CZ-NH1	6.98	123.79	120.30
85	AA	460	U	C6-N1-C2	-6.98	116.81	121.00
34	BA	449	G	C8-N9-C4	6.98	109.19	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	634	U	P-O5'-C5'	6.98	132.06	120.90
35	BB	1165	A	C4-C5-C6	-6.98	113.51	117.00
38	BE	183	C	O4'-C1'-N1	6.98	113.78	108.20
40	BG	166	C	N3-C2-O2	-6.98	117.02	121.90
41	BH	28	U	P-O5'-C5'	6.98	132.06	120.90
41	BH	86	G	OP1-P-OP2	-6.98	109.13	119.60
56	BW	111	MET	CG-SD-CE	-6.98	89.03	100.20
85	AA	710	A	C4-N9-C1'	6.98	138.86	126.30
85	AA	1156	A	N1-C6-N6	6.98	122.79	118.60
85	AA	1383	C	O4'-C1'-N1	6.98	113.78	108.20
35	BB	1076	U	N3-C2-O2	-6.98	117.32	122.20
35	BB	1352	C	C6-N1-C2	-6.98	117.51	120.30
42	BI	25	TYR	CB-CG-CD2	-6.98	116.81	121.00
85	AA	42	G	C6-N1-C2	-6.98	120.92	125.10
85	AA	1985	C	C6-N1-C2	-6.98	117.51	120.30
85	AA	2112	G	C5'-C4'-C3'	-6.98	104.84	116.00
34	BA	462	C	C1'-O4'-C4'	-6.97	104.32	109.90
34	BA	462	C	C6-N1-C1'	-6.97	112.43	120.80
34	BA	822	U	O4'-C1'-N1	6.97	113.78	108.20
34	BA	976	C	N3-C4-N4	-6.97	113.12	118.00
34	BA	1315	C	C5'-C4'-C3'	-6.97	104.84	116.00
35	BB	61	A	OP1-P-OP2	-6.97	109.14	119.60
35	BB	465	C	C3'-C2'-C1'	-6.97	95.92	101.50
35	BB	1508	G	O3'-P-O5'	-6.97	90.75	104.00
36	BC	124	A	C6-C5-N7	-6.97	127.42	132.30
37	BD	4	U	C5-C6-N1	-6.97	119.21	122.70
38	BE	107	U	O4'-C4'-C3'	-6.97	97.03	104.00
40	BG	36	G	P-O5'-C5'	-6.97	109.74	120.90
48	BO	201	ARG	NE-CZ-NH1	6.97	123.79	120.30
85	AA	172	A	N1-C6-N6	6.97	122.78	118.60
85	AA	548	G	C3'-C2'-C1'	-6.97	95.92	101.50
85	AA	1495	G	N3-C2-N2	6.97	124.78	119.90
85	AA	1549	G	N1-C6-O6	6.97	124.08	119.90
34	BA	604	G	P-O3'-C3'	6.97	128.07	119.70
34	BA	635	G	OP1-P-OP2	-6.97	109.14	119.60
34	BA	1222	C	C5'-C4'-C3'	-6.97	104.84	116.00
34	BA	1809	G	C5-C6-N1	6.97	114.99	111.50
35	BB	1019	C	O4'-C1'-N1	6.97	113.78	108.20
37	BD	32	A	O4'-C1'-N9	6.97	113.78	108.20
85	AA	864	C	C3'-C2'-C1'	-6.97	95.92	101.50
34	BA	1732	A	N3-C4-C5	-6.97	121.92	126.80
70	Bk	110	ARG	NE-CZ-NH1	6.97	123.78	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2215	C	N3-C2-O2	-6.97	117.02	121.90
34	BA	129	U	C4'-C3'-C2'	6.97	109.57	102.60
34	BA	1491	U	P-O5'-C5'	-6.97	109.75	120.90
34	BA	1794	A	C5'-C4'-C3'	6.97	127.15	116.00
35	BB	743	C	C4'-C3'-C2'	-6.97	95.63	102.60
35	BB	1298	C	C5-C4-N4	6.97	125.08	120.20
36	BC	145	G	C5-C6-O6	6.97	132.78	128.60
40	BG	35	G	C8-N9-C1'	6.97	136.06	127.00
85	AA	735	G	C8-N9-C1'	6.97	136.06	127.00
85	AA	881	C	N3-C2-O2	-6.97	117.02	121.90
34	BA	1207	A	N1-C6-N6	6.97	122.78	118.60
34	BA	1366	C	C6-N1-C2	-6.97	117.51	120.30
35	BB	680	A	O4'-C1'-C2'	6.97	113.87	107.60
35	BB	1196	A	N1-C2-N3	-6.97	125.82	129.30
85	AA	771	A	C6-N1-C2	-6.97	114.42	118.60
85	AA	854	A	C1'-O4'-C4'	-6.97	104.33	109.90
85	AA	2089	G	C8-N9-C1'	6.97	136.06	127.00
34	BA	263	G	C3'-C2'-C1'	-6.97	95.93	101.50
38	BE	93	U	C5'-C4'-O4'	6.97	117.46	109.10
42	BI	104	ALA	C-N-CA	6.97	139.12	121.70
85	AA	306	C	C6-N1-C2	-6.97	117.51	120.30
85	AA	431	G	C1'-O4'-C4'	-6.97	104.33	109.90
85	AA	1335	C	O4'-C1'-N1	6.97	113.77	108.20
34	BA	671	C	O4'-C1'-N1	6.96	113.77	108.20
35	BB	391	G	C8-N9-C1'	6.96	136.05	127.00
35	BB	620	G	N9-C1'-C2'	-6.96	104.34	112.00
49	BP	23	ARG	NE-CZ-NH1	6.96	123.78	120.30
54	BU	159	TYR	CB-CG-CD1	-6.96	116.82	121.00
85	AA	571	G	C5'-C4'-C3'	6.96	127.14	116.00
85	AA	588	G	C4'-C3'-C2'	-6.96	95.64	102.60
85	AA	594	C	O4'-C1'-N1	6.96	113.77	108.20
85	AA	1879	U	N3-C2-O2	-6.96	117.32	122.20
34	BA	344	G	C6-N1-C2	-6.96	120.92	125.10
35	BB	390	G	N1-C6-O6	-6.96	115.72	119.90
35	BB	392	G	C4-N9-C1'	-6.96	117.45	126.50
35	BB	474	G	C1'-O4'-C4'	-6.96	104.33	109.90
35	BB	660	G	N1-C6-O6	6.96	124.08	119.90
37	BD	13	A	P-O3'-C3'	-6.96	111.34	119.70
40	BG	15	G	C4-N9-C1'	-6.96	117.45	126.50
85	AA	249	C	N1-C2-O2	6.96	123.08	118.90
85	AA	1085	U	O4'-C1'-N1	6.96	113.77	108.20
85	AA	2058	C	C2-N3-C4	-6.96	116.42	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	16	U	O4'-C4'-C3'	-6.96	97.04	104.00
1	A0	107	MET	CG-SD-CE	-6.96	89.06	100.20
34	BA	856	G	C5-C6-N1	6.96	114.98	111.50
34	BA	1411	C	P-O5'-C5'	6.96	132.04	120.90
34	BA	1819	U	O5'-P-OP1	-6.96	99.44	105.70
35	BB	514	G	C8-N9-C1'	6.96	136.05	127.00
35	BB	741	A	C5'-C4'-C3'	-6.96	104.86	116.00
35	BB	971	A	OP1-P-O3'	6.96	120.52	105.20
35	BB	1419	G	C8-N9-C4	-6.96	103.61	106.40
40	BG	43	U	C5'-C4'-C3'	6.96	127.14	116.00
85	AA	351	C	C5'-C4'-C3'	-6.96	104.86	116.00
85	AA	1495	G	N9-C1'-C2'	-6.96	104.34	112.00
34	BA	54	A	P-O3'-C3'	-6.96	111.35	119.70
35	BB	1483	A	C5'-C4'-O4'	6.96	117.45	109.10
3	A2	48	ARG	NE-CZ-NH1	6.96	123.78	120.30
34	BA	78	U	P-O3'-C3'	-6.96	111.35	119.70
34	BA	652	C	O4'-C1'-N1	6.96	113.77	108.20
34	BA	843	G	C8-N9-C4	-6.96	103.62	106.40
34	BA	1563	G	O5'-P-OP2	-6.96	99.44	105.70
35	BB	357	C	O4'-C1'-N1	6.96	113.77	108.20
35	BB	823	G	C8-N9-C4	-6.96	103.62	106.40
35	BB	1505	U	C5'-C4'-C3'	-6.96	104.87	116.00
36	BC	86	U	C1'-O4'-C4'	-6.96	104.33	109.90
39	BF	2	G	C5-C6-O6	-6.96	124.42	128.60
85	AA	390	U	C4'-C3'-C2'	-6.96	95.64	102.60
85	AA	1952	C	C6-N1-C2	-6.96	117.52	120.30
13	AE	76	PHE	CB-CG-CD1	6.96	125.67	120.80
34	BA	745	A	C5'-C4'-C3'	-6.96	104.87	116.00
35	BB	46	U	C4'-C3'-C2'	6.96	109.56	102.60
35	BB	363	A	P-O5'-C5'	6.96	132.03	120.90
38	BE	8	G	P-O3'-C3'	-6.96	111.35	119.70
38	BE	73	A	C5'-C4'-C3'	6.96	127.13	116.00
47	BN	19	PRO	N-CA-C	6.96	130.19	112.10
85	AA	748	C	C3'-C2'-C1'	-6.96	95.93	101.50
34	BA	355	U	N3-C2-O2	-6.96	117.33	122.20
34	BA	692	U	C4-C5-C6	-6.96	115.53	119.70
34	BA	1049	G	C5-N7-C8	-6.96	100.82	104.30
35	BB	90	G	C8-N9-C1'	6.96	136.04	127.00
35	BB	1510	G	N9-C1'-C2'	-6.96	104.35	112.00
85	AA	307	G	C4'-C3'-O3'	6.96	126.91	113.00
85	AA	1404	G	O4'-C1'-N9	6.96	113.76	108.20
34	BA	615	A	C3'-C2'-C1'	-6.95	95.94	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	641	U	N3-C4-O4	-6.95	114.53	119.40
34	BA	825	G	C1'-O4'-C4'	-6.95	104.34	109.90
34	BA	1652	G	P-O3'-C3'	-6.95	111.36	119.70
35	BB	1065	G	C8-N9-C1'	-6.95	117.96	127.00
39	BF	54	U	C3'-C2'-C1'	6.95	107.06	101.50
85	AA	650	G	C6-N1-C2	-6.95	120.93	125.10
85	AA	788	G	O4'-C4'-C3'	-6.95	97.05	104.00
85	AA	838	G	N3-C4-N9	6.95	130.17	126.00
85	AA	1516	A	O4'-C1'-C2'	6.95	113.86	107.60
85	AA	1536	C	P-O5'-C5'	-6.95	109.78	120.90
85	AA	2132	A	O4'-C1'-N9	6.95	113.76	108.20
34	BA	794	G	C1'-O4'-C4'	-6.95	104.34	109.90
35	BB	168	U	O4'-C1'-N1	6.95	113.76	108.20
85	AA	519	A	C4'-C3'-C2'	-6.95	95.65	102.60
4	A3	210	ARG	NE-CZ-NH2	-6.95	116.83	120.30
5	A4	3	ALA	N-CA-CB	6.95	119.83	110.10
35	BB	63	A	C1'-O4'-C4'	-6.95	104.34	109.90
35	BB	1306	G	O5'-C5'-C4'	-6.95	98.49	111.70
38	BE	17	U	C1'-O4'-C4'	-6.95	104.34	109.90
40	BG	2	U	C2-N3-C4	-6.95	122.83	127.00
85	AA	644	A	N1-C6-N6	6.95	122.77	118.60
34	BA	110	C	N3-C4-C5	-6.95	119.12	121.90
35	BB	137	A	C4-N9-C1'	-6.95	113.79	126.30
35	BB	1071	G	P-O3'-C3'	-6.95	111.36	119.70
38	BE	39	U	C2-N3-C4	-6.95	122.83	127.00
40	BG	135	C	O4'-C1'-N1	6.95	113.76	108.20
59	BZ	75	TRP	CB-CG-CD2	-6.95	117.57	126.60
85	AA	1217	U	N3-C2-O2	-6.95	117.34	122.20
85	AA	2074	G	C4-N9-C1'	-6.95	117.47	126.50
34	BA	1331	G	P-O3'-C3'	-6.95	111.36	119.70
35	BB	518	G	C5'-C4'-O4'	6.95	117.44	109.10
36	BC	15	G	C5-C6-O6	-6.95	124.43	128.60
85	AA	430	G	C5-C6-O6	-6.95	124.43	128.60
85	AA	2154	C	P-O5'-C5'	-6.95	109.78	120.90
8	A7	211	CYS	N-CA-C	-6.95	92.25	111.00
34	BA	87	G	C1'-O4'-C4'	-6.95	104.34	109.90
34	BA	89	G	C8-N9-C1'	6.95	136.03	127.00
34	BA	103	G	P-O5'-C5'	-6.95	109.79	120.90
34	BA	114	U	P-O5'-C5'	-6.95	109.79	120.90
34	BA	702	G	O3'-P-O5'	6.95	117.19	104.00
34	BA	994	G	C5'-C4'-C3'	-6.95	104.89	116.00
35	BB	315	C	O4'-C1'-N1	6.95	113.76	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1015	U	O4'-C1'-N1	6.95	113.76	108.20
35	BB	1016	C	C3'-C2'-C1'	-6.95	95.94	101.50
37	BD	78	C	O4'-C1'-N1	6.95	113.76	108.20
39	BF	34	C	C4'-C3'-C2'	-6.95	95.65	102.60
85	AA	1056	C	C6-N1-C1'	-6.95	112.47	120.80
85	AA	1440	C	C3'-C2'-C1'	-6.95	95.94	101.50
85	AA	1632	G	C5-C6-O6	-6.95	124.43	128.60
34	BA	1538	G	N1-C2-N2	-6.94	109.95	116.20
35	BB	255	A	O4'-C1'-N9	6.94	113.76	108.20
85	AA	764	U	P-O3'-C3'	6.94	128.03	119.70
34	BA	310	C	C5'-C4'-C3'	-6.94	104.89	116.00
34	BA	639	U	O4'-C1'-N1	6.94	113.75	108.20
40	BG	13	A	C5'-C4'-O4'	6.94	117.43	109.10
67	Bh	50	ARG	NE-CZ-NH2	6.94	123.77	120.30
85	AA	456	A	C4-N9-C1'	-6.94	113.81	126.30
34	BA	89	G	N1-C6-O6	6.94	124.06	119.90
34	BA	679	U	P-O3'-C3'	6.94	128.03	119.70
34	BA	1544	G	C1'-O4'-C4'	-6.94	104.35	109.90
34	BA	1603	A	C4'-C3'-C2'	-6.94	95.66	102.60
35	BB	374	A	C6-N1-C2	-6.94	114.44	118.60
35	BB	1054	G	C5-C6-O6	-6.94	124.44	128.60
41	BH	100	A	OP1-P-O3'	6.94	120.47	105.20
85	AA	265	A	C1'-O4'-C4'	-6.94	104.35	109.90
38	BE	96	G	N1-C6-O6	-6.94	115.74	119.90
34	BA	277	A	O4'-C1'-N9	6.94	113.75	108.20
34	BA	308	C	O4'-C1'-N1	6.94	113.75	108.20
34	BA	1282	G	C5'-C4'-C3'	-6.94	104.90	116.00
34	BA	1636	C	C6-N1-C2	6.94	123.08	120.30
35	BB	509	A	C8-N9-C4	6.94	108.58	105.80
35	BB	1000	U	P-O5'-C5'	-6.94	109.80	120.90
38	BE	146	U	C5-C6-N1	-6.94	119.23	122.70
67	Bh	163	ARG	NE-CZ-NH1	6.94	123.77	120.30
77	Br	99	ARG	CB-CA-C	-6.94	96.52	110.40
85	AA	189	G	O4'-C1'-N9	6.94	113.75	108.20
85	AA	236	G	O4'-C1'-N9	6.94	113.75	108.20
86	AB	67	C	C6-N1-C2	-6.94	117.53	120.30
53	BT	75	HIS	CA-CB-CG	-6.94	101.81	113.60
85	AA	722	G	C4-N9-C1'	-6.94	117.48	126.50
85	AA	859	G	C2-N3-C4	6.94	115.37	111.90
85	AA	1291	A	O4'-C1'-N9	6.94	113.75	108.20
21	AM	41	PHE	CB-CG-CD2	-6.93	115.95	120.80
34	BA	54	A	C3'-C2'-C1'	-6.93	95.95	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	167	U	O4'-C1'-N1	6.93	113.75	108.20
34	BA	195	G	C8-N9-C4	-6.93	103.63	106.40
34	BA	1676	A	O5'-P-OP2	-6.93	99.46	105.70
35	BB	284	C	O4'-C1'-N1	6.93	113.75	108.20
35	BB	1022	C	N1-C2-O2	6.93	123.06	118.90
35	BB	1441	C	C5'-C4'-O4'	6.93	117.42	109.10
35	BB	1467	A	C5'-C4'-C3'	6.93	127.09	116.00
40	BG	166	C	C5-C4-N4	6.93	125.05	120.20
85	AA	1514	A	C5-C6-N6	-6.93	118.15	123.70
85	AA	1636	C	C6-N1-C2	-6.93	117.53	120.30
34	BA	568	G	C4'-C3'-C2'	-6.93	95.67	102.60
35	BB	1506	C	C6-N1-C1'	-6.93	112.48	120.80
34	BA	290	G	N3-C2-N2	6.93	124.75	119.90
35	BB	507	G	N9-C1'-C2'	-6.93	104.38	112.00
35	BB	530	C	O4'-C1'-N1	6.93	113.75	108.20
39	BF	56	C	O5'-P-OP2	-6.93	99.46	105.70
41	BH	113	G	C1'-O4'-C4'	-6.93	104.36	109.90
85	AA	1765	G	O4'-C1'-N9	6.93	113.75	108.20
34	BA	155	U	N1-C2-N3	6.93	119.06	114.90
34	BA	435	U	C5'-C4'-C3'	6.93	127.09	116.00
34	BA	571	G	C8-N9-C4	-6.93	103.63	106.40
34	BA	1202	G	N9-C1'-C2'	-6.93	104.38	112.00
34	BA	1504	A	P-O5'-C5'	6.93	131.99	120.90
35	BB	871	C	C1'-O4'-C4'	-6.93	104.36	109.90
38	BE	33	C	O4'-C1'-N1	6.93	113.74	108.20
38	BE	82	C	C1'-O4'-C4'	-6.93	104.36	109.90
40	BG	7	U	O4'-C1'-N1	6.93	113.74	108.20
41	BH	15	A	C4'-C3'-C2'	-6.93	95.67	102.60
47	BN	187	TYR	N-CA-C	-6.93	92.29	111.00
65	Bf	150	HIS	CA-CB-CG	-6.93	101.82	113.60
85	AA	128	U	C3'-C2'-C1'	-6.93	95.96	101.50
85	AA	231	G	O4'-C1'-N9	6.93	113.74	108.20
34	BA	1733	G	O4'-C1'-N9	6.93	113.74	108.20
85	AA	1790	G	N9-C4-C5	-6.93	102.63	105.40
85	AA	2164	G	C3'-C2'-C1'	-6.93	95.96	101.50
22	AO	95	ARG	NE-CZ-NH1	6.93	123.76	120.30
34	BA	814	C	N1-C1'-C2'	-6.93	104.38	112.00
34	BA	1542	A	N9-C1'-C2'	-6.93	104.38	112.00
35	BB	1504	U	P-O3'-C3'	6.93	128.01	119.70
41	BH	133	U	O4'-C1'-N1	6.93	113.74	108.20
85	AA	2109	G	O4'-C1'-N9	6.93	113.74	108.20
85	AA	2202	G	C5-C6-O6	-6.93	124.44	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	231	U	P-O5'-C5'	6.92	131.98	120.90
34	BA	383	G	N3-C2-N2	-6.92	115.05	119.90
35	BB	93	A	O4'-C1'-N9	6.92	113.74	108.20
39	BF	60	C	C6-N1-C2	-6.92	117.53	120.30
85	AA	210	G	N9-C1'-C2'	-6.92	104.38	112.00
85	AA	1399	U	O4'-C1'-N1	6.92	113.74	108.20
7	A6	53	ARG	NE-CZ-NH1	6.92	123.76	120.30
34	BA	102	G	C3'-C2'-C1'	-6.92	95.96	101.50
34	BA	1286	C	C5-C4-N4	6.92	125.05	120.20
34	BA	1472	G	C8-N9-C4	6.92	109.17	106.40
35	BB	624	A	P-O3'-C3'	-6.92	111.39	119.70
41	BH	111	U	C5-C4-O4	6.92	130.05	125.90
85	AA	878	U	O4'-C1'-N1	6.92	113.74	108.20
85	AA	1091	C	O4'-C1'-N1	6.92	113.74	108.20
34	BA	930	A	C5'-C4'-C3'	-6.92	104.92	116.00
34	BA	1294	C	O4'-C4'-C3'	6.92	111.64	106.10
34	BA	1485	U	C6-N1-C1'	6.92	130.89	121.20
34	BA	1659	G	O4'-C1'-N9	6.92	113.74	108.20
35	BB	449	C	C6-N1-C2	-6.92	117.53	120.30
35	BB	942	G	P-O3'-C3'	6.92	128.01	119.70
40	BG	46	G	P-O3'-C3'	-6.92	111.39	119.70
41	BH	11	C	N1-C2-O2	6.92	123.05	118.90
59	BZ	78	HIS	CB-CA-C	-6.92	96.56	110.40
85	AA	633	C	C2-N1-C1'	6.92	126.41	118.80
85	AA	1833	C	C5'-C4'-O4'	6.92	117.41	109.10
34	BA	286	C	C5'-C4'-C3'	6.92	127.07	116.00
34	BA	687	G	C6-N1-C2	-6.92	120.95	125.10
34	BA	1025	A	P-O3'-C3'	-6.92	111.40	119.70
40	BG	13	A	C1'-O4'-C4'	-6.92	104.36	109.90
85	AA	25	C	OP1-P-OP2	-6.92	109.22	119.60
85	AA	457	G	C1'-O4'-C4'	-6.92	104.36	109.90
86	AB	58	A	C5'-C4'-C3'	-6.92	104.93	116.00
27	AT	34	HIS	CA-CB-CG	6.92	125.36	113.60
34	BA	12	G	C5-N7-C8	-6.92	100.84	104.30
35	BB	887	G	P-O3'-C3'	-6.92	111.40	119.70
38	BE	17	U	C5'-C4'-C3'	-6.92	104.93	116.00
85	AA	164	G	C5'-C4'-C3'	-6.92	104.93	116.00
85	AA	590	U	O4'-C1'-N1	6.92	113.73	108.20
85	AA	657	C	C6-N1-C2	-6.92	117.53	120.30
85	AA	688	C	C5'-C4'-O4'	6.92	117.40	109.10
85	AA	1612	C	P-O5'-C5'	-6.92	109.83	120.90
34	BA	440	A	O5'-C5'-C4'	-6.92	98.56	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	939	C	N3-C2-O2	-6.92	117.06	121.90
35	BB	76	C	P-O5'-C5'	6.92	131.97	120.90
37	BD	81	C	N1-C2-O2	6.92	123.05	118.90
44	BK	9	TYR	CB-CG-CD1	-6.92	116.85	121.00
65	Bf	398	ARG	NE-CZ-NH2	-6.92	116.84	120.30
74	Bo	56	ARG	NE-CZ-NH1	6.92	123.76	120.30
85	AA	79	G	C8-N9-C1'	6.92	135.99	127.00
85	AA	819	G	N3-C4-C5	-6.92	125.14	128.60
85	AA	823	C	C3'-C2'-C1'	-6.92	95.97	101.50
25	AR	43	ASN	CA-CB-CG	-6.92	98.19	113.40
34	BA	1577	U	C6-N1-C1'	6.92	130.88	121.20
35	BB	752	A	N1-C6-N6	-6.92	114.45	118.60
35	BB	1170	U	C1'-O4'-C4'	-6.92	104.37	109.90
35	BB	1254	G	N1-C6-O6	-6.92	115.75	119.90
38	BE	193	A	C5-N7-C8	-6.92	100.44	103.90
34	BA	13	U	C1'-O4'-C4'	-6.91	104.37	109.90
34	BA	591	G	C4-N9-C1'	-6.91	117.51	126.50
34	BA	1433	U	C1'-O4'-C4'	-6.91	104.37	109.90
35	BB	574	G	C8-N9-C1'	6.91	135.99	127.00
77	Br	237	HIS	CA-CB-CG	-6.91	101.85	113.60
85	AA	130	G	O3'-P-O5'	-6.91	90.86	104.00
34	BA	455	A	P-O5'-C5'	-6.91	109.84	120.90
35	BB	976	U	N1-C2-N3	6.91	119.05	114.90
85	AA	492	C	C2-N1-C1'	-6.91	111.20	118.80
85	AA	2235	C	O4'-C1'-N1	6.91	113.73	108.20
19	AK	135	TRP	CB-CG-CD2	-6.91	117.62	126.60
34	BA	875	G	C3'-C2'-C1'	-6.91	95.97	101.50
34	BA	1814	U	C6-N1-C1'	6.91	130.87	121.20
35	BB	860	U	C2-N1-C1'	6.91	125.99	117.70
38	BE	155	C	C6-N1-C1'	-6.91	112.51	120.80
74	Bo	14	TYR	CB-CG-CD2	-6.91	116.85	121.00
85	AA	1505	G	C5-C6-O6	6.91	132.75	128.60
4	A3	62	PHE	CB-CG-CD2	-6.91	115.97	120.80
19	AK	34	ASN	C-N-CA	6.91	138.97	121.70
34	BA	42	A	C4-N9-C1'	-6.91	113.86	126.30
34	BA	300	C	O3'-P-O5'	-6.91	90.88	104.00
34	BA	558	C	OP2-P-O3'	6.91	120.40	105.20
34	BA	1433	U	P-O5'-C5'	-6.91	109.85	120.90
35	BB	713	U	C6-N1-C1'	6.91	130.87	121.20
38	BE	27	A	P-O5'-C5'	-6.91	109.85	120.90
34	BA	185	A	C5-C6-N6	-6.91	118.17	123.70
37	BD	98	G	C3'-C2'-C1'	-6.91	95.97	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	63	G	C6-C5-N7	-6.91	126.26	130.40
47	BN	207	ALA	CB-CA-C	-6.91	99.74	110.10
85	AA	105	A	O4'-C1'-C2'	6.91	113.82	107.60
3	A2	41	ARG	NE-CZ-NH2	6.91	123.75	120.30
34	BA	123	C	N1-C2-O2	-6.91	114.76	118.90
34	BA	306	G	C8-N9-C1'	6.91	135.98	127.00
34	BA	1234	U	C3'-C2'-C1'	-6.91	95.98	101.50
47	BN	205	ARG	NE-CZ-NH1	6.91	123.75	120.30
85	AA	31	C	C3'-C2'-C1'	-6.91	95.98	101.50
85	AA	617	C	C2'-C3'-O3'	6.91	124.75	113.70
6	A5	61	ASP	N-CA-CB	-6.90	98.17	110.60
13	AE	54	ILE	C-N-CA	6.90	136.80	122.30
34	BA	13	U	P-O3'-C3'	-6.90	111.42	119.70
34	BA	1490	U	O4'-C1'-C2'	6.90	113.81	107.60
35	BB	20	U	C2-N3-C4	-6.90	122.86	127.00
35	BB	830	G	O4'-C4'-C3'	-6.90	97.10	104.00
64	Be	174	ARG	NE-CZ-NH1	6.90	123.75	120.30
85	AA	1358	A	C1'-O4'-C4'	-6.90	104.38	109.90
34	BA	625	U	C4'-C3'-C2'	-6.90	95.70	102.60
34	BA	1439	C	O4'-C1'-N1	6.90	113.72	108.20
34	BA	1723	U	O5'-C5'-C4'	6.90	124.81	111.70
34	BA	1845	G	N1-C2-N2	6.90	122.41	116.20
35	BB	38	C	N3-C2-O2	-6.90	117.07	121.90
35	BB	434	A	N1-C6-N6	6.90	122.74	118.60
35	BB	578	G	C6-N1-C2	-6.90	120.96	125.10
35	BB	880	G	P-O3'-C3'	-6.90	111.42	119.70
35	BB	1047	C	N3-C2-O2	-6.90	117.07	121.90
35	BB	1199	A	O3'-P-O5'	-6.90	90.89	104.00
58	BY	64	ARG	CA-CB-CG	6.90	128.59	113.40
85	AA	76	G	C3'-C2'-C1'	-6.90	95.98	101.50
85	AA	261	U	C2-N1-C1'	-6.90	109.42	117.70
85	AA	537	G	C5-C6-O6	-6.90	124.46	128.60
85	AA	1003	G	C8-N9-C1'	6.90	135.97	127.00
29	AV	5	ARG	NE-CZ-NH1	6.90	123.75	120.30
34	BA	18	G	C5-C6-N1	6.90	114.95	111.50
34	BA	765	U	C3'-C2'-C1'	6.90	107.02	101.50
34	BA	1685	C	C2-N1-C1'	-6.90	111.21	118.80
35	BB	817	C	C1'-O4'-C4'	-6.90	104.38	109.90
35	BB	824	C	N1-C1'-C2'	-6.90	104.41	112.00
35	BB	960	C	O4'-C1'-N1	6.90	113.72	108.20
35	BB	1257	A	N1-C6-N6	6.90	122.74	118.60
40	BG	130	G	P-O3'-C3'	-6.90	111.42	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	98	U	N1-C1'-C2'	-6.90	104.41	112.00
85	AA	607	U	C3'-C2'-C1'	-6.90	95.98	101.50
85	AA	986	U	C6-N1-C2	-6.90	116.86	121.00
34	BA	367	G	C8-N9-C1'	-6.90	118.03	127.00
34	BA	934	G	C4-N9-C1'	-6.90	117.53	126.50
58	BY	19	ARG	NE-CZ-NH1	6.90	123.75	120.30
9	A8	33	ARG	NE-CZ-NH2	-6.90	116.85	120.30
34	BA	162	G	N3-C4-N9	-6.90	121.86	126.00
34	BA	592	G	O3'-P-O5'	6.90	117.11	104.00
34	BA	1087	A	O4'-C1'-N9	6.90	113.72	108.20
35	BB	134	G	N3-C2-N2	6.90	124.73	119.90
35	BB	758	A	C4'-C3'-C2'	-6.90	95.70	102.60
35	BB	1369	A	C5'-C4'-O4'	-6.90	100.82	109.10
38	BE	156	C	O4'-C1'-N1	6.90	113.72	108.20
38	BE	171	U	C5-C6-N1	-6.90	119.25	122.70
40	BG	24	A	C5-C6-N6	6.90	129.22	123.70
85	AA	161	A	C2-N3-C4	-6.90	107.15	110.60
85	AA	726	U	C5'-C4'-C3'	-6.90	104.96	116.00
85	AA	831	C	C3'-C2'-C1'	-6.90	95.98	101.50
85	AA	1204	A	N9-C4-C5	6.90	108.56	105.80
21	AM	143	ARG	NE-CZ-NH2	-6.90	116.85	120.30
25	AR	17	TYR	CB-CG-CD1	6.90	125.14	121.00
34	BA	25	C	C6-N1-C2	-6.90	117.54	120.30
34	BA	258	C	C3'-C2'-C1'	-6.90	95.98	101.50
34	BA	467	A	N1-C6-N6	6.90	122.74	118.60
35	BB	412	A	C8-N9-C4	6.90	108.56	105.80
85	AA	1645	G	C8-N9-C1'	6.90	135.96	127.00
34	BA	565	U	O4'-C1'-N1	6.89	113.72	108.20
34	BA	765	U	N3-C2-O2	-6.89	117.37	122.20
35	BB	781	U	C5'-C4'-C3'	6.89	127.03	116.00
36	BC	108	A	N1-C6-N6	-6.89	114.46	118.60
38	BE	51	C	C2-N3-C4	-6.89	116.45	119.90
41	BH	109	G	C5'-C4'-C3'	-6.89	104.97	116.00
85	AA	2117	U	C4-C5-C6	-6.89	115.56	119.70
34	BA	730	C	C1'-O4'-C4'	-6.89	104.39	109.90
34	BA	936	A	O5'-P-OP2	-6.89	99.50	105.70
34	BA	1038	U	O4'-C1'-N1	6.89	113.71	108.20
34	BA	1178	U	P-O3'-C3'	-6.89	111.43	119.70
34	BA	1728	G	C8-N9-C1'	-6.89	118.04	127.00
72	Bm	103	ARG	NE-CZ-NH1	6.89	123.75	120.30
85	AA	141	A	O4'-C1'-N9	6.89	113.71	108.20
85	AA	1232	U	P-O5'-C5'	6.89	131.93	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1346	C	O4'-C1'-N1	6.89	113.71	108.20
85	AA	1928	A	P-O3'-C3'	-6.89	111.43	119.70
35	BB	858	U	P-O5'-C5'	6.89	131.93	120.90
35	BB	1233	U	O4'-C1'-N1	6.89	113.71	108.20
85	AA	551	C	O4'-C1'-N1	6.89	113.71	108.20
8	A7	187	ARG	CG-CD-NE	-6.89	97.33	111.80
34	BA	519	G	C5'-C4'-O4'	6.89	117.37	109.10
34	BA	890	G	C4-N9-C1'	-6.89	117.54	126.50
34	BA	1183	U	P-O3'-C3'	-6.89	111.43	119.70
34	BA	1564	A	P-O5'-C5'	-6.89	109.88	120.90
34	BA	1653	G	O4'-C1'-C2'	6.89	113.80	107.60
35	BB	567	G	N1-C6-O6	-6.89	115.77	119.90
35	BB	1003	G	C1'-O4'-C4'	-6.89	104.39	109.90
35	BB	1099	U	C4'-C3'-C2'	-6.89	95.71	102.60
35	BB	1223	A	N1-C2-N3	-6.89	125.86	129.30
36	BC	26	U	O4'-C1'-N1	-6.89	102.69	108.20
85	AA	232	U	O4'-C1'-N1	6.89	113.71	108.20
85	AA	2129	U	O4'-C4'-C3'	-6.89	97.11	104.00
35	BB	783	U	P-O3'-C3'	6.89	127.97	119.70
35	BB	969	C	C6-N1-C2	-6.89	117.55	120.30
36	BC	70	C	C1'-O4'-C4'	-6.89	104.39	109.90
34	BA	1127	U	P-O5'-C5'	6.89	131.92	120.90
34	BA	1277	G	C5'-C4'-C3'	-6.89	104.98	116.00
34	BA	1540	C	N1-C2-O2	-6.89	114.77	118.90
35	BB	347	G	N1-C6-O6	6.89	124.03	119.90
36	BC	91	G	N9-C1'-C2'	-6.89	104.42	112.00
85	AA	1368	G	N3-C4-C5	-6.89	125.16	128.60
85	AA	1440	C	O3'-P-O5'	6.89	117.08	104.00
85	AA	1457	C	O3'-P-O5'	6.89	117.08	104.00
85	AA	1683	U	O4'-C1'-N1	6.89	113.71	108.20
85	AA	1731	G	C5-C6-N1	6.89	114.94	111.50
85	AA	2141	G	N1-C2-N3	-6.89	119.77	123.90
2	A1	241	GLY	N-CA-C	6.88	130.31	113.10
34	BA	487	A	P-O5'-C5'	6.88	131.91	120.90
34	BA	793	A	C5'-C4'-C3'	-6.88	104.98	116.00
34	BA	965	A	C3'-C2'-C1'	-6.88	95.99	101.50
34	BA	1695	G	N3-C2-N2	6.88	124.72	119.90
35	BB	831	C	C3'-C2'-C1'	-6.88	95.99	101.50
35	BB	970	C	O3'-P-O5'	6.88	117.08	104.00
35	BB	1410	G	P-O3'-C3'	-6.88	111.44	119.70
36	BC	123	G	N3-C2-N2	-6.88	115.08	119.90
40	BG	25	G	C8-N9-C1'	6.88	135.95	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
73	Bn	28	HIS	N-CA-CB	6.88	122.99	110.60
85	AA	2152	C	C5'-C4'-C3'	-6.88	104.98	116.00
35	BB	1252	G	C8-N9-C4	6.88	109.15	106.40
40	BG	43	U	C2-N3-C4	-6.88	122.87	127.00
40	BG	172	C	C5-C6-N1	-6.88	117.56	121.00
85	AA	240	A	C4'-C3'-C2'	-6.88	95.72	102.60
85	AA	372	U	P-O3'-C3'	-6.88	111.44	119.70
85	AA	1151	G	C5'-C4'-C3'	6.88	127.01	116.00
85	AA	1860	A	P-O5'-C5'	6.88	131.91	120.90
85	AA	1906	C	O4'-C1'-N1	6.88	113.71	108.20
15	AG	129	TYR	CA-CB-CG	-6.88	100.33	113.40
34	BA	444	A	N1-C6-N6	6.88	122.73	118.60
34	BA	930	A	N9-C1'-C2'	-6.88	104.43	112.00
34	BA	1311	G	N1-C2-N2	-6.88	110.01	116.20
35	BB	654	C	N1-C1'-C2'	-6.88	104.43	112.00
35	BB	1387	C	N3-C2-O2	-6.88	117.08	121.90
36	BC	153	C	C4'-C3'-C2'	6.88	109.48	102.60
37	BD	78	C	P-O5'-C5'	-6.88	109.89	120.90
85	AA	476	C	C3'-C2'-C1'	-6.88	96.00	101.50
85	AA	847	G	C6-N1-C2	-6.88	120.97	125.10
34	BA	399	G	C4-C5-N7	6.88	113.55	110.80
35	BB	1281	G	C8-N9-C4	6.88	109.15	106.40
37	BD	11	A	C5-C6-N1	6.88	121.14	117.70
65	Bf	350	ASN	C-N-CA	6.88	138.90	121.70
85	AA	330	C	OP1-P-OP2	-6.88	109.28	119.60
85	AA	768	C	C5'-C4'-C3'	-6.88	104.99	116.00
85	AA	1948	A	C3'-C2'-C1'	-6.88	96.00	101.50
34	BA	445	C	C3'-C2'-C1'	-6.88	96.00	101.50
34	BA	516	U	OP1-P-O3'	6.88	120.33	105.20
34	BA	747	G	P-O3'-C3'	-6.88	111.45	119.70
34	BA	842	U	C4-C5-C6	-6.88	115.57	119.70
35	BB	63	A	O4'-C1'-C2'	-6.88	98.92	105.80
35	BB	1105	G	O3'-P-O5'	-6.88	90.93	104.00
35	BB	1427	A	C5-C6-N1	6.88	121.14	117.70
35	BB	1497	C	C6-N1-C2	-6.88	117.55	120.30
39	BF	12	U	P-O5'-C5'	-6.88	109.90	120.90
43	BJ	127	ARG	NE-CZ-NH1	6.88	123.74	120.30
83	Bx	81	ARG	NE-CZ-NH1	6.88	123.74	120.30
85	AA	47	A	C5-C6-N1	6.88	121.14	117.70
85	AA	537	G	C1'-O4'-C4'	-6.88	104.40	109.90
85	AA	1215	A	O3'-P-O5'	-6.88	90.93	104.00
85	AA	1464	G	C6-N1-C2	-6.88	120.97	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1513	U	P-O3'-C3'	-6.88	111.45	119.70
85	AA	1668	G	N7-C8-N9	6.88	116.54	113.10
34	BA	139	U	C6-N1-C2	-6.88	116.87	121.00
34	BA	517	A	O5'-C5'-C4'	-6.88	98.63	111.70
34	BA	747	G	N7-C8-N9	6.88	116.54	113.10
34	BA	1123	G	P-O5'-C5'	6.88	131.90	120.90
34	BA	1697	U	C5'-C4'-O4'	-6.88	100.85	109.10
34	BA	255	G	C4-N9-C1'	-6.88	117.56	126.50
34	BA	1746	G	O4'-C1'-N9	6.88	113.70	108.20
38	BE	28	C	C5'-C4'-C3'	-6.88	105.00	116.00
61	Bb	76	ASP	CB-CG-OD1	-6.88	112.11	118.30
85	AA	75	U	C5-C6-N1	6.88	126.14	122.70
85	AA	909	C	C1'-O4'-C4'	-6.88	104.40	109.90
85	AA	1371	C	P-O5'-C5'	6.88	131.90	120.90
34	BA	118	C	O4'-C1'-N1	6.87	113.70	108.20
34	BA	867	C	O5'-P-OP2	6.87	118.95	110.70
34	BA	1541	G	C8-N9-C4	6.87	109.15	106.40
35	BB	815	G	C8-N9-C1'	6.87	135.93	127.00
47	BN	81	SER	N-CA-CB	6.87	120.81	110.50
85	AA	32	U	C1'-O4'-C4'	-6.87	104.40	109.90
23	AP	151	ARG	NE-CZ-NH1	6.87	123.74	120.30
34	BA	363	G	P-O5'-C5'	-6.87	109.91	120.90
34	BA	437	G	P-O3'-C3'	6.87	127.95	119.70
34	BA	1691	G	C5-C6-O6	-6.87	124.48	128.60
38	BE	89	G	C5'-C4'-C3'	6.87	127.00	116.00
39	BF	52	A	P-O3'-C3'	-6.87	111.45	119.70
39	BF	55	A	C5-C6-N6	-6.87	118.20	123.70
40	BG	9	G	N3-C2-N2	-6.87	115.09	119.90
85	AA	773	G	C3'-C2'-C1'	-6.87	96.00	101.50
85	AA	1955	U	C5'-C4'-C3'	-6.87	105.00	116.00
34	BA	371	U	O4'-C1'-N1	6.87	113.70	108.20
35	BB	1014	U	O4'-C1'-N1	6.87	113.70	108.20
34	BA	306	G	C5-C6-O6	6.87	132.72	128.60
34	BA	1011	G	O5'-C5'-C4'	-6.87	98.65	111.70
34	BA	1505	G	P-O5'-C5'	-6.87	109.91	120.90
35	BB	128	C	N3-C2-O2	-6.87	117.09	121.90
35	BB	135	C	C6-N1-C1'	6.87	129.04	120.80
35	BB	703	U	O5'-C5'-C4'	-6.87	98.65	111.70
35	BB	801	G	O3'-P-O5'	-6.87	90.95	104.00
35	BB	1474	A	P-O3'-C3'	6.87	127.94	119.70
38	BE	149	A	C5-C6-N1	-6.87	114.27	117.70
39	BF	43	U	C1'-O4'-C4'	-6.87	104.41	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	12	U	N3-C4-O4	6.87	124.21	119.40
41	BH	103	C	O4'-C4'-C3'	-6.87	97.13	104.00
53	BT	121	ARG	NE-CZ-NH2	-6.87	116.86	120.30
85	AA	552	C	N3-C4-N4	6.87	122.81	118.00
85	AA	1103	A	P-O5'-C5'	6.87	131.89	120.90
85	AA	1483	A	O4'-C1'-N9	6.87	113.69	108.20
85	AA	2131	C	O4'-C1'-N1	6.87	113.69	108.20
34	BA	8	G	P-O3'-C3'	-6.87	111.46	119.70
34	BA	165	C	C6-N1-C2	-6.87	117.55	120.30
34	BA	900	A	P-O5'-C5'	-6.87	109.91	120.90
35	BB	561	C	C4'-C3'-C2'	-6.87	95.73	102.60
35	BB	1346	A	P-O5'-C5'	6.87	131.89	120.90
36	BC	33	U	C5-C4-O4	-6.87	121.78	125.90
40	BG	128	U	C5-C6-N1	-6.87	119.27	122.70
85	AA	320	U	O4'-C1'-N1	6.87	113.69	108.20
85	AA	943	U	C2-N3-C4	-6.87	122.88	127.00
85	AA	1136	A	C5'-C4'-C3'	6.87	126.99	116.00
22	AO	95	ARG	NE-CZ-NH2	-6.87	116.87	120.30
34	BA	94	G	P-O3'-C3'	-6.87	111.46	119.70
35	BB	662	G	C8-N9-C4	6.87	109.15	106.40
35	BB	801	G	N9-C1'-C2'	-6.87	104.45	112.00
35	BB	825	U	O4'-C4'-C3'	-6.87	97.13	104.00
36	BC	141	C	C2-N3-C4	-6.87	116.47	119.90
39	BF	11	C	O4'-C4'-C3'	-6.87	97.14	104.00
39	BF	52	A	O4'-C1'-N9	6.87	113.69	108.20
40	BG	112	C	C5'-C4'-O4'	6.87	117.34	109.10
41	BH	84	A	OP1-P-OP2	-6.87	109.30	119.60
61	Bb	88	MET	CG-SD-CE	-6.87	89.22	100.20
69	Bj	10	ARG	NE-CZ-NH2	-6.87	116.87	120.30
85	AA	633	C	C5'-C4'-O4'	6.87	117.34	109.10
85	AA	1876	U	O4'-C1'-N1	6.87	113.69	108.20
34	BA	583	G	C5'-C4'-C3'	-6.86	105.02	116.00
34	BA	721	A	O3'-P-O5'	-6.86	90.96	104.00
34	BA	1256	A	C3'-C2'-C1'	-6.86	96.01	101.50
34	BA	1280	A	C4'-C3'-C2'	6.86	109.46	102.60
35	BB	676	G	C6-N1-C2	-6.86	120.98	125.10
36	BC	75	G	P-O5'-C5'	6.86	131.88	120.90
40	BG	38	A	C4-N9-C1'	-6.86	113.95	126.30
85	AA	436	G	C8-N9-C4	6.86	109.14	106.40
85	AA	1921	G	C5-N7-C8	6.86	107.73	104.30
34	BA	12	G	C4-C5-C6	-6.86	114.68	118.80
35	BB	1071	G	C5-C6-N1	6.86	114.93	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	716	G	C4-N9-C1'	-6.86	117.58	126.50
34	BA	297	A	C5'-C4'-C3'	-6.86	105.02	116.00
35	BB	1443	C	P-O3'-C3'	-6.86	111.47	119.70
36	BC	6	G	C6-N1-C2	-6.86	120.98	125.10
37	BD	31	U	P-O3'-C3'	-6.86	111.47	119.70
85	AA	1004	G	C4-N9-C1'	-6.86	117.58	126.50
85	AA	1855	U	C2'-C3'-O3'	6.86	124.68	113.70
85	AA	2150	G	C6-N1-C2	-6.86	120.98	125.10
85	AA	2230	U	O3'-P-O5'	6.86	117.03	104.00
34	BA	1085	G	N3-C4-C5	-6.86	125.17	128.60
45	BL	36	ARG	NE-CZ-NH2	-6.86	116.87	120.30
85	AA	249	C	C5'-C4'-C3'	-6.86	105.03	116.00
85	AA	865	G	C1'-O4'-C4'	-6.86	104.41	109.90
85	AA	1676	G	C4-N9-C1'	-6.86	117.58	126.50
34	BA	650	C	C4'-C3'-C2'	-6.86	95.74	102.60
34	BA	685	C	C2-N1-C1'	6.86	126.34	118.80
34	BA	1043	C	O4'-C1'-N1	6.86	113.69	108.20
34	BA	1442	A	C5-N7-C8	6.86	107.33	103.90
34	BA	1800	G	C5'-C4'-O4'	6.86	117.33	109.10
35	BB	352	C	O4'-C1'-N1	6.86	113.69	108.20
59	BZ	13	ARG	NE-CZ-NH1	6.86	123.73	120.30
62	Bc	18	PHE	N-CA-CB	-6.86	98.25	110.60
85	AA	245	A	O4'-C1'-N9	6.86	113.69	108.20
85	AA	1339	C	C4'-C3'-C2'	-6.86	95.74	102.60
34	BA	919	A	O5'-C5'-C4'	-6.86	98.67	111.70
34	BA	1676	A	O4'-C1'-N9	6.86	113.68	108.20
35	BB	280	C	O4'-C1'-N1	6.86	113.68	108.20
35	BB	731	U	O4'-C1'-N1	6.86	113.68	108.20
36	BC	124	A	N3-C4-C5	-6.86	122.00	126.80
85	AA	710	A	C8-N9-C1'	-6.86	115.36	127.70
85	AA	1118	U	O4'-C1'-N1	6.86	113.68	108.20
85	AA	1816	C	P-O3'-C3'	6.86	127.93	119.70
34	BA	53	G	C5-C6-O6	-6.85	124.49	128.60
34	BA	1718	C	C5-C4-N4	-6.85	115.40	120.20
35	BB	14	C	O4'-C1'-N1	6.85	113.68	108.20
50	BQ	90	ARG	NE-CZ-NH1	6.85	123.73	120.30
2	A1	66	LEU	N-CA-C	6.85	129.50	111.00
34	BA	689	C	N3-C2-O2	-6.85	117.10	121.90
34	BA	894	G	N3-C2-N2	6.85	124.70	119.90
34	BA	934	G	C6-N1-C2	-6.85	120.99	125.10
34	BA	1797	A	C1'-O4'-C4'	-6.85	104.42	109.90
35	BB	311	C	O4'-C1'-N1	6.85	113.68	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	831	C	O3'-P-O5'	-6.85	90.98	104.00
35	BB	1514	G	C8-N9-C4	-6.85	103.66	106.40
39	BF	68	C	P-O3'-C3'	-6.85	111.48	119.70
61	Bb	131	ASP	N-CA-CB	-6.85	98.27	110.60
82	Bw	124	ARG	NE-CZ-NH2	6.85	123.73	120.30
85	AA	611	G	C2-N3-C4	6.85	115.33	111.90
85	AA	945	A	C5'-C4'-O4'	6.85	117.32	109.10
85	AA	1373	U	P-O3'-C3'	6.85	127.92	119.70
40	BG	10	U	N1-C2-N3	6.85	119.01	114.90
41	BH	5	G	P-O3'-C3'	-6.85	111.48	119.70
85	AA	182	C	C5'-C4'-O4'	6.85	117.32	109.10
85	AA	849	A	P-O3'-C3'	-6.85	111.48	119.70
34	BA	470	C	C5-C4-N4	6.85	125.00	120.20
34	BA	554	A	P-O3'-C3'	6.85	127.92	119.70
34	BA	584	A	N1-C6-N6	-6.85	114.49	118.60
34	BA	920	U	C2-N3-C4	-6.85	122.89	127.00
34	BA	1325	G	C6-N1-C2	-6.85	120.99	125.10
34	BA	1471	U	O4'-C1'-N1	6.85	113.68	108.20
34	BA	1611	A	C8-N9-C1'	6.85	140.03	127.70
34	BA	1842	U	C1'-O4'-C4'	-6.85	104.42	109.90
35	BB	440	U	O4'-C1'-N1	6.85	113.68	108.20
35	BB	642	G	C8-N9-C1'	-6.85	118.10	127.00
38	BE	49	A	P-O5'-C5'	-6.85	109.94	120.90
38	BE	74	U	O4'-C1'-N1	6.85	113.68	108.20
85	AA	44	C	N3-C4-N4	6.85	122.80	118.00
85	AA	577	U	P-O5'-C5'	6.85	131.86	120.90
85	AA	2009	A	C3'-C2'-C1'	-6.85	96.02	101.50
85	AA	2075	C	P-O3'-C3'	-6.85	111.48	119.70
34	BA	189	G	C8-N9-C4	-6.85	103.66	106.40
34	BA	926	A	C8-N9-C4	6.85	108.54	105.80
35	BB	367	C	C5-C6-N1	6.85	124.42	121.00
35	BB	974	C	C2-N1-C1'	6.85	126.33	118.80
38	BE	127	G	P-O3'-C3'	6.85	127.92	119.70
65	Bf	240	ASN	N-CA-CB	6.85	122.93	110.60
85	AA	725	G	O4'-C1'-N9	6.85	113.68	108.20
85	AA	905	C	C5'-C4'-O4'	6.85	117.32	109.10
2	A1	100	TYR	CB-CA-C	-6.85	96.71	110.40
35	BB	1001	G	C4'-C3'-C2'	-6.85	95.75	102.60
41	BH	16	A	C5-N7-C8	-6.85	100.48	103.90
41	BH	33	G	C8-N9-C1'	6.85	135.90	127.00
3	A2	27	TYR	CB-CG-CD1	6.84	125.11	121.00
34	BA	155	U	P-O3'-C3'	-6.84	111.49	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	214	A	O4'-C1'-N9	6.84	113.67	108.20
34	BA	790	G	P-O5'-C5'	-6.84	109.95	120.90
35	BB	569	G	N3-C2-N2	6.84	124.69	119.90
35	BB	601	U	C6-N1-C2	-6.84	116.89	121.00
35	BB	710	A	N1-C6-N6	-6.84	114.49	118.60
35	BB	970	C	C1'-O4'-C4'	-6.84	104.42	109.90
37	BD	77	A	C1'-O4'-C4'	-6.84	104.42	109.90
40	BG	164	U	P-O5'-C5'	6.84	131.85	120.90
40	BG	166	C	C6-N1-C1'	6.84	129.01	120.80
48	BO	148	TYR	CB-CG-CD1	6.84	125.11	121.00
85	AA	54	C	C1'-O4'-C4'	-6.84	104.42	109.90
85	AA	517	A	N9-C1'-C2'	-6.84	104.47	112.00
85	AA	822	U	C6-N1-C2	-6.84	116.89	121.00
85	AA	923	A	O4'-C4'-C3'	-6.84	97.16	104.00
85	AA	1135	U	C1'-O4'-C4'	-6.84	104.42	109.90
85	AA	1814	U	O4'-C1'-N1	6.84	113.67	108.20
86	AB	11	C	C6-N1-C2	-6.84	117.56	120.30
34	BA	192	G	N3-C2-N2	6.84	124.69	119.90
34	BA	1527	G	P-O5'-C5'	-6.84	109.95	120.90
35	BB	1038	G	N1-C6-O6	6.84	124.01	119.90
85	AA	1505	G	C5-C6-N1	6.84	114.92	111.50
8	A7	36	ARG	NE-CZ-NH1	6.84	123.72	120.30
34	BA	124	G	C5-C6-O6	-6.84	124.50	128.60
34	BA	465	A	P-O5'-C5'	-6.84	109.95	120.90
34	BA	785	G	C1'-O4'-C4'	-6.84	104.43	109.90
34	BA	897	U	C3'-C2'-C1'	-6.84	96.03	101.50
34	BA	1011	G	C6-N1-C2	-6.84	121.00	125.10
35	BB	545	C	N1-C1'-C2'	6.84	122.89	114.00
35	BB	623	A	C8-N9-C1'	6.84	140.01	127.70
85	AA	718	C	O4'-C1'-N1	6.84	113.67	108.20
34	BA	217	C	C2-N1-C1'	6.84	126.32	118.80
34	BA	696	A	C5-C6-N6	-6.84	118.23	123.70
34	BA	1325	G	C3'-C2'-C1'	-6.84	96.03	101.50
34	BA	1503	U	C2-N1-C1'	-6.84	109.49	117.70
35	BB	1064	U	C6-N1-C1'	-6.84	111.63	121.20
36	BC	3	C	O5'-C5'-C4'	-6.84	98.70	111.70
77	Br	158	ARG	NE-CZ-NH2	-6.84	116.88	120.30
85	AA	1014	U	C5'-C4'-O4'	6.84	117.31	109.10
85	AA	1105	G	C8-N9-C1'	-6.84	118.11	127.00
85	AA	1437	G	C5'-C4'-C3'	6.84	126.94	116.00
85	AA	1932	C	C2-N1-C1'	6.84	126.32	118.80
85	AA	2148	C	C1'-O4'-C4'	-6.84	104.43	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	24	C	O4'-C1'-N1	6.84	113.67	108.20
35	BB	684	U	N1-C2-N3	6.84	119.00	114.90
37	BD	77	A	N1-C2-N3	-6.84	125.88	129.30
85	AA	156	G	N3-C2-N2	6.84	124.69	119.90
85	AA	557	G	C8-N9-C1'	6.84	135.89	127.00
85	AA	915	G	N1-C6-O6	6.84	124.00	119.90
85	AA	1427	A	C5'-C4'-C3'	6.84	126.94	116.00
85	AA	1499	G	C5-C6-O6	-6.84	124.50	128.60
85	AA	1681	G	C5-C6-O6	-6.84	124.50	128.60
85	AA	2162	G	O4'-C1'-N9	6.84	113.67	108.20
34	BA	27	G	C3'-C2'-C1'	-6.84	96.03	101.50
34	BA	160	G	C5'-C4'-O4'	6.84	117.30	109.10
34	BA	680	C	O4'-C4'-C3'	-6.84	97.16	104.00
34	BA	1476	G	C1'-O4'-C4'	-6.84	104.43	109.90
34	BA	1565	U	O4'-C4'-C3'	-6.84	97.16	104.00
38	BE	163	A	C6-N1-C2	6.84	122.70	118.60
40	BG	34	A	N7-C8-N9	-6.84	110.38	113.80
77	Br	289	ARG	NE-CZ-NH1	6.84	123.72	120.30
85	AA	336	C	C2-N3-C4	-6.84	116.48	119.90
85	AA	1203	G	N1-C6-O6	6.84	124.00	119.90
34	BA	1256	A	C1'-O4'-C4'	-6.83	104.43	109.90
35	BB	834	U	O5'-C5'-C4'	6.83	124.69	111.70
35	BB	1370	G	C5-C6-O6	6.83	132.70	128.60
36	BC	17	U	C4-C5-C6	-6.83	115.60	119.70
37	BD	78	C	O3'-P-O5'	-6.83	91.01	104.00
85	AA	58	C	O4'-C1'-N1	6.83	113.67	108.20
85	AA	156	G	C5'-C4'-C3'	-6.83	105.06	116.00
85	AA	534	A	C4-C5-C6	-6.83	113.58	117.00
85	AA	769	C	C2-N3-C4	-6.83	116.48	119.90
85	AA	830	A	N9-C1'-C2'	-6.83	104.48	112.00
85	AA	1295	G	P-O3'-C3'	-6.83	111.50	119.70
34	BA	267	G	C6-N1-C2	-6.83	121.00	125.10
34	BA	291	C	P-O5'-C5'	-6.83	109.97	120.90
34	BA	557	U	C3'-C2'-O2'	6.83	133.12	113.30
34	BA	1719	G	P-O3'-C3'	-6.83	111.50	119.70
35	BB	15	C	N3-C2-O2	-6.83	117.12	121.90
35	BB	661	G	C8-N9-C1'	6.83	135.88	127.00
35	BB	1304	U	C5'-C4'-C3'	-6.83	105.07	116.00
42	BI	29	LEU	CB-CA-C	-6.83	97.22	110.20
85	AA	167	A	C3'-C2'-C1'	-6.83	96.03	101.50
85	AA	625	G	C5-C6-O6	-6.83	124.50	128.60
85	AA	799	G	N1-C6-O6	6.83	124.00	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	168	U	C2-N1-C1'	-6.83	109.50	117.70
34	BA	349	G	C5'-C4'-C3'	6.83	126.93	116.00
34	BA	600	G	O4'-C1'-N9	6.83	113.67	108.20
34	BA	1599	A	C1'-O4'-C4'	-6.83	104.44	109.90
35	BB	259	U	O4'-C1'-N1	6.83	113.67	108.20
35	BB	261	C	O3'-P-O5'	-6.83	91.02	104.00
38	BE	129	G	C6-C5-N7	-6.83	126.30	130.40
41	BH	76	G	O4'-C1'-N9	6.83	113.67	108.20
41	BH	84	A	C2-N3-C4	6.83	114.02	110.60
85	AA	1211	C	O3'-P-O5'	-6.83	91.02	104.00
85	AA	1265	C	C6-N1-C2	-6.83	117.57	120.30
85	AA	1809	G	O4'-C1'-N9	6.83	113.67	108.20
2	A1	96	PHE	N-CA-CB	6.83	122.89	110.60
6	A5	78	ILE	CB-CA-C	-6.83	97.94	111.60
34	BA	1202	G	C5-C6-N1	6.83	114.92	111.50
34	BA	1540	C	C5'-C4'-C3'	-6.83	105.07	116.00
35	BB	972	C	O4'-C1'-N1	6.83	113.66	108.20
85	AA	160	A	P-O3'-C3'	-6.83	111.50	119.70
85	AA	463	G	C5'-C4'-C3'	-6.83	105.07	116.00
34	BA	278	U	C5'-C4'-O4'	6.83	117.29	109.10
35	BB	1082	A	O3'-P-O5'	6.83	116.97	104.00
85	AA	492	C	C6-N1-C1'	6.83	128.99	120.80
2	A1	186	ARG	NE-CZ-NH2	-6.83	116.89	120.30
34	BA	610	A	O4'-C4'-C3'	-6.83	97.17	104.00
35	BB	844	G	C4-N9-C1'	-6.83	117.62	126.50
35	BB	1120	A	C1'-O4'-C4'	-6.83	104.44	109.90
77	Br	313	TYR	CB-CG-CD1	6.83	125.10	121.00
85	AA	8	U	N3-C4-O4	6.83	124.18	119.40
35	BB	70	A	P-O3'-C3'	-6.83	111.51	119.70
35	BB	1326	U	C4'-C3'-C2'	6.83	109.42	102.60
36	BC	45	C	O4'-C1'-N1	6.83	113.66	108.20
85	AA	172	A	N9-C1'-C2'	-6.83	104.49	112.00
85	AA	1163	G	C8-N9-C1'	6.83	135.87	127.00
85	AA	2044	A	C5'-C4'-O4'	6.83	117.29	109.10
85	AA	2134	U	P-O3'-C3'	-6.83	111.51	119.70
85	AA	2150	G	C3'-C2'-C1'	-6.83	96.04	101.50
22	AO	79	ARG	NE-CZ-NH2	-6.82	116.89	120.30
27	AT	111	ARG	NE-CZ-NH1	-6.82	116.89	120.30
34	BA	438	A	C5-C6-N6	6.82	129.16	123.70
35	BB	1485	G	C8-N9-C4	-6.82	103.67	106.40
41	BH	29	G	P-O3'-C3'	-6.82	111.51	119.70
41	BH	68	G	O4'-C1'-N9	6.82	113.66	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	100	A	O5'-P-OP1	6.82	118.89	110.70
85	AA	330	C	C4'-C3'-C2'	-6.82	95.78	102.60
85	AA	334	A	C1'-O4'-C4'	-6.82	104.44	109.90
85	AA	605	A	P-O3'-C3'	-6.82	111.51	119.70
85	AA	928	U	C5'-C4'-C3'	6.82	126.92	116.00
35	BB	1348	C	O4'-C1'-N1	6.82	113.66	108.20
85	AA	794	A	C2-N3-C4	-6.82	107.19	110.60
85	AA	898	A	P-O3'-C3'	6.82	127.89	119.70
34	BA	816	G	C5'-C4'-C3'	6.82	126.91	116.00
34	BA	1567	G	C5'-C4'-O4'	-6.82	100.92	109.10
35	BB	23	U	O4'-C1'-N1	6.82	113.66	108.20
35	BB	1262	A	C5-C6-N6	-6.82	118.24	123.70
85	AA	86	G	C4-N9-C1'	-6.82	117.63	126.50
85	AA	1125	G	O4'-C1'-N9	6.82	113.66	108.20
85	AA	1174	G	N1-C6-O6	6.82	123.99	119.90
85	AA	1213	U	P-O3'-C3'	6.82	127.88	119.70
85	AA	1258	U	O4'-C1'-N1	6.82	113.66	108.20
85	AA	1361	A	O3'-P-O5'	-6.82	91.04	104.00
85	AA	2196	G	C5-C6-O6	6.82	132.69	128.60
85	AA	2245	A	P-O5'-C5'	6.82	131.81	120.90
34	BA	487	A	C3'-C2'-C1'	-6.82	96.05	101.50
34	BA	520	G	C6-C5-N7	-6.82	126.31	130.40
34	BA	701	G	C5-C6-O6	-6.82	124.51	128.60
36	BC	147	G	N3-C2-N2	6.82	124.67	119.90
85	AA	2030	U	O3'-P-O5'	-6.82	91.04	104.00
85	AA	2033	C	C1'-O4'-C4'	-6.82	104.44	109.90
34	BA	231	U	C6-N1-C2	-6.82	116.91	121.00
34	BA	316	G	C8-N9-C1'	6.82	135.86	127.00
34	BA	1004	U	C5'-C4'-C3'	6.82	126.91	116.00
34	BA	1017	C	N1-C2-O2	6.82	122.99	118.90
34	BA	1101	A	C5'-C4'-O4'	6.82	117.28	109.10
34	BA	1432	C	C2-N3-C4	-6.82	116.49	119.90
35	BB	5	A	C2-N3-C4	-6.82	107.19	110.60
35	BB	1489	A	O3'-P-O5'	6.82	116.95	104.00
36	BC	108	A	C5'-C4'-C3'	-6.82	105.09	116.00
44	BK	119	TYR	CB-CG-CD2	-6.82	116.91	121.00
85	AA	119	G	C5'-C4'-C3'	6.82	126.91	116.00
85	AA	763	U	C2-N1-C1'	-6.82	109.52	117.70
85	AA	1242	A	C8-N9-C4	-6.82	103.07	105.80
85	AA	1644	G	N3-C4-C5	-6.82	125.19	128.60
1	A0	168	MET	CG-SD-CE	-6.82	89.30	100.20
2	A1	38	CYS	CA-CB-SG	6.82	126.27	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	269	G	C5'-C4'-C3'	-6.82	105.10	116.00
34	BA	652	C	C5'-C4'-C3'	-6.82	105.10	116.00
34	BA	1038	U	N1-C1'-C2'	-6.82	104.50	112.00
34	BA	1741	G	N3-C2-N2	6.82	124.67	119.90
35	BB	454	U	C6-N1-C1'	6.82	130.74	121.20
35	BB	490	G	C6-N1-C2	-6.82	121.01	125.10
38	BE	8	G	N1-C2-N2	-6.82	110.07	116.20
54	BU	15	PHE	CB-CG-CD2	-6.82	116.03	120.80
85	AA	210	G	C4-C5-C6	-6.82	114.71	118.80
85	AA	277	G	C5-C6-O6	-6.82	124.51	128.60
85	AA	977	U	C1'-O4'-C4'	-6.82	104.45	109.90
85	AA	1451	U	C2-N1-C1'	6.82	125.88	117.70
85	AA	2163	G	C8-N9-C4	-6.82	103.67	106.40
85	AA	2199	G	C3'-C2'-C1'	-6.82	96.05	101.50
34	BA	195	G	C4'-C3'-C2'	6.81	109.41	102.60
85	AA	1197	U	N3-C4-O4	-6.81	114.63	119.40
85	AA	1879	U	C4'-C3'-C2'	6.81	109.41	102.60
85	AA	1984	A	C5-C6-N6	-6.81	118.25	123.70
86	AB	2	C	P-O5'-C5'	6.81	131.80	120.90
34	BA	430	A	O4'-C1'-N9	6.81	113.65	108.20
35	BB	888	U	P-O5'-C5'	6.81	131.80	120.90
35	BB	1441	C	C5'-C4'-C3'	-6.81	105.10	116.00
61	Bb	122	LYS	N-CA-C	-6.81	92.61	111.00
85	AA	24	U	C2-N1-C1'	-6.81	109.53	117.70
85	AA	241	U	O4'-C1'-N1	6.81	113.65	108.20
85	AA	902	A	O4'-C1'-C2'	-6.81	98.99	105.80
85	AA	1438	C	C6-N1-C2	-6.81	117.58	120.30
85	AA	2034	G	C6-C5-N7	-6.81	126.31	130.40
85	AA	2200	A	P-O5'-C5'	-6.81	110.00	120.90
34	BA	144	C	C6-N1-C1'	-6.81	112.63	120.80
37	BD	73	U	C2'-C3'-O3'	6.81	124.60	113.70
38	BE	165	U	C2-N3-C4	-6.81	122.91	127.00
85	AA	620	U	P-O5'-C5'	-6.81	110.00	120.90
85	AA	2249	U	O4'-C1'-C2'	-6.81	98.99	105.80
34	BA	112	C	N1-C2-O2	6.81	122.99	118.90
34	BA	330	A	C5-C6-N6	-6.81	118.25	123.70
34	BA	911	G	C8-N9-C1'	6.81	135.85	127.00
34	BA	1008	A	N1-C6-N6	6.81	122.69	118.60
34	BA	1172	C	C2-N1-C1'	-6.81	111.31	118.80
34	BA	1487	U	C2-N1-C1'	-6.81	109.53	117.70
34	BA	1707	C	C6-N1-C2	-6.81	117.58	120.30
34	BA	1737	A	C3'-C2'-C1'	-6.81	96.05	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	380	G	C6-N1-C2	-6.81	121.01	125.10
35	BB	694	C	C4'-C3'-C2'	6.81	109.41	102.60
35	BB	1135	U	C5'-C4'-O4'	6.81	117.27	109.10
35	BB	1479	C	C1'-O4'-C4'	-6.81	104.45	109.90
41	BH	63	G	N7-C8-N9	6.81	116.50	113.10
85	AA	238	C	C2-N1-C1'	6.81	126.29	118.80
85	AA	869	A	C5'-C4'-O4'	6.81	117.27	109.10
85	AA	1540	A	N1-C6-N6	6.81	122.69	118.60
34	BA	21	C	O5'-C5'-C4'	-6.81	98.77	111.70
34	BA	80	U	C5-C6-N1	-6.81	119.30	122.70
34	BA	384	U	N3-C2-O2	-6.81	117.44	122.20
34	BA	943	G	C5-C6-O6	-6.81	124.52	128.60
34	BA	1287	G	C6-N1-C2	-6.81	121.02	125.10
35	BB	1144	A	C6-N1-C2	-6.81	114.52	118.60
35	BB	1282	G	O4'-C1'-N9	6.81	113.65	108.20
38	BE	123	A	N9-C4-C5	-6.81	103.08	105.80
39	BF	56	C	N3-C4-C5	-6.81	119.18	121.90
85	AA	372	U	O4'-C1'-N1	6.81	113.64	108.20
85	AA	553	G	C1'-O4'-C4'	-6.81	104.45	109.90
85	AA	596	A	N1-C6-N6	6.81	122.68	118.60
85	AA	891	G	C8-N9-C1'	-6.81	118.15	127.00
34	BA	447	U	O4'-C1'-N1	6.81	113.64	108.20
34	BA	1692	U	P-O3'-C3'	-6.81	111.53	119.70
38	BE	10	G	N9-C1'-C2'	6.81	122.85	114.00
85	AA	869	A	N1-C2-N3	6.81	132.70	129.30
34	BA	411	C	C2-N1-C1'	6.80	126.29	118.80
34	BA	624	G	C5'-C4'-O4'	6.80	117.27	109.10
34	BA	766	A	C1'-O4'-C4'	-6.80	104.46	109.90
34	BA	1294	C	N3-C4-C5	-6.80	119.18	121.90
34	BA	1670	A	C5-C6-N6	6.80	129.14	123.70
35	BB	27	C	C3'-C2'-C1'	-6.80	96.06	101.50
35	BB	369	A	C5'-C4'-O4'	6.80	117.27	109.10
35	BB	414	C	O4'-C1'-N1	6.80	113.64	108.20
35	BB	1403	G	O4'-C1'-N9	6.80	113.64	108.20
39	BF	17	U	C5-C6-N1	-6.80	119.30	122.70
40	BG	102	G	N3-C4-N9	6.80	130.08	126.00
85	AA	750	A	C4'-C3'-C2'	-6.80	95.80	102.60
85	AA	809	A	N1-C6-N6	6.80	122.68	118.60
34	BA	1738	G	C3'-C2'-C1'	-6.80	96.06	101.50
38	BE	136	G	C1'-O4'-C4'	-6.80	104.46	109.90
4	A3	214	ARG	NE-CZ-NH1	6.80	123.70	120.30
34	BA	76	U	O4'-C1'-N1	6.80	113.64	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	154	A	P-O5'-C5'	6.80	131.78	120.90
34	BA	764	G	O3'-P-O5'	6.80	116.92	104.00
34	BA	1041	U	N1-C1'-C2'	-6.80	104.52	112.00
34	BA	1189	A	C8-N9-C4	-6.80	103.08	105.80
34	BA	1440	C	C4'-C3'-C2'	-6.80	95.80	102.60
34	BA	1551	G	N9-C1'-C2'	-6.80	104.52	112.00
35	BB	406	A	C3'-C2'-C1'	-6.80	96.06	101.50
35	BB	1030	U	P-O3'-C3'	6.80	127.86	119.70
40	BG	40	G	C1'-O4'-C4'	-6.80	104.46	109.90
40	BG	156	G	C5'-C4'-O4'	6.80	117.26	109.10
85	AA	1227	A	C5'-C4'-C3'	-6.80	105.12	116.00
86	AB	65	G	C4-N9-C1'	6.80	135.34	126.50
34	BA	611	A	C3'-C2'-C1'	-6.80	96.06	101.50
35	BB	893	U	O3'-P-O5'	6.80	116.92	104.00
35	BB	1141	A	P-O3'-C3'	-6.80	111.54	119.70
35	BB	1455	A	N1-C6-N6	6.80	122.68	118.60
37	BD	102	C	C3'-C2'-C1'	-6.80	96.06	101.50
40	BG	65	C	C2-N1-C1'	-6.80	111.32	118.80
85	AA	54	C	C6-N1-C1'	6.80	128.96	120.80
85	AA	1678	U	C3'-C2'-C1'	-6.80	96.06	101.50
85	AA	2099	C	P-O5'-C5'	6.80	131.78	120.90
85	AA	2186	U	C4'-C3'-C2'	-6.80	95.80	102.60
85	AA	2249	U	C5'-C4'-O4'	6.80	117.26	109.10
34	BA	248	G	C6-C5-N7	-6.80	126.32	130.40
35	BB	1317	U	O4'-C1'-N1	6.80	113.64	108.20
38	BE	111	C	P-O5'-C5'	-6.80	110.02	120.90
38	BE	176	G	C4'-C3'-C2'	6.80	109.40	102.60
39	BF	12	U	N1-C1'-C2'	-6.80	104.52	112.00
34	BA	272	A	C3'-C2'-C1'	-6.80	96.06	101.50
34	BA	594	G	C1'-O4'-C4'	-6.80	104.46	109.90
35	BB	425	G	C3'-C2'-C1'	-6.80	96.06	101.50
85	AA	467	U	N1-C2-O2	6.80	127.56	122.80
85	AA	626	G	O3'-P-O5'	-6.80	91.09	104.00
85	AA	877	G	N9-C1'-C2'	-6.80	104.52	112.00
85	AA	1126	G	O4'-C1'-N9	6.80	113.64	108.20
85	AA	1712	A	O4'-C1'-C2'	6.80	113.72	107.60
85	AA	2211	G	C3'-C2'-C1'	-6.80	96.06	101.50
24	AQ	48	ARG	NE-CZ-NH2	-6.79	116.90	120.30
35	BB	941	C	O4'-C1'-N1	6.79	113.64	108.20
35	BB	971	A	C2'-C3'-O3'	6.79	124.57	113.70
38	BE	46	G	O5'-P-OP1	-6.79	99.58	105.70
85	AA	1174	G	C4-N9-C1'	-6.79	117.67	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	445	C	C1'-O4'-C4'	-6.79	104.47	109.90
34	BA	687	G	C5-C6-O6	6.79	132.68	128.60
34	BA	926	A	C4'-C3'-C2'	6.79	109.39	102.60
35	BB	41	A	C5-C6-N1	6.79	121.10	117.70
35	BB	133	G	C5-C6-O6	-6.79	124.52	128.60
35	BB	1205	A	C5'-C4'-C3'	-6.79	105.13	116.00
41	BH	91	G	O4'-C1'-N9	6.79	113.64	108.20
42	BI	111	ARG	NE-CZ-NH2	6.79	123.70	120.30
77	Br	109	ARG	NE-CZ-NH1	6.79	123.70	120.30
80	Bu	253	HIS	CA-CB-CG	-6.79	102.05	113.60
85	AA	31	C	P-O3'-C3'	-6.79	111.55	119.70
85	AA	117	C	O4'-C1'-N1	6.79	113.64	108.20
85	AA	461	G	C4-C5-N7	6.79	113.52	110.80
85	AA	958	C	P-O5'-C5'	-6.79	110.03	120.90
85	AA	1373	U	C1'-O4'-C4'	-6.79	104.47	109.90
24	AQ	65	THR	N-CA-CB	-6.79	97.39	110.30
34	BA	544	U	O4'-C1'-N1	6.79	113.63	108.20
34	BA	684	G	N1-C2-N2	-6.79	110.09	116.20
34	BA	1275	G	C5'-C4'-C3'	-6.79	105.14	116.00
34	BA	1553	G	C4'-C3'-C2'	-6.79	95.81	102.60
34	BA	1787	U	C6-N1-C2	-6.79	116.93	121.00
37	BD	14	C	C2-N1-C1'	-6.79	111.33	118.80
61	Bb	32	ARG	NE-CZ-NH2	-6.79	116.90	120.30
85	AA	537	G	C8-N9-C4	6.79	109.12	106.40
85	AA	555	C	O4'-C1'-N1	6.79	113.63	108.20
85	AA	1471	G	N9-C4-C5	-6.79	102.68	105.40
35	BB	990	G	P-O3'-C3'	6.79	127.85	119.70
36	BC	26	U	N1-C2-O2	-6.79	118.05	122.80
85	AA	975	G	N3-C2-N2	6.79	124.65	119.90
34	BA	21	C	P-O3'-C3'	6.79	127.85	119.70
34	BA	1329	U	P-O3'-C3'	-6.79	111.55	119.70
34	BA	1711	G	C5-C6-N1	6.79	114.89	111.50
34	BA	1719	G	C5-C6-O6	6.79	132.67	128.60
35	BB	1537	C	O4'-C1'-N1	6.79	113.63	108.20
85	AA	523	U	C6-N1-C2	-6.79	116.93	121.00
85	AA	525	C	P-O3'-C3'	-6.79	111.55	119.70
11	AC	201	ASN	CA-CB-CG	-6.79	98.47	113.40
34	BA	906	A	C8-N9-C1'	-6.79	115.48	127.70
41	BH	123	G	O4'-C1'-N9	6.79	113.63	108.20
85	AA	338	G	C4'-C3'-C2'	-6.79	95.81	102.60
85	AA	1146	C	C5-C4-N4	6.79	124.95	120.20
34	BA	202	A	N1-C2-N3	-6.79	125.91	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	203	U	C5'-C4'-C3'	-6.79	105.14	116.00
34	BA	436	U	C4'-C3'-C2'	-6.79	95.81	102.60
34	BA	1622	U	C5'-C4'-O4'	6.79	117.24	109.10
35	BB	1066	G	O4'-C1'-N9	6.79	113.63	108.20
35	BB	1539	C	C6-N1-C1'	6.79	128.94	120.80
36	BC	35	C	C6-N1-C2	-6.79	117.59	120.30
38	BE	140	G	C6-N1-C2	-6.79	121.03	125.10
41	BH	114	G	C8-N9-C4	6.79	109.11	106.40
64	Be	227	ARG	NE-CZ-NH2	6.79	123.69	120.30
85	AA	814	G	N3-C2-N2	-6.79	115.15	119.90
85	AA	1277	C	O3'-P-O5'	6.79	116.89	104.00
85	AA	1448	A	C5-C6-N6	-6.79	118.27	123.70
85	AA	1531	G	C8-N9-C1'	6.79	135.82	127.00
85	AA	1692	U	N1-C2-N3	6.79	118.97	114.90
34	BA	806	U	O4'-C4'-C3'	-6.78	97.22	104.00
34	BA	1460	U	C2-N3-C4	-6.78	122.93	127.00
34	BA	1802	C	O4'-C1'-N1	6.78	113.63	108.20
35	BB	439	G	C5-C6-N1	6.78	114.89	111.50
35	BB	768	A	C4-N9-C1'	-6.78	114.09	126.30
35	BB	1441	C	C1'-O4'-C4'	-6.78	104.47	109.90
38	BE	32	U	C1'-O4'-C4'	6.78	115.33	109.90
38	BE	75	C	C2-N1-C1'	-6.78	111.34	118.80
85	AA	735	G	C5-C6-N1	6.78	114.89	111.50
85	AA	1347	C	O4'-C1'-N1	6.78	113.63	108.20
34	BA	506	U	C4'-C3'-C2'	6.78	109.38	102.60
34	BA	551	U	OP1-P-OP2	-6.78	109.43	119.60
38	BE	104	G	C3'-C2'-C1'	6.78	106.93	101.50
85	AA	327	G	C4'-C3'-C2'	-6.78	95.82	102.60
85	AA	1408	U	O4'-C1'-N1	6.78	113.62	108.20
85	AA	1843	A	P-O5'-C5'	-6.78	110.05	120.90
85	AA	1921	G	C8-N9-C4	-6.78	103.69	106.40
3	A2	174	TYR	CB-CG-CD1	6.78	125.07	121.00
34	BA	617	G	O4'-C1'-N9	-6.78	102.78	108.20
34	BA	1739	G	C3'-C2'-C1'	-6.78	96.08	101.50
35	BB	569	G	N1-C2-N2	-6.78	110.10	116.20
36	BC	6	G	C8-N9-C4	6.78	109.11	106.40
36	BC	90	U	C3'-C2'-C1'	-6.78	96.08	101.50
38	BE	96	G	C1'-O4'-C4'	-6.78	104.48	109.90
41	BH	25	A	O3'-P-O5'	-6.78	91.12	104.00
56	BW	134	HIS	CA-CB-CG	-6.78	102.07	113.60
85	AA	209	C	C5'-C4'-C3'	6.78	126.85	116.00
85	AA	548	G	C4-N9-C1'	-6.78	117.69	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	601	A	C3'-C2'-C1'	-6.78	96.08	101.50
85	AA	925	G	C3'-C2'-C1'	-6.78	96.08	101.50
85	AA	1699	A	N1-C6-N6	6.78	122.67	118.60
85	AA	1995	U	C5-C4-O4	6.78	129.97	125.90
85	AA	2107	C	O5'-C5'-C4'	-6.78	98.82	111.70
85	AA	2245	A	C2'-C3'-O3'	6.78	124.55	113.70
34	BA	222	C	C6-N1-C2	-6.78	117.59	120.30
34	BA	400	A	C5-C6-N6	6.78	129.12	123.70
36	BC	10	C	O5'-P-OP2	6.78	118.83	110.70
40	BG	167	C	C3'-C2'-C1'	-6.78	96.08	101.50
57	BX	87	TYR	CA-CB-CG	6.78	126.28	113.40
85	AA	47	A	C4-N9-C1'	6.78	138.50	126.30
85	AA	767	A	C5'-C4'-C3'	6.78	126.85	116.00
86	AB	23	A	C8-N9-C4	-6.78	103.09	105.80
26	AS	131	TYR	N-CA-CB	-6.78	98.40	110.60
34	BA	395	G	C4-N9-C1'	-6.78	117.69	126.50
34	BA	517	A	N7-C8-N9	-6.78	110.41	113.80
34	BA	1310	C	C6-N1-C2	-6.78	117.59	120.30
34	BA	1526	C	C1'-O4'-C4'	-6.78	104.48	109.90
35	BB	769	C	C6-N1-C2	-6.78	117.59	120.30
53	BT	163	ARG	NE-CZ-NH1	6.78	123.69	120.30
85	AA	188	G	C4-N9-C1'	-6.78	117.69	126.50
85	AA	209	C	C4'-C3'-C2'	-6.78	95.82	102.60
85	AA	1227	A	O4'-C1'-N9	6.78	113.62	108.20
34	BA	274	C	C1'-O4'-C4'	-6.78	104.48	109.90
34	BA	1186	U	C6-N1-C2	-6.78	116.94	121.00
35	BB	857	G	N1-C6-O6	6.78	123.97	119.90
35	BB	893	U	C4'-C3'-C2'	-6.78	95.83	102.60
35	BB	1073	A	C4'-C3'-C2'	6.78	109.38	102.60
35	BB	1517	G	O4'-C1'-N9	6.78	113.62	108.20
47	BN	155	ARG	NE-CZ-NH1	6.78	123.69	120.30
85	AA	1154	A	P-O3'-C3'	6.78	127.83	119.70
85	AA	2092	A	C4'-C3'-C2'	6.78	109.38	102.60
34	BA	366	G	C1'-O4'-C4'	-6.77	104.48	109.90
34	BA	647	U	C1'-O4'-C4'	-6.77	104.48	109.90
35	BB	1105	G	C5-C6-O6	-6.77	124.54	128.60
40	BG	172	C	C5-C4-N4	-6.77	115.46	120.20
41	BH	34	G	C4'-C3'-C2'	-6.77	95.83	102.60
85	AA	1892	G	C8-N9-C1'	6.77	135.81	127.00
34	BA	295	G	P-O3'-C3'	-6.77	111.57	119.70
34	BA	424	U	C5'-C4'-C3'	-6.77	105.16	116.00
34	BA	638	U	N3-C4-O4	-6.77	114.66	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	865	C	N3-C2-O2	-6.77	117.16	121.90
34	BA	1302	C	P-O3'-C3'	-6.77	111.57	119.70
34	BA	1701	U	C2-N3-C4	-6.77	122.94	127.00
34	BA	1836	A	C3'-C2'-C1'	-6.77	96.08	101.50
35	BB	474	G	C4-N9-C1'	-6.77	117.69	126.50
35	BB	483	C	C1'-O4'-C4'	-6.77	104.48	109.90
35	BB	1219	A	C4'-C3'-C2'	-6.77	95.83	102.60
35	BB	1475	U	O5'-P-OP2	-6.77	99.61	105.70
36	BC	60	U	C5'-C4'-C3'	-6.77	105.16	116.00
85	AA	850	U	C4'-C3'-C2'	-6.77	95.83	102.60
6	A5	77	ARG	CB-CA-C	-6.77	96.86	110.40
34	BA	926	A	N7-C8-N9	-6.77	110.41	113.80
40	BG	6	A	N1-C6-N6	6.77	122.66	118.60
85	AA	1348	C	O4'-C1'-N1	6.77	113.62	108.20
86	AB	56	C	P-O5'-C5'	-6.77	110.07	120.90
34	BA	327	G	O4'-C1'-N9	6.77	113.62	108.20
34	BA	569	C	C3'-C2'-C1'	-6.77	96.08	101.50
34	BA	1227	U	C5'-C4'-O4'	-6.77	100.98	109.10
35	BB	429	C	N3-C2-O2	-6.77	117.16	121.90
35	BB	999	G	C4-N9-C1'	-6.77	117.70	126.50
40	BG	123	C	C3'-C2'-C1'	-6.77	96.08	101.50
40	BG	167	C	C4'-C3'-C2'	6.77	109.37	102.60
41	BH	6	U	N1-C2-N3	6.77	118.96	114.90
77	Br	352	ARG	N-CA-CB	-6.77	98.42	110.60
85	AA	552	C	P-O5'-C5'	-6.77	110.07	120.90
85	AA	680	U	N1-C2-O2	6.77	127.54	122.80
85	AA	685	U	N3-C2-O2	-6.77	117.46	122.20
85	AA	2073	U	C2-N1-C1'	-6.77	109.58	117.70
1	A0	203	CYS	CA-CB-SG	-6.77	101.82	114.00
34	BA	1204	U	N3-C2-O2	-6.77	117.46	122.20
34	BA	1654	G	C4-N9-C1'	-6.77	117.70	126.50
35	BB	529	A	C8-N9-C4	-6.77	103.09	105.80
38	BE	136	G	C5'-C4'-C3'	6.77	126.83	116.00
40	BG	80	G	C5-C6-O6	6.77	132.66	128.60
40	BG	115	C	N1-C1'-C2'	-6.77	104.56	112.00
85	AA	774	C	C1'-O4'-C4'	-6.77	104.49	109.90
85	AA	867	G	C5-N7-C8	-6.77	100.92	104.30
85	AA	364	C	P-O5'-C5'	6.77	131.72	120.90
85	AA	1085	U	P-O3'-C3'	6.77	127.82	119.70
85	AA	1176	C	N3-C2-O2	-6.77	117.16	121.90
85	AA	1340	C	C6-N1-C2	-6.77	117.59	120.30
85	AA	2206	A	O4'-C1'-N9	6.77	113.61	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	530	A	C2-N3-C4	-6.76	107.22	110.60
34	BA	541	C	O4'-C1'-N1	6.76	113.61	108.20
34	BA	1579	G	C5-C6-O6	6.76	132.66	128.60
35	BB	1071	G	N3-C4-C5	-6.76	125.22	128.60
41	BH	48	G	C4-N9-C1'	6.76	135.29	126.50
85	AA	57	G	N3-C4-C5	-6.76	125.22	128.60
85	AA	488	G	C6-N1-C2	-6.76	121.04	125.10
85	AA	1259	U	O5'-C5'-C4'	-6.76	98.85	111.70
85	AA	1864	G	N1-C6-O6	6.76	123.96	119.90
34	BA	398	G	C8-N9-C4	6.76	109.11	106.40
34	BA	1196	C	C5-C4-N4	6.76	124.93	120.20
34	BA	1200	U	C2'-C3'-O3'	6.76	124.52	113.70
34	BA	1693	U	O3'-P-O5'	-6.76	91.15	104.00
35	BB	777	C	O4'-C1'-N1	6.76	113.61	108.20
37	BD	48	G	C5-C6-N1	6.76	114.88	111.50
80	Bu	66	HIS	CB-CA-C	-6.76	96.87	110.40
85	AA	258	G	C8-N9-C4	6.76	109.11	106.40
23	AP	120	HIS	CA-CB-CG	-6.76	102.11	113.60
34	BA	177	G	C5-C6-O6	-6.76	124.54	128.60
34	BA	703	U	C3'-C2'-C1'	-6.76	96.09	101.50
39	BF	56	C	C5-C6-N1	-6.76	117.62	121.00
52	BS	44	TRP	CB-CG-CD2	-6.76	117.81	126.60
53	BT	9	ARG	NE-CZ-NH1	6.76	123.68	120.30
85	AA	674	U	C2-N1-C1'	-6.76	109.59	117.70
7	A6	56	ARG	NE-CZ-NH1	6.76	123.68	120.30
34	BA	1043	C	P-O3'-C3'	-6.76	111.59	119.70
34	BA	1085	G	C8-N9-C4	-6.76	103.70	106.40
34	BA	1497	A	N1-C6-N6	-6.76	114.54	118.60
35	BB	441	G	C8-N9-C1'	6.76	135.79	127.00
35	BB	532	C	C6-N1-C1'	6.76	128.91	120.80
36	BC	44	A	C5-C6-N1	6.76	121.08	117.70
36	BC	123	G	C4-N9-C1'	-6.76	117.71	126.50
56	BW	81	ILE	CB-CA-C	6.76	125.12	111.60
85	AA	1647	G	O5'-C5'-C4'	6.76	124.55	111.70
85	AA	1935	G	C5'-C4'-C3'	-6.76	105.18	116.00
5	A4	46	PHE	CB-CG-CD2	-6.76	116.07	120.80
34	BA	587	U	C4'-C3'-O3'	-6.76	95.21	109.40
34	BA	1562	G	C5-C6-N1	6.76	114.88	111.50
36	BC	136	G	P-O3'-C3'	-6.76	111.59	119.70
41	BH	30	C	C2-N1-C1'	6.76	126.23	118.80
68	Bi	22	ARG	NE-CZ-NH1	6.76	123.68	120.30
85	AA	2152	C	C6-N1-C1'	6.76	128.91	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	247	U	O3'-P-O5'	6.76	116.84	104.00
34	BA	517	A	O5'-P-OP2	-6.76	99.62	105.70
34	BA	796	G	C5-N7-C8	-6.76	100.92	104.30
34	BA	813	C	C5'-C4'-C3'	6.76	126.81	116.00
36	BC	62	A	N3-C4-C5	6.76	131.53	126.80
38	BE	105	A	C5-C6-N6	-6.76	118.30	123.70
77	Br	199	ARG	NE-CZ-NH1	6.76	123.68	120.30
85	AA	157	G	C4-N9-C1'	-6.76	117.72	126.50
34	BA	329	G	O4'-C1'-N9	-6.75	102.80	108.20
34	BA	1508	C	P-O3'-C3'	-6.75	111.59	119.70
35	BB	1258	G	C5'-C4'-O4'	6.75	117.21	109.10
38	BE	26	G	OP1-P-OP2	-6.75	109.47	119.60
51	BR	57	CYS	CB-CA-C	-6.75	96.89	110.40
85	AA	898	A	O3'-P-O5'	6.75	116.83	104.00
85	AA	1270	C	O4'-C1'-N1	6.75	113.60	108.20
34	BA	81	C	O4'-C1'-N1	6.75	113.60	108.20
34	BA	198	U	O4'-C1'-C2'	6.75	113.68	107.60
35	BB	822	G	C4-N9-C1'	6.75	135.28	126.50
85	AA	5	U	P-O3'-C3'	-6.75	111.60	119.70
85	AA	1101	C	C6-N1-C2	-6.75	117.60	120.30
85	AA	1264	U	C5'-C4'-C3'	-6.75	105.19	116.00
85	AA	1729	C	N3-C4-N4	6.75	122.73	118.00
85	AA	2087	C	C4'-C3'-C2'	6.75	109.35	102.60
85	AA	2183	U	C6-N1-C2	-6.75	116.95	121.00
85	AA	2204	A	P-O3'-C3'	-6.75	111.60	119.70
34	BA	111	U	O5'-P-OP2	-6.75	99.62	105.70
34	BA	441	A	N1-C6-N6	6.75	122.65	118.60
34	BA	676	G	P-O5'-C5'	-6.75	110.10	120.90
34	BA	802	G	N9-C1'-C2'	-6.75	104.57	112.00
34	BA	1146	U	C4'-C3'-C2'	-6.75	95.85	102.60
34	BA	1287	G	C4'-C3'-C2'	6.75	109.35	102.60
35	BB	1027	U	O4'-C1'-N1	6.75	113.60	108.20
40	BG	20	U	C6-N1-C1'	6.75	130.65	121.20
40	BG	102	G	N1-C6-O6	-6.75	115.85	119.90
85	AA	1109	G	P-O5'-C5'	6.75	131.70	120.90
85	AA	1190	G	N9-C4-C5	-6.75	102.70	105.40
85	AA	1454	U	N1-C2-N3	6.75	118.95	114.90
85	AA	2005	U	C5'-C4'-O4'	6.75	117.20	109.10
85	AA	2120	C	C5'-C4'-C3'	-6.75	105.20	116.00
85	AA	2202	G	C5-C6-N1	6.75	114.88	111.50
34	BA	806	U	O3'-P-O5'	6.75	116.83	104.00
71	Bl	95	TRP	O-C-N	-6.75	111.90	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1250	A	N1-C6-N6	-6.75	114.55	118.60
86	AB	47	U	P-O5'-C5'	6.75	131.70	120.90
34	BA	1517	U	O4'-C1'-N1	6.75	113.60	108.20
35	BB	7	C	O4'-C1'-N1	6.75	113.60	108.20
35	BB	876	G	P-O5'-C5'	6.75	131.70	120.90
35	BB	1176	G	O4'-C1'-N9	6.75	113.60	108.20
38	BE	41	C	P-O5'-C5'	-6.75	110.10	120.90
40	BG	156	G	C2'-C3'-O3'	6.75	124.50	113.70
78	Bs	14	ASN	CB-CA-C	-6.75	96.90	110.40
8	A7	92	ARG	NE-CZ-NH1	6.75	123.67	120.30
34	BA	201	A	O4'-C4'-C3'	-6.75	97.25	104.00
34	BA	285	C	O4'-C1'-N1	6.75	113.60	108.20
34	BA	362	G	O4'-C1'-C2'	6.75	113.67	107.60
34	BA	1816	G	N1-C6-O6	6.75	123.95	119.90
35	BB	738	G	N1-C6-O6	6.75	123.95	119.90
35	BB	840	C	O3'-P-O5'	6.75	116.82	104.00
38	BE	171	U	C2'-C3'-O3'	6.75	124.49	113.70
85	AA	122	A	C5-C6-N6	-6.75	118.30	123.70
85	AA	2170	G	C8-N9-C1'	6.75	135.77	127.00
35	BB	308	C	O4'-C1'-N1	6.75	113.60	108.20
35	BB	652	G	C5-C6-O6	-6.75	124.55	128.60
35	BB	677	U	C1'-O4'-C4'	-6.75	104.50	109.90
35	BB	1379	U	O3'-P-O5'	6.75	116.81	104.00
41	BH	1	U	O4'-C1'-N1	6.75	113.60	108.20
85	AA	616	A	C5'-C4'-C3'	-6.75	105.21	116.00
85	AA	2045	U	C2-N1-C1'	-6.75	109.61	117.70
34	BA	1285	G	N1-C6-O6	-6.74	115.85	119.90
34	BA	1541	G	N9-C4-C5	-6.74	102.70	105.40
35	BB	373	C	P-O3'-C3'	-6.74	111.61	119.70
35	BB	519	A	C5-C6-N6	-6.74	118.31	123.70
35	BB	742	G	O4'-C1'-N9	6.74	113.59	108.20
35	BB	1154	C	O4'-C1'-C2'	6.74	113.67	107.60
38	BE	62	C	C6-N1-C1'	6.74	128.89	120.80
74	Bo	85	ARG	NE-CZ-NH1	6.74	123.67	120.30
85	AA	387	U	P-O3'-C3'	6.74	127.79	119.70
85	AA	523	U	O4'-C1'-N1	6.74	113.59	108.20
85	AA	686	U	P-O5'-C5'	-6.74	110.11	120.90
85	AA	1216	A	O4'-C1'-N9	6.74	113.59	108.20
13	AE	88	ARG	NE-CZ-NH1	6.74	123.67	120.30
34	BA	1539	A	P-O3'-C3'	-6.74	111.61	119.70
35	BB	5	A	C8-N9-C4	-6.74	103.10	105.80
49	BP	158	PRO	N-CA-C	-6.74	94.57	112.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	113	G	C3'-C2'-C1'	-6.74	96.11	101.50
34	BA	691	A	P-O5'-C5'	-6.74	110.12	120.90
34	BA	906	A	C8-N9-C4	6.74	108.50	105.80
34	BA	1106	A	C5'-C4'-O4'	-6.74	101.01	109.10
34	BA	1710	C	C3'-C2'-C1'	-6.74	96.11	101.50
35	BB	472	C	O3'-P-O5'	6.74	116.81	104.00
36	BC	12	A	C6-N1-C2	-6.74	114.56	118.60
72	Bm	36	ASP	CA-CB-CG	-6.74	98.57	113.40
85	AA	443	A	C4'-C3'-C2'	6.74	109.34	102.60
85	AA	774	C	O4'-C1'-N1	6.74	113.59	108.20
85	AA	991	G	O4'-C1'-N9	6.74	113.59	108.20
85	AA	1646	U	C1'-O4'-C4'	-6.74	104.51	109.90
85	AA	2060	G	N1-C6-O6	6.74	123.94	119.90
6	A5	117	TYR	CB-CG-CD2	-6.74	116.96	121.00
34	BA	223	U	O4'-C1'-N1	6.74	113.59	108.20
35	BB	1091	C	N1-C2-N3	6.74	123.92	119.20
36	BC	18	G	C4-N9-C1'	-6.74	117.74	126.50
40	BG	73	U	N3-C2-O2	-6.74	117.48	122.20
41	BH	51	C	C6-N1-C2	-6.74	117.60	120.30
85	AA	632	U	O4'-C1'-N1	6.74	113.59	108.20
85	AA	1954	C	O4'-C1'-N1	6.74	113.59	108.20
85	AA	2060	G	C8-N9-C1'	6.74	135.76	127.00
34	BA	339	G	OP1-P-O3'	6.74	120.02	105.20
35	BB	1263	A	C5-C6-N1	6.74	121.07	117.70
32	AY	37	ARG	N-CA-CB	-6.74	98.47	110.60
34	BA	1700	C	O5'-P-OP2	-6.74	99.64	105.70
35	BB	1086	G	P-O5'-C5'	-6.74	110.12	120.90
35	BB	1093	C	C2-N3-C4	-6.74	116.53	119.90
35	BB	1177	U	P-O3'-C3'	-6.74	111.62	119.70
35	BB	1455	A	P-O3'-C3'	6.74	127.78	119.70
37	BD	93	G	C8-N9-C1'	6.74	135.76	127.00
38	BE	160	C	C5'-C4'-C3'	-6.74	105.22	116.00
39	BF	7	G	O4'-C1'-N9	6.74	113.59	108.20
85	AA	8	U	P-O5'-C5'	-6.74	110.12	120.90
85	AA	51	A	N1-C6-N6	6.74	122.64	118.60
85	AA	379	U	P-O3'-C3'	-6.74	111.62	119.70
85	AA	1107	A	N1-C6-N6	-6.74	114.56	118.60
2	A1	240	GLN	C-N-CA	6.73	136.44	122.30
34	BA	585	G	C4-N9-C1'	6.73	135.25	126.50
34	BA	851	C	C6-N1-C1'	6.73	128.88	120.80
34	BA	1208	U	C2-N1-C1'	-6.73	109.62	117.70
35	BB	4	C	C3'-C2'-C1'	6.73	106.89	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	35	G	C4-N9-C1'	-6.73	117.75	126.50
35	BB	51	U	C1'-O4'-C4'	-6.73	104.51	109.90
40	BG	20	U	C5'-C4'-C3'	-6.73	105.23	116.00
41	BH	96	G	O4'-C1'-N9	6.73	113.59	108.20
85	AA	1978	G	P-O5'-C5'	-6.73	110.13	120.90
34	BA	170	U	C2-N3-C4	-6.73	122.96	127.00
34	BA	563	A	C5-C6-N6	6.73	129.09	123.70
34	BA	763	U	C1'-O4'-C4'	6.73	115.29	109.90
34	BA	1001	G	C6-N1-C2	-6.73	121.06	125.10
34	BA	1281	U	O3'-P-O5'	6.73	116.79	104.00
35	BB	312	U	C4'-C3'-C2'	-6.73	95.87	102.60
35	BB	1132	A	N1-C2-N3	-6.73	125.93	129.30
85	AA	523	U	C5'-C4'-C3'	-6.73	105.23	116.00
85	AA	1593	C	P-O3'-C3'	6.73	127.78	119.70
85	AA	2224	U	O4'-C1'-N1	6.73	113.59	108.20
86	AB	46	G	P-O3'-C3'	-6.73	111.62	119.70
5	A4	91	PHE	CA-CB-CG	-6.73	97.75	113.90
34	BA	700	G	C4-N9-C1'	-6.73	117.75	126.50
34	BA	1528	U	C4'-C3'-C2'	6.73	109.33	102.60
35	BB	459	U	O4'-C1'-N1	6.73	113.58	108.20
35	BB	502	C	C2-N1-C1'	-6.73	111.40	118.80
36	BC	153	C	C2'-C3'-O3'	6.73	124.47	113.70
40	BG	9	G	C3'-C2'-C1'	-6.73	96.11	101.50
40	BG	171	A	P-O5'-C5'	6.73	131.67	120.90
77	Br	33	HIS	CA-CB-CG	-6.73	102.16	113.60
85	AA	535	G	C8-N9-C4	6.73	109.09	106.40
34	BA	126	G	N1-C6-O6	6.73	123.94	119.90
34	BA	333	A	C5-C6-N6	-6.73	118.32	123.70
34	BA	568	G	O4'-C1'-N9	6.73	113.58	108.20
35	BB	657	A	C3'-C2'-C1'	6.73	106.88	101.50
37	BD	115	A	N1-C6-N6	6.73	122.64	118.60
38	BE	96	G	N3-C2-N2	-6.73	115.19	119.90
39	BF	19	A	O4'-C4'-C3'	-6.73	97.27	104.00
85	AA	267	U	C4'-C3'-C2'	6.73	109.33	102.60
85	AA	1251	G	C5'-C4'-C3'	-6.73	105.23	116.00
85	AA	1934	A	C5'-C4'-O4'	6.73	117.17	109.10
34	BA	744	G	N3-C4-C5	-6.73	125.24	128.60
34	BA	1636	C	C2-N3-C4	6.73	123.26	119.90
34	BA	1675	C	C6-N1-C2	-6.73	117.61	120.30
35	BB	830	G	O4'-C1'-N9	6.73	113.58	108.20
54	BU	108	LYS	N-CA-CB	-6.73	98.49	110.60
80	Bu	75	VAL	CG1-CB-CG2	6.73	121.66	110.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	425	G	C8-N9-C4	6.73	109.09	106.40
85	AA	825	U	C2-N3-C4	-6.73	122.96	127.00
85	AA	875	C	N1-C1'-C2'	-6.73	104.60	112.00
34	BA	401	A	C3'-C2'-C1'	-6.73	96.12	101.50
34	BA	516	U	C2-N1-C1'	-6.73	109.63	117.70
34	BA	1692	U	C2-N1-C1'	6.73	125.77	117.70
34	BA	1735	G	N3-C4-N9	-6.73	121.96	126.00
35	BB	772	U	C2-N1-C1'	-6.73	109.63	117.70
35	BB	780	U	C3'-C2'-C1'	6.73	106.88	101.50
35	BB	869	G	C8-N9-C1'	6.73	135.74	127.00
85	AA	288	G	C4-N9-C1'	-6.73	117.76	126.50
34	BA	1101	A	N9-C1'-C2'	-6.72	104.60	112.00
34	BA	1470	G	C5-C6-O6	-6.72	124.57	128.60
34	BA	1631	U	C6-N1-C2	-6.72	116.97	121.00
34	BA	1692	U	C6-N1-C2	-6.72	116.97	121.00
40	BG	27	C	C1'-O4'-C4'	-6.72	104.52	109.90
40	BG	38	A	N1-C6-N6	6.72	122.63	118.60
41	BH	101	A	N1-C2-N3	-6.72	125.94	129.30
58	BY	73	ARG	NE-CZ-NH2	-6.72	116.94	120.30
67	Bh	71	ALA	N-CA-CB	-6.72	100.69	110.10
85	AA	1674	G	C8-N9-C4	6.72	109.09	106.40
34	BA	327	G	C5-C6-O6	-6.72	124.57	128.60
34	BA	712	C	P-O5'-C5'	6.72	131.66	120.90
34	BA	733	G	C5-C6-O6	-6.72	124.57	128.60
34	BA	892	C	C2-N1-C1'	-6.72	111.40	118.80
34	BA	1218	G	C5'-C4'-C3'	6.72	126.76	116.00
35	BB	787	A	C8-N9-C1'	6.72	139.80	127.70
35	BB	812	G	N9-C1'-C2'	-6.72	104.60	112.00
41	BH	46	C	C2-N1-C1'	-6.72	111.41	118.80
85	AA	380	C	P-O3'-C3'	-6.72	111.63	119.70
85	AA	1371	C	C2-N1-C1'	-6.72	111.41	118.80
85	AA	1864	G	C1'-O4'-C4'	-6.72	104.52	109.90
86	AB	13	C	C1'-O4'-C4'	-6.72	104.52	109.90
34	BA	758	G	C5-C6-N1	6.72	114.86	111.50
35	BB	1159	U	N3-C2-O2	-6.72	117.50	122.20
85	AA	732	G	C8-N9-C4	-6.72	103.71	106.40
34	BA	47	U	N1-C2-N3	6.72	118.93	114.90
34	BA	508	C	C1'-O4'-C4'	-6.72	104.52	109.90
34	BA	1523	U	C2-N3-C4	-6.72	122.97	127.00
85	AA	130	G	C8-N9-C1'	6.72	135.74	127.00
85	AA	838	G	C6-N1-C2	-6.72	121.07	125.10
85	AA	968	U	P-O5'-C5'	-6.72	110.15	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1128	G	C5-C6-N1	6.72	114.86	111.50
34	BA	280	A	C5-C6-N6	-6.72	118.33	123.70
34	BA	1808	A	C6-N1-C2	-6.72	114.57	118.60
35	BB	1288	G	C8-N9-C4	6.72	109.09	106.40
69	Bj	3	CYS	N-CA-CB	6.72	122.69	110.60
85	AA	753	U	O5'-C5'-C4'	6.72	124.46	111.70
85	AA	1215	A	C5'-C4'-C3'	-6.72	105.25	116.00
85	AA	1949	U	O4'-C1'-N1	6.72	113.57	108.20
5	A4	67	LEU	CB-CA-C	6.72	122.96	110.20
34	BA	288	U	O4'-C1'-N1	6.72	113.57	108.20
34	BA	1176	C	C2'-C3'-O3'	6.72	124.45	113.70
34	BA	1597	G	C5-C6-O6	-6.72	124.57	128.60
35	BB	1000	U	O4'-C4'-C3'	-6.72	97.28	104.00
35	BB	1201	G	C5-C6-O6	-6.72	124.57	128.60
35	BB	1220	A	N9-C1'-C2'	-6.72	104.61	112.00
35	BB	1254	G	O5'-C5'-C4'	-6.72	98.94	111.70
40	BG	108	G	N9-C1'-C2'	-6.72	104.61	112.00
52	BS	156	ARG	C-N-CA	6.72	138.49	121.70
59	BZ	66	ARG	NE-CZ-NH1	6.72	123.66	120.30
85	AA	476	C	P-O5'-C5'	-6.72	110.15	120.90
85	AA	515	C	P-O5'-C5'	-6.72	110.15	120.90
85	AA	607	U	C1'-O4'-C4'	-6.72	104.53	109.90
85	AA	902	A	N1-C6-N6	6.72	122.63	118.60
85	AA	1038	U	O4'-C1'-N1	6.72	113.57	108.20
85	AA	1044	G	O4'-C1'-N9	6.72	113.57	108.20
85	AA	1191	G	N1-C6-O6	-6.72	115.87	119.90
85	AA	2004	U	C6-N1-C2	-6.72	116.97	121.00
85	AA	2074	G	C8-N9-C1'	6.72	135.73	127.00
4	A3	221	ARG	NE-CZ-NH1	6.71	123.66	120.30
34	BA	13	U	C5'-C4'-O4'	6.71	117.16	109.10
34	BA	545	U	C5'-C4'-O4'	6.71	117.16	109.10
34	BA	1026	C	C6-N1-C2	-6.71	117.61	120.30
34	BA	1248	A	C1'-O4'-C4'	-6.71	104.53	109.90
35	BB	1115	G	C1'-O4'-C4'	-6.71	104.53	109.90
36	BC	18	G	N3-C4-N9	-6.71	121.97	126.00
36	BC	87	C	C5'-C4'-O4'	6.71	117.16	109.10
53	BT	85	ARG	NE-CZ-NH1	6.71	123.66	120.30
85	AA	100	A	C1'-O4'-C4'	-6.71	104.53	109.90
85	AA	863	C	C1'-O4'-C4'	-6.71	104.53	109.90
85	AA	1460	G	C4'-C3'-C2'	-6.71	95.89	102.60
85	AA	1538	C	C5'-C4'-O4'	6.71	117.16	109.10
34	BA	1296	U	C2-N3-C4	-6.71	122.97	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1394	U	O4'-C1'-N1	6.71	113.57	108.20
35	BB	1074	U	O4'-C1'-N1	6.71	113.57	108.20
40	BG	162	A	N1-C6-N6	6.71	122.63	118.60
47	BN	5	ASN	C-N-CA	6.71	138.48	121.70
85	AA	83	U	O5'-C5'-C4'	-6.71	98.94	111.70
85	AA	336	C	C3'-C2'-C1'	-6.71	96.13	101.50
34	BA	1483	U	N3-C2-O2	-6.71	117.50	122.20
34	BA	1529	G	N1-C6-O6	-6.71	115.87	119.90
34	BA	1746	G	C5-C6-O6	-6.71	124.57	128.60
35	BB	1374	U	C2-N3-C4	-6.71	122.97	127.00
38	BE	25	U	C3'-C2'-C1'	-6.71	96.13	101.50
85	AA	962	U	O4'-C1'-N1	6.71	113.57	108.20
85	AA	969	U	P-O3'-C3'	6.71	127.75	119.70
85	AA	1226	A	C5'-C4'-O4'	-6.71	101.05	109.10
86	AB	19	G	O4'-C1'-N9	6.71	113.57	108.20
34	BA	680	C	O4'-C1'-N1	6.71	113.57	108.20
35	BB	364	U	C2-N1-C1'	6.71	125.75	117.70
37	BD	16	U	C5'-C4'-C3'	-6.71	105.26	116.00
85	AA	627	A	P-O5'-C5'	-6.71	110.17	120.90
34	BA	815	C	C2-N1-C1'	6.71	126.18	118.80
34	BA	1485	U	O5'-P-OP2	-6.71	99.66	105.70
34	BA	1651	C	C5'-C4'-C3'	-6.71	105.27	116.00
34	BA	1662	U	C4'-C3'-C2'	-6.71	95.89	102.60
34	BA	1776	G	P-O3'-C3'	6.71	127.75	119.70
35	BB	57	G	C8-N9-C4	-6.71	103.72	106.40
35	BB	434	A	P-O5'-C5'	-6.71	110.17	120.90
35	BB	853	U	OP1-P-OP2	-6.71	109.54	119.60
35	BB	1162	A	C6-N1-C2	-6.71	114.58	118.60
35	BB	1285	U	N1-C2-N3	-6.71	110.88	114.90
39	BF	25	G	N1-C6-O6	6.71	123.92	119.90
52	BS	95	GLU	N-CA-CB	-6.71	98.53	110.60
74	Bo	84	ARG	NE-CZ-NH1	6.71	123.65	120.30
85	AA	498	C	C6-N1-C2	-6.71	117.62	120.30
85	AA	827	C	C6-N1-C2	-6.71	117.62	120.30
85	AA	2141	G	C8-N9-C1'	6.71	135.72	127.00
26	AS	142	ARG	NE-CZ-NH1	6.71	123.65	120.30
34	BA	16	C	C6-N1-C2	-6.71	117.62	120.30
34	BA	335	C	N3-C2-O2	-6.71	117.21	121.90
34	BA	924	U	N3-C2-O2	-6.71	117.50	122.20
34	BA	1005	C	O4'-C1'-C2'	-6.71	99.09	105.80
34	BA	1019	C	N1-C1'-C2'	-6.71	104.62	112.00
35	BB	1087	A	C1'-O4'-C4'	-6.71	104.53	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1469	A	C5-C6-N1	6.71	121.05	117.70
38	BE	176	G	C3'-C2'-C1'	-6.71	96.14	101.50
40	BG	105	A	C2'-C3'-O3'	6.71	124.43	113.70
47	BN	66	PRO	N-CA-C	6.71	129.54	112.10
85	AA	597	A	C1'-O4'-C4'	-6.71	104.53	109.90
85	AA	623	G	C5-C6-O6	-6.71	124.58	128.60
5	A4	3	ALA	N-CA-C	-6.71	92.90	111.00
34	BA	448	U	C6-N1-C2	-6.71	116.98	121.00
41	BH	5	G	C4-N9-C1'	-6.71	117.78	126.50
85	AA	316	C	N3-C2-O2	-6.71	117.21	121.90
33	AZ	39	GLN	N-CA-C	-6.70	92.90	111.00
34	BA	313	C	P-O5'-C5'	6.70	131.62	120.90
34	BA	568	G	C4-N9-C1'	-6.70	117.78	126.50
34	BA	613	A	N1-C6-N6	-6.70	114.58	118.60
34	BA	961	C	C5'-C4'-C3'	-6.70	105.28	116.00
34	BA	1203	G	N9-C1'-C2'	-6.70	104.63	112.00
34	BA	1454	G	C5-N7-C8	-6.70	100.95	104.30
35	BB	996	G	C4-N9-C1'	-6.70	117.79	126.50
35	BB	1421	C	N1-C2-O2	6.70	122.92	118.90
36	BC	145	G	C8-N9-C4	-6.70	103.72	106.40
37	BD	81	C	C2-N3-C4	-6.70	116.55	119.90
37	BD	95	G	C5'-C4'-C3'	6.70	126.73	116.00
38	BE	198	A	O4'-C1'-N9	6.70	113.56	108.20
41	BH	72	G	P-O5'-C5'	-6.70	110.17	120.90
41	BH	129	G	C5'-C4'-O4'	6.70	117.14	109.10
80	Bu	169	LEU	CB-CG-CD2	6.70	122.39	111.00
85	AA	471	U	P-O3'-C3'	-6.70	111.66	119.70
85	AA	910	G	O4'-C1'-N9	6.70	113.56	108.20
85	AA	1229	G	N3-C2-N2	6.70	124.59	119.90
85	AA	1561	A	N1-C6-N6	-6.70	114.58	118.60
85	AA	1715	C	C1'-O4'-C4'	-6.70	104.54	109.90
85	AA	1816	C	P-O5'-C5'	6.70	131.62	120.90
85	AA	1878	C	P-O3'-C3'	-6.70	111.66	119.70
34	BA	1420	A	C8-N9-C4	-6.70	103.12	105.80
35	BB	1130	U	O5'-C5'-C4'	-6.70	98.97	111.70
65	Bf	312	HIS	CA-CB-CG	-6.70	102.21	113.60
84	By	121	ARG	NE-CZ-NH1	6.70	123.65	120.30
85	AA	2203	C	O4'-C1'-N1	6.70	113.56	108.20
20	AL	60	ARG	NE-CZ-NH1	6.70	123.65	120.30
34	BA	190	U	C4'-C3'-C2'	-6.70	95.90	102.60
34	BA	760	G	C4-N9-C1'	-6.70	117.79	126.50
34	BA	1073	G	C5-C6-N1	6.70	114.85	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1704	G	O3'-P-O5'	6.70	116.73	104.00
35	BB	1371	G	C1'-O4'-C4'	-6.70	104.54	109.90
36	BC	10	C	N1-C1'-C2'	-6.70	104.63	112.00
41	BH	12	U	O4'-C1'-N1	6.70	113.56	108.20
50	BQ	50	GLN	N-CA-C	6.70	129.09	111.00
59	BZ	41	ASN	CA-CB-CG	-6.70	98.66	113.40
65	Bf	303	ARG	NE-CZ-NH1	6.70	123.65	120.30
85	AA	1794	U	C6-N1-C2	-6.70	116.98	121.00
85	AA	2121	G	N9-C4-C5	6.70	108.08	105.40
34	BA	214	A	N3-C4-N9	6.70	132.76	127.40
34	BA	449	G	P-O3'-C3'	-6.70	111.66	119.70
34	BA	774	A	O4'-C1'-N9	6.70	113.56	108.20
35	BB	823	G	N1-C6-O6	-6.70	115.88	119.90
35	BB	1033	U	P-O3'-C3'	-6.70	111.66	119.70
38	BE	203	C	C4-C5-C6	-6.70	114.05	117.40
82	Bw	8	LEU	CB-CA-C	-6.70	97.47	110.20
85	AA	307	G	O4'-C4'-C3'	-6.70	97.30	104.00
85	AA	850	U	C2-N3-C4	-6.70	122.98	127.00
85	AA	991	G	C2'-C3'-O3'	6.70	124.42	113.70
85	AA	1211	C	C5-C4-N4	-6.70	115.51	120.20
85	AA	1247	A	C4'-C3'-C2'	-6.70	95.90	102.60
85	AA	1381	C	O4'-C1'-N1	6.70	113.56	108.20
34	BA	971	G	C5-C6-O6	-6.70	124.58	128.60
35	BB	1523	U	O5'-P-OP2	-6.70	99.67	105.70
36	BC	138	C	C3'-C2'-C1'	-6.70	96.14	101.50
84	By	74	HIS	CB-CA-C	6.70	123.79	110.40
85	AA	1480	C	C5'-C4'-C3'	-6.70	105.28	116.00
85	AA	2237	G	O4'-C1'-C2'	6.70	113.63	107.60
34	BA	223	U	P-O5'-C5'	-6.70	110.19	120.90
34	BA	684	G	C8-N9-C4	-6.70	103.72	106.40
35	BB	128	C	O4'-C1'-N1	6.70	113.56	108.20
35	BB	795	A	C1'-O4'-C4'	-6.70	104.54	109.90
35	BB	1024	G	O5'-C5'-C4'	-6.70	98.98	111.70
35	BB	1262	A	C6-N1-C2	-6.70	114.58	118.60
37	BD	75	G	C5'-C4'-C3'	-6.70	105.29	116.00
38	BE	40	C	C2'-C3'-O3'	6.70	124.41	113.70
40	BG	2	U	C1'-O4'-C4'	-6.70	104.54	109.90
41	BH	20	A	C8-N9-C1'	6.70	139.75	127.70
85	AA	1537	A	C3'-C2'-C1'	6.70	106.86	101.50
85	AA	2015	U	C2-N1-C1'	-6.70	109.67	117.70
69	Bj	70	ARG	NE-CZ-NH2	-6.69	116.95	120.30
85	AA	860	C	O4'-C1'-C2'	6.69	113.62	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2170	G	P-O3'-C3'	-6.69	111.67	119.70
31	AX	188	MET	CG-SD-CE	-6.69	89.49	100.20
34	BA	1306	U	C5'-C4'-O4'	6.69	117.13	109.10
34	BA	1561	C	C2-N1-C1'	-6.69	111.44	118.80
35	BB	1327	U	O4'-C1'-N1	6.69	113.55	108.20
37	BD	119	U	C2-N1-C1'	6.69	125.73	117.70
38	BE	196	C	N1-C2-O2	6.69	122.92	118.90
40	BG	86	U	C1'-O4'-C4'	-6.69	104.55	109.90
41	BH	128	G	O5'-P-OP2	6.69	118.73	110.70
83	Bx	118	ARG	NE-CZ-NH1	6.69	123.65	120.30
85	AA	405	C	O4'-C1'-N1	6.69	113.55	108.20
85	AA	474	C	O4'-C1'-N1	6.69	113.55	108.20
85	AA	1504	A	N9-C1'-C2'	-6.69	104.64	112.00
17	AI	137	ARG	NE-CZ-NH1	6.69	123.64	120.30
34	BA	223	U	C2-N1-C1'	-6.69	109.67	117.70
34	BA	305	C	O4'-C1'-N1	6.69	113.55	108.20
34	BA	717	U	C1'-O4'-C4'	-6.69	104.55	109.90
34	BA	1061	A	P-O3'-C3'	-6.69	111.67	119.70
34	BA	1224	A	C2'-C3'-O3'	6.69	124.40	113.70
34	BA	1300	G	O4'-C1'-C2'	6.69	113.62	107.60
35	BB	375	G	C5-C6-N1	6.69	114.84	111.50
85	AA	816	A	C8-N9-C4	-6.69	103.12	105.80
85	AA	1016	G	O3'-P-O5'	-6.69	91.29	104.00
85	AA	1339	C	O4'-C1'-N1	6.69	113.55	108.20
85	AA	2163	G	C8-N9-C1'	6.69	135.70	127.00
34	BA	239	C	O4'-C1'-N1	6.69	113.55	108.20
34	BA	247	U	C1'-O4'-C4'	-6.69	104.55	109.90
41	BH	31	A	C2'-C3'-O3'	6.69	124.40	113.70
64	Be	70	ARG	CD-NE-CZ	6.69	132.96	123.60
85	AA	663	C	C3'-C2'-C1'	-6.69	96.15	101.50
85	AA	1185	G	N1-C6-O6	6.69	123.91	119.90
34	BA	62	A	P-O3'-C3'	6.69	127.73	119.70
34	BA	218	G	C1'-O4'-C4'	-6.69	104.55	109.90
34	BA	1825	U	C5-C4-O4	-6.69	121.89	125.90
35	BB	27	C	C2-N1-C1'	-6.69	111.44	118.80
35	BB	647	U	N3-C2-O2	-6.69	117.52	122.20
35	BB	1179	C	C6-N1-C2	-6.69	117.62	120.30
35	BB	1258	G	OP1-P-OP2	-6.69	109.57	119.60
36	BC	116	C	C6-N1-C1'	6.69	128.83	120.80
39	BF	49	C	C5-C6-N1	6.69	124.34	121.00
40	BG	16	G	O4'-C4'-C3'	-6.69	97.31	104.00
41	BH	34	G	C5'-C4'-O4'	6.69	117.13	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	336	C	C6-N1-C2	-6.69	117.62	120.30
85	AA	846	U	O4'-C1'-N1	6.69	113.55	108.20
85	AA	960	G	C8-N9-C4	-6.69	103.72	106.40
85	AA	1254	A	C1'-O4'-C4'	-6.69	104.55	109.90
85	AA	1801	U	O3'-P-O5'	-6.69	91.30	104.00
85	AA	1876	U	C5'-C4'-C3'	6.69	126.70	116.00
36	BC	108	A	N9-C1'-C2'	-6.69	104.65	112.00
85	AA	391	G	C8-N9-C1'	6.69	135.69	127.00
34	BA	19	G	C8-N9-C1'	6.68	135.69	127.00
34	BA	907	A	N9-C1'-C2'	-6.68	104.65	112.00
34	BA	1210	A	N1-C2-N3	-6.68	125.96	129.30
34	BA	1367	G	C5'-C4'-C3'	-6.68	105.31	116.00
35	BB	347	G	C5-C6-O6	-6.68	124.59	128.60
35	BB	384	A	C6-C5-N7	-6.68	127.62	132.30
35	BB	743	C	C5-C4-N4	-6.68	115.52	120.20
35	BB	1083	C	N3-C2-O2	-6.68	117.22	121.90
35	BB	1360	A	C4-C5-C6	-6.68	113.66	117.00
40	BG	128	U	N1-C1'-C2'	-6.68	104.65	112.00
51	BR	63	TYR	CA-CB-CG	-6.68	100.70	113.40
58	BY	14	HIS	CB-CA-C	6.68	123.77	110.40
85	AA	23	G	C5'-C4'-C3'	6.68	126.70	116.00
85	AA	258	G	C5-C6-N1	6.68	114.84	111.50
85	AA	416	U	P-O3'-C3'	6.68	127.72	119.70
85	AA	1254	A	C4-N9-C1'	-6.68	114.27	126.30
85	AA	1560	A	C5-N7-C8	6.68	107.24	103.90
85	AA	2141	G	C4'-C3'-C2'	6.68	109.28	102.60
34	BA	139	U	N3-C2-O2	-6.68	117.52	122.20
38	BE	31	A	N7-C8-N9	6.68	117.14	113.80
85	AA	416	U	C6-N1-C2	-6.68	116.99	121.00
85	AA	1585	A	P-O5'-C5'	-6.68	110.21	120.90
34	BA	788	C	C4'-C3'-C2'	6.68	109.28	102.60
35	BB	566	A	O4'-C1'-N9	6.68	113.55	108.20
34	BA	6	C	N3-C2-O2	-6.68	117.22	121.90
34	BA	39	C	P-O5'-C5'	6.68	131.59	120.90
34	BA	212	A	C8-N9-C1'	6.68	139.72	127.70
34	BA	429	G	C6-N1-C2	-6.68	121.09	125.10
34	BA	843	G	C1'-O4'-C4'	-6.68	104.56	109.90
34	BA	964	U	C4'-C3'-C2'	-6.68	95.92	102.60
34	BA	1641	G	C5-C6-O6	-6.68	124.59	128.60
35	BB	423	G	P-O3'-C3'	-6.68	111.69	119.70
38	BE	81	C	O4'-C1'-N1	6.68	113.54	108.20
39	BF	5	U	C4'-C3'-C2'	-6.68	95.92	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
73	Bn	33	ARG	NE-CZ-NH1	6.68	123.64	120.30
85	AA	18	C	C6-N1-C2	-6.68	117.63	120.30
85	AA	1302	A	O5'-C5'-C4'	6.68	124.39	111.70
85	AA	1465	C	C2-N1-C1'	-6.68	111.45	118.80
85	AA	1593	C	O4'-C1'-C2'	6.68	113.61	107.60
85	AA	1900	C	N3-C4-N4	-6.68	113.32	118.00
24	AQ	21	MET	CG-SD-CE	-6.68	89.52	100.20
35	BB	1507	U	O5'-C5'-C4'	-6.68	99.01	111.70
38	BE	112	G	C8-N9-C4	-6.68	103.73	106.40
65	Bf	72	ARG	N-CA-CB	6.68	122.62	110.60
85	AA	1379	A	O4'-C1'-N9	6.68	113.54	108.20
85	AA	1931	C	P-O3'-C3'	6.68	127.71	119.70
85	AA	2215	C	O5'-P-OP1	-6.68	99.69	105.70
34	BA	25	C	O5'-P-OP2	6.68	118.71	110.70
34	BA	319	C	C4'-C3'-C2'	-6.68	95.92	102.60
34	BA	1284	G	P-O5'-C5'	-6.68	110.22	120.90
35	BB	706	G	C4-N9-C1'	-6.68	117.82	126.50
35	BB	969	C	P-O3'-C3'	-6.68	111.69	119.70
35	BB	1508	G	O4'-C4'-C3'	-6.68	97.32	104.00
50	BQ	58	ARG	NE-CZ-NH1	6.68	123.64	120.30
65	Bf	390	ARG	NE-CZ-NH1	6.68	123.64	120.30
77	Br	160	TYR	CB-CG-CD1	6.68	125.01	121.00
85	AA	438	G	O5'-C5'-C4'	-6.68	99.02	111.70
85	AA	449	G	N9-C1'-C2'	-6.68	104.66	112.00
85	AA	467	U	C6-N1-C1'	-6.68	111.85	121.20
85	AA	589	A	OP1-P-OP2	-6.68	109.59	119.60
85	AA	1466	U	O5'-C5'-C4'	6.68	124.39	111.70
85	AA	1917	G	C4-N9-C1'	-6.68	117.82	126.50
86	AB	15	G	C8-N9-C4	-6.68	103.73	106.40
34	BA	196	A	C5-C6-N6	-6.67	118.36	123.70
34	BA	1220	C	O5'-C5'-C4'	-6.67	99.02	111.70
34	BA	1368	G	P-O5'-C5'	6.67	131.58	120.90
35	BB	498	G	C5-C6-N1	6.67	114.84	111.50
35	BB	574	G	C4-N9-C1'	-6.67	117.82	126.50
35	BB	689	C	C5'-C4'-C3'	6.67	126.68	116.00
35	BB	1266	A	C3'-C2'-C1'	-6.67	96.16	101.50
36	BC	125	A	P-O3'-C3'	-6.67	111.69	119.70
39	BF	57	C	O5'-P-OP2	-6.67	99.69	105.70
39	BF	63	U	O4'-C1'-N1	6.67	113.54	108.20
40	BG	125	C	N3-C2-O2	-6.67	117.23	121.90
67	Bh	89	SER	N-CA-CB	-6.67	100.49	110.50
82	Bw	70	TYR	CA-CB-CG	-6.67	100.72	113.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	30	G	C4-N9-C1'	-6.67	117.82	126.50
85	AA	874	A	O5'-C5'-C4'	6.67	124.38	111.70
34	BA	770	G	C2-N3-C4	-6.67	108.56	111.90
34	BA	1710	C	N3-C2-O2	-6.67	117.23	121.90
35	BB	1453	G	C4-N9-C1'	-6.67	117.83	126.50
85	AA	973	U	O3'-P-O5'	6.67	116.68	104.00
85	AA	1650	G	O4'-C1'-N9	6.67	113.54	108.20
85	AA	1897	A	C2-N3-C4	6.67	113.94	110.60
34	BA	36	A	C4'-C3'-C2'	-6.67	95.93	102.60
34	BA	316	G	C5-C6-N1	6.67	114.84	111.50
34	BA	1221	A	C5'-C4'-C3'	6.67	126.67	116.00
34	BA	1425	G	O3'-P-O5'	-6.67	91.33	104.00
35	BB	715	G	C5'-C4'-O4'	6.67	117.11	109.10
35	BB	835	C	C2'-C3'-O3'	6.67	124.38	113.70
35	BB	1439	U	C5'-C4'-C3'	6.67	126.68	116.00
38	BE	74	U	C2-N3-C4	-6.67	123.00	127.00
41	BH	69	C	O4'-C1'-N1	6.67	113.54	108.20
85	AA	66	U	C6-N1-C2	-6.67	117.00	121.00
85	AA	270	A	C3'-C2'-C1'	-6.67	96.16	101.50
85	AA	800	A	C5-N7-C8	-6.67	100.56	103.90
85	AA	1544	G	C8-N9-C1'	6.67	135.67	127.00
85	AA	2033	C	C5'-C4'-C3'	-6.67	105.33	116.00
34	BA	128	C	C4'-C3'-C2'	6.67	109.27	102.60
34	BA	208	A	C5-C6-N6	6.67	129.03	123.70
34	BA	342	U	P-O3'-C3'	-6.67	111.70	119.70
34	BA	573	U	OP1-P-O3'	6.67	119.87	105.20
34	BA	1829	A	O4'-C1'-N9	6.67	113.53	108.20
35	BB	822	G	N7-C8-N9	6.67	116.43	113.10
40	BG	23	C	C4'-C3'-C2'	-6.67	95.93	102.60
57	BX	101	ASN	CA-CB-CG	-6.67	98.73	113.40
80	Bu	269	ARG	NE-CZ-NH1	6.67	123.63	120.30
85	AA	264	A	N9-C4-C5	-6.67	103.13	105.80
85	AA	399	A	O4'-C1'-N9	6.67	113.54	108.20
85	AA	834	U	C3'-C2'-C1'	-6.67	96.17	101.50
34	BA	304	G	O3'-P-O5'	6.67	116.67	104.00
34	BA	506	U	C2-N3-C4	-6.67	123.00	127.00
34	BA	614	A	O4'-C1'-N9	6.67	113.53	108.20
34	BA	664	C	C3'-C2'-C1'	-6.67	96.17	101.50
36	BC	33	U	C6-N1-C2	-6.67	117.00	121.00
65	Bf	165	PHE	N-CA-CB	-6.67	98.60	110.60
85	AA	25	C	O3'-P-O5'	6.67	116.67	104.00
85	AA	455	G	C8-N9-C4	6.67	109.07	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	490	A	O4'-C1'-N9	6.67	113.53	108.20
85	AA	777	U	P-O3'-C3'	-6.67	111.70	119.70
85	AA	1080	A	O4'-C1'-N9	6.67	113.53	108.20
85	AA	1525	C	O4'-C1'-N1	6.67	113.53	108.20
85	AA	2036	A	C8-N9-C1'	-6.67	115.70	127.70
34	BA	1694	C	C5'-C4'-C3'	-6.67	105.33	116.00
35	BB	1538	G	O4'-C1'-N9	6.67	113.53	108.20
85	AA	560	C	C5'-C4'-C3'	6.67	126.66	116.00
86	AB	5	G	O4'-C1'-N9	6.67	113.53	108.20
34	BA	88	C	C6-N1-C2	-6.66	117.64	120.30
34	BA	103	G	C5'-C4'-C3'	-6.66	105.34	116.00
34	BA	294	C	N1-C2-N3	6.66	123.86	119.20
34	BA	345	G	P-O3'-C3'	-6.66	111.70	119.70
34	BA	584	A	C5-C6-N1	6.66	121.03	117.70
34	BA	835	U	O4'-C1'-N1	6.66	113.53	108.20
34	BA	938	C	O4'-C1'-C2'	6.66	113.60	107.60
34	BA	1114	G	N1-C2-N2	-6.66	110.20	116.20
34	BA	1710	C	C1'-O4'-C4'	-6.66	104.57	109.90
35	BB	78	C	N3-C2-O2	-6.66	117.23	121.90
58	BY	74	ARG	NE-CZ-NH1	6.66	123.63	120.30
85	AA	182	C	O4'-C1'-N1	6.66	113.53	108.20
85	AA	836	A	O4'-C1'-N9	6.66	113.53	108.20
85	AA	868	A	N1-C6-N6	6.66	122.60	118.60
85	AA	1540	A	N7-C8-N9	6.66	117.13	113.80
85	AA	1818	C	C5-C6-N1	6.66	124.33	121.00
34	BA	596	G	N3-C4-N9	6.66	130.00	126.00
34	BA	1497	A	N3-C4-N9	-6.66	122.07	127.40
41	BH	34	G	C5-C6-N1	6.66	114.83	111.50
48	BO	146	ASN	CA-CB-CG	-6.66	98.74	113.40
82	Bw	82	ARG	NE-CZ-NH2	-6.66	116.97	120.30
85	AA	293	A	C8-N9-C4	-6.66	103.14	105.80
85	AA	1199	C	C1'-O4'-C4'	-6.66	104.57	109.90
21	AM	151	ARG	NE-CZ-NH1	6.66	123.63	120.30
34	BA	58	A	C4'-C3'-C2'	-6.66	95.94	102.60
34	BA	375	C	N1-C2-O2	6.66	122.90	118.90
34	BA	644	C	N3-C4-C5	-6.66	119.24	121.90
34	BA	953	G	P-O3'-C3'	6.66	127.69	119.70
34	BA	1439	C	C2-N3-C4	-6.66	116.57	119.90
34	BA	1563	G	C4'-C3'-C2'	6.66	109.26	102.60
34	BA	1591	G	N1-C6-O6	-6.66	115.90	119.90
34	BA	1786	C	C6-N1-C1'	6.66	128.79	120.80
35	BB	855	G	C8-N9-C1'	6.66	135.66	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1356	G	N1-C2-N3	6.66	127.90	123.90
37	BD	87	G	C8-N9-C4	6.66	109.06	106.40
40	BG	11	G	C6-C5-N7	-6.66	126.40	130.40
40	BG	165	C	C6-N1-C2	-6.66	117.64	120.30
41	BH	111	U	O5'-C5'-C4'	-6.66	99.05	111.70
85	AA	354	C	C1'-O4'-C4'	-6.66	104.57	109.90
85	AA	1035	C	C2-N3-C4	-6.66	116.57	119.90
34	BA	441	A	C5-C6-N6	-6.66	118.37	123.70
34	BA	1712	U	P-O3'-C3'	-6.66	111.71	119.70
35	BB	1303	A	O4'-C1'-N9	6.66	113.53	108.20
36	BC	144	C	P-O3'-C3'	-6.66	111.71	119.70
40	BG	54	G	C5'-C4'-C3'	-6.66	105.34	116.00
48	BO	201	ARG	NE-CZ-NH2	-6.66	116.97	120.30
74	Bo	14	TYR	CA-CB-CG	-6.66	100.75	113.40
85	AA	109	G	C8-N9-C4	-6.66	103.74	106.40
85	AA	413	G	C4-N9-C1'	-6.66	117.84	126.50
85	AA	760	U	C1'-O4'-C4'	-6.66	104.57	109.90
34	BA	1663	U	N1-C2-N3	6.66	118.89	114.90
36	BC	17	U	C5'-C4'-O4'	6.66	117.09	109.10
39	BF	45	G	P-O5'-C5'	6.66	131.55	120.90
40	BG	1	G	P-O3'-C3'	-6.66	111.71	119.70
34	BA	14	G	P-O3'-C3'	-6.66	111.71	119.70
34	BA	1311	G	C5-C6-N1	6.66	114.83	111.50
35	BB	706	G	P-O5'-C5'	6.66	131.55	120.90
36	BC	58	G	C4-N9-C1'	-6.66	117.85	126.50
36	BC	125	A	C8-N9-C4	-6.66	103.14	105.80
38	BE	16	C	C3'-C2'-C1'	-6.66	96.17	101.50
40	BG	83	U	C2-N3-C4	-6.66	123.01	127.00
85	AA	1182	A	P-O5'-C5'	6.66	131.55	120.90
85	AA	1281	G	N9-C4-C5	-6.66	102.74	105.40
35	BB	21	C	O5'-C5'-C4'	-6.65	99.06	111.70
38	BE	195	G	C5'-C4'-O4'	6.65	117.08	109.10
41	BH	1	U	C3'-C2'-C1'	-6.65	96.18	101.50
85	AA	544	A	O5'-P-OP1	-6.65	99.71	105.70
85	AA	755	G	O5'-C5'-C4'	-6.65	99.06	111.70
85	AA	770	C	C5-C6-N1	6.65	124.33	121.00
85	AA	963	U	C3'-C2'-C1'	-6.65	96.18	101.50
85	AA	1681	G	C8-N9-C1'	6.65	135.65	127.00
34	BA	139	U	N1-C2-O2	6.65	127.46	122.80
34	BA	1277	G	N1-C6-O6	-6.65	115.91	119.90
34	BA	1841	A	C5-C6-N1	6.65	121.03	117.70
35	BB	269	A	O4'-C1'-N9	6.65	113.52	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1539	C	P-O5'-C5'	-6.65	110.26	120.90
48	BO	75	PHE	CB-CG-CD1	6.65	125.46	120.80
85	AA	1521	U	C5-C6-N1	6.65	126.03	122.70
34	BA	1435	A	O5'-C5'-C4'	-6.65	99.06	111.70
34	BA	1748	G	C8-N9-C1'	6.65	135.65	127.00
35	BB	267	C	O4'-C1'-N1	6.65	113.52	108.20
35	BB	827	U	N3-C2-O2	-6.65	117.55	122.20
36	BC	124	A	C8-N9-C1'	-6.65	115.73	127.70
41	BH	102	C	N1-C2-O2	6.65	122.89	118.90
81	Bv	66	MET	CG-SD-CE	-6.65	89.56	100.20
85	AA	177	A	C4'-C3'-C2'	6.65	109.25	102.60
85	AA	429	G	O4'-C1'-C2'	6.65	113.59	107.60
85	AA	736	U	C4'-C3'-C2'	6.65	109.25	102.60
85	AA	1826	U	C2'-C3'-O3'	6.65	124.34	113.70
85	AA	2091	C	C5'-C4'-C3'	6.65	126.64	116.00
34	BA	34	U	P-O5'-C5'	-6.65	110.26	120.90
34	BA	288	U	C4'-C3'-C2'	-6.65	95.95	102.60
34	BA	1241	U	O5'-P-OP2	6.65	118.68	110.70
65	Bf	350	ASN	N-CA-CB	6.65	122.57	110.60
85	AA	1116	G	N1-C2-N2	-6.65	110.22	116.20
20	AL	80	ARG	NE-CZ-NH1	6.65	123.62	120.30
34	BA	941	G	C8-N9-C4	6.65	109.06	106.40
34	BA	999	G	P-O3'-C3'	-6.65	111.72	119.70
34	BA	1304	C	N3-C4-N4	-6.65	113.35	118.00
34	BA	1694	C	C1'-O4'-C4'	-6.65	104.58	109.90
35	BB	388	C	P-O3'-C3'	6.65	127.68	119.70
35	BB	788	U	P-O3'-C3'	-6.65	111.72	119.70
85	AA	147	G	C3'-C2'-C1'	-6.65	96.18	101.50
85	AA	570	U	C2-N3-C4	-6.65	123.01	127.00
85	AA	717	G	O4'-C1'-N9	6.65	113.52	108.20
85	AA	857	G	C5-C6-O6	-6.65	124.61	128.60
85	AA	1181	U	P-O5'-C5'	6.65	131.54	120.90
85	AA	1525	C	C6-N1-C1'	6.65	128.78	120.80
85	AA	1796	C	C3'-C2'-C1'	-6.65	96.18	101.50
85	AA	1963	G	P-O3'-C3'	6.65	127.68	119.70
85	AA	2234	C	C1'-O4'-C4'	-6.65	104.58	109.90
3	A2	51	ARG	NE-CZ-NH2	-6.65	116.98	120.30
34	BA	94	G	C4-N9-C1'	-6.65	117.86	126.50
34	BA	492	G	N1-C6-O6	-6.65	115.91	119.90
34	BA	862	C	O4'-C1'-N1	6.65	113.52	108.20
34	BA	1521	C	O4'-C1'-N1	6.65	113.52	108.20
34	BA	1695	G	C5'-C4'-C3'	-6.65	105.37	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	458	U	N1-C2-N3	6.65	118.89	114.90
35	BB	1313	C	P-O5'-C5'	-6.65	110.27	120.90
39	BF	56	C	O5'-C5'-C4'	-6.65	99.07	111.70
41	BH	41	A	N7-C8-N9	-6.65	110.48	113.80
85	AA	349	C	O4'-C1'-N1	6.65	113.52	108.20
85	AA	881	C	C1'-O4'-C4'	-6.65	104.58	109.90
85	AA	927	A	C6-N1-C2	-6.65	114.61	118.60
34	BA	610	A	N1-C6-N6	6.64	122.59	118.60
34	BA	683	C	C1'-O4'-C4'	-6.64	104.58	109.90
34	BA	868	C	N3-C2-O2	-6.64	117.25	121.90
34	BA	1522	G	C5-C6-N1	6.64	114.82	111.50
36	BC	32	U	C1'-O4'-C4'	-6.64	104.58	109.90
36	BC	68	A	C8-N9-C4	-6.64	103.14	105.80
36	BC	151	G	P-O3'-C3'	6.64	127.67	119.70
38	BE	41	C	O4'-C1'-N1	6.64	113.52	108.20
41	BH	77	G	OP1-P-O3'	6.64	119.82	105.20
85	AA	76	G	O4'-C4'-C3'	-6.64	97.36	104.00
85	AA	210	G	C6-N1-C2	-6.64	121.11	125.10
85	AA	515	C	N3-C2-O2	-6.64	117.25	121.90
85	AA	976	G	C1'-O4'-C4'	-6.64	104.58	109.90
85	AA	1960	C	C5-C6-N1	6.64	124.32	121.00
34	BA	1739	G	C6-N1-C2	-6.64	121.11	125.10
35	BB	48	G	P-O3'-C3'	6.64	127.67	119.70
35	BB	1293	C	N1-C2-O2	6.64	122.89	118.90
85	AA	730	G	N3-C4-C5	-6.64	125.28	128.60
85	AA	731	U	O3'-P-O5'	-6.64	91.38	104.00
85	AA	1934	A	P-O5'-C5'	6.64	131.53	120.90
1	A0	101	GLN	C-N-CA	6.64	138.30	121.70
4	A3	212	TYR	CB-CG-CD1	-6.64	117.02	121.00
34	BA	862	C	C1'-O4'-C4'	-6.64	104.59	109.90
35	BB	1094	A	C4-N9-C1'	-6.64	114.35	126.30
41	BH	4	U	C4'-C3'-C2'	6.64	109.24	102.60
41	BH	77	G	O5'-P-OP2	-6.64	99.72	105.70
85	AA	186	U	P-O5'-C5'	-6.64	110.28	120.90
85	AA	1396	C	O4'-C1'-N1	6.64	113.51	108.20
85	AA	1505	G	C4-N9-C1'	-6.64	117.87	126.50
34	BA	542	A	O4'-C1'-N9	6.64	113.51	108.20
34	BA	560	U	O4'-C1'-N1	6.64	113.51	108.20
34	BA	817	U	C1'-O4'-C4'	-6.64	104.59	109.90
34	BA	853	A	N1-C6-N6	-6.64	114.62	118.60
34	BA	1149	C	C5-C4-N4	6.64	124.85	120.20
34	BA	1186	U	C5-C6-N1	6.64	126.02	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1484	A	C5-C6-N1	6.64	121.02	117.70
34	BA	1538	G	C5-C6-O6	-6.64	124.62	128.60
34	BA	1650	G	C1'-O4'-C4'	-6.64	104.59	109.90
35	BB	1497	C	N3-C2-O2	-6.64	117.25	121.90
38	BE	22	A	O4'-C1'-N9	6.64	113.51	108.20
40	BG	64	C	O4'-C1'-N1	6.64	113.51	108.20
85	AA	308	U	C5'-C4'-O4'	6.64	117.07	109.10
85	AA	788	G	C4-N9-C1'	-6.64	117.87	126.50
85	AA	856	G	O4'-C4'-C3'	-6.64	97.36	104.00
85	AA	941	C	N3-C2-O2	-6.64	117.25	121.90
85	AA	2158	U	C5'-C4'-C3'	-6.64	105.38	116.00
7	A6	2	ARG	NE-CZ-NH1	6.64	123.62	120.30
34	BA	280	A	C5'-C4'-C3'	-6.64	105.38	116.00
34	BA	518	C	P-O3'-C3'	6.64	127.67	119.70
34	BA	1305	A	C5'-C4'-C3'	-6.64	105.38	116.00
34	BA	1416	C	C4'-C3'-C2'	6.64	109.24	102.60
34	BA	1534	U	O5'-P-OP1	-6.64	99.73	105.70
78	Bs	46	ARG	NE-CZ-NH1	6.64	123.62	120.30
85	AA	493	A	N1-C6-N6	-6.64	114.62	118.60
85	AA	1976	G	C1'-O4'-C4'	-6.64	104.59	109.90
34	BA	494	A	O4'-C1'-N9	6.64	113.51	108.20
34	BA	867	C	N3-C2-O2	-6.64	117.25	121.90
35	BB	52	G	C4'-C3'-C2'	6.64	109.24	102.60
35	BB	488	G	C4-N9-C1'	-6.64	117.87	126.50
35	BB	1185	G	C1'-O4'-C4'	-6.64	104.59	109.90
38	BE	139	U	C4'-C3'-C2'	-6.64	95.96	102.60
40	BG	1	G	C1'-O4'-C4'	-6.64	104.59	109.90
41	BH	102	C	N3-C4-C5	6.64	124.56	121.90
81	Bv	179	PHE	N-CA-CB	6.64	122.55	110.60
85	AA	649	C	C3'-C2'-C1'	-6.64	96.19	101.50
85	AA	919	U	O3'-P-O5'	6.64	116.61	104.00
85	AA	1288	A	C8-N9-C4	-6.64	103.14	105.80
85	AA	2201	A	O4'-C1'-N9	6.64	113.51	108.20
23	AP	87	ARG	N-CA-CB	6.63	122.54	110.60
34	BA	688	G	C1'-O4'-C4'	-6.63	104.59	109.90
34	BA	1611	A	C5'-C4'-C3'	-6.63	105.38	116.00
35	BB	81	A	P-O5'-C5'	6.63	131.51	120.90
35	BB	713	U	C2'-C3'-O3'	6.63	124.31	113.70
36	BC	65	G	C1'-O4'-C4'	-6.63	104.59	109.90
47	BN	44	ARG	N-CA-CB	-6.63	98.66	110.60
85	AA	335	G	C4-N9-C1'	-6.63	117.88	126.50
85	AA	368	C	N1-C2-N3	6.63	123.84	119.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	451	G	C5'-C4'-C3'	-6.63	105.39	116.00
85	AA	867	G	C1'-O4'-C4'	-6.63	104.59	109.90
85	AA	1384	C	O4'-C1'-N1	6.63	113.51	108.20
85	AA	2085	C	O5'-C5'-C4'	-6.63	99.09	111.70
34	BA	1472	G	N7-C8-N9	-6.63	109.78	113.10
34	BA	1798	G	O4'-C1'-N9	6.63	113.51	108.20
85	AA	1634	U	N1-C2-O2	6.63	127.44	122.80
15	AG	69	ARG	NE-CZ-NH1	6.63	123.62	120.30
34	BA	60	A	C5-C6-N6	-6.63	118.39	123.70
34	BA	90	G	O4'-C1'-N9	6.63	113.51	108.20
34	BA	103	G	N3-C2-N2	6.63	124.54	119.90
34	BA	899	G	C3'-C2'-C1'	-6.63	96.19	101.50
34	BA	1137	U	C2-N1-C1'	-6.63	109.74	117.70
34	BA	1161	G	N3-C2-N2	6.63	124.54	119.90
34	BA	1282	G	O5'-P-OP2	-6.63	99.73	105.70
34	BA	1412	G	C5'-C4'-C3'	-6.63	105.39	116.00
34	BA	1699	A	O3'-P-O5'	6.63	116.60	104.00
35	BB	486	G	C5'-C4'-C3'	6.63	126.61	116.00
35	BB	660	G	P-O3'-C3'	6.63	127.66	119.70
35	BB	1348	C	C3'-C2'-C1'	-6.63	96.19	101.50
40	BG	151	A	C1'-O4'-C4'	-6.63	104.59	109.90
65	Bf	351	GLN	N-CA-CB	6.63	122.54	110.60
85	AA	126	U	C4'-C3'-C2'	-6.63	95.97	102.60
85	AA	678	A	O3'-P-O5'	-6.63	91.40	104.00
85	AA	1076	U	O4'-C1'-N1	6.63	113.50	108.20
34	BA	103	G	N9-C1'-C2'	-6.63	104.71	112.00
41	BH	76	G	N1-C6-O6	-6.63	115.92	119.90
85	AA	1496	U	N3-C2-O2	-6.63	117.56	122.20
34	BA	1345	U	O4'-C1'-N1	6.63	113.50	108.20
34	BA	1573	C	C4'-C3'-C2'	6.63	109.23	102.60
35	BB	901	U	O4'-C4'-C3'	-6.63	97.37	104.00
38	BE	146	U	C2-N1-C1'	-6.63	109.75	117.70
39	BF	60	C	C5'-C4'-O4'	6.63	117.05	109.10
54	BU	116	ARG	NE-CZ-NH1	6.63	123.61	120.30
57	BX	125	TYR	CA-CB-CG	-6.63	100.81	113.40
62	Bc	26	ARG	NE-CZ-NH1	6.63	123.61	120.30
85	AA	273	C	C1'-O4'-C4'	-6.63	104.60	109.90
85	AA	1114	A	C4-C5-C6	-6.63	113.69	117.00
85	AA	1210	U	C4-C5-C6	-6.63	115.72	119.70
85	AA	1516	A	C5'-C4'-O4'	6.63	117.05	109.10
85	AA	1934	A	C5-C6-N1	6.63	121.01	117.70
34	BA	489	A	C1'-O4'-C4'	-6.63	104.60	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	784	C	C5'-C4'-C3'	6.63	126.60	116.00
34	BA	1035	A	P-O3'-C3'	6.63	127.65	119.70
35	BB	119	G	C4-N9-C1'	-6.63	117.88	126.50
35	BB	1024	G	C8-N9-C1'	6.63	135.62	127.00
37	BD	75	G	P-O3'-C3'	6.63	127.65	119.70
85	AA	974	U	P-O5'-C5'	6.63	131.50	120.90
85	AA	1794	U	C2-N1-C1'	-6.63	109.75	117.70
86	AB	55	U	P-O3'-C3'	-6.63	111.75	119.70
3	A2	57	ARG	NE-CZ-NH1	6.62	123.61	120.30
34	BA	180	G	N1-C6-O6	6.62	123.88	119.90
34	BA	527	C	C2-N3-C4	-6.62	116.59	119.90
34	BA	847	U	C6-N1-C2	-6.62	117.03	121.00
35	BB	857	G	O3'-P-O5'	6.62	116.59	104.00
35	BB	1346	A	C5-C6-N1	6.62	121.01	117.70
35	BB	1480	G	N1-C2-N3	-6.62	119.92	123.90
36	BC	169	G	O4'-C1'-N9	6.62	113.50	108.20
37	BD	48	G	N1-C2-N3	-6.62	119.92	123.90
85	AA	739	C	O4'-C1'-N1	6.62	113.50	108.20
34	BA	159	U	C6-N1-C2	-6.62	117.03	121.00
34	BA	573	U	C6-N1-C2	-6.62	117.03	121.00
34	BA	590	U	C3'-C2'-C1'	-6.62	96.20	101.50
34	BA	1041	U	O5'-C5'-C4'	-6.62	99.12	111.70
34	BA	1412	G	O4'-C1'-N9	6.62	113.50	108.20
35	BB	4	C	O5'-P-OP1	6.62	118.65	110.70
35	BB	1053	G	N3-C4-C5	-6.62	125.29	128.60
35	BB	1383	C	P-O3'-C3'	-6.62	111.75	119.70
35	BB	1484	A	P-O3'-C3'	-6.62	111.75	119.70
41	BH	58	C	C2-N1-C1'	-6.62	111.51	118.80
45	BL	138	LEU	C-N-CA	6.62	136.21	122.30
85	AA	1687	U	C3'-C2'-C1'	-6.62	96.20	101.50
86	AB	68	C	C6-N1-C1'	6.62	128.75	120.80
26	AS	107	ARG	NE-CZ-NH2	-6.62	116.99	120.30
34	BA	250	G	OP1-P-OP2	-6.62	109.67	119.60
34	BA	1371	U	C5'-C4'-C3'	-6.62	105.41	116.00
35	BB	481	A	C5'-C4'-O4'	6.62	117.05	109.10
35	BB	912	C	O4'-C1'-N1	6.62	113.50	108.20
35	BB	916	U	O4'-C1'-N1	6.62	113.50	108.20
35	BB	1476	C	C2-N1-C1'	-6.62	111.52	118.80
35	BB	1477	C	N3-C2-O2	-6.62	117.27	121.90
37	BD	48	G	N9-C1'-C2'	-6.62	104.72	112.00
85	AA	469	G	C4-N9-C1'	-6.62	117.89	126.50
85	AA	568	C	C6-N1-C1'	6.62	128.75	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1280	U	O4'-C1'-N1	6.62	113.50	108.20
2	A1	110	ARG	CB-CA-C	-6.62	97.16	110.40
34	BA	1832	A	C1'-O4'-C4'	-6.62	104.60	109.90
77	Br	216	THR	N-CA-CB	6.62	122.88	110.30
85	AA	109	G	P-O3'-C3'	-6.62	111.76	119.70
85	AA	572	G	C4'-C3'-C2'	-6.62	95.98	102.60
85	AA	1649	U	P-O3'-C3'	6.62	127.64	119.70
85	AA	2167	A	C8-N9-C1'	6.62	139.62	127.70
34	BA	136	A	C1'-O4'-C4'	-6.62	104.61	109.90
34	BA	243	C	C6-N1-C2	-6.62	117.65	120.30
34	BA	260	A	C1'-O4'-C4'	-6.62	104.61	109.90
34	BA	576	C	C2-N1-C1'	6.62	126.08	118.80
34	BA	909	G	C5-C6-N1	6.62	114.81	111.50
34	BA	1084	A	O5'-C5'-C4'	6.62	124.28	111.70
34	BA	1354	G	N1-C6-O6	6.62	123.87	119.90
34	BA	1563	G	C5'-C4'-C3'	-6.62	105.41	116.00
39	BF	44	C	O4'-C1'-N1	6.62	113.50	108.20
40	BG	140	G	C8-N9-C4	6.62	109.05	106.40
85	AA	38	C	O4'-C1'-N1	6.62	113.50	108.20
85	AA	171	U	C2'-C3'-O3'	6.62	124.29	113.70
85	AA	207	G	N7-C8-N9	6.62	116.41	113.10
85	AA	853	G	O5'-P-OP1	6.62	118.64	110.70
85	AA	863	C	N1-C2-N3	6.62	123.83	119.20
85	AA	1252	A	O4'-C1'-N9	6.62	113.50	108.20
85	AA	1495	G	C4-N9-C1'	-6.62	117.90	126.50
85	AA	2015	U	P-O5'-C5'	-6.62	110.31	120.90
34	BA	1146	U	C1'-O4'-C4'	-6.62	104.61	109.90
35	BB	509	A	C5-C6-N6	-6.62	118.41	123.70
59	BZ	111	ARG	NE-CZ-NH2	-6.62	116.99	120.30
66	Bg	107	GLY	N-CA-C	-6.62	96.56	113.10
85	AA	577	U	C6-N1-C2	-6.62	117.03	121.00
5	A4	75	LEU	N-CA-CB	-6.62	97.17	110.40
34	BA	59	A	C5'-C4'-O4'	6.62	117.04	109.10
34	BA	117	C	C5'-C4'-O4'	6.62	117.04	109.10
34	BA	253	U	P-O3'-C3'	-6.62	111.76	119.70
34	BA	1503	U	C5-C6-N1	-6.62	119.39	122.70
35	BB	4	C	C5'-C4'-C3'	6.62	126.58	116.00
35	BB	95	A	C3'-C2'-C1'	-6.62	96.21	101.50
35	BB	1541	G	N3-C2-N2	6.62	124.53	119.90
36	BC	66	G	C5-C6-O6	-6.62	124.63	128.60
40	BG	73	U	C2-N1-C1'	-6.62	109.76	117.70
66	Bg	70	TYR	CB-CG-CD1	-6.62	117.03	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	438	G	C1'-O4'-C4'	-6.62	104.61	109.90
85	AA	1258	U	P-O5'-C5'	6.62	131.49	120.90
34	BA	51	C	P-O3'-C3'	-6.61	111.77	119.70
34	BA	103	G	N1-C6-O6	-6.61	115.93	119.90
34	BA	114	U	P-O3'-C3'	6.61	127.64	119.70
34	BA	125	G	C5-C6-N1	6.61	114.81	111.50
34	BA	489	A	O4'-C4'-C3'	-6.61	97.39	104.00
34	BA	1126	U	C5'-C4'-C3'	6.61	126.58	116.00
34	BA	1239	G	N1-C6-O6	-6.61	115.93	119.90
35	BB	1224	C	C5'-C4'-O4'	6.61	117.04	109.10
38	BE	97	G	P-O5'-C5'	-6.61	110.32	120.90
38	BE	109	C	P-O5'-C5'	6.61	131.48	120.90
40	BG	182	G	O4'-C1'-N9	6.61	113.49	108.20
41	BH	120	C	P-O5'-C5'	-6.61	110.32	120.90
66	Bg	34	TYR	CB-CG-CD1	-6.61	117.03	121.00
73	Bn	66	TYR	N-CA-CB	-6.61	98.69	110.60
85	AA	1944	C	O4'-C1'-N1	6.61	113.49	108.20
35	BB	728	A	C8-N9-C4	-6.61	103.16	105.80
40	BG	129	G	N9-C1'-C2'	-6.61	104.73	112.00
85	AA	150	U	C2-N1-C1'	-6.61	109.77	117.70
85	AA	1363	U	C4'-C3'-C2'	-6.61	95.99	102.60
35	BB	513	G	C4-N9-C1'	-6.61	117.91	126.50
35	BB	976	U	C1'-O4'-C4'	-6.61	104.61	109.90
35	BB	1440	A	C8-N9-C4	6.61	108.44	105.80
45	BL	64	PHE	CB-CG-CD2	-6.61	116.17	120.80
59	BZ	57	ARG	NE-CZ-NH1	6.61	123.61	120.30
85	AA	68	A	P-O5'-C5'	-6.61	110.32	120.90
85	AA	1058	G	O4'-C1'-N9	6.61	113.49	108.20
85	AA	2237	G	C4-N9-C1'	-6.61	117.91	126.50
34	BA	135	G	C1'-O4'-C4'	-6.61	104.61	109.90
34	BA	374	U	C1'-O4'-C4'	-6.61	104.61	109.90
34	BA	501	U	C4'-C3'-C2'	-6.61	95.99	102.60
34	BA	1379	G	C5-C6-O6	-6.61	124.64	128.60
34	BA	1649	A	N9-C1'-C2'	6.61	122.59	114.00
34	BA	1736	A	P-O5'-C5'	6.61	131.47	120.90
35	BB	1096	G	C6-N1-C2	-6.61	121.14	125.10
36	BC	130	U	N3-C2-O2	-6.61	117.57	122.20
37	BD	112	U	C6-N1-C1'	6.61	130.45	121.20
40	BG	115	C	C1'-O4'-C4'	-6.61	104.61	109.90
85	AA	1776	C	O4'-C1'-N1	6.61	113.49	108.20
34	BA	69	C	N3-C4-N4	-6.61	113.37	118.00
34	BA	383	G	N1-C6-O6	6.61	123.86	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	127	U	C2-N3-C4	-6.61	123.04	127.00
35	BB	1151	A	N9-C4-C5	6.61	108.44	105.80
35	BB	1162	A	C4-N9-C1'	6.61	138.19	126.30
35	BB	1508	G	N9-C1'-C2'	-6.61	104.73	112.00
85	AA	354	C	P-O5'-C5'	-6.61	110.33	120.90
85	AA	930	G	O4'-C1'-N9	6.61	113.48	108.20
85	AA	1480	C	N1-C1'-C2'	-6.61	104.73	112.00
85	AA	1675	U	O4'-C1'-N1	6.61	113.49	108.20
86	AB	11	C	O4'-C1'-N1	6.61	113.49	108.20
34	BA	9	A	C1'-O4'-C4'	-6.61	104.61	109.90
34	BA	571	G	C5'-C4'-C3'	6.61	126.57	116.00
36	BC	139	A	N7-C8-N9	6.61	117.10	113.80
85	AA	273	C	O4'-C1'-N1	6.61	113.48	108.20
85	AA	292	C	C5'-C4'-C3'	6.61	126.57	116.00
85	AA	814	G	N9-C1'-C2'	-6.61	104.73	112.00
85	AA	1229	G	N1-C6-O6	-6.61	115.94	119.90
85	AA	1238	U	O4'-C1'-N1	6.61	113.48	108.20
86	AB	24	G	O4'-C1'-N9	6.61	113.48	108.20
34	BA	567	U	C5'-C4'-C3'	-6.60	105.43	116.00
34	BA	1377	A	O4'-C1'-N9	6.60	113.48	108.20
35	BB	710	A	C4-N9-C1'	-6.60	114.41	126.30
35	BB	795	A	N1-C6-N6	-6.60	114.64	118.60
38	BE	17	U	N1-C1'-C2'	-6.60	104.74	112.00
83	Bx	205	ARG	NE-CZ-NH1	6.60	123.60	120.30
85	AA	259	A	C4-N9-C1'	-6.60	114.41	126.30
85	AA	436	G	O4'-C1'-N9	6.60	113.48	108.20
34	BA	460	G	C8-N9-C1'	6.60	135.58	127.00
34	BA	686	U	C6-N1-C2	-6.60	117.04	121.00
34	BA	1171	C	C6-N1-C2	-6.60	117.66	120.30
34	BA	1699	A	C4'-C3'-O3'	6.60	126.20	113.00
35	BB	775	U	P-O5'-C5'	6.60	131.47	120.90
35	BB	859	U	C1'-O4'-C4'	-6.60	104.62	109.90
36	BC	122	A	C5-C6-N6	-6.60	118.42	123.70
37	BD	42	A	C5'-C4'-C3'	-6.60	105.44	116.00
40	BG	176	G	P-O5'-C5'	6.60	131.46	120.90
50	BQ	137	TRP	CB-CG-CD2	-6.60	118.02	126.60
73	Bn	57	ARG	NE-CZ-NH2	-6.60	117.00	120.30
85	AA	993	G	C5'-C4'-C3'	-6.60	105.44	116.00
85	AA	1055	U	O4'-C1'-N1	6.60	113.48	108.20
85	AA	1454	U	N3-C2-O2	-6.60	117.58	122.20
85	AA	1806	C	C2-N1-C1'	6.60	126.06	118.80
85	AA	1905	A	C6-N1-C2	-6.60	114.64	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	3	C	N3-C4-N4	6.60	122.62	118.00
32	AY	18	THR	N-CA-CB	6.60	122.84	110.30
35	BB	1290	C	C3'-C2'-C1'	-6.60	96.22	101.50
85	AA	271	A	C3'-C2'-C1'	-6.60	96.22	101.50
85	AA	851	G	P-O5'-C5'	-6.60	110.34	120.90
85	AA	1183	C	C6-N1-C1'	-6.60	112.88	120.80
85	AA	1713	A	N1-C6-N6	6.60	122.56	118.60
85	AA	1719	C	C6-N1-C2	-6.60	117.66	120.30
34	BA	136	A	C5'-C4'-C3'	-6.60	105.44	116.00
34	BA	327	G	C6-N1-C2	-6.60	121.14	125.10
34	BA	869	C	P-O5'-C5'	-6.60	110.34	120.90
35	BB	653	G	N9-C1'-C2'	-6.60	104.74	112.00
35	BB	1134	G	C5-N7-C8	-6.60	101.00	104.30
35	BB	1332	G	O4'-C1'-N9	6.60	113.48	108.20
35	BB	1533	U	C6-N1-C1'	-6.60	111.96	121.20
40	BG	159	A	C5-C6-N1	6.60	121.00	117.70
43	BJ	57	ARG	NE-CZ-NH2	-6.60	117.00	120.30
45	BL	159	HIS	CA-CB-CG	-6.60	102.38	113.60
85	AA	185	A	N1-C6-N6	6.60	122.56	118.60
85	AA	1109	G	C4'-C3'-C2'	6.60	109.20	102.60
85	AA	2124	G	C5-C6-O6	-6.60	124.64	128.60
12	AD	72	TYR	CB-CG-CD2	-6.60	117.04	121.00
34	BA	295	G	C5'-C4'-C3'	-6.60	105.44	116.00
34	BA	877	U	C5-C6-N1	-6.60	119.40	122.70
34	BA	1170	A	C3'-C2'-C1'	-6.60	96.22	101.50
35	BB	1092	G	P-O5'-C5'	6.60	131.46	120.90
38	BE	147	G	C5-C6-N1	6.60	114.80	111.50
38	BE	193	A	N7-C8-N9	6.60	117.10	113.80
40	BG	50	G	C8-N9-C4	6.60	109.04	106.40
40	BG	139	U	C5-C4-O4	6.60	129.86	125.90
41	BH	29	G	C2-N3-C4	-6.60	108.60	111.90
85	AA	318	A	O4'-C1'-N9	-6.60	102.92	108.20
85	AA	523	U	C5-C4-O4	-6.60	121.94	125.90
85	AA	2125	A	C4'-C3'-C2'	6.60	109.20	102.60
6	A5	204	GLU	C-N-CA	6.60	136.15	122.30
35	BB	458	U	C1'-O4'-C4'	-6.60	104.62	109.90
41	BH	44	A	C5-C6-N6	-6.60	118.42	123.70
85	AA	59	C	N3-C4-N4	6.60	122.62	118.00
85	AA	527	A	C8-N9-C4	6.60	108.44	105.80
85	AA	1134	G	C5-C6-N1	6.60	114.80	111.50
85	AA	1551	G	C8-N9-C1'	-6.60	118.42	127.00
35	BB	135	C	C2-N1-C1'	-6.59	111.55	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	713	U	C2-N1-C1'	-6.59	109.79	117.70
35	BB	1448	U	O5'-C5'-C4'	-6.59	99.17	111.70
83	Bx	265	ARG	NE-CZ-NH1	6.59	123.60	120.30
85	AA	315	U	C3'-C2'-C1'	-6.59	96.22	101.50
85	AA	523	U	N3-C4-O4	6.59	124.02	119.40
85	AA	803	C	O4'-C1'-N1	6.59	113.48	108.20
85	AA	1856	G	P-O3'-C3'	-6.59	111.79	119.70
85	AA	2248	A	P-O3'-C3'	6.59	127.61	119.70
34	BA	106	U	P-O3'-C3'	-6.59	111.79	119.70
35	BB	599	U	C5'-C4'-O4'	6.59	117.01	109.10
40	BG	13	A	P-O5'-C5'	-6.59	110.35	120.90
41	BH	9	C	C6-N1-C2	-6.59	117.66	120.30
41	BH	99	G	O4'-C1'-N9	6.59	113.47	108.20
85	AA	815	G	C1'-O4'-C4'	-6.59	104.63	109.90
85	AA	818	C	C4-C5-C6	-6.59	114.10	117.40
34	BA	798	G	O3'-P-O5'	6.59	116.53	104.00
34	BA	1049	G	C5-C6-O6	-6.59	124.64	128.60
34	BA	1735	G	C5-C6-N1	6.59	114.80	111.50
35	BB	496	C	C3'-C2'-C1'	-6.59	96.23	101.50
35	BB	1446	C	O4'-C1'-C2'	6.59	113.53	107.60
40	BG	95	U	C6-N1-C1'	6.59	130.43	121.20
85	AA	939	A	C4'-C3'-C2'	-6.59	96.01	102.60
34	BA	537	C	C6-N1-C1'	6.59	128.71	120.80
34	BA	594	G	C5'-C4'-C3'	6.59	126.54	116.00
34	BA	1072	U	N1-C1'-C2'	-6.59	104.75	112.00
35	BB	80	C	C5-C4-N4	6.59	124.81	120.20
35	BB	893	U	N1-C1'-C2'	-6.59	104.75	112.00
35	BB	1464	G	P-O5'-C5'	6.59	131.44	120.90
37	BD	26	C	C5'-C4'-O4'	6.59	117.01	109.10
38	BE	119	U	C6-N1-C2	-6.59	117.05	121.00
39	BF	36	G	C4-N9-C1'	-6.59	117.93	126.50
47	BN	74	ARG	NE-CZ-NH2	-6.59	117.01	120.30
85	AA	568	C	N3-C2-O2	-6.59	117.29	121.90
85	AA	650	G	P-O3'-C3'	6.59	127.61	119.70
85	AA	690	G	N1-C2-N2	-6.59	110.27	116.20
85	AA	941	C	OP1-P-OP2	-6.59	109.72	119.60
85	AA	1004	G	N7-C8-N9	6.59	116.39	113.10
85	AA	2100	A	C6-N1-C2	-6.59	114.65	118.60
85	AA	2107	C	P-O5'-C5'	6.59	131.44	120.90
85	AA	2227	A	C4-N9-C1'	-6.59	114.44	126.30
34	BA	185	A	C1'-O4'-C4'	-6.59	104.63	109.90
34	BA	1502	G	C5-C6-O6	-6.59	124.65	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1798	G	OP1-P-OP2	-6.59	109.72	119.60
35	BB	1328	C	O4'-C1'-N1	6.59	113.47	108.20
36	BC	155	C	C5'-C4'-C3'	-6.59	105.46	116.00
44	BK	75	TYR	CB-CG-CD1	-6.59	117.05	121.00
49	BP	163	ARG	NE-CZ-NH2	-6.59	117.01	120.30
81	Bv	176	HIS	CA-CB-CG	-6.59	102.40	113.60
85	AA	879	G	C8-N9-C4	6.59	109.03	106.40
85	AA	1211	C	N1-C1'-C2'	-6.59	104.75	112.00
34	BA	134	U	C2-N1-C1'	-6.59	109.80	117.70
34	BA	817	U	C3'-C2'-C1'	-6.59	96.23	101.50
34	BA	1070	G	C5-N7-C8	-6.59	101.01	104.30
34	BA	1608	C	P-O3'-C3'	-6.59	111.80	119.70
34	BA	1609	U	C2'-C3'-O3'	6.59	124.24	113.70
34	BA	1722	U	P-O3'-C3'	-6.59	111.80	119.70
35	BB	420	U	C3'-C2'-C1'	-6.59	96.23	101.50
35	BB	1487	G	C5-C6-N1	6.59	114.79	111.50
38	BE	128	G	C4'-C3'-C2'	6.59	109.19	102.60
39	BF	13	U	C4'-C3'-C2'	-6.59	96.01	102.60
40	BG	20	U	C2-N1-C1'	-6.59	109.80	117.70
41	BH	127	A	C6-N1-C2	-6.59	114.65	118.60
51	BR	30	PHE	CB-CG-CD1	6.59	125.41	120.80
85	AA	19	A	P-O5'-C5'	6.59	131.44	120.90
85	AA	117	C	C3'-C2'-C1'	-6.59	96.23	101.50
85	AA	252	G	N9-C1'-C2'	6.59	122.56	114.00
85	AA	557	G	N9-C1'-C2'	-6.59	104.75	112.00
85	AA	650	G	C1'-O4'-C4'	-6.59	104.63	109.90
85	AA	1471	G	C8-N9-C4	6.59	109.03	106.40
85	AA	1661	U	N1-C2-N3	6.59	118.85	114.90
34	BA	736	G	O5'-P-OP2	6.58	118.60	110.70
34	BA	1156	U	C5-C4-O4	6.58	129.85	125.90
34	BA	1724	G	O3'-P-O5'	6.58	116.51	104.00
35	BB	1133	C	C2-N1-C1'	-6.58	111.56	118.80
85	AA	48	G	N1-C6-O6	-6.58	115.95	119.90
34	BA	858	C	N1-C2-O2	6.58	122.85	118.90
35	BB	29	C	C5'-C4'-C3'	6.58	126.53	116.00
35	BB	417	A	C5-C6-N6	-6.58	118.43	123.70
35	BB	498	G	C4-N9-C1'	-6.58	117.94	126.50
35	BB	1442	C	P-O3'-C3'	-6.58	111.80	119.70
35	BB	1448	U	N1-C2-N3	6.58	118.85	114.90
37	BD	73	U	C2-N3-C4	-6.58	123.05	127.00
38	BE	27	A	N9-C1'-C2'	-6.58	104.76	112.00
85	AA	701	C	C5'-C4'-C3'	6.58	126.53	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1167	G	C8-N9-C4	-6.58	103.77	106.40
85	AA	1259	U	O4'-C4'-C3'	-6.58	97.42	104.00
85	AA	1406	U	P-O3'-C3'	6.58	127.60	119.70
85	AA	1938	G	C4-N9-C1'	-6.58	117.94	126.50
21	AM	84	PHE	CB-CG-CD2	-6.58	116.19	120.80
23	AP	82	ARG	NE-CZ-NH1	6.58	123.59	120.30
34	BA	130	U	O5'-C5'-C4'	-6.58	99.19	111.70
34	BA	817	U	P-O3'-C3'	-6.58	111.80	119.70
34	BA	1072	U	C3'-C2'-C1'	-6.58	96.24	101.50
34	BA	1406	U	O4'-C1'-N1	6.58	113.47	108.20
35	BB	598	C	C1'-O4'-C4'	-6.58	104.64	109.90
35	BB	893	U	P-O3'-C3'	6.58	127.60	119.70
38	BE	198	A	C5-C6-N1	6.58	120.99	117.70
85	AA	400	G	C8-N9-C1'	6.58	135.56	127.00
85	AA	483	G	C8-N9-C1'	-6.58	118.44	127.00
85	AA	720	A	C3'-C2'-C1'	-6.58	96.23	101.50
85	AA	1270	C	C5-C6-N1	-6.58	117.71	121.00
85	AA	1787	G	C8-N9-C1'	6.58	135.56	127.00
34	BA	1705	C	P-O3'-C3'	6.58	127.60	119.70
35	BB	400	C	O4'-C1'-N1	6.58	113.46	108.20
35	BB	1425	A	C4-N9-C1'	-6.58	114.46	126.30
37	BD	19	C	C1'-O4'-C4'	-6.58	104.64	109.90
40	BG	17	A	N1-C6-N6	6.58	122.55	118.60
52	BS	42	ARG	NE-CZ-NH2	-6.58	117.01	120.30
52	BS	119	ARG	NE-CZ-NH1	6.58	123.59	120.30
85	AA	1162	A	O4'-C1'-C2'	6.58	113.52	107.60
85	AA	1283	C	N3-C2-O2	-6.58	117.29	121.90
22	AO	60	ARG	NE-CZ-NH2	-6.58	117.01	120.30
27	AT	117	ARG	N-CA-CB	6.58	122.44	110.60
34	BA	522	C	N1-C1'-C2'	-6.58	104.76	112.00
34	BA	1227	U	C2-N3-C4	-6.58	123.05	127.00
34	BA	1656	A	C4-C5-C6	-6.58	113.71	117.00
35	BB	104	G	O4'-C1'-N9	6.58	113.46	108.20
35	BB	481	A	C1'-O4'-C4'	-6.58	104.64	109.90
35	BB	794	G	N1-C6-O6	-6.58	115.95	119.90
35	BB	1098	G	C8-N9-C4	6.58	109.03	106.40
35	BB	1488	G	N3-C2-N2	6.58	124.50	119.90
36	BC	14	G	P-O3'-C3'	-6.58	111.81	119.70
36	BC	136	G	C8-N9-C1'	6.58	135.55	127.00
40	BG	4	A	N9-C4-C5	6.58	108.43	105.80
44	BK	157	MET	CG-SD-CE	6.58	110.72	100.20
85	AA	164	G	C1'-O4'-C4'	-6.58	104.64	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	411	U	C2-N1-C1'	-6.58	109.81	117.70
85	AA	1202	G	N3-C2-N2	6.58	124.50	119.90
85	AA	1676	G	C6-N1-C2	-6.58	121.15	125.10
85	AA	1725	G	P-O3'-C3'	6.58	127.59	119.70
85	AA	2218	G	C8-N9-C4	6.58	109.03	106.40
34	BA	252	A	C4'-C3'-C2'	-6.58	96.02	102.60
35	BB	1337	C	N3-C2-O2	-6.58	117.30	121.90
36	BC	69	U	C4'-C3'-C2'	6.58	109.18	102.60
38	BE	155	C	O4'-C1'-C2'	-6.58	99.22	105.80
85	AA	1520	A	C4'-C3'-C2'	-6.58	96.02	102.60
2	A1	27	ARG	NE-CZ-NH1	6.58	123.59	120.30
34	BA	344	G	C5-C6-O6	-6.58	124.66	128.60
34	BA	668	G	C5-C6-O6	-6.58	124.66	128.60
34	BA	690	G	C5-C6-N1	6.58	114.79	111.50
34	BA	1073	G	N9-C1'-C2'	-6.58	104.77	112.00
34	BA	1724	G	C5-C6-O6	6.58	132.55	128.60
35	BB	661	G	P-O3'-C3'	-6.58	111.81	119.70
35	BB	709	G	P-O3'-C3'	6.58	127.59	119.70
35	BB	1129	C	C6-N1-C2	-6.58	117.67	120.30
35	BB	1371	G	C5'-C4'-C3'	-6.58	105.48	116.00
83	Bx	111	ALA	N-CA-CB	-6.58	100.89	110.10
85	AA	2050	C	O4'-C1'-C2'	-6.58	99.22	105.80
34	BA	1176	C	N3-C4-N4	-6.57	113.40	118.00
34	BA	1331	G	C5-C6-O6	-6.57	124.66	128.60
34	BA	1585	A	C2'-C3'-O3'	6.57	124.22	113.70
34	BA	1593	U	N1-C2-N3	6.57	118.84	114.90
35	BB	582	G	P-O3'-C3'	-6.57	111.81	119.70
35	BB	1100	C	O4'-C1'-N1	6.57	113.46	108.20
35	BB	1460	G	P-O3'-C3'	6.57	127.59	119.70
38	BE	100	U	O3'-P-O5'	6.57	116.49	104.00
40	BG	137	G	P-O3'-C3'	-6.57	111.81	119.70
65	Bf	111	VAL	CB-CA-C	-6.57	98.91	111.40
85	AA	1574	C	C4'-C3'-C2'	-6.57	96.03	102.60
85	AA	1649	U	C6-N1-C1'	6.57	130.40	121.20
85	AA	1903	G	C5'-C4'-C3'	-6.57	105.48	116.00
85	AA	2065	U	O4'-C1'-N1	6.57	113.46	108.20
85	AA	2093	U	O4'-C1'-N1	6.57	113.46	108.20
24	AQ	84	ARG	NE-CZ-NH1	6.57	123.59	120.30
34	BA	683	C	C6-N1-C1'	-6.57	112.91	120.80
39	BF	40	U	C2'-C3'-O3'	6.57	124.22	113.70
44	BK	119	TYR	C-N-CA	6.57	136.10	122.30
62	Bc	66	SER	N-CA-C	-6.57	93.26	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	26	C	O4'-C1'-N1	6.57	113.46	108.20
34	BA	421	G	N1-C6-O6	-6.57	115.96	119.90
34	BA	1500	G	P-O3'-C3'	-6.57	111.81	119.70
35	BB	23	U	O3'-P-O5'	6.57	116.48	104.00
35	BB	808	U	C5'-C4'-C3'	6.57	126.51	116.00
35	BB	1001	G	C1'-O4'-C4'	-6.57	104.64	109.90
35	BB	1236	A	C5-C6-N6	-6.57	118.44	123.70
35	BB	1288	G	C5-C6-O6	-6.57	124.66	128.60
37	BD	41	G	C8-N9-C1'	6.57	135.54	127.00
38	BE	172	U	N1-C2-N3	-6.57	110.96	114.90
85	AA	1007	G	O4'-C1'-N9	6.57	113.46	108.20
85	AA	1439	A	O4'-C1'-N9	6.57	113.46	108.20
85	AA	1462	A	OP1-P-OP2	-6.57	109.74	119.60
85	AA	1574	C	O4'-C1'-N1	6.57	113.46	108.20
85	AA	1842	C	C5'-C4'-C3'	6.57	126.51	116.00
34	BA	9	A	C3'-C2'-C1'	-6.57	96.25	101.50
34	BA	124	G	N9-C1'-C2'	-6.57	104.77	112.00
34	BA	544	U	C5'-C4'-C3'	-6.57	105.49	116.00
34	BA	1054	U	C4'-C3'-C2'	-6.57	96.03	102.60
34	BA	1476	G	C4-N9-C1'	-6.57	117.96	126.50
34	BA	1601	C	P-O5'-C5'	-6.57	110.39	120.90
40	BG	75	C	C2'-C3'-O3'	6.57	124.21	113.70
41	BH	32	U	N3-C4-C5	6.57	118.54	114.60
47	BN	114	MET	CG-SD-CE	6.57	110.71	100.20
54	BU	122	GLY	N-CA-C	-6.57	96.68	113.10
1	A0	210	ARG	NE-CZ-NH2	6.57	123.58	120.30
34	BA	12	G	O5'-C5'-C4'	-6.57	99.22	111.70
34	BA	479	U	O4'-C1'-N1	6.57	113.45	108.20
34	BA	511	U	C2-N1-C1'	6.57	125.58	117.70
35	BB	610	U	C6-N1-C2	-6.57	117.06	121.00
35	BB	710	A	C8-N9-C1'	6.57	139.52	127.70
38	BE	7	U	O3'-P-O5'	6.57	116.48	104.00
47	BN	69	ARG	CG-CD-NE	-6.57	98.01	111.80
54	BU	150	THR	CA-CB-CG2	6.57	121.59	112.40
85	AA	48	G	C5-C6-O6	6.57	132.54	128.60
85	AA	227	A	C5-C6-N6	6.57	128.95	123.70
85	AA	2071	U	C6-N1-C1'	6.57	130.39	121.20
85	AA	2238	C	C1'-O4'-C4'	-6.57	104.65	109.90
34	BA	863	G	O5'-P-OP1	-6.57	99.79	105.70
34	BA	1737	A	C5'-C4'-C3'	-6.57	105.50	116.00
35	BB	390	G	C4'-C3'-C2'	-6.57	96.03	102.60
35	BB	474	G	C4'-C3'-C2'	-6.57	96.03	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1022	C	N3-C2-O2	-6.57	117.30	121.90
35	BB	1251	G	P-O3'-C3'	-6.57	111.82	119.70
35	BB	1534	U	C6-N1-C1'	6.57	130.39	121.20
38	BE	117	A	C6-C5-N7	-6.57	127.70	132.30
40	BG	53	C	C1'-O4'-C4'	-6.57	104.65	109.90
41	BH	29	G	C1'-O4'-C4'	-6.57	104.65	109.90
73	Bn	49	TRP	CB-CG-CD2	-6.57	118.06	126.60
85	AA	241	U	C6-N1-C2	-6.57	117.06	121.00
85	AA	634	U	P-O3'-C3'	-6.57	111.82	119.70
85	AA	635	G	O5'-C5'-C4'	-6.57	99.23	111.70
85	AA	1469	G	N1-C2-N3	-6.57	119.96	123.90
2	A1	143	ARG	NE-CZ-NH2	-6.56	117.02	120.30
34	BA	1739	G	N1-C2-N2	6.56	122.11	116.20
35	BB	774	C	O4'-C1'-C2'	6.56	113.51	107.60
62	Bc	12	VAL	N-CA-C	6.56	128.72	111.00
5	A4	35	SER	CA-C-N	-6.56	102.76	117.20
34	BA	304	G	P-O5'-C5'	-6.56	110.40	120.90
34	BA	784	C	P-O3'-C3'	-6.56	111.83	119.70
34	BA	853	A	O4'-C1'-N9	6.56	113.45	108.20
34	BA	866	C	O5'-P-OP1	-6.56	99.79	105.70
34	BA	1265	G	N1-C6-O6	6.56	123.84	119.90
34	BA	1401	C	C3'-C2'-C1'	-6.56	96.25	101.50
34	BA	1404	A	C8-N9-C1'	6.56	139.51	127.70
34	BA	1505	G	C4-C5-N7	-6.56	108.17	110.80
34	BA	1818	A	P-O3'-C3'	-6.56	111.82	119.70
35	BB	369	A	O4'-C1'-N9	6.56	113.45	108.20
35	BB	706	G	C8-N9-C1'	6.56	135.53	127.00
35	BB	962	U	O4'-C1'-N1	6.56	113.45	108.20
35	BB	1393	C	C6-N1-C1'	-6.56	112.93	120.80
37	BD	97	U	C2'-C3'-O3'	6.56	124.20	113.70
38	BE	129	G	N1-C2-N2	-6.56	110.29	116.20
43	BJ	23	ARG	NE-CZ-NH1	6.56	123.58	120.30
85	AA	531	G	N1-C6-O6	-6.56	115.96	119.90
85	AA	1460	G	C6-N1-C2	-6.56	121.16	125.10
85	AA	1852	U	C2-N3-C4	-6.56	123.06	127.00
35	BB	843	G	C3'-C2'-C1'	-6.56	96.25	101.50
35	BB	1165	A	C4-N9-C1'	-6.56	114.49	126.30
35	BB	1186	A	C4-N9-C1'	-6.56	114.49	126.30
85	AA	1288	A	C3'-C2'-C1'	-6.56	96.25	101.50
85	AA	1862	C	C6-N1-C2	-6.56	117.68	120.30
85	AA	1864	G	C5'-C4'-O4'	6.56	116.97	109.10
85	AA	2170	G	C4-N9-C1'	-6.56	117.97	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	557	U	P-O3'-C3'	6.56	127.57	119.70
34	BA	824	C	C1'-O4'-C4'	-6.56	104.65	109.90
34	BA	908	G	N9-C1'-C2'	-6.56	104.78	112.00
34	BA	1696	G	OP2-P-O3'	6.56	119.63	105.20
35	BB	144	G	C6-N1-C2	-6.56	121.17	125.10
35	BB	311	C	C6-N1-C2	-6.56	117.68	120.30
35	BB	335	C	O4'-C1'-N1	6.56	113.45	108.20
35	BB	1027	U	C2-N3-C4	-6.56	123.06	127.00
35	BB	1117	G	C1'-O4'-C4'	-6.56	104.65	109.90
35	BB	1263	A	N1-C6-N6	-6.56	114.66	118.60
35	BB	1486	C	C5'-C4'-C3'	-6.56	105.51	116.00
37	BD	56	G	C5-C6-O6	-6.56	124.66	128.60
85	AA	523	U	C2-N3-C4	-6.56	123.06	127.00
85	AA	550	G	O4'-C1'-N9	6.56	113.45	108.20
85	AA	1156	A	P-O3'-C3'	-6.56	111.83	119.70
85	AA	2020	C	C2'-C3'-O3'	6.56	124.19	113.70
27	AT	74	GLY	C-N-CA	6.56	138.09	121.70
34	BA	1541	G	C4-C5-C6	-6.56	114.87	118.80
35	BB	611	U	C2-N1-C1'	-6.56	109.83	117.70
35	BB	1460	G	O4'-C1'-N9	6.56	113.45	108.20
36	BC	5	U	P-O3'-C3'	-6.56	111.83	119.70
36	BC	47	C	P-O3'-C3'	6.56	127.57	119.70
85	AA	1004	G	C3'-C2'-C1'	-6.56	96.25	101.50
34	BA	333	A	C5'-C4'-O4'	6.56	116.97	109.10
34	BA	599	U	O4'-C4'-C3'	-6.56	97.44	104.00
34	BA	898	G	O4'-C1'-C2'	-6.56	99.24	105.80
34	BA	1793	G	C8-N9-C4	-6.56	103.78	106.40
35	BB	676	G	C5-C6-O6	-6.56	124.67	128.60
35	BB	836	U	P-O3'-C3'	6.56	127.57	119.70
40	BG	54	G	C8-N9-C4	6.56	109.02	106.40
85	AA	436	G	P-O3'-C3'	-6.56	111.83	119.70
85	AA	1571	A	P-O3'-C3'	-6.56	111.83	119.70
34	BA	286	C	N1-C1'-C2'	-6.55	104.79	112.00
35	BB	9	G	C5-C6-O6	-6.55	124.67	128.60
35	BB	823	G	C1'-O4'-C4'	-6.55	104.66	109.90
35	BB	1363	A	C5'-C4'-C3'	-6.55	105.51	116.00
36	BC	6	G	N1-C6-O6	6.55	123.83	119.90
39	BF	35	C	C5-C4-N4	-6.55	115.61	120.20
41	BH	77	G	OP1-P-OP2	-6.55	109.77	119.60
77	Br	262	PHE	CA-CB-CG	-6.55	98.17	113.90
85	AA	486	G	C2'-C3'-O3'	6.55	124.19	113.70
85	AA	1637	C	C5'-C4'-C3'	-6.55	105.51	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	AM	125	ARG	NE-CZ-NH2	-6.55	117.02	120.30
34	BA	1196	C	C5'-C4'-C3'	-6.55	105.52	116.00
34	BA	1399	A	C5'-C4'-O4'	6.55	116.96	109.10
85	AA	249	C	O4'-C1'-N1	6.55	113.44	108.20
85	AA	283	A	C2-N3-C4	-6.55	107.32	110.60
85	AA	860	C	C5-C4-N4	6.55	124.79	120.20
85	AA	1211	C	N3-C4-C5	6.55	124.52	121.90
34	BA	697	A	P-O3'-C3'	-6.55	111.84	119.70
34	BA	1342	C	O4'-C1'-N1	6.55	113.44	108.20
34	BA	1486	U	O4'-C1'-N1	6.55	113.44	108.20
34	BA	1629	A	N1-C6-N6	-6.55	114.67	118.60
35	BB	1304	U	C1'-O4'-C4'	-6.55	104.66	109.90
39	BF	42	G	C8-N9-C4	-6.55	103.78	106.40
68	Bi	119	ARG	NE-CZ-NH1	6.55	123.58	120.30
85	AA	627	A	C5'-C4'-C3'	-6.55	105.52	116.00
34	BA	407	A	C5'-C4'-C3'	-6.55	105.52	116.00
34	BA	1232	C	C5'-C4'-C3'	-6.55	105.52	116.00
34	BA	1500	G	C5-N7-C8	6.55	107.57	104.30
35	BB	517	G	C8-N9-C1'	6.55	135.51	127.00
35	BB	611	U	O5'-P-OP2	-6.55	99.81	105.70
35	BB	1152	U	C3'-C2'-C1'	-6.55	96.26	101.50
36	BC	26	U	C5'-C4'-C3'	6.55	126.48	116.00
38	BE	201	A	N9-C4-C5	-6.55	103.18	105.80
40	BG	57	A	C5'-C4'-C3'	-6.55	105.52	116.00
85	AA	708	G	C6-N1-C2	-6.55	121.17	125.10
85	AA	1124	G	C1'-O4'-C4'	-6.55	104.66	109.90
85	AA	1380	U	O4'-C1'-N1	6.55	113.44	108.20
85	AA	1483	A	N1-C2-N3	-6.55	126.03	129.30
85	AA	2002	A	N3-C4-C5	-6.55	122.22	126.80
34	BA	1350	C	O4'-C1'-N1	6.55	113.44	108.20
34	BA	1700	C	O4'-C1'-N1	6.55	113.44	108.20
35	BB	765	G	O4'-C1'-N9	6.55	113.44	108.20
79	Bt	90	HIS	CA-CB-CG	-6.55	102.47	113.60
85	AA	2119	C	C5'-C4'-O4'	-6.55	101.24	109.10
31	AX	161	THR	CA-CB-CG2	-6.55	103.23	112.40
34	BA	204	U	P-O3'-C3'	6.55	127.56	119.70
34	BA	259	C	C6-N1-C2	-6.55	117.68	120.30
34	BA	523	A	C5-C6-N6	6.55	128.94	123.70
34	BA	707	C	C2-N3-C4	-6.55	116.63	119.90
34	BA	1551	G	C5-C6-N1	6.55	114.77	111.50
34	BA	1625	C	C1'-O4'-C4'	-6.55	104.66	109.90
35	BB	765	G	C8-N9-C1'	6.55	135.51	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	45	G	C5-C6-O6	-6.55	124.67	128.60
40	BG	30	C	C2-N1-C1'	-6.55	111.60	118.80
40	BG	90	G	N1-C6-O6	-6.55	115.97	119.90
70	Bk	59	MET	N-CA-CB	-6.55	98.82	110.60
85	AA	1226	A	N9-C1'-C2'	-6.55	104.80	112.00
85	AA	2020	C	C6-N1-C1'	-6.55	112.94	120.80
85	AA	2163	G	C3'-C2'-C1'	-6.55	96.26	101.50
4	A3	2	LYS	N-CA-CB	-6.54	98.82	110.60
34	BA	42	A	C8-N9-C1'	6.54	139.48	127.70
34	BA	734	G	C5-C6-N1	6.54	114.77	111.50
35	BB	28	G	C5-C6-O6	-6.54	124.67	128.60
35	BB	404	A	C5-N7-C8	-6.54	100.63	103.90
35	BB	898	U	P-O5'-C5'	6.54	131.37	120.90
38	BE	146	U	C6-N1-C1'	6.54	130.36	121.20
85	AA	260	A	C8-N9-C4	6.54	108.42	105.80
85	AA	1946	C	O4'-C1'-N1	6.54	113.44	108.20
85	AA	2231	G	O4'-C1'-N9	6.54	113.44	108.20
24	AQ	89	ARG	NE-CZ-NH2	-6.54	117.03	120.30
34	BA	282	A	C2-N3-C4	6.54	113.87	110.60
34	BA	1411	C	O4'-C4'-C3'	-6.54	97.46	104.00
34	BA	1836	A	C1'-O4'-C4'	-6.54	104.67	109.90
35	BB	401	U	P-O5'-C5'	-6.54	110.43	120.90
35	BB	528	G	C5'-C4'-C3'	-6.54	105.53	116.00
35	BB	1352	C	C4'-C3'-C2'	-6.54	96.06	102.60
41	BH	15	A	O4'-C1'-N9	6.54	113.43	108.20
85	AA	57	G	O5'-C5'-C4'	-6.54	99.27	111.70
34	BA	135	G	C3'-C2'-C1'	-6.54	96.27	101.50
34	BA	765	U	C5'-C4'-C3'	6.54	126.47	116.00
34	BA	1142	C	C6-N1-C2	-6.54	117.68	120.30
35	BB	1196	A	C4-C5-C6	-6.54	113.73	117.00
36	BC	53	A	C5'-C4'-C3'	-6.54	105.53	116.00
40	BG	70	C	O4'-C1'-N1	6.54	113.43	108.20
41	BH	2	U	C4'-C3'-C2'	-6.54	96.06	102.60
41	BH	79	A	C5-C6-N6	-6.54	118.47	123.70
85	AA	1269	A	N1-C6-N6	-6.54	114.67	118.60
85	AA	1440	C	P-O5'-C5'	-6.54	110.43	120.90
85	AA	1632	G	P-O3'-C3'	6.54	127.55	119.70
34	BA	206	C	O5'-C5'-C4'	6.54	124.13	111.70
34	BA	607	C	C6-N1-C1'	-6.54	112.95	120.80
35	BB	5	A	C3'-C2'-C1'	-6.54	96.27	101.50
80	Bu	145	ARG	N-CA-C	-6.54	93.34	111.00
85	AA	1816	C	C2-N1-C1'	6.54	125.99	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1838	C	O4'-C1'-N1	6.54	113.43	108.20
85	AA	2039	G	C8-N9-C1'	6.54	135.50	127.00
34	BA	605	G	N1-C6-O6	-6.54	115.98	119.90
34	BA	1194	G	C5'-C4'-C3'	6.54	126.46	116.00
35	BB	508	U	C1'-O4'-C4'	-6.54	104.67	109.90
35	BB	702	G	N9-C1'-C2'	-6.54	104.81	112.00
37	BD	8	A	O5'-C5'-C4'	-6.54	99.28	111.70
81	Bv	44	ARG	NE-CZ-NH1	6.54	123.57	120.30
85	AA	121	C	O4'-C1'-N1	6.54	113.43	108.20
85	AA	213	G	C5-C6-O6	-6.54	124.68	128.60
85	AA	493	A	C8-N9-C1'	6.54	139.47	127.70
85	AA	721	C	O4'-C1'-N1	6.54	113.43	108.20
85	AA	1616	U	P-O3'-C3'	6.54	127.55	119.70
85	AA	1784	G	O4'-C1'-N9	6.54	113.43	108.20
85	AA	2172	A	C1'-O4'-C4'	-6.54	104.67	109.90
34	BA	91	C	C6-N1-C2	-6.54	117.69	120.30
34	BA	1614	G	C8-N9-C4	6.54	109.02	106.40
34	BA	1660	A	P-O5'-C5'	-6.54	110.44	120.90
35	BB	106	A	C5-C6-N6	-6.54	118.47	123.70
36	BC	44	A	C5'-C4'-C3'	-6.54	105.54	116.00
39	BF	49	C	C3'-C2'-C1'	-6.54	96.27	101.50
40	BG	6	A	C5-C6-N6	-6.54	118.47	123.70
85	AA	527	A	N9-C1'-C2'	-6.54	104.81	112.00
85	AA	1535	C	OP2-P-O3'	-6.54	90.82	105.20
4	A3	191	ARG	NE-CZ-NH1	6.54	123.57	120.30
23	AP	200	PHE	CB-CG-CD2	-6.54	116.22	120.80
34	BA	686	U	P-O5'-C5'	-6.54	110.44	120.90
34	BA	944	G	C1'-O4'-C4'	-6.54	104.67	109.90
34	BA	1545	C	C2-N1-C1'	-6.54	111.61	118.80
34	BA	1685	C	C6-N1-C1'	6.54	128.64	120.80
36	BC	137	C	C5'-C4'-C3'	-6.54	105.54	116.00
38	BE	147	G	C6-N1-C2	-6.54	121.18	125.10
77	Br	232	SER	N-CA-C	-6.54	93.35	111.00
85	AA	57	G	N1-C6-O6	-6.54	115.98	119.90
85	AA	1093	C	C6-N1-C2	-6.54	117.69	120.30
85	AA	2133	A	C1'-O4'-C4'	-6.54	104.67	109.90
34	BA	977	G	C5'-C4'-O4'	6.53	116.94	109.10
34	BA	1014	A	C5'-C4'-C3'	-6.53	105.55	116.00
34	BA	1256	A	O4'-C1'-N9	6.53	113.43	108.20
35	BB	120	C	C3'-C2'-C1'	-6.53	96.27	101.50
35	BB	815	G	P-O5'-C5'	-6.53	110.44	120.90
37	BD	69	U	N1-C2-N3	6.53	118.82	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	48	G	O4'-C1'-N9	6.53	113.43	108.20
84	By	92	PHE	CB-CG-CD1	6.53	125.37	120.80
85	AA	38	C	C3'-C2'-C1'	-6.53	96.27	101.50
85	AA	51	A	O4'-C1'-N9	6.53	113.43	108.20
85	AA	178	U	C5'-C4'-C3'	-6.53	105.55	116.00
85	AA	1359	U	C2-N1-C1'	6.53	125.54	117.70
85	AA	1622	G	O3'-P-O5'	6.53	116.41	104.00
34	BA	25	C	P-O5'-C5'	-6.53	110.45	120.90
34	BA	181	G	C3'-C2'-C1'	-6.53	96.28	101.50
34	BA	602	G	P-O3'-C3'	-6.53	111.86	119.70
34	BA	1436	A	O3'-P-O5'	6.53	116.41	104.00
35	BB	776	U	O3'-P-O5'	-6.53	91.59	104.00
35	BB	1474	A	O4'-C1'-N9	6.53	113.43	108.20
85	AA	1366	A	C4-N9-C1'	-6.53	114.54	126.30
85	AA	1371	C	C4'-C3'-C2'	-6.53	96.07	102.60
85	AA	2007	G	N1-C6-O6	-6.53	115.98	119.90
34	BA	122	U	C5'-C4'-C3'	-6.53	105.55	116.00
34	BA	383	G	C1'-O4'-C4'	-6.53	104.67	109.90
34	BA	499	C	P-O3'-C3'	-6.53	111.86	119.70
34	BA	1234	U	C1'-O4'-C4'	-6.53	104.68	109.90
34	BA	1616	A	O3'-P-O5'	-6.53	91.59	104.00
35	BB	689	C	C6-N1-C2	-6.53	117.69	120.30
38	BE	133	C	C1'-O4'-C4'	-6.53	104.68	109.90
39	BF	22	U	O4'-C1'-N1	6.53	113.42	108.20
85	AA	7	G	C4-N9-C1'	-6.53	118.01	126.50
85	AA	416	U	O4'-C1'-N1	-6.53	102.98	108.20
85	AA	717	G	P-O3'-C3'	-6.53	111.86	119.70
85	AA	881	C	P-O3'-C3'	6.53	127.54	119.70
85	AA	1171	C	O4'-C1'-N1	6.53	113.42	108.20
85	AA	1355	U	C6-N1-C2	-6.53	117.08	121.00
85	AA	1991	C	C4-C5-C6	6.53	120.67	117.40
34	BA	1217	A	P-O3'-C3'	-6.53	111.87	119.70
35	BB	963	G	C1'-O4'-C4'	-6.53	104.68	109.90
35	BB	1152	U	C1'-O4'-C4'	-6.53	104.68	109.90
40	BG	104	A	N7-C8-N9	6.53	117.06	113.80
85	AA	1845	G	C5-C6-N1	6.53	114.77	111.50
85	AA	2149	C	P-O3'-C3'	-6.53	111.87	119.70
19	AK	118	TYR	CB-CG-CD2	-6.53	117.08	121.00
34	BA	353	U	C2-N3-C4	-6.53	123.08	127.00
34	BA	1346	U	C1'-O4'-C4'	-6.53	104.68	109.90
34	BA	1558	C	N3-C2-O2	-6.53	117.33	121.90
35	BB	851	U	C5'-C4'-O4'	6.53	116.93	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1333	U	C5'-C4'-C3'	-6.53	105.56	116.00
37	BD	44	U	C3'-C2'-C1'	-6.53	96.28	101.50
38	BE	61	A	C4-N9-C1'	-6.53	114.55	126.30
39	BF	35	C	C5'-C4'-C3'	-6.53	105.56	116.00
39	BF	53	G	C1'-O4'-C4'	-6.53	104.68	109.90
65	Bf	448	ARG	NE-CZ-NH2	-6.53	117.04	120.30
85	AA	64	A	C1'-O4'-C4'	-6.53	104.68	109.90
85	AA	1626	U	C5'-C4'-C3'	-6.53	105.56	116.00
85	AA	1950	G	C8-N9-C1'	6.53	135.49	127.00
34	BA	29	U	C5'-C4'-C3'	6.53	126.44	116.00
34	BA	102	G	N3-C4-C5	-6.53	125.34	128.60
34	BA	194	G	O4'-C1'-N9	6.53	113.42	108.20
34	BA	793	A	O4'-C1'-N9	6.53	113.42	108.20
34	BA	810	A	C4-N9-C1'	-6.53	114.56	126.30
35	BB	668	A	O4'-C1'-N9	6.53	113.42	108.20
38	BE	136	G	N9-C1'-C2'	6.53	122.48	114.00
40	BG	135	C	C2-N3-C4	-6.53	116.64	119.90
41	BH	96	G	O5'-P-OP1	6.53	118.53	110.70
49	BP	167	ALA	CB-CA-C	-6.53	100.31	110.10
36	BC	109	A	P-O3'-C3'	-6.52	111.87	119.70
60	Ba	115	PHE	CB-CG-CD1	6.52	125.37	120.80
66	Bg	43	LYS	CB-CA-C	-6.52	97.35	110.40
85	AA	972	G	N1-C6-O6	6.52	123.81	119.90
85	AA	1148	G	N1-C6-O6	6.52	123.81	119.90
85	AA	1578	G	C5'-C4'-O4'	6.52	116.93	109.10
86	AB	42	C	C6-N1-C2	-6.52	117.69	120.30
34	BA	268	U	O4'-C1'-C2'	6.52	113.47	107.60
34	BA	688	G	C4-N9-C1'	-6.52	118.02	126.50
34	BA	1204	U	C6-N1-C1'	6.52	130.33	121.20
34	BA	1344	G	C5'-C4'-C3'	-6.52	105.56	116.00
35	BB	585	U	C5-C6-N1	-6.52	119.44	122.70
35	BB	669	A	O4'-C1'-N9	6.52	113.42	108.20
35	BB	1265	U	P-O3'-C3'	-6.52	111.87	119.70
36	BC	11	G	N1-C2-N2	-6.52	110.33	116.20
38	BE	139	U	O5'-C5'-C4'	-6.52	99.31	111.70
41	BH	18	C	P-O3'-C3'	-6.52	111.87	119.70
85	AA	228	C	O4'-C1'-N1	6.52	113.42	108.20
85	AA	2169	C	C1'-O4'-C4'	-6.52	104.68	109.90
34	BA	880	G	P-O5'-C5'	-6.52	110.47	120.90
34	BA	1062	G	O4'-C1'-N9	6.52	113.42	108.20
35	BB	1417	C	C6-N1-C2	-6.52	117.69	120.30
35	BB	1496	C	C1'-O4'-C4'	-6.52	104.68	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	3	C	P-O3'-C3'	-6.52	111.88	119.70
85	AA	927	A	N9-C1'-C2'	-6.52	104.83	112.00
85	AA	1006	C	O4'-C1'-N1	6.52	113.42	108.20
34	BA	60	A	N1-C6-N6	6.52	122.51	118.60
35	BB	353	G	O4'-C1'-N9	6.52	113.42	108.20
35	BB	1114	A	C4'-C3'-C2'	-6.52	96.08	102.60
35	BB	1144	A	O4'-C1'-N9	6.52	113.42	108.20
35	BB	1446	C	N3-C4-N4	-6.52	113.44	118.00
40	BG	75	C	C2-N1-C1'	-6.52	111.63	118.80
85	AA	44	C	C3'-C2'-C1'	-6.52	96.28	101.50
85	AA	300	C	O5'-C5'-C4'	-6.52	99.31	111.70
85	AA	622	G	C8-N9-C1'	6.52	135.47	127.00
85	AA	691	U	C2-N3-C4	-6.52	123.09	127.00
85	AA	1103	A	N7-C8-N9	6.52	117.06	113.80
34	BA	19	G	C3'-C2'-C1'	-6.52	96.29	101.50
34	BA	383	G	C6-N1-C2	-6.52	121.19	125.10
34	BA	424	U	C2-N3-C4	-6.52	123.09	127.00
34	BA	961	C	N1-C2-O2	6.52	122.81	118.90
34	BA	1722	U	C3'-C2'-C1'	-6.52	96.29	101.50
35	BB	322	G	C8-N9-C1'	-6.52	118.53	127.00
35	BB	916	U	P-O3'-C3'	-6.52	111.88	119.70
35	BB	1255	U	O4'-C1'-N1	6.52	113.41	108.20
35	BB	1452	U	C4-C5-C6	-6.52	115.79	119.70
38	BE	16	C	C5-C4-N4	-6.52	115.64	120.20
59	BZ	43	ARG	NE-CZ-NH1	6.52	123.56	120.30
67	Bh	74	PHE	N-CA-CB	6.52	122.33	110.60
85	AA	750	A	C8-N9-C4	6.52	108.41	105.80
85	AA	886	A	O4'-C4'-C3'	-6.52	97.48	104.00
85	AA	1182	A	O4'-C1'-N9	6.52	113.41	108.20
85	AA	1471	G	N3-C2-N2	6.52	124.46	119.90
85	AA	1832	G	P-O3'-C3'	-6.52	111.88	119.70
35	BB	1492	C	C5'-C4'-C3'	-6.52	105.58	116.00
41	BH	4	U	C2-N1-C1'	-6.52	109.88	117.70
85	AA	1097	G	C4-N9-C1'	-6.52	118.03	126.50
86	AB	13	C	O4'-C1'-N1	6.52	113.41	108.20
34	BA	1597	G	N7-C8-N9	-6.51	109.84	113.10
35	BB	277	C	O4'-C1'-N1	6.51	113.41	108.20
35	BB	507	G	C5'-C4'-C3'	6.51	126.42	116.00
35	BB	818	U	C3'-C2'-C1'	-6.51	96.29	101.50
35	BB	1414	A	O4'-C1'-N9	6.51	113.41	108.20
64	Be	69	TYR	CB-CG-CD1	6.51	124.91	121.00
85	AA	1284	A	C1'-O4'-C4'	-6.51	104.69	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1687	U	C2-N1-C1'	-6.51	109.88	117.70
35	BB	1058	U	P-O3'-C3'	-6.51	111.89	119.70
36	BC	20	C	O4'-C1'-N1	6.51	113.41	108.20
68	Bi	103	ALA	N-CA-C	6.51	128.59	111.00
70	Bk	67	ARG	CD-NE-CZ	-6.51	114.48	123.60
85	AA	246	C	N3-C2-O2	-6.51	117.34	121.90
85	AA	560	C	O4'-C1'-N1	6.51	113.41	108.20
85	AA	1004	G	O4'-C1'-N9	6.51	113.41	108.20
85	AA	1490	A	N9-C4-C5	-6.51	103.19	105.80
85	AA	2095	U	C2-N1-C1'	-6.51	109.89	117.70
34	BA	12	G	O4'-C1'-N9	6.51	113.41	108.20
34	BA	1017	C	P-O5'-C5'	6.51	131.32	120.90
34	BA	1034	U	C3'-C2'-C1'	-6.51	96.29	101.50
34	BA	1196	C	C6-N1-C2	-6.51	117.70	120.30
34	BA	1408	C	C1'-O4'-C4'	-6.51	104.69	109.90
34	BA	1792	U	C5'-C4'-O4'	-6.51	101.28	109.10
35	BB	165	C	O4'-C1'-N1	6.51	113.41	108.20
79	Bt	32	LYS	C-N-CA	6.51	137.98	121.70
85	AA	267	U	C6-N1-C1'	6.51	130.32	121.20
85	AA	413	G	N3-C2-N2	6.51	124.46	119.90
85	AA	1009	G	C4-N9-C1'	-6.51	118.04	126.50
85	AA	1091	C	C5-C6-N1	6.51	124.26	121.00
85	AA	1538	C	C6-N1-C1'	6.51	128.62	120.80
85	AA	1885	A	O4'-C1'-C2'	6.51	113.46	107.60
85	AA	2150	G	C6-C5-N7	-6.51	126.49	130.40
86	AB	8	U	O4'-C1'-N1	6.51	113.41	108.20
34	BA	245	U	N3-C2-O2	-6.51	117.64	122.20
34	BA	646	C	O4'-C1'-N1	6.51	113.41	108.20
34	BA	1222	C	C5-C4-N4	-6.51	115.64	120.20
34	BA	1739	G	N3-C2-N2	-6.51	115.34	119.90
35	BB	705	C	O4'-C1'-N1	6.51	113.41	108.20
35	BB	1053	G	C6-C5-N7	-6.51	126.49	130.40
35	BB	1509	G	O4'-C1'-N9	6.51	113.41	108.20
40	BG	105	A	O5'-C5'-C4'	-6.51	99.33	111.70
40	BG	108	G	N1-C6-O6	-6.51	115.99	119.90
57	BX	87	TYR	N-CA-CB	6.51	122.32	110.60
64	Be	147	ARG	NE-CZ-NH1	6.51	123.56	120.30
85	AA	471	U	C2'-C3'-O3'	6.51	124.11	113.70
85	AA	548	G	O4'-C1'-N9	6.51	113.41	108.20
85	AA	574	U	C3'-C2'-C1'	-6.51	96.29	101.50
85	AA	1466	U	N1-C2-N3	-6.51	111.00	114.90
85	AA	1695	G	C6-N1-C2	-6.51	121.19	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2211	G	C1'-O4'-C4'	-6.51	104.69	109.90
34	BA	180	G	N3-C4-C5	-6.51	125.35	128.60
34	BA	280	A	C5'-C4'-O4'	6.51	116.91	109.10
35	BB	737	C	O4'-C1'-N1	6.51	113.41	108.20
38	BE	66	A	C5-C6-N6	-6.51	118.49	123.70
40	BG	44	G	C1'-O4'-C4'	-6.51	104.69	109.90
51	BR	140	MET	CG-SD-CE	6.51	110.61	100.20
34	BA	34	U	N1-C2-N3	6.51	118.80	114.90
34	BA	593	G	O3'-P-O5'	-6.51	91.64	104.00
34	BA	1334	G	C1'-O4'-C4'	-6.51	104.69	109.90
35	BB	711	C	N3-C2-O2	-6.51	117.35	121.90
35	BB	958	C	C5'-C4'-O4'	-6.51	101.29	109.10
39	BF	10	A	O4'-C1'-N9	6.51	113.41	108.20
47	BN	121	LEU	N-CA-CB	-6.51	97.39	110.40
85	AA	147	G	C1'-O4'-C4'	-6.51	104.69	109.90
85	AA	1407	C	O4'-C1'-N1	6.51	113.41	108.20
85	AA	1858	G	C1'-O4'-C4'	-6.51	104.69	109.90
85	AA	1913	G	P-O5'-C5'	6.51	131.31	120.90
34	BA	297	A	O4'-C1'-N9	-6.50	103.00	108.20
34	BA	857	C	OP1-P-O3'	6.50	119.51	105.20
35	BB	129	U	O4'-C1'-N1	6.50	113.40	108.20
35	BB	1202	G	O3'-P-O5'	-6.50	91.64	104.00
40	BG	101	G	P-O3'-C3'	-6.50	111.89	119.70
85	AA	628	C	P-O3'-C3'	6.50	127.51	119.70
85	AA	1147	A	C8-N9-C4	-6.50	103.20	105.80
85	AA	1454	U	C6-N1-C2	-6.50	117.10	121.00
85	AA	1721	A	C1'-O4'-C4'	-6.50	104.70	109.90
35	BB	793	A	C5'-C4'-O4'	6.50	116.90	109.10
35	BB	1023	G	C4'-C3'-C2'	-6.50	96.10	102.60
35	BB	1458	U	C4-C5-C6	-6.50	115.80	119.70
62	Bc	45	PHE	CB-CG-CD2	-6.50	116.25	120.80
85	AA	896	C	C4'-C3'-C2'	6.50	109.10	102.60
85	AA	982	G	C4-C5-C6	-6.50	114.90	118.80
85	AA	1355	U	OP1-P-O3'	6.50	119.51	105.20
85	AA	2192	A	N1-C6-N6	-6.50	114.70	118.60
35	BB	879	G	C1'-O4'-C4'	6.50	115.10	109.90
35	BB	959	C	O4'-C1'-N1	6.50	113.40	108.20
35	BB	1102	U	C5-C6-N1	-6.50	119.45	122.70
38	BE	18	U	N1-C2-O2	6.50	127.35	122.80
40	BG	30	C	N3-C2-O2	-6.50	117.35	121.90
47	BN	198	ARG	NE-CZ-NH1	6.50	123.55	120.30
85	AA	692	U	O5'-P-OP2	-6.50	99.85	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1044	G	P-O5'-C5'	6.50	131.30	120.90
85	AA	1077	U	O4'-C1'-N1	6.50	113.40	108.20
85	AA	1109	G	O4'-C4'-C3'	-6.50	97.50	104.00
85	AA	1148	G	P-O3'-C3'	-6.50	111.90	119.70
85	AA	1293	U	P-O3'-C3'	-6.50	111.90	119.70
85	AA	1503	G	C5-C6-O6	-6.50	124.70	128.60
8	A7	13	ARG	NE-CZ-NH1	6.50	123.55	120.30
34	BA	616	G	O5'-P-OP1	6.50	118.50	110.70
34	BA	1215	U	C5-C6-N1	-6.50	119.45	122.70
35	BB	599	U	N1-C2-N3	6.50	118.80	114.90
35	BB	1259	A	C8-N9-C4	6.50	108.40	105.80
85	AA	3	U	C6-N1-C1'	-6.50	112.10	121.20
85	AA	229	U	C5'-C4'-C3'	-6.50	105.60	116.00
85	AA	1920	A	O4'-C1'-N9	6.50	113.40	108.20
34	BA	452	A	C5-C6-N6	-6.50	118.50	123.70
34	BA	1438	C	P-O3'-C3'	-6.50	111.90	119.70
38	BE	112	G	C4'-C3'-O3'	6.50	126.00	113.00
71	Bl	138	ARG	NE-CZ-NH2	-6.50	117.05	120.30
85	AA	371	C	O4'-C1'-N1	6.50	113.40	108.20
85	AA	619	A	P-O5'-C5'	6.50	131.30	120.90
85	AA	932	A	P-O5'-C5'	6.50	131.30	120.90
85	AA	1149	A	C1'-O4'-C4'	-6.50	104.70	109.90
34	BA	181	G	C6-N1-C2	-6.50	121.20	125.10
34	BA	496	G	C8-N9-C1'	6.50	135.45	127.00
34	BA	841	G	O4'-C1'-N9	6.50	113.40	108.20
34	BA	1497	A	C2-N3-C4	-6.50	107.35	110.60
35	BB	767	A	C4'-C3'-C2'	-6.50	96.10	102.60
35	BB	1052	G	O4'-C1'-C2'	6.50	113.45	107.60
35	BB	1069	C	P-O3'-C3'	-6.50	111.90	119.70
38	BE	27	A	P-O3'-C3'	-6.50	111.90	119.70
38	BE	149	A	N1-C6-N6	6.50	122.50	118.60
38	BE	199	A	C5-C6-N1	6.50	120.95	117.70
41	BH	72	G	C8-N9-C1'	6.50	135.44	127.00
65	Bf	157	TRP	N-CA-C	6.50	128.54	111.00
83	Bx	115	ARG	NE-CZ-NH1	6.50	123.55	120.30
85	AA	127	U	C4'-C3'-C2'	-6.50	96.10	102.60
85	AA	446	C	O4'-C1'-N1	6.50	113.40	108.20
85	AA	1816	C	C2-N3-C4	-6.50	116.65	119.90
35	BB	503	G	N9-C1'-C2'	-6.50	104.86	112.00
35	BB	843	G	C1'-O4'-C4'	-6.50	104.70	109.90
35	BB	1183	U	O4'-C1'-N1	6.50	113.40	108.20
37	BD	55	A	C3'-C2'-C1'	-6.50	96.30	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	66	A	O4'-C1'-N9	6.50	113.40	108.20
40	BG	153	C	C2-N1-C1'	-6.50	111.66	118.80
56	BW	96	TYR	CB-CG-CD2	-6.50	117.10	121.00
85	AA	1496	U	N3-C4-C5	6.50	118.50	114.60
34	BA	251	U	P-O5'-C5'	6.49	131.29	120.90
34	BA	736	G	C5-N7-C8	-6.49	101.05	104.30
34	BA	841	G	C8-N9-C1'	6.49	135.44	127.00
34	BA	861	C	P-O5'-C5'	-6.49	110.51	120.90
34	BA	1732	A	C5'-C4'-C3'	-6.49	105.61	116.00
34	BA	1732	A	C5-N7-C8	6.49	107.15	103.90
35	BB	483	C	C4'-C3'-C2'	-6.49	96.11	102.60
38	BE	199	A	N9-C1'-C2'	-6.49	104.86	112.00
85	AA	341	C	P-O5'-C5'	6.49	131.29	120.90
85	AA	869	A	C6-N1-C2	-6.49	114.70	118.60
85	AA	1644	G	C6-C5-N7	-6.49	126.50	130.40
85	AA	1956	C	P-O3'-C3'	6.49	127.49	119.70
85	AA	1976	G	C5'-C4'-O4'	6.49	116.89	109.10
85	AA	2210	C	C3'-C2'-C1'	-6.49	96.31	101.50
34	BA	395	G	C5'-C4'-C3'	-6.49	105.61	116.00
34	BA	457	A	P-O3'-C3'	-6.49	111.91	119.70
35	BB	434	A	C5-C6-N6	-6.49	118.51	123.70
35	BB	840	C	C2-N3-C4	-6.49	116.65	119.90
35	BB	1410	G	N1-C6-O6	6.49	123.80	119.90
39	BF	4	A	O4'-C1'-N9	6.49	113.39	108.20
40	BG	14	G	O4'-C1'-N9	6.49	113.39	108.20
41	BH	40	C	P-O5'-C5'	6.49	131.29	120.90
85	AA	247	G	C4-N9-C1'	-6.49	118.06	126.50
85	AA	945	A	C1'-O4'-C4'	-6.49	104.71	109.90
85	AA	1181	U	O5'-P-OP1	6.49	118.49	110.70
85	AA	1271	U	C1'-O4'-C4'	6.49	115.09	109.90
34	BA	351	A	P-O3'-C3'	-6.49	111.91	119.70
34	BA	395	G	C5-C6-O6	6.49	132.50	128.60
34	BA	794	G	P-O5'-C5'	-6.49	110.51	120.90
34	BA	869	C	N3-C4-C5	6.49	124.50	121.90
34	BA	1211	G	C5'-C4'-C3'	6.49	126.39	116.00
34	BA	1775	U	O4'-C1'-N1	6.49	113.39	108.20
35	BB	3	C	C5-C4-N4	-6.49	115.66	120.20
35	BB	468	U	O5'-C5'-C4'	-6.49	99.37	111.70
35	BB	566	A	O3'-P-O5'	-6.49	91.67	104.00
35	BB	659	C	C2-N1-C1'	-6.49	111.66	118.80
35	BB	1331	U	C5'-C4'-O4'	6.49	116.89	109.10
38	BE	96	G	C4-N9-C1'	-6.49	118.06	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	755	G	N3-C4-C5	-6.49	125.35	128.60
85	AA	1634	U	C2-N1-C1'	6.49	125.49	117.70
85	AA	2129	U	C4'-C3'-C2'	-6.49	96.11	102.60
34	BA	881	C	C2'-C3'-O3'	6.49	124.08	113.70
34	BA	1511	C	O4'-C1'-C2'	6.49	113.44	107.60
34	BA	1834	A	O4'-C1'-N9	6.49	113.39	108.20
35	BB	650	A	C5'-C4'-O4'	6.49	116.89	109.10
35	BB	1137	G	O4'-C1'-N9	6.49	113.39	108.20
35	BB	1455	A	N7-C8-N9	6.49	117.05	113.80
36	BC	41	A	N1-C6-N6	6.49	122.49	118.60
62	Bc	133	LYS	N-CA-CB	6.49	122.28	110.60
85	AA	238	C	C6-N1-C1'	-6.49	113.01	120.80
85	AA	611	G	C4-N9-C1'	-6.49	118.07	126.50
85	AA	1982	C	OP1-P-OP2	-6.49	109.87	119.60
35	BB	997	G	O4'-C1'-N9	6.49	113.39	108.20
38	BE	70	C	C5-C4-N4	6.49	124.74	120.20
85	AA	107	A	N9-C1'-C2'	-6.49	104.86	112.00
85	AA	575	G	C2'-C3'-O3'	6.49	124.08	113.70
85	AA	1979	A	C4'-C3'-C2'	-6.49	96.11	102.60
85	AA	2027	U	O4'-C1'-N1	6.49	113.39	108.20
25	AR	26	ASN	CA-CB-CG	-6.49	99.13	113.40
34	BA	321	G	C4'-C3'-C2'	-6.49	96.11	102.60
34	BA	1382	G	P-O3'-C3'	-6.49	111.92	119.70
34	BA	1600	G	C4'-C3'-C2'	-6.49	96.11	102.60
34	BA	1654	G	C1'-O4'-C4'	-6.49	104.71	109.90
34	BA	1752	A	C8-N9-C1'	6.49	139.37	127.70
35	BB	1492	C	C4'-C3'-C2'	-6.49	96.11	102.60
37	BD	42	A	P-O3'-C3'	-6.49	111.92	119.70
40	BG	116	G	N9-C1'-C2'	-6.49	104.86	112.00
41	BH	129	G	O4'-C1'-C2'	6.49	113.44	107.60
84	By	169	ASP	N-CA-C	6.49	128.51	111.00
85	AA	249	C	N3-C2-O2	-6.49	117.36	121.90
85	AA	488	G	C5-C6-O6	-6.49	124.71	128.60
85	AA	1047	G	O4'-C1'-N9	6.49	113.39	108.20
85	AA	1831	U	C4'-C3'-C2'	-6.49	96.11	102.60
86	AB	18	G	O4'-C1'-N9	6.49	113.39	108.20
34	BA	608	G	C5-C6-O6	-6.48	124.71	128.60
34	BA	1320	A	P-O3'-C3'	-6.48	111.92	119.70
35	BB	981	A	C8-N9-C1'	-6.48	116.03	127.70
34	BA	33	C	C3'-C2'-C1'	-6.48	96.31	101.50
34	BA	696	A	C8-N9-C4	6.48	108.39	105.80
34	BA	868	C	N3-C4-N4	-6.48	113.46	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1658	G	C1'-O4'-C4'	-6.48	104.71	109.90
35	BB	457	U	C1'-O4'-C4'	-6.48	104.71	109.90
35	BB	769	C	O4'-C1'-N1	6.48	113.39	108.20
35	BB	1442	C	C6-N1-C2	-6.48	117.71	120.30
37	BD	79	G	C5-C6-N1	6.48	114.74	111.50
37	BD	81	C	O4'-C1'-N1	6.48	113.39	108.20
41	BH	12	U	C2-N3-C4	-6.48	123.11	127.00
68	Bi	129	ARG	CG-CD-NE	-6.48	98.19	111.80
85	AA	676	U	C2-N3-C4	-6.48	123.11	127.00
85	AA	791	C	C6-N1-C2	-6.48	117.71	120.30
85	AA	929	G	C6-N1-C2	-6.48	121.21	125.10
85	AA	964	C	O4'-C1'-N1	6.48	113.39	108.20
85	AA	1608	U	O4'-C1'-N1	6.48	113.39	108.20
85	AA	1687	U	N1-C2-O2	6.48	127.34	122.80
34	BA	711	C	C6-N1-C2	-6.48	117.71	120.30
34	BA	824	C	O4'-C1'-N1	6.48	113.38	108.20
34	BA	829	U	P-O3'-C3'	-6.48	111.92	119.70
35	BB	1383	C	O4'-C1'-N1	6.48	113.39	108.20
55	BV	60	ARG	C-N-CA	6.48	137.90	121.70
85	AA	642	G	C8-N9-C1'	6.48	135.42	127.00
85	AA	1955	U	C3'-C2'-C1'	-6.48	96.32	101.50
85	AA	2183	U	C2-N3-C4	-6.48	123.11	127.00
34	BA	737	U	O4'-C1'-N1	6.48	113.38	108.20
34	BA	997	U	N1-C2-O2	6.48	127.33	122.80
34	BA	1212	A	N1-C6-N6	-6.48	114.71	118.60
40	BG	103	C	C3'-C2'-C1'	-6.48	96.32	101.50
85	AA	159	G	C8-N9-C4	6.48	108.99	106.40
34	BA	573	U	N1-C2-N3	6.48	118.79	114.90
34	BA	1305	A	C8-N9-C1'	-6.48	116.04	127.70
34	BA	1379	G	N1-C6-O6	6.48	123.79	119.90
35	BB	1255	U	C5'-C4'-C3'	-6.48	105.64	116.00
35	BB	1548	C	C4'-C3'-C2'	-6.48	96.12	102.60
36	BC	162	C	P-O3'-C3'	6.48	127.47	119.70
39	BF	50	C	C4'-C3'-C2'	6.48	109.08	102.60
40	BG	62	C	P-O3'-C3'	6.48	127.47	119.70
85	AA	571	G	C4-N9-C1'	-6.48	118.08	126.50
85	AA	578	U	C4'-C3'-C2'	-6.48	96.12	102.60
85	AA	579	U	C2-N1-C1'	-6.48	109.93	117.70
85	AA	1301	C	C2-N1-C1'	-6.48	111.67	118.80
85	AA	2148	C	P-O3'-C3'	-6.48	111.93	119.70
85	AA	2196	G	N9-C1'-C2'	-6.48	104.88	112.00
34	BA	24	C	C1'-O4'-C4'	-6.48	104.72	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	155	G	O4'-C1'-N9	6.48	113.38	108.20
35	BB	502	C	C6-N1-C1'	6.48	128.57	120.80
35	BB	1025	A	C8-N9-C4	-6.48	103.21	105.80
37	BD	86	A	C5'-C4'-O4'	-6.48	101.33	109.10
82	Bw	91	TYR	CB-CG-CD1	-6.48	117.11	121.00
85	AA	890	U	C6-N1-C2	-6.48	117.11	121.00
85	AA	1797	U	C5-C4-O4	-6.48	122.01	125.90
85	AA	1869	U	O4'-C1'-N1	6.48	113.38	108.20
27	AT	43	ALA	N-CA-CB	-6.47	101.03	110.10
34	BA	843	G	N3-C4-N9	6.47	129.88	126.00
34	BA	1721	U	O3'-P-O5'	-6.47	91.70	104.00
35	BB	40	C	C6-N1-C1'	6.47	128.57	120.80
35	BB	454	U	C1'-O4'-C4'	-6.47	104.72	109.90
35	BB	1152	U	P-O5'-C5'	-6.47	110.54	120.90
35	BB	1210	U	C1'-O4'-C4'	-6.47	104.72	109.90
36	BC	97	U	C3'-C2'-C1'	-6.47	96.32	101.50
40	BG	41	U	C1'-O4'-C4'	-6.47	104.72	109.90
50	BQ	79	PHE	CA-CB-CG	-6.47	98.36	113.90
85	AA	661	C	P-O5'-C5'	6.47	131.26	120.90
85	AA	1280	U	C4-C5-C6	-6.47	115.81	119.70
85	AA	1798	U	P-O3'-C3'	-6.47	111.93	119.70
86	AB	8	U	O5'-C5'-C4'	6.47	124.00	111.70
34	BA	514	U	C2-N3-C4	-6.47	123.12	127.00
34	BA	590	U	C6-N1-C2	-6.47	117.12	121.00
34	BA	618	G	N9-C4-C5	6.47	107.99	105.40
34	BA	972	C	O4'-C1'-N1	6.47	113.38	108.20
34	BA	1110	A	C5-C6-N1	6.47	120.94	117.70
34	BA	1314	A	C5'-C4'-C3'	-6.47	105.64	116.00
34	BA	1439	C	N3-C2-O2	-6.47	117.37	121.90
35	BB	73	G	C5-C6-O6	-6.47	124.72	128.60
35	BB	807	U	O3'-P-O5'	6.47	116.30	104.00
36	BC	74	U	P-O3'-C3'	-6.47	111.93	119.70
37	BD	115	A	C4'-C3'-C2'	-6.47	96.13	102.60
38	BE	100	U	N1-C2-N3	6.47	118.78	114.90
39	BF	47	C	C3'-C2'-C1'	-6.47	96.32	101.50
41	BH	46	C	C6-N1-C2	-6.47	117.71	120.30
57	BX	87	TYR	N-CA-C	-6.47	93.53	111.00
85	AA	638	G	P-O3'-C3'	-6.47	111.93	119.70
85	AA	929	G	C5-C6-N1	6.47	114.74	111.50
85	AA	1173	A	C5-C6-N6	-6.47	118.52	123.70
85	AA	1243	G	P-O3'-C3'	-6.47	111.93	119.70
85	AA	1280	U	C1'-O4'-C4'	-6.47	104.72	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1502	A	P-O5'-C5'	6.47	131.26	120.90
85	AA	2199	G	C1'-O4'-C4'	-6.47	104.72	109.90
31	AX	166	PHE	CB-CG-CD1	-6.47	116.27	120.80
34	BA	271	C	O4'-C1'-N1	6.47	113.38	108.20
34	BA	404	C	O4'-C1'-N1	6.47	113.38	108.20
34	BA	1425	G	C1'-O4'-C4'	-6.47	104.72	109.90
85	AA	283	A	P-O3'-C3'	6.47	127.47	119.70
85	AA	972	G	C4'-C3'-C2'	6.47	109.07	102.60
85	AA	1177	G	O4'-C1'-N9	6.47	113.38	108.20
85	AA	1785	U	P-O5'-C5'	6.47	131.25	120.90
34	BA	28	C	O4'-C1'-N1	6.47	113.38	108.20
34	BA	147	U	C5-C6-N1	-6.47	119.47	122.70
34	BA	663	U	C2-N3-C4	-6.47	123.12	127.00
34	BA	1238	C	C3'-C2'-C1'	-6.47	96.32	101.50
34	BA	1527	G	N9-C1'-C2'	-6.47	104.88	112.00
34	BA	1579	G	C4-N9-C1'	6.47	134.91	126.50
35	BB	883	G	C1'-O4'-C4'	-6.47	104.72	109.90
35	BB	1384	A	O4'-C1'-N9	6.47	113.38	108.20
38	BE	195	G	C1'-O4'-C4'	-6.47	104.72	109.90
39	BF	16	C	O5'-C5'-C4'	6.47	123.99	111.70
41	BH	130	G	C5'-C4'-C3'	-6.47	105.65	116.00
85	AA	260	A	O4'-C1'-N9	6.47	113.38	108.20
85	AA	1157	U	C5'-C4'-C3'	-6.47	105.65	116.00
85	AA	1191	G	P-O5'-C5'	6.47	131.25	120.90
34	BA	784	C	N1-C1'-C2'	-6.47	104.89	112.00
35	BB	322	G	C4-N9-C1'	6.47	134.91	126.50
35	BB	1202	G	O4'-C4'-C3'	-6.47	97.53	104.00
36	BC	37	U	C2-N1-C1'	6.47	125.46	117.70
40	BG	97	G	C5-C6-O6	-6.47	124.72	128.60
62	Bc	124	CYS	N-CA-CB	6.47	122.24	110.60
85	AA	508	C	O4'-C4'-C3'	-6.47	97.53	104.00
85	AA	1435	C	C3'-C2'-C1'	-6.47	96.33	101.50
34	BA	168	U	N1-C2-N3	6.47	118.78	114.90
34	BA	431	A	C6-N1-C2	-6.47	114.72	118.60
34	BA	542	A	C5'-C4'-C3'	-6.47	105.65	116.00
34	BA	590	U	N3-C2-O2	-6.47	117.67	122.20
35	BB	858	U	O4'-C1'-N1	6.47	113.37	108.20
40	BG	9	G	C4-C5-C6	-6.47	114.92	118.80
48	BO	153	ARG	NE-CZ-NH1	6.47	123.53	120.30
85	AA	244	G	N9-C1'-C2'	-6.47	104.89	112.00
85	AA	541	A	C4'-C3'-C2'	-6.47	96.13	102.60
85	AA	591	A	P-O3'-C3'	-6.47	111.94	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	688	C	C3'-C2'-C1'	-6.47	96.33	101.50
85	AA	1268	C	C1'-O4'-C4'	-6.47	104.73	109.90
22	AO	35	CYS	CA-CB-SG	-6.46	102.36	114.00
30	AW	34	MET	CG-SD-CE	-6.46	89.86	100.20
34	BA	109	A	P-O5'-C5'	-6.46	110.56	120.90
34	BA	1594	G	N9-C1'-C2'	-6.46	104.89	112.00
35	BB	279	A	O4'-C1'-N9	6.46	113.37	108.20
35	BB	1132	A	C4-C5-C6	-6.46	113.77	117.00
36	BC	104	A	C5'-C4'-C3'	-6.46	105.66	116.00
85	AA	15	U	C5'-C4'-C3'	-6.46	105.66	116.00
85	AA	817	G	C4'-C3'-C2'	6.46	109.06	102.60
85	AA	1586	C	C6-N1-C2	-6.46	117.71	120.30
85	AA	1894	G	C5'-C4'-O4'	-6.46	101.34	109.10
3	A2	14	TYR	CB-CG-CD2	-6.46	117.12	121.00
34	BA	262	A	C2-N3-C4	-6.46	107.37	110.60
34	BA	861	C	P-O3'-C3'	-6.46	111.94	119.70
37	BD	6	C	P-O5'-C5'	-6.46	110.56	120.90
39	BF	40	U	P-O3'-C3'	-6.46	111.94	119.70
85	AA	58	C	C5-C6-N1	6.46	124.23	121.00
85	AA	1231	G	C5'-C4'-C3'	6.46	126.34	116.00
34	BA	301	U	O4'-C4'-C3'	-6.46	97.54	104.00
34	BA	1487	U	C6-N1-C2	-6.46	117.12	121.00
34	BA	1668	C	C6-N1-C2	-6.46	117.72	120.30
34	BA	1749	C	C6-N1-C2	-6.46	117.72	120.30
34	BA	1795	A	C8-N9-C4	6.46	108.39	105.80
35	BB	589	U	O4'-C1'-N1	6.46	113.37	108.20
35	BB	998	G	C5-C6-N1	6.46	114.73	111.50
35	BB	1130	U	C2-N3-C4	-6.46	123.12	127.00
41	BH	78	C	OP1-P-O3'	6.46	119.42	105.20
69	Bj	101	GLN	CB-CA-C	-6.46	97.48	110.40
85	AA	24	U	C2-N3-C4	-6.46	123.12	127.00
85	AA	155	U	N1-C1'-C2'	-6.46	104.89	112.00
85	AA	1510	A	P-O5'-C5'	6.46	131.24	120.90
85	AA	1937	G	O4'-C1'-N9	6.46	113.37	108.20
34	BA	1506	C	C6-N1-C1'	6.46	128.55	120.80
36	BC	162	C	N3-C2-O2	-6.46	117.38	121.90
50	BQ	219	ARG	NE-CZ-NH1	6.46	123.53	120.30
85	AA	1459	C	C2-N1-C1'	-6.46	111.69	118.80
85	AA	1561	A	C8-N9-C4	6.46	108.38	105.80
34	BA	500	C	O4'-C1'-N1	6.46	113.37	108.20
34	BA	675	C	P-O5'-C5'	6.46	131.23	120.90
34	BA	1588	U	C4'-C3'-C2'	6.46	109.06	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1735	G	C2'-C3'-O3'	6.46	124.03	113.70
34	BA	1798	G	C4-N9-C1'	6.46	134.90	126.50
35	BB	764	C	C2-N1-C1'	-6.46	111.70	118.80
35	BB	1508	G	N1-C6-O6	-6.46	116.03	119.90
36	BC	137	C	C3'-C2'-C1'	-6.46	96.33	101.50
38	BE	98	C	P-O3'-C3'	-6.46	111.95	119.70
38	BE	167	U	C2-N1-C1'	-6.46	109.95	117.70
82	Bw	33	PHE	CB-CG-CD2	-6.46	116.28	120.80
85	AA	115	U	C5-C6-N1	-6.46	119.47	122.70
4	A3	226	ARG	NE-CZ-NH1	6.46	123.53	120.30
34	BA	865	C	C2-N3-C4	-6.46	116.67	119.90
34	BA	1088	G	N9-C1'-C2'	-6.46	104.90	112.00
34	BA	1229	G	N1-C6-O6	6.46	123.77	119.90
34	BA	1757	C	O4'-C1'-N1	6.46	113.37	108.20
35	BB	160	A	P-O3'-C3'	6.46	127.45	119.70
35	BB	1122	C	C5'-C4'-C3'	-6.46	105.67	116.00
35	BB	1292	G	O5'-C5'-C4'	-6.46	99.43	111.70
36	BC	120	G	C6-C5-N7	-6.46	126.53	130.40
41	BH	86	G	N1-C2-N3	-6.46	120.03	123.90
85	AA	426	C	P-O3'-C3'	-6.46	111.95	119.70
85	AA	577	U	N1-C2-N3	6.46	118.77	114.90
85	AA	750	A	N1-C6-N6	6.46	122.47	118.60
85	AA	795	C	C2-N1-C1'	-6.46	111.70	118.80
85	AA	1513	U	O4'-C1'-N1	6.46	113.36	108.20
85	AA	1541	G	C3'-C2'-C1'	-6.46	96.33	101.50
85	AA	1800	U	O4'-C1'-N1	6.46	113.36	108.20
85	AA	2128	G	P-O5'-C5'	-6.46	110.57	120.90
85	AA	2146	G	O4'-C1'-N9	6.46	113.36	108.20
35	BB	795	A	C4-N9-C1'	-6.46	114.68	126.30
35	BB	1188	A	C5-C6-N6	6.46	128.86	123.70
85	AA	315	U	C5'-C4'-O4'	6.46	116.85	109.10
85	AA	725	G	C1'-O4'-C4'	-6.46	104.74	109.90
34	BA	829	U	P-O5'-C5'	6.45	131.23	120.90
34	BA	837	U	C5'-C4'-O4'	6.45	116.84	109.10
34	BA	1191	C	C2-N1-C1'	-6.45	111.70	118.80
34	BA	1336	U	P-O3'-C3'	-6.45	111.95	119.70
34	BA	1464	C	C6-N1-C1'	6.45	128.54	120.80
34	BA	1748	G	C4-N9-C1'	-6.45	118.11	126.50
34	BA	1780	U	O4'-C1'-N1	6.45	113.36	108.20
35	BB	633	C	C1'-O4'-C4'	-6.45	104.74	109.90
35	BB	1185	G	C5-C6-O6	-6.45	124.73	128.60
37	BD	96	C	O4'-C1'-N1	6.45	113.36	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	107	U	P-O5'-C5'	-6.45	110.57	120.90
53	BT	153	ARG	NE-CZ-NH1	6.45	123.53	120.30
80	Bu	263	ARG	NE-CZ-NH2	-6.45	117.07	120.30
85	AA	423	G	C3'-C2'-C1'	-6.45	96.34	101.50
85	AA	651	G	P-O3'-C3'	-6.45	111.95	119.70
85	AA	1140	G	N3-C2-N2	6.45	124.42	119.90
85	AA	1987	G	O4'-C1'-N9	6.45	113.36	108.20
35	BB	1045	G	C5-C6-O6	-6.45	124.73	128.60
35	BB	1290	C	C1'-O4'-C4'	-6.45	104.74	109.90
38	BE	58	U	C4-C5-C6	-6.45	115.83	119.70
85	AA	84	C	C5-C6-N1	6.45	124.23	121.00
85	AA	301	U	C5'-C4'-O4'	6.45	116.84	109.10
85	AA	820	G	P-O5'-C5'	-6.45	110.58	120.90
85	AA	926	C	N1-C1'-C2'	-6.45	104.90	112.00
85	AA	1502	A	O4'-C1'-N9	6.45	113.36	108.20
85	AA	1800	U	C4'-C3'-C2'	-6.45	96.15	102.60
34	BA	803	U	O3'-P-O5'	-6.45	91.74	104.00
34	BA	1840	C	O4'-C1'-N1	6.45	113.36	108.20
35	BB	1285	U	C5-C6-N1	-6.45	119.47	122.70
36	BC	112	G	P-O5'-C5'	6.45	131.22	120.90
85	AA	159	G	C1'-O4'-C4'	-6.45	104.74	109.90
34	BA	138	C	C5-C6-N1	6.45	124.22	121.00
34	BA	884	G	O4'-C1'-N9	6.45	113.36	108.20
35	BB	625	A	O4'-C1'-N9	6.45	113.36	108.20
35	BB	1011	C	O4'-C1'-N1	6.45	113.36	108.20
36	BC	162	C	C6-N1-C2	-6.45	117.72	120.30
41	BH	129	G	C8-N9-C1'	6.45	135.38	127.00
85	AA	508	C	C3'-C2'-C1'	-6.45	96.34	101.50
85	AA	921	C	O4'-C1'-N1	6.45	113.36	108.20
35	BB	126	C	C3'-C2'-C1'	-6.45	96.34	101.50
85	AA	54	C	P-O3'-C3'	-6.45	111.96	119.70
85	AA	508	C	O4'-C1'-N1	6.45	113.36	108.20
85	AA	1248	U	N1-C2-N3	6.45	118.77	114.90
85	AA	2238	C	O4'-C1'-N1	6.45	113.36	108.20
34	BA	18	G	N1-C6-O6	6.45	123.77	119.90
34	BA	204	U	C4'-C3'-C2'	-6.45	96.16	102.60
34	BA	792	A	P-O3'-C3'	6.45	127.43	119.70
34	BA	1083	A	C2'-C3'-O3'	6.45	124.01	113.70
34	BA	1435	A	N9-C1'-C2'	-6.45	104.91	112.00
34	BA	1616	A	C8-N9-C4	-6.45	103.22	105.80
35	BB	733	G	N1-C6-O6	6.45	123.77	119.90
35	BB	836	U	C5-C4-O4	-6.45	122.03	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1118	G	C3'-C2'-C1'	-6.45	96.34	101.50
35	BB	1494	G	O3'-P-O5'	6.45	116.25	104.00
36	BC	55	U	C5'-C4'-O4'	6.45	116.83	109.10
37	BD	113	G	C5'-C4'-C3'	6.45	126.31	116.00
53	BT	109	TYR	CB-CG-CD1	6.45	124.87	121.00
58	BY	55	TRP	CB-CG-CD2	-6.45	118.22	126.60
85	AA	369	A	O4'-C1'-N9	6.45	113.36	108.20
85	AA	561	C	C5-C4-N4	6.45	124.71	120.20
85	AA	1858	G	P-O3'-C3'	-6.45	111.96	119.70
32	AY	60	GLN	CA-CB-CG	6.44	127.58	113.40
34	BA	230	A	P-O3'-C3'	6.44	127.43	119.70
34	BA	258	C	N3-C2-O2	-6.44	117.39	121.90
34	BA	842	U	O3'-P-O5'	-6.44	91.76	104.00
34	BA	1314	A	O3'-P-O5'	-6.44	91.76	104.00
35	BB	1027	U	P-O5'-C5'	6.44	131.21	120.90
85	AA	368	C	O3'-P-O5'	6.44	116.24	104.00
34	BA	6	C	C4'-C3'-C2'	6.44	109.04	102.60
34	BA	410	G	C5'-C4'-O4'	6.44	116.83	109.10
34	BA	495	A	C4-N9-C1'	-6.44	114.70	126.30
34	BA	512	U	N3-C2-O2	-6.44	117.69	122.20
34	BA	761	U	C5-C6-N1	-6.44	119.48	122.70
34	BA	1270	G	C4-N9-C1'	-6.44	118.12	126.50
34	BA	1425	G	P-O5'-C5'	6.44	131.21	120.90
35	BB	104	G	C4-N9-C1'	-6.44	118.12	126.50
35	BB	893	U	O4'-C1'-N1	6.44	113.35	108.20
37	BD	40	C	OP2-P-O3'	6.44	119.37	105.20
39	BF	47	C	C6-N1-C2	-6.44	117.72	120.30
44	BK	139	ARG	CD-NE-CZ	-6.44	114.58	123.60
62	Bc	7	LEU	N-CA-CB	-6.44	97.51	110.40
85	AA	97	A	C1'-O4'-C4'	-6.44	104.75	109.90
85	AA	273	C	C5-C6-N1	6.44	124.22	121.00
85	AA	508	C	P-O5'-C5'	6.44	131.21	120.90
85	AA	1891	U	N3-C2-O2	-6.44	117.69	122.20
21	AM	126	HIS	CA-CB-CG	6.44	124.55	113.60
27	AT	34	HIS	C-N-CD	-6.44	106.43	120.60
34	BA	231	U	O4'-C1'-N1	6.44	113.35	108.20
34	BA	960	C	P-O5'-C5'	-6.44	110.59	120.90
35	BB	132	G	O4'-C1'-C2'	6.44	113.40	107.60
82	Bw	164	TYR	CB-CG-CD1	-6.44	117.14	121.00
85	AA	243	A	C5'-C4'-C3'	6.44	126.31	116.00
85	AA	289	G	O4'-C4'-C3'	-6.44	97.56	104.00
85	AA	742	U	C4'-C3'-C2'	-6.44	96.16	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1209	U	N3-C2-O2	-6.44	117.69	122.20
34	BA	400	A	O4'-C1'-C2'	6.44	113.39	107.60
34	BA	851	C	O5'-C5'-C4'	-6.44	99.47	111.70
35	BB	24	C	C1'-O4'-C4'	-6.44	104.75	109.90
35	BB	1507	U	N1-C2-O2	6.44	127.31	122.80
35	BB	1542	C	P-O3'-C3'	-6.44	111.97	119.70
64	Be	161	SER	C-N-CA	6.44	137.80	121.70
85	AA	195	C	O4'-C1'-N1	6.44	113.35	108.20
85	AA	1593	C	C6-N1-C2	-6.44	117.72	120.30
85	AA	1799	C	C5-C6-N1	6.44	124.22	121.00
6	A5	77	ARG	NE-CZ-NH1	6.44	123.52	120.30
34	BA	1535	G	C4'-C3'-C2'	-6.44	96.16	102.60
34	BA	1786	C	O4'-C1'-N1	6.44	113.35	108.20
35	BB	405	U	O4'-C1'-N1	6.44	113.35	108.20
64	Be	163	ARG	CA-CB-CG	6.44	127.56	113.40
85	AA	42	G	P-O3'-C3'	6.44	127.42	119.70
85	AA	1018	G	C1'-O4'-C4'	-6.44	104.75	109.90
85	AA	1453	U	C4'-C3'-C2'	-6.44	96.16	102.60
85	AA	2105	G	P-O3'-C3'	-6.44	111.98	119.70
31	AX	123	ARG	NE-CZ-NH2	-6.44	117.08	120.30
34	BA	322	U	N3-C2-O2	-6.44	117.69	122.20
85	AA	177	A	O4'-C1'-C2'	6.44	113.39	107.60
85	AA	541	A	C3'-C2'-C1'	-6.44	96.35	101.50
34	BA	429	G	N3-C4-N9	-6.43	122.14	126.00
35	BB	845	C	C1'-O4'-C4'	-6.43	104.75	109.90
40	BG	82	U	C6-N1-C1'	6.43	130.21	121.20
40	BG	112	C	N1-C1'-C2'	-6.43	104.92	112.00
40	BG	145	C	C3'-C2'-C1'	-6.43	96.35	101.50
40	BG	172	C	P-O5'-C5'	-6.43	110.61	120.90
41	BH	26	C	C5'-C4'-C3'	6.43	126.30	116.00
41	BH	55	C	P-O3'-C3'	-6.43	111.98	119.70
85	AA	301	U	P-O3'-C3'	6.43	127.42	119.70
85	AA	350	U	O4'-C1'-N1	6.43	113.35	108.20
85	AA	386	G	C3'-C2'-C1'	-6.43	96.35	101.50
85	AA	665	A	N1-C6-N6	-6.43	114.74	118.60
85	AA	867	G	P-O5'-C5'	-6.43	110.60	120.90
85	AA	1096	G	C1'-O4'-C4'	-6.43	104.75	109.90
85	AA	1929	G	C5-C6-O6	-6.43	124.74	128.60
34	BA	252	A	P-O5'-C5'	-6.43	110.61	120.90
34	BA	615	A	C5'-C4'-O4'	6.43	116.82	109.10
34	BA	1078	U	C4-C5-C6	-6.43	115.84	119.70
34	BA	1724	G	N9-C1'-C2'	-6.43	104.92	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1218	G	O4'-C1'-N9	6.43	113.35	108.20
36	BC	25	C	C2'-C3'-O3'	6.43	123.99	113.70
36	BC	124	A	N3-C4-N9	6.43	132.55	127.40
38	BE	23	G	N1-C6-O6	-6.43	116.04	119.90
41	BH	39	G	O4'-C4'-C3'	-6.43	97.57	104.00
85	AA	118	C	O4'-C1'-C2'	6.43	113.39	107.60
85	AA	393	C	C1'-O4'-C4'	-6.43	104.75	109.90
34	BA	602	G	C5-C6-O6	-6.43	124.74	128.60
34	BA	815	C	C6-N1-C1'	-6.43	113.08	120.80
34	BA	1368	G	O4'-C1'-N9	6.43	113.34	108.20
34	BA	1558	C	O5'-C5'-C4'	-6.43	99.48	111.70
34	BA	1775	U	C4'-C3'-C2'	-6.43	96.17	102.60
85	AA	1777	C	O4'-C1'-N1	6.43	113.34	108.20
85	AA	1946	C	P-O3'-C3'	6.43	127.42	119.70
85	AA	2110	U	O4'-C1'-C2'	6.43	113.39	107.60
1	A0	24	MET	CG-SD-CE	-6.43	89.91	100.20
34	BA	544	U	N1-C2-N3	-6.43	111.04	114.90
34	BA	693	G	N9-C1'-C2'	6.43	122.36	114.00
34	BA	1546	C	N3-C2-O2	-6.43	117.40	121.90
34	BA	1648	G	O5'-P-OP1	6.43	118.42	110.70
34	BA	1713	U	P-O3'-C3'	6.43	127.42	119.70
35	BB	85	A	C8-N9-C4	6.43	108.37	105.80
35	BB	696	G	P-O3'-C3'	-6.43	111.98	119.70
35	BB	1142	C	C3'-C2'-C1'	-6.43	96.36	101.50
35	BB	1361	A	C5-C6-N1	6.43	120.91	117.70
35	BB	1381	U	O4'-C1'-N1	6.43	113.34	108.20
36	BC	92	C	C4'-C3'-C2'	-6.43	96.17	102.60
85	AA	171	U	O4'-C1'-N1	-6.43	103.06	108.20
85	AA	852	C	N1-C1'-C2'	-6.43	104.93	112.00
85	AA	1337	A	O4'-C1'-N9	6.43	113.34	108.20
85	AA	1508	A	O4'-C1'-N9	6.43	113.34	108.20
85	AA	1805	A	P-O5'-C5'	6.43	131.19	120.90
85	AA	2241	C	C2-N1-C1'	-6.43	111.73	118.80
19	AK	46	LEU	CA-C-N	6.43	135.10	117.10
20	AL	63	ARG	NE-CZ-NH1	6.43	123.51	120.30
34	BA	67	A	C8-N9-C4	6.43	108.37	105.80
35	BB	306	U	P-O5'-C5'	6.43	131.18	120.90
35	BB	652	G	C3'-C2'-C1'	6.43	106.64	101.50
37	BD	64	A	P-O3'-C3'	6.43	127.41	119.70
40	BG	63	U	O3'-P-O5'	-6.43	91.79	104.00
85	AA	2241	C	O4'-C1'-C2'	6.43	113.39	107.60
34	BA	850	C	C4'-C3'-C2'	-6.43	96.17	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	885	A	P-O3'-C3'	6.43	127.41	119.70
34	BA	976	C	C1'-O4'-C4'	-6.43	104.76	109.90
34	BA	1250	C	N1-C1'-C2'	-6.43	104.93	112.00
34	BA	1407	C	C6-N1-C2	-6.43	117.73	120.30
35	BB	452	A	P-O5'-C5'	-6.43	110.62	120.90
35	BB	526	A	C5-C6-N1	-6.43	114.49	117.70
51	BR	155	GLU	N-CA-C	-6.43	93.65	111.00
80	Bu	34	ARG	NE-CZ-NH1	6.43	123.51	120.30
85	AA	43	A	O5'-C5'-C4'	-6.43	99.49	111.70
85	AA	444	U	C4-C5-C6	-6.43	115.84	119.70
85	AA	564	A	O4'-C1'-N9	6.43	113.34	108.20
85	AA	622	G	N3-C2-N2	6.43	124.40	119.90
85	AA	2019	G	C4-N9-C1'	-6.43	118.15	126.50
85	AA	2165	C	C2-N3-C4	6.43	123.11	119.90
4	A3	89	ARG	NE-CZ-NH1	6.42	123.51	120.30
34	BA	1316	G	C4-N9-C1'	6.42	134.85	126.50
35	BB	5	A	N1-C2-N3	-6.42	126.09	129.30
35	BB	36	U	C2-N3-C4	-6.42	123.15	127.00
35	BB	857	G	C3'-C2'-C1'	-6.42	96.36	101.50
36	BC	55	U	O4'-C1'-N1	6.42	113.34	108.20
36	BC	89	U	C5'-C4'-C3'	-6.42	105.72	116.00
37	BD	74	A	C8-N9-C4	6.42	108.37	105.80
38	BE	137	A	C4-N9-C1'	-6.42	114.73	126.30
38	BE	204	U	C2-N3-C4	-6.42	123.15	127.00
39	BF	36	G	C5-C6-O6	-6.42	124.75	128.60
40	BG	34	A	C4-N9-C1'	-6.42	114.74	126.30
41	BH	38	G	C5-C6-N1	6.42	114.71	111.50
85	AA	65	A	O4'-C4'-C3'	-6.42	97.58	104.00
85	AA	1263	G	C5'-C4'-C3'	-6.42	105.72	116.00
85	AA	1710	C	C4'-C3'-C2'	6.42	109.03	102.60
85	AA	1736	U	O4'-C1'-N1	6.42	113.34	108.20
85	AA	2022	A	C5-C6-N1	6.42	120.91	117.70
85	AA	2169	C	C6-N1-C2	-6.42	117.73	120.30
34	BA	612	U	C3'-C2'-C1'	-6.42	96.36	101.50
34	BA	1683	C	C5'-C4'-C3'	6.42	126.28	116.00
38	BE	188	C	C3'-C2'-C1'	-6.42	96.36	101.50
40	BG	116	G	C8-N9-C1'	6.42	135.35	127.00
85	AA	278	C	C1'-O4'-C4'	-6.42	104.76	109.90
85	AA	1211	C	C3'-C2'-C1'	-6.42	96.36	101.50
85	AA	1507	G	P-O5'-C5'	-6.42	110.62	120.90
85	AA	2148	C	C3'-C2'-C1'	-6.42	96.36	101.50
34	BA	141	G	C6-N1-C2	-6.42	121.25	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	183	G	C3'-C2'-C1'	-6.42	96.36	101.50
34	BA	322	U	C6-N1-C1'	6.42	130.19	121.20
34	BA	809	U	C5-C4-O4	6.42	129.75	125.90
34	BA	874	G	N1-C2-N3	-6.42	120.05	123.90
34	BA	907	A	C5-C6-N6	6.42	128.84	123.70
34	BA	1059	U	C5'-C4'-C3'	-6.42	105.73	116.00
34	BA	1494	G	O4'-C1'-N9	6.42	113.34	108.20
35	BB	359	A	C5'-C4'-C3'	-6.42	105.72	116.00
35	BB	690	C	P-O5'-C5'	6.42	131.17	120.90
38	BE	2	G	C5-C6-N1	6.42	114.71	111.50
38	BE	68	U	C6-N1-C1'	6.42	130.19	121.20
39	BF	15	U	O4'-C1'-C2'	-6.42	99.38	105.80
61	Bb	124	ARG	NE-CZ-NH1	6.42	123.51	120.30
65	Bf	346	SER	CA-C-N	6.42	131.33	117.20
85	AA	641	A	C5-C6-N6	6.42	128.84	123.70
85	AA	1006	C	C6-N1-C2	-6.42	117.73	120.30
85	AA	1474	U	C6-N1-C2	-6.42	117.15	121.00
85	AA	2169	C	C5-C4-N4	6.42	124.69	120.20
34	BA	420	A	C5'-C4'-C3'	-6.42	105.73	116.00
34	BA	1307	U	C1'-O4'-C4'	-6.42	104.76	109.90
35	BB	1215	U	P-O3'-C3'	-6.42	112.00	119.70
37	BD	14	C	C2'-C3'-O3'	6.42	123.97	113.70
85	AA	381	A	P-O5'-C5'	-6.42	110.63	120.90
85	AA	1973	G	P-O5'-C5'	-6.42	110.63	120.90
34	BA	420	A	N9-C4-C5	-6.42	103.23	105.80
35	BB	385	C	P-O3'-C3'	-6.42	112.00	119.70
35	BB	682	U	P-O3'-C3'	-6.42	112.00	119.70
35	BB	1111	C	C1'-O4'-C4'	-6.42	104.77	109.90
40	BG	92	U	C4'-C3'-C2'	6.42	109.02	102.60
41	BH	29	G	C4-C5-N7	6.42	113.37	110.80
85	AA	937	G	P-O3'-C3'	-6.42	112.00	119.70
85	AA	1491	G	C2-N3-C4	-6.42	108.69	111.90
34	BA	727	G	C6-N1-C2	-6.42	121.25	125.10
34	BA	917	C	N3-C4-N4	-6.42	113.51	118.00
34	BA	1607	U	C2-N3-C4	-6.42	123.15	127.00
35	BB	1485	G	P-O3'-C3'	-6.42	112.00	119.70
53	BT	38	ARG	N-CA-CB	-6.42	99.05	110.60
85	AA	101	C	C5'-C4'-O4'	-6.42	101.40	109.10
85	AA	513	G	C5'-C4'-C3'	-6.42	105.73	116.00
85	AA	1244	A	C8-N9-C4	6.42	108.37	105.80
34	BA	1581	G	C6-N1-C2	-6.42	121.25	125.10
85	AA	191	C	O4'-C1'-N1	6.42	113.33	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	543	A	C5'-C4'-C3'	6.42	126.27	116.00
85	AA	1176	C	N1-C2-N3	6.42	123.69	119.20
34	BA	176	G	O4'-C1'-N9	6.41	113.33	108.20
34	BA	201	A	O3'-P-O5'	-6.41	91.81	104.00
34	BA	271	C	C4'-C3'-C2'	-6.41	96.19	102.60
34	BA	609	G	C4-N9-C1'	-6.41	118.16	126.50
34	BA	979	G	O3'-P-O5'	6.41	116.18	104.00
34	BA	997	U	P-O5'-C5'	-6.41	110.64	120.90
34	BA	1778	U	C1'-O4'-C4'	-6.41	104.77	109.90
35	BB	440	U	P-O5'-C5'	6.41	131.16	120.90
35	BB	481	A	N9-C1'-C2'	-6.41	104.95	112.00
35	BB	918	C	O4'-C1'-N1	6.41	113.33	108.20
72	Bm	12	ARG	NE-CZ-NH1	6.41	123.51	120.30
85	AA	526	G	C4'-C3'-C2'	-6.41	96.19	102.60
85	AA	661	C	O4'-C1'-N1	6.41	113.33	108.20
85	AA	2110	U	O4'-C1'-N1	6.41	113.33	108.20
86	AB	70	G	C4'-C3'-C2'	-6.41	96.19	102.60
34	BA	314	A	C5'-C4'-C3'	-6.41	105.74	116.00
34	BA	1204	U	N1-C2-N3	6.41	118.75	114.90
34	BA	1328	U	P-O3'-C3'	-6.41	112.01	119.70
35	BB	145	G	C5'-C4'-O4'	-6.41	101.41	109.10
39	BF	51	C	OP2-P-O3'	6.41	119.31	105.20
85	AA	277	G	N9-C1'-C2'	-6.41	104.95	112.00
85	AA	750	A	O3'-P-O5'	6.41	116.18	104.00
85	AA	2056	C	C6-N1-C2	-6.41	117.73	120.30
25	AR	58	CYS	CA-CB-SG	-6.41	102.46	114.00
34	BA	351	A	C5-C6-N1	6.41	120.91	117.70
34	BA	1435	A	P-O3'-C3'	-6.41	112.01	119.70
34	BA	1549	U	O5'-C5'-C4'	-6.41	99.52	111.70
34	BA	1718	C	N3-C4-N4	6.41	122.49	118.00
35	BB	555	G	P-O5'-C5'	-6.41	110.64	120.90
35	BB	806	U	C6-N1-C2	-6.41	117.15	121.00
35	BB	1110	G	C5-C6-O6	-6.41	124.75	128.60
35	BB	1212	C	N3-C2-O2	-6.41	117.41	121.90
35	BB	1228	A	P-O3'-C3'	-6.41	112.01	119.70
36	BC	71	A	O4'-C1'-N9	6.41	113.33	108.20
38	BE	130	G	N3-C4-N9	6.41	129.85	126.00
39	BF	13	U	C6-N1-C1'	6.41	130.17	121.20
39	BF	36	G	C8-N9-C1'	6.41	135.33	127.00
71	Bl	75	TYR	CB-CG-CD2	-6.41	117.15	121.00
71	Bl	93	VAL	CA-CB-CG1	6.41	120.52	110.90
77	Br	204	ARG	NE-CZ-NH2	-6.41	117.09	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	315	U	N3-C2-O2	-6.41	117.71	122.20
85	AA	541	A	O5'-C5'-C4'	-6.41	99.52	111.70
85	AA	889	G	C5'-C4'-C3'	-6.41	105.74	116.00
34	BA	691	A	C4-C5-C6	-6.41	113.80	117.00
35	BB	824	C	C4'-C3'-C2'	-6.41	96.19	102.60
35	BB	1371	G	C3'-C2'-C1'	-6.41	96.37	101.50
35	BB	1510	G	C1'-O4'-C4'	-6.41	104.77	109.90
36	BC	15	G	C3'-C2'-C1'	-6.41	96.37	101.50
38	BE	176	G	N1-C2-N3	6.41	127.75	123.90
40	BG	24	A	O4'-C1'-N9	6.41	113.33	108.20
85	AA	49	C	O4'-C1'-N1	6.41	113.33	108.20
85	AA	974	U	O4'-C1'-C2'	6.41	113.37	107.60
26	AS	11	ARG	NE-CZ-NH1	6.41	123.50	120.30
34	BA	717	U	C4-C5-C6	-6.41	115.86	119.70
35	BB	1144	A	C5'-C4'-C3'	-6.41	105.75	116.00
48	BO	149	ARG	NE-CZ-NH1	6.41	123.50	120.30
85	AA	315	U	P-O3'-C3'	6.41	127.39	119.70
86	AB	1	G	C5-C6-O6	-6.41	124.76	128.60
86	AB	22	G	C4-N9-C1'	6.41	134.83	126.50
34	BA	323	C	O3'-P-O5'	-6.41	91.83	104.00
34	BA	867	C	O3'-P-O5'	-6.41	91.83	104.00
34	BA	1321	A	C3'-C2'-C1'	-6.41	96.38	101.50
34	BA	1639	U	N3-C2-O2	-6.41	117.72	122.20
35	BB	40	C	C2-N1-C1'	-6.41	111.75	118.80
35	BB	363	A	C2-N3-C4	6.41	113.80	110.60
35	BB	1093	C	C5'-C4'-O4'	6.41	116.79	109.10
40	BG	119	A	C4'-C3'-C2'	-6.41	96.19	102.60
85	AA	985	G	C4'-C3'-C2'	-6.41	96.19	102.60
85	AA	1106	A	N1-C2-N3	6.41	132.50	129.30
85	AA	1141	U	C5'-C4'-C3'	-6.41	105.75	116.00
22	AO	145	ARG	NE-CZ-NH2	-6.40	117.10	120.30
34	BA	4	A	P-O5'-C5'	6.40	131.15	120.90
34	BA	1215	U	C4'-C3'-C2'	6.40	109.00	102.60
34	BA	1603	A	P-O5'-C5'	-6.40	110.65	120.90
35	BB	461	U	C5-C4-O4	-6.40	122.06	125.90
51	BR	3	HIS	CA-CB-CG	-6.40	102.71	113.60
85	AA	177	A	C1'-O4'-C4'	-6.40	104.78	109.90
85	AA	617	C	C6-N1-C1'	-6.40	113.12	120.80
5	A4	146	ARG	CB-CA-C	-6.40	97.59	110.40
34	BA	71	G	C5-C6-N1	6.40	114.70	111.50
34	BA	471	U	N1-C2-N3	6.40	118.74	114.90
34	BA	1366	C	O4'-C1'-N1	6.40	113.32	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1543	A	C1'-O4'-C4'	-6.40	104.78	109.90
35	BB	56	U	N3-C2-O2	-6.40	117.72	122.20
35	BB	616	U	C5'-C4'-C3'	-6.40	105.76	116.00
38	BE	135	A	O4'-C1'-C2'	-6.40	99.40	105.80
38	BE	148	C	O4'-C1'-N1	6.40	113.32	108.20
38	BE	196	C	N3-C2-O2	-6.40	117.42	121.90
38	BE	200	A	N1-C6-N6	6.40	122.44	118.60
54	BU	149	ARG	NE-CZ-NH1	6.40	123.50	120.30
85	AA	761	G	C8-N9-C1'	-6.40	118.68	127.00
85	AA	974	U	O3'-P-O5'	6.40	116.17	104.00
85	AA	1179	A	O4'-C1'-N9	6.40	113.32	108.20
85	AA	1992	A	C8-N9-C4	-6.40	103.24	105.80
23	AP	253	HIS	CA-CB-CG	-6.40	102.72	113.60
34	BA	115	U	C5'-C4'-O4'	6.40	116.78	109.10
34	BA	301	U	O5'-C5'-C4'	6.40	123.86	111.70
34	BA	488	C	C6-N1-C2	-6.40	117.74	120.30
35	BB	735	A	O4'-C1'-N9	6.40	113.32	108.20
35	BB	1376	G	P-O3'-C3'	-6.40	112.02	119.70
35	BB	1421	C	C6-N1-C2	-6.40	117.74	120.30
36	BC	23	G	C8-N9-C4	-6.40	103.84	106.40
36	BC	48	A	P-O3'-C3'	6.40	127.38	119.70
36	BC	92	C	O3'-P-O5'	-6.40	91.84	104.00
40	BG	26	G	C3'-C2'-C1'	-6.40	96.38	101.50
40	BG	79	U	P-O3'-C3'	-6.40	112.02	119.70
41	BH	25	A	C2'-C3'-O3'	6.40	123.94	113.70
85	AA	857	G	O4'-C1'-N9	6.40	113.32	108.20
85	AA	2107	C	O3'-P-O5'	6.40	116.16	104.00
85	AA	2227	A	O5'-C5'-C4'	-6.40	99.54	111.70
34	BA	143	A	C3'-C2'-C1'	-6.40	96.38	101.50
34	BA	605	G	C6-C5-N7	-6.40	126.56	130.40
34	BA	736	G	C4'-C3'-C2'	-6.40	96.20	102.60
35	BB	836	U	C5-C6-N1	6.40	125.90	122.70
37	BD	49	A	O4'-C1'-N9	6.40	113.32	108.20
37	BD	57	C	C5'-C4'-O4'	6.40	116.78	109.10
85	AA	466	A	C6-C5-N7	6.40	136.78	132.30
85	AA	510	A	O5'-C5'-C4'	6.40	123.86	111.70
8	A7	5	TYR	CB-CG-CD1	6.40	124.84	121.00
34	BA	1001	G	C3'-C2'-C1'	-6.40	96.38	101.50
35	BB	125	G	C8-N9-C4	-6.40	103.84	106.40
35	BB	994	A	C3'-C2'-C1'	6.40	106.62	101.50
36	BC	88	A	C6-N1-C2	-6.40	114.76	118.60
41	BH	92	A	N1-C2-N3	6.40	132.50	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	131	C	O3'-P-O5'	-6.40	91.85	104.00
85	AA	486	G	P-O5'-C5'	-6.40	110.66	120.90
85	AA	1473	U	O4'-C1'-N1	6.40	113.32	108.20
85	AA	1786	G	C8-N9-C4	-6.40	103.84	106.40
85	AA	2048	C	N3-C2-O2	-6.40	117.42	121.90
34	BA	483	A	O5'-P-OP1	-6.40	99.94	105.70
34	BA	815	C	C6-N1-C2	-6.40	117.74	120.30
35	BB	796	C	O4'-C4'-C3'	-6.40	97.60	104.00
36	BC	87	C	P-O5'-C5'	-6.40	110.67	120.90
64	Be	69	TYR	CB-CG-CD2	-6.40	117.16	121.00
34	BA	639	U	OP1-P-OP2	-6.39	110.01	119.60
34	BA	1190	A	C4-N9-C1'	-6.39	114.79	126.30
34	BA	1197	U	C5'-C4'-O4'	-6.39	101.43	109.10
35	BB	118	A	C5-C6-N6	-6.39	118.58	123.70
35	BB	1291	G	N9-C1'-C2'	-6.39	104.97	112.00
35	BB	1532	C	N3-C2-O2	-6.39	117.42	121.90
38	BE	96	G	N9-C4-C5	6.39	107.96	105.40
85	AA	937	G	N1-C6-O6	6.39	123.74	119.90
86	AB	65	G	P-O3'-C3'	-6.39	112.03	119.70
34	BA	486	G	C1'-O4'-C4'	6.39	115.01	109.90
34	BA	1778	U	C5'-C4'-C3'	-6.39	105.77	116.00
35	BB	1236	A	C4'-C3'-C2'	-6.39	96.21	102.60
35	BB	1536	G	N1-C6-O6	6.39	123.74	119.90
36	BC	14	G	O4'-C1'-N9	6.39	113.31	108.20
37	BD	36	C	P-O3'-C3'	-6.39	112.03	119.70
40	BG	20	U	C4'-C3'-C2'	-6.39	96.21	102.60
67	Bh	38	TRP	CB-CG-CD2	-6.39	118.29	126.60
85	AA	31	C	C1'-O4'-C4'	-6.39	104.79	109.90
85	AA	151	A	N1-C6-N6	-6.39	114.76	118.60
85	AA	2043	A	P-O3'-C3'	-6.39	112.03	119.70
85	AA	2152	C	C6-N1-C2	-6.39	117.74	120.30
86	AB	53	G	C5'-C4'-C3'	-6.39	105.77	116.00
34	BA	868	C	O5'-P-OP2	-6.39	99.95	105.70
34	BA	1163	G	N1-C2-N2	-6.39	110.45	116.20
84	By	180	GLN	N-CA-CB	-6.39	99.10	110.60
85	AA	100	A	C4-N9-C1'	-6.39	114.80	126.30
85	AA	1458	G	C4-C5-N7	-6.39	108.24	110.80
85	AA	1760	C	C6-N1-C2	-6.39	117.74	120.30
34	BA	16	C	C2-N3-C4	-6.39	116.70	119.90
34	BA	316	G	O5'-C5'-C4'	-6.39	99.56	111.70
34	BA	353	U	C5'-C4'-C3'	-6.39	105.78	116.00
34	BA	1726	U	C5-C4-O4	6.39	129.73	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	487	A	C8-N9-C4	6.39	108.36	105.80
40	BG	140	G	C1'-O4'-C4'	-6.39	104.79	109.90
85	AA	88	G	O4'-C1'-N9	6.39	113.31	108.20
85	AA	110	U	C4'-C3'-C2'	6.39	108.99	102.60
85	AA	631	G	N3-C2-N2	6.39	124.37	119.90
85	AA	986	U	N1-C2-N3	6.39	118.73	114.90
8	A7	148	ARG	NE-CZ-NH2	-6.39	117.11	120.30
34	BA	1291	A	O4'-C1'-C2'	6.39	113.35	107.60
38	BE	104	G	N3-C2-N2	6.39	124.37	119.90
85	AA	574	U	C1'-O4'-C4'	-6.39	104.79	109.90
85	AA	771	A	P-O3'-C3'	6.39	127.36	119.70
85	AA	978	U	N1-C1'-C2'	-6.39	104.97	112.00
85	AA	1449	C	C2-N1-C1'	-6.39	111.77	118.80
25	AR	64	ARG	NE-CZ-NH1	6.39	123.49	120.30
26	AS	13	LEU	CB-CA-C	-6.39	98.06	110.20
34	BA	22	C	C6-N1-C2	-6.39	117.75	120.30
34	BA	1018	U	C5-C4-O4	6.39	129.73	125.90
34	BA	1521	C	N3-C2-O2	-6.39	117.43	121.90
35	BB	23	U	P-O3'-C3'	6.39	127.36	119.70
35	BB	472	C	C2-N3-C4	-6.39	116.71	119.90
35	BB	642	G	N3-C4-N9	6.39	129.83	126.00
35	BB	1183	U	C1'-O4'-C4'	-6.39	104.79	109.90
36	BC	12	A	P-O3'-C3'	-6.39	112.04	119.70
37	BD	86	A	C4-N9-C1'	-6.39	114.81	126.30
39	BF	42	G	C8-N9-C1'	6.39	135.30	127.00
53	BT	187	ARG	NE-CZ-NH1	6.39	123.49	120.30
85	AA	157	G	N7-C8-N9	-6.39	109.91	113.10
85	AA	258	G	C3'-C2'-C1'	-6.39	96.39	101.50
85	AA	967	C	C4'-C3'-C2'	-6.39	96.21	102.60
85	AA	1120	G	C3'-C2'-C1'	-6.39	96.39	101.50
85	AA	1186	C	P-O3'-C3'	-6.39	112.04	119.70
85	AA	1256	C	C5'-C4'-C3'	-6.39	105.78	116.00
85	AA	1985	C	C3'-C2'-C1'	-6.39	96.39	101.50
34	BA	295	G	C4-N9-C1'	6.38	134.80	126.50
34	BA	904	G	C8-N9-C4	6.38	108.95	106.40
34	BA	1306	U	O4'-C1'-C2'	-6.38	99.42	105.80
34	BA	1613	G	C5-N7-C8	-6.38	101.11	104.30
35	BB	130	G	N9-C1'-C2'	-6.38	104.98	112.00
37	BD	58	G	C8-N9-C1'	6.38	135.30	127.00
38	BE	192	A	C5-C6-N6	6.38	128.81	123.70
41	BH	33	G	P-O5'-C5'	6.38	131.12	120.90
85	AA	373	G	C8-N9-C4	6.38	108.95	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	552	C	C5-C6-N1	6.38	124.19	121.00
85	AA	596	A	C6-N1-C2	-6.38	114.77	118.60
34	BA	808	U	O4'-C1'-N1	6.38	113.31	108.20
38	BE	25	U	C5'-C4'-C3'	6.38	126.21	116.00
85	AA	110	U	O3'-P-O5'	6.38	116.13	104.00
85	AA	251	A	C5'-C4'-O4'	6.38	116.76	109.10
85	AA	565	G	P-O3'-C3'	-6.38	112.04	119.70
85	AA	806	G	O5'-C5'-C4'	-6.38	99.57	111.70
85	AA	1993	C	O4'-C1'-N1	6.38	113.31	108.20
25	AR	84	ARG	NE-CZ-NH1	6.38	123.49	120.30
34	BA	234	A	C8-N9-C1'	6.38	139.19	127.70
34	BA	471	U	C4'-C3'-C2'	6.38	108.98	102.60
34	BA	1343	A	C3'-C2'-C1'	-6.38	96.39	101.50
34	BA	1560	U	P-O3'-C3'	-6.38	112.04	119.70
35	BB	442	U	C1'-O4'-C4'	-6.38	104.80	109.90
35	BB	463	C	O4'-C1'-N1	6.38	113.31	108.20
35	BB	496	C	C1'-O4'-C4'	-6.38	104.79	109.90
35	BB	1298	C	O4'-C1'-N1	6.38	113.31	108.20
36	BC	123	G	N3-C4-N9	-6.38	122.17	126.00
37	BD	95	G	C6-N1-C2	-6.38	121.27	125.10
38	BE	25	U	OP1-P-OP2	-6.38	110.03	119.60
41	BH	67	G	C8-N9-C1'	6.38	135.30	127.00
58	BY	81	ARG	NE-CZ-NH1	6.38	123.49	120.30
82	Bw	150	ALA	N-CA-CB	6.38	119.03	110.10
85	AA	429	G	C1'-O4'-C4'	-6.38	104.80	109.90
85	AA	2058	C	C5'-C4'-C3'	6.38	126.21	116.00
34	BA	1477	C	C6-N1-C1'	-6.38	113.14	120.80
35	BB	972	C	C1'-O4'-C4'	-6.38	104.80	109.90
35	BB	1485	G	C4-N9-C1'	-6.38	118.21	126.50
39	BF	14	C	C6-N1-C1'	-6.38	113.14	120.80
49	BP	11	ARG	NE-CZ-NH2	-6.38	117.11	120.30
85	AA	144	A	P-O3'-C3'	6.38	127.36	119.70
85	AA	271	A	N1-C2-N3	-6.38	126.11	129.30
85	AA	789	A	O5'-P-OP2	6.38	118.36	110.70
85	AA	1288	A	C2-N3-C4	6.38	113.79	110.60
34	BA	755	G	C1'-O4'-C4'	-6.38	104.80	109.90
34	BA	1322	A	C1'-O4'-C4'	-6.38	104.80	109.90
34	BA	1392	A	P-O3'-C3'	-6.38	112.05	119.70
35	BB	356	C	C1'-O4'-C4'	-6.38	104.80	109.90
35	BB	475	A	P-O5'-C5'	6.38	131.11	120.90
35	BB	1065	G	C8-N9-C4	6.38	108.95	106.40
35	BB	1330	A	P-O3'-C3'	-6.38	112.05	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	39	A	P-O5'-C5'	-6.38	110.69	120.90
85	AA	964	C	P-O5'-C5'	-6.38	110.69	120.90
85	AA	975	G	C3'-C2'-C1'	-6.38	96.40	101.50
85	AA	1993	C	P-O3'-C3'	-6.38	112.05	119.70
34	BA	134	U	C1'-O4'-C4'	-6.38	104.80	109.90
34	BA	632	U	C5'-C4'-C3'	-6.38	105.80	116.00
34	BA	952	G	C5'-C4'-C3'	6.38	126.20	116.00
34	BA	995	A	O4'-C1'-N9	6.38	113.30	108.20
34	BA	1267	A	O4'-C4'-C3'	6.38	111.20	106.10
34	BA	1353	U	O4'-C1'-N1	6.38	113.30	108.20
34	BA	1836	A	O4'-C1'-C2'	6.38	113.34	107.60
35	BB	441	G	C8-N9-C4	6.38	108.95	106.40
35	BB	998	G	N9-C1'-C2'	-6.38	104.99	112.00
35	BB	1004	A	C8-N9-C1'	6.38	139.18	127.70
35	BB	1413	U	N1-C2-N3	6.38	118.73	114.90
39	BF	35	C	P-O3'-C3'	-6.38	112.05	119.70
85	AA	657	C	N3-C2-O2	-6.38	117.44	121.90
85	AA	725	G	C4'-C3'-C2'	-6.38	96.22	102.60
85	AA	844	C	O4'-C1'-N1	6.38	113.30	108.20
34	BA	1679	C	N3-C2-O2	-6.38	117.44	121.90
35	BB	111	C	C5'-C4'-O4'	6.38	116.75	109.10
35	BB	958	C	C6-N1-C2	-6.38	117.75	120.30
35	BB	1059	U	P-O3'-C3'	-6.38	112.05	119.70
77	Br	306	GLN	N-CA-CB	6.38	122.07	110.60
85	AA	1130	G	N1-C6-O6	-6.38	116.08	119.90
85	AA	1367	C	C4'-C3'-C2'	-6.38	96.22	102.60
34	BA	684	G	C8-N9-C1'	-6.37	118.71	127.00
34	BA	939	C	C1'-O4'-C4'	-6.37	104.80	109.90
34	BA	1013	A	N1-C6-N6	6.37	122.42	118.60
40	BG	62	C	C5'-C4'-C3'	6.37	126.20	116.00
85	AA	86	G	N1-C6-O6	6.37	123.72	119.90
85	AA	393	C	N3-C2-O2	-6.37	117.44	121.90
85	AA	728	U	O3'-P-O5'	-6.37	91.89	104.00
85	AA	1795	C	O4'-C1'-N1	6.37	113.30	108.20
85	AA	2105	G	C6-N1-C2	-6.37	121.28	125.10
35	BB	928	C	O4'-C1'-N1	6.37	113.30	108.20
35	BB	936	U	O4'-C1'-N1	6.37	113.30	108.20
71	Bl	47	TYR	CB-CG-CD2	-6.37	117.18	121.00
85	AA	347	U	O4'-C1'-N1	6.37	113.30	108.20
85	AA	420	C	C1'-O4'-C4'	-6.37	104.80	109.90
85	AA	662	U	P-O3'-C3'	-6.37	112.05	119.70
85	AA	1478	G	N3-C2-N2	6.37	124.36	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1694	C	C6-N1-C2	-6.37	117.75	120.30
85	AA	1978	G	O4'-C1'-N9	6.37	113.30	108.20
34	BA	562	C	N1-C1'-C2'	-6.37	104.99	112.00
34	BA	1629	A	C6-N1-C2	-6.37	114.78	118.60
39	BF	30	C	C5'-C4'-C3'	-6.37	105.81	116.00
85	AA	1657	C	C4'-C3'-C2'	6.37	108.97	102.60
34	BA	160	G	C5'-C4'-C3'	-6.37	105.81	116.00
34	BA	312	U	N3-C2-O2	-6.37	117.74	122.20
34	BA	631	G	O4'-C1'-N9	6.37	113.30	108.20
34	BA	818	G	C3'-C2'-C1'	-6.37	96.40	101.50
34	BA	890	G	C8-N9-C1'	6.37	135.28	127.00
35	BB	615	A	N1-C6-N6	6.37	122.42	118.60
35	BB	746	A	C5'-C4'-O4'	6.37	116.74	109.10
38	BE	95	G	C4-N9-C1'	-6.37	118.22	126.50
62	Bc	108	ARG	NE-CZ-NH2	-6.37	117.12	120.30
85	AA	660	G	C5-C6-O6	-6.37	124.78	128.60
85	AA	1289	U	C2-N3-C4	-6.37	123.18	127.00
85	AA	1406	U	O4'-C1'-N1	6.37	113.30	108.20
34	BA	564	C	C5-C6-N1	-6.37	117.82	121.00
34	BA	1785	G	O4'-C1'-N9	6.37	113.29	108.20
35	BB	976	U	C6-N1-C2	-6.37	117.18	121.00
40	BG	181	C	O5'-C5'-C4'	-6.37	99.60	111.70
85	AA	33	U	C4'-C3'-C2'	-6.37	96.23	102.60
85	AA	134	U	O4'-C1'-N1	6.37	113.29	108.20
85	AA	268	A	C5'-C4'-C3'	-6.37	105.81	116.00
85	AA	1762	G	P-O3'-C3'	6.37	127.34	119.70
3	A2	184	ARG	NE-CZ-NH1	6.37	123.48	120.30
34	BA	253	U	C5-C4-O4	6.37	129.72	125.90
34	BA	818	G	C5-C6-O6	-6.37	124.78	128.60
35	BB	654	C	C2-N1-C1'	-6.37	111.80	118.80
35	BB	1313	C	N3-C2-O2	-6.37	117.44	121.90
35	BB	1434	G	P-O3'-C3'	-6.37	112.06	119.70
38	BE	95	G	C5-C6-N1	6.37	114.68	111.50
85	AA	176	C	O4'-C1'-N1	6.37	113.29	108.20
85	AA	847	G	C1'-O4'-C4'	-6.37	104.81	109.90
85	AA	869	A	C2-N3-C4	-6.37	107.42	110.60
85	AA	1175	A	P-O3'-C3'	-6.37	112.06	119.70
85	AA	1267	A	C1'-O4'-C4'	-6.37	104.81	109.90
34	BA	687	G	N1-C2-N2	-6.36	110.47	116.20
35	BB	15	C	O4'-C1'-N1	6.36	113.29	108.20
35	BB	80	C	O4'-C1'-N1	6.36	113.29	108.20
35	BB	98	A	O5'-P-OP2	-6.36	99.97	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	136	A	P-O5'-C5'	6.36	131.08	120.90
35	BB	779	C	C6-N1-C2	6.36	122.84	120.30
35	BB	854	G	C3'-C2'-C1'	-6.36	96.41	101.50
85	AA	577	U	C4'-C3'-C2'	-6.36	96.24	102.60
85	AA	865	G	C4'-C3'-C2'	-6.36	96.24	102.60
85	AA	894	A	C1'-O4'-C4'	-6.36	104.81	109.90
85	AA	925	G	P-O3'-C3'	-6.36	112.06	119.70
85	AA	1832	G	O4'-C1'-N9	6.36	113.29	108.20
35	BB	11	A	P-O3'-C3'	6.36	127.33	119.70
35	BB	137	A	C8-N9-C4	6.36	108.34	105.80
85	AA	924	A	C5'-C4'-O4'	6.36	116.73	109.10
85	AA	1615	A	C1'-O4'-C4'	-6.36	104.81	109.90
34	BA	89	G	O5'-C5'-C4'	-6.36	99.61	111.70
34	BA	719	G	C4'-C3'-C2'	-6.36	96.24	102.60
34	BA	856	G	O4'-C1'-N9	6.36	113.29	108.20
34	BA	1283	U	C2-N1-C1'	-6.36	110.07	117.70
34	BA	1625	C	C5'-C4'-O4'	6.36	116.73	109.10
35	BB	659	C	C2'-C3'-O3'	6.36	123.88	113.70
36	BC	36	G	C5-C6-O6	-6.36	124.78	128.60
53	BT	113	LYS	N-CA-CB	6.36	122.05	110.60
85	AA	155	U	P-O5'-C5'	-6.36	110.72	120.90
85	AA	680	U	O4'-C1'-C2'	-6.36	99.44	105.80
85	AA	1161	U	O4'-C1'-N1	6.36	113.29	108.20
85	AA	2041	G	N1-C2-N2	-6.36	110.48	116.20
34	BA	177	G	C6-N1-C2	-6.36	121.28	125.10
34	BA	515	U	O4'-C1'-C2'	6.36	113.32	107.60
34	BA	1841	A	P-O5'-C5'	6.36	131.07	120.90
35	BB	116	G	P-O5'-C5'	-6.36	110.72	120.90
61	Bb	19	TYR	CA-CB-CG	6.36	125.48	113.40
80	Bu	57	ASN	C-N-CA	6.36	137.60	121.70
85	AA	899	A	C3'-C2'-C1'	-6.36	96.41	101.50
34	BA	109	A	P-O3'-C3'	-6.36	112.07	119.70
34	BA	297	A	N9-C1'-C2'	6.36	122.26	114.00
34	BA	321	G	C5-N7-C8	6.36	107.48	104.30
34	BA	747	G	C6-N1-C2	-6.36	121.28	125.10
34	BA	856	G	P-O3'-C3'	-6.36	112.07	119.70
34	BA	1756	C	O4'-C1'-N1	6.36	113.29	108.20
35	BB	1333	U	C2-N3-C4	-6.36	123.19	127.00
37	BD	58	G	C4-N9-C1'	-6.36	118.23	126.50
40	BG	8	U	C5-C6-N1	-6.36	119.52	122.70
40	BG	103	C	C4'-C3'-C2'	-6.36	96.24	102.60
81	Bv	42	ARG	CG-CD-NE	-6.36	98.45	111.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	305	A	P-O3'-C3'	-6.36	112.07	119.70
85	AA	447	C	N1-C2-O2	6.36	122.72	118.90
85	AA	705	G	C4-N9-C1'	-6.36	118.23	126.50
85	AA	1115	G	C3'-C2'-C1'	-6.36	96.41	101.50
85	AA	1575	G	O4'-C1'-N9	6.36	113.29	108.20
85	AA	1579	A	P-O3'-C3'	-6.36	112.07	119.70
85	AA	1934	A	C1'-O4'-C4'	-6.36	104.81	109.90
5	A4	158	PHE	C-N-CA	6.36	137.59	121.70
34	BA	490	A	C8-N9-C4	6.36	108.34	105.80
34	BA	627	U	O4'-C1'-N1	6.36	113.28	108.20
34	BA	1000	G	C8-N9-C1'	6.36	135.26	127.00
35	BB	968	C	P-O3'-C3'	-6.36	112.07	119.70
38	BE	82	C	C2'-C3'-O3'	6.36	123.87	113.70
72	Bm	76	ARG	NE-CZ-NH2	6.36	123.48	120.30
85	AA	860	C	C6-N1-C2	-6.36	117.76	120.30
85	AA	906	U	N1-C1'-C2'	-6.36	105.01	112.00
85	AA	1466	U	O3'-P-O5'	6.36	116.07	104.00
20	AL	58	MET	N-CA-CB	6.35	122.04	110.60
34	BA	123	C	P-O3'-C3'	-6.35	112.08	119.70
34	BA	1285	G	N3-C2-N2	6.35	124.35	119.90
34	BA	1818	A	O3'-P-O5'	6.35	116.07	104.00
53	BT	88	ARG	NE-CZ-NH2	-6.35	117.12	120.30
85	AA	2225	G	C5-C6-N1	6.35	114.68	111.50
18	AJ	23	ARG	NE-CZ-NH1	6.35	123.48	120.30
34	BA	167	U	C5-C4-O4	6.35	129.71	125.90
34	BA	1001	G	C1'-O4'-C4'	-6.35	104.82	109.90
34	BA	1330	G	C4-N9-C1'	-6.35	118.24	126.50
34	BA	1459	U	P-O5'-C5'	-6.35	110.74	120.90
34	BA	1692	U	O4'-C1'-N1	6.35	113.28	108.20
34	BA	1801	G	O4'-C4'-C3'	-6.35	97.65	104.00
35	BB	845	C	C3'-C2'-C1'	-6.35	96.42	101.50
35	BB	1350	A	C6-N1-C2	-6.35	114.79	118.60
38	BE	51	C	C4'-C3'-C2'	6.35	108.95	102.60
39	BF	12	U	N1-C2-N3	-6.35	111.09	114.90
43	BJ	177	ARG	NE-CZ-NH1	6.35	123.48	120.30
62	Bc	137	HIS	CB-CA-C	-6.35	97.69	110.40
80	Bu	98	ALA	N-CA-CB	-6.35	101.21	110.10
85	AA	327	G	C4'-C3'-O3'	6.35	125.70	113.00
85	AA	684	G	OP2-P-O3'	6.35	119.18	105.20
85	AA	705	G	C1'-O4'-C4'	-6.35	104.82	109.90
85	AA	1267	A	O4'-C1'-N9	6.35	113.28	108.20
85	AA	1579	A	N1-C6-N6	-6.35	114.79	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1728	G	N9-C4-C5	-6.35	102.86	105.40
34	BA	1796	A	C3'-C2'-C1'	-6.35	96.42	101.50
35	BB	1251	G	C5-C6-N1	6.35	114.68	111.50
35	BB	1533	U	C2-N1-C1'	6.35	125.32	117.70
41	BH	119	U	C5-C6-N1	-6.35	119.52	122.70
85	AA	189	G	C5-C6-N1	6.35	114.68	111.50
85	AA	767	A	N1-C2-N3	-6.35	126.12	129.30
85	AA	2218	G	N7-C8-N9	-6.35	109.92	113.10
2	A1	46	ARG	NE-CZ-NH1	6.35	123.47	120.30
34	BA	522	C	O4'-C1'-N1	6.35	113.28	108.20
34	BA	645	U	O4'-C1'-N1	6.35	113.28	108.20
34	BA	823	G	N1-C6-O6	-6.35	116.09	119.90
34	BA	1026	C	C5'-C4'-C3'	6.35	126.16	116.00
34	BA	1069	U	P-O3'-C3'	-6.35	112.08	119.70
35	BB	48	G	O4'-C1'-N9	6.35	113.28	108.20
35	BB	494	C	P-O3'-C3'	6.35	127.32	119.70
35	BB	1102	U	C2-N1-C1'	-6.35	110.08	117.70
35	BB	1171	U	P-O5'-C5'	6.35	131.06	120.90
36	BC	10	C	P-O3'-C3'	-6.35	112.08	119.70
36	BC	91	G	N1-C6-O6	-6.35	116.09	119.90
62	Bc	142	ARG	N-CA-C	6.35	128.14	111.00
85	AA	159	G	O4'-C1'-C2'	6.35	113.31	107.60
85	AA	1135	U	C2-N1-C1'	-6.35	110.08	117.70
85	AA	1491	G	C6-N1-C2	-6.35	121.29	125.10
5	A4	47	HIS	CA-CB-CG	-6.35	102.81	113.60
34	BA	1143	U	O5'-C5'-C4'	-6.35	99.64	111.70
34	BA	1495	A	C2'-C3'-O3'	6.35	123.86	113.70
35	BB	1365	G	P-O5'-C5'	-6.35	110.75	120.90
38	BE	64	A	C8-N9-C4	6.35	108.34	105.80
66	Bg	94	ARG	NE-CZ-NH2	-6.35	117.13	120.30
85	AA	1291	A	C5'-C4'-O4'	6.35	116.72	109.10
85	AA	1522	U	C4-C5-C6	-6.35	115.89	119.70
85	AA	1829	C	N3-C2-O2	-6.35	117.46	121.90
85	AA	2044	A	C5'-C4'-C3'	-6.35	105.84	116.00
34	BA	1039	G	C5'-C4'-C3'	-6.35	105.85	116.00
35	BB	420	U	C4'-C3'-C2'	-6.35	96.25	102.60
35	BB	760	C	O4'-C1'-N1	6.35	113.28	108.20
35	BB	1197	G	C8-N9-C4	6.35	108.94	106.40
85	AA	484	G	P-O3'-C3'	-6.35	112.08	119.70
85	AA	936	C	C2-N3-C4	-6.35	116.73	119.90
34	BA	418	G	C3'-C2'-C1'	-6.34	96.42	101.50
34	BA	561	U	OP1-P-OP2	-6.34	110.08	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	583	G	N3-C2-N2	6.34	124.34	119.90
34	BA	634	U	O4'-C1'-N1	6.34	113.28	108.20
34	BA	1522	G	C1'-O4'-C4'	-6.34	104.82	109.90
34	BA	1774	G	C8-N9-C1'	6.34	135.25	127.00
35	BB	298	G	O4'-C1'-N9	6.34	113.28	108.20
35	BB	954	G	O4'-C1'-N9	6.34	113.28	108.20
38	BE	85	G	O4'-C1'-N9	6.34	113.28	108.20
40	BG	163	G	C2-N3-C4	-6.34	108.73	111.90
41	BH	86	G	N1-C6-O6	6.34	123.71	119.90
52	BS	82	TYR	CB-CG-CD2	-6.34	117.19	121.00
80	Bu	190	ASN	N-CA-CB	6.34	122.02	110.60
85	AA	1095	C	C6-N1-C1'	6.34	128.41	120.80
85	AA	1458	G	C5'-C4'-O4'	-6.34	101.49	109.10
85	AA	1544	G	N9-C1'-C2'	-6.34	105.02	112.00
34	BA	325	A	C2-N3-C4	-6.34	107.43	110.60
34	BA	622	G	N1-C6-O6	6.34	123.71	119.90
35	BB	568	A	C8-N9-C4	6.34	108.34	105.80
35	BB	954	G	C3'-C2'-C1'	-6.34	96.43	101.50
38	BE	47	U	P-O5'-C5'	-6.34	110.75	120.90
39	BF	46	G	O4'-C1'-N9	6.34	113.28	108.20
40	BG	9	G	C4-C5-N7	-6.34	108.26	110.80
85	AA	1210	U	P-O3'-C3'	-6.34	112.09	119.70
85	AA	1365	U	C2-N3-C4	-6.34	123.19	127.00
85	AA	1797	U	C5'-C4'-C3'	-6.34	105.85	116.00
85	AA	2069	A	C3'-C2'-C1'	-6.34	96.43	101.50
31	AX	123	ARG	NE-CZ-NH1	6.34	123.47	120.30
34	BA	492	G	O4'-C1'-N9	6.34	113.27	108.20
34	BA	828	A	O5'-P-OP2	-6.34	99.99	105.70
34	BA	1415	C	C5'-C4'-C3'	-6.34	105.85	116.00
35	BB	546	A	C5-C6-N1	6.34	120.87	117.70
35	BB	1239	A	N1-C6-N6	-6.34	114.80	118.60
65	Bf	410	GLU	C-N-CA	6.34	137.55	121.70
65	Bf	448	ARG	NE-CZ-NH1	6.34	123.47	120.30
85	AA	281	C	P-O5'-C5'	-6.34	110.75	120.90
85	AA	423	G	C5'-C4'-C3'	-6.34	105.86	116.00
85	AA	768	C	C2'-C3'-O3'	6.34	123.85	113.70
85	AA	980	U	P-O5'-C5'	-6.34	110.75	120.90
3	A2	57	ARG	NE-CZ-NH2	-6.34	117.13	120.30
34	BA	137	C	C6-N1-C2	-6.34	117.76	120.30
34	BA	1001	G	C5-C6-N1	6.34	114.67	111.50
34	BA	1203	G	C1'-O4'-C4'	-6.34	104.83	109.90
34	BA	1210	A	C2-N3-C4	6.34	113.77	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	423	G	C5-C6-N1	6.34	114.67	111.50
37	BD	116	C	C1'-O4'-C4'	-6.34	104.83	109.90
39	BF	62	U	C4'-C3'-C2'	-6.34	96.26	102.60
85	AA	438	G	N9-C1'-C2'	-6.34	105.03	112.00
85	AA	1489	G	N9-C4-C5	-6.34	102.86	105.40
85	AA	1790	G	C5'-C4'-C3'	6.34	126.14	116.00
85	AA	1840	C	O5'-C5'-C4'	-6.34	99.65	111.70
85	AA	1881	C	C1'-O4'-C4'	-6.34	104.83	109.90
85	AA	1916	A	C5'-C4'-C3'	-6.34	105.86	116.00
34	BA	348	U	N3-C2-O2	-6.34	117.76	122.20
34	BA	1545	C	C5-C6-N1	-6.34	117.83	121.00
34	BA	1642	A	C1'-O4'-C4'	-6.34	104.83	109.90
34	BA	1696	G	C5-C6-N1	6.34	114.67	111.50
35	BB	673	C	P-O5'-C5'	6.34	131.04	120.90
35	BB	899	C	N3-C2-O2	-6.34	117.46	121.90
35	BB	1422	G	C6-C5-N7	-6.34	126.60	130.40
36	BC	8	C	C2'-C3'-O3'	6.34	123.84	113.70
36	BC	24	G	C5-C6-O6	-6.34	124.80	128.60
37	BD	93	G	O3'-P-O5'	-6.34	91.96	104.00
39	BF	32	G	C6-N1-C2	-6.34	121.30	125.10
85	AA	1057	G	C1'-O4'-C4'	-6.34	104.83	109.90
85	AA	1261	U	C3'-C2'-C1'	-6.34	96.43	101.50
85	AA	1574	C	O3'-P-O5'	-6.34	91.96	104.00
13	AE	113	TYR	CB-CG-CD1	-6.34	117.20	121.00
34	BA	84	U	C2'-C3'-O3'	6.34	123.84	113.70
34	BA	1295	U	C4'-C3'-O3'	6.34	125.67	113.00
34	BA	1336	U	C4'-C3'-C2'	-6.34	96.26	102.60
34	BA	1589	U	C6-N1-C2	-6.34	117.20	121.00
40	BG	15	G	C4'-C3'-C2'	6.34	108.94	102.60
85	AA	313	A	N9-C4-C5	-6.34	103.27	105.80
2	A1	48	ARG	NE-CZ-NH2	-6.33	117.13	120.30
34	BA	942	G	P-O3'-C3'	-6.33	112.10	119.70
34	BA	1158	A	C3'-C2'-C1'	-6.33	96.43	101.50
34	BA	1488	C	C4-C5-C6	-6.33	114.23	117.40
35	BB	368	C	N1-C2-O2	6.33	122.70	118.90
35	BB	828	G	N3-C2-N2	6.33	124.33	119.90
35	BB	974	C	C6-N1-C1'	-6.33	113.20	120.80
85	AA	213	G	N1-C6-O6	6.33	123.70	119.90
85	AA	260	A	C4-C5-C6	-6.33	113.83	117.00
85	AA	293	A	N1-C6-N6	6.33	122.40	118.60
85	AA	1159	C	C2-N1-C1'	-6.33	111.83	118.80
85	AA	1472	G	C1'-O4'-C4'	-6.33	104.83	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	AV	17	ARG	N-CA-CB	6.33	122.00	110.60
34	BA	800	G	C5-C6-N1	6.33	114.67	111.50
34	BA	890	G	P-O5'-C5'	-6.33	110.77	120.90
34	BA	983	A	N1-C6-N6	6.33	122.40	118.60
34	BA	1347	G	C4-N9-C1'	-6.33	118.27	126.50
35	BB	373	C	N3-C2-O2	-6.33	117.47	121.90
35	BB	992	C	O4'-C1'-N1	6.33	113.27	108.20
36	BC	22	U	P-O3'-C3'	6.33	127.30	119.70
38	BE	147	G	C4-N9-C1'	-6.33	118.27	126.50
40	BG	171	A	C5'-C4'-O4'	6.33	116.70	109.10
85	AA	82	A	C3'-C2'-C1'	-6.33	96.43	101.50
85	AA	820	G	C2'-C3'-O3'	6.33	123.83	113.70
85	AA	1874	G	C1'-O4'-C4'	-6.33	104.83	109.90
34	BA	624	G	C5-C6-O6	-6.33	124.80	128.60
34	BA	893	U	O4'-C1'-N1	6.33	113.27	108.20
34	BA	1724	G	C6-N1-C2	-6.33	121.30	125.10
35	BB	1215	U	C6-N1-C1'	6.33	130.06	121.20
35	BB	1221	G	C4-C5-C6	-6.33	115.00	118.80
42	BI	191	ARG	NE-CZ-NH1	6.33	123.47	120.30
34	BA	746	C	N1-C2-O2	6.33	122.70	118.90
34	BA	1150	A	N1-C6-N6	-6.33	114.80	118.60
34	BA	1247	G	C8-N9-C4	6.33	108.93	106.40
34	BA	1801	G	O4'-C1'-C2'	6.33	113.30	107.60
35	BB	1408	G	C5-C6-O6	-6.33	124.80	128.60
37	BD	101	A	C5-N7-C8	-6.33	100.73	103.90
76	Bq	30	ARG	NE-CZ-NH1	6.33	123.47	120.30
85	AA	1386	C	O4'-C1'-N1	6.33	113.26	108.20
85	AA	2190	U	C6-N1-C1'	6.33	130.06	121.20
34	BA	33	C	P-O3'-C3'	-6.33	112.11	119.70
34	BA	585	G	C8-N9-C4	-6.33	103.87	106.40
34	BA	862	C	C5-C4-N4	-6.33	115.77	120.20
34	BA	1733	G	C8-N9-C1'	6.33	135.23	127.00
35	BB	478	G	P-O5'-C5'	6.33	131.03	120.90
35	BB	795	A	C8-N9-C1'	6.33	139.09	127.70
38	BE	44	C	O4'-C1'-N1	6.33	113.26	108.20
41	BH	47	G	C1'-O4'-C4'	-6.33	104.84	109.90
85	AA	448	G	N3-C2-N2	6.33	124.33	119.90
85	AA	521	A	C2-N3-C4	-6.33	107.44	110.60
85	AA	698	G	C3'-C2'-C1'	-6.33	96.44	101.50
85	AA	778	C	O4'-C1'-N1	6.33	113.26	108.20
85	AA	1969	A	O4'-C1'-N9	6.33	113.26	108.20
34	BA	897	U	P-O5'-C5'	6.33	131.02	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1555	G	C8-N9-C1'	6.33	135.22	127.00
35	BB	517	G	C4-N9-C1'	-6.33	118.28	126.50
38	BE	115	U	C4'-C3'-C2'	-6.33	96.27	102.60
38	BE	170	U	O3'-P-O5'	-6.33	91.98	104.00
85	AA	838	G	N3-C4-C5	-6.33	125.44	128.60
85	AA	1903	G	C4'-C3'-C2'	-6.33	96.27	102.60
34	BA	53	G	C4'-C3'-C2'	6.33	108.93	102.60
34	BA	208	A	C8-N9-C4	-6.33	103.27	105.80
34	BA	287	U	P-O5'-C5'	-6.33	110.78	120.90
34	BA	506	U	P-O3'-C3'	-6.33	112.11	119.70
34	BA	808	U	P-O3'-C3'	-6.33	112.11	119.70
34	BA	926	A	C4-C5-C6	-6.33	113.84	117.00
35	BB	274	U	O4'-C1'-N1	6.33	113.26	108.20
35	BB	598	C	C3'-C2'-C1'	-6.33	96.44	101.50
35	BB	1404	A	C8-N9-C4	6.33	108.33	105.80
41	BH	35	G	C1'-O4'-C4'	-6.33	104.84	109.90
41	BH	101	A	C5'-C4'-C3'	6.33	126.12	116.00
85	AA	854	A	N1-C6-N6	-6.33	114.80	118.60
85	AA	1240	A	N9-C1'-C2'	-6.33	105.04	112.00
85	AA	1863	A	C1'-O4'-C4'	-6.33	104.84	109.90
85	AA	1963	G	N1-C6-O6	6.33	123.70	119.90
85	AA	1985	C	P-O5'-C5'	-6.33	110.78	120.90
85	AA	2136	C	O4'-C1'-N1	6.33	113.26	108.20
34	BA	446	U	P-O3'-C3'	-6.32	112.11	119.70
34	BA	1077	G	O4'-C1'-N9	6.32	113.26	108.20
34	BA	1348	G	C4'-C3'-C2'	-6.32	96.28	102.60
34	BA	1355	G	C5-C6-O6	-6.32	124.81	128.60
35	BB	390	G	P-O3'-C3'	-6.32	112.11	119.70
35	BB	823	G	N9-C1'-C2'	-6.32	105.04	112.00
35	BB	1417	C	C2-N1-C1'	-6.32	111.84	118.80
35	BB	1508	G	C5'-C4'-C3'	6.32	126.12	116.00
37	BD	54	A	P-O3'-C3'	-6.32	112.11	119.70
37	BD	71	G	C5-C6-N1	6.32	114.66	111.50
39	BF	69	A	C8-N9-C4	6.32	108.33	105.80
51	BR	53	ALA	N-CA-CB	6.32	118.95	110.10
61	Bb	35	ALA	N-CA-CB	6.32	118.95	110.10
65	Bf	451	ARG	NE-CZ-NH2	-6.32	117.14	120.30
83	Bx	110	TYR	CA-CB-CG	-6.32	101.39	113.40
85	AA	398	U	C2-N3-C4	-6.32	123.21	127.00
34	BA	99	G	C8-N9-C4	-6.32	103.87	106.40
34	BA	940	C	C6-N1-C2	-6.32	117.77	120.30
34	BA	1535	G	C5-C6-N1	6.32	114.66	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	360	C	C4'-C3'-C2'	6.32	108.92	102.60
85	AA	1903	G	O4'-C1'-N9	6.32	113.26	108.20
85	AA	2122	A	C8-N9-C1'	6.32	139.08	127.70
85	AA	2201	A	N9-C1'-C2'	-6.32	105.05	112.00
34	BA	855	C	C6-N1-C1'	-6.32	113.22	120.80
34	BA	1107	A	O4'-C1'-N9	6.32	113.26	108.20
34	BA	1193	A	P-O3'-C3'	6.32	127.28	119.70
34	BA	1489	U	O4'-C1'-N1	6.32	113.26	108.20
34	BA	1728	G	N9-C1'-C2'	-6.32	105.05	112.00
34	BA	1736	A	O3'-P-O5'	-6.32	91.99	104.00
35	BB	816	U	O3'-P-O5'	-6.32	91.99	104.00
35	BB	1015	U	C2-N3-C4	-6.32	123.21	127.00
36	BC	35	C	O4'-C1'-N1	6.32	113.26	108.20
85	AA	378	A	C5-C6-N1	6.32	120.86	117.70
85	AA	1450	U	N3-C4-C5	6.32	118.39	114.60
85	AA	2214	A	C1'-O4'-C4'	-6.32	104.84	109.90
86	AB	31	A	P-O3'-C3'	6.32	127.28	119.70
34	BA	754	G	C5-C6-N1	6.32	114.66	111.50
34	BA	1605	G	O5'-P-OP1	6.32	118.28	110.70
35	BB	503	G	P-O5'-C5'	-6.32	110.79	120.90
35	BB	957	A	O4'-C4'-C3'	-6.32	97.68	104.00
38	BE	200	A	C5-C6-N6	-6.32	118.64	123.70
85	AA	17	C	C5'-C4'-C3'	-6.32	105.89	116.00
85	AA	1022	G	C8-N9-C4	-6.32	103.87	106.40
85	AA	1293	U	P-O5'-C5'	6.32	131.01	120.90
85	AA	2103	C	C5'-C4'-C3'	-6.32	105.89	116.00
34	BA	344	G	P-O3'-C3'	6.32	127.28	119.70
34	BA	753	G	N1-C6-O6	6.32	123.69	119.90
34	BA	854	A	N1-C6-N6	6.32	122.39	118.60
34	BA	958	G	O3'-P-O5'	6.32	116.00	104.00
34	BA	1721	U	C5-C4-O4	6.32	129.69	125.90
35	BB	1354	C	C5'-C4'-C3'	-6.32	105.89	116.00
39	BF	2	G	O4'-C1'-N9	6.32	113.25	108.20
45	BL	130	TYR	CB-CG-CD1	6.32	124.79	121.00
85	AA	35	U	C1'-O4'-C4'	-6.32	104.85	109.90
85	AA	1205	U	C2-N3-C4	-6.32	123.21	127.00
85	AA	1557	U	C2-N1-C1'	-6.32	110.12	117.70
85	AA	1894	G	N1-C6-O6	6.32	123.69	119.90
24	AQ	56	ARG	NE-CZ-NH1	6.32	123.46	120.30
31	AX	116	ARG	NE-CZ-NH1	6.32	123.46	120.30
34	BA	379	C	P-O5'-C5'	6.32	131.01	120.90
34	BA	565	U	C1'-O4'-C4'	-6.32	104.85	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	965	A	O4'-C1'-N9	6.32	113.25	108.20
34	BA	1200	U	N3-C4-O4	6.32	123.82	119.40
34	BA	1637	G	N1-C6-O6	-6.32	116.11	119.90
34	BA	1679	C	C6-N1-C2	-6.32	117.77	120.30
35	BB	366	G	P-O5'-C5'	-6.32	110.80	120.90
35	BB	833	G	C6-N1-C2	-6.32	121.31	125.10
35	BB	864	U	C5'-C4'-C3'	-6.32	105.89	116.00
35	BB	1291	G	N1-C6-O6	-6.32	116.11	119.90
38	BE	72	C	C2-N3-C4	-6.32	116.74	119.90
40	BG	9	G	C4'-C3'-O3'	-6.32	96.14	109.40
57	BX	65	THR	N-CA-CB	6.32	122.30	110.30
85	AA	337	C	C3'-C2'-C1'	-6.32	96.45	101.50
85	AA	528	U	C2-N3-C4	-6.32	123.21	127.00
85	AA	1564	U	O4'-C1'-N1	6.32	113.25	108.20
85	AA	2056	C	N3-C2-O2	-6.32	117.48	121.90
34	BA	210	G	C5-C6-N1	6.31	114.66	111.50
35	BB	922	C	O4'-C1'-N1	6.31	113.25	108.20
37	BD	32	A	O4'-C1'-C2'	6.31	113.28	107.60
85	AA	544	A	C8-N9-C4	6.31	108.33	105.80
85	AA	736	U	O4'-C4'-C3'	-6.31	97.69	104.00
85	AA	863	C	C5'-C4'-C3'	-6.31	105.90	116.00
34	BA	512	U	C5'-C4'-O4'	6.31	116.67	109.10
34	BA	736	G	C4-C5-N7	6.31	113.33	110.80
34	BA	1459	U	P-O3'-C3'	6.31	127.28	119.70
34	BA	1737	A	C4'-C3'-C2'	-6.31	96.29	102.60
34	BA	1774	G	C4-N9-C1'	-6.31	118.29	126.50
35	BB	967	G	C1'-O4'-C4'	-6.31	104.85	109.90
35	BB	1156	U	C6-N1-C2	-6.31	117.21	121.00
35	BB	1185	G	C6-N1-C2	-6.31	121.31	125.10
35	BB	1246	C	C3'-C2'-C1'	-6.31	96.45	101.50
74	Bo	69	TYR	CB-CG-CD2	-6.31	117.21	121.00
85	AA	442	G	C1'-O4'-C4'	-6.31	104.85	109.90
85	AA	1117	G	O4'-C1'-N9	6.31	113.25	108.20
85	AA	1139	G	P-O5'-C5'	-6.31	110.80	120.90
85	AA	1200	A	O3'-P-O5'	6.31	115.99	104.00
85	AA	1935	G	N1-C6-O6	6.31	123.69	119.90
85	AA	2038	C	C6-N1-C2	-6.31	117.78	120.30
34	BA	739	A	N9-C1'-C2'	6.31	122.20	114.00
34	BA	1485	U	C2-N3-C4	-6.31	123.21	127.00
34	BA	1739	G	C1'-O4'-C4'	-6.31	104.85	109.90
34	BA	1792	U	C5'-C4'-C3'	6.31	126.10	116.00
35	BB	133	G	C5'-C4'-C3'	-6.31	105.90	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	BD	110	G	C5'-C4'-C3'	6.31	126.10	116.00
85	AA	658	C	C1'-O4'-C4'	-6.31	104.85	109.90
85	AA	785	C	C2'-C3'-O3'	6.31	123.80	113.70
85	AA	939	A	C3'-C2'-C1'	-6.31	96.45	101.50
85	AA	2195	A	N9-C1'-C2'	-6.31	105.06	112.00
20	AL	33	ARG	NE-CZ-NH1	6.31	123.45	120.30
34	BA	138	C	C5'-C4'-O4'	6.31	116.67	109.10
34	BA	519	G	C8-N9-C1'	-6.31	118.80	127.00
34	BA	1078	U	C6-N1-C2	-6.31	117.22	121.00
35	BB	373	C	C6-N1-C2	-6.31	117.78	120.30
35	BB	621	C	C1'-O4'-C4'	-6.31	104.85	109.90
35	BB	701	U	C3'-C2'-C1'	-6.31	96.45	101.50
36	BC	151	G	C4-N9-C1'	-6.31	118.30	126.50
50	BQ	211	ARG	NE-CZ-NH1	6.31	123.45	120.30
82	Bw	164	TYR	CA-CB-CG	-6.31	101.41	113.40
85	AA	1303	U	C2-N1-C1'	6.31	125.27	117.70
85	AA	1676	G	N9-C4-C5	6.31	107.92	105.40
34	BA	25	C	C2-N3-C4	-6.31	116.75	119.90
34	BA	233	U	C2-N1-C1'	-6.31	110.13	117.70
34	BA	806	U	C2-N3-C4	-6.31	123.22	127.00
35	BB	381	C	C5'-C4'-C3'	6.31	126.09	116.00
35	BB	453	C	C2-N3-C4	-6.31	116.75	119.90
35	BB	539	G	C5-C6-O6	-6.31	124.82	128.60
35	BB	986	C	P-O3'-C3'	6.31	127.27	119.70
35	BB	1222	A	C3'-C2'-C1'	-6.31	96.45	101.50
35	BB	1281	G	P-O3'-C3'	-6.31	112.13	119.70
41	BH	22	A	C8-N9-C4	6.31	108.32	105.80
59	BZ	67	VAL	CB-CA-C	-6.31	99.42	111.40
85	AA	717	G	N3-C2-N2	6.31	124.32	119.90
85	AA	1372	C	P-O5'-C5'	6.31	130.99	120.90
85	AA	1573	A	C2-N3-C4	-6.31	107.45	110.60
85	AA	1731	G	P-O3'-C3'	-6.31	112.13	119.70
85	AA	2050	C	P-O3'-C3'	6.31	127.27	119.70
34	BA	120	A	P-O3'-C3'	6.31	127.27	119.70
34	BA	159	U	N1-C1'-C2'	-6.31	105.06	112.00
34	BA	288	U	P-O3'-C3'	-6.31	112.13	119.70
35	BB	1406	C	O4'-C1'-N1	6.31	113.25	108.20
36	BC	116	C	N1-C2-O2	6.31	122.68	118.90
45	BL	124	ASP	CA-CB-CG	-6.31	99.53	113.40
85	AA	443	A	O4'-C1'-C2'	6.31	113.28	107.60
85	AA	639	C	P-O5'-C5'	-6.31	110.81	120.90
85	AA	805	A	P-O5'-C5'	6.31	130.99	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	835	C	C3'-C2'-C1'	-6.31	96.45	101.50
85	AA	977	U	C3'-C2'-C1'	-6.31	96.45	101.50
85	AA	1818	C	O4'-C1'-N1	6.31	113.25	108.20
34	BA	84	U	C6-N1-C2	-6.30	117.22	121.00
34	BA	561	U	N3-C4-O4	6.30	123.81	119.40
34	BA	1655	G	C5-C6-N1	6.30	114.65	111.50
38	BE	32	U	O4'-C4'-C3'	6.30	111.14	106.10
85	AA	2165	C	O4'-C1'-N1	6.30	113.24	108.20
85	AA	2189	U	C5'-C4'-O4'	6.30	116.67	109.10
35	BB	1444	U	C5'-C4'-C3'	-6.30	105.92	116.00
85	AA	572	G	P-O5'-C5'	-6.30	110.81	120.90
21	AM	125	ARG	CD-NE-CZ	-6.30	114.78	123.60
34	BA	718	U	N1-C1'-C2'	-6.30	105.07	112.00
34	BA	1289	C	C4'-C3'-C2'	-6.30	96.30	102.60
35	BB	1231	U	O3'-P-O5'	6.30	115.97	104.00
35	BB	1523	U	C5'-C4'-C3'	-6.30	105.92	116.00
37	BD	80	G	C8-N9-C4	6.30	108.92	106.40
38	BE	192	A	N1-C6-N6	-6.30	114.82	118.60
85	AA	979	U	C2-N3-C4	-6.30	123.22	127.00
85	AA	1409	U	O4'-C1'-N1	6.30	113.24	108.20
85	AA	2133	A	C8-N9-C4	6.30	108.32	105.80
34	BA	321	G	C8-N9-C1'	6.30	135.19	127.00
34	BA	590	U	C2-N1-C1'	-6.30	110.14	117.70
34	BA	992	A	C4'-C3'-C2'	6.30	108.90	102.60
34	BA	1308	C	O5'-P-OP2	-6.30	100.03	105.70
85	AA	44	C	C2'-C3'-O3'	6.30	123.78	113.70
85	AA	272	C	C3'-C2'-C1'	-6.30	96.46	101.50
85	AA	478	U	C5-C6-N1	-6.30	119.55	122.70
85	AA	713	G	P-O3'-C3'	-6.30	112.14	119.70
85	AA	983	A	O3'-P-O5'	-6.30	92.03	104.00
30	AW	5	ASP	N-CA-C	6.30	128.00	111.00
34	BA	180	G	C8-N9-C4	-6.30	103.88	106.40
35	BB	72	G	C1'-O4'-C4'	-6.30	104.86	109.90
35	BB	857	G	C5'-C4'-C3'	-6.30	105.92	116.00
35	BB	1396	G	C8-N9-C4	6.30	108.92	106.40
38	BE	114	G	C5'-C4'-O4'	6.30	116.66	109.10
34	BA	299	C	O4'-C1'-N1	6.30	113.24	108.20
34	BA	1099	U	P-O3'-C3'	-6.30	112.14	119.70
34	BA	1509	U	N3-C2-O2	-6.30	117.79	122.20
34	BA	1696	G	O3'-P-O5'	6.30	115.96	104.00
34	BA	1708	A	C8-N9-C4	6.30	108.32	105.80
35	BB	140	U	O4'-C1'-N1	6.30	113.24	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	449	C	N3-C4-N4	-6.30	113.59	118.00
35	BB	1109	A	P-O3'-C3'	-6.30	112.14	119.70
38	BE	38	C	C5'-C4'-C3'	-6.30	105.93	116.00
40	BG	45	G	C1'-O4'-C4'	-6.30	104.86	109.90
84	By	24	VAL	CB-CA-C	-6.30	99.44	111.40
85	AA	869	A	C5-N7-C8	-6.30	100.75	103.90
85	AA	1459	C	N1-C2-N3	6.30	123.61	119.20
85	AA	1716	U	N3-C2-O2	-6.30	117.79	122.20
85	AA	1816	C	N1-C2-O2	6.30	122.68	118.90
85	AA	1917	G	O4'-C1'-N9	6.30	113.24	108.20
85	AA	1955	U	N3-C2-O2	-6.30	117.79	122.20
34	BA	1173	C	P-O5'-C5'	6.29	130.97	120.90
34	BA	1446	G	N1-C6-O6	-6.29	116.12	119.90
35	BB	807	U	C6-N1-C1'	6.29	130.01	121.20
38	BE	77	C	C2-N1-C1'	6.29	125.72	118.80
41	BH	39	G	C2-N3-C4	6.29	115.05	111.90
85	AA	204	U	O4'-C1'-N1	6.29	113.24	108.20
85	AA	785	C	C4'-C3'-C2'	6.29	108.89	102.60
85	AA	887	A	O4'-C1'-N9	6.29	113.24	108.20
6	A5	180	ARG	NE-CZ-NH1	6.29	123.45	120.30
34	BA	265	A	O5'-C5'-C4'	-6.29	99.74	111.70
34	BA	691	A	O3'-P-O5'	6.29	115.96	104.00
34	BA	1484	A	C5'-C4'-C3'	6.29	126.07	116.00
35	BB	642	G	C3'-C2'-C1'	-6.29	96.47	101.50
35	BB	1514	G	C1'-O4'-C4'	-6.29	104.87	109.90
40	BG	145	C	P-O5'-C5'	-6.29	110.83	120.90
85	AA	47	A	C5'-C4'-O4'	6.29	116.65	109.10
85	AA	589	A	C8-N9-C4	-6.29	103.28	105.80
85	AA	1157	U	C2-N1-C1'	-6.29	110.15	117.70
34	BA	485	C	O3'-P-O5'	6.29	115.95	104.00
34	BA	1335	A	C3'-C2'-C1'	-6.29	96.47	101.50
34	BA	1561	C	C2-N3-C4	-6.29	116.75	119.90
34	BA	1694	C	C2-N1-C1'	-6.29	111.88	118.80
34	BA	1733	G	N9-C1'-C2'	-6.29	105.08	112.00
35	BB	24	C	C3'-C2'-C1'	-6.29	96.47	101.50
35	BB	314	A	C5-C6-N6	-6.29	118.67	123.70
35	BB	1014	U	C2-N1-C1'	-6.29	110.15	117.70
35	BB	1251	G	N9-C4-C5	-6.29	102.88	105.40
35	BB	1520	C	P-O3'-C3'	6.29	127.25	119.70
49	BP	102	PHE	CB-CG-CD2	-6.29	116.40	120.80
68	Bi	62	ARG	NE-CZ-NH2	-6.29	117.15	120.30
85	AA	414	C	N1-C2-N3	6.29	123.60	119.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	962	U	C1'-O4'-C4'	-6.29	104.87	109.90
85	AA	1191	G	O4'-C1'-N9	6.29	113.23	108.20
35	BB	870	C	N3-C4-C5	-6.29	119.38	121.90
85	AA	47	A	C8-N9-C1'	-6.29	116.38	127.70
85	AA	718	C	N3-C2-O2	-6.29	117.50	121.90
34	BA	195	G	N1-C6-O6	-6.29	116.13	119.90
34	BA	1367	G	C5'-C4'-O4'	6.29	116.65	109.10
34	BA	1499	A	C4-N9-C1'	-6.29	114.98	126.30
34	BA	1550	G	P-O3'-C3'	6.29	127.25	119.70
34	BA	1751	C	O4'-C1'-N1	6.29	113.23	108.20
35	BB	130	G	C4-N9-C1'	-6.29	118.33	126.50
35	BB	791	A	C8-N9-C4	-6.29	103.28	105.80
35	BB	1347	C	C5'-C4'-C3'	-6.29	105.94	116.00
37	BD	45	U	C2-N3-C4	6.29	130.77	127.00
37	BD	71	G	C4'-C3'-C2'	-6.29	96.31	102.60
38	BE	20	C	OP1-P-OP2	-6.29	110.17	119.60
85	AA	112	A	C8-N9-C4	6.29	108.31	105.80
85	AA	811	A	P-O5'-C5'	6.29	130.96	120.90
85	AA	1506	U	N1-C2-N3	6.29	118.67	114.90
85	AA	1691	U	O4'-C1'-N1	6.29	113.23	108.20
85	AA	2147	A	P-O3'-C3'	-6.29	112.15	119.70
85	AA	2155	U	C2'-C3'-O3'	6.29	123.76	113.70
85	AA	2196	G	O5'-C5'-C4'	-6.29	99.75	111.70
34	BA	68	A	C2-N3-C4	-6.29	107.46	110.60
34	BA	290	G	C1'-O4'-C4'	-6.29	104.87	109.90
34	BA	909	G	C8-N9-C4	6.29	108.92	106.40
34	BA	1271	C	P-O3'-C3'	-6.29	112.16	119.70
34	BA	1519	G	C5'-C4'-C3'	-6.29	105.94	116.00
35	BB	426	A	P-O5'-C5'	6.29	130.96	120.90
35	BB	431	U	C5'-C4'-O4'	6.29	116.64	109.10
35	BB	804	U	C6-N1-C2	-6.29	117.23	121.00
74	Bo	48	ARG	NE-CZ-NH1	6.29	123.44	120.30
85	AA	127	U	C5'-C4'-O4'	6.29	116.64	109.10
85	AA	1995	U	N3-C4-O4	-6.29	115.00	119.40
34	BA	602	G	O4'-C1'-N9	6.29	113.23	108.20
34	BA	673	U	C5'-C4'-O4'	-6.29	101.56	109.10
34	BA	1058	C	C2-N3-C4	-6.29	116.76	119.90
34	BA	1491	U	C5-C4-O4	6.29	129.67	125.90
34	BA	1782	C	N3-C2-O2	-6.29	117.50	121.90
35	BB	440	U	N1-C2-N3	6.29	118.67	114.90
35	BB	627	G	C3'-C2'-C1'	6.29	106.53	101.50
35	BB	743	C	C2-N3-C4	-6.29	116.76	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1272	G	C5-C6-O6	-6.29	124.83	128.60
35	BB	1438	U	C4-C5-C6	-6.29	115.93	119.70
53	BT	136	ARG	N-CA-CB	-6.29	99.29	110.60
58	BY	61	ARG	NE-CZ-NH1	6.29	123.44	120.30
67	Bh	91	HIS	CA-CB-CG	-6.29	102.92	113.60
85	AA	796	U	P-O3'-C3'	6.29	127.24	119.70
85	AA	1594	G	C5'-C4'-C3'	-6.29	105.94	116.00
85	AA	2047	U	C5'-C4'-C3'	-6.29	105.94	116.00
85	AA	2177	C	O4'-C1'-N1	6.29	113.23	108.20
85	AA	2187	G	C8-N9-C4	6.29	108.91	106.40
35	BB	635	A	O3'-P-O5'	6.28	115.94	104.00
35	BB	1098	G	N1-C6-O6	-6.28	116.13	119.90
36	BC	27	U	C6-N1-C2	-6.28	117.23	121.00
38	BE	26	G	C3'-C2'-C1'	-6.28	96.47	101.50
40	BG	11	G	C8-N9-C4	-6.28	103.89	106.40
41	BH	9	C	C6-N1-C1'	-6.28	113.26	120.80
42	BI	71	MET	N-CA-CB	-6.28	99.29	110.60
59	BZ	14	ARG	NE-CZ-NH1	6.28	123.44	120.30
85	AA	1190	G	C8-N9-C1'	6.28	135.17	127.00
85	AA	1900	C	C2-N1-C1'	-6.28	111.89	118.80
85	AA	1978	G	C5-C6-N1	6.28	114.64	111.50
85	AA	2069	A	C5'-C4'-O4'	6.28	116.64	109.10
85	AA	2075	C	C1'-O4'-C4'	-6.28	104.87	109.90
34	BA	255	G	C8-N9-C1'	6.28	135.17	127.00
34	BA	321	G	C1'-O4'-C4'	-6.28	104.88	109.90
34	BA	1177	C	N3-C4-N4	6.28	122.40	118.00
34	BA	1222	C	N3-C4-N4	6.28	122.40	118.00
34	BA	1721	U	C5-C6-N1	-6.28	119.56	122.70
35	BB	1293	C	N3-C2-O2	-6.28	117.50	121.90
66	Bg	20	THR	N-CA-CB	6.28	122.23	110.30
85	AA	145	C	P-O3'-C3'	-6.28	112.16	119.70
85	AA	1675	U	P-O3'-C3'	-6.28	112.16	119.70
34	BA	267	G	C5-C6-O6	-6.28	124.83	128.60
34	BA	1513	G	N1-C6-O6	-6.28	116.13	119.90
35	BB	465	C	O4'-C1'-N1	6.28	113.22	108.20
35	BB	679	G	N1-C2-N2	-6.28	110.55	116.20
35	BB	1199	A	N1-C2-N3	-6.28	126.16	129.30
35	BB	1257	A	C5-C6-N6	-6.28	118.67	123.70
35	BB	1454	G	C1'-O4'-C4'	-6.28	104.88	109.90
36	BC	16	A	C1'-O4'-C4'	-6.28	104.88	109.90
85	AA	889	G	P-O5'-C5'	6.28	130.95	120.90
85	AA	1033	C	O4'-C1'-N1	6.28	113.22	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1995	U	O4'-C1'-N1	6.28	113.22	108.20
34	BA	277	A	N1-C6-N6	-6.28	114.83	118.60
34	BA	729	C	N1-C2-O2	6.28	122.67	118.90
34	BA	816	G	C4'-C3'-C2'	6.28	108.88	102.60
34	BA	941	G	N9-C4-C5	-6.28	102.89	105.40
34	BA	951	C	P-O3'-C3'	-6.28	112.17	119.70
34	BA	1157	A	N9-C1'-C2'	-6.28	105.09	112.00
34	BA	1475	G	N9-C1'-C2'	-6.28	105.09	112.00
34	BA	1511	C	C4'-C3'-C2'	6.28	108.88	102.60
36	BC	17	U	N1-C2-N3	-6.28	111.13	114.90
53	BT	3	SER	N-CA-CB	6.28	119.92	110.50
64	Be	204	ARG	NE-CZ-NH1	6.28	123.44	120.30
85	AA	472	A	C8-N9-C4	-6.28	103.29	105.80
85	AA	1150	G	C5-C6-O6	-6.28	124.83	128.60
85	AA	1348	C	C6-N1-C2	-6.28	117.79	120.30
13	AE	71	ASP	CB-CG-OD2	-6.28	112.65	118.30
34	BA	106	U	C3'-C2'-C1'	-6.28	96.48	101.50
34	BA	433	G	O5'-C5'-C4'	6.28	123.63	111.70
34	BA	1014	A	C5-C6-N6	6.28	128.72	123.70
34	BA	1725	U	P-O5'-C5'	-6.28	110.86	120.90
34	BA	1758	C	O4'-C1'-N1	6.28	113.22	108.20
34	BA	1793	G	N3-C2-N2	-6.28	115.51	119.90
35	BB	27	C	O4'-C1'-N1	6.28	113.22	108.20
35	BB	112	G	N3-C4-C5	-6.28	125.46	128.60
35	BB	524	C	C5'-C4'-O4'	6.28	116.63	109.10
35	BB	693	U	C1'-O4'-C4'	-6.28	104.88	109.90
35	BB	746	A	C1'-O4'-C4'	-6.28	104.88	109.90
35	BB	798	A	OP1-P-OP2	-6.28	110.18	119.60
35	BB	1527	A	C5'-C4'-C3'	-6.28	105.96	116.00
36	BC	7	U	P-O5'-C5'	-6.28	110.86	120.90
37	BD	16	U	C2-N1-C1'	-6.28	110.17	117.70
38	BE	120	C	O4'-C1'-N1	6.28	113.22	108.20
41	BH	49	C	C3'-C2'-C1'	-6.28	96.48	101.50
65	Bf	193	GLY	N-CA-C	-6.28	97.41	113.10
75	Bp	17	ARG	NE-CZ-NH1	6.28	123.44	120.30
85	AA	158	C	C5'-C4'-C3'	-6.28	105.95	116.00
85	AA	1492	U	N1-C1'-C2'	-6.28	105.09	112.00
34	BA	210	G	N1-C6-O6	-6.28	116.14	119.90
34	BA	626	G	N3-C4-C5	-6.28	125.46	128.60
34	BA	649	A	P-O3'-C3'	-6.28	112.17	119.70
34	BA	1697	U	N3-C2-O2	6.28	126.59	122.20
35	BB	1517	G	P-O3'-C3'	-6.28	112.17	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	89	U	O5'-C5'-C4'	-6.28	99.78	111.70
36	BC	140	U	C6-N1-C1'	-6.28	112.42	121.20
40	BG	166	C	O4'-C1'-N1	6.28	113.22	108.20
77	Br	225	LEU	N-CA-C	-6.28	94.05	111.00
85	AA	548	G	C1'-O4'-C4'	-6.28	104.88	109.90
85	AA	1160	U	P-O3'-C3'	6.28	127.23	119.70
34	BA	575	U	OP2-P-O3'	6.27	119.00	105.20
34	BA	682	A	P-O5'-C5'	-6.27	110.86	120.90
34	BA	1202	G	C5'-C4'-O4'	6.27	116.63	109.10
34	BA	1333	G	N9-C1'-C2'	-6.27	105.10	112.00
34	BA	1347	G	C8-N9-C1'	6.27	135.16	127.00
35	BB	894	A	O3'-P-O5'	6.27	115.92	104.00
35	BB	1548	C	C6-N1-C1'	6.27	128.33	120.80
36	BC	139	A	C5'-C4'-O4'	-6.27	101.57	109.10
37	BD	45	U	C3'-C2'-C1'	-6.27	96.48	101.50
85	AA	962	U	N1-C1'-C2'	-6.27	105.10	112.00
34	BA	477	C	O4'-C1'-N1	6.27	113.22	108.20
34	BA	810	A	O4'-C1'-N9	6.27	113.22	108.20
34	BA	1425	G	O4'-C1'-C2'	-6.27	99.53	105.80
34	BA	1660	A	P-O3'-C3'	-6.27	112.17	119.70
35	BB	448	G	C6-N1-C2	-6.27	121.34	125.10
35	BB	479	U	C6-N1-C1'	6.27	129.98	121.20
35	BB	1228	A	O5'-C5'-C4'	6.27	123.62	111.70
35	BB	1475	U	OP1-P-OP2	-6.27	110.19	119.60
36	BC	74	U	O4'-C1'-N1	6.27	113.22	108.20
37	BD	48	G	C2-N3-C4	-6.27	108.76	111.90
37	BD	92	G	C6-N1-C2	-6.27	121.34	125.10
62	Bc	21	LYS	N-CA-CB	-6.27	99.31	110.60
77	Br	195	MET	CA-CB-CG	6.27	123.97	113.30
85	AA	314	C	C5'-C4'-C3'	6.27	126.04	116.00
34	BA	869	C	N3-C2-O2	-6.27	117.51	121.90
35	BB	879	G	OP1-P-OP2	-6.27	110.19	119.60
36	BC	44	A	C6-N1-C2	-6.27	114.84	118.60
85	AA	744	C	C2-N1-C1'	6.27	125.70	118.80
85	AA	1916	A	N3-C4-C5	6.27	131.19	126.80
85	AA	2067	A	P-O5'-C5'	-6.27	110.87	120.90
15	AG	32	ASP	CA-CB-CG	-6.27	99.61	113.40
34	BA	128	C	C6-N1-C2	-6.27	117.79	120.30
34	BA	683	C	C5'-C4'-O4'	-6.27	101.58	109.10
34	BA	1215	U	O5'-C5'-C4'	-6.27	99.79	111.70
34	BA	1262	A	N7-C8-N9	-6.27	110.67	113.80
34	BA	1447	C	C1'-O4'-C4'	-6.27	104.88	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1674	G	C4-N9-C1'	6.27	134.65	126.50
35	BB	59	U	O5'-C5'-C4'	-6.27	99.79	111.70
35	BB	1237	C	O4'-C1'-C2'	6.27	113.24	107.60
35	BB	1520	C	C4'-C3'-C2'	-6.27	96.33	102.60
38	BE	59	U	P-O3'-C3'	-6.27	112.18	119.70
62	Bc	78	LYS	CG-CD-CE	6.27	130.71	111.90
85	AA	365	G	N7-C8-N9	-6.27	109.97	113.10
85	AA	1503	G	O4'-C1'-C2'	6.27	113.24	107.60
85	AA	1609	U	C4'-C3'-C2'	-6.27	96.33	102.60
85	AA	2228	G	C6-N1-C2	-6.27	121.34	125.10
86	AB	41	C	O4'-C1'-N1	6.27	113.22	108.20
34	BA	761	U	N3-C2-O2	-6.27	117.81	122.20
34	BA	974	G	P-O3'-C3'	-6.27	112.18	119.70
34	BA	1344	G	C5-C6-O6	-6.27	124.84	128.60
35	BB	662	G	N3-C2-N2	6.27	124.29	119.90
35	BB	1299	G	C5-N7-C8	-6.27	101.17	104.30
35	BB	1387	C	N3-C4-N4	-6.27	113.61	118.00
38	BE	140	G	C5-C6-N1	6.27	114.63	111.50
39	BF	11	C	C5'-C4'-O4'	6.27	116.62	109.10
39	BF	53	G	C3'-C2'-C1'	-6.27	96.49	101.50
40	BG	94	G	C5'-C4'-C3'	6.27	126.03	116.00
34	BA	335	C	O5'-C5'-C4'	-6.27	99.80	111.70
34	BA	658	C	C2-N1-C1'	-6.27	111.91	118.80
34	BA	984	U	C5-C6-N1	-6.27	119.57	122.70
34	BA	1011	G	C8-N9-C1'	6.27	135.15	127.00
34	BA	1172	C	C6-N1-C1'	6.27	128.32	120.80
85	AA	2042	G	C8-N9-C4	6.27	108.91	106.40
20	AL	108	MET	CA-CB-CG	6.26	123.95	113.30
34	BA	405	C	C2-N1-C1'	-6.26	111.91	118.80
34	BA	784	C	N1-C2-O2	6.26	122.66	118.90
34	BA	867	C	C2-N3-C4	-6.26	116.77	119.90
34	BA	1206	C	C6-N1-C1'	-6.26	113.28	120.80
34	BA	1286	C	P-O3'-C3'	-6.26	112.18	119.70
35	BB	710	A	P-O3'-C3'	-6.26	112.18	119.70
38	BE	144	A	N1-C6-N6	-6.26	114.84	118.60
75	Bp	69	VAL	CA-CB-CG2	-6.26	101.50	110.90
85	AA	320	U	P-O3'-C3'	-6.26	112.18	119.70
85	AA	340	G	C1'-O4'-C4'	-6.26	104.89	109.90
85	AA	764	U	O3'-P-O5'	6.26	115.90	104.00
85	AA	858	G	C1'-O4'-C4'	-6.26	104.89	109.90
85	AA	1727	U	C2-N1-C1'	-6.26	110.18	117.70
34	BA	768	G	OP2-P-O3'	6.26	118.98	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1106	A	P-O5'-C5'	6.26	130.92	120.90
34	BA	1505	G	C8-N9-C1'	-6.26	118.86	127.00
34	BA	1622	U	P-O3'-C3'	-6.26	112.19	119.70
35	BB	313	C	C6-N1-C2	-6.26	117.80	120.30
34	BA	305	C	N1-C1'-C2'	-6.26	105.11	112.00
34	BA	371	U	C5-C6-N1	-6.26	119.57	122.70
34	BA	589	A	C3'-C2'-C1'	-6.26	96.49	101.50
34	BA	739	A	O3'-P-O5'	6.26	115.90	104.00
34	BA	767	U	O4'-C1'-N1	6.26	113.21	108.20
34	BA	922	C	C3'-C2'-C1'	-6.26	96.49	101.50
34	BA	922	C	P-O5'-C5'	-6.26	110.88	120.90
34	BA	1218	G	N3-C4-C5	-6.26	125.47	128.60
34	BA	1332	U	N1-C1'-C2'	-6.26	105.11	112.00
35	BB	843	G	N9-C1'-C2'	-6.26	105.11	112.00
35	BB	1187	G	O4'-C1'-N9	6.26	113.21	108.20
35	BB	1429	A	P-O3'-C3'	-6.26	112.19	119.70
35	BB	1461	C	N3-C4-N4	6.26	122.38	118.00
35	BB	1521	G	C6-N1-C2	-6.26	121.34	125.10
35	BB	1538	G	N9-C1'-C2'	-6.26	105.11	112.00
36	BC	19	A	C3'-C2'-C1'	6.26	106.51	101.50
36	BC	72	A	C5'-C4'-C3'	-6.26	105.98	116.00
36	BC	111	C	N3-C4-N4	6.26	122.38	118.00
38	BE	201	A	C5'-C4'-O4'	6.26	116.61	109.10
85	AA	1761	C	O4'-C1'-N1	6.26	113.21	108.20
34	BA	184	C	C1'-O4'-C4'	-6.26	104.89	109.90
34	BA	201	A	N1-C2-N3	-6.26	126.17	129.30
34	BA	572	G	C5'-C4'-C3'	6.26	126.02	116.00
34	BA	721	A	C5'-C4'-C3'	-6.26	105.98	116.00
34	BA	1135	U	C2-N1-C1'	-6.26	110.19	117.70
34	BA	1234	U	C2-N1-C1'	-6.26	110.19	117.70
35	BB	657	A	O4'-C1'-N9	6.26	113.21	108.20
35	BB	802	G	C5-C6-N1	6.26	114.63	111.50
41	BH	23	G	C5-N7-C8	-6.26	101.17	104.30
67	Bh	33	ARG	NE-CZ-NH1	6.26	123.43	120.30
70	Bk	80	ARG	NE-CZ-NH1	6.26	123.43	120.30
85	AA	160	A	C5'-C4'-C3'	-6.26	105.98	116.00
85	AA	164	G	C3'-C2'-C1'	-6.26	96.49	101.50
85	AA	335	G	C8-N9-C1'	6.26	135.14	127.00
85	AA	932	A	C5-N7-C8	-6.26	100.77	103.90
85	AA	1848	G	C2-N3-C4	-6.26	108.77	111.90
17	AI	60	ALA	CA-C-N	6.26	134.62	117.10
35	BB	368	C	C4'-C3'-C2'	6.26	108.86	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1424	G	C8-N9-C1'	6.26	135.14	127.00
37	BD	64	A	O5'-C5'-C4'	-6.26	99.81	111.70
40	BG	43	U	O4'-C1'-N1	6.26	113.21	108.20
41	BH	65	G	O4'-C1'-N9	6.26	113.21	108.20
85	AA	216	U	O4'-C1'-N1	6.26	113.21	108.20
85	AA	2092	A	P-O3'-C3'	-6.26	112.19	119.70
7	A6	145	PHE	CB-CG-CD2	-6.26	116.42	120.80
34	BA	13	U	C6-N1-C1'	-6.26	112.44	121.20
34	BA	78	U	C2-N1-C1'	-6.26	110.19	117.70
34	BA	1066	A	P-O3'-C3'	-6.26	112.19	119.70
34	BA	1079	C	C4-C5-C6	-6.26	114.27	117.40
34	BA	1708	A	O4'-C4'-C3'	-6.26	97.74	104.00
41	BH	62	C	C5'-C4'-O4'	6.26	116.61	109.10
85	AA	706	U	O5'-C5'-C4'	-6.26	99.81	111.70
85	AA	773	G	C4-N9-C1'	-6.26	118.37	126.50
85	AA	857	G	C1'-O4'-C4'	-6.26	104.89	109.90
85	AA	2176	U	O5'-C5'-C4'	-6.26	99.81	111.70
34	BA	511	U	P-O3'-C3'	6.25	127.21	119.70
34	BA	546	U	C5'-C4'-O4'	6.25	116.61	109.10
34	BA	940	C	N3-C2-O2	-6.25	117.52	121.90
34	BA	1146	U	P-O3'-C3'	6.25	127.21	119.70
37	BD	12	U	C3'-C2'-C1'	-6.25	96.50	101.50
40	BG	137	G	C5'-C4'-C3'	-6.25	105.99	116.00
85	AA	239	G	P-O5'-C5'	-6.25	110.89	120.90
85	AA	267	U	O4'-C4'-C3'	-6.25	97.75	104.00
85	AA	1254	A	O4'-C1'-N9	6.25	113.20	108.20
30	AW	24	ARG	NE-CZ-NH1	6.25	123.43	120.30
32	AY	47	LYS	N-CA-CB	-6.25	99.34	110.60
34	BA	7	U	C5'-C4'-C3'	6.25	126.01	116.00
34	BA	47	U	C4'-C3'-C2'	6.25	108.85	102.60
34	BA	756	A	P-O3'-C3'	-6.25	112.20	119.70
34	BA	859	G	C5-C6-O6	-6.25	124.85	128.60
34	BA	1084	A	O4'-C1'-N9	6.25	113.20	108.20
35	BB	464	C	O4'-C1'-N1	6.25	113.20	108.20
35	BB	1495	U	C6-N1-C1'	6.25	129.95	121.20
37	BD	98	G	N3-C2-N2	6.25	124.28	119.90
38	BE	60	C	C6-N1-C2	-6.25	117.80	120.30
73	Bn	14	ARG	C-N-CA	6.25	137.34	121.70
85	AA	1223	A	P-O3'-C3'	-6.25	112.20	119.70
85	AA	1788	U	P-O5'-C5'	6.25	130.91	120.90
85	AA	1966	C	O4'-C1'-N1	6.25	113.20	108.20
85	AA	1985	C	P-O3'-C3'	-6.25	112.20	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A2	64	PHE	CA-CB-CG	-6.25	98.90	113.90
4	A3	7	TYR	N-CA-CB	-6.25	99.35	110.60
27	AT	101	LEU	C-N-CA	6.25	135.43	122.30
34	BA	296	G	C8-N9-C4	-6.25	103.90	106.40
34	BA	730	C	C3'-C2'-C1'	-6.25	96.50	101.50
34	BA	1173	C	C2-N1-C1'	-6.25	111.92	118.80
34	BA	1435	A	C8-N9-C4	6.25	108.30	105.80
35	BB	356	C	P-O3'-C3'	-6.25	112.20	119.70
35	BB	405	U	C5-C4-O4	-6.25	122.15	125.90
35	BB	494	C	N1-C1'-C2'	-6.25	105.12	112.00
35	BB	1170	U	P-O5'-C5'	-6.25	110.90	120.90
35	BB	1307	C	O4'-C1'-N1	6.25	113.20	108.20
35	BB	1548	C	N1-C1'-C2'	-6.25	105.12	112.00
36	BC	136	G	N3-C2-N2	6.25	124.28	119.90
37	BD	51	G	C2'-C3'-O3'	6.25	123.70	113.70
64	Be	83	PHE	CB-CA-C	6.25	122.90	110.40
85	AA	25	C	C6-N1-C2	-6.25	117.80	120.30
85	AA	70	U	C5'-C4'-O4'	6.25	116.60	109.10
85	AA	1453	U	P-O3'-C3'	6.25	127.20	119.70
85	AA	2054	G	O4'-C1'-N9	6.25	113.20	108.20
34	BA	572	G	N9-C1'-C2'	6.25	122.12	114.00
34	BA	1000	G	C4-N9-C1'	-6.25	118.38	126.50
35	BB	412	A	C1'-O4'-C4'	-6.25	104.90	109.90
35	BB	995	C	C4'-C3'-C2'	6.25	108.85	102.60
36	BC	158	U	P-O3'-C3'	-6.25	112.20	119.70
39	BF	51	C	C2-N3-C4	-6.25	116.78	119.90
40	BG	122	G	C6-N1-C2	-6.25	121.35	125.10
85	AA	48	G	C5'-C4'-C3'	-6.25	106.00	116.00
85	AA	112	A	O5'-C5'-C4'	6.25	123.58	111.70
85	AA	600	C	P-O3'-C3'	6.25	127.20	119.70
85	AA	1234	G	P-O5'-C5'	6.25	130.90	120.90
85	AA	1467	U	C2-N3-C4	-6.25	123.25	127.00
85	AA	2199	G	P-O3'-C3'	6.25	127.20	119.70
7	A6	52	ARG	NE-CZ-NH1	6.25	123.42	120.30
34	BA	450	G	N1-C6-O6	6.25	123.65	119.90
34	BA	861	C	N1-C1'-C2'	-6.25	105.13	112.00
34	BA	1247	G	C5'-C4'-O4'	6.25	116.60	109.10
34	BA	1609	U	C6-N1-C1'	6.25	129.95	121.20
34	BA	1807	G	C3'-C2'-C1'	-6.25	96.50	101.50
35	BB	398	A	P-O3'-C3'	6.25	127.20	119.70
35	BB	642	G	C5-C6-O6	6.25	132.35	128.60
35	BB	1208	G	C6-C5-N7	-6.25	126.65	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1209	A	C5-C6-N6	-6.25	118.70	123.70
35	BB	1517	G	C5-N7-C8	-6.25	101.18	104.30
37	BD	63	C	P-O5'-C5'	-6.25	110.90	120.90
37	BD	91	U	C5'-C4'-C3'	-6.25	106.00	116.00
38	BE	204	U	C5'-C4'-C3'	-6.25	106.00	116.00
85	AA	809	A	O5'-C5'-C4'	-6.25	99.83	111.70
85	AA	899	A	N9-C1'-C2'	-6.25	105.13	112.00
85	AA	1858	G	C8-N9-C1'	6.25	135.12	127.00
85	AA	1985	C	O4'-C1'-N1	6.25	113.20	108.20
85	AA	2171	A	O4'-C1'-N9	6.25	113.20	108.20
34	BA	764	G	OP1-P-OP2	-6.25	110.23	119.60
34	BA	1189	A	O4'-C1'-N9	6.25	113.20	108.20
34	BA	1702	G	C4-N9-C1'	-6.25	118.38	126.50
35	BB	505	G	O5'-C5'-C4'	-6.25	99.83	111.70
35	BB	667	G	O5'-C5'-C4'	-6.25	99.83	111.70
35	BB	1434	G	N3-C2-N2	6.25	124.27	119.90
35	BB	1518	U	C5'-C4'-C3'	6.25	126.00	116.00
39	BF	64	U	OP1-P-OP2	-6.25	110.23	119.60
49	BP	120	ARG	NE-CZ-NH1	6.25	123.42	120.30
85	AA	1661	U	O4'-C1'-N1	6.25	113.20	108.20
34	BA	604	G	OP1-P-OP2	-6.25	110.23	119.60
35	BB	512	C	O4'-C1'-N1	6.25	113.20	108.20
38	BE	202	C	C5'-C4'-C3'	6.25	125.99	116.00
39	BF	47	C	C5'-C4'-C3'	-6.25	106.01	116.00
40	BG	108	G	C5'-C4'-C3'	-6.25	106.01	116.00
41	BH	11	C	P-O3'-C3'	6.25	127.19	119.70
41	BH	12	U	C5-C6-N1	-6.25	119.58	122.70
41	BH	110	C	C5'-C4'-O4'	6.25	116.59	109.10
85	AA	90	A	C5'-C4'-O4'	6.25	116.59	109.10
85	AA	161	A	C5-C6-N1	-6.25	114.58	117.70
85	AA	889	G	N1-C2-N2	6.25	121.82	116.20
34	BA	69	C	C5-C4-N4	6.24	124.57	120.20
34	BA	480	G	N9-C4-C5	-6.24	102.90	105.40
34	BA	620	C	C6-N1-C2	-6.24	117.80	120.30
34	BA	1037	C	P-O5'-C5'	-6.24	110.91	120.90
34	BA	1253	G	O3'-P-O5'	-6.24	92.14	104.00
35	BB	318	C	O4'-C1'-N1	6.24	113.19	108.20
35	BB	667	G	C3'-C2'-C1'	-6.24	96.50	101.50
36	BC	122	A	C1'-O4'-C4'	-6.24	104.91	109.90
37	BD	93	G	N9-C1'-C2'	-6.24	105.13	112.00
41	BH	63	G	C5'-C4'-O4'	6.24	116.59	109.10
42	BI	58	ASN	CA-CB-CG	-6.24	99.66	113.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
49	BP	114	ARG	NE-CZ-NH1	6.24	123.42	120.30
71	Bl	42	GLY	N-CA-C	-6.24	97.49	113.10
85	AA	61	C	C2-N1-C1'	-6.24	111.93	118.80
85	AA	71	G	O3'-P-O5'	-6.24	92.14	104.00
85	AA	273	C	C3'-C2'-C1'	-6.24	96.50	101.50
85	AA	442	G	C6-N1-C2	-6.24	121.35	125.10
85	AA	689	U	C5'-C4'-C3'	-6.24	106.01	116.00
85	AA	2236	U	C2-N3-C4	-6.24	123.25	127.00
34	BA	1108	U	C5'-C4'-C3'	-6.24	106.01	116.00
34	BA	1226	G	C5-C6-O6	-6.24	124.86	128.60
34	BA	1748	G	N1-C6-O6	6.24	123.64	119.90
35	BB	764	C	C6-N1-C1'	6.24	128.29	120.80
35	BB	1396	G	C3'-C2'-C1'	-6.24	96.51	101.50
35	BB	1491	G	P-O5'-C5'	-6.24	110.91	120.90
36	BC	76	C	P-O3'-C3'	-6.24	112.21	119.70
85	AA	367	A	C5-C6-N6	6.24	128.69	123.70
85	AA	1138	U	P-O3'-C3'	-6.24	112.21	119.70
85	AA	1287	C	N1-C2-N3	6.24	123.57	119.20
85	AA	1456	A	N1-C6-N6	6.24	122.34	118.60
85	AA	1572	C	P-O3'-C3'	-6.24	112.21	119.70
34	BA	22	C	O4'-C1'-N1	6.24	113.19	108.20
34	BA	43	U	P-O3'-C3'	6.24	127.19	119.70
34	BA	116	G	O4'-C1'-N9	6.24	113.19	108.20
34	BA	428	C	O4'-C1'-N1	6.24	113.19	108.20
37	BD	74	A	P-O3'-C3'	-6.24	112.21	119.70
38	BE	12	A	P-O3'-C3'	-6.24	112.21	119.70
40	BG	40	G	C4'-C3'-C2'	-6.24	96.36	102.60
85	AA	553	G	N7-C8-N9	6.24	116.22	113.10
85	AA	682	C	N1-C1'-C2'	-6.24	105.14	112.00
85	AA	1515	A	O5'-C5'-C4'	-6.24	99.84	111.70
85	AA	1679	U	C2-N3-C4	-6.24	123.26	127.00
85	AA	1699	A	O5'-C5'-C4'	-6.24	99.84	111.70
85	AA	1926	A	P-O3'-C3'	6.24	127.19	119.70
8	A7	148	ARG	NE-CZ-NH1	6.24	123.42	120.30
34	BA	131	A	C2'-C3'-O3'	6.24	123.68	113.70
34	BA	218	G	C8-N9-C4	6.24	108.90	106.40
34	BA	807	U	C2-N1-C1'	-6.24	110.21	117.70
34	BA	856	G	N9-C1'-C2'	-6.24	105.14	112.00
34	BA	862	C	O3'-P-O5'	-6.24	92.14	104.00
34	BA	1488	C	P-O5'-C5'	-6.24	110.92	120.90
34	BA	1513	G	C5-C6-N1	6.24	114.62	111.50
34	BA	1572	G	P-O3'-C3'	6.24	127.19	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	5	U	C6-N1-C1'	-6.24	112.47	121.20
41	BH	55	C	C6-N1-C2	-6.24	117.81	120.30
41	BH	132	C	O4'-C1'-N1	6.24	113.19	108.20
85	AA	96	C	P-O5'-C5'	6.24	130.88	120.90
85	AA	497	G	C5'-C4'-C3'	-6.24	106.02	116.00
85	AA	1934	A	C5'-C4'-C3'	-6.24	106.02	116.00
34	BA	1259	C	C1'-O4'-C4'	-6.24	104.91	109.90
34	BA	1295	U	C2-N3-C4	-6.24	123.26	127.00
34	BA	1521	C	C1'-O4'-C4'	-6.24	104.91	109.90
35	BB	375	G	C1'-O4'-C4'	-6.24	104.91	109.90
35	BB	651	G	N7-C8-N9	6.24	116.22	113.10
54	BU	81	ARG	NE-CZ-NH1	6.24	123.42	120.30
85	AA	570	U	N3-C2-O2	-6.24	117.83	122.20
7	A6	108	LEU	N-CA-CB	-6.24	97.93	110.40
34	BA	967	C	C6-N1-C1'	6.24	128.28	120.80
34	BA	1716	A	O5'-C5'-C4'	6.24	123.55	111.70
35	BB	131	A	C4-N9-C1'	-6.24	115.08	126.30
35	BB	429	C	C2-N3-C4	-6.24	116.78	119.90
35	BB	604	C	O4'-C1'-N1	6.24	113.19	108.20
35	BB	888	U	C4'-C3'-C2'	-6.24	96.36	102.60
35	BB	1063	C	C2-N3-C4	-6.24	116.78	119.90
35	BB	1388	A	C1'-O4'-C4'	-6.24	104.91	109.90
38	BE	21	C	P-O3'-C3'	6.24	127.18	119.70
85	AA	424	A	C8-N9-C4	-6.24	103.31	105.80
85	AA	1368	G	C2-N3-C4	-6.24	108.78	111.90
34	BA	26	C	C5-C4-N4	-6.23	115.84	120.20
34	BA	824	C	C3'-C2'-C1'	-6.23	96.51	101.50
34	BA	957	A	C5'-C4'-C3'	-6.23	106.03	116.00
34	BA	1006	G	O5'-C5'-C4'	-6.23	99.86	111.70
34	BA	1549	U	N1-C2-O2	6.23	127.16	122.80
35	BB	427	U	C3'-C2'-C1'	-6.23	96.51	101.50
35	BB	1190	U	N3-C2-O2	-6.23	117.84	122.20
41	BH	23	G	N9-C1'-C2'	6.23	122.11	114.00
85	AA	147	G	O4'-C1'-N9	6.23	113.19	108.20
2	A1	32	PRO	N-CA-C	6.23	128.30	112.10
34	BA	8	G	N3-C2-N2	6.23	124.26	119.90
34	BA	759	A	C8-N9-C4	6.23	108.29	105.80
34	BA	1592	U	C1'-O4'-C4'	-6.23	104.91	109.90
35	BB	448	G	C8-N9-C1'	6.23	135.10	127.00
35	BB	1012	G	C5-C6-O6	-6.23	124.86	128.60
35	BB	1185	G	C4-N9-C1'	-6.23	118.40	126.50
38	BE	16	C	C5'-C4'-O4'	6.23	116.58	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	95	C	N3-C4-N4	6.23	122.36	118.00
45	BL	150	ARG	NE-CZ-NH1	6.23	123.42	120.30
49	BP	14	ARG	NE-CZ-NH1	6.23	123.42	120.30
85	AA	282	C	C5'-C4'-O4'	6.23	116.58	109.10
85	AA	1807	A	O4'-C1'-N9	6.23	113.19	108.20
85	AA	2047	U	C6-N1-C2	-6.23	117.26	121.00
34	BA	49	A	C4-N9-C1'	-6.23	115.09	126.30
34	BA	449	G	C5-C6-N1	6.23	114.61	111.50
34	BA	609	G	P-O3'-C3'	-6.23	112.22	119.70
34	BA	1049	G	O3'-P-O5'	6.23	115.84	104.00
34	BA	1137	U	C2'-C3'-O3'	6.23	123.67	113.70
34	BA	1144	A	C8-N9-C4	-6.23	103.31	105.80
34	BA	1152	A	C8-N9-C4	6.23	108.29	105.80
35	BB	433	C	C4'-C3'-C2'	6.23	108.83	102.60
35	BB	566	A	C4'-C3'-C2'	-6.23	96.37	102.60
35	BB	984	U	O4'-C1'-N1	6.23	113.18	108.20
35	BB	1031	G	C5-C6-O6	-6.23	124.86	128.60
35	BB	1195	A	C5'-C4'-C3'	-6.23	106.03	116.00
38	BE	141	A	C5-C6-N6	-6.23	118.72	123.70
41	BH	19	G	O4'-C1'-N9	6.23	113.18	108.20
54	BU	57	TYR	CB-CG-CD1	-6.23	117.26	121.00
85	AA	256	A	C8-N9-C4	6.23	108.29	105.80
85	AA	312	G	N3-C2-N2	6.23	124.26	119.90
85	AA	970	U	O5'-C5'-C4'	-6.23	99.86	111.70
85	AA	1181	U	P-O3'-C3'	-6.23	112.22	119.70
85	AA	1353	U	C5'-C4'-O4'	6.23	116.58	109.10
85	AA	1822	G	C1'-O4'-C4'	-6.23	104.92	109.90
85	AA	2143	U	C1'-O4'-C4'	-6.23	104.92	109.90
35	BB	1103	A	C3'-C2'-C1'	-6.23	96.52	101.50
37	BD	105	G	C1'-O4'-C4'	-6.23	104.92	109.90
52	BS	162	ARG	NE-CZ-NH1	6.23	123.41	120.30
85	AA	342	C	C1'-O4'-C4'	-6.23	104.92	109.90
14	AF	116	ARG	NE-CZ-NH1	6.23	123.41	120.30
31	AX	119	MET	N-CA-CB	-6.23	99.39	110.60
34	BA	988	U	C5'-C4'-C3'	-6.23	106.03	116.00
34	BA	1502	G	O4'-C4'-C3'	-6.23	97.77	104.00
34	BA	1791	C	O4'-C1'-N1	6.23	113.18	108.20
35	BB	36	U	C4'-C3'-C2'	-6.23	96.37	102.60
35	BB	90	G	N7-C8-N9	-6.23	109.99	113.10
35	BB	384	A	OP1-P-OP2	-6.23	110.26	119.60
35	BB	802	G	N1-C2-N2	-6.23	110.59	116.20
35	BB	837	A	C4-N9-C1'	-6.23	115.09	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1187	G	C4-C5-N7	-6.23	108.31	110.80
36	BC	9	G	O3'-P-O5'	-6.23	92.17	104.00
38	BE	59	U	C5-C6-N1	-6.23	119.59	122.70
38	BE	140	G	P-O5'-C5'	-6.23	110.94	120.90
39	BF	70	A	C3'-C2'-C1'	-6.23	96.52	101.50
40	BG	19	C	N3-C2-O2	-6.23	117.54	121.90
41	BH	67	G	N1-C6-O6	-6.23	116.16	119.90
77	Br	226	ASP	CA-CB-CG	-6.23	99.70	113.40
85	AA	843	U	O4'-C1'-N1	6.23	113.18	108.20
85	AA	975	G	N3-C4-C5	-6.23	125.49	128.60
85	AA	1315	C	O4'-C1'-N1	6.23	113.18	108.20
34	BA	516	U	N1-C2-O2	6.23	127.16	122.80
34	BA	610	A	P-O5'-C5'	-6.23	110.94	120.90
34	BA	861	C	N3-C2-O2	-6.23	117.54	121.90
34	BA	932	G	N9-C1'-C2'	-6.23	105.15	112.00
34	BA	1048	C	N1-C1'-C2'	-6.23	105.15	112.00
34	BA	1724	G	N3-C4-N9	-6.23	122.27	126.00
35	BB	582	G	N9-C1'-C2'	-6.23	105.15	112.00
35	BB	839	G	O5'-C5'-C4'	6.23	123.53	111.70
85	AA	389	A	N9-C1'-C2'	-6.23	105.15	112.00
85	AA	1268	C	N3-C2-O2	-6.23	117.54	121.90
85	AA	1848	G	C5-C6-O6	-6.23	124.86	128.60
34	BA	795	G	C3'-C2'-C1'	-6.22	96.52	101.50
34	BA	1847	G	P-O5'-C5'	-6.22	110.94	120.90
35	BB	881	G	C5-C6-O6	-6.22	124.86	128.60
37	BD	50	A	N9-C1'-C2'	-6.22	105.15	112.00
39	BF	4	A	O5'-P-OP1	-6.22	100.10	105.70
40	BG	46	G	C6-N1-C2	-6.22	121.37	125.10
40	BG	89	A	P-O3'-C3'	-6.22	112.23	119.70
41	BH	17	A	C5'-C4'-C3'	-6.22	106.04	116.00
78	Bs	30	ARG	N-CA-CB	-6.22	99.39	110.60
85	AA	422	G	O5'-C5'-C4'	-6.22	99.87	111.70
85	AA	1184	A	N1-C2-N3	6.22	132.41	129.30
85	AA	2032	G	O4'-C1'-N9	6.22	113.18	108.20
85	AA	2055	G	N1-C6-O6	6.22	123.64	119.90
4	A3	60	GLU	N-CA-CB	6.22	121.80	110.60
34	BA	125	G	C6-N1-C2	-6.22	121.37	125.10
34	BA	389	U	N3-C2-O2	-6.22	117.84	122.20
34	BA	417	A	O3'-P-O5'	-6.22	92.18	104.00
34	BA	484	A	C4-N9-C1'	-6.22	115.10	126.30
34	BA	1211	G	C3'-C2'-C1'	6.22	106.48	101.50
34	BA	1555	G	O4'-C1'-N9	6.22	113.18	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1566	G	C8-N9-C1'	6.22	135.09	127.00
35	BB	440	U	N3-C2-O2	-6.22	117.84	122.20
35	BB	1252	G	C2-N3-C4	-6.22	108.79	111.90
35	BB	1415	G	N1-C6-O6	-6.22	116.17	119.90
36	BC	88	A	C4'-C3'-C2'	-6.22	96.38	102.60
38	BE	203	C	C5'-C4'-O4'	-6.22	101.63	109.10
68	Bi	114	HIS	CA-CB-CG	6.22	124.18	113.60
83	Bx	162	ARG	NE-CZ-NH1	6.22	123.41	120.30
85	AA	514	U	C5-C4-O4	-6.22	122.17	125.90
85	AA	605	A	OP1-P-OP2	-6.22	110.27	119.60
34	BA	582	U	C5-C6-N1	-6.22	119.59	122.70
34	BA	747	G	C5-C6-N1	6.22	114.61	111.50
35	BB	483	C	C6-N1-C2	6.22	122.79	120.30
35	BB	779	C	O4'-C1'-C2'	6.22	113.20	107.60
36	BC	148	C	O4'-C1'-N1	6.22	113.18	108.20
39	BF	32	G	C2-N3-C4	-6.22	108.79	111.90
40	BG	180	C	O4'-C4'-C3'	6.22	111.08	106.10
34	BA	102	G	O4'-C4'-C3'	-6.22	97.78	104.00
34	BA	308	C	N3-C2-O2	-6.22	117.55	121.90
34	BA	571	G	O4'-C1'-N9	6.22	113.18	108.20
34	BA	1270	G	N9-C1'-C2'	-6.22	105.16	112.00
34	BA	1496	G	P-O3'-C3'	6.22	127.16	119.70
34	BA	1565	U	C4'-C3'-C2'	6.22	108.82	102.60
34	BA	1641	G	P-O3'-C3'	-6.22	112.24	119.70
35	BB	1205	A	C6-C5-N7	-6.22	127.95	132.30
37	BD	75	G	O4'-C1'-N9	6.22	113.18	108.20
41	BH	104	U	N1-C1'-C2'	-6.22	105.16	112.00
70	Bk	71	ARG	NE-CZ-NH2	-6.22	117.19	120.30
85	AA	442	G	P-O3'-C3'	6.22	127.16	119.70
85	AA	469	G	C5-C6-O6	6.22	132.33	128.60
85	AA	1194	U	C2-N3-C4	-6.22	123.27	127.00
85	AA	1591	U	C5'-C4'-C3'	6.22	125.95	116.00
5	A4	4	GLN	N-CA-CB	6.22	121.79	110.60
34	BA	683	C	O4'-C1'-N1	6.22	113.17	108.20
35	BB	1354	C	P-O5'-C5'	-6.22	110.95	120.90
38	BE	193	A	C8-N9-C4	-6.22	103.31	105.80
85	AA	27	U	C5'-C4'-C3'	6.22	125.95	116.00
85	AA	349	C	C2-N1-C1'	-6.22	111.96	118.80
85	AA	1182	A	C5'-C4'-C3'	-6.22	106.05	116.00
85	AA	1279	A	C4'-C3'-C2'	-6.22	96.38	102.60
85	AA	1571	A	C3'-C2'-C1'	-6.22	96.53	101.50
8	A7	251	TRP	N-CA-C	-6.22	94.21	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	91	C	O4'-C1'-C2'	-6.22	99.58	105.80
34	BA	477	C	N3-C2-O2	-6.22	117.55	121.90
34	BA	554	A	OP2-P-O3'	6.22	118.88	105.20
34	BA	679	U	C6-N1-C1'	6.22	129.90	121.20
34	BA	987	C	C6-N1-C2	-6.22	117.81	120.30
34	BA	1709	A	O4'-C1'-C2'	6.22	113.19	107.60
35	BB	605	C	C2'-C3'-O3'	6.22	123.65	113.70
38	BE	65	U	C5'-C4'-C3'	-6.22	106.06	116.00
49	BP	98	ALA	CB-CA-C	6.22	119.43	110.10
67	Bh	142	GLY	N-CA-C	-6.22	97.56	113.10
85	AA	812	C	N1-C1'-C2'	-6.22	105.16	112.00
85	AA	844	C	C5-C6-N1	6.22	124.11	121.00
85	AA	1983	C	O3'-P-O5'	-6.22	92.19	104.00
34	BA	78	U	N3-C2-O2	-6.21	117.85	122.20
34	BA	120	A	C4'-C3'-C2'	-6.21	96.39	102.60
34	BA	1048	C	C2-N1-C1'	-6.21	111.96	118.80
34	BA	1144	A	P-O3'-C3'	-6.21	112.24	119.70
34	BA	1183	U	O4'-C1'-N1	6.21	113.17	108.20
34	BA	1591	G	C4'-C3'-C2'	6.21	108.81	102.60
35	BB	47	C	O4'-C1'-N1	6.21	113.17	108.20
35	BB	1205	A	N9-C4-C5	-6.21	103.31	105.80
85	AA	967	C	C2-N1-C1'	-6.21	111.97	118.80
85	AA	1277	C	C4'-C3'-C2'	-6.21	96.39	102.60
85	AA	1934	A	C6-N1-C2	-6.21	114.87	118.60
85	AA	2196	G	N1-C6-O6	-6.21	116.17	119.90
34	BA	189	G	C3'-C2'-C1'	-6.21	96.53	101.50
34	BA	927	A	O4'-C1'-C2'	6.21	113.19	107.60
35	BB	356	C	P-O5'-C5'	6.21	130.84	120.90
37	BD	16	U	C1'-O4'-C4'	-6.21	104.93	109.90
85	AA	439	U	C2-N3-C4	-6.21	123.27	127.00
85	AA	2146	G	C6-N1-C2	-6.21	121.37	125.10
86	AB	62	C	C1'-O4'-C4'	-6.21	104.93	109.90
6	A5	191	ARG	NE-CZ-NH2	-6.21	117.19	120.30
24	AQ	77	TYR	CB-CG-CD1	-6.21	117.27	121.00
34	BA	270	U	C6-N1-C1'	-6.21	112.50	121.20
34	BA	808	U	C5'-C4'-O4'	6.21	116.55	109.10
36	BC	52	A	P-O5'-C5'	-6.21	110.96	120.90
37	BD	49	A	N7-C8-N9	6.21	116.91	113.80
38	BE	79	G	N1-C6-O6	6.21	123.63	119.90
38	BE	188	C	O4'-C1'-C2'	-6.21	99.59	105.80
39	BF	52	A	C5'-C4'-O4'	6.21	116.55	109.10
40	BG	44	G	N9-C1'-C2'	-6.21	105.17	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
59	BZ	63	ARG	NE-CZ-NH2	-6.21	117.19	120.30
69	Bj	3	CYS	C-N-CD	-6.21	106.93	120.60
73	Bn	41	ARG	NE-CZ-NH1	6.21	123.41	120.30
85	AA	271	A	C5-C6-N6	-6.21	118.73	123.70
85	AA	313	A	N1-C6-N6	-6.21	114.87	118.60
85	AA	505	U	P-O5'-C5'	-6.21	110.96	120.90
85	AA	717	G	C8-N9-C1'	6.21	135.08	127.00
85	AA	1006	C	C2-N3-C4	-6.21	116.79	119.90
85	AA	1074	U	C1'-O4'-C4'	-6.21	104.93	109.90
85	AA	1660	U	N1-C1'-C2'	-6.21	105.17	112.00
85	AA	1690	A	P-O5'-C5'	-6.21	110.96	120.90
85	AA	2103	C	P-O5'-C5'	-6.21	110.96	120.90
34	BA	270	U	P-O3'-C3'	6.21	127.15	119.70
34	BA	765	U	C6-N1-C1'	-6.21	112.51	121.20
34	BA	1824	U	O5'-C5'-C4'	6.21	123.50	111.70
35	BB	750	G	C8-N9-C1'	6.21	135.07	127.00
35	BB	1118	G	C5'-C4'-C3'	-6.21	106.06	116.00
35	BB	1433	U	C2'-C3'-O3'	6.21	123.64	113.70
37	BD	16	U	C3'-C2'-C1'	-6.21	96.53	101.50
65	Bf	145	GLN	N-CA-CB	6.21	121.78	110.60
85	AA	704	A	C8-N9-C4	6.21	108.28	105.80
34	BA	169	C	O4'-C1'-N1	6.21	113.17	108.20
34	BA	959	G	O5'-P-OP2	-6.21	100.11	105.70
34	BA	1149	C	N1-C2-O2	6.21	122.63	118.90
34	BA	1519	G	C4'-C3'-C2'	-6.21	96.39	102.60
34	BA	1711	G	O5'-P-OP2	-6.21	100.11	105.70
35	BB	694	C	C2-N3-C4	6.21	123.00	119.90
35	BB	1018	U	C4-C5-C6	-6.21	115.97	119.70
39	BF	5	U	O4'-C1'-N1	6.21	113.17	108.20
41	BH	51	C	C2-N1-C1'	-6.21	111.97	118.80
85	AA	769	C	N1-C2-N3	6.21	123.55	119.20
34	BA	1241	U	O5'-P-OP1	-6.21	100.11	105.70
35	BB	567	G	C5-C6-N1	6.21	114.60	111.50
35	BB	1304	U	C6-N1-C1'	-6.21	112.51	121.20
36	BC	8	C	C2-N3-C4	6.21	123.00	119.90
39	BF	11	C	C6-N1-C1'	-6.21	113.35	120.80
39	BF	14	C	C5-C4-N4	-6.21	115.86	120.20
40	BG	68	U	C2-N3-C4	-6.21	123.28	127.00
40	BG	139	U	C6-N1-C1'	6.21	129.89	121.20
41	BH	48	G	P-O5'-C5'	-6.21	110.97	120.90
59	BZ	60	PHE	CB-CA-C	-6.21	97.99	110.40
85	AA	376	C	N3-C2-O2	-6.21	117.56	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2122	A	N9-C4-C5	6.21	108.28	105.80
85	AA	2199	G	N3-C2-N2	6.21	124.25	119.90
34	BA	587	U	C5-C4-O4	-6.21	122.18	125.90
35	BB	268	G	N1-C6-O6	6.21	123.62	119.90
38	BE	100	U	C1'-O4'-C4'	-6.21	104.94	109.90
41	BH	33	G	O4'-C1'-N9	6.21	113.16	108.20
85	AA	1617	G	C5-C6-O6	-6.21	124.88	128.60
85	AA	1700	C	C6-N1-C1'	6.21	128.25	120.80
34	BA	1702	G	C8-N9-C1'	6.20	135.06	127.00
35	BB	490	G	C5-C6-N1	6.20	114.60	111.50
35	BB	1250	A	C5-C6-N6	6.20	128.66	123.70
35	BB	1369	A	C5'-C4'-C3'	6.20	125.92	116.00
35	BB	1410	G	C5'-C4'-O4'	6.20	116.55	109.10
58	BY	5	ASP	N-CA-C	-6.20	94.25	111.00
85	AA	448	G	C4-N9-C1'	-6.20	118.44	126.50
85	AA	598	C	C5'-C4'-O4'	6.20	116.54	109.10
34	BA	291	C	C3'-C2'-C1'	-6.20	96.54	101.50
34	BA	306	G	C3'-C2'-C1'	-6.20	96.54	101.50
34	BA	1619	U	C2-N1-C1'	-6.20	110.26	117.70
34	BA	1805	C	P-O5'-C5'	6.20	130.82	120.90
37	BD	76	U	O4'-C1'-N1	6.20	113.16	108.20
38	BE	85	G	C2-N3-C4	6.20	115.00	111.90
68	Bi	45	ARG	NE-CZ-NH1	6.20	123.40	120.30
85	AA	1455	C	O4'-C4'-C3'	-6.20	97.80	104.00
85	AA	2074	G	C2'-C3'-O3'	6.20	123.62	113.70
34	BA	249	A	P-O3'-C3'	-6.20	112.26	119.70
34	BA	1498	A	C2'-C3'-O3'	6.20	123.62	113.70
36	BC	139	A	C1'-O4'-C4'	-6.20	104.94	109.90
37	BD	78	C	C6-N1-C1'	6.20	128.24	120.80
37	BD	92	G	N3-C4-C5	-6.20	125.50	128.60
39	BF	21	C	O3'-P-O5'	-6.20	92.22	104.00
39	BF	46	G	P-O3'-C3'	-6.20	112.26	119.70
67	Bh	34	ASP	CB-CA-C	6.20	122.80	110.40
77	Br	196	ARG	NE-CZ-NH2	6.20	123.40	120.30
85	AA	488	G	C8-N9-C4	-6.20	103.92	106.40
85	AA	924	A	C3'-C2'-C1'	-6.20	96.54	101.50
85	AA	1291	A	N1-C6-N6	-6.20	114.88	118.60
85	AA	1896	G	C6-C5-N7	-6.20	126.68	130.40
85	AA	2088	U	O4'-C1'-N1	6.20	113.16	108.20
34	BA	615	A	C1'-O4'-C4'	-6.20	104.94	109.90
34	BA	1242	A	C5'-C4'-C3'	-6.20	106.08	116.00
35	BB	34	G	C5-C6-N1	6.20	114.60	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	39	C	C5'-C4'-C3'	-6.20	106.08	116.00
35	BB	1057	G	O4'-C1'-N9	6.20	113.16	108.20
85	AA	1488	G	C4'-C3'-C2'	-6.20	96.40	102.60
85	AA	1644	G	C4-C5-C6	6.20	122.52	118.80
85	AA	2021	A	C5'-C4'-C3'	-6.20	106.08	116.00
16	AH	34	PHE	CB-CG-CD1	6.20	125.14	120.80
34	BA	629	G	N9-C4-C5	6.20	107.88	105.40
34	BA	963	G	O4'-C1'-N9	6.20	113.16	108.20
34	BA	1054	U	P-O5'-C5'	-6.20	110.98	120.90
35	BB	1416	A	P-O3'-C3'	-6.20	112.26	119.70
41	BH	100	A	C4-C5-N7	6.20	113.80	110.70
85	AA	116	G	C5-C6-O6	-6.20	124.88	128.60
85	AA	391	G	C5-C6-O6	6.20	132.32	128.60
15	AG	19	ARG	NE-CZ-NH1	6.20	123.40	120.30
34	BA	23	A	C5'-C4'-O4'	6.20	116.53	109.10
34	BA	954	U	N3-C2-O2	-6.20	117.86	122.20
34	BA	1197	U	C2-N3-C4	-6.20	123.28	127.00
35	BB	374	A	C8-N9-C4	6.20	108.28	105.80
35	BB	1191	G	C8-N9-C1'	6.20	135.05	127.00
36	BC	121	G	C3'-C2'-C1'	-6.20	96.54	101.50
39	BF	35	C	N3-C4-N4	6.20	122.34	118.00
85	AA	1371	C	N3-C2-O2	-6.20	117.56	121.90
85	AA	1615	A	C5'-C4'-C3'	-6.20	106.09	116.00
10	A9	118	ARG	NE-CZ-NH1	6.19	123.40	120.30
34	BA	186	G	N1-C6-O6	-6.19	116.18	119.90
39	BF	13	U	O4'-C4'-C3'	-6.19	97.81	104.00
39	BF	47	C	P-O3'-C3'	-6.19	112.27	119.70
45	BL	148	ARG	NE-CZ-NH1	6.19	123.40	120.30
85	AA	1291	A	O3'-P-O5'	-6.19	92.23	104.00
85	AA	1405	U	O4'-C1'-N1	6.19	113.16	108.20
2	A1	215	PHE	CB-CG-CD2	-6.19	116.47	120.80
34	BA	185	A	O5'-C5'-C4'	-6.19	99.94	111.70
34	BA	390	A	C5'-C4'-C3'	6.19	125.91	116.00
34	BA	1672	C	P-O5'-C5'	6.19	130.81	120.90
34	BA	1806	A	C8-N9-C4	6.19	108.28	105.80
35	BB	567	G	P-O5'-C5'	-6.19	110.99	120.90
40	BG	58	G	N1-C6-O6	6.19	123.61	119.90
55	BV	73	ARG	NE-CZ-NH2	-6.19	117.20	120.30
85	AA	116	G	C4-N9-C1'	-6.19	118.45	126.50
85	AA	769	C	N3-C4-C5	-6.19	119.42	121.90
34	BA	360	C	O4'-C1'-C2'	-6.19	99.61	105.80
34	BA	691	A	C4'-C3'-C2'	6.19	108.79	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	715	U	C4'-C3'-C2'	6.19	108.79	102.60
34	BA	863	G	N3-C4-C5	-6.19	125.50	128.60
34	BA	1019	C	C1'-O4'-C4'	-6.19	104.95	109.90
34	BA	1297	G	C3'-C2'-C1'	6.19	106.45	101.50
34	BA	1343	A	C5'-C4'-O4'	6.19	116.53	109.10
34	BA	1709	A	N1-C2-N3	-6.19	126.20	129.30
35	BB	866	A	N1-C6-N6	6.19	122.31	118.60
35	BB	1020	U	C6-N1-C2	-6.19	117.29	121.00
35	BB	1321	G	C5'-C4'-C3'	-6.19	106.09	116.00
42	BI	66	ARG	NE-CZ-NH1	6.19	123.39	120.30
53	BT	19	ARG	NE-CZ-NH1	6.19	123.39	120.30
85	AA	771	A	O5'-C5'-C4'	-6.19	99.94	111.70
85	AA	878	U	O5'-C5'-C4'	-6.19	99.94	111.70
85	AA	1436	A	C5'-C4'-C3'	6.19	125.91	116.00
85	AA	1576	G	O4'-C1'-N9	6.19	113.15	108.20
85	AA	1992	A	C2-N3-C4	-6.19	107.50	110.60
85	AA	2095	U	C6-N1-C1'	6.19	129.87	121.20
86	AB	5	G	C4-N9-C1'	-6.19	118.45	126.50
34	BA	1191	C	N3-C2-O2	-6.19	117.57	121.90
34	BA	1281	U	C6-N1-C2	-6.19	117.29	121.00
34	BA	1720	U	N1-C2-O2	6.19	127.13	122.80
39	BF	29	U	C3'-C2'-C1'	-6.19	96.55	101.50
41	BH	13	C	C1'-O4'-C4'	-6.19	104.95	109.90
53	BT	75	HIS	N-CA-CB	6.19	121.74	110.60
71	Bl	96	SER	C-N-CA	6.19	137.17	121.70
85	AA	1855	U	C2-N1-C1'	-6.19	110.27	117.70
34	BA	61	G	C3'-C2'-C1'	-6.19	96.55	101.50
34	BA	377	G	N1-C2-N2	-6.19	110.63	116.20
34	BA	577	U	O5'-P-OP2	6.19	118.13	110.70
34	BA	595	U	O4'-C1'-N1	6.19	113.15	108.20
34	BA	753	G	N3-C2-N2	6.19	124.23	119.90
34	BA	1711	G	P-O5'-C5'	-6.19	111.00	120.90
35	BB	292	U	O4'-C1'-N1	6.19	113.15	108.20
35	BB	476	A	O4'-C1'-N9	6.19	113.15	108.20
35	BB	1246	C	O4'-C1'-N1	6.19	113.15	108.20
35	BB	1540	U	P-O5'-C5'	-6.19	111.00	120.90
36	BC	63	G	C5'-C4'-O4'	6.19	116.53	109.10
36	BC	115	G	C3'-C2'-C1'	-6.19	96.55	101.50
38	BE	137	A	C8-N9-C1'	6.19	138.84	127.70
85	AA	205	A	O4'-C1'-N9	6.19	113.15	108.20
85	AA	1410	C	O4'-C1'-N1	6.19	113.15	108.20
85	AA	1483	A	N1-C6-N6	-6.19	114.89	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	237	A	N7-C8-N9	-6.19	110.71	113.80
34	BA	488	C	OP1-P-OP2	-6.19	110.32	119.60
34	BA	772	G	C5'-C4'-O4'	6.19	116.52	109.10
34	BA	1271	C	O4'-C1'-N1	6.19	113.15	108.20
34	BA	1297	G	P-O3'-C3'	6.19	127.12	119.70
85	AA	1492	U	C1'-O4'-C4'	-6.19	104.95	109.90
13	AE	104	ARG	NE-CZ-NH1	-6.18	117.21	120.30
34	BA	1201	G	OP2-P-O3'	6.18	118.81	105.20
34	BA	1513	G	P-O3'-C3'	-6.18	112.28	119.70
35	BB	584	A	P-O3'-C3'	-6.18	112.28	119.70
35	BB	637	G	N9-C4-C5	6.18	107.87	105.40
35	BB	1408	G	O4'-C1'-N9	6.18	113.15	108.20
35	BB	1438	U	C3'-C2'-C1'	-6.18	96.55	101.50
37	BD	94	C	C5'-C4'-C3'	-6.18	106.10	116.00
40	BG	23	C	N1-C1'-C2'	-6.18	105.20	112.00
40	BG	25	G	N9-C1'-C2'	-6.18	105.20	112.00
40	BG	161	C	P-O3'-C3'	6.18	127.12	119.70
85	AA	292	C	C3'-C2'-C1'	-6.18	96.55	101.50
85	AA	422	G	N3-C4-C5	-6.18	125.51	128.60
85	AA	923	A	C5'-C4'-O4'	6.18	116.52	109.10
85	AA	1209	U	C4'-C3'-C2'	6.18	108.78	102.60
85	AA	1274	A	P-O3'-C3'	-6.18	112.28	119.70
85	AA	1359	U	O4'-C1'-N1	6.18	113.15	108.20
2	A1	96	PHE	CB-CG-CD2	-6.18	116.47	120.80
34	BA	47	U	C6-N1-C1'	6.18	129.85	121.20
34	BA	218	G	N1-C6-O6	6.18	123.61	119.90
34	BA	1210	A	C5'-C4'-C3'	-6.18	106.11	116.00
34	BA	1419	A	C6-N1-C2	-6.18	114.89	118.60
34	BA	1425	G	C5-C6-O6	-6.18	124.89	128.60
34	BA	1627	U	P-O5'-C5'	6.18	130.79	120.90
35	BB	1035	C	P-O5'-C5'	-6.18	111.01	120.90
35	BB	1040	C	OP1-P-OP2	-6.18	110.33	119.60
35	BB	1107	C	C6-N1-C2	-6.18	117.83	120.30
38	BE	20	C	N1-C2-O2	6.18	122.61	118.90
65	Bf	449	GLU	N-CA-CB	6.18	121.73	110.60
85	AA	291	G	O4'-C1'-N9	6.18	113.15	108.20
85	AA	339	A	C1'-O4'-C4'	-6.18	104.95	109.90
85	AA	687	G	N9-C1'-C2'	-6.18	105.20	112.00
85	AA	822	U	P-O5'-C5'	6.18	130.79	120.90
85	AA	2042	G	O4'-C1'-C2'	6.18	113.16	107.60
86	AB	31	A	O4'-C1'-N9	6.18	113.15	108.20
34	BA	983	A	C8-N9-C4	6.18	108.27	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	717	A	C8-N9-C4	-6.18	103.33	105.80
40	BG	139	U	C2-N1-C1'	-6.18	110.28	117.70
77	Br	137	MET	CG-SD-CE	-6.18	90.31	100.20
85	AA	196	U	C5-C6-N1	-6.18	119.61	122.70
85	AA	1256	C	C1'-O4'-C4'	-6.18	104.96	109.90
34	BA	187	G	N3-C2-N2	6.18	124.23	119.90
35	BB	102	G	O5'-P-OP2	6.18	118.11	110.70
35	BB	119	G	C1'-O4'-C4'	-6.18	104.96	109.90
35	BB	329	U	C2-N1-C1'	6.18	125.11	117.70
35	BB	1183	U	O3'-P-O5'	6.18	115.74	104.00
85	AA	18	C	C5'-C4'-C3'	6.18	125.89	116.00
85	AA	210	G	O3'-P-O5'	-6.18	92.26	104.00
85	AA	271	A	O4'-C1'-N9	6.18	113.14	108.20
85	AA	722	G	O3'-P-O5'	-6.18	92.26	104.00
85	AA	995	G	O4'-C1'-N9	6.18	113.14	108.20
85	AA	1542	A	C4'-C3'-C2'	-6.18	96.42	102.60
85	AA	1794	U	C4'-C3'-C2'	-6.18	96.42	102.60
85	AA	1934	A	C8-N9-C4	-6.18	103.33	105.80
1	A0	213	ARG	CD-NE-CZ	-6.18	114.95	123.60
5	A4	163	ASP	CA-CB-CG	-6.18	99.81	113.40
34	BA	16	C	N1-C2-N3	6.18	123.53	119.20
34	BA	278	U	C5'-C4'-C3'	-6.18	106.12	116.00
34	BA	1514	A	C4-N9-C1'	-6.18	115.18	126.30
35	BB	144	G	C4-N9-C1'	-6.18	118.47	126.50
77	Br	210	MET	CG-SD-CE	-6.18	90.31	100.20
85	AA	1458	G	C2-N3-C4	6.18	114.99	111.90
85	AA	1698	A	C1'-O4'-C4'	-6.18	104.96	109.90
34	BA	964	U	N1-C2-O2	-6.18	118.48	122.80
34	BA	1052	G	O4'-C1'-C2'	6.18	113.16	107.60
34	BA	1754	C	O4'-C1'-N1	6.18	113.14	108.20
35	BB	1230	A	C8-N9-C1'	6.18	138.82	127.70
36	BC	147	G	N1-C2-N2	-6.18	110.64	116.20
38	BE	62	C	O4'-C1'-C2'	6.18	113.16	107.60
39	BF	65	U	O5'-C5'-C4'	-6.18	99.96	111.70
85	AA	588	G	C5'-C4'-C3'	-6.18	106.12	116.00
85	AA	1432	C	C6-N1-C2	-6.18	117.83	120.30
85	AA	1557	U	P-O5'-C5'	6.18	130.78	120.90
8	A7	264	ASP	CB-CG-OD1	6.17	123.86	118.30
34	BA	50	G	N7-C8-N9	-6.17	110.01	113.10
34	BA	764	G	P-O5'-C5'	6.17	130.78	120.90
34	BA	1221	A	C4-N9-C1'	-6.17	115.19	126.30
35	BB	84	G	O4'-C1'-N9	6.17	113.14	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	618	U	C2-N1-C1'	-6.17	110.29	117.70
35	BB	768	A	C8-N9-C1'	6.17	138.82	127.70
36	BC	154	A	N9-C1'-C2'	-6.17	105.21	112.00
38	BE	38	C	O4'-C1'-N1	6.17	113.14	108.20
38	BE	160	C	O4'-C1'-N1	6.17	113.14	108.20
57	BX	101	ASN	CB-CA-C	6.17	122.75	110.40
62	Bc	111	ARG	NE-CZ-NH1	6.17	123.39	120.30
85	AA	412	G	P-O5'-C5'	-6.17	111.02	120.90
85	AA	1142	G	C1'-O4'-C4'	-6.17	104.96	109.90
85	AA	2077	G	C8-N9-C1'	6.17	135.03	127.00
85	AA	2102	A	C4'-C3'-C2'	-6.17	96.43	102.60
85	AA	2211	G	O3'-P-O5'	6.17	115.73	104.00
34	BA	375	C	N3-C2-O2	-6.17	117.58	121.90
81	Bv	75	ARG	NE-CZ-NH1	6.17	123.39	120.30
85	AA	485	A	C5'-C4'-O4'	6.17	116.51	109.10
34	BA	14	G	C4-N9-C1'	-6.17	118.48	126.50
34	BA	160	G	C4-C5-C6	-6.17	115.10	118.80
34	BA	648	C	C6-N1-C2	-6.17	117.83	120.30
34	BA	723	C	C6-N1-C2	-6.17	117.83	120.30
34	BA	1097	G	P-O5'-C5'	6.17	130.77	120.90
34	BA	1488	C	O5'-C5'-C4'	6.17	123.43	111.70
35	BB	64	U	O4'-C1'-N1	6.17	113.14	108.20
35	BB	368	C	O4'-C1'-C2'	6.17	113.15	107.60
35	BB	830	G	C1'-O4'-C4'	-6.17	104.96	109.90
35	BB	1003	G	C4-N9-C1'	-6.17	118.48	126.50
35	BB	1016	C	N3-C2-O2	-6.17	117.58	121.90
35	BB	1342	C	C4'-C3'-C2'	6.17	108.77	102.60
40	BG	9	G	N1-C6-O6	-6.17	116.20	119.90
85	AA	95	U	C2-N1-C1'	-6.17	110.29	117.70
85	AA	139	G	N3-C2-N2	6.17	124.22	119.90
85	AA	992	G	C5'-C4'-C3'	-6.17	106.13	116.00
85	AA	1092	G	O4'-C1'-N9	6.17	113.14	108.20
85	AA	2043	A	C4'-C3'-C2'	-6.17	96.43	102.60
85	AA	2114	U	C6-N1-C1'	6.17	129.84	121.20
86	AB	15	G	O5'-C5'-C4'	6.17	123.42	111.70
86	AB	56	C	C5'-C4'-O4'	6.17	116.51	109.10
34	BA	29	U	C5-C4-O4	6.17	129.60	125.90
34	BA	825	G	C5-C6-N1	6.17	114.58	111.50
34	BA	1163	G	N3-C2-N2	6.17	124.22	119.90
34	BA	1326	U	C6-N1-C1'	-6.17	112.56	121.20
34	BA	1538	G	O4'-C1'-N9	6.17	113.14	108.20
35	BB	892	U	C4'-C3'-C2'	-6.17	96.43	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	75	G	C3'-C2'-C1'	-6.17	96.56	101.50
62	Bc	136	PHE	CB-CG-CD1	6.17	125.12	120.80
85	AA	496	C	N1-C2-O2	6.17	122.60	118.90
85	AA	1796	C	O4'-C4'-C3'	-6.17	97.83	104.00
34	BA	38	G	O4'-C1'-N9	6.17	113.14	108.20
34	BA	740	A	P-O3'-C3'	-6.17	112.30	119.70
34	BA	1173	C	O4'-C1'-C2'	6.17	113.15	107.60
34	BA	1400	A	N1-C6-N6	6.17	122.30	118.60
36	BC	76	C	C4'-C3'-C2'	-6.17	96.43	102.60
37	BD	26	C	C2-N3-C4	-6.17	116.82	119.90
37	BD	44	U	N3-C2-O2	-6.17	117.88	122.20
40	BG	23	C	O5'-P-OP2	6.17	118.10	110.70
40	BG	101	G	N9-C1'-C2'	-6.17	105.22	112.00
41	BH	66	G	C1'-O4'-C4'	-6.17	104.96	109.90
41	BH	76	G	C5-C6-O6	6.17	132.30	128.60
48	BO	75	PHE	CB-CG-CD2	-6.17	116.48	120.80
76	Bq	45	ARG	N-CA-CB	6.17	121.70	110.60
85	AA	340	G	C5'-C4'-C3'	-6.17	106.13	116.00
85	AA	718	C	C2-N1-C1'	-6.17	112.02	118.80
34	BA	514	U	P-O3'-C3'	-6.17	112.30	119.70
34	BA	712	C	OP1-P-OP2	-6.17	110.35	119.60
34	BA	769	U	O4'-C1'-N1	6.17	113.13	108.20
34	BA	1215	U	N1-C2-O2	6.17	127.12	122.80
34	BA	1332	U	C2-N3-C4	-6.17	123.30	127.00
34	BA	1456	C	O4'-C1'-N1	6.17	113.13	108.20
34	BA	1739	G	C4-C5-C6	-6.17	115.10	118.80
35	BB	449	C	N3-C4-C5	6.17	124.37	121.90
35	BB	950	G	C5-C6-O6	-6.17	124.90	128.60
35	BB	1177	U	O4'-C4'-C3'	-6.17	97.83	104.00
35	BB	1385	C	C1'-O4'-C4'	-6.17	104.97	109.90
38	BE	114	G	C1'-O4'-C4'	-6.17	104.97	109.90
83	Bx	91	PHE	CB-CG-CD1	-6.17	116.48	120.80
85	AA	349	C	C6-N1-C1'	6.17	128.20	120.80
85	AA	862	U	C2'-C3'-O3'	6.17	123.57	113.70
34	BA	250	G	O4'-C1'-N9	6.17	113.13	108.20
35	BB	351	G	N1-C6-O6	6.17	123.60	119.90
35	BB	714	U	N1-C2-N3	6.17	118.60	114.90
35	BB	1247	C	O4'-C1'-N1	6.17	113.13	108.20
35	BB	1297	G	O4'-C1'-N9	6.17	113.13	108.20
35	BB	1485	G	N3-C2-N2	6.17	124.22	119.90
36	BC	60	U	N3-C4-O4	6.17	123.72	119.40
41	BH	119	U	OP1-P-O3'	6.17	118.76	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1496	U	N1-C1'-C2'	-6.17	105.22	112.00
85	AA	2054	G	P-O5'-C5'	6.17	130.76	120.90
34	BA	960	C	N3-C2-O2	-6.16	117.59	121.90
34	BA	1019	C	C6-N1-C1'	-6.16	113.40	120.80
34	BA	1156	U	C5-C6-N1	-6.16	119.62	122.70
34	BA	1254	C	C6-N1-C2	-6.16	117.83	120.30
34	BA	1273	U	O3'-P-O5'	-6.16	92.29	104.00
34	BA	1532	G	N1-C2-N2	-6.16	110.65	116.20
34	BA	1796	A	C1'-O4'-C4'	-6.16	104.97	109.90
34	BA	1804	A	C1'-O4'-C4'	-6.16	104.97	109.90
35	BB	491	A	P-O3'-C3'	6.16	127.10	119.70
35	BB	844	G	O4'-C1'-N9	6.16	113.13	108.20
41	BH	12	U	N3-C4-C5	-6.16	110.90	114.60
65	Bf	360	ALA	N-CA-CB	6.16	118.73	110.10
85	AA	30	G	P-O3'-C3'	-6.16	112.31	119.70
85	AA	440	U	N1-C1'-C2'	-6.16	105.22	112.00
85	AA	929	G	N1-C6-O6	6.16	123.60	119.90
85	AA	1100	U	O3'-P-O5'	-6.16	92.29	104.00
85	AA	1862	C	O4'-C1'-N1	6.16	113.13	108.20
34	BA	161	U	C2'-C3'-O3'	6.16	123.56	113.70
34	BA	383	G	C2'-C3'-O3'	6.16	123.56	113.70
34	BA	1370	A	C5'-C4'-C3'	-6.16	106.14	116.00
35	BB	1048	A	C8-N9-C4	6.16	108.27	105.80
85	AA	25	C	O5'-P-OP1	6.16	118.09	110.70
85	AA	1184	A	P-O3'-C3'	-6.16	112.31	119.70
34	BA	526	C	C5-C4-N4	-6.16	115.89	120.20
34	BA	649	A	O4'-C1'-N9	6.16	113.13	108.20
34	BA	774	A	C5'-C4'-O4'	-6.16	101.71	109.10
34	BA	1120	U	C3'-C2'-C1'	-6.16	96.57	101.50
34	BA	1162	U	C6-N1-C2	-6.16	117.30	121.00
34	BA	1691	G	O3'-P-O5'	6.16	115.71	104.00
35	BB	257	G	C5-C6-O6	-6.16	124.90	128.60
35	BB	770	G	P-O5'-C5'	6.16	130.76	120.90
35	BB	1023	G	C1'-O4'-C4'	-6.16	104.97	109.90
36	BC	15	G	N1-C6-O6	6.16	123.60	119.90
67	Bh	53	TYR	CB-CG-CD2	-6.16	117.30	121.00
85	AA	36	U	C2-N3-C4	-6.16	123.30	127.00
85	AA	1116	G	C4-N9-C1'	-6.16	118.49	126.50
85	AA	1560	A	O4'-C4'-C3'	-6.16	97.84	104.00
16	AH	13	SER	O-C-N	-6.16	112.85	122.70
34	BA	119	G	O4'-C1'-N9	6.16	113.13	108.20
34	BA	367	G	N1-C6-O6	6.16	123.59	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1000	G	N1-C6-O6	-6.16	116.20	119.90
34	BA	1034	U	C5'-C4'-C3'	-6.16	106.15	116.00
34	BA	1704	G	C8-N9-C1'	6.16	135.00	127.00
35	BB	837	A	O3'-P-O5'	6.16	115.70	104.00
39	BF	9	C	N3-C4-N4	-6.16	113.69	118.00
40	BG	85	C	C2-N1-C1'	6.16	125.57	118.80
51	BR	16	LYS	CB-CA-C	-6.16	98.08	110.40
53	BT	6	LEU	N-CA-CB	6.16	122.72	110.40
64	Be	83	PHE	CB-CG-CD1	-6.16	116.49	120.80
85	AA	100	A	C8-N9-C4	-6.16	103.34	105.80
85	AA	245	A	C4-N9-C1'	-6.16	115.21	126.30
85	AA	424	A	C5-N7-C8	-6.16	100.82	103.90
85	AA	587	G	C8-N9-C1'	6.16	135.01	127.00
85	AA	768	C	P-O5'-C5'	-6.16	111.05	120.90
85	AA	938	A	P-O5'-C5'	6.16	130.75	120.90
34	BA	712	C	N3-C2-O2	-6.16	117.59	121.90
34	BA	1300	G	N1-C6-O6	6.16	123.59	119.90
35	BB	60	A	O4'-C1'-N9	6.16	113.13	108.20
35	BB	1045	G	C6-N1-C2	-6.16	121.41	125.10
85	AA	1694	C	O4'-C1'-N1	6.16	113.12	108.20
5	A4	127	PHE	CB-CG-CD1	6.16	125.11	120.80
34	BA	218	G	O4'-C1'-N9	6.16	113.12	108.20
34	BA	304	G	N9-C1'-C2'	-6.16	105.23	112.00
34	BA	657	C	O4'-C1'-N1	6.16	113.12	108.20
34	BA	1211	G	C4-N9-C1'	6.16	134.50	126.50
35	BB	419	G	C5-C6-O6	-6.16	124.91	128.60
35	BB	509	A	N1-C6-N6	6.16	122.29	118.60
35	BB	1452	U	P-O5'-C5'	6.16	130.75	120.90
37	BD	83	A	C8-N9-C4	6.16	108.26	105.80
38	BE	10	G	N3-C2-N2	-6.16	115.59	119.90
85	AA	826	C	C5'-C4'-O4'	6.16	116.49	109.10
85	AA	1063	U	O5'-C5'-C4'	6.16	123.39	111.70
85	AA	1704	C	N3-C4-C5	6.16	124.36	121.90
85	AA	2056	C	C5'-C4'-C3'	-6.16	106.15	116.00
34	BA	605	G	C5-C6-N1	6.15	114.58	111.50
34	BA	652	C	C4'-C3'-C2'	-6.15	96.45	102.60
35	BB	508	U	C2-N1-C1'	-6.15	110.32	117.70
35	BB	1259	A	C1'-O4'-C4'	-6.15	104.98	109.90
85	AA	154	U	O4'-C1'-N1	6.15	113.12	108.20
85	AA	338	G	C4-N9-C1'	-6.15	118.50	126.50
85	AA	691	U	O4'-C1'-N1	6.15	113.12	108.20
85	AA	2194	U	O4'-C1'-C2'	6.15	113.14	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	147	U	O5'-C5'-C4'	-6.15	100.01	111.70
34	BA	280	A	C4'-C3'-C2'	6.15	108.75	102.60
34	BA	833	U	C4'-C3'-C2'	-6.15	96.45	102.60
34	BA	1185	U	O4'-C1'-N1	6.15	113.12	108.20
34	BA	1731	A	O4'-C4'-C3'	-6.15	97.85	104.00
35	BB	955	U	O4'-C1'-N1	6.15	113.12	108.20
37	BD	51	G	C8-N9-C4	6.15	108.86	106.40
40	BG	46	G	N1-C6-O6	6.15	123.59	119.90
57	BX	60	TYR	CB-CG-CD1	-6.15	117.31	121.00
85	AA	83	U	P-O5'-C5'	6.15	130.74	120.90
85	AA	521	A	P-O5'-C5'	6.15	130.75	120.90
31	AX	55	ARG	NE-CZ-NH2	-6.15	117.22	120.30
34	BA	13	U	C4-C5-C6	-6.15	116.01	119.70
34	BA	47	U	C2-N3-C4	-6.15	123.31	127.00
34	BA	605	G	O4'-C1'-N9	6.15	113.12	108.20
34	BA	768	G	C5'-C4'-C3'	-6.15	106.16	116.00
34	BA	851	C	C5'-C4'-C3'	6.15	125.84	116.00
34	BA	1212	A	C5-C6-N1	6.15	120.78	117.70
34	BA	1341	A	O4'-C4'-C3'	-6.15	97.85	104.00
35	BB	501	G	N3-C2-N2	6.15	124.21	119.90
35	BB	638	G	C4-N9-C1'	-6.15	118.50	126.50
36	BC	135	A	C5'-C4'-C3'	-6.15	106.16	116.00
37	BD	30	A	C8-N9-C4	6.15	108.26	105.80
39	BF	64	U	C6-N1-C1'	-6.15	112.59	121.20
41	BH	21	G	N9-C1'-C2'	-6.15	105.23	112.00
43	BJ	57	ARG	NE-CZ-NH1	6.15	123.38	120.30
85	AA	1259	U	P-O3'-C3'	-6.15	112.32	119.70
85	AA	1535	C	C3'-C2'-C1'	6.15	106.42	101.50
85	AA	1568	U	O4'-C1'-N1	6.15	113.12	108.20
27	AT	10	VAL	CB-CA-C	-6.15	99.72	111.40
34	BA	89	G	C5'-C4'-C3'	6.15	125.84	116.00
34	BA	503	C	P-O3'-C3'	-6.15	112.32	119.70
34	BA	767	U	P-O5'-C5'	-6.15	111.06	120.90
34	BA	1295	U	O4'-C1'-C2'	6.15	113.13	107.60
34	BA	1747	C	O3'-P-O5'	6.15	115.68	104.00
35	BB	1268	C	N3-C2-O2	-6.15	117.59	121.90
38	BE	106	C	N3-C4-N4	6.15	122.30	118.00
85	AA	305	A	O4'-C4'-C3'	-6.15	97.85	104.00
85	AA	2086	C	O4'-C1'-N1	6.15	113.12	108.20
34	BA	289	A	C5-C6-N1	6.15	120.77	117.70
34	BA	1087	A	C6-N1-C2	-6.15	114.91	118.60
35	BB	1452	U	C5'-C4'-C3'	6.15	125.84	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	BL	167	MET	CG-SD-CE	-6.15	90.36	100.20
59	BZ	40	TYR	CA-CB-CG	-6.15	101.72	113.40
69	Bj	10	ARG	NE-CZ-NH1	6.15	123.37	120.30
85	AA	163	C	C6-N1-C1'	-6.15	113.42	120.80
85	AA	338	G	C1'-O4'-C4'	-6.15	104.98	109.90
85	AA	447	C	C4'-C3'-C2'	6.15	108.75	102.60
85	AA	451	G	N3-C2-N2	6.15	124.20	119.90
85	AA	667	A	P-O5'-C5'	-6.15	111.06	120.90
85	AA	692	U	C2-N3-C4	-6.15	123.31	127.00
85	AA	1526	G	C8-N9-C4	-6.15	103.94	106.40
85	AA	1909	C	C4'-C3'-C2'	-6.15	96.45	102.60
85	AA	1932	C	C6-N1-C1'	-6.15	113.42	120.80
85	AA	1974	C	P-O5'-C5'	-6.15	111.06	120.90
29	AV	53	ARG	NE-CZ-NH1	6.15	123.37	120.30
34	BA	217	C	C5'-C4'-C3'	-6.15	106.17	116.00
34	BA	703	U	C6-N1-C2	-6.15	117.31	121.00
36	BC	108	A	C5'-C4'-O4'	6.15	116.47	109.10
37	BD	93	G	C5-C6-N1	6.15	114.57	111.50
34	BA	3	G	O4'-C1'-N9	6.14	113.11	108.20
34	BA	277	A	N7-C8-N9	-6.14	110.73	113.80
34	BA	449	G	N7-C8-N9	-6.14	110.03	113.10
34	BA	1087	A	C5'-C4'-C3'	-6.14	106.17	116.00
34	BA	1515	U	N1-C1'-C2'	-6.14	105.24	112.00
34	BA	1583	A	C8-N9-C4	6.14	108.26	105.80
34	BA	1711	G	C2-N3-C4	6.14	114.97	111.90
35	BB	474	G	C5-C6-O6	-6.14	124.91	128.60
35	BB	557	C	N3-C2-O2	-6.14	117.60	121.90
35	BB	857	G	C5-C6-O6	-6.14	124.91	128.60
35	BB	1346	A	C5'-C4'-C3'	6.14	125.83	116.00
37	BD	80	G	C5-C6-N1	6.14	114.57	111.50
41	BH	22	A	C4'-C3'-O3'	-6.14	96.50	109.40
41	BH	52	G	C5-C6-O6	-6.14	124.91	128.60
65	Bf	268	SER	N-CA-CB	6.14	119.72	110.50
85	AA	145	C	C1'-O4'-C4'	-6.14	104.98	109.90
85	AA	536	C	C1'-O4'-C4'	-6.14	104.98	109.90
85	AA	551	C	C6-N1-C2	-6.14	117.84	120.30
85	AA	1198	U	C2-N3-C4	-6.14	123.31	127.00
34	BA	513	U	O4'-C1'-C2'	6.14	113.13	107.60
34	BA	1325	G	C5-C6-N1	6.14	114.57	111.50
34	BA	1666	U	C1'-O4'-C4'	-6.14	104.99	109.90
34	BA	1726	U	C6-N1-C2	-6.14	117.31	121.00
35	BB	552	C	O4'-C1'-C2'	6.14	113.13	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	106	C	P-O3'-C3'	-6.14	112.33	119.70
41	BH	63	G	C2'-C3'-O3'	6.14	123.53	113.70
41	BH	103	C	C5'-C4'-C3'	6.14	125.83	116.00
71	Bl	93	VAL	N-CA-CB	-6.14	97.99	111.50
85	AA	114	C	P-O3'-C3'	-6.14	112.33	119.70
85	AA	383	C	C5-C4-N4	6.14	124.50	120.20
85	AA	1218	C	O4'-C1'-N1	6.14	113.11	108.20
85	AA	2046	G	C1'-O4'-C4'	-6.14	104.99	109.90
34	BA	1187	U	C3'-C2'-C1'	-6.14	96.59	101.50
35	BB	1070	G	C3'-C2'-C1'	-6.14	96.59	101.50
35	BB	1319	U	P-O3'-C3'	-6.14	112.33	119.70
85	AA	764	U	C2-N3-C4	6.14	130.68	127.00
85	AA	1378	U	C2-N3-C4	-6.14	123.31	127.00
85	AA	2204	A	C8-N9-C4	6.14	108.26	105.80
3	A2	39	ALA	N-CA-CB	6.14	118.70	110.10
34	BA	917	C	N1-C2-O2	6.14	122.58	118.90
34	BA	1030	C	C1'-O4'-C4'	-6.14	104.99	109.90
34	BA	1052	G	C8-N9-C4	6.14	108.86	106.40
34	BA	1231	C	C3'-C2'-C1'	-6.14	96.59	101.50
34	BA	1233	U	O4'-C1'-C2'	6.14	113.13	107.60
34	BA	1365	U	O3'-P-O5'	-6.14	92.34	104.00
34	BA	1472	G	C5-C6-N1	6.14	114.57	111.50
34	BA	1788	U	P-O5'-C5'	6.14	130.72	120.90
34	BA	1807	G	N3-C2-N2	6.14	124.20	119.90
35	BB	126	C	C1'-O4'-C4'	-6.14	104.99	109.90
35	BB	130	G	O4'-C1'-C2'	6.14	113.12	107.60
35	BB	775	U	N3-C4-O4	-6.14	115.10	119.40
35	BB	1063	C	C5'-C4'-C3'	-6.14	106.18	116.00
35	BB	1397	G	C5'-C4'-C3'	-6.14	106.17	116.00
40	BG	119	A	P-O3'-C3'	6.14	127.07	119.70
44	BK	10	ARG	NE-CZ-NH2	-6.14	117.23	120.30
73	Bn	57	ARG	NE-CZ-NH1	6.14	123.37	120.30
74	Bo	47	PHE	CA-CB-CG	-6.14	99.16	113.90
85	AA	115	U	O3'-P-O5'	-6.14	92.34	104.00
85	AA	217	G	N1-C6-O6	6.14	123.58	119.90
85	AA	258	G	C4-N9-C1'	-6.14	118.52	126.50
85	AA	744	C	C4-C5-C6	-6.14	114.33	117.40
85	AA	1106	A	C4'-C3'-C2'	-6.14	96.46	102.60
85	AA	1521	U	O3'-P-O5'	-6.14	92.34	104.00
85	AA	2145	G	C5-C6-N1	6.14	114.57	111.50
34	BA	549	G	N1-C6-O6	-6.14	116.22	119.90
34	BA	1418	G	O5'-P-OP2	-6.14	100.18	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1091	C	C5'-C4'-C3'	-6.14	106.18	116.00
37	BD	68	C	N3-C2-O2	-6.14	117.60	121.90
37	BD	79	G	C4'-C3'-C2'	6.14	108.74	102.60
38	BE	154	A	N9-C1'-C2'	-6.14	105.25	112.00
85	AA	161	A	C5'-C4'-O4'	6.14	116.47	109.10
85	AA	1576	G	C4-C5-C6	-6.14	115.12	118.80
85	AA	2102	A	N9-C1'-C2'	-6.14	105.25	112.00
23	AP	136	ARG	NE-CZ-NH2	-6.14	117.23	120.30
34	BA	33	C	O4'-C1'-N1	6.14	113.11	108.20
34	BA	294	C	O3'-P-O5'	6.14	115.66	104.00
34	BA	401	A	C4-C5-C6	-6.14	113.93	117.00
34	BA	523	A	C4-N9-C1'	-6.14	115.25	126.30
34	BA	617	G	P-O3'-C3'	-6.14	112.34	119.70
34	BA	678	C	C3'-C2'-C1'	6.14	106.41	101.50
34	BA	1058	C	O3'-P-O5'	6.14	115.66	104.00
35	BB	829	C	P-O3'-C3'	6.14	127.06	119.70
35	BB	1143	A	P-O3'-C3'	-6.14	112.34	119.70
35	BB	1442	C	C5'-C4'-C3'	6.14	125.82	116.00
39	BF	17	U	C4-C5-C6	-6.14	116.02	119.70
41	BH	49	C	C6-N1-C1'	-6.14	113.44	120.80
65	Bf	416	PHE	CB-CG-CD2	-6.14	116.50	120.80
84	By	173	PHE	CB-CG-CD1	6.14	125.10	120.80
85	AA	389	A	C1'-O4'-C4'	-6.14	104.99	109.90
85	AA	478	U	C5'-C4'-C3'	-6.14	106.18	116.00
85	AA	719	C	C3'-C2'-C1'	-6.14	96.59	101.50
85	AA	1251	G	C5-C6-O6	-6.14	124.92	128.60
24	AQ	42	ASN	CA-CB-CG	-6.13	99.91	113.40
34	BA	1176	C	N3-C2-O2	-6.13	117.61	121.90
35	BB	1315	C	O5'-C5'-C4'	-6.13	100.05	111.70
35	BB	1379	U	P-O3'-C3'	6.13	127.06	119.70
35	BB	1487	G	P-O5'-C5'	-6.13	111.08	120.90
40	BG	29	U	N3-C4-C5	6.13	118.28	114.60
63	Bd	29	MET	CG-SD-CE	-6.13	90.39	100.20
71	Bl	139	ARG	NE-CZ-NH1	6.13	123.37	120.30
85	AA	146	U	P-O5'-C5'	-6.13	111.08	120.90
85	AA	909	C	C2'-C3'-O3'	6.13	123.51	113.70
85	AA	920	A	N9-C1'-C2'	-6.13	105.25	112.00
85	AA	922	A	C5'-C4'-O4'	6.13	116.46	109.10
85	AA	2096	G	C8-N9-C4	-6.13	103.95	106.40
34	BA	1011	G	N1-C6-O6	6.13	123.58	119.90
34	BA	1260	G	C8-N9-C1'	6.13	134.97	127.00
35	BB	148	C	P-O3'-C3'	6.13	127.06	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	386	G	C5-C6-O6	-6.13	124.92	128.60
35	BB	604	C	C1'-O4'-C4'	-6.13	104.99	109.90
35	BB	1164	U	C6-N1-C1'	6.13	129.79	121.20
38	BE	102	U	C6-N1-C1'	-6.13	112.61	121.20
41	BH	130	G	C5-C6-N1	6.13	114.57	111.50
85	AA	862	U	C6-N1-C2	-6.13	117.32	121.00
85	AA	1616	U	O4'-C1'-N1	6.13	113.11	108.20
86	AB	32	U	C2-N1-C1'	-6.13	110.34	117.70
34	BA	686	U	N3-C2-O2	-6.13	117.91	122.20
34	BA	881	C	N1-C2-O2	6.13	122.58	118.90
34	BA	1240	G	C4-C5-N7	6.13	113.25	110.80
34	BA	1540	C	C5-C4-N4	-6.13	115.91	120.20
35	BB	506	G	N3-C2-N2	6.13	124.19	119.90
35	BB	985	A	P-O3'-C3'	6.13	127.06	119.70
35	BB	1077	C	N3-C4-N4	-6.13	113.71	118.00
35	BB	1279	C	C5'-C4'-C3'	6.13	125.81	116.00
35	BB	1300	U	O4'-C1'-N1	6.13	113.11	108.20
35	BB	1481	C	O5'-C5'-C4'	6.13	123.35	111.70
36	BC	168	C	C5'-C4'-O4'	6.13	116.46	109.10
40	BG	172	C	C5'-C4'-O4'	6.13	116.46	109.10
41	BH	101	A	C1'-O4'-C4'	-6.13	105.00	109.90
85	AA	80	G	C4'-C3'-C2'	-6.13	96.47	102.60
85	AA	643	C	C3'-C2'-C1'	-6.13	96.59	101.50
85	AA	717	G	C4-N9-C1'	-6.13	118.53	126.50
85	AA	739	C	C4'-C3'-C2'	-6.13	96.47	102.60
85	AA	1016	G	C5'-C4'-C3'	-6.13	106.19	116.00
85	AA	1303	U	O4'-C1'-N1	6.13	113.11	108.20
85	AA	1874	G	P-O3'-C3'	6.13	127.06	119.70
35	BB	650	A	C4'-C3'-C2'	-6.13	96.47	102.60
35	BB	1335	G	N1-C6-O6	-6.13	116.22	119.90
77	Br	75	ILE	N-CA-C	6.13	127.55	111.00
85	AA	354	C	N3-C2-O2	-6.13	117.61	121.90
85	AA	913	U	O4'-C1'-N1	6.13	113.10	108.20
85	AA	1338	C	O4'-C1'-N1	6.13	113.10	108.20
29	AV	99	ARG	NE-CZ-NH1	6.13	123.36	120.30
34	BA	449	G	C4-C5-C6	-6.13	115.12	118.80
34	BA	511	U	C6-N1-C1'	-6.13	112.62	121.20
39	BF	38	C	N3-C4-C5	-6.13	119.45	121.90
85	AA	762	U	C5'-C4'-C3'	6.13	125.81	116.00
85	AA	787	U	C1'-O4'-C4'	-6.13	105.00	109.90
85	AA	1549	G	C5-N7-C8	-6.13	101.24	104.30
86	AB	10	G	O3'-P-O5'	-6.13	92.36	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	AV	68	MET	N-CA-C	-6.13	94.46	111.00
34	BA	120	A	P-O5'-C5'	6.13	130.70	120.90
34	BA	898	G	C4'-C3'-C2'	-6.13	96.47	102.60
34	BA	1062	G	P-O3'-C3'	6.13	127.05	119.70
34	BA	1329	U	N1-C2-O2	6.13	127.09	122.80
34	BA	1354	G	O4'-C1'-N9	6.13	113.10	108.20
35	BB	419	G	C3'-C2'-C1'	-6.13	96.60	101.50
35	BB	1539	C	N3-C2-O2	-6.13	117.61	121.90
36	BC	103	A	P-O5'-C5'	-6.13	111.10	120.90
38	BE	146	U	O4'-C1'-N1	6.13	113.10	108.20
39	BF	58	U	OP2-P-O3'	6.13	118.68	105.20
40	BG	117	C	O4'-C1'-N1	6.13	113.10	108.20
85	AA	836	A	C5-C6-N1	6.13	120.76	117.70
85	AA	2197	A	O5'-C5'-C4'	6.13	123.34	111.70
85	AA	2237	G	C8-N9-C1'	6.13	134.97	127.00
30	AW	48	TYR	CA-CB-CG	-6.12	101.76	113.40
34	BA	298	G	C4-N9-C1'	6.12	134.46	126.50
34	BA	1784	G	P-O5'-C5'	6.12	130.70	120.90
35	BB	557	C	O4'-C1'-N1	6.12	113.10	108.20
35	BB	814	A	N1-C6-N6	-6.12	114.92	118.60
35	BB	1288	G	C4-N9-C1'	-6.12	118.54	126.50
35	BB	1539	C	N1-C1'-C2'	-6.12	105.26	112.00
37	BD	66	G	N9-C1'-C2'	-6.12	105.26	112.00
80	Bu	244	ASP	C-N-CA	6.12	137.01	121.70
85	AA	266	U	C6-N1-C2	-6.12	117.33	121.00
85	AA	858	G	C6-N1-C2	-6.12	121.42	125.10
25	AR	51	ASP	N-CA-C	-6.12	94.47	111.00
34	BA	535	G	C5-C6-N1	6.12	114.56	111.50
34	BA	1184	A	C6-N1-C2	-6.12	114.93	118.60
34	BA	1673	G	C5-C6-N1	6.12	114.56	111.50
34	BA	1709	A	C1'-O4'-C4'	-6.12	105.00	109.90
34	BA	1737	A	P-O3'-C3'	-6.12	112.35	119.70
35	BB	257	G	N1-C6-O6	6.12	123.57	119.90
35	BB	296	G	O4'-C1'-N9	6.12	113.10	108.20
35	BB	441	G	O4'-C1'-C2'	6.12	113.11	107.60
35	BB	535	U	C2-N1-C1'	-6.12	110.35	117.70
35	BB	1314	G	C3'-C2'-C1'	-6.12	96.60	101.50
38	BE	166	G	O5'-P-OP2	-6.12	100.19	105.70
41	BH	132	C	C5-C6-N1	6.12	124.06	121.00
45	BL	157	PHE	CB-CG-CD2	-6.12	116.51	120.80
84	By	112	ASN	CB-CA-C	-6.12	98.15	110.40
85	AA	559	G	C4'-C3'-C2'	-6.12	96.48	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1154	A	O4'-C1'-N9	6.12	113.10	108.20
85	AA	1236	G	O4'-C4'-C3'	-6.12	97.88	104.00
85	AA	1240	A	C5-C6-N6	-6.12	118.80	123.70
34	BA	504	A	O5'-C5'-C4'	-6.12	100.07	111.70
34	BA	1026	C	N3-C4-C5	6.12	124.35	121.90
34	BA	1167	A	C8-N9-C4	6.12	108.25	105.80
34	BA	1305	A	C6-N1-C2	-6.12	114.93	118.60
35	BB	95	A	O4'-C1'-N9	6.12	113.10	108.20
35	BB	977	G	C4'-C3'-C2'	-6.12	96.48	102.60
35	BB	1283	C	C3'-C2'-C1'	-6.12	96.60	101.50
35	BB	1419	G	C6-N1-C2	-6.12	121.43	125.10
69	Bj	109	LYS	N-CA-CB	-6.12	99.58	110.60
85	AA	189	G	C3'-C2'-C1'	-6.12	96.60	101.50
85	AA	532	G	C4'-C3'-C2'	-6.12	96.48	102.60
85	AA	1670	U	C5-C4-O4	6.12	129.57	125.90
85	AA	1730	C	C6-N1-C2	-6.12	117.85	120.30
85	AA	2099	C	O4'-C1'-N1	6.12	113.10	108.20
19	AK	126	ASP	CA-CB-CG	-6.12	99.94	113.40
34	BA	904	G	C1'-O4'-C4'	-6.12	105.00	109.90
34	BA	1390	C	O4'-C1'-N1	6.12	113.10	108.20
41	BH	57	A	O4'-C1'-N9	6.12	113.10	108.20
85	AA	1784	G	C4'-C3'-C2'	-6.12	96.48	102.60
85	AA	2100	A	C4'-C3'-C2'	6.12	108.72	102.60
34	BA	10	G	C1'-O4'-C4'	-6.12	105.00	109.90
34	BA	170	U	O4'-C1'-N1	6.12	113.09	108.20
34	BA	175	G	C4-N9-C1'	-6.12	118.55	126.50
34	BA	187	G	C4-N9-C1'	-6.12	118.55	126.50
34	BA	241	U	O3'-P-O5'	-6.12	92.38	104.00
34	BA	603	U	O5'-P-OP2	-6.12	100.19	105.70
34	BA	667	U	C2-N1-C1'	-6.12	110.36	117.70
34	BA	691	A	C8-N9-C4	6.12	108.25	105.80
34	BA	1211	G	N3-C2-N2	-6.12	115.62	119.90
34	BA	1668	C	O4'-C1'-N1	6.12	113.09	108.20
34	BA	1675	C	N3-C2-O2	-6.12	117.62	121.90
35	BB	155	G	C5-C6-O6	-6.12	124.93	128.60
38	BE	172	U	N1-C2-O2	6.12	127.08	122.80
41	BH	20	A	O4'-C1'-N9	6.12	113.09	108.20
41	BH	23	G	C1'-O4'-C4'	6.12	114.80	109.90
48	BO	168	TYR	CA-CB-CG	-6.12	101.77	113.40
85	AA	600	C	O5'-P-OP1	-6.12	100.19	105.70
85	AA	703	U	P-O3'-C3'	-6.12	112.36	119.70
85	AA	1848	G	N9-C4-C5	-6.12	102.95	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
5	A4	91	PHE	CB-CG-CD2	-6.12	116.52	120.80
34	BA	1204	U	C2-N1-C1'	-6.12	110.36	117.70
35	BB	811	C	O4'-C1'-N1	6.12	113.09	108.20
37	BD	5	A	O4'-C1'-N9	6.12	113.09	108.20
85	AA	397	G	C5-C6-N1	6.12	114.56	111.50
85	AA	718	C	C6-N1-C1'	6.12	128.14	120.80
85	AA	972	G	O4'-C1'-C2'	6.12	113.11	107.60
34	BA	813	C	N1-C2-O2	6.12	122.57	118.90
34	BA	892	C	C6-N1-C1'	6.12	128.14	120.80
34	BA	916	A	P-O3'-C3'	-6.12	112.36	119.70
34	BA	1639	U	C6-N1-C1'	6.12	129.76	121.20
34	BA	1779	U	O4'-C1'-N1	6.12	113.09	108.20
35	BB	139	G	P-O3'-C3'	-6.12	112.36	119.70
35	BB	788	U	C5-C4-O4	6.12	129.57	125.90
35	BB	1476	C	N3-C2-O2	-6.12	117.62	121.90
38	BE	96	G	C4-C5-N7	-6.12	108.35	110.80
38	BE	194	A	C6-C5-N7	-6.12	128.02	132.30
49	BP	32	ASP	CB-CA-C	6.12	122.63	110.40
67	Bh	128	ASP	N-CA-CB	-6.12	99.59	110.60
85	AA	1174	G	P-O3'-C3'	-6.12	112.36	119.70
85	AA	1313	C	C6-N1-C2	-6.12	117.85	120.30
85	AA	1337	A	C3'-C2'-C1'	-6.12	96.61	101.50
85	AA	1647	G	N3-C4-C5	-6.12	125.54	128.60
85	AA	1731	G	N3-C4-C5	-6.12	125.54	128.60
85	AA	1792	C	C3'-C2'-C1'	6.12	106.39	101.50
86	AB	65	G	C8-N9-C1'	-6.12	119.05	127.00
34	BA	8	G	C3'-C2'-C1'	-6.11	96.61	101.50
34	BA	50	G	C5-C6-N1	6.11	114.56	111.50
34	BA	257	G	C3'-C2'-C1'	-6.11	96.61	101.50
34	BA	261	A	N1-C6-N6	6.11	122.27	118.60
34	BA	418	G	O4'-C1'-C2'	-6.11	99.69	105.80
34	BA	422	C	C2-N3-C4	-6.11	116.84	119.90
34	BA	797	A	C3'-C2'-C1'	6.11	106.39	101.50
34	BA	933	U	C5-C6-N1	-6.11	119.64	122.70
34	BA	1218	G	C5-C6-O6	6.11	132.27	128.60
34	BA	1281	U	N3-C2-O2	-6.11	117.92	122.20
34	BA	1417	C	O4'-C1'-N1	6.11	113.09	108.20
40	BG	28	A	P-O3'-C3'	-6.11	112.36	119.70
41	BH	65	G	C8-N9-C1'	6.11	134.95	127.00
85	AA	601	A	O4'-C4'-C3'	-6.11	97.89	104.00
85	AA	1651	C	N1-C1'-C2'	-6.11	105.28	112.00
24	AQ	101	PHE	CB-CG-CD1	6.11	125.08	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1442	A	N1-C2-N3	6.11	132.36	129.30
85	AA	306	C	P-O5'-C5'	-6.11	111.12	120.90
85	AA	1876	U	P-O3'-C3'	6.11	127.03	119.70
85	AA	1971	G	O4'-C1'-N9	6.11	113.09	108.20
34	BA	469	C	OP2-P-O3'	6.11	118.64	105.20
34	BA	1674	G	P-O5'-C5'	-6.11	111.12	120.90
35	BB	405	U	C3'-C2'-C1'	6.11	106.39	101.50
35	BB	1187	G	P-O5'-C5'	-6.11	111.12	120.90
56	BW	37	TYR	CB-CG-CD1	6.11	124.67	121.00
67	Bh	77	THR	N-CA-CB	6.11	121.91	110.30
77	Br	217	ARG	N-CA-CB	6.11	121.60	110.60
77	Br	250	TRP	CA-CB-CG	-6.11	102.09	113.70
85	AA	852	C	C5-C4-N4	-6.11	115.92	120.20
85	AA	1206	A	C1'-O4'-C4'	-6.11	105.01	109.90
85	AA	1441	G	C5-C6-N1	6.11	114.56	111.50
85	AA	1668	G	C1'-O4'-C4'	-6.11	105.01	109.90
85	AA	1992	A	N7-C8-N9	6.11	116.86	113.80
34	BA	5	C	C2-N3-C4	6.11	122.95	119.90
34	BA	15	G	C5-N7-C8	-6.11	101.25	104.30
34	BA	320	G	N1-C6-O6	-6.11	116.23	119.90
34	BA	894	G	C5-C6-O6	6.11	132.26	128.60
35	BB	3	C	N3-C2-O2	-6.11	117.62	121.90
38	BE	13	A	O4'-C4'-C3'	-6.11	97.89	104.00
41	BH	52	G	C8-N9-C4	-6.11	103.96	106.40
58	BY	75	ARG	NE-CZ-NH1	6.11	123.35	120.30
85	AA	2106	C	C5'-C4'-C3'	-6.11	106.23	116.00
7	A6	140	VAL	N-CA-C	-6.11	94.51	111.00
34	BA	1251	A	O4'-C1'-C2'	6.11	113.10	107.60
34	BA	1295	U	O5'-C5'-C4'	6.11	123.31	111.70
35	BB	472	C	C6-N1-C1'	6.11	128.13	120.80
35	BB	1066	G	N1-C6-O6	6.11	123.56	119.90
35	BB	1304	U	P-O5'-C5'	-6.11	111.13	120.90
35	BB	1428	C	C5'-C4'-C3'	-6.11	106.23	116.00
35	BB	1485	G	C5'-C4'-O4'	6.11	116.43	109.10
36	BC	144	C	C6-N1-C2	-6.11	117.86	120.30
37	BD	85	C	C6-N1-C1'	6.11	128.13	120.80
38	BE	133	C	P-O5'-C5'	-6.11	111.13	120.90
65	Bf	447	ARG	NE-CZ-NH1	6.11	123.35	120.30
85	AA	245	A	C3'-C2'-C1'	-6.11	96.61	101.50
85	AA	867	G	N9-C1'-C2'	-6.11	105.28	112.00
85	AA	1509	A	N7-C8-N9	6.11	116.85	113.80
85	AA	1865	C	P-O3'-C3'	6.11	127.03	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2089	G	O4'-C1'-C2'	6.11	113.10	107.60
85	AA	2215	C	C4'-C3'-C2'	6.11	108.71	102.60
6	A5	78	ILE	N-CA-C	-6.11	94.52	111.00
34	BA	153	C	O4'-C1'-N1	6.11	113.08	108.20
34	BA	521	C	C2-N3-C4	-6.11	116.85	119.90
34	BA	847	U	O4'-C1'-C2'	-6.11	99.69	105.80
35	BB	961	G	C5'-C4'-C3'	-6.11	106.23	116.00
35	BB	1383	C	N3-C4-N4	6.11	122.27	118.00
38	BE	184	G	C4-N9-C1'	-6.11	118.56	126.50
85	AA	369	A	O5'-C5'-C4'	-6.11	100.10	111.70
85	AA	868	A	C5-C6-N6	-6.11	118.81	123.70
85	AA	1595	G	P-O3'-C3'	-6.11	112.37	119.70
85	AA	1979	A	N9-C1'-C2'	-6.11	105.28	112.00
85	AA	2008	G	P-O3'-C3'	-6.11	112.37	119.70
10	A9	152	MET	CG-SD-CE	-6.10	90.43	100.20
34	BA	1379	G	O4'-C1'-N9	6.10	113.08	108.20
35	BB	818	U	C1'-O4'-C4'	-6.10	105.02	109.90
35	BB	1461	C	OP1-P-OP2	-6.10	110.44	119.60
82	Bw	82	ARG	CD-NE-CZ	6.10	132.15	123.60
85	AA	162	A	C5-C6-N6	-6.10	118.82	123.70
85	AA	748	C	C6-N1-C2	-6.10	117.86	120.30
85	AA	1458	G	C5'-C4'-C3'	-6.10	106.23	116.00
7	A6	113	PHE	CB-CG-CD2	-6.10	116.53	120.80
19	AK	68	ARG	NE-CZ-NH2	-6.10	117.25	120.30
34	BA	72	U	N3-C2-O2	-6.10	117.93	122.20
34	BA	371	U	C2-N1-C1'	-6.10	110.38	117.70
34	BA	975	A	C5'-C4'-C3'	-6.10	106.24	116.00
34	BA	1701	U	N1-C2-N3	6.10	118.56	114.90
35	BB	483	C	O5'-C5'-C4'	-6.10	100.11	111.70
35	BB	631	G	C4-N9-C1'	6.10	134.43	126.50
65	Bf	277	THR	N-CA-CB	6.10	121.89	110.30
34	BA	23	A	C5'-C4'-C3'	-6.10	106.24	116.00
35	BB	1040	C	O5'-P-OP2	6.10	118.02	110.70
36	BC	82	C	C2-N1-C1'	6.10	125.51	118.80
38	BE	85	G	C5'-C4'-O4'	-6.10	101.78	109.10
39	BF	52	A	O3'-P-O5'	-6.10	92.41	104.00
47	BN	206	ARG	NE-CZ-NH2	-6.10	117.25	120.30
85	AA	543	A	O4'-C1'-N9	6.10	113.08	108.20
85	AA	660	G	C8-N9-C4	6.10	108.84	106.40
85	AA	663	C	P-O5'-C5'	-6.10	111.14	120.90
85	AA	986	U	O4'-C1'-N1	6.10	113.08	108.20
85	AA	1456	A	C5-C6-N6	-6.10	118.82	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	244	A	P-O3'-C3'	-6.10	112.38	119.70
34	BA	479	U	C5'-C4'-C3'	-6.10	106.24	116.00
34	BA	765	U	C5'-C4'-O4'	6.10	116.42	109.10
34	BA	882	G	P-O5'-C5'	-6.10	111.14	120.90
35	BB	505	G	N9-C1'-C2'	-6.10	105.29	112.00
35	BB	860	U	C6-N1-C1'	-6.10	112.66	121.20
35	BB	1357	C	N1-C2-O2	6.10	122.56	118.90
38	BE	110	U	C4'-C3'-C2'	-6.10	96.50	102.60
39	BF	57	C	N1-C1'-C2'	-6.10	105.29	112.00
40	BG	181	C	C2-N3-C4	-6.10	116.85	119.90
63	Bd	14	ARG	NE-CZ-NH1	6.10	123.35	120.30
85	AA	165	C	C1'-O4'-C4'	-6.10	105.02	109.90
85	AA	169	G	C8-N9-C1'	6.10	134.93	127.00
85	AA	491	G	N1-C2-N2	-6.10	110.71	116.20
85	AA	558	U	P-O5'-C5'	-6.10	111.14	120.90
85	AA	1135	U	C5'-C4'-C3'	-6.10	106.24	116.00
85	AA	2179	C	P-O5'-C5'	6.10	130.66	120.90
85	AA	2182	A	C5-C6-N6	-6.10	118.82	123.70
26	AS	67	ARG	NE-CZ-NH2	-6.10	117.25	120.30
34	BA	600	G	C4-N9-C1'	-6.10	118.57	126.50
34	BA	624	G	C6-N1-C2	-6.10	121.44	125.10
34	BA	841	G	C5'-C4'-C3'	-6.10	106.25	116.00
34	BA	1200	U	C5-C4-O4	-6.10	122.24	125.90
34	BA	1797	A	N1-C2-N3	-6.10	126.25	129.30
35	BB	416	U	O4'-C1'-C2'	6.10	113.09	107.60
35	BB	1269	A	C3'-C2'-C1'	-6.10	96.62	101.50
35	BB	1334	C	P-O3'-C3'	-6.10	112.38	119.70
38	BE	96	G	C8-N9-C1'	6.10	134.93	127.00
41	BH	110	C	C2-N1-C1'	6.10	125.51	118.80
85	AA	524	A	P-O5'-C5'	6.10	130.66	120.90
85	AA	572	G	C1'-O4'-C4'	-6.10	105.02	109.90
85	AA	884	A	P-O3'-C3'	6.10	127.02	119.70
85	AA	1127	G	C4-N9-C1'	-6.10	118.57	126.50
85	AA	1854	U	C6-N1-C1'	-6.10	112.66	121.20
34	BA	337	C	C5'-C4'-C3'	-6.10	106.25	116.00
34	BA	996	U	C5-C6-N1	-6.10	119.65	122.70
34	BA	1071	G	N1-C6-O6	6.10	123.56	119.90
34	BA	1769	U	O4'-C1'-N1	6.10	113.08	108.20
35	BB	631	G	C8-N9-C1'	-6.10	119.08	127.00
85	AA	1453	U	O4'-C1'-N1	6.10	113.08	108.20
7	A6	34	GLY	C-N-CA	6.09	136.94	121.70
34	BA	103	G	C8-N9-C4	6.09	108.84	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	156	U	O4'-C1'-N1	6.09	113.08	108.20
34	BA	539	C	N3-C2-O2	-6.09	117.63	121.90
34	BA	620	C	O4'-C4'-C3'	-6.09	97.91	104.00
34	BA	701	G	C5'-C4'-O4'	6.09	116.41	109.10
34	BA	1012	A	C1'-O4'-C4'	-6.09	105.03	109.90
34	BA	1550	G	C5'-C4'-C3'	6.09	125.75	116.00
34	BA	1737	A	P-O5'-C5'	6.09	130.65	120.90
35	BB	571	C	N1-C1'-C2'	-6.09	105.30	112.00
35	BB	869	G	C5'-C4'-O4'	6.09	116.41	109.10
35	BB	1081	U	C5-C6-N1	-6.09	119.65	122.70
35	BB	1195	A	P-O5'-C5'	6.09	130.65	120.90
35	BB	1270	C	N3-C2-O2	-6.09	117.63	121.90
35	BB	1532	C	N3-C4-C5	6.09	124.34	121.90
36	BC	35	C	O4'-C1'-C2'	6.09	113.08	107.60
37	BD	40	C	O5'-P-OP2	6.09	118.01	110.70
37	BD	68	C	O4'-C1'-N1	6.09	113.08	108.20
38	BE	87	U	C3'-C2'-C1'	-6.09	96.62	101.50
85	AA	1168	C	O4'-C1'-N1	6.09	113.08	108.20
85	AA	1668	G	C8-N9-C4	-6.09	103.96	106.40
85	AA	2190	U	O4'-C1'-N1	6.09	113.08	108.20
86	AB	62	C	O4'-C1'-C2'	6.09	113.08	107.60
34	BA	470	C	C5'-C4'-O4'	6.09	116.41	109.10
34	BA	797	A	N1-C6-N6	-6.09	114.94	118.60
41	BH	100	A	C5-C6-N6	-6.09	118.83	123.70
85	AA	39	A	O4'-C1'-N9	6.09	113.08	108.20
85	AA	804	A	N1-C6-N6	-6.09	114.94	118.60
34	BA	454	G	N1-C6-O6	-6.09	116.25	119.90
34	BA	1505	G	N9-C1'-C2'	6.09	121.92	114.00
34	BA	1579	G	N3-C2-N2	6.09	124.16	119.90
34	BA	1732	A	C1'-O4'-C4'	-6.09	105.03	109.90
34	BA	1749	C	O4'-C1'-N1	6.09	113.07	108.20
34	BA	1823	A	O4'-C4'-C3'	-6.09	97.91	104.00
35	BB	64	U	C5'-C4'-C3'	-6.09	106.25	116.00
35	BB	484	G	C8-N9-C4	-6.09	103.96	106.40
35	BB	531	U	C3'-C2'-C1'	6.09	106.37	101.50
35	BB	965	G	C5-C6-O6	-6.09	124.94	128.60
35	BB	1100	C	N3-C4-N4	-6.09	113.74	118.00
36	BC	35	C	N3-C2-O2	-6.09	117.64	121.90
40	BG	9	G	N1-C2-N2	6.09	121.68	116.20
40	BG	123	C	C2-N1-C1'	-6.09	112.10	118.80
53	BT	132	PHE	CB-CG-CD2	-6.09	116.54	120.80
85	AA	239	G	C2-N3-C4	6.09	114.95	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	380	C	C5'-C4'-C3'	-6.09	106.25	116.00
85	AA	687	G	C8-N9-C1'	-6.09	119.08	127.00
85	AA	932	A	C4'-C3'-C2'	6.09	108.69	102.60
34	BA	124	G	C4'-C3'-C2'	6.09	108.69	102.60
34	BA	159	U	C5-C4-O4	6.09	129.55	125.90
34	BA	295	G	O4'-C1'-N9	6.09	113.07	108.20
34	BA	754	G	N9-C1'-C2'	-6.09	105.30	112.00
34	BA	783	U	C4'-C3'-C2'	6.09	108.69	102.60
34	BA	1211	G	O4'-C4'-C3'	6.09	110.97	106.10
34	BA	1330	G	C1'-O4'-C4'	-6.09	105.03	109.90
35	BB	148	C	N1-C2-O2	6.09	122.55	118.90
35	BB	482	A	O4'-C1'-N9	6.09	113.07	108.20
35	BB	569	G	C5-C6-N1	6.09	114.55	111.50
36	BC	104	A	C4'-C3'-C2'	-6.09	96.51	102.60
44	BK	94	PHE	CB-CA-C	-6.09	98.22	110.40
44	BK	153	ARG	NE-CZ-NH1	6.09	123.34	120.30
69	Bj	106	ARG	NE-CZ-NH1	6.09	123.34	120.30
85	AA	361	U	O4'-C4'-C3'	-6.09	97.91	104.00
85	AA	696	G	N1-C2-N2	6.09	121.68	116.20
85	AA	789	A	O4'-C4'-C3'	-6.09	97.91	104.00
85	AA	871	U	C4'-C3'-C2'	6.09	108.69	102.60
85	AA	879	G	O4'-C1'-N9	6.09	113.07	108.20
85	AA	984	A	C5'-C4'-O4'	6.09	116.41	109.10
85	AA	1241	A	C5-C6-N6	-6.09	118.83	123.70
85	AA	1671	G	C3'-C2'-C1'	-6.09	96.63	101.50
15	AG	50	ILE	CB-CA-C	-6.09	99.42	111.60
34	BA	1273	U	C5-C6-N1	-6.09	119.66	122.70
35	BB	470	C	N3-C2-O2	-6.09	117.64	121.90
35	BB	1051	U	N3-C2-O2	-6.09	117.94	122.20
36	BC	2	A	OP1-P-O3'	6.09	118.59	105.20
41	BH	17	A	O4'-C1'-N9	6.09	113.07	108.20
73	Bn	24	ARG	CA-CB-CG	6.09	126.79	113.40
35	BB	560	C	C2'-C3'-O3'	6.09	123.44	113.70
35	BB	661	G	C3'-C2'-C1'	-6.09	96.63	101.50
35	BB	838	G	O3'-P-O5'	6.09	115.57	104.00
35	BB	879	G	N3-C4-C5	-6.09	125.56	128.60
35	BB	1018	U	C5'-C4'-O4'	6.09	116.40	109.10
35	BB	1202	G	C5-C6-N1	6.09	114.54	111.50
35	BB	1323	U	C5'-C4'-C3'	-6.09	106.26	116.00
35	BB	1434	G	C8-N9-C4	6.09	108.83	106.40
79	Bt	16	ARG	NE-CZ-NH2	-6.09	117.26	120.30
81	Bv	63	SER	N-CA-CB	6.09	119.63	110.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	718	C	C3'-C2'-C1'	-6.09	96.63	101.50
85	AA	853	G	OP1-P-OP2	-6.09	110.47	119.60
85	AA	866	U	N1-C1'-C2'	-6.09	105.31	112.00
85	AA	1220	A	P-O3'-C3'	-6.09	112.40	119.70
85	AA	1481	U	O4'-C1'-N1	6.09	113.07	108.20
27	AT	97	ARG	CG-CD-NE	-6.08	99.02	111.80
34	BA	1107	A	C6-N1-C2	-6.08	114.95	118.60
60	Ba	59	ARG	CB-CA-C	6.08	122.57	110.40
85	AA	232	U	C4'-C3'-C2'	-6.08	96.52	102.60
85	AA	2219	G	P-O5'-C5'	-6.08	111.16	120.90
34	BA	651	U	C1'-O4'-C4'	-6.08	105.03	109.90
34	BA	688	G	C8-N9-C1'	6.08	134.91	127.00
34	BA	1663	U	C2-N3-C4	-6.08	123.35	127.00
35	BB	79	U	C1'-O4'-C4'	-6.08	105.03	109.90
35	BB	472	C	O4'-C1'-C2'	6.08	113.08	107.60
35	BB	662	G	C5-C6-N1	6.08	114.54	111.50
35	BB	1044	U	C6-N1-C2	-6.08	117.35	121.00
37	BD	57	C	C3'-C2'-C1'	-6.08	96.63	101.50
39	BF	15	U	C3'-C2'-C1'	-6.08	96.63	101.50
39	BF	44	C	C6-N1-C2	-6.08	117.87	120.30
39	BF	50	C	C6-N1-C2	-6.08	117.87	120.30
41	BH	24	U	O4'-C1'-C2'	-6.08	99.72	105.80
41	BH	106	G	C5-C6-N1	6.08	114.54	111.50
41	BH	122	U	C2-N1-C1'	6.08	125.00	117.70
62	Bc	88	SER	N-CA-C	6.08	127.42	111.00
63	Bd	14	ARG	N-CA-CB	-6.08	99.65	110.60
65	Bf	112	ASP	CA-CB-CG	-6.08	100.01	113.40
85	AA	166	C	O4'-C1'-N1	6.08	113.07	108.20
85	AA	597	A	C5'-C4'-O4'	6.08	116.40	109.10
85	AA	994	A	O4'-C1'-N9	6.08	113.07	108.20
85	AA	1809	G	P-O5'-C5'	6.08	130.63	120.90
85	AA	1955	U	O3'-P-O5'	6.08	115.56	104.00
19	AK	137	ARG	CD-NE-CZ	-6.08	115.09	123.60
34	BA	141	G	C5-N7-C8	-6.08	101.26	104.30
34	BA	482	C	C1'-O4'-C4'	6.08	114.77	109.90
34	BA	793	A	C4'-C3'-C2'	-6.08	96.52	102.60
34	BA	993	C	P-O3'-C3'	6.08	127.00	119.70
34	BA	1606	A	C5'-C4'-C3'	-6.08	106.27	116.00
35	BB	367	C	C1'-O4'-C4'	-6.08	105.03	109.90
35	BB	992	C	C6-N1-C2	-6.08	117.87	120.30
35	BB	1420	U	P-O3'-C3'	-6.08	112.40	119.70
65	Bf	167	ARG	NE-CZ-NH2	-6.08	117.26	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	42	G	O4'-C1'-N9	6.08	113.06	108.20
85	AA	178	U	C1'-O4'-C4'	-6.08	105.03	109.90
85	AA	600	C	P-O5'-C5'	6.08	130.63	120.90
85	AA	742	U	O5'-C5'-C4'	-6.08	100.14	111.70
85	AA	1257	A	C8-N9-C4	6.08	108.23	105.80
85	AA	1526	G	N7-C8-N9	6.08	116.14	113.10
85	AA	2008	G	N1-C2-N2	6.08	121.67	116.20
34	BA	1638	U	C2'-C3'-O3'	6.08	123.43	113.70
34	BA	1677	C	P-O3'-C3'	-6.08	112.40	119.70
35	BB	550	G	C4-N9-C1'	-6.08	118.59	126.50
35	BB	1097	U	C3'-C2'-C1'	-6.08	96.64	101.50
35	BB	1339	C	C5'-C4'-C3'	-6.08	106.27	116.00
55	BV	121	TYR	CB-CG-CD1	6.08	124.65	121.00
85	AA	479	C	C6-N1-C2	-6.08	117.87	120.30
85	AA	2005	U	P-O5'-C5'	-6.08	111.17	120.90
21	AM	15	ARG	NE-CZ-NH1	6.08	123.34	120.30
34	BA	952	G	N9-C1'-C2'	-6.08	105.31	112.00
34	BA	992	A	N1-C6-N6	-6.08	114.95	118.60
35	BB	743	C	N3-C4-N4	6.08	122.26	118.00
35	BB	1418	C	P-O5'-C5'	-6.08	111.17	120.90
35	BB	1530	U	N3-C2-O2	-6.08	117.95	122.20
38	BE	101	C	O4'-C1'-N1	6.08	113.06	108.20
40	BG	56	G	C5'-C4'-C3'	-6.08	106.28	116.00
71	Bl	62	ARG	NE-CZ-NH2	-6.08	117.26	120.30
75	Bp	36	VAL	CB-CA-C	-6.08	99.85	111.40
85	AA	130	G	C4-N9-C1'	-6.08	118.60	126.50
85	AA	1281	G	N7-C8-N9	-6.08	110.06	113.10
85	AA	1366	A	C4'-C3'-C2'	-6.08	96.52	102.60
85	AA	1610	G	C5'-C4'-C3'	-6.08	106.27	116.00
85	AA	2102	A	C8-N9-C4	6.08	108.23	105.80
85	AA	2118	U	O3'-P-O5'	6.08	115.55	104.00
86	AB	8	U	O3'-P-O5'	-6.08	92.45	104.00
34	BA	570	G	C5'-C4'-C3'	-6.08	106.28	116.00
34	BA	807	U	C3'-C2'-C1'	-6.08	96.64	101.50
34	BA	1085	G	C4'-C3'-C2'	-6.08	96.52	102.60
35	BB	846	A	C4'-C3'-C2'	-6.08	96.52	102.60
35	BB	1462	G	C5'-C4'-C3'	6.08	125.72	116.00
38	BE	177	U	C5-C4-O4	6.08	129.55	125.90
41	BH	45	G	N9-C1'-C2'	-6.08	105.32	112.00
85	AA	212	G	O4'-C1'-N9	6.08	113.06	108.20
85	AA	2161	C	C6-N1-C1'	6.08	128.09	120.80
34	BA	18	G	C8-N9-C1'	6.08	134.90	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	514	U	C4'-C3'-C2'	6.08	108.67	102.60
34	BA	983	A	C5-C6-N6	-6.08	118.84	123.70
34	BA	1135	U	C6-N1-C1'	6.08	129.71	121.20
35	BB	103	C	P-O3'-C3'	-6.08	112.41	119.70
35	BB	540	G	P-O5'-C5'	-6.08	111.18	120.90
35	BB	1408	G	N1-C6-O6	6.08	123.55	119.90
35	BB	1458	U	OP2-P-O3'	6.08	118.57	105.20
41	BH	114	G	O4'-C1'-C2'	6.08	113.07	107.60
44	BK	139	ARG	NE-CZ-NH2	-6.08	117.26	120.30
54	BU	7	TYR	CA-CB-CG	6.08	124.94	113.40
85	AA	70	U	O4'-C1'-N1	6.08	113.06	108.20
85	AA	245	A	C8-N9-C1'	6.08	138.64	127.70
85	AA	258	G	C5-C6-O6	-6.08	124.95	128.60
85	AA	384	C	C5'-C4'-C3'	-6.08	106.28	116.00
85	AA	1548	A	C3'-C2'-C1'	-6.08	96.64	101.50
5	A4	190	PHE	CB-CG-CD2	-6.07	116.55	120.80
34	BA	300	C	N3-C4-N4	-6.07	113.75	118.00
34	BA	423	G	P-O5'-C5'	-6.07	111.18	120.90
34	BA	470	C	N3-C4-N4	-6.07	113.75	118.00
34	BA	1551	G	O4'-C1'-N9	6.07	113.06	108.20
34	BA	1607	U	C3'-C2'-C1'	-6.07	96.64	101.50
35	BB	378	C	C5'-C4'-O4'	6.07	116.39	109.10
35	BB	1199	A	C8-N9-C4	6.07	108.23	105.80
35	BB	1385	C	C3'-C2'-C1'	-6.07	96.64	101.50
35	BB	1393	C	O3'-P-O5'	-6.07	92.46	104.00
37	BD	11	A	C4-C5-C6	-6.07	113.96	117.00
37	BD	85	C	C4'-C3'-C2'	6.07	108.67	102.60
38	BE	32	U	C4'-C3'-O3'	-6.07	96.65	109.40
42	BI	106	ARG	NE-CZ-NH1	6.07	123.34	120.30
60	Ba	84	TYR	CB-CG-CD2	-6.07	117.36	121.00
69	Bj	114	VAL	CA-C-N	6.07	130.56	117.20
75	Bp	25	VAL	CA-CB-CG2	-6.07	101.79	110.90
85	AA	473	C	O3'-P-O5'	6.07	115.54	104.00
85	AA	1288	A	C1'-O4'-C4'	-6.07	105.04	109.90
85	AA	1488	G	P-O5'-C5'	-6.07	111.18	120.90
85	AA	1803	U	C5'-C4'-O4'	6.07	116.39	109.10
85	AA	1897	A	C5-C6-N1	6.07	120.74	117.70
86	AB	15	G	P-O3'-C3'	6.07	126.99	119.70
5	A4	102	ARG	NE-CZ-NH1	6.07	123.34	120.30
25	AR	62	TYR	CA-CB-CG	-6.07	101.86	113.40
34	BA	1499	A	C5-C6-N1	6.07	120.74	117.70
82	Bw	192	ASP	CA-CB-CG	-6.07	100.04	113.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	866	U	O4'-C4'-C3'	-6.07	97.93	104.00
85	AA	1140	G	O5'-C5'-C4'	6.07	123.24	111.70
85	AA	1835	U	C2-N3-C4	-6.07	123.36	127.00
6	A5	160	ARG	NE-CZ-NH1	6.07	123.33	120.30
24	AQ	64	LYS	C-N-CA	6.07	136.87	121.70
34	BA	473	A	P-O3'-C3'	-6.07	112.42	119.70
34	BA	661	C	C3'-C2'-C1'	-6.07	96.64	101.50
34	BA	882	G	C6-N1-C2	-6.07	121.46	125.10
34	BA	1127	U	C5-C4-O4	-6.07	122.26	125.90
35	BB	7	C	C2-N1-C1'	-6.07	112.12	118.80
35	BB	505	G	C5-C6-N1	6.07	114.54	111.50
35	BB	619	A	C4-N9-C1'	-6.07	115.37	126.30
35	BB	857	G	C8-N9-C1'	6.07	134.89	127.00
35	BB	858	U	C3'-C2'-C1'	-6.07	96.64	101.50
35	BB	1003	G	C8-N9-C4	-6.07	103.97	106.40
35	BB	1157	G	C8-N9-C1'	6.07	134.89	127.00
35	BB	1387	C	C6-N1-C2	-6.07	117.87	120.30
39	BF	30	C	C6-N1-C2	6.07	122.73	120.30
40	BG	15	G	C3'-C2'-C1'	-6.07	96.64	101.50
40	BG	98	A	C1'-O4'-C4'	-6.07	105.04	109.90
85	AA	83	U	O5'-P-OP1	6.07	117.98	110.70
85	AA	157	G	P-O5'-C5'	-6.07	111.19	120.90
85	AA	349	C	P-O3'-C3'	-6.07	112.42	119.70
85	AA	1728	G	O4'-C1'-N9	6.07	113.06	108.20
85	AA	1907	U	O4'-C1'-N1	6.07	113.06	108.20
85	AA	1937	G	C8-N9-C1'	6.07	134.89	127.00
85	AA	1983	C	O4'-C1'-N1	6.07	113.06	108.20
85	AA	2061	C	P-O5'-C5'	-6.07	111.19	120.90
34	BA	862	C	N3-C4-C5	6.07	124.33	121.90
34	BA	1344	G	N3-C2-N2	6.07	124.15	119.90
34	BA	1401	C	C6-N1-C2	-6.07	117.87	120.30
34	BA	1803	A	N9-C1'-C2'	-6.07	105.32	112.00
35	BB	694	C	C5-C4-N4	6.07	124.45	120.20
36	BC	67	U	C6-N1-C1'	6.07	129.70	121.20
39	BF	5	U	C1'-O4'-C4'	-6.07	105.05	109.90
85	AA	224	C	O4'-C1'-N1	6.07	113.06	108.20
85	AA	400	G	OP1-P-OP2	-6.07	110.50	119.60
85	AA	1201	A	N9-C1'-C2'	-6.07	105.32	112.00
85	AA	1897	A	C8-N9-C4	-6.07	103.37	105.80
34	BA	397	A	C8-N9-C4	6.07	108.23	105.80
34	BA	687	G	N3-C4-C5	-6.07	125.57	128.60
34	BA	700	G	C6-N1-C2	-6.07	121.46	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	891	C	O4'-C1'-N1	6.07	113.05	108.20
34	BA	1099	U	C1'-O4'-C4'	-6.07	105.05	109.90
34	BA	1738	G	N9-C1'-C2'	-6.07	105.33	112.00
35	BB	818	U	C5'-C4'-C3'	-6.07	106.29	116.00
38	BE	193	A	N1-C6-N6	6.07	122.24	118.60
40	BG	105	A	C8-N9-C4	6.07	108.23	105.80
41	BH	127	A	C5'-C4'-C3'	6.07	125.71	116.00
48	BO	136	ARG	NH1-CZ-NH2	-6.07	112.73	119.40
63	Bd	61	GLU	CG-CD-OE2	-6.07	106.17	118.30
85	AA	158	C	O4'-C1'-C2'	6.07	113.06	107.60
85	AA	196	U	C4'-C3'-C2'	-6.07	96.53	102.60
85	AA	456	A	N7-C8-N9	-6.07	110.77	113.80
85	AA	666	A	P-O3'-C3'	-6.07	112.42	119.70
85	AA	1010	U	C2-N3-C4	-6.07	123.36	127.00
85	AA	1237	A	O4'-C4'-C3'	-6.07	97.93	104.00
85	AA	1961	U	C5'-C4'-C3'	-6.07	106.29	116.00
85	AA	2108	C	P-O3'-C3'	-6.07	112.42	119.70
85	AA	2112	G	C1'-O4'-C4'	-6.07	105.05	109.90
20	AL	5	ARG	NE-CZ-NH1	6.07	123.33	120.30
32	AY	3	LYS	C-N-CA	6.07	136.86	121.70
34	BA	53	G	O4'-C1'-C2'	6.07	113.06	107.60
34	BA	608	G	C3'-C2'-C1'	6.07	106.35	101.50
34	BA	958	G	P-O3'-C3'	6.07	126.98	119.70
34	BA	1309	U	O5'-C5'-C4'	6.07	123.23	111.70
34	BA	1384	G	C5-C6-O6	-6.07	124.96	128.60
35	BB	391	G	O5'-P-OP2	-6.07	100.24	105.70
35	BB	670	G	P-O5'-C5'	6.07	130.60	120.90
35	BB	691	A	P-O3'-C3'	-6.07	112.42	119.70
35	BB	812	G	C5-C6-N1	6.07	114.53	111.50
35	BB	851	U	C4'-C3'-C2'	-6.07	96.53	102.60
35	BB	995	C	C5'-C4'-O4'	6.07	116.38	109.10
35	BB	1077	C	C5-C4-N4	6.07	124.44	120.20
39	BF	16	C	C6-N1-C1'	-6.07	113.52	120.80
67	Bh	66	ARG	NE-CZ-NH1	6.07	123.33	120.30
75	Bp	63	SER	N-CA-CB	-6.07	101.40	110.50
77	Br	27	PHE	CB-CG-CD2	-6.07	116.56	120.80
77	Br	184	ASP	CB-CG-OD1	6.07	123.76	118.30
85	AA	838	G	N9-C1'-C2'	-6.07	105.33	112.00
85	AA	2198	G	C4'-C3'-C2'	-6.07	96.53	102.60
86	AB	42	C	C5'-C4'-C3'	-6.07	106.29	116.00
86	AB	67	C	P-O5'-C5'	-6.07	111.20	120.90
34	BA	1820	G	O4'-C1'-C2'	6.06	113.06	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	BD	116	C	C6-N1-C2	-6.06	117.88	120.30
85	AA	114	C	C4'-C3'-C2'	-6.06	96.54	102.60
85	AA	341	C	O5'-C5'-C4'	-6.06	100.18	111.70
85	AA	673	A	C5-C6-N6	6.06	128.55	123.70
18	AJ	76	CYS	CB-CA-C	-6.06	98.28	110.40
23	AP	261	ARG	N-CA-C	-6.06	94.63	111.00
34	BA	90	G	P-O3'-C3'	6.06	126.97	119.70
34	BA	581	U	N1-C1'-C2'	6.06	121.88	114.00
35	BB	22	A	C4-N9-C1'	-6.06	115.39	126.30
35	BB	73	G	P-O5'-C5'	-6.06	111.20	120.90
35	BB	79	U	C2-N1-C1'	-6.06	110.42	117.70
35	BB	493	U	N3-C2-O2	-6.06	117.96	122.20
38	BE	30	C	C4-C5-C6	-6.06	114.37	117.40
38	BE	48	G	N3-C4-C5	-6.06	125.57	128.60
38	BE	118	C	P-O3'-C3'	6.06	126.97	119.70
39	BF	48	G	C5'-C4'-O4'	6.06	116.37	109.10
85	AA	23	G	C8-N9-C4	-6.06	103.97	106.40
85	AA	357	C	C3'-C2'-C1'	6.06	106.35	101.50
85	AA	571	G	O4'-C4'-C3'	-6.06	97.94	104.00
85	AA	713	G	C8-N9-C1'	-6.06	119.12	127.00
85	AA	723	U	O5'-C5'-C4'	-6.06	100.18	111.70
85	AA	781	G	N9-C1'-C2'	-6.06	105.33	112.00
85	AA	960	G	N1-C6-O6	6.06	123.54	119.90
85	AA	975	G	N1-C2-N2	-6.06	110.74	116.20
85	AA	1398	U	O4'-C1'-N1	6.06	113.05	108.20
85	AA	1558	U	C4'-C3'-C2'	6.06	108.66	102.60
85	AA	2039	G	C5'-C4'-C3'	-6.06	106.30	116.00
34	BA	216	C	C2-N1-C1'	-6.06	112.13	118.80
34	BA	1108	U	C3'-C2'-C1'	-6.06	96.65	101.50
35	BB	730	G	P-O3'-C3'	-6.06	112.43	119.70
36	BC	69	U	C2-N3-C4	-6.06	123.36	127.00
38	BE	112	G	C3'-C2'-C1'	-6.06	96.65	101.50
85	AA	1983	C	OP1-P-OP2	-6.06	110.51	119.60
85	AA	2083	G	C1'-O4'-C4'	6.06	114.75	109.90
34	BA	28	C	N3-C2-O2	-6.06	117.66	121.90
34	BA	299	C	C2-N3-C4	-6.06	116.87	119.90
34	BA	586	G	C5-C6-O6	-6.06	124.96	128.60
34	BA	769	U	C2'-C3'-O3'	6.06	123.39	113.70
34	BA	815	C	O4'-C4'-C3'	-6.06	97.94	104.00
34	BA	1191	C	N1-C1'-C2'	-6.06	105.33	112.00
35	BB	392	G	N3-C2-N2	6.06	124.14	119.90
35	BB	395	U	P-O3'-C3'	-6.06	112.43	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	783	U	O4'-C1'-N1	6.06	113.05	108.20
35	BB	1213	U	C6-N1-C2	-6.06	117.36	121.00
37	BD	61	C	O4'-C1'-N1	6.06	113.05	108.20
38	BE	195	G	C8-N9-C4	-6.06	103.98	106.40
82	Bw	172	ASN	N-CA-CB	6.06	121.51	110.60
85	AA	77	C	C4'-C3'-C2'	-6.06	96.54	102.60
85	AA	189	G	C5'-C4'-C3'	-6.06	106.30	116.00
85	AA	268	A	O4'-C4'-C3'	-6.06	97.94	104.00
85	AA	569	A	N9-C4-C5	6.06	108.22	105.80
85	AA	572	G	C5'-C4'-C3'	-6.06	106.31	116.00
85	AA	1825	A	O4'-C1'-N9	6.06	113.05	108.20
33	AZ	101	ARG	NE-CZ-NH1	6.06	123.33	120.30
34	BA	1220	C	N3-C4-C5	6.06	124.32	121.90
34	BA	1404	A	C5'-C4'-C3'	-6.06	106.31	116.00
35	BB	112	G	C2-N3-C4	6.06	114.93	111.90
35	BB	375	G	C8-N9-C4	6.06	108.82	106.40
35	BB	542	A	C5-C6-N6	-6.06	118.85	123.70
35	BB	577	U	C5-C6-N1	-6.06	119.67	122.70
35	BB	696	G	C3'-C2'-C1'	-6.06	96.65	101.50
35	BB	904	C	C5'-C4'-C3'	6.06	125.69	116.00
65	Bf	288	ARG	NE-CZ-NH1	6.06	123.33	120.30
85	AA	200	U	P-O3'-C3'	-6.06	112.43	119.70
85	AA	535	G	P-O3'-C3'	-6.06	112.43	119.70
85	AA	574	U	C6-N1-C2	-6.06	117.37	121.00
85	AA	1125	G	N7-C8-N9	-6.06	110.07	113.10
85	AA	1899	A	C5-C6-N6	-6.06	118.86	123.70
85	AA	2001	C	N1-C1'-C2'	6.06	121.88	114.00
34	BA	108	A	N9-C1'-C2'	-6.06	105.34	112.00
35	BB	855	G	C4-N9-C1'	-6.06	118.63	126.50
35	BB	1473	U	C2-N3-C4	-6.06	123.37	127.00
80	Bu	251	ARG	NE-CZ-NH1	6.06	123.33	120.30
85	AA	201	U	C5'-C4'-C3'	-6.06	106.31	116.00
85	AA	766	G	O4'-C1'-N9	6.06	113.05	108.20
85	AA	829	C	P-O3'-C3'	6.06	126.97	119.70
85	AA	1943	U	P-O3'-C3'	6.06	126.97	119.70
34	BA	813	C	N1-C2-N3	6.05	123.44	119.20
34	BA	933	U	C6-N1-C1'	6.05	129.68	121.20
34	BA	1101	A	C5'-C4'-C3'	-6.05	106.31	116.00
34	BA	1300	G	C6-N1-C2	-6.05	121.47	125.10
35	BB	1098	G	C4-N9-C1'	-6.05	118.63	126.50
37	BD	35	C	O5'-C5'-C4'	-6.05	100.19	111.70
85	AA	474	C	C6-N1-C2	-6.05	117.88	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	655	U	C5'-C4'-C3'	-6.05	106.31	116.00
85	AA	787	U	OP2-P-O3'	6.05	118.52	105.20
85	AA	1094	G	C4-N9-C1'	-6.05	118.63	126.50
85	AA	1156	A	O4'-C1'-N9	6.05	113.04	108.20
85	AA	1311	U	P-O3'-C3'	-6.05	112.44	119.70
85	AA	1967	A	C4'-C3'-C2'	6.05	108.66	102.60
34	BA	445	C	O4'-C1'-N1	6.05	113.04	108.20
34	BA	1353	U	C2-N1-C1'	6.05	124.96	117.70
34	BA	1571	C	C2-N3-C4	-6.05	116.87	119.90
35	BB	835	C	C5'-C4'-C3'	-6.05	106.32	116.00
35	BB	1123	A	N1-C6-N6	-6.05	114.97	118.60
35	BB	1482	A	C5'-C4'-O4'	6.05	116.36	109.10
85	AA	69	C	C1'-O4'-C4'	-6.05	105.06	109.90
34	BA	1	C	N3-C4-C5	-6.05	119.48	121.90
34	BA	458	G	C1'-O4'-C4'	-6.05	105.06	109.90
34	BA	818	G	C1'-O4'-C4'	-6.05	105.06	109.90
34	BA	1566	G	C4-N9-C1'	-6.05	118.63	126.50
34	BA	1696	G	N3-C2-N2	6.05	124.14	119.90
34	BA	1747	C	O4'-C1'-N1	6.05	113.04	108.20
35	BB	1000	U	C3'-C2'-C1'	-6.05	96.66	101.50
35	BB	1517	G	C2-N3-C4	-6.05	108.87	111.90
37	BD	91	U	C6-N1-C1'	6.05	129.67	121.20
41	BH	98	U	O4'-C1'-N1	6.05	113.04	108.20
68	Bi	46	ARG	NE-CZ-NH1	6.05	123.33	120.30
82	Bw	42	LYS	N-CA-C	-6.05	94.66	111.00
85	AA	76	G	C4-N9-C1'	-6.05	118.63	126.50
85	AA	1094	G	C4'-C3'-C2'	-6.05	96.55	102.60
85	AA	1277	C	C1'-O4'-C4'	-6.05	105.06	109.90
85	AA	2093	U	P-O3'-C3'	-6.05	112.44	119.70
85	AA	2229	G	C8-N9-C4	-6.05	103.98	106.40
34	BA	1472	G	C4'-C3'-C2'	6.05	108.65	102.60
34	BA	1516	G	C5'-C4'-O4'	6.05	116.36	109.10
34	BA	1727	A	N1-C6-N6	-6.05	114.97	118.60
35	BB	610	U	N1-C2-N3	6.05	118.53	114.90
35	BB	1400	C	C4'-C3'-C2'	6.05	108.65	102.60
36	BC	151	G	C8-N9-C1'	6.05	134.87	127.00
37	BD	18	G	C5-C6-O6	-6.05	124.97	128.60
40	BG	26	G	C4-N9-C1'	-6.05	118.64	126.50
41	BH	86	G	O4'-C1'-N9	6.05	113.04	108.20
59	BZ	37	ARG	NE-CZ-NH2	-6.05	117.28	120.30
83	Bx	47	PHE	N-CA-CB	-6.05	99.71	110.60
85	AA	257	U	P-O5'-C5'	-6.05	111.22	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	355	G	P-O5'-C5'	-6.05	111.22	120.90
85	AA	2030	U	C5'-C4'-C3'	-6.05	106.32	116.00
7	A6	33	TYR	CA-CB-CG	-6.05	101.91	113.40
34	BA	180	G	C4-N9-C1'	6.05	134.36	126.50
34	BA	188	C	C4'-C3'-C2'	-6.05	96.55	102.60
34	BA	541	C	N3-C2-O2	-6.05	117.67	121.90
35	BB	139	G	C8-N9-C1'	6.05	134.86	127.00
35	BB	428	G	C6-N1-C2	-6.05	121.47	125.10
35	BB	508	U	C6-N1-C2	-6.05	117.37	121.00
35	BB	626	C	O4'-C1'-N1	6.05	113.04	108.20
85	AA	1579	A	O4'-C1'-N9	6.05	113.04	108.20
85	AA	2150	G	P-O5'-C5'	6.05	130.58	120.90
3	A2	72	LYS	N-CA-CB	-6.05	99.72	110.60
34	BA	721	A	C6-N1-C2	-6.05	114.97	118.60
34	BA	1330	G	C8-N9-C1'	6.05	134.86	127.00
35	BB	164	U	O4'-C1'-N1	6.05	113.04	108.20
37	BD	110	G	C3'-C2'-C1'	-6.05	96.66	101.50
40	BG	43	U	C3'-C2'-C1'	-6.05	96.66	101.50
85	AA	314	C	C5-C6-N1	6.05	124.02	121.00
85	AA	702	G	C4-N9-C1'	-6.05	118.64	126.50
85	AA	739	C	C3'-C2'-C1'	-6.05	96.66	101.50
85	AA	767	A	O3'-P-O5'	-6.05	92.51	104.00
85	AA	1302	A	C5-C6-N6	-6.05	118.86	123.70
34	BA	254	U	N1-C2-N3	-6.04	111.27	114.90
34	BA	539	C	P-O3'-C3'	-6.04	112.45	119.70
34	BA	1396	A	C4'-C3'-C2'	-6.04	96.56	102.60
34	BA	1478	G	P-O5'-C5'	6.04	130.57	120.90
34	BA	1532	G	N1-C6-O6	-6.04	116.27	119.90
34	BA	1697	U	C6-N1-C2	-6.04	117.37	121.00
35	BB	445	G	O4'-C1'-N9	6.04	113.04	108.20
35	BB	1529	G	O4'-C1'-N9	6.04	113.04	108.20
40	BG	176	G	O5'-C5'-C4'	-6.04	100.21	111.70
85	AA	410	A	N1-C6-N6	-6.04	114.97	118.60
85	AA	633	C	N1-C2-O2	6.04	122.53	118.90
85	AA	1018	G	P-O3'-C3'	-6.04	112.45	119.70
85	AA	1635	C	O4'-C1'-N1	6.04	113.04	108.20
35	BB	718	G	O4'-C1'-N9	6.04	113.03	108.20
35	BB	1064	U	C2-N1-C1'	6.04	124.95	117.70
35	BB	1071	G	C4-N9-C1'	-6.04	118.64	126.50
38	BE	150	G	C5'-C4'-C3'	-6.04	106.33	116.00
64	Be	156	LYS	N-CA-C	-6.04	94.68	111.00
85	AA	586	G	O4'-C1'-N9	6.04	113.03	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	730	G	P-O3'-C3'	-6.04	112.45	119.70
85	AA	1129	A	P-O3'-C3'	6.04	126.95	119.70
85	AA	1375	U	P-O5'-C5'	-6.04	111.23	120.90
85	AA	1444	U	C2-N1-C1'	-6.04	110.45	117.70
85	AA	1934	A	N1-C6-N6	6.04	122.23	118.60
86	AB	48	C	N3-C2-O2	-6.04	117.67	121.90
7	A6	176	ARG	CB-CA-C	-6.04	98.32	110.40
34	BA	589	A	O3'-P-O5'	-6.04	92.52	104.00
34	BA	1215	U	C5'-C4'-C3'	-6.04	106.33	116.00
34	BA	1579	G	N3-C4-C5	-6.04	125.58	128.60
35	BB	371	C	P-O3'-C3'	6.04	126.95	119.70
35	BB	573	C	O4'-C1'-N1	6.04	113.03	108.20
35	BB	722	U	P-O5'-C5'	6.04	130.57	120.90
35	BB	1148	U	O5'-C5'-C4'	-6.04	100.22	111.70
35	BB	1179	C	O4'-C1'-N1	6.04	113.03	108.20
85	AA	159	G	C4'-C3'-C2'	6.04	108.64	102.60
85	AA	244	G	C4-N9-C1'	-6.04	118.65	126.50
85	AA	366	A	C5'-C4'-C3'	-6.04	106.33	116.00
85	AA	719	C	N3-C2-O2	-6.04	117.67	121.90
85	AA	1917	G	P-O5'-C5'	-6.04	111.23	120.90
34	BA	909	G	C4-N9-C1'	-6.04	118.65	126.50
34	BA	1333	G	C1'-O4'-C4'	-6.04	105.07	109.90
36	BC	145	G	C5'-C4'-O4'	6.04	116.35	109.10
38	BE	17	U	P-O3'-C3'	6.04	126.95	119.70
49	BP	8	ARG	CD-NE-CZ	-6.04	115.14	123.60
57	BX	156	ASP	CB-CG-OD2	6.04	123.74	118.30
85	AA	318	A	N1-C6-N6	-6.04	114.98	118.60
85	AA	367	A	N9-C4-C5	6.04	108.22	105.80
85	AA	1483	A	N7-C8-N9	-6.04	110.78	113.80
5	A4	127	PHE	CB-CG-CD2	-6.04	116.57	120.80
34	BA	791	A	C3'-C2'-C1'	-6.04	96.67	101.50
34	BA	934	G	N3-C2-N2	-6.04	115.67	119.90
34	BA	1609	U	N1-C2-N3	6.04	118.52	114.90
34	BA	1675	C	C3'-C2'-C1'	-6.04	96.67	101.50
35	BB	264	U	O4'-C1'-N1	6.04	113.03	108.20
35	BB	824	C	P-O5'-C5'	-6.04	111.24	120.90
35	BB	1484	A	P-O5'-C5'	6.04	130.56	120.90
37	BD	69	U	C5-C4-O4	-6.04	122.28	125.90
40	BG	106	G	O4'-C1'-C2'	-6.04	99.76	105.80
41	BH	111	U	C4'-C3'-C2'	-6.04	96.56	102.60
41	BH	118	U	C2'-C3'-O3'	6.04	123.36	113.70
41	BH	135	U	N1-C2-N3	6.04	118.52	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1154	A	C3'-C2'-C1'	6.04	106.33	101.50
85	AA	2002	A	C8-N9-C1'	-6.04	116.83	127.70
85	AA	2168	C	C1'-O4'-C4'	-6.04	105.07	109.90
12	AD	81	ASN	CA-CB-CG	-6.04	100.12	113.40
34	BA	928	C	O4'-C1'-N1	6.04	113.03	108.20
38	BE	105	A	O5'-P-OP2	-6.04	100.27	105.70
40	BG	75	C	C5-C4-N4	-6.04	115.97	120.20
85	AA	250	C	C5'-C4'-C3'	6.04	125.66	116.00
85	AA	396	U	N3-C2-O2	-6.04	117.97	122.20
85	AA	1027	U	O4'-C1'-N1	6.04	113.03	108.20
85	AA	2042	G	C4-N9-C1'	-6.04	118.65	126.50
34	BA	183	G	N3-C2-N2	6.04	124.12	119.90
34	BA	1101	A	N1-C6-N6	-6.04	114.98	118.60
34	BA	1732	A	P-O3'-C3'	-6.04	112.46	119.70
35	BB	33	A	C8-N9-C4	6.04	108.21	105.80
35	BB	109	U	C2-N1-C1'	-6.04	110.46	117.70
35	BB	540	G	C2-N3-C4	-6.04	108.88	111.90
35	BB	1029	U	C4'-C3'-C2'	6.04	108.64	102.60
37	BD	29	C	P-O3'-C3'	-6.04	112.46	119.70
40	BG	152	G	C5-C6-O6	6.04	132.22	128.60
85	AA	17	C	C4'-C3'-C2'	-6.04	96.56	102.60
85	AA	478	U	C5'-C4'-O4'	6.04	116.34	109.10
85	AA	620	U	C3'-C2'-C1'	-6.04	96.67	101.50
85	AA	1684	U	O4'-C4'-C3'	-6.04	97.96	104.00
34	BA	10	G	C3'-C2'-C1'	-6.03	96.67	101.50
34	BA	89	G	OP2-P-O3'	6.03	118.47	105.20
34	BA	118	C	C5-C4-N4	6.03	124.42	120.20
34	BA	248	G	C3'-C2'-C1'	-6.03	96.67	101.50
34	BA	759	A	C5-N7-C8	-6.03	100.88	103.90
35	BB	154	A	N1-C6-N6	6.03	122.22	118.60
35	BB	473	U	C5'-C4'-C3'	-6.03	106.34	116.00
35	BB	514	G	C4-N9-C1'	-6.03	118.66	126.50
35	BB	1202	G	N3-C4-C5	-6.03	125.58	128.60
54	BU	77	ASN	CA-CB-CG	-6.03	100.13	113.40
85	AA	963	U	C4-C5-C6	-6.03	116.08	119.70
85	AA	1858	G	C4-N9-C1'	-6.03	118.66	126.50
85	AA	1923	A	OP1-P-OP2	-6.03	110.55	119.60
85	AA	2075	C	O4'-C1'-C2'	6.03	113.03	107.60
34	BA	294	C	C4'-C3'-C2'	6.03	108.63	102.60
34	BA	407	A	C8-N9-C4	-6.03	103.39	105.80
34	BA	1470	G	C4-N9-C1'	-6.03	118.66	126.50
34	BA	1611	A	O4'-C1'-N9	6.03	113.03	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1826	C	N3-C4-C5	6.03	124.31	121.90
35	BB	134	G	N1-C2-N2	-6.03	110.77	116.20
35	BB	587	A	C5'-C4'-C3'	6.03	125.65	116.00
38	BE	13	A	P-O3'-C3'	6.03	126.94	119.70
38	BE	127	G	C4-C5-C6	-6.03	115.18	118.80
40	BG	149	U	O4'-C1'-N1	6.03	113.03	108.20
70	Bk	77	LYS	N-CA-C	-6.03	94.71	111.00
85	AA	527	A	C1'-O4'-C4'	-6.03	105.07	109.90
34	BA	56	G	C3'-C2'-C1'	-6.03	96.67	101.50
34	BA	301	U	N3-C2-O2	-6.03	117.98	122.20
34	BA	581	U	C2'-C3'-O3'	6.03	123.35	113.70
34	BA	739	A	C5'-C4'-C3'	6.03	125.65	116.00
34	BA	950	C	O4'-C1'-N1	6.03	113.02	108.20
34	BA	1469	G	N1-C6-O6	6.03	123.52	119.90
34	BA	1712	U	C5-C4-O4	-6.03	122.28	125.90
34	BA	1765	G	O4'-C1'-N9	6.03	113.03	108.20
35	BB	64	U	P-O3'-C3'	-6.03	112.46	119.70
35	BB	320	G	O4'-C1'-N9	6.03	113.02	108.20
35	BB	822	G	C3'-C2'-C1'	-6.03	96.68	101.50
35	BB	1491	G	N9-C1'-C2'	-6.03	105.37	112.00
38	BE	87	U	C4'-C3'-C2'	-6.03	96.57	102.60
40	BG	38	A	O4'-C1'-C2'	6.03	113.03	107.60
57	BX	87	TYR	CB-CG-CD2	6.03	124.62	121.00
58	BY	70	ARG	NE-CZ-NH2	-6.03	117.28	120.30
85	AA	15	U	N1-C2-N3	6.03	118.52	114.90
85	AA	441	C	C1'-O4'-C4'	-6.03	105.08	109.90
85	AA	472	A	C4-N9-C1'	-6.03	115.45	126.30
85	AA	986	U	C2-N3-C4	-6.03	123.38	127.00
34	BA	1232	C	C1'-O4'-C4'	-6.03	105.08	109.90
34	BA	1451	A	O4'-C1'-N9	6.03	113.02	108.20
35	BB	1493	A	C8-N9-C1'	6.03	138.55	127.70
38	BE	94	U	N3-C4-C5	6.03	118.22	114.60
41	BH	10	U	N3-C4-O4	-6.03	115.18	119.40
85	AA	613	G	C4'-C3'-C2'	-6.03	96.57	102.60
85	AA	1592	C	C5-C6-N1	6.03	124.01	121.00
34	BA	640	U	C6-N1-C1'	-6.03	112.76	121.20
34	BA	756	A	C5'-C4'-O4'	-6.03	101.87	109.10
35	BB	337	U	P-O5'-C5'	6.03	130.54	120.90
35	BB	1292	G	C6-N1-C2	-6.03	121.48	125.10
35	BB	1440	A	P-O3'-C3'	-6.03	112.47	119.70
35	BB	1463	A	C4-N9-C1'	6.03	137.15	126.30
38	BE	14	C	O3'-P-O5'	-6.03	92.55	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	BN	33	GLN	CA-C-N	6.03	133.98	117.10
85	AA	71	G	P-O3'-C3'	6.03	126.93	119.70
85	AA	209	C	P-O5'-C5'	-6.03	111.25	120.90
85	AA	583	U	P-O3'-C3'	-6.03	112.47	119.70
85	AA	1113	G	O4'-C1'-N9	6.03	113.02	108.20
85	AA	1293	U	O4'-C4'-C3'	-6.03	97.97	104.00
85	AA	2115	G	C5-C6-O6	-6.03	124.98	128.60
34	BA	307	C	N1-C1'-C2'	-6.03	105.37	112.00
34	BA	649	A	C5'-C4'-C3'	-6.03	106.36	116.00
34	BA	683	C	C4-C5-C6	6.03	120.41	117.40
34	BA	1631	U	O5'-C5'-C4'	6.03	123.15	111.70
35	BB	751	A	C5-C6-N6	6.03	128.52	123.70
35	BB	1363	A	C4-N9-C1'	-6.03	115.45	126.30
35	BB	1537	C	P-O3'-C3'	-6.03	112.47	119.70
37	BD	19	C	C4'-C3'-C2'	-6.03	96.57	102.60
37	BD	59	G	C4-N9-C1'	-6.03	118.67	126.50
40	BG	30	C	C6-N1-C1'	6.03	128.03	120.80
41	BH	58	C	N3-C2-O2	-6.03	117.68	121.90
85	AA	357	C	N3-C4-N4	6.03	122.22	118.00
85	AA	837	C	O4'-C1'-N1	6.03	113.02	108.20
85	AA	1082	U	C2-N1-C1'	6.03	124.93	117.70
85	AA	1874	G	C5'-C4'-O4'	6.03	116.33	109.10
34	BA	1165	A	C8-N9-C1'	6.02	138.54	127.70
85	AA	36	U	C2-N1-C1'	-6.02	110.47	117.70
85	AA	75	U	P-O5'-C5'	6.02	130.54	120.90
85	AA	605	A	N9-C1'-C2'	-6.02	105.37	112.00
85	AA	1929	G	C4-N9-C1'	-6.02	118.67	126.50
85	AA	2162	G	C8-N9-C1'	6.02	134.83	127.00
13	AE	145	ARG	NE-CZ-NH1	6.02	123.31	120.30
34	BA	155	U	C2-N3-C4	-6.02	123.39	127.00
34	BA	318	U	P-O3'-C3'	-6.02	112.47	119.70
34	BA	1258	G	O4'-C1'-N9	6.02	113.02	108.20
34	BA	1413	G	C5-C6-N1	6.02	114.51	111.50
35	BB	430	A	C8-N9-C4	6.02	108.21	105.80
35	BB	839	G	N9-C4-C5	6.02	107.81	105.40
35	BB	913	C	O4'-C1'-N1	6.02	113.02	108.20
35	BB	1434	G	N9-C4-C5	-6.02	102.99	105.40
36	BC	139	A	P-O5'-C5'	-6.02	111.27	120.90
38	BE	180	G	N3-C2-N2	-6.02	115.68	119.90
40	BG	172	C	P-O3'-C3'	-6.02	112.47	119.70
41	BH	48	G	OP1-P-O3'	6.02	118.45	105.20
85	AA	335	G	C1'-O4'-C4'	-6.02	105.08	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1996	A	C2-N3-C4	-6.02	107.59	110.60
86	AB	1	G	N1-C6-O6	6.02	123.51	119.90
34	BA	891	C	C2-N1-C1'	-6.02	112.18	118.80
34	BA	1038	U	C1'-O4'-C4'	-6.02	105.08	109.90
41	BH	89	C	OP1-P-OP2	-6.02	110.57	119.60
85	AA	83	U	N1-C1'-C2'	-6.02	105.38	112.00
85	AA	530	A	P-O3'-C3'	-6.02	112.47	119.70
85	AA	657	C	C3'-C2'-C1'	-6.02	96.68	101.50
85	AA	820	G	P-O3'-C3'	6.02	126.92	119.70
85	AA	1449	C	C2'-C3'-O3'	6.02	123.33	113.70
85	AA	2045	U	N3-C2-O2	-6.02	117.98	122.20
85	AA	2180	C	O4'-C1'-N1	6.02	113.02	108.20
34	BA	143	A	P-O3'-C3'	-6.02	112.48	119.70
34	BA	749	G	C1'-O4'-C4'	-6.02	105.08	109.90
34	BA	801	U	O4'-C1'-N1	6.02	113.02	108.20
34	BA	1165	A	C4-N9-C1'	-6.02	115.47	126.30
34	BA	1707	C	C5'-C4'-C3'	6.02	125.63	116.00
34	BA	1800	G	N1-C6-O6	6.02	123.51	119.90
35	BB	1413	U	OP1-P-O3'	6.02	118.44	105.20
38	BE	199	A	O4'-C4'-C3'	-6.02	97.98	104.00
40	BG	141	A	C8-N9-C4	6.02	108.21	105.80
52	BS	82	TYR	CB-CG-CD1	6.02	124.61	121.00
85	AA	671	G	P-O3'-C3'	-6.02	112.48	119.70
85	AA	887	A	C5'-C4'-O4'	6.02	116.32	109.10
34	BA	228	A	C5-C6-N6	-6.02	118.89	123.70
34	BA	304	G	C4-N9-C1'	-6.02	118.68	126.50
34	BA	1048	C	C6-N1-C1'	6.02	128.02	120.80
34	BA	1571	C	C5'-C4'-C3'	6.02	125.63	116.00
35	BB	18	A	C5'-C4'-C3'	6.02	125.63	116.00
35	BB	321	C	O4'-C1'-N1	6.02	113.01	108.20
35	BB	442	U	C3'-C2'-C1'	-6.02	96.69	101.50
35	BB	497	C	N1-C1'-C2'	-6.02	105.38	112.00
35	BB	518	G	C1'-O4'-C4'	-6.02	105.08	109.90
35	BB	1166	A	C4'-C3'-C2'	-6.02	96.58	102.60
35	BB	1196	A	C3'-C2'-C1'	-6.02	96.69	101.50
38	BE	19	G	OP1-P-OP2	-6.02	110.57	119.60
38	BE	101	C	P-O3'-C3'	-6.02	112.48	119.70
85	AA	575	G	N1-C6-O6	6.02	123.51	119.90
85	AA	1613	A	C5'-C4'-C3'	-6.02	106.37	116.00
85	AA	2154	C	C4'-C3'-C2'	-6.02	96.58	102.60
34	BA	1633	C	P-O5'-C5'	6.02	130.53	120.90
34	BA	1816	G	C6-N1-C2	-6.02	121.49	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	126	C	O4'-C1'-N1	6.02	113.01	108.20
35	BB	694	C	N3-C4-N4	-6.02	113.79	118.00
35	BB	1131	C	P-O5'-C5'	-6.02	111.28	120.90
36	BC	51	A	C5-N7-C8	-6.02	100.89	103.90
39	BF	39	C	C5-C6-N1	6.02	124.01	121.00
40	BG	84	U	O4'-C1'-C2'	-6.02	99.78	105.80
2	A1	215	PHE	CB-CG-CD1	6.01	125.01	120.80
34	BA	1268	C	C5'-C4'-C3'	-6.01	106.38	116.00
35	BB	351	G	C5-C6-O6	-6.01	124.99	128.60
35	BB	820	C	O4'-C1'-N1	6.01	113.01	108.20
38	BE	6	A	C5-C6-N1	6.01	120.71	117.70
38	BE	116	U	N1-C2-O2	6.01	127.01	122.80
41	BH	135	U	C5-C4-O4	6.01	129.51	125.90
72	Bm	27	ARG	NE-CZ-NH1	6.01	123.31	120.30
85	AA	203	C	C4'-C3'-C2'	-6.01	96.59	102.60
85	AA	309	G	C4'-C3'-C2'	6.01	108.61	102.60
85	AA	375	C	C5'-C4'-C3'	6.01	125.62	116.00
85	AA	535	G	P-O5'-C5'	6.01	130.52	120.90
85	AA	711	C	C3'-C2'-C1'	6.01	106.31	101.50
85	AA	1290	G	C4'-C3'-C2'	-6.01	96.58	102.60
7	A6	4	TYR	N-CA-C	6.01	127.23	111.00
7	A6	145	PHE	CB-CG-CD1	6.01	125.01	120.80
34	BA	304	G	C8-N9-C1'	6.01	134.82	127.00
34	BA	1734	U	C6-N1-C1'	6.01	129.62	121.20
35	BB	58	G	O4'-C1'-N9	6.01	113.01	108.20
35	BB	1174	C	C6-N1-C1'	-6.01	113.58	120.80
49	BP	160	ALA	N-CA-C	6.01	127.24	111.00
85	AA	2029	G	P-O3'-C3'	-6.01	112.48	119.70
34	BA	216	C	C5'-C4'-C3'	-6.01	106.38	116.00
34	BA	244	A	C1'-O4'-C4'	-6.01	105.09	109.90
34	BA	295	G	N3-C4-C5	-6.01	125.59	128.60
34	BA	304	G	C5-C6-O6	-6.01	124.99	128.60
34	BA	859	G	O4'-C1'-N9	6.01	113.01	108.20
34	BA	1043	C	C6-N1-C2	-6.01	117.89	120.30
34	BA	1236	U	C2-N3-C4	-6.01	123.39	127.00
35	BB	1250	A	O4'-C1'-C2'	6.01	113.01	107.60
36	BC	134	G	N1-C2-N2	-6.01	110.79	116.20
38	BE	19	G	O3'-P-O5'	6.01	115.42	104.00
67	Bh	23	ARG	NE-CZ-NH1	6.01	123.31	120.30
85	AA	334	A	C3'-C2'-C1'	-6.01	96.69	101.50
85	AA	448	G	N1-C6-O6	-6.01	116.29	119.90
85	AA	753	U	C5'-C4'-C3'	6.01	125.62	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1197	U	P-O3'-C3'	-6.01	112.48	119.70
85	AA	2120	C	P-O3'-C3'	-6.01	112.49	119.70
31	AX	172	ARG	NE-CZ-NH2	-6.01	117.30	120.30
34	BA	604	G	C3'-C2'-C1'	-6.01	96.69	101.50
34	BA	1153	C	C5-C4-N4	-6.01	115.99	120.20
35	BB	650	A	O4'-C1'-N9	6.01	113.01	108.20
35	BB	1045	G	N9-C1'-C2'	-6.01	105.39	112.00
35	BB	1140	C	P-O5'-C5'	6.01	130.51	120.90
38	BE	130	G	N1-C2-N2	-6.01	110.79	116.20
40	BG	104	A	C1'-O4'-C4'	-6.01	105.09	109.90
41	BH	41	A	N9-C4-C5	-6.01	103.40	105.80
68	Bi	128	ARG	NE-CZ-NH1	6.01	123.31	120.30
71	Bl	95	TRP	N-CA-C	-6.01	94.77	111.00
85	AA	353	G	C8-N9-C1'	6.01	134.81	127.00
85	AA	616	A	C5'-C4'-O4'	6.01	116.31	109.10
85	AA	1840	C	C2-N3-C4	-6.01	116.89	119.90
85	AA	2118	U	C6-N1-C2	-6.01	117.39	121.00
34	BA	777	C	C2-N3-C4	-6.01	116.90	119.90
34	BA	1342	C	C5-C4-N4	-6.01	115.99	120.20
34	BA	1642	A	C4-N9-C1'	-6.01	115.48	126.30
35	BB	512	C	C4'-C3'-C2'	-6.01	96.59	102.60
35	BB	832	C	P-O3'-C3'	-6.01	112.49	119.70
35	BB	1036	G	P-O3'-C3'	-6.01	112.49	119.70
35	BB	1096	G	C5-C6-N1	6.01	114.50	111.50
85	AA	862	U	C6-N1-C1'	6.01	129.61	121.20
85	AA	2096	G	C6-N1-C2	-6.01	121.50	125.10
34	BA	478	G	C3'-C2'-C1'	-6.01	96.69	101.50
34	BA	1573	C	N3-C2-O2	-6.01	117.69	121.90
34	BA	1804	A	O4'-C1'-C2'	-6.01	99.79	105.80
35	BB	1054	G	C5'-C4'-C3'	-6.01	106.39	116.00
39	BF	53	G	O3'-P-O5'	-6.01	92.59	104.00
41	BH	2	U	P-O5'-C5'	6.01	130.51	120.90
81	Bv	175	LEU	N-CA-CB	-6.01	98.39	110.40
85	AA	146	U	O5'-C5'-C4'	6.01	123.11	111.70
85	AA	1648	G	P-O5'-C5'	-6.01	111.29	120.90
34	BA	253	U	N3-C2-O2	-6.00	118.00	122.20
34	BA	733	G	C4'-C3'-C2'	6.00	108.61	102.60
35	BB	895	U	N1-C2-N3	6.00	118.50	114.90
72	Bm	23	TYR	CB-CG-CD2	-6.00	117.40	121.00
85	AA	171	U	O4'-C4'-C3'	6.00	110.90	106.10
34	BA	110	C	C4'-C3'-C2'	-6.00	96.60	102.60
34	BA	313	C	O4'-C1'-N1	6.00	113.00	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	440	A	C5'-C4'-C3'	-6.00	106.39	116.00
34	BA	630	U	O3'-P-O5'	-6.00	92.59	104.00
34	BA	1008	A	P-O3'-C3'	-6.00	112.50	119.70
34	BA	1154	U	P-O3'-C3'	-6.00	112.50	119.70
34	BA	1274	A	O3'-P-O5'	-6.00	92.59	104.00
34	BA	1322	A	N1-C6-N6	6.00	122.20	118.60
34	BA	1421	A	C8-N9-C4	6.00	108.20	105.80
34	BA	1529	G	P-O3'-C3'	-6.00	112.50	119.70
34	BA	1545	C	N1-C2-N3	6.00	123.40	119.20
34	BA	1742	G	N1-C6-O6	6.00	123.50	119.90
35	BB	338	C	O4'-C1'-N1	6.00	113.00	108.20
35	BB	423	G	C8-N9-C1'	6.00	134.81	127.00
35	BB	474	G	N9-C4-C5	-6.00	103.00	105.40
35	BB	1544	A	O4'-C1'-N9	6.00	113.00	108.20
36	BC	169	G	N3-C4-N9	-6.00	122.40	126.00
40	BG	84	U	C5'-C4'-O4'	6.00	116.30	109.10
40	BG	122	G	C8-N9-C4	6.00	108.80	106.40
47	BN	13	GLN	N-CA-CB	6.00	121.41	110.60
52	BS	71	LEU	N-CA-C	6.00	127.21	111.00
85	AA	351	C	N3-C2-O2	-6.00	117.70	121.90
85	AA	959	C	P-O3'-C3'	-6.00	112.50	119.70
85	AA	1670	U	C4'-C3'-C2'	-6.00	96.60	102.60
85	AA	2054	G	C4-N9-C1'	-6.00	118.70	126.50
85	AA	2187	G	O4'-C1'-N9	6.00	113.00	108.20
34	BA	758	G	C5-C6-O6	-6.00	125.00	128.60
34	BA	1125	G	O3'-P-O5'	-6.00	92.60	104.00
34	BA	1499	A	C8-N9-C4	-6.00	103.40	105.80
35	BB	995	C	C2'-C3'-O3'	6.00	123.30	113.70
35	BB	1236	A	O4'-C1'-N9	6.00	113.00	108.20
36	BC	17	U	O4'-C1'-N1	6.00	113.00	108.20
40	BG	133	C	O3'-P-O5'	-6.00	92.60	104.00
40	BG	133	C	P-O3'-C3'	6.00	126.90	119.70
52	BS	109	TYR	CB-CG-CD1	6.00	124.60	121.00
85	AA	38	C	C4'-C3'-C2'	-6.00	96.60	102.60
85	AA	370	A	C4'-C3'-C2'	6.00	108.60	102.60
85	AA	492	C	C5-C4-N4	6.00	124.40	120.20
85	AA	587	G	N9-C1'-C2'	-6.00	105.40	112.00
85	AA	964	C	C6-N1-C2	-6.00	117.90	120.30
34	BA	611	A	C6-C5-N7	-6.00	128.10	132.30
34	BA	1295	U	P-O3'-C3'	6.00	126.90	119.70
35	BB	568	A	O4'-C1'-N9	6.00	113.00	108.20
35	BB	1446	C	N3-C4-C5	6.00	124.30	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A0	49	ASN	CB-CA-C	-6.00	98.40	110.40
34	BA	192	G	O3'-P-O5'	6.00	115.40	104.00
34	BA	454	G	P-O5'-C5'	-6.00	111.30	120.90
34	BA	1487	U	P-O5'-C5'	6.00	130.50	120.90
34	BA	1535	G	C5'-C4'-O4'	6.00	116.30	109.10
35	BB	1390	U	C5-C6-N1	-6.00	119.70	122.70
48	BO	154	PHE	N-CA-CB	6.00	121.40	110.60
84	By	53	PHE	CB-CG-CD1	6.00	125.00	120.80
85	AA	128	U	C5'-C4'-C3'	-6.00	106.40	116.00
85	AA	186	U	C6-N1-C1'	6.00	129.60	121.20
85	AA	282	C	O4'-C4'-C3'	6.00	110.90	106.10
85	AA	459	C	C1'-O4'-C4'	-6.00	105.10	109.90
85	AA	487	G	C4-N9-C1'	-6.00	118.70	126.50
85	AA	703	U	O4'-C1'-N1	6.00	113.00	108.20
85	AA	1088	C	C6-N1-C2	-6.00	117.90	120.30
85	AA	1153	G	C4'-C3'-C2'	-6.00	96.60	102.60
85	AA	1577	G	O4'-C1'-N9	6.00	113.00	108.20
34	BA	1407	C	C2-N1-C1'	-6.00	112.20	118.80
35	BB	93	A	N1-C6-N6	6.00	122.20	118.60
35	BB	144	G	C8-N9-C1'	6.00	134.80	127.00
35	BB	642	G	N3-C4-C5	-6.00	125.60	128.60
35	BB	1133	C	C5-C4-N4	6.00	124.40	120.20
78	Bs	51	LEU	CB-CG-CD2	6.00	121.19	111.00
85	AA	862	U	C2-N3-C4	-6.00	123.40	127.00
34	BA	1324	G	O5'-C5'-C4'	-6.00	100.31	111.70
34	BA	1484	A	C1'-O4'-C4'	-6.00	105.10	109.90
35	BB	1279	C	N1-C1'-C2'	-6.00	105.41	112.00
41	BH	119	U	O5'-C5'-C4'	-6.00	100.31	111.70
80	Bu	20	TYR	CA-CB-CG	-6.00	102.01	113.40
85	AA	628	C	C2-N3-C4	-6.00	116.90	119.90
2	A1	46	ARG	N-CA-CB	-5.99	99.81	110.60
34	BA	313	C	O4'-C1'-C2'	5.99	112.99	107.60
34	BA	387	A	C8-N9-C4	-5.99	103.40	105.80
34	BA	535	G	N3-C2-N2	5.99	124.10	119.90
34	BA	565	U	C2-N1-C1'	-5.99	110.51	117.70
34	BA	671	C	C2-N3-C4	-5.99	116.90	119.90
34	BA	748	C	P-O5'-C5'	5.99	130.49	120.90
35	BB	1479	C	N3-C2-O2	-5.99	117.70	121.90
36	BC	10	C	O5'-C5'-C4'	5.99	123.09	111.70
42	BI	69	VAL	CG1-CB-CG2	-5.99	101.31	110.90
82	Bw	252	PHE	N-CA-CB	-5.99	99.81	110.60
85	AA	470	C	N3-C2-O2	-5.99	117.70	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	716	G	P-O5'-C5'	-5.99	111.31	120.90
85	AA	771	A	N3-C4-C5	-5.99	122.60	126.80
85	AA	944	C	P-O5'-C5'	5.99	130.49	120.90
85	AA	1674	G	N1-C6-O6	5.99	123.50	119.90
85	AA	1711	C	C1'-O4'-C4'	-5.99	105.11	109.90
35	BB	1068	G	C2-N3-C4	-5.99	108.90	111.90
35	BB	1176	G	C3'-C2'-C1'	-5.99	96.71	101.50
38	BE	72	C	O5'-C5'-C4'	5.99	123.08	111.70
38	BE	94	U	N3-C2-O2	-5.99	118.01	122.20
38	BE	153	C	P-O5'-C5'	-5.99	111.31	120.90
44	BK	115	MET	CG-SD-CE	-5.99	90.61	100.20
85	AA	162	A	C5-N7-C8	-5.99	100.90	103.90
85	AA	425	G	N7-C8-N9	-5.99	110.10	113.10
85	AA	1370	G	C4-N9-C1'	-5.99	118.71	126.50
85	AA	2066	C	P-O5'-C5'	-5.99	111.31	120.90
34	BA	14	G	P-O5'-C5'	-5.99	111.32	120.90
34	BA	183	G	C5-C6-O6	-5.99	125.01	128.60
34	BA	478	G	N9-C1'-C2'	-5.99	105.41	112.00
34	BA	486	G	N9-C1'-C2'	5.99	121.79	114.00
34	BA	523	A	O4'-C1'-N9	-5.99	103.41	108.20
34	BA	551	U	C5-C4-O4	5.99	129.49	125.90
34	BA	574	U	C1'-O4'-C4'	5.99	114.69	109.90
34	BA	681	G	C5-N7-C8	-5.99	101.31	104.30
34	BA	857	C	C6-N1-C1'	5.99	127.99	120.80
34	BA	951	C	N1-C1'-C2'	-5.99	105.41	112.00
34	BA	1243	A	C4'-C3'-C2'	-5.99	96.61	102.60
34	BA	1457	C	O4'-C1'-C2'	5.99	112.99	107.60
35	BB	93	A	O5'-C5'-C4'	-5.99	100.32	111.70
35	BB	1256	C	N3-C2-O2	-5.99	117.71	121.90
37	BD	100	A	C5'-C4'-O4'	-5.99	101.91	109.10
38	BE	194	A	C5-N7-C8	-5.99	100.91	103.90
40	BG	20	U	N1-C2-N3	5.99	118.49	114.90
41	BH	76	G	OP1-P-O3'	5.99	118.38	105.20
25	AR	54	THR	N-CA-C	-5.99	94.83	111.00
34	BA	1071	G	C6-N1-C2	-5.99	121.51	125.10
34	BA	1094	U	C5-C6-N1	-5.99	119.71	122.70
34	BA	1344	G	N7-C8-N9	5.99	116.09	113.10
34	BA	1438	C	N3-C2-O2	-5.99	117.71	121.90
34	BA	1632	G	C5-C6-O6	-5.99	125.01	128.60
34	BA	1685	C	C2-N3-C4	-5.99	116.91	119.90
34	BA	1747	C	C2'-C3'-O3'	5.99	123.28	113.70
35	BB	55	C	P-O5'-C5'	-5.99	111.32	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	147	C	C2-N1-C1'	5.99	125.39	118.80
35	BB	490	G	N1-C6-O6	5.99	123.49	119.90
35	BB	828	G	N9-C1'-C2'	-5.99	105.41	112.00
35	BB	1072	C	C1'-O4'-C4'	-5.99	105.11	109.90
40	BG	168	A	P-O5'-C5'	-5.99	111.32	120.90
60	Ba	101	ALA	N-CA-CB	5.99	118.48	110.10
85	AA	99	U	C5'-C4'-C3'	-5.99	106.42	116.00
85	AA	385	A	P-O5'-C5'	-5.99	111.32	120.90
34	BA	780	U	O5'-C5'-C4'	5.99	123.08	111.70
34	BA	955	G	O4'-C1'-N9	5.99	112.99	108.20
34	BA	1703	A	N1-C6-N6	5.99	122.19	118.60
35	BB	428	G	O4'-C1'-N9	5.99	112.99	108.20
40	BG	152	G	C4-N9-C1'	-5.99	118.72	126.50
83	Bx	40	PHE	CB-CA-C	-5.99	98.43	110.40
85	AA	488	G	C8-N9-C1'	5.99	134.78	127.00
85	AA	1526	G	C2'-C3'-O3'	5.99	123.28	113.70
85	AA	1721	A	C4-N9-C1'	-5.99	115.52	126.30
85	AA	1854	U	C2-N1-C1'	-5.99	110.52	117.70
34	BA	1403	G	N9-C1'-C2'	-5.99	105.42	112.00
34	BA	1499	A	O5'-C5'-C4'	-5.99	100.33	111.70
35	BB	412	A	P-O5'-C5'	-5.99	111.32	120.90
35	BB	969	C	N3-C2-O2	-5.99	117.71	121.90
35	BB	1234	G	N3-C2-N2	5.99	124.09	119.90
35	BB	1354	C	C5'-C4'-O4'	-5.99	101.92	109.10
35	BB	1466	A	C5-N7-C8	-5.99	100.91	103.90
40	BG	130	G	C4-N9-C1'	-5.99	118.72	126.50
51	BR	81	GLY	N-CA-C	-5.99	98.14	113.10
85	AA	910	G	C8-N9-C1'	5.99	134.78	127.00
85	AA	934	A	P-O3'-C3'	-5.99	112.52	119.70
85	AA	2113	U	C2-N1-C1'	-5.99	110.52	117.70
34	BA	1569	C	C5'-C4'-O4'	5.98	116.28	109.10
35	BB	540	G	N1-C6-O6	5.98	123.49	119.90
38	BE	134	A	N1-C2-N3	-5.98	126.31	129.30
38	BE	141	A	O4'-C1'-N9	5.98	112.99	108.20
41	BH	24	U	P-O3'-C3'	-5.98	112.52	119.70
34	BA	339	G	N1-C6-O6	-5.98	116.31	119.90
34	BA	595	U	C3'-C2'-C1'	5.98	106.29	101.50
34	BA	1220	C	C2'-C3'-O3'	5.98	123.27	113.70
35	BB	2	C	O4'-C4'-C3'	-5.98	98.02	104.00
35	BB	545	C	C3'-C2'-C1'	5.98	106.29	101.50
35	BB	1162	A	C8-N9-C1'	-5.98	116.93	127.70
35	BB	1540	U	O4'-C1'-N1	5.98	112.99	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	70	A	O4'-C1'-N9	5.98	112.98	108.20
40	BG	181	C	C1'-O4'-C4'	-5.98	105.11	109.90
82	Bw	220	PHE	CB-CG-CD1	5.98	124.99	120.80
85	AA	715	G	N9-C1'-C2'	-5.98	105.42	112.00
85	AA	1981	A	P-O5'-C5'	-5.98	111.33	120.90
5	A4	162	ARG	N-CA-CB	5.98	121.36	110.60
34	BA	276	C	N3-C2-O2	-5.98	117.71	121.90
34	BA	403	A	C4'-C3'-C2'	-5.98	96.62	102.60
34	BA	420	A	C1'-O4'-C4'	-5.98	105.12	109.90
34	BA	1223	C	C6-N1-C1'	-5.98	113.62	120.80
34	BA	1653	G	N1-C6-O6	-5.98	116.31	119.90
34	BA	1737	A	O3'-P-O5'	-5.98	92.64	104.00
40	BG	141	A	C3'-C2'-C1'	5.98	106.28	101.50
73	Bn	66	TYR	CB-CA-C	5.98	122.36	110.40
85	AA	132	G	O4'-C1'-N9	5.98	112.98	108.20
85	AA	1342	C	O4'-C1'-N1	5.98	112.98	108.20
85	AA	1900	C	C6-N1-C1'	5.98	127.98	120.80
85	AA	2125	A	C4-C5-C6	-5.98	114.01	117.00
85	AA	2139	G	C5-N7-C8	-5.98	101.31	104.30
85	AA	2192	A	C5-C6-N1	5.98	120.69	117.70
34	BA	743	A	N7-C8-N9	-5.98	110.81	113.80
34	BA	1030	C	C3'-C2'-C1'	-5.98	96.72	101.50
34	BA	1289	C	C6-N1-C2	-5.98	117.91	120.30
35	BB	41	A	O4'-C4'-C3'	5.98	110.88	106.10
35	BB	458	U	C2-N3-C4	-5.98	123.41	127.00
35	BB	1306	G	C4-N9-C1'	-5.98	118.73	126.50
37	BD	65	G	C8-N9-C4	5.98	108.79	106.40
38	BE	14	C	C4'-C3'-C2'	5.98	108.58	102.60
41	BH	56	C	O4'-C1'-N1	5.98	112.98	108.20
85	AA	96	C	N1-C2-O2	5.98	122.49	118.90
85	AA	1212	C	C2-N3-C4	-5.98	116.91	119.90
85	AA	1826	U	C6-N1-C1'	-5.98	112.83	121.20
34	BA	612	U	C5'-C4'-C3'	-5.98	106.44	116.00
34	BA	884	G	C2-N3-C4	-5.98	108.91	111.90
34	BA	1286	C	O4'-C1'-N1	5.98	112.98	108.20
34	BA	1762	U	O4'-C1'-N1	5.98	112.98	108.20
35	BB	8	U	C2'-C3'-O3'	5.98	123.27	113.70
35	BB	382	U	N1-C2-N3	5.98	118.49	114.90
35	BB	620	G	C3'-C2'-C1'	-5.98	96.72	101.50
35	BB	1206	G	C1'-O4'-C4'	-5.98	105.12	109.90
36	BC	155	C	P-O5'-C5'	-5.98	111.34	120.90
38	BE	163	A	C3'-C2'-C1'	-5.98	96.72	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
65	Bf	144	ARG	NE-CZ-NH1	5.98	123.29	120.30
84	By	44	PHE	CB-CG-CD2	-5.98	116.61	120.80
85	AA	707	U	C4'-C3'-C2'	-5.98	96.62	102.60
85	AA	901	C	C2-N3-C4	-5.98	116.91	119.90
85	AA	1505	G	C8-N9-C1'	5.98	134.77	127.00
35	BB	1337	C	C5'-C4'-O4'	5.98	116.27	109.10
35	BB	1386	C	C5'-C4'-C3'	-5.98	106.44	116.00
35	BB	1452	U	C2'-C3'-O3'	5.98	123.26	113.70
35	BB	1528	U	C3'-C2'-C1'	-5.98	96.72	101.50
36	BC	88	A	O3'-P-O5'	-5.98	92.64	104.00
80	Bu	264	ARG	NE-CZ-NH2	-5.98	117.31	120.30
85	AA	264	A	O4'-C1'-N9	5.98	112.98	108.20
85	AA	656	U	P-O5'-C5'	-5.98	111.34	120.90
85	AA	1223	A	N1-C2-N3	-5.98	126.31	129.30
34	BA	373	G	C5-C6-O6	-5.97	125.02	128.60
34	BA	750	C	P-O3'-C3'	-5.97	112.53	119.70
34	BA	897	U	C2'-C3'-O3'	5.97	123.26	113.70
34	BA	1129	U	O4'-C1'-N1	5.97	112.98	108.20
34	BA	1680	G	C5'-C4'-C3'	5.97	125.56	116.00
34	BA	1822	U	C5'-C4'-C3'	5.97	125.56	116.00
36	BC	146	U	C2'-C3'-O3'	5.97	123.26	113.70
67	Bh	35	ASP	CA-CB-CG	5.97	126.54	113.40
85	AA	338	G	N3-C2-N2	5.97	124.08	119.90
85	AA	875	C	C1'-O4'-C4'	-5.97	105.12	109.90
85	AA	1055	U	C2-N1-C1'	5.97	124.87	117.70
85	AA	1301	C	P-O5'-C5'	-5.97	111.34	120.90
85	AA	1812	C	O5'-C5'-C4'	-5.97	100.35	111.70
85	AA	2045	U	C6-N1-C1'	5.97	129.56	121.20
85	AA	2119	C	O4'-C1'-N1	5.97	112.98	108.20
85	AA	2241	C	C6-N1-C1'	5.97	127.97	120.80
86	AB	3	C	C6-N1-C2	-5.97	117.91	120.30
34	BA	275	C	P-O3'-C3'	-5.97	112.53	119.70
34	BA	684	G	N3-C4-N9	5.97	129.58	126.00
35	BB	1147	G	N1-C6-O6	5.97	123.48	119.90
74	Bo	13	ARG	NH1-CZ-NH2	-5.97	112.83	119.40
77	Br	250	TRP	CB-CA-C	-5.97	98.45	110.40
85	AA	31	C	O4'-C1'-N1	5.97	112.98	108.20
85	AA	115	U	C4-C5-C6	-5.97	116.12	119.70
85	AA	168	A	O5'-C5'-C4'	-5.97	100.35	111.70
85	AA	652	U	C2-N3-C4	-5.97	123.42	127.00
85	AA	847	G	C5-C6-N1	5.97	114.49	111.50
85	AA	1709	U	P-O3'-C3'	5.97	126.87	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2096	G	P-O5'-C5'	5.97	130.46	120.90
85	AA	2204	A	C5'-C4'-O4'	5.97	116.27	109.10
29	AV	53	ARG	NE-CZ-NH2	-5.97	117.31	120.30
34	BA	656	U	P-O5'-C5'	5.97	130.45	120.90
34	BA	1295	U	C5'-C4'-O4'	-5.97	101.94	109.10
38	BE	184	G	N1-C6-O6	-5.97	116.32	119.90
85	AA	916	A	O4'-C1'-N9	5.97	112.98	108.20
34	BA	83	G	C6-C5-N7	-5.97	126.82	130.40
34	BA	103	G	N7-C8-N9	-5.97	110.11	113.10
34	BA	307	C	O4'-C1'-N1	5.97	112.97	108.20
34	BA	700	G	C3'-C2'-C1'	-5.97	96.72	101.50
34	BA	736	G	O4'-C1'-N9	5.97	112.98	108.20
34	BA	806	U	C5'-C4'-C3'	5.97	125.55	116.00
34	BA	1430	C	P-O3'-C3'	5.97	126.86	119.70
34	BA	1637	G	C8-N9-C1'	5.97	134.76	127.00
34	BA	1733	G	C4-N9-C1'	-5.97	118.74	126.50
35	BB	1106	G	N1-C6-O6	-5.97	116.32	119.90
37	BD	23	A	N9-C1'-C2'	-5.97	105.43	112.00
38	BE	168	C	N1-C2-N3	5.97	123.38	119.20
40	BG	12	A	O4'-C1'-C2'	-5.97	99.83	105.80
85	AA	646	C	N3-C2-O2	-5.97	117.72	121.90
85	AA	1176	C	C2-N1-C1'	-5.97	112.23	118.80
85	AA	1482	C	C5-C6-N1	5.97	123.98	121.00
34	BA	713	C	P-O5'-C5'	-5.97	111.35	120.90
37	BD	62	A	C4-N9-C1'	-5.97	115.56	126.30
38	BE	23	G	C5-C6-N1	5.97	114.48	111.50
38	BE	181	U	O5'-P-OP2	-5.97	100.33	105.70
40	BG	140	G	O4'-C1'-C2'	5.97	112.97	107.60
41	BH	22	A	C5-C6-N6	-5.97	118.92	123.70
69	Bj	11	ARG	NE-CZ-NH1	5.97	123.28	120.30
85	AA	365	G	O4'-C1'-N9	5.97	112.97	108.20
85	AA	386	G	N1-C2-N2	-5.97	110.83	116.20
85	AA	453	G	N3-C4-N9	5.97	129.58	126.00
85	AA	760	U	O4'-C4'-C3'	-5.97	98.03	104.00
34	BA	75	U	N1-C1'-C2'	-5.97	105.44	112.00
34	BA	260	A	C5-C6-N6	5.97	128.47	123.70
34	BA	399	G	N3-C4-C5	5.97	131.58	128.60
34	BA	541	C	C6-N1-C1'	5.97	127.96	120.80
34	BA	1222	C	P-O3'-C3'	-5.97	112.54	119.70
34	BA	1280	A	C5-C6-N1	5.97	120.68	117.70
35	BB	364	U	C6-N1-C2	-5.97	117.42	121.00
35	BB	798	A	C8-N9-C4	-5.97	103.41	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	812	G	C1'-O4'-C4'	-5.97	105.13	109.90
35	BB	1485	G	C5'-C4'-C3'	-5.97	106.45	116.00
85	AA	355	G	C4-N9-C1'	-5.97	118.74	126.50
85	AA	769	C	N3-C2-O2	-5.97	117.72	121.90
85	AA	906	U	C5'-C4'-O4'	5.97	116.26	109.10
85	AA	1275	A	C5-C6-N6	5.97	128.47	123.70
34	BA	665	C	C3'-C2'-C1'	-5.96	96.73	101.50
34	BA	783	U	N3-C2-O2	-5.96	118.03	122.20
34	BA	1061	A	C8-N9-C4	5.96	108.19	105.80
34	BA	1476	G	C5'-C4'-C3'	-5.96	106.46	116.00
34	BA	1559	C	O4'-C1'-N1	5.96	112.97	108.20
34	BA	1774	G	O4'-C1'-N9	5.96	112.97	108.20
35	BB	260	A	C5'-C4'-C3'	-5.96	106.46	116.00
38	BE	18	U	O5'-C5'-C4'	-5.96	100.37	111.70
40	BG	18	U	C5'-C4'-C3'	-5.96	106.46	116.00
40	BG	85	C	C5'-C4'-C3'	5.96	125.54	116.00
42	BI	172	HIS	CA-CB-CG	5.96	123.74	113.60
80	Bu	208	ARG	NE-CZ-NH1	5.96	123.28	120.30
85	AA	276	C	C5'-C4'-O4'	-5.96	101.94	109.10
85	AA	803	C	C5-C4-N4	-5.96	116.02	120.20
85	AA	1304	C	P-O5'-C5'	-5.96	111.36	120.90
85	AA	1557	U	O3'-P-O5'	5.96	115.33	104.00
85	AA	1917	G	C8-N9-C1'	5.96	134.75	127.00
35	BB	397	C	N3-C2-O2	-5.96	117.73	121.90
38	BE	199	A	C8-N9-C4	5.96	108.19	105.80
82	Bw	62	ARG	NE-CZ-NH2	-5.96	117.32	120.30
85	AA	2227	A	N7-C8-N9	-5.96	110.82	113.80
34	BA	411	C	C6-N1-C1'	-5.96	113.65	120.80
34	BA	526	C	P-O5'-C5'	-5.96	111.36	120.90
34	BA	535	G	N3-C4-N9	5.96	129.58	126.00
34	BA	621	G	N1-C6-O6	5.96	123.48	119.90
34	BA	747	G	N3-C4-C5	-5.96	125.62	128.60
34	BA	751	A	C2-N3-C4	5.96	113.58	110.60
34	BA	1581	G	N3-C4-C5	-5.96	125.62	128.60
34	BA	1611	A	C4'-C3'-C2'	-5.96	96.64	102.60
34	BA	1711	G	N1-C2-N3	-5.96	120.32	123.90
35	BB	516	G	C4'-C3'-C2'	-5.96	96.64	102.60
35	BB	1462	G	N3-C4-C5	-5.96	125.62	128.60
37	BD	87	G	N7-C8-N9	-5.96	110.12	113.10
85	AA	338	G	N9-C1'-C2'	-5.96	105.44	112.00
85	AA	1116	G	N9-C1'-C2'	-5.96	105.44	112.00
34	BA	46	C	P-O3'-C3'	-5.96	112.55	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	986	G	C4-N9-C1'	-5.96	118.75	126.50
34	BA	1060	C	O4'-C1'-C2'	-5.96	99.84	105.80
34	BA	1491	U	OP1-P-O3'	5.96	118.31	105.20
34	BA	1640	G	C5'-C4'-C3'	5.96	125.54	116.00
35	BB	1259	A	O4'-C1'-N9	5.96	112.97	108.20
35	BB	1379	U	C5-C6-N1	-5.96	119.72	122.70
40	BG	9	G	O4'-C1'-C2'	5.96	112.96	107.60
85	AA	1137	C	O4'-C1'-N1	5.96	112.97	108.20
85	AA	1668	G	C5-C6-O6	5.96	132.18	128.60
34	BA	596	G	N3-C2-N2	5.96	124.07	119.90
34	BA	662	U	P-O5'-C5'	5.96	130.43	120.90
34	BA	1846	G	P-O3'-C3'	-5.96	112.55	119.70
35	BB	835	C	P-O3'-C3'	5.96	126.85	119.70
35	BB	981	A	N9-C1'-C2'	5.96	121.75	114.00
35	BB	993	A	N3-C4-C5	5.96	130.97	126.80
35	BB	1283	C	O4'-C1'-N1	5.96	112.97	108.20
41	BH	9	C	O5'-P-OP2	5.96	117.85	110.70
85	AA	408	C	C5-C4-N4	5.96	124.37	120.20
85	AA	976	G	O4'-C1'-N9	5.96	112.97	108.20
85	AA	1195	U	C2-N3-C4	-5.96	123.42	127.00
85	AA	1476	C	C2-N3-C4	-5.96	116.92	119.90
34	BA	57	A	C3'-C2'-C1'	-5.96	96.73	101.50
34	BA	87	G	P-O5'-C5'	-5.96	111.37	120.90
34	BA	955	G	C5-C6-N1	5.96	114.48	111.50
34	BA	1100	A	C5'-C4'-C3'	-5.96	106.47	116.00
34	BA	1219	G	C4'-C3'-C2'	-5.96	96.64	102.60
34	BA	1220	C	N1-C1'-C2'	-5.96	105.45	112.00
34	BA	1521	C	P-O3'-C3'	-5.96	112.55	119.70
34	BA	1845	G	N9-C4-C5	5.96	107.78	105.40
35	BB	300	U	O4'-C1'-N1	5.96	112.96	108.20
35	BB	968	C	O4'-C4'-C3'	-5.96	98.04	104.00
35	BB	1165	A	OP1-P-O3'	5.96	118.31	105.20
35	BB	1536	G	C8-N9-C4	-5.96	104.02	106.40
37	BD	49	A	C3'-C2'-C1'	-5.96	96.73	101.50
37	BD	60	C	C6-N1-C2	-5.96	117.92	120.30
37	BD	108	G	C8-N9-C4	-5.96	104.02	106.40
41	BH	36	C	O4'-C1'-N1	5.96	112.97	108.20
42	BI	74	ARG	NE-CZ-NH1	5.96	123.28	120.30
85	AA	107	A	C5-N7-C8	-5.96	100.92	103.90
85	AA	535	G	C5'-C4'-C3'	-5.96	106.47	116.00
85	AA	1263	G	C5'-C4'-O4'	5.96	116.25	109.10
85	AA	1519	A	N1-C6-N6	-5.96	115.03	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	24	G	C8-N9-C1'	5.96	134.74	127.00
39	BF	42	G	P-O3'-C3'	-5.96	112.55	119.70
40	BG	92	U	O4'-C1'-C2'	5.96	112.96	107.60
40	BG	111	C	P-O5'-C5'	5.96	130.43	120.90
85	AA	1839	G	C4-N9-C1'	-5.96	118.76	126.50
85	AA	2139	G	N1-C6-O6	5.96	123.47	119.90
34	BA	568	G	O3'-P-O5'	5.95	115.31	104.00
34	BA	613	A	C1'-O4'-C4'	-5.95	105.14	109.90
34	BA	635	G	C2-N3-C4	5.95	114.88	111.90
34	BA	1415	C	N3-C2-O2	-5.95	117.73	121.90
35	BB	645	C	P-O5'-C5'	5.95	130.43	120.90
35	BB	852	G	N3-C2-N2	5.95	124.07	119.90
36	BC	147	G	N1-C6-O6	5.95	123.47	119.90
36	BC	148	C	O5'-C5'-C4'	-5.95	100.39	111.70
39	BF	5	U	N1-C2-O2	5.95	126.97	122.80
41	BH	116	A	C5'-C4'-C3'	5.95	125.53	116.00
65	Bf	282	PHE	CB-CG-CD2	-5.95	116.63	120.80
85	AA	1975	G	O4'-C1'-N9	5.95	112.96	108.20
85	AA	2169	C	C6-N1-C1'	5.95	127.94	120.80
34	BA	447	U	O5'-C5'-C4'	-5.95	100.39	111.70
34	BA	1735	G	N9-C1'-C2'	-5.95	105.45	112.00
37	BD	9	C	N3-C2-O2	-5.95	117.73	121.90
40	BG	54	G	N1-C2-N2	5.95	121.56	116.20
85	AA	485	A	N1-C6-N6	5.95	122.17	118.60
85	AA	1494	C	O4'-C1'-N1	5.95	112.96	108.20
85	AA	2007	G	C8-N9-C1'	5.95	134.74	127.00
34	BA	1314	A	O4'-C1'-N9	5.95	112.96	108.20
34	BA	1432	C	N3-C4-C5	5.95	124.28	121.90
34	BA	1719	G	O4'-C1'-N9	5.95	112.96	108.20
34	BA	1793	G	C5-C6-O6	-5.95	125.03	128.60
35	BB	472	C	P-O5'-C5'	5.95	130.42	120.90
35	BB	855	G	P-O3'-C3'	-5.95	112.56	119.70
35	BB	1153	G	C5-C6-O6	-5.95	125.03	128.60
35	BB	1291	G	C8-N9-C1'	5.95	134.74	127.00
36	BC	116	C	C2-N1-C1'	-5.95	112.25	118.80
37	BD	105	G	P-O3'-C3'	-5.95	112.56	119.70
38	BE	134	A	C5'-C4'-C3'	5.95	125.52	116.00
74	Bo	45	PHE	CB-CG-CD2	-5.95	116.63	120.80
85	AA	1074	U	C5'-C4'-O4'	5.95	116.24	109.10
6	A5	75	ARG	NE-CZ-NH2	5.95	123.27	120.30
34	BA	1130	U	O4'-C1'-N1	5.95	112.96	108.20
34	BA	1211	G	N1-C2-N3	5.95	127.47	123.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1270	G	C8-N9-C4	5.95	108.78	106.40
34	BA	1673	G	C5-C6-O6	5.95	132.17	128.60
35	BB	1035	C	O4'-C1'-N1	5.95	112.96	108.20
35	BB	1097	U	O4'-C1'-N1	5.95	112.96	108.20
36	BC	88	A	C3'-C2'-C1'	-5.95	96.74	101.50
72	Bm	76	ARG	CA-C-N	-5.95	104.11	117.20
85	AA	526	G	C5-C6-N1	5.95	114.47	111.50
85	AA	1272	G	C5'-C4'-C3'	-5.95	106.48	116.00
85	AA	1547	G	O4'-C1'-N9	5.95	112.96	108.20
34	BA	113	G	P-O5'-C5'	-5.95	111.39	120.90
34	BA	575	U	O4'-C1'-N1	5.95	112.96	108.20
34	BA	645	U	N3-C4-O4	5.95	123.56	119.40
34	BA	728	A	O4'-C4'-C3'	-5.95	98.05	104.00
34	BA	772	G	C4-N9-C1'	5.95	134.23	126.50
35	BB	1029	U	N1-C1'-C2'	-5.95	105.46	112.00
35	BB	1242	C	O4'-C1'-N1	5.95	112.96	108.20
35	BB	1311	G	C3'-C2'-C1'	-5.95	96.74	101.50
35	BB	1489	A	P-O5'-C5'	5.95	130.41	120.90
40	BG	95	U	O5'-P-OP2	-5.95	100.35	105.70
85	AA	53	G	C1'-O4'-C4'	-5.95	105.14	109.90
85	AA	261	U	C2-N3-C4	-5.95	123.43	127.00
85	AA	413	G	C1'-O4'-C4'	-5.95	105.14	109.90
85	AA	982	G	P-O5'-C5'	-5.95	111.38	120.90
85	AA	1451	U	N3-C4-C5	5.95	118.17	114.60
85	AA	1712	A	N1-C6-N6	-5.95	115.03	118.60
3	A2	145	ARG	NE-CZ-NH1	5.95	123.27	120.30
34	BA	36	A	C5'-C4'-C3'	-5.95	106.49	116.00
34	BA	421	G	C1'-O4'-C4'	-5.95	105.14	109.90
34	BA	790	G	C2-N3-C4	-5.95	108.93	111.90
34	BA	888	G	P-O3'-C3'	-5.95	112.57	119.70
34	BA	1070	G	C5-C6-O6	-5.95	125.03	128.60
34	BA	1078	U	C2-N3-C4	-5.95	123.43	127.00
34	BA	1625	C	C4'-C3'-C2'	-5.95	96.66	102.60
34	BA	1660	A	O4'-C1'-N9	5.95	112.96	108.20
35	BB	591	A	C6-N1-C2	-5.95	115.03	118.60
35	BB	1251	G	C8-N9-C1'	5.95	134.73	127.00
36	BC	6	G	P-O5'-C5'	-5.95	111.39	120.90
38	BE	96	G	P-O3'-C3'	-5.95	112.56	119.70
38	BE	100	U	P-O3'-C3'	5.95	126.83	119.70
38	BE	123	A	C5-C6-N6	-5.95	118.94	123.70
38	BE	135	A	O3'-P-O5'	5.95	115.30	104.00
77	Br	200	TYR	CB-CG-CD2	-5.95	117.43	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	737	G	P-O5'-C5'	5.95	130.41	120.90
85	AA	758	C	C6-N1-C1'	5.95	127.93	120.80
85	AA	1113	G	P-O3'-C3'	-5.95	112.56	119.70
85	AA	1375	U	C6-N1-C2	-5.95	117.43	121.00
34	BA	181	G	C1'-O4'-C4'	-5.94	105.14	109.90
34	BA	1251	A	C5-C6-N1	5.94	120.67	117.70
35	BB	1165	A	O4'-C1'-C2'	5.94	112.95	107.60
37	BD	48	G	C4'-C3'-O3'	5.94	124.89	113.00
82	Bw	82	ARG	N-CA-CB	-5.94	99.90	110.60
85	AA	594	C	C6-N1-C2	5.94	122.68	120.30
11	AC	222	ARG	NE-CZ-NH1	5.94	123.27	120.30
34	BA	31	A	N1-C6-N6	-5.94	115.03	118.60
34	BA	1564	A	C4-N9-C1'	-5.94	115.60	126.30
34	BA	1732	A	C4-N9-C1'	5.94	137.00	126.30
35	BB	43	G	O4'-C1'-N9	5.94	112.95	108.20
35	BB	70	A	O3'-P-O5'	5.94	115.29	104.00
35	BB	103	C	N3-C2-O2	-5.94	117.74	121.90
35	BB	647	U	C2-N3-C4	-5.94	123.43	127.00
38	BE	174	U	N1-C2-N3	5.94	118.47	114.90
41	BH	61	C	C3'-C2'-C1'	-5.94	96.75	101.50
79	Bt	28	TYR	CB-CG-CD1	-5.94	117.44	121.00
85	AA	342	C	C5'-C4'-O4'	5.94	116.23	109.10
85	AA	470	C	C1'-O4'-C4'	-5.94	105.15	109.90
85	AA	710	A	C5'-C4'-O4'	-5.94	101.97	109.10
85	AA	1842	C	C4'-C3'-C2'	5.94	108.54	102.60
85	AA	2166	G	C4'-C3'-C2'	-5.94	96.66	102.60
85	AA	2217	A	N9-C1'-C2'	-5.94	105.46	112.00
34	BA	207	A	C8-N9-C4	5.94	108.18	105.80
34	BA	224	G	O3'-P-O5'	-5.94	92.71	104.00
34	BA	721	A	P-O3'-C3'	-5.94	112.57	119.70
34	BA	1483	U	O4'-C1'-C2'	5.94	112.95	107.60
34	BA	1594	G	O4'-C1'-N9	5.94	112.95	108.20
35	BB	166	C	O4'-C1'-N1	5.94	112.95	108.20
35	BB	520	G	N1-C6-O6	5.94	123.46	119.90
35	BB	562	A	O4'-C1'-N9	5.94	112.95	108.20
35	BB	631	G	C5-N7-C8	-5.94	101.33	104.30
36	BC	36	G	C3'-C2'-C1'	-5.94	96.75	101.50
36	BC	132	U	O3'-P-O5'	5.94	115.29	104.00
41	BH	39	G	O5'-P-OP1	5.94	117.83	110.70
85	AA	198	U	C6-N1-C2	-5.94	117.44	121.00
85	AA	514	U	N3-C4-C5	5.94	118.17	114.60
85	AA	1172	A	C5-C6-N6	-5.94	118.95	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1773	U	O4'-C1'-N1	5.94	112.95	108.20
31	AX	44	ARG	NE-CZ-NH1	5.94	123.27	120.30
34	BA	1418	G	C4-N9-C1'	-5.94	118.78	126.50
34	BA	1499	A	C6-N1-C2	-5.94	115.04	118.60
34	BA	1507	C	C5'-C4'-C3'	5.94	125.50	116.00
35	BB	1011	C	C6-N1-C2	-5.94	117.92	120.30
35	BB	1026	G	C4-N9-C1'	-5.94	118.78	126.50
40	BG	65	C	C6-N1-C2	-5.94	117.92	120.30
40	BG	176	G	P-O3'-C3'	-5.94	112.57	119.70
41	BH	6	U	O4'-C1'-N1	5.94	112.95	108.20
85	AA	1640	G	N1-C6-O6	5.94	123.46	119.90
85	AA	1671	G	C5'-C4'-C3'	-5.94	106.50	116.00
34	BA	99	G	N1-C2-N3	5.94	127.46	123.90
34	BA	826	C	C6-N1-C2	-5.94	117.92	120.30
34	BA	1205	A	C4'-C3'-C2'	-5.94	96.66	102.60
34	BA	1587	C	C5'-C4'-C3'	-5.94	106.50	116.00
34	BA	1776	G	C5'-C4'-C3'	5.94	125.50	116.00
35	BB	698	C	C6-N1-C2	-5.94	117.92	120.30
35	BB	775	U	O4'-C1'-C2'	5.94	112.94	107.60
35	BB	1335	G	C4-N9-C1'	-5.94	118.78	126.50
35	BB	1389	C	C5'-C4'-C3'	-5.94	106.50	116.00
35	BB	1536	G	C3'-C2'-C1'	-5.94	96.75	101.50
85	AA	749	C	N3-C4-N4	-5.94	113.84	118.00
85	AA	1487	G	N7-C8-N9	-5.94	110.13	113.10
85	AA	1932	C	C5'-C4'-C3'	5.94	125.50	116.00
85	AA	2158	U	O4'-C1'-N1	5.94	112.95	108.20
1	A0	138	ARG	NE-CZ-NH2	-5.94	117.33	120.30
8	A7	34	THR	CA-CB-CG2	-5.94	104.09	112.40
34	BA	144	C	C5'-C4'-C3'	-5.94	106.50	116.00
35	BB	1202	G	N1-C6-O6	5.94	123.46	119.90
36	BC	95	A	P-O5'-C5'	5.94	130.40	120.90
36	BC	118	U	N3-C2-O2	-5.94	118.05	122.20
38	BE	23	G	O4'-C4'-C3'	-5.94	98.06	104.00
41	BH	41	A	C8-N9-C1'	5.94	138.38	127.70
85	AA	1029	G	O4'-C1'-N9	5.94	112.95	108.20
85	AA	1644	G	N1-C2-N2	-5.94	110.86	116.20
34	BA	734	G	C4'-C3'-C2'	-5.93	96.67	102.60
35	BB	380	G	N9-C4-C5	5.93	107.77	105.40
35	BB	880	G	P-O5'-C5'	5.93	130.39	120.90
35	BB	1371	G	C4-N9-C1'	-5.93	118.78	126.50
35	BB	1452	U	O5'-P-OP2	-5.93	100.36	105.70
35	BB	1505	U	P-O3'-C3'	-5.93	112.58	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	39	G	P-O3'-C3'	-5.93	112.58	119.70
38	BE	136	G	O4'-C1'-C2'	5.93	112.94	107.60
85	AA	71	G	N1-C6-O6	-5.93	116.34	119.90
85	AA	708	G	C8-N9-C1'	5.93	134.72	127.00
85	AA	1478	G	C5'-C4'-C3'	-5.93	106.51	116.00
85	AA	2113	U	C4'-C3'-C2'	-5.93	96.67	102.60
30	AW	5	ASP	N-CA-CB	-5.93	99.92	110.60
34	BA	15	G	N9-C4-C5	-5.93	103.03	105.40
34	BA	499	C	C5'-C4'-C3'	5.93	125.49	116.00
34	BA	665	C	N1-C2-N3	5.93	123.35	119.20
34	BA	1646	U	C6-N1-C2	-5.93	117.44	121.00
35	BB	983	C	C6-N1-C1'	-5.93	113.68	120.80
35	BB	1202	G	N9-C4-C5	-5.93	103.03	105.40
40	BG	5	G	N3-C2-N2	5.93	124.05	119.90
40	BG	59	G	P-O3'-C3'	-5.93	112.58	119.70
40	BG	75	C	P-O3'-C3'	5.93	126.82	119.70
40	BG	116	G	C4-N9-C1'	-5.93	118.79	126.50
85	AA	585	G	C5-C6-N1	5.93	114.47	111.50
85	AA	784	C	C5-C6-N1	5.93	123.97	121.00
85	AA	1214	C	P-O5'-C5'	5.93	130.39	120.90
85	AA	1415	G	C4'-C3'-C2'	5.93	108.53	102.60
85	AA	1484	G	C5-C6-N1	5.93	114.47	111.50
85	AA	1921	G	N3-C4-C5	-5.93	125.63	128.60
86	AB	45	U	O4'-C1'-N1	5.93	112.95	108.20
34	BA	768	G	N9-C4-C5	-5.93	103.03	105.40
34	BA	1600	G	C4-N9-C1'	-5.93	118.79	126.50
35	BB	625	A	C1'-O4'-C4'	-5.93	105.16	109.90
35	BB	1022	C	N1-C1'-C2'	-5.93	105.48	112.00
35	BB	1145	G	C5-N7-C8	-5.93	101.33	104.30
35	BB	1369	A	C6-N1-C2	-5.93	115.04	118.60
36	BC	17	U	C6-N1-C1'	-5.93	112.90	121.20
53	BT	52	ARG	NE-CZ-NH2	-5.93	117.33	120.30
85	AA	1855	U	C5'-C4'-C3'	-5.93	106.51	116.00
3	A2	80	ARG	NE-CZ-NH1	5.93	123.27	120.30
34	BA	832	C	C5'-C4'-C3'	-5.93	106.51	116.00
34	BA	1232	C	C6-N1-C2	-5.93	117.93	120.30
35	BB	558	U	C5'-C4'-C3'	-5.93	106.51	116.00
35	BB	1316	U	O4'-C1'-N1	5.93	112.94	108.20
35	BB	1334	C	N1-C1'-C2'	-5.93	105.48	112.00
36	BC	74	U	C2-N3-C4	-5.93	123.44	127.00
38	BE	180	G	OP2-P-O3'	5.93	118.25	105.20
38	BE	185	G	O4'-C1'-N9	5.93	112.94	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	BN	54	ARG	NE-CZ-NH1	5.93	123.27	120.30
81	Bv	6	LYS	C-N-CA	5.93	136.52	121.70
85	AA	20	G	N3-C2-N2	-5.93	115.75	119.90
85	AA	32	U	C4-C5-C6	-5.93	116.14	119.70
85	AA	552	C	C5-C4-N4	-5.93	116.05	120.20
85	AA	920	A	C5'-C4'-C3'	5.93	125.49	116.00
34	BA	12	G	N3-C2-N2	-5.93	115.75	119.90
34	BA	133	A	C5'-C4'-C3'	5.93	125.49	116.00
34	BA	1525	G	O4'-C4'-C3'	-5.93	98.07	104.00
34	BA	1808	A	C5'-C4'-O4'	5.93	116.21	109.10
35	BB	438	G	C3'-C2'-C1'	-5.93	96.76	101.50
35	BB	778	A	O5'-C5'-C4'	-5.93	100.44	111.70
35	BB	1202	G	C8-N9-C1'	-5.93	119.29	127.00
35	BB	1444	U	O4'-C1'-N1	5.93	112.94	108.20
41	BH	9	C	N3-C2-O2	-5.93	117.75	121.90
41	BH	124	C	C3'-C2'-C1'	-5.93	96.76	101.50
85	AA	273	C	C2-N1-C1'	-5.93	112.28	118.80
85	AA	1313	C	O4'-C1'-N1	5.93	112.94	108.20
86	AB	62	C	C2-N1-C1'	-5.93	112.28	118.80
34	BA	47	U	N3-C2-O2	-5.93	118.05	122.20
34	BA	1495	A	O4'-C4'-C3'	-5.93	98.07	104.00
35	BB	590	G	N1-C2-N2	-5.93	110.87	116.20
35	BB	728	A	N9-C4-C5	5.93	108.17	105.80
35	BB	1099	U	O3'-P-O5'	-5.93	92.74	104.00
35	BB	1196	A	C2-N3-C4	-5.93	107.64	110.60
37	BD	11	A	N1-C6-N6	-5.93	115.05	118.60
38	BE	74	U	C5'-C4'-C3'	-5.93	106.52	116.00
65	Bf	459	ARG	NE-CZ-NH1	5.93	123.26	120.30
85	AA	493	A	C3'-C2'-C1'	-5.93	96.76	101.50
85	AA	735	G	C6-N1-C2	-5.93	121.54	125.10
85	AA	1246	G	C6-N1-C2	-5.93	121.54	125.10
85	AA	1987	G	N1-C6-O6	5.93	123.46	119.90
34	BA	101	G	N1-C2-N2	-5.92	110.87	116.20
34	BA	209	A	C4-N9-C1'	-5.92	115.63	126.30
34	BA	903	C	C6-N1-C2	-5.92	117.93	120.30
34	BA	1061	A	C1'-O4'-C4'	-5.92	105.16	109.90
34	BA	1454	G	N3-C2-N2	-5.92	115.75	119.90
34	BA	1739	G	O4'-C1'-N9	5.92	112.94	108.20
34	BA	1782	C	N1-C2-O2	5.92	122.45	118.90
35	BB	1026	G	N9-C4-C5	-5.92	103.03	105.40
35	BB	1078	U	P-O5'-C5'	-5.92	111.42	120.90
35	BB	1152	U	C5-C6-N1	-5.92	119.74	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	24	G	C5-C6-O6	-5.92	125.05	128.60
38	BE	28	C	C1'-O4'-C4'	-5.92	105.16	109.90
40	BG	138	C	C6-N1-C2	-5.92	117.93	120.30
41	BH	111	U	P-O5'-C5'	5.92	130.38	120.90
85	AA	14	C	N3-C2-O2	-5.92	117.75	121.90
85	AA	561	C	C6-N1-C2	-5.92	117.93	120.30
85	AA	687	G	N3-C4-N9	5.92	129.55	126.00
85	AA	836	A	C1'-O4'-C4'	-5.92	105.16	109.90
85	AA	841	U	O4'-C1'-N1	5.92	112.94	108.20
85	AA	917	A	O4'-C1'-N9	5.92	112.94	108.20
85	AA	1185	G	N9-C1'-C2'	-5.92	105.48	112.00
85	AA	1211	C	N1-C2-N3	5.92	123.35	119.20
85	AA	1289	U	N3-C2-O2	-5.92	118.05	122.20
85	AA	1298	G	C8-N9-C1'	5.92	134.70	127.00
85	AA	1665	G	P-O3'-C3'	-5.92	112.59	119.70
85	AA	1729	C	C1'-O4'-C4'	-5.92	105.16	109.90
85	AA	1998	A	C5-C6-N6	-5.92	118.96	123.70
85	AA	2054	G	C8-N9-C1'	5.92	134.70	127.00
85	AA	2137	A	N1-C6-N6	5.92	122.16	118.60
86	AB	15	G	O3'-P-O5'	5.92	115.25	104.00
34	BA	49	A	N9-C1'-C2'	-5.92	105.48	112.00
35	BB	17	U	O5'-C5'-C4'	-5.92	100.45	111.70
35	BB	1432	U	N3-C2-O2	-5.92	118.05	122.20
80	Bu	194	ASP	N-CA-CB	-5.92	99.94	110.60
85	AA	714	U	O4'-C1'-N1	5.92	112.94	108.20
85	AA	1145	U	O4'-C1'-N1	5.92	112.94	108.20
85	AA	1751	G	C2'-C3'-O3'	5.92	123.18	113.70
6	A5	121	LEU	CB-CA-C	-5.92	98.95	110.20
11	AC	77	ARG	NE-CZ-NH1	5.92	123.26	120.30
34	BA	14	G	C4-C5-C6	-5.92	115.25	118.80
34	BA	210	G	C5'-C4'-O4'	-5.92	102.00	109.10
34	BA	322	U	C1'-O4'-C4'	-5.92	105.16	109.90
34	BA	671	C	N3-C2-O2	-5.92	117.75	121.90
34	BA	888	G	O4'-C1'-C2'	5.92	112.93	107.60
34	BA	938	C	C2-N3-C4	-5.92	116.94	119.90
35	BB	513	G	C8-N9-C1'	5.92	134.70	127.00
35	BB	529	A	C5'-C4'-C3'	5.92	125.47	116.00
35	BB	546	A	C2-N3-C4	5.92	113.56	110.60
35	BB	599	U	N1-C1'-C2'	-5.92	105.49	112.00
35	BB	626	C	C5'-C4'-C3'	-5.92	106.52	116.00
36	BC	12	A	C5-C6-N6	-5.92	118.96	123.70
36	BC	87	C	P-O3'-C3'	-5.92	112.59	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	20	C	C6-N1-C2	5.92	122.67	120.30
38	BE	206	G	C4-N9-C1'	-5.92	118.80	126.50
40	BG	90	G	C5-C6-N1	5.92	114.46	111.50
65	Bf	405	SER	N-CA-C	5.92	126.99	111.00
77	Br	303	LYS	C-N-CA	5.92	136.50	121.70
85	AA	28	A	C8-N9-C4	5.92	108.17	105.80
85	AA	682	C	C5'-C4'-O4'	5.92	116.21	109.10
85	AA	748	C	C1'-O4'-C4'	-5.92	105.16	109.90
85	AA	982	G	N7-C8-N9	-5.92	110.14	113.10
85	AA	1466	U	C5-C6-N1	-5.92	119.74	122.70
34	BA	779	U	C6-N1-C2	-5.92	117.45	121.00
34	BA	1091	U	C2-N1-C1'	-5.92	110.59	117.70
35	BB	104	G	C1'-O4'-C4'	-5.92	105.16	109.90
35	BB	1315	C	N3-C4-C5	5.92	124.27	121.90
38	BE	168	C	N3-C2-O2	-5.92	117.76	121.90
85	AA	97	A	N7-C8-N9	-5.92	110.84	113.80
34	BA	647	U	OP1-P-O3'	5.92	118.22	105.20
34	BA	888	G	O4'-C1'-N9	5.92	112.94	108.20
34	BA	1057	C	P-O3'-C3'	5.92	126.80	119.70
34	BA	1398	C	O4'-C1'-N1	5.92	112.93	108.20
34	BA	1846	G	O4'-C1'-N9	5.92	112.94	108.20
35	BB	35	G	C5-C6-N1	5.92	114.46	111.50
35	BB	963	G	O4'-C1'-N9	5.92	112.94	108.20
38	BE	23	G	P-O3'-C3'	-5.92	112.60	119.70
38	BE	34	C	P-O3'-C3'	-5.92	112.60	119.70
40	BG	154	C	P-O3'-C3'	-5.92	112.60	119.70
62	Bc	97	ASP	CB-CA-C	5.92	122.24	110.40
85	AA	90	A	C5'-C4'-C3'	-5.92	106.53	116.00
85	AA	1023	U	C6-N1-C2	-5.92	117.45	121.00
85	AA	1728	G	C5-C6-O6	-5.92	125.05	128.60
34	BA	1488	C	C2'-C3'-O3'	5.92	123.17	113.70
35	BB	1523	U	C2-N1-C1'	-5.92	110.60	117.70
38	BE	37	C	C5-C6-N1	5.92	123.96	121.00
40	BG	106	G	C2-N3-C4	-5.92	108.94	111.90
65	Bf	113	ARG	NE-CZ-NH1	5.92	123.26	120.30
70	Bk	59	MET	CA-CB-CG	5.92	123.36	113.30
85	AA	1970	A	C8-N9-C1'	5.92	138.35	127.70
34	BA	115	U	O5'-C5'-C4'	-5.92	100.46	111.70
34	BA	306	G	C4-N9-C1'	-5.92	118.81	126.50
34	BA	1226	G	P-O5'-C5'	-5.92	111.44	120.90
34	BA	1680	G	C4'-C3'-C2'	5.92	108.52	102.60
35	BB	706	G	C3'-C2'-C1'	-5.92	96.77	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	767	A	P-O5'-C5'	-5.92	111.44	120.90
37	BD	75	G	N1-C2-N2	5.92	121.52	116.20
37	BD	83	A	N9-C1'-C2'	5.92	121.69	114.00
59	BZ	7	ARG	NE-CZ-NH1	5.92	123.26	120.30
85	AA	283	A	P-O5'-C5'	5.92	130.36	120.90
85	AA	475	A	N1-C6-N6	-5.92	115.05	118.60
85	AA	1551	G	C4-N9-C1'	5.92	134.19	126.50
85	AA	1583	U	P-O3'-C3'	5.92	126.80	119.70
85	AA	2169	C	C2-N1-C1'	-5.92	112.29	118.80
34	BA	237	A	C4-N9-C1'	-5.91	115.66	126.30
34	BA	271	C	C2-N1-C1'	5.91	125.30	118.80
34	BA	840	U	C3'-C2'-C1'	-5.91	96.77	101.50
34	BA	1147	C	C2'-C3'-O3'	5.91	123.16	113.70
34	BA	1378	A	P-O5'-C5'	-5.91	111.44	120.90
35	BB	428	G	O4'-C4'-C3'	-5.91	98.09	104.00
35	BB	796	C	C5'-C4'-O4'	5.91	116.20	109.10
35	BB	897	C	C5'-C4'-O4'	5.91	116.20	109.10
35	BB	1018	U	O4'-C1'-N1	5.91	112.93	108.20
36	BC	37	U	C6-N1-C1'	-5.91	112.92	121.20
36	BC	141	C	C1'-O4'-C4'	-5.91	105.17	109.90
40	BG	47	G	O4'-C1'-N9	5.91	112.93	108.20
40	BG	139	U	C1'-O4'-C4'	-5.91	105.17	109.90
40	BG	180	C	P-O5'-C5'	-5.91	111.44	120.90
41	BH	60	A	P-O3'-C3'	5.91	126.80	119.70
77	Br	113	LYS	CB-CA-C	-5.91	98.57	110.40
83	Bx	61	ARG	NE-CZ-NH1	5.91	123.26	120.30
85	AA	426	C	N1-C2-O2	5.91	122.45	118.90
85	AA	1272	G	P-O3'-C3'	-5.91	112.60	119.70
85	AA	2208	G	C8-N9-C1'	5.91	134.69	127.00
17	AI	122	TYR	CB-CG-CD1	-5.91	117.45	121.00
34	BA	426	A	OP1-P-OP2	-5.91	110.73	119.60
34	BA	738	C	C6-N1-C2	-5.91	117.94	120.30
38	BE	169	C	P-O3'-C3'	5.91	126.80	119.70
40	BG	112	C	C6-N1-C1'	5.91	127.89	120.80
42	BI	39	ARG	NE-CZ-NH1	5.91	123.26	120.30
70	Bk	59	MET	CG-SD-CE	-5.91	90.74	100.20
4	A3	95	ARG	N-CA-C	-5.91	95.04	111.00
34	BA	180	G	O4'-C1'-N9	5.91	112.93	108.20
34	BA	619	U	O4'-C1'-N1	5.91	112.93	108.20
34	BA	871	G	P-O5'-C5'	-5.91	111.44	120.90
34	BA	1060	C	C1'-O4'-C4'	-5.91	105.17	109.90
34	BA	1262	A	O5'-C5'-C4'	-5.91	100.47	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	139	G	C8-N9-C4	5.91	108.76	106.40
35	BB	273	G	O4'-C1'-N9	5.91	112.93	108.20
35	BB	1459	U	P-O3'-C3'	-5.91	112.61	119.70
38	BE	140	G	N1-C6-O6	-5.91	116.35	119.90
39	BF	71	G	P-O3'-C3'	-5.91	112.61	119.70
49	BP	167	ALA	N-CA-CB	5.91	118.38	110.10
85	AA	130	G	C5'-C4'-C3'	-5.91	106.54	116.00
85	AA	996	A	P-O5'-C5'	-5.91	111.44	120.90
85	AA	1277	C	N3-C4-N4	5.91	122.14	118.00
85	AA	1560	A	C2'-C3'-O3'	5.91	123.16	113.70
85	AA	1794	U	N3-C2-O2	-5.91	118.06	122.20
34	BA	296	G	C4-C5-C6	-5.91	115.25	118.80
34	BA	530	A	C4'-C3'-C2'	5.91	108.51	102.60
34	BA	628	U	N3-C2-O2	-5.91	118.06	122.20
34	BA	669	U	O4'-C4'-C3'	-5.91	98.09	104.00
34	BA	921	G	C5'-C4'-C3'	-5.91	106.55	116.00
34	BA	1112	U	C2-N1-C1'	-5.91	110.61	117.70
34	BA	1404	A	C4-N9-C1'	-5.91	115.66	126.30
34	BA	1839	G	C5'-C4'-C3'	-5.91	106.55	116.00
35	BB	742	G	C8-N9-C4	-5.91	104.04	106.40
35	BB	1187	G	N3-C4-C5	-5.91	125.65	128.60
85	AA	362	G	C8-N9-C4	5.91	108.76	106.40
85	AA	398	U	C5'-C4'-C3'	5.91	125.45	116.00
85	AA	744	C	O4'-C1'-N1	5.91	112.93	108.20
85	AA	982	G	N3-C4-C5	5.91	131.55	128.60
85	AA	1524	A	N1-C6-N6	5.91	122.14	118.60
85	AA	1658	G	C6-N1-C2	-5.91	121.55	125.10
85	AA	2017	U	C5'-C4'-C3'	-5.91	106.55	116.00
85	AA	2239	A	C4'-C3'-C2'	-5.91	96.69	102.60
34	BA	160	G	C4'-C3'-C2'	-5.91	96.69	102.60
34	BA	633	G	C1'-O4'-C4'	-5.91	105.17	109.90
34	BA	1383	U	P-O5'-C5'	-5.91	111.45	120.90
35	BB	1254	G	C1'-O4'-C4'	5.91	114.63	109.90
41	BH	101	A	C5-C6-N1	5.91	120.65	117.70
85	AA	250	C	C2-N1-C1'	-5.91	112.30	118.80
85	AA	2085	C	N1-C1'-C2'	-5.91	105.50	112.00
34	BA	904	G	O4'-C1'-C2'	5.91	112.92	107.60
34	BA	1244	G	P-O3'-C3'	-5.91	112.61	119.70
34	BA	1613	G	C4-C5-N7	5.91	113.16	110.80
34	BA	1667	G	P-O5'-C5'	-5.91	111.45	120.90
35	BB	638	G	O5'-C5'-C4'	-5.91	100.48	111.70
35	BB	1183	U	C2-N3-C4	-5.91	123.46	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	BD	116	C	C5'-C4'-C3'	-5.91	106.55	116.00
38	BE	58	U	O3'-P-O5'	-5.91	92.78	104.00
38	BE	114	G	O4'-C1'-N9	5.91	112.92	108.20
38	BE	177	U	C2-N3-C4	5.91	130.54	127.00
40	BG	86	U	O4'-C1'-N1	5.91	112.92	108.20
14	AF	59	ARG	NE-CZ-NH1	5.90	123.25	120.30
34	BA	184	C	C4'-C3'-C2'	-5.90	96.70	102.60
34	BA	608	G	P-O5'-C5'	-5.90	111.45	120.90
34	BA	1562	G	N1-C6-O6	-5.90	116.36	119.90
35	BB	1024	G	C5-C6-O6	-5.90	125.06	128.60
35	BB	1168	G	C3'-C2'-C1'	-5.90	96.78	101.50
40	BG	76	C	C6-N1-C1'	-5.90	113.72	120.80
40	BG	162	A	C2'-C3'-O3'	5.90	123.15	113.70
85	AA	1677	A	O4'-C1'-N9	5.90	112.92	108.20
34	BA	492	G	C4'-C3'-C2'	5.90	108.50	102.60
34	BA	1043	C	N3-C2-O2	-5.90	117.77	121.90
34	BA	1172	C	N1-C2-O2	5.90	122.44	118.90
34	BA	1309	U	C2-N1-C1'	5.90	124.78	117.70
34	BA	1462	U	P-O5'-C5'	-5.90	111.46	120.90
35	BB	681	G	C3'-C2'-C1'	-5.90	96.78	101.50
37	BD	17	G	C1'-O4'-C4'	-5.90	105.18	109.90
40	BG	10	U	N1-C1'-C2'	-5.90	105.51	112.00
40	BG	41	U	C5'-C4'-C3'	-5.90	106.56	116.00
41	BH	108	U	C5-C4-O4	-5.90	122.36	125.90
85	AA	216	U	C2-N1-C1'	5.90	124.78	117.70
85	AA	388	G	N1-C6-O6	5.90	123.44	119.90
85	AA	449	G	O4'-C1'-N9	5.90	112.92	108.20
85	AA	981	A	C4'-C3'-C2'	-5.90	96.70	102.60
85	AA	1432	C	C6-N1-C1'	-5.90	113.72	120.80
85	AA	1488	G	N9-C1'-C2'	-5.90	105.51	112.00
85	AA	1566	A	P-O3'-C3'	-5.90	112.62	119.70
34	BA	247	U	P-O5'-C5'	5.90	130.34	120.90
34	BA	688	G	C5-C6-N1	5.90	114.45	111.50
34	BA	1211	G	N3-C4-N9	-5.90	122.46	126.00
34	BA	1266	A	P-O5'-C5'	5.90	130.34	120.90
35	BB	368	C	P-O3'-C3'	-5.90	112.62	119.70
35	BB	1524	G	N9-C1'-C2'	-5.90	105.51	112.00
36	BC	169	G	N3-C2-N2	-5.90	115.77	119.90
39	BF	34	C	N1-C1'-C2'	-5.90	105.51	112.00
40	BG	24	A	N3-C4-N9	-5.90	122.68	127.40
75	Bp	28	ASN	N-CA-C	-5.90	95.07	111.00
85	AA	122	A	O4'-C4'-C3'	-5.90	98.10	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	306	C	N3-C4-N4	5.90	122.13	118.00
85	AA	338	G	O4'-C1'-N9	5.90	112.92	108.20
85	AA	2037	A	C5'-C4'-C3'	-5.90	106.56	116.00
85	AA	2116	U	C2-N3-C4	-5.90	123.46	127.00
85	AA	2241	C	N3-C2-O2	-5.90	117.77	121.90
86	AB	52	G	C3'-C2'-C1'	-5.90	96.78	101.50
34	BA	202	A	C8-N9-C4	5.90	108.16	105.80
34	BA	586	G	C4'-C3'-C2'	-5.90	96.70	102.60
34	BA	668	G	C6-N1-C2	-5.90	121.56	125.10
34	BA	1070	G	O5'-P-OP2	-5.90	100.39	105.70
34	BA	1094	U	N3-C2-O2	-5.90	118.07	122.20
34	BA	1451	A	C6-N1-C2	-5.90	115.06	118.60
34	BA	1554	C	P-O5'-C5'	-5.90	111.46	120.90
34	BA	1778	U	C6-N1-C2	-5.90	117.46	121.00
34	BA	1818	A	N1-C6-N6	-5.90	115.06	118.60
35	BB	553	U	C2-N3-C4	-5.90	123.46	127.00
35	BB	1540	U	C5'-C4'-C3'	5.90	125.44	116.00
37	BD	100	A	C5-N7-C8	-5.90	100.95	103.90
41	BH	127	A	OP1-P-O3'	5.90	118.18	105.20
85	AA	1355	U	C6-N1-C1'	5.90	129.46	121.20
85	AA	1586	C	C1'-O4'-C4'	-5.90	105.18	109.90
8	A7	34	THR	N-CA-CB	5.90	121.51	110.30
17	AI	133	VAL	CA-CB-CG1	5.90	119.75	110.90
34	BA	590	U	C6-N1-C1'	5.90	129.46	121.20
34	BA	701	G	C3'-C2'-C1'	-5.90	96.78	101.50
34	BA	760	G	O3'-P-O5'	-5.90	92.80	104.00
34	BA	1673	G	N1-C2-N2	-5.90	110.89	116.20
35	BB	153	G	C5-C6-O6	-5.90	125.06	128.60
35	BB	610	U	N3-C2-O2	-5.90	118.07	122.20
35	BB	1222	A	C2-N3-C4	-5.90	107.65	110.60
35	BB	1246	C	C5'-C4'-C3'	-5.90	106.56	116.00
35	BB	1290	C	O4'-C1'-N1	5.90	112.92	108.20
35	BB	1546	C	C1'-O4'-C4'	-5.90	105.18	109.90
36	BC	140	U	C4'-C3'-C2'	5.90	108.50	102.60
85	AA	734	C	C6-N1-C2	-5.90	117.94	120.30
85	AA	767	A	O4'-C4'-C3'	-5.90	98.10	104.00
85	AA	1657	C	C2-N1-C1'	-5.90	112.31	118.80
85	AA	2100	A	C5-C6-N1	5.90	120.65	117.70
11	AC	211	TYR	N-CA-CB	-5.90	99.99	110.60
34	BA	1486	U	N3-C4-C5	5.90	118.14	114.60
35	BB	545	C	C6-N1-C1'	-5.90	113.72	120.80
38	BE	95	G	C8-N9-C1'	5.90	134.66	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	569	A	C8-N9-C4	-5.90	103.44	105.80
85	AA	1028	C	O4'-C1'-N1	5.90	112.92	108.20
85	AA	2110	U	N1-C1'-C2'	-5.90	105.51	112.00
34	BA	162	G	C8-N9-C4	5.89	108.76	106.40
34	BA	838	U	O5'-C5'-C4'	-5.89	100.50	111.70
34	BA	1134	A	P-O3'-C3'	-5.89	112.63	119.70
34	BA	1153	C	O4'-C1'-N1	5.89	112.92	108.20
34	BA	1166	A	C8-N9-C1'	5.89	138.31	127.70
38	BE	73	A	C5-C6-N6	-5.89	118.98	123.70
38	BE	182	U	C5'-C4'-O4'	5.89	116.17	109.10
39	BF	9	C	C2-N3-C4	-5.89	116.95	119.90
52	BS	69	ARG	NE-CZ-NH2	-5.89	117.35	120.30
61	Bb	61	TYR	CB-CG-CD2	-5.89	117.46	121.00
67	Bh	38	TRP	CA-CB-CG	5.89	124.90	113.70
85	AA	95	U	C1'-O4'-C4'	-5.89	105.18	109.90
85	AA	113	U	O4'-C1'-N1	5.89	112.92	108.20
85	AA	120	C	N1-C1'-C2'	-5.89	105.52	112.00
85	AA	148	G	N3-C2-N2	5.89	124.03	119.90
85	AA	442	G	C2-N3-C4	5.89	114.85	111.90
85	AA	517	A	C5-N7-C8	-5.89	100.95	103.90
85	AA	749	C	P-O5'-C5'	5.89	130.33	120.90
85	AA	1354	A	C4'-C3'-C2'	-5.89	96.71	102.60
85	AA	1822	G	N3-C2-N2	5.89	124.03	119.90
85	AA	2148	C	O4'-C1'-N1	5.89	112.92	108.20
25	AR	7	TYR	N-CA-C	-5.89	95.09	111.00
34	BA	77	C	N3-C2-O2	-5.89	117.78	121.90
34	BA	851	C	C6-N1-C2	-5.89	117.94	120.30
34	BA	1411	C	N3-C2-O2	-5.89	117.78	121.90
34	BA	1667	G	C4-N9-C1'	-5.89	118.84	126.50
35	BB	458	U	N3-C2-O2	-5.89	118.08	122.20
35	BB	522	A	O4'-C1'-N9	-5.89	103.49	108.20
35	BB	589	U	P-O3'-C3'	-5.89	112.63	119.70
35	BB	799	A	C5-C6-N1	5.89	120.65	117.70
35	BB	1021	C	P-O3'-C3'	-5.89	112.63	119.70
35	BB	1467	A	N9-C1'-C2'	-5.89	105.52	112.00
40	BG	164	U	C5'-C4'-O4'	5.89	116.17	109.10
85	AA	15	U	C6-N1-C2	-5.89	117.46	121.00
85	AA	23	G	C1'-O4'-C4'	-5.89	105.19	109.90
85	AA	542	G	O4'-C1'-N9	5.89	112.91	108.20
85	AA	869	A	N1-C6-N6	5.89	122.14	118.60
34	BA	36	A	C8-N9-C1'	5.89	138.30	127.70
34	BA	203	U	P-O3'-C3'	-5.89	112.63	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	211	C	N1-C2-N3	5.89	123.32	119.20
34	BA	480	G	N3-C4-C5	5.89	131.54	128.60
34	BA	863	G	OP1-P-OP2	-5.89	110.76	119.60
34	BA	1246	G	P-O5'-C5'	-5.89	111.47	120.90
34	BA	1729	G	C4'-C3'-C2'	-5.89	96.71	102.60
35	BB	778	A	N1-C2-N3	-5.89	126.36	129.30
38	BE	107	U	N3-C2-O2	5.89	126.32	122.20
41	BH	31	A	O4'-C1'-C2'	5.89	112.90	107.60
85	AA	1797	U	N3-C4-C5	5.89	118.14	114.60
86	AB	53	G	C1'-O4'-C4'	-5.89	105.19	109.90
34	BA	23	A	C6-N1-C2	-5.89	115.07	118.60
34	BA	165	C	C4-C5-C6	5.89	120.34	117.40
34	BA	277	A	C1'-O4'-C4'	-5.89	105.19	109.90
34	BA	290	G	O5'-C5'-C4'	-5.89	100.51	111.70
34	BA	1314	A	N1-C6-N6	5.89	122.13	118.60
35	BB	1290	C	O3'-P-O5'	5.89	115.19	104.00
35	BB	1515	C	N1-C2-N3	5.89	123.32	119.20
36	BC	81	U	O5'-C5'-C4'	-5.89	100.51	111.70
37	BD	18	G	C3'-C2'-C1'	-5.89	96.79	101.50
37	BD	113	G	C5-C6-O6	-5.89	125.07	128.60
38	BE	79	G	P-O3'-C3'	-5.89	112.63	119.70
38	BE	143	A	N1-C6-N6	-5.89	115.07	118.60
41	BH	54	U	P-O3'-C3'	-5.89	112.63	119.70
42	BI	20	TYR	CB-CG-CD1	-5.89	117.47	121.00
85	AA	65	A	C5'-C4'-O4'	-5.89	102.03	109.10
85	AA	283	A	C5-N7-C8	-5.89	100.95	103.90
85	AA	921	C	N1-C2-O2	5.89	122.43	118.90
85	AA	1447	U	N3-C2-O2	-5.89	118.08	122.20
85	AA	1533	C	O4'-C1'-N1	5.89	112.91	108.20
85	AA	1606	G	O4'-C1'-N9	5.89	112.91	108.20
86	AB	69	G	C4-N9-C1'	-5.89	118.84	126.50
2	A1	64	GLN	C-N-CA	5.89	134.66	122.30
35	BB	854	G	O4'-C1'-N9	5.89	112.91	108.20
35	BB	1433	U	O5'-C5'-C4'	-5.89	100.51	111.70
35	BB	1456	G	O4'-C1'-N9	5.89	112.91	108.20
36	BC	7	U	C5'-C4'-C3'	-5.89	106.58	116.00
36	BC	56	G	N3-C2-N2	5.89	124.02	119.90
36	BC	155	C	C5'-C4'-O4'	5.89	116.17	109.10
41	BH	6	U	C6-N1-C1'	5.89	129.44	121.20
48	BO	36	ARG	CG-CD-NE	-5.89	99.43	111.80
85	AA	389	A	C5-C6-N1	5.89	120.64	117.70
85	AA	913	U	C5-C4-O4	-5.89	122.37	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	959	C	C6-N1-C2	-5.89	117.94	120.30
32	AY	40	TYR	CB-CG-CD2	-5.89	117.47	121.00
34	BA	405	C	N3-C2-O2	-5.89	117.78	121.90
34	BA	583	G	N1-C6-O6	-5.89	116.37	119.90
34	BA	586	G	O4'-C1'-N9	5.89	112.91	108.20
34	BA	1812	C	P-O3'-C3'	-5.89	112.64	119.70
35	BB	89	C	P-O5'-C5'	-5.89	111.48	120.90
35	BB	419	G	O4'-C1'-C2'	5.89	112.90	107.60
35	BB	573	C	C6-N1-C1'	-5.89	113.74	120.80
35	BB	1492	C	C2'-C3'-O3'	5.89	123.12	113.70
40	BG	136	G	N9-C1'-C2'	-5.89	105.53	112.00
52	BS	120	TYR	CB-CG-CD1	5.89	124.53	121.00
82	Bw	245	ARG	NE-CZ-NH1	5.89	123.24	120.30
85	AA	201	U	O4'-C1'-N1	5.89	112.91	108.20
85	AA	250	C	C5-C6-N1	5.89	123.94	121.00
85	AA	464	A	C8-N9-C4	-5.89	103.44	105.80
85	AA	743	C	C3'-C2'-C1'	-5.89	96.79	101.50
85	AA	884	A	C4-C5-C6	-5.89	114.06	117.00
85	AA	1241	A	N1-C6-N6	5.89	122.13	118.60
34	BA	298	G	OP1-P-O3'	5.88	118.14	105.20
34	BA	1088	G	C4-N9-C1'	-5.88	118.85	126.50
35	BB	498	G	C3'-C2'-C1'	-5.88	96.79	101.50
36	BC	4	G	C5'-C4'-C3'	5.88	125.41	116.00
38	BE	31	A	O3'-P-O5'	-5.88	92.82	104.00
38	BE	196	C	C6-N1-C2	-5.88	117.95	120.30
85	AA	92	G	O3'-P-O5'	-5.88	92.82	104.00
85	AA	440	U	O4'-C1'-N1	5.88	112.91	108.20
85	AA	1114	A	P-O5'-C5'	-5.88	111.48	120.90
85	AA	1483	A	C8-N9-C4	5.88	108.15	105.80
20	AL	5	ARG	CD-NE-CZ	5.88	131.84	123.60
34	BA	1314	A	N9-C1'-C2'	-5.88	105.53	112.00
34	BA	1538	G	N3-C4-C5	-5.88	125.66	128.60
35	BB	996	G	N1-C2-N3	-5.88	120.37	123.90
35	BB	999	G	C3'-C2'-C1'	-5.88	96.79	101.50
36	BC	29	C	N1-C2-O2	-5.88	115.37	118.90
36	BC	152	C	C2'-C3'-O3'	5.88	123.11	113.70
38	BE	15	A	C4'-C3'-C2'	-5.88	96.72	102.60
48	BO	219	PHE	CB-CG-CD2	-5.88	116.68	120.80
55	BV	50	TYR	CB-CG-CD2	-5.88	117.47	121.00
77	Br	217	ARG	NE-CZ-NH2	-5.88	117.36	120.30
85	AA	83	U	C2-N1-C1'	-5.88	110.64	117.70
85	AA	1931	C	C5'-C4'-O4'	-5.88	102.04	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2034	G	C5-C6-O6	-5.88	125.07	128.60
34	BA	947	A	C5'-C4'-C3'	-5.88	106.59	116.00
35	BB	487	A	C5-C6-N6	-5.88	119.00	123.70
35	BB	1495	U	C5'-C4'-C3'	-5.88	106.59	116.00
36	BC	21	U	P-O3'-C3'	-5.88	112.64	119.70
36	BC	95	A	O4'-C4'-C3'	-5.88	98.12	104.00
36	BC	118	U	O4'-C1'-N1	5.88	112.91	108.20
37	BD	27	A	C3'-C2'-C1'	-5.88	96.80	101.50
40	BG	49	A	C3'-C2'-C1'	-5.88	96.80	101.50
85	AA	854	A	OP1-P-OP2	-5.88	110.78	119.60
85	AA	881	C	C3'-C2'-C1'	-5.88	96.80	101.50
85	AA	1092	G	N7-C8-N9	5.88	116.04	113.10
85	AA	1174	G	C8-N9-C1'	5.88	134.65	127.00
85	AA	1368	G	C5'-C4'-O4'	5.88	116.16	109.10
85	AA	1591	U	O4'-C1'-N1	5.88	112.91	108.20
85	AA	2077	G	O3'-P-O5'	5.88	115.18	104.00
85	AA	2107	C	P-O3'-C3'	5.88	126.76	119.70
85	AA	2151	U	C2-N1-C1'	-5.88	110.64	117.70
34	BA	12	G	O5'-P-OP2	-5.88	100.41	105.70
34	BA	1104	C	C2-N3-C4	-5.88	116.96	119.90
34	BA	1450	G	OP1-P-OP2	-5.88	110.78	119.60
35	BB	74	U	C2-N3-C4	-5.88	123.47	127.00
37	BD	105	G	N9-C1'-C2'	-5.88	105.53	112.00
85	AA	1242	A	C1'-O4'-C4'	-5.88	105.20	109.90
15	AG	26	LEU	CB-CA-C	-5.88	99.03	110.20
34	BA	184	C	N3-C2-O2	-5.88	117.78	121.90
34	BA	250	G	C8-N9-C4	-5.88	104.05	106.40
34	BA	300	C	N1-C1'-C2'	-5.88	105.53	112.00
34	BA	311	C	O4'-C1'-N1	5.88	112.90	108.20
34	BA	404	C	C4'-C3'-C2'	5.88	108.48	102.60
34	BA	466	G	O5'-C5'-C4'	5.88	122.87	111.70
34	BA	752	A	P-O3'-C3'	5.88	126.75	119.70
34	BA	807	U	O4'-C1'-N1	5.88	112.90	108.20
34	BA	903	C	O4'-C1'-C2'	5.88	112.89	107.60
34	BA	970	U	O4'-C1'-N1	5.88	112.90	108.20
34	BA	1644	A	C5'-C4'-O4'	5.88	116.15	109.10
35	BB	661	G	N9-C4-C5	5.88	107.75	105.40
35	BB	768	A	C3'-C2'-C1'	-5.88	96.80	101.50
35	BB	792	G	C5-N7-C8	-5.88	101.36	104.30
37	BD	28	C	O4'-C1'-N1	5.88	112.90	108.20
40	BG	144	G	P-O3'-C3'	-5.88	112.65	119.70
85	AA	1129	A	P-O5'-C5'	-5.88	111.50	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1538	C	C5'-C4'-C3'	-5.88	106.59	116.00
85	AA	1733	G	C5'-C4'-C3'	5.88	125.41	116.00
34	BA	19	G	C5-C6-O6	5.88	132.13	128.60
34	BA	731	A	C3'-C2'-C1'	-5.88	96.80	101.50
34	BA	748	C	N3-C2-O2	-5.88	117.79	121.90
34	BA	843	G	N1-C2-N2	-5.88	110.91	116.20
34	BA	1108	U	C2-N1-C1'	-5.88	110.65	117.70
35	BB	156	G	N1-C6-O6	5.88	123.43	119.90
35	BB	1534	U	C2'-C3'-O3'	5.88	123.10	113.70
41	BH	32	U	C4'-C3'-C2'	-5.88	96.72	102.60
85	AA	1000	U	O4'-C1'-N1	-5.88	103.50	108.20
85	AA	1366	A	C5'-C4'-C3'	-5.88	106.60	116.00
85	AA	1905	A	C5-C6-N1	5.88	120.64	117.70
34	BA	170	U	P-O3'-C3'	-5.88	112.65	119.70
34	BA	713	C	O4'-C1'-N1	5.88	112.90	108.20
34	BA	1470	G	O4'-C1'-N9	5.88	112.90	108.20
34	BA	1503	U	C2-N3-C4	-5.88	123.47	127.00
35	BB	1044	U	O4'-C1'-N1	5.88	112.90	108.20
38	BE	51	C	C5-C4-N4	-5.88	116.09	120.20
38	BE	141	A	O5'-C5'-C4'	-5.88	100.54	111.70
41	BH	63	G	N3-C4-C5	-5.88	125.66	128.60
65	Bf	146	THR	CA-CB-CG2	-5.88	104.17	112.40
69	Bj	94	ARG	NE-CZ-NH1	5.88	123.24	120.30
77	Br	105	LYS	N-CA-C	5.88	126.86	111.00
77	Br	231	ASN	CA-CB-CG	-5.88	100.48	113.40
85	AA	669	G	O4'-C1'-N9	5.88	112.90	108.20
85	AA	997	U	C5'-C4'-C3'	-5.88	106.60	116.00
85	AA	2056	C	P-O3'-C3'	-5.88	112.65	119.70
85	AA	2112	G	P-O5'-C5'	5.88	130.30	120.90
34	BA	15	G	C4-C5-N7	5.87	113.15	110.80
34	BA	204	U	C1'-O4'-C4'	-5.87	105.20	109.90
34	BA	255	G	P-O5'-C5'	5.87	130.30	120.90
34	BA	921	G	C1'-O4'-C4'	-5.87	105.20	109.90
34	BA	1087	A	P-O3'-C3'	-5.87	112.65	119.70
34	BA	1289	C	P-O5'-C5'	-5.87	111.50	120.90
35	BB	421	U	C5'-C4'-C3'	5.87	125.40	116.00
35	BB	429	C	C4'-C3'-C2'	-5.87	96.73	102.60
35	BB	796	C	C1'-O4'-C4'	-5.87	105.20	109.90
35	BB	1196	A	O4'-C1'-C2'	5.87	112.89	107.60
35	BB	1283	C	C2-N3-C4	-5.87	116.96	119.90
36	BC	92	C	O4'-C1'-N1	5.87	112.90	108.20
36	BC	141	C	O3'-P-O5'	-5.87	92.84	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	BD	10	C	N3-C4-N4	5.87	122.11	118.00
39	BF	64	U	O4'-C1'-N1	5.87	112.90	108.20
40	BG	25	G	C1'-O4'-C4'	-5.87	105.20	109.90
41	BH	29	G	C5-C6-N1	5.87	114.44	111.50
41	BH	39	G	O4'-C1'-N9	5.87	112.90	108.20
50	BQ	202	ARG	NE-CZ-NH1	5.87	123.24	120.30
63	Bd	21	ILE	N-CA-CB	5.87	124.31	110.80
85	AA	319	U	C5'-C4'-C3'	5.87	125.40	116.00
85	AA	1016	G	C4-N9-C1'	-5.87	118.86	126.50
85	AA	1247	A	C4-C5-C6	-5.87	114.06	117.00
85	AA	2172	A	N7-C8-N9	5.87	116.74	113.80
34	BA	1119	A	C1'-O4'-C4'	-5.87	105.20	109.90
34	BA	1503	U	C3'-C2'-C1'	-5.87	96.80	101.50
34	BA	1620	U	C2-N3-C4	-5.87	123.48	127.00
35	BB	380	G	N3-C2-N2	-5.87	115.79	119.90
35	BB	572	G	O4'-C1'-N9	5.87	112.90	108.20
35	BB	1066	G	C1'-O4'-C4'	-5.87	105.20	109.90
36	BC	136	G	C4-N9-C1'	-5.87	118.87	126.50
37	BD	2	G	C5'-C4'-C3'	5.87	125.39	116.00
37	BD	116	C	C3'-C2'-C1'	-5.87	96.80	101.50
44	BK	83	GLU	CB-CA-C	-5.87	98.66	110.40
50	BQ	174	LYS	C-N-CA	5.87	136.38	121.70
85	AA	138	C	C5'-C4'-C3'	5.87	125.39	116.00
85	AA	156	G	C8-N9-C1'	5.87	134.63	127.00
85	AA	168	A	N7-C8-N9	5.87	116.74	113.80
85	AA	185	A	C5-C6-N6	-5.87	119.00	123.70
85	AA	317	A	C5'-C4'-C3'	5.87	125.39	116.00
85	AA	400	G	P-O5'-C5'	-5.87	111.50	120.90
85	AA	530	A	C8-N9-C4	-5.87	103.45	105.80
85	AA	584	G	C6-C5-N7	-5.87	126.88	130.40
85	AA	758	C	C2-N1-C1'	-5.87	112.34	118.80
85	AA	1357	U	O4'-C1'-N1	5.87	112.90	108.20
85	AA	1672	G	C8-N9-C4	5.87	108.75	106.40
34	BA	1066	A	O4'-C1'-N9	5.87	112.90	108.20
35	BB	634	A	C3'-C2'-C1'	-5.87	96.80	101.50
36	BC	84	U	C6-N1-C1'	-5.87	112.98	121.20
48	BO	144	ARG	NE-CZ-NH1	5.87	123.23	120.30
85	AA	496	C	C3'-C2'-C1'	-5.87	96.80	101.50
85	AA	631	G	P-O3'-C3'	5.87	126.74	119.70
85	AA	1122	U	C1'-O4'-C4'	-5.87	105.20	109.90
34	BA	704	G	C8-N9-C1'	5.87	134.63	127.00
34	BA	915	A	C5'-C4'-C3'	5.87	125.39	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	953	G	C5-C6-N1	5.87	114.44	111.50
34	BA	1043	C	O5'-C5'-C4'	5.87	122.85	111.70
34	BA	1097	G	C1'-O4'-C4'	5.87	114.59	109.90
34	BA	1448	G	C1'-O4'-C4'	-5.87	105.21	109.90
35	BB	597	C	N3-C2-O2	-5.87	117.79	121.90
35	BB	1251	G	O4'-C1'-N9	5.87	112.89	108.20
37	BD	76	U	C5-C4-O4	-5.87	122.38	125.90
39	BF	14	C	N3-C4-N4	5.87	122.11	118.00
40	BG	123	C	C1'-O4'-C4'	-5.87	105.21	109.90
85	AA	386	G	N3-C4-N9	5.87	129.52	126.00
85	AA	454	G	C2'-C3'-O3'	5.87	123.09	113.70
85	AA	493	A	P-O3'-C3'	-5.87	112.66	119.70
85	AA	902	A	P-O3'-C3'	-5.87	112.66	119.70
85	AA	938	A	C5-C6-N6	-5.87	119.01	123.70
85	AA	1145	U	O4'-C4'-C3'	-5.87	98.13	104.00
85	AA	1256	C	C5-C4-N4	-5.87	116.09	120.20
85	AA	1678	U	C1'-O4'-C4'	-5.87	105.20	109.90
85	AA	1951	U	O4'-C1'-N1	5.87	112.89	108.20
85	AA	2213	A	C5'-C4'-O4'	-5.87	102.06	109.10
34	BA	519	G	C5-C6-O6	-5.87	125.08	128.60
34	BA	687	G	C2'-C3'-O3'	5.87	123.09	113.70
34	BA	1006	G	N1-C6-O6	-5.87	116.38	119.90
35	BB	501	G	N1-C2-N2	-5.87	110.92	116.20
35	BB	584	A	C6-N1-C2	-5.87	115.08	118.60
35	BB	1351	G	O4'-C1'-C2'	5.87	112.88	107.60
35	BB	1356	G	N3-C2-N2	5.87	124.01	119.90
38	BE	16	C	OP1-P-OP2	-5.87	110.80	119.60
39	BF	34	C	C2-N3-C4	-5.87	116.97	119.90
39	BF	56	C	C2'-C3'-O3'	5.87	123.09	113.70
62	Bc	40	LYS	C-N-CA	5.87	136.37	121.70
85	AA	501	A	C4-N9-C1'	-5.87	115.74	126.30
85	AA	1001	G	P-O3'-C3'	-5.87	112.66	119.70
85	AA	1113	G	C4-N9-C1'	5.87	134.13	126.50
34	BA	278	U	P-O3'-C3'	-5.87	112.66	119.70
34	BA	705	C	C6-N1-C2	-5.87	117.95	120.30
34	BA	827	A	C4'-C3'-C2'	-5.87	96.73	102.60
34	BA	1841	A	O3'-P-O5'	5.87	115.14	104.00
35	BB	162	U	O4'-C1'-N1	5.87	112.89	108.20
35	BB	837	A	N1-C6-N6	-5.87	115.08	118.60
41	BH	82	U	O4'-C1'-N1	5.87	112.89	108.20
85	AA	21	U	P-O3'-C3'	-5.87	112.66	119.70
85	AA	991	G	P-O3'-C3'	-5.87	112.66	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1001	G	O4'-C1'-N9	5.87	112.89	108.20
85	AA	1267	A	N9-C1'-C2'	-5.87	105.55	112.00
85	AA	1357	U	C5'-C4'-C3'	-5.87	106.61	116.00
85	AA	1555	G	C5'-C4'-C3'	-5.87	106.61	116.00
85	AA	1851	A	O4'-C1'-C2'	5.87	112.88	107.60
21	AM	33	MET	C-N-CA	5.86	136.36	121.70
21	AM	40	ARG	NE-CZ-NH2	-5.86	117.37	120.30
34	BA	146	G	N1-C6-O6	-5.86	116.38	119.90
34	BA	179	U	C6-N1-C2	-5.86	117.48	121.00
34	BA	749	G	C4-N9-C1'	-5.86	118.88	126.50
34	BA	1220	C	C2-N3-C4	-5.86	116.97	119.90
34	BA	1294	C	C6-N1-C2	-5.86	117.95	120.30
35	BB	582	G	O4'-C1'-N9	5.86	112.89	108.20
35	BB	796	C	C3'-C2'-C1'	-5.86	96.81	101.50
36	BC	9	G	P-O3'-C3'	5.86	126.74	119.70
38	BE	21	C	C2-N3-C4	-5.86	116.97	119.90
41	BH	33	G	N9-C4-C5	-5.86	103.06	105.40
41	BH	84	A	O4'-C1'-N9	5.86	112.89	108.20
80	Bu	278	TYR	CB-CG-CD2	-5.86	117.48	121.00
85	AA	95	U	C6-N1-C2	-5.86	117.48	121.00
85	AA	794	A	C3'-C2'-C1'	-5.86	96.81	101.50
85	AA	812	C	P-O3'-C3'	-5.86	112.67	119.70
34	BA	405	C	C2-N3-C4	-5.86	116.97	119.90
34	BA	856	G	N3-C2-N2	5.86	124.00	119.90
34	BA	1301	G	C5-C6-N1	5.86	114.43	111.50
34	BA	1397	C	C4'-C3'-C2'	-5.86	96.74	102.60
35	BB	84	G	P-O5'-C5'	-5.86	111.52	120.90
36	BC	98	C	O3'-P-O5'	-5.86	92.86	104.00
38	BE	77	C	C5'-C4'-C3'	5.86	125.38	116.00
65	Bf	463	ARG	NE-CZ-NH1	5.86	123.23	120.30
85	AA	1961	U	C1'-O4'-C4'	-5.86	105.21	109.90
15	AG	45	LEU	CB-CA-C	-5.86	99.06	110.20
34	BA	12	G	N9-C4-C5	-5.86	103.06	105.40
34	BA	187	G	P-O5'-C5'	-5.86	111.52	120.90
34	BA	321	G	C4-N9-C1'	-5.86	118.88	126.50
34	BA	756	A	O5'-P-OP2	-5.86	100.43	105.70
34	BA	1427	U	C5'-C4'-O4'	5.86	116.13	109.10
35	BB	258	C	O4'-C1'-N1	5.86	112.89	108.20
35	BB	1118	G	C5-C6-N1	5.86	114.43	111.50
35	BB	1498	G	O4'-C1'-N9	5.86	112.89	108.20
36	BC	54	G	C5'-C4'-C3'	5.86	125.38	116.00
38	BE	107	U	P-O3'-C3'	-5.86	112.67	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	44	G	O3'-P-O5'	5.86	115.13	104.00
85	AA	957	A	C1'-O4'-C4'	-5.86	105.21	109.90
85	AA	991	G	C4-C5-C6	-5.86	115.28	118.80
85	AA	2125	A	C8-N9-C4	5.86	108.14	105.80
34	BA	629	G	C4-C5-N7	-5.86	108.46	110.80
34	BA	1532	G	C5-C6-N1	5.86	114.43	111.50
34	BA	1680	G	C4-N9-C1'	5.86	134.12	126.50
35	BB	316	U	O4'-C1'-N1	5.86	112.89	108.20
35	BB	795	A	C5'-C4'-C3'	5.86	125.37	116.00
40	BG	115	C	P-O3'-C3'	-5.86	112.67	119.70
85	AA	806	G	C8-N9-C4	-5.86	104.06	106.40
85	AA	2157	G	N9-C4-C5	5.86	107.74	105.40
1	A0	213	ARG	NE-CZ-NH1	5.86	123.23	120.30
34	BA	383	G	C3'-C2'-C1'	-5.86	96.81	101.50
34	BA	613	A	C5'-C4'-C3'	-5.86	106.63	116.00
34	BA	937	G	C3'-C2'-C1'	-5.86	96.81	101.50
34	BA	1177	C	N3-C4-C5	-5.86	119.56	121.90
34	BA	1569	C	N3-C2-O2	-5.86	117.80	121.90
34	BA	1590	G	N9-C1'-C2'	-5.86	105.56	112.00
41	BH	77	G	C2-N3-C4	5.86	114.83	111.90
67	Bh	75	LYS	N-CA-C	5.86	126.81	111.00
85	AA	313	A	C5-C6-N1	5.86	120.63	117.70
85	AA	1222	A	C5'-C4'-C3'	-5.86	106.63	116.00
85	AA	1424	G	P-O3'-C3'	-5.86	112.67	119.70
85	AA	1611	A	P-O3'-C3'	5.86	126.73	119.70
34	BA	195	G	C5-C6-N1	5.86	114.43	111.50
34	BA	297	A	O4'-C4'-C3'	5.86	110.78	106.10
34	BA	780	U	C5'-C4'-O4'	5.86	116.13	109.10
34	BA	1156	U	N1-C1'-C2'	-5.86	105.56	112.00
34	BA	1798	G	N1-C6-O6	5.86	123.41	119.90
35	BB	391	G	C3'-C2'-C1'	-5.86	96.82	101.50
35	BB	558	U	C1'-O4'-C4'	-5.86	105.22	109.90
35	BB	657	A	C2-N3-C4	5.86	113.53	110.60
35	BB	852	G	C8-N9-C4	-5.86	104.06	106.40
35	BB	906	G	C5-C6-O6	-5.86	125.09	128.60
35	BB	1012	G	P-O3'-C3'	-5.86	112.67	119.70
41	BH	59	G	N7-C8-N9	5.86	116.03	113.10
47	BN	46	LEU	N-CA-CB	-5.86	98.69	110.40
57	BX	125	TYR	CB-CG-CD1	-5.86	117.49	121.00
85	AA	117	C	C2-N3-C4	-5.86	116.97	119.90
85	AA	277	G	N3-C2-N2	5.86	124.00	119.90
85	AA	492	C	O4'-C1'-N1	5.86	112.89	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	533	C	C1'-O4'-C4'	-5.86	105.22	109.90
85	AA	901	C	C6-N1-C1'	-5.86	113.77	120.80
85	AA	1094	G	C5'-C4'-C3'	-5.86	106.63	116.00
34	BA	931	G	OP1-P-O3'	5.85	118.08	105.20
34	BA	932	G	C5-C6-O6	-5.85	125.09	128.60
34	BA	1372	C	C6-N1-C2	-5.85	117.96	120.30
35	BB	374	A	N7-C8-N9	-5.85	110.87	113.80
35	BB	946	G	O4'-C1'-N9	5.85	112.88	108.20
35	BB	1313	C	P-O3'-C3'	-5.85	112.67	119.70
85	AA	740	A	P-O3'-C3'	-5.85	112.67	119.70
85	AA	867	G	N7-C8-N9	5.85	116.03	113.10
85	AA	1043	U	O4'-C1'-N1	5.85	112.88	108.20
34	BA	247	U	N1-C1'-C2'	5.85	121.61	114.00
34	BA	355	U	C2-N1-C1'	-5.85	110.68	117.70
34	BA	761	U	N1-C2-O2	5.85	126.90	122.80
34	BA	888	G	O5'-C5'-C4'	-5.85	100.58	111.70
34	BA	895	U	P-O5'-C5'	-5.85	111.54	120.90
34	BA	964	U	C5'-C4'-O4'	5.85	116.12	109.10
34	BA	1220	C	C5-C6-N1	-5.85	118.07	121.00
34	BA	1559	C	C5'-C4'-O4'	5.85	116.12	109.10
35	BB	89	C	O5'-P-OP2	-5.85	100.43	105.70
35	BB	102	G	C5-C6-O6	-5.85	125.09	128.60
35	BB	505	G	N3-C4-N9	5.85	129.51	126.00
35	BB	887	G	C5-C6-O6	-5.85	125.09	128.60
35	BB	1262	A	C5-C6-N1	5.85	120.63	117.70
36	BC	41	A	P-O3'-C3'	5.85	126.72	119.70
38	BE	136	G	O5'-P-OP2	-5.85	100.43	105.70
38	BE	139	U	C5'-C4'-C3'	-5.85	106.64	116.00
38	BE	143	A	P-O3'-C3'	5.85	126.72	119.70
65	Bf	342	GLY	N-CA-C	5.85	127.73	113.10
85	AA	6	G	P-O5'-C5'	5.85	130.26	120.90
85	AA	187	C	P-O5'-C5'	5.85	130.26	120.90
85	AA	575	G	C1'-O4'-C4'	-5.85	105.22	109.90
85	AA	736	U	C4'-C3'-O3'	5.85	124.71	113.00
85	AA	786	G	N1-C6-O6	5.85	123.41	119.90
85	AA	894	A	O5'-C5'-C4'	5.85	122.82	111.70
85	AA	998	U	O4'-C1'-N1	5.85	112.88	108.20
85	AA	1723	U	C5'-C4'-C3'	-5.85	106.64	116.00
85	AA	1736	U	C5'-C4'-C3'	-5.85	106.64	116.00
34	BA	688	G	P-O3'-C3'	-5.85	112.68	119.70
34	BA	783	U	N1-C2-O2	5.85	126.90	122.80
34	BA	1375	C	O4'-C1'-N1	5.85	112.88	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1830	A	C2'-C3'-O3'	5.85	123.06	113.70
35	BB	480	C	P-O5'-C5'	-5.85	111.54	120.90
35	BB	995	C	O3'-P-O5'	5.85	115.12	104.00
69	Bj	91	ARG	NE-CZ-NH1	5.85	123.23	120.30
77	Br	92	GLY	CA-C-N	-5.85	104.33	117.20
85	AA	1451	U	C5-C4-O4	-5.85	122.39	125.90
85	AA	1752	C	O5'-C5'-C4'	-5.85	100.58	111.70
34	BA	6	C	O5'-C5'-C4'	-5.85	100.58	111.70
34	BA	810	A	P-O5'-C5'	5.85	130.26	120.90
34	BA	1041	U	C4-C5-C6	-5.85	116.19	119.70
35	BB	70	A	O4'-C1'-N9	5.85	112.88	108.20
35	BB	770	G	O4'-C1'-N9	5.85	112.88	108.20
35	BB	790	A	N9-C4-C5	-5.85	103.46	105.80
35	BB	1203	C	P-O3'-C3'	-5.85	112.68	119.70
35	BB	1548	C	C2-N1-C1'	-5.85	112.37	118.80
37	BD	44	U	C6-N1-C2	-5.85	117.49	121.00
40	BG	60	A	O4'-C4'-C3'	-5.85	98.15	104.00
40	BG	95	U	N3-C4-O4	-5.85	115.31	119.40
85	AA	564	A	C5'-C4'-C3'	5.85	125.36	116.00
85	AA	681	G	C5-N7-C8	-5.85	101.38	104.30
85	AA	730	G	N9-C4-C5	5.85	107.74	105.40
85	AA	846	U	C5'-C4'-C3'	5.85	125.36	116.00
85	AA	1273	C	C4'-C3'-C2'	-5.85	96.75	102.60
34	BA	887	U	C2-N1-C1'	-5.85	110.68	117.70
34	BA	990	G	C5'-C4'-O4'	5.85	116.12	109.10
34	BA	1456	C	O3'-P-O5'	5.85	115.11	104.00
34	BA	1525	G	C5-C6-N1	5.85	114.42	111.50
34	BA	1642	A	C8-N9-C1'	5.85	138.23	127.70
35	BB	111	C	C4'-C3'-C2'	-5.85	96.75	102.60
35	BB	1429	A	N1-C6-N6	-5.85	115.09	118.60
41	BH	67	G	C8-N9-C4	-5.85	104.06	106.40
56	BW	58	ASP	CB-CA-C	5.85	122.09	110.40
74	Bo	13	ARG	NE-CZ-NH2	5.85	123.22	120.30
85	AA	241	U	N1-C2-N3	5.85	118.41	114.90
85	AA	411	U	C5-C6-N1	-5.85	119.78	122.70
85	AA	941	C	C6-N1-C2	-5.85	117.96	120.30
85	AA	2199	G	N9-C1'-C2'	-5.85	105.57	112.00
34	BA	1416	C	O4'-C1'-C2'	5.85	112.86	107.60
35	BB	77	A	C8-N9-C1'	5.85	138.22	127.70
35	BB	1053	G	N3-C2-N2	5.85	123.99	119.90
35	BB	1415	G	C5-C6-O6	5.85	132.11	128.60
38	BE	70	C	N3-C4-N4	-5.85	113.91	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	59	C	P-O3'-C3'	5.85	126.72	119.70
85	AA	519	A	O4'-C4'-C3'	-5.85	98.15	104.00
34	BA	18	G	P-O5'-C5'	-5.84	111.55	120.90
34	BA	418	G	OP2-P-O3'	5.84	118.06	105.20
34	BA	568	G	C3'-C2'-C1'	-5.84	96.83	101.50
34	BA	852	C	C2'-C3'-O3'	5.84	123.05	113.70
34	BA	914	G	C5-C6-O6	-5.84	125.09	128.60
35	BB	815	G	C8-N9-C4	5.84	108.74	106.40
35	BB	1146	C	C2-N1-C1'	5.84	125.23	118.80
35	BB	1312	U	C5-C6-N1	-5.84	119.78	122.70
35	BB	1496	C	C3'-C2'-C1'	-5.84	96.82	101.50
36	BC	159	U	O4'-C1'-N1	5.84	112.88	108.20
38	BE	117	A	N3-C4-C5	-5.84	122.71	126.80
38	BE	132	U	P-O5'-C5'	5.84	130.25	120.90
41	BH	92	A	C5-C6-N6	-5.84	119.02	123.70
62	Bc	127	ARG	CB-CA-C	-5.84	98.71	110.40
85	AA	582	A	C3'-C2'-C1'	-5.84	96.82	101.50
85	AA	688	C	C6-N1-C2	-5.84	117.96	120.30
85	AA	934	A	O4'-C1'-N9	5.84	112.88	108.20
85	AA	976	G	C3'-C2'-C1'	-5.84	96.83	101.50
85	AA	1280	U	N3-C4-C5	-5.84	111.09	114.60
85	AA	1503	G	C8-N9-C1'	5.84	134.60	127.00
2	A1	10	TYR	N-CA-CB	5.84	121.12	110.60
34	BA	412	G	C5-C6-O6	-5.84	125.09	128.60
34	BA	530	A	N1-C2-N3	-5.84	126.38	129.30
34	BA	1454	G	C4-C5-N7	5.84	113.14	110.80
35	BB	1544	A	N1-C6-N6	-5.84	115.09	118.60
85	AA	108	C	O4'-C4'-C3'	-5.84	98.16	104.00
85	AA	122	A	O4'-C1'-N9	5.84	112.87	108.20
24	AQ	101	PHE	CB-CG-CD2	-5.84	116.71	120.80
34	BA	500	C	C1'-O4'-C4'	-5.84	105.23	109.90
34	BA	908	G	C5-C6-O6	-5.84	125.09	128.60
34	BA	1097	G	C8-N9-C4	-5.84	104.06	106.40
34	BA	1703	A	C5-C6-N6	-5.84	119.03	123.70
35	BB	478	G	C1'-O4'-C4'	-5.84	105.23	109.90
35	BB	1129	C	C4'-C3'-C2'	-5.84	96.76	102.60
35	BB	1135	U	P-O3'-C3'	-5.84	112.69	119.70
38	BE	143	A	P-O5'-C5'	-5.84	111.55	120.90
40	BG	37	G	C5'-C4'-C3'	-5.84	106.65	116.00
40	BG	73	U	C2'-C3'-O3'	5.84	123.05	113.70
40	BG	169	A	C8-N9-C1'	-5.84	117.19	127.70
65	Bf	354	THR	CA-CB-CG2	-5.84	104.22	112.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
71	Bl	111	ARG	N-CA-CB	-5.84	100.09	110.60
85	AA	443	A	O4'-C1'-N9	-5.84	103.53	108.20
85	AA	885	A	O4'-C1'-N9	5.84	112.87	108.20
85	AA	1226	A	P-O3'-C3'	-5.84	112.69	119.70
85	AA	1483	A	C2-N3-C4	5.84	113.52	110.60
34	BA	217	C	N3-C4-N4	5.84	122.09	118.00
35	BB	17	U	C4'-C3'-C2'	-5.84	96.76	102.60
35	BB	499	A	O4'-C1'-N9	5.84	112.87	108.20
35	BB	970	C	C5'-C4'-C3'	-5.84	106.66	116.00
35	BB	1063	C	N1-C2-O2	5.84	122.40	118.90
35	BB	1139	A	C6-N1-C2	-5.84	115.10	118.60
40	BG	22	G	N1-C6-O6	5.84	123.40	119.90
41	BH	76	G	OP2-P-O3'	5.84	118.05	105.20
85	AA	178	U	P-O5'-C5'	-5.84	111.56	120.90
85	AA	190	A	C4-N9-C1'	-5.84	115.79	126.30
85	AA	534	A	C5-C6-N6	5.84	128.37	123.70
85	AA	629	A	N7-C8-N9	-5.84	110.88	113.80
85	AA	833	U	C2-N1-C1'	5.84	124.71	117.70
85	AA	1368	G	C6-C5-N7	-5.84	126.90	130.40
85	AA	1412	G	O4'-C1'-N9	5.84	112.87	108.20
85	AA	2215	C	C6-N1-C2	-5.84	117.96	120.30
34	BA	382	G	C8-N9-C1'	5.84	134.59	127.00
34	BA	570	G	OP2-P-O3'	5.84	118.04	105.20
35	BB	11	A	C5'-C4'-C3'	5.84	125.34	116.00
35	BB	1020	U	O4'-C1'-N1	5.84	112.87	108.20
35	BB	1230	A	C6-N1-C2	-5.84	115.10	118.60
85	AA	169	G	O4'-C1'-N9	5.84	112.87	108.20
85	AA	545	A	N9-C1'-C2'	-5.84	105.58	112.00
85	AA	778	C	N3-C2-O2	-5.84	117.81	121.90
85	AA	1008	C	C5'-C4'-C3'	5.84	125.34	116.00
34	BA	49	A	C8-N9-C1'	5.84	138.21	127.70
34	BA	88	C	C5-C4-N4	5.84	124.28	120.20
34	BA	425	G	O4'-C1'-N9	5.84	112.87	108.20
34	BA	430	A	O3'-P-O5'	-5.84	92.91	104.00
34	BA	453	A	P-O5'-C5'	-5.84	111.56	120.90
34	BA	675	C	OP2-P-O3'	5.84	118.04	105.20
34	BA	807	U	N3-C2-O2	5.84	126.29	122.20
34	BA	1311	G	N1-C6-O6	-5.84	116.40	119.90
34	BA	1513	G	C8-N9-C4	-5.84	104.06	106.40
35	BB	533	U	C5'-C4'-O4'	5.84	116.11	109.10
35	BB	898	U	O4'-C1'-N1	5.84	112.87	108.20
35	BB	1538	G	C5-C6-O6	-5.84	125.10	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	152	U	P-O3'-C3'	5.84	126.70	119.70
81	Bv	75	ARG	CD-NE-CZ	5.84	131.77	123.60
85	AA	210	G	O5'-P-OP1	5.84	117.70	110.70
85	AA	362	G	N1-C6-O6	5.84	123.40	119.90
85	AA	519	A	O4'-C1'-N9	5.84	112.87	108.20
85	AA	852	C	P-O3'-C3'	5.84	126.70	119.70
85	AA	992	G	N9-C1'-C2'	-5.84	105.58	112.00
85	AA	1203	G	C5-C6-O6	-5.84	125.10	128.60
85	AA	1246	G	N1-C6-O6	5.84	123.40	119.90
85	AA	1266	C	P-O3'-C3'	-5.84	112.70	119.70
85	AA	1732	G	P-O3'-C3'	-5.84	112.70	119.70
85	AA	1856	G	C8-N9-C1'	5.84	134.59	127.00
85	AA	1975	G	C4-N9-C1'	-5.84	118.91	126.50
85	AA	2002	A	C3'-C2'-C1'	-5.84	96.83	101.50
34	BA	476	U	C1'-O4'-C4'	-5.83	105.23	109.90
35	BB	506	G	P-O5'-C5'	-5.83	111.56	120.90
35	BB	1086	G	C5-C6-N1	5.83	114.42	111.50
85	AA	692	U	C3'-C2'-C1'	-5.83	96.83	101.50
30	AW	13	VAL	N-CA-CB	-5.83	98.67	111.50
34	BA	64	A	C2'-C3'-O3'	5.83	123.03	113.70
34	BA	1254	C	C3'-C2'-C1'	-5.83	96.83	101.50
34	BA	1292	A	P-O5'-C5'	-5.83	111.57	120.90
35	BB	254	A	P-O5'-C5'	5.83	130.23	120.90
35	BB	596	C	O4'-C1'-N1	5.83	112.87	108.20
35	BB	1309	A	O4'-C1'-N9	5.83	112.87	108.20
35	BB	1513	U	O4'-C1'-N1	5.83	112.87	108.20
37	BD	68	C	P-O3'-C3'	-5.83	112.70	119.70
65	Bf	343	ARG	NE-CZ-NH2	-5.83	117.38	120.30
85	AA	306	C	O3'-P-O5'	5.83	115.08	104.00
85	AA	381	A	N9-C1'-C2'	-5.83	105.58	112.00
85	AA	1695	G	C5-C6-O6	-5.83	125.10	128.60
5	A4	118	GLN	CB-CA-C	-5.83	98.74	110.40
34	BA	921	G	C4'-C3'-C2'	-5.83	96.77	102.60
34	BA	988	U	P-O3'-C3'	-5.83	112.70	119.70
34	BA	1041	U	O4'-C1'-C2'	5.83	112.85	107.60
34	BA	1559	C	C4'-C3'-C2'	5.83	108.43	102.60
34	BA	1738	G	P-O3'-C3'	-5.83	112.70	119.70
35	BB	498	G	O4'-C1'-N9	5.83	112.87	108.20
35	BB	676	G	O3'-P-O5'	5.83	115.08	104.00
36	BC	11	G	O4'-C1'-N9	5.83	112.86	108.20
40	BG	108	G	N3-C2-N2	5.83	123.98	119.90
84	By	3	ILE	N-CA-C	5.83	126.75	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1008	C	C6-N1-C2	-5.83	117.97	120.30
85	AA	1167	G	C5-C6-O6	5.83	132.10	128.60
34	BA	36	A	C4-N9-C1'	-5.83	115.81	126.30
34	BA	217	C	O5'-C5'-C4'	-5.83	100.62	111.70
34	BA	348	U	C2-N3-C4	-5.83	123.50	127.00
34	BA	685	C	O5'-C5'-C4'	5.83	122.78	111.70
36	BC	97	U	C4'-C3'-C2'	5.83	108.43	102.60
37	BD	36	C	C1'-O4'-C4'	-5.83	105.24	109.90
40	BG	32	U	C5'-C4'-O4'	5.83	116.10	109.10
40	BG	44	G	C5'-C4'-O4'	5.83	116.10	109.10
85	AA	372	U	N3-C2-O2	-5.83	118.12	122.20
85	AA	423	G	C5-C6-N1	5.83	114.42	111.50
34	BA	78	U	C5'-C4'-C3'	-5.83	106.67	116.00
34	BA	399	G	C2-N3-C4	-5.83	108.98	111.90
34	BA	960	C	O5'-P-OP1	5.83	117.69	110.70
34	BA	1200	U	O4'-C1'-C2'	-5.83	99.97	105.80
34	BA	1809	G	C6-C5-N7	-5.83	126.90	130.40
35	BB	128	C	N1-C2-O2	5.83	122.40	118.90
35	BB	577	U	N3-C2-O2	-5.83	118.12	122.20
35	BB	610	U	C2'-C3'-O3'	5.83	123.03	113.70
38	BE	151	C	C5'-C4'-O4'	5.83	116.09	109.10
41	BH	127	A	C8-N9-C4	5.83	108.13	105.80
56	BW	84	ARG	NE-CZ-NH2	-5.83	117.39	120.30
84	By	173	PHE	CB-CA-C	-5.83	98.74	110.40
85	AA	307	G	C4'-C3'-C2'	-5.83	96.77	102.60
85	AA	595	A	C8-N9-C4	5.83	108.13	105.80
85	AA	711	C	O4'-C1'-N1	5.83	112.86	108.20
85	AA	1218	C	C2-N3-C4	-5.83	116.99	119.90
85	AA	1594	G	C5'-C4'-O4'	5.83	116.09	109.10
85	AA	2244	G	C4'-C3'-C2'	5.83	108.43	102.60
86	AB	19	G	P-O3'-C3'	-5.83	112.71	119.70
34	BA	446	U	C3'-C2'-C1'	-5.83	96.84	101.50
34	BA	552	C	N3-C4-C5	5.83	124.23	121.90
35	BB	126	C	C5-C6-N1	5.83	123.91	121.00
35	BB	383	U	C4'-C3'-C2'	-5.83	96.77	102.60
35	BB	1146	C	P-O5'-C5'	-5.83	111.58	120.90
35	BB	1166	A	N9-C4-C5	-5.83	103.47	105.80
40	BG	174	G	C4-N9-C1'	-5.83	118.92	126.50
85	AA	331	G	C4-N9-C1'	5.83	134.07	126.50
85	AA	611	G	N3-C4-C5	-5.83	125.69	128.60
85	AA	661	C	C6-N1-C1'	5.83	127.79	120.80
85	AA	1494	C	P-O5'-C5'	-5.83	111.58	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1829	C	O4'-C1'-N1	5.83	112.86	108.20
26	AS	15	ARG	NE-CZ-NH1	5.83	123.21	120.30
34	BA	172	A	P-O5'-C5'	-5.83	111.58	120.90
34	BA	603	U	N3-C4-O4	5.83	123.48	119.40
34	BA	662	U	N3-C2-O2	-5.83	118.12	122.20
34	BA	800	G	C5'-C4'-C3'	5.83	125.32	116.00
34	BA	884	G	P-O3'-C3'	-5.83	112.71	119.70
35	BB	565	U	C5'-C4'-C3'	-5.83	106.68	116.00
35	BB	664	A	P-O5'-C5'	-5.83	111.58	120.90
35	BB	679	G	C5'-C4'-O4'	5.83	116.09	109.10
35	BB	1234	G	C5'-C4'-C3'	-5.83	106.68	116.00
38	BE	113	C	P-O3'-C3'	-5.83	112.71	119.70
41	BH	78	C	OP1-P-OP2	-5.83	110.86	119.60
64	Be	150	LEU	CB-CG-CD2	5.83	120.90	111.00
85	AA	207	G	C6-C5-N7	-5.83	126.91	130.40
85	AA	286	C	C3'-C2'-C1'	-5.83	96.84	101.50
85	AA	316	C	P-O5'-C5'	5.83	130.22	120.90
85	AA	507	C	C3'-C2'-C1'	-5.83	96.84	101.50
85	AA	507	C	C4'-C3'-C2'	5.83	108.42	102.60
85	AA	652	U	P-O5'-C5'	-5.83	111.58	120.90
85	AA	658	C	C5'-C4'-O4'	5.83	116.09	109.10
85	AA	1923	A	O5'-P-OP2	-5.83	100.46	105.70
86	AB	15	G	N1-C2-N2	-5.83	110.96	116.20
34	BA	489	A	P-O3'-C3'	-5.82	112.71	119.70
35	BB	504	C	C5-C6-N1	5.82	123.91	121.00
35	BB	1031	G	C5'-C4'-C3'	5.82	125.32	116.00
35	BB	1224	C	N1-C1'-C2'	-5.82	105.59	112.00
38	BE	165	U	C6-N1-C2	-5.82	117.51	121.00
40	BG	38	A	C3'-C2'-C1'	-5.82	96.84	101.50
85	AA	108	C	P-O5'-C5'	-5.82	111.58	120.90
85	AA	327	G	P-O3'-C3'	-5.82	112.71	119.70
85	AA	547	A	C3'-C2'-C1'	-5.82	96.84	101.50
85	AA	1467	U	P-O3'-C3'	-5.82	112.71	119.70
85	AA	1662	U	P-O5'-C5'	5.82	130.22	120.90
85	AA	1933	G	C5'-C4'-C3'	5.82	125.32	116.00
85	AA	2055	G	C5'-C4'-C3'	5.82	125.32	116.00
85	AA	2063	C	O5'-C5'-C4'	-5.82	100.64	111.70
34	BA	501	U	C1'-O4'-C4'	-5.82	105.24	109.90
35	BB	365	U	C2-N1-C1'	-5.82	110.71	117.70
35	BB	1102	U	C4'-C3'-C2'	-5.82	96.78	102.60
35	BB	1200	A	O3'-P-O5'	5.82	115.06	104.00
39	BF	1	C	C4-C5-C6	5.82	120.31	117.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	100	A	N1-C6-N6	-5.82	115.11	118.60
85	AA	1483	A	O4'-C1'-C2'	5.82	112.84	107.60
4	A3	71	PRO	C-N-CA	5.82	136.25	121.70
34	BA	552	C	O5'-P-OP2	5.82	117.69	110.70
34	BA	729	C	P-O5'-C5'	-5.82	111.59	120.90
34	BA	1557	G	C1'-O4'-C4'	-5.82	105.24	109.90
35	BB	315	C	P-O5'-C5'	5.82	130.21	120.90
35	BB	841	U	O4'-C1'-C2'	5.82	112.84	107.60
41	BH	23	G	N1-C6-O6	5.82	123.39	119.90
49	BP	23	ARG	CG-CD-NE	-5.82	99.58	111.80
73	Bn	39	TYR	N-CA-C	-5.82	95.28	111.00
85	AA	602	U	C5'-C4'-C3'	-5.82	106.69	116.00
85	AA	809	A	C5-C6-N6	-5.82	119.04	123.70
85	AA	842	G	C4'-C3'-C2'	-5.82	96.78	102.60
85	AA	1484	G	N1-C2-N2	-5.82	110.96	116.20
85	AA	1497	U	O4'-C4'-C3'	-5.82	98.18	104.00
85	AA	1599	G	O4'-C1'-N9	5.82	112.86	108.20
34	BA	725	C	C1'-O4'-C4'	-5.82	105.25	109.90
35	BB	738	G	C8-N9-C1'	5.82	134.56	127.00
39	BF	2	G	N1-C6-O6	5.82	123.39	119.90
41	BH	128	G	C5-C6-N1	5.82	114.41	111.50
85	AA	412	G	C1'-O4'-C4'	-5.82	105.25	109.90
85	AA	692	U	C6-N1-C2	-5.82	117.51	121.00
85	AA	1204	A	C3'-C2'-C1'	-5.82	96.84	101.50
85	AA	2110	U	C1'-O4'-C4'	-5.82	105.25	109.90
5	A4	191	MET	N-CA-C	5.82	126.71	111.00
34	BA	269	G	N9-C4-C5	5.82	107.73	105.40
34	BA	818	G	C5-C6-N1	5.82	114.41	111.50
34	BA	926	A	N9-C1'-C2'	-5.82	105.60	112.00
34	BA	1500	G	N1-C2-N2	-5.82	110.97	116.20
35	BB	16	G	O4'-C1'-N9	-5.82	103.55	108.20
35	BB	498	G	C5'-C4'-C3'	-5.82	106.69	116.00
36	BC	160	C	C6-N1-C2	-5.82	117.97	120.30
39	BF	27	G	C8-N9-C4	5.82	108.73	106.40
62	Bc	109	ARG	CD-NE-CZ	-5.82	115.46	123.60
85	AA	417	U	P-O3'-C3'	-5.82	112.72	119.70
85	AA	501	A	C8-N9-C1'	5.82	138.17	127.70
85	AA	643	C	N3-C2-O2	-5.82	117.83	121.90
85	AA	2209	U	N1-C2-N3	5.82	118.39	114.90
35	BB	37	C	O3'-P-O5'	5.82	115.05	104.00
35	BB	434	A	N7-C8-N9	-5.82	110.89	113.80
35	BB	1470	G	C6-C5-N7	-5.82	126.91	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1492	C	O4'-C1'-N1	5.82	112.85	108.20
38	BE	89	G	N3-C4-C5	-5.82	125.69	128.60
40	BG	80	G	C4-N9-C1'	-5.82	118.94	126.50
40	BG	176	G	N9-C1'-C2'	-5.82	105.60	112.00
85	AA	179	G	C4-N9-C1'	5.82	134.06	126.50
85	AA	443	A	C3'-C2'-C1'	-5.82	96.85	101.50
85	AA	1666	U	O4'-C4'-C3'	-5.82	98.18	104.00
34	BA	544	U	C5'-C4'-O4'	5.81	116.08	109.10
34	BA	1473	A	N1-C6-N6	-5.81	115.11	118.60
34	BA	1845	G	N1-C6-O6	5.81	123.39	119.90
35	BB	5	A	OP1-P-OP2	-5.81	110.88	119.60
35	BB	100	A	O4'-C1'-N9	5.81	112.85	108.20
35	BB	491	A	O4'-C1'-N9	5.81	112.85	108.20
35	BB	696	G	O4'-C1'-N9	5.81	112.85	108.20
36	BC	110	A	O4'-C1'-N9	5.81	112.85	108.20
38	BE	91	G	O5'-C5'-C4'	-5.81	100.65	111.70
40	BG	22	G	C5'-C4'-O4'	5.81	116.08	109.10
41	BH	117	U	C6-N1-C2	-5.81	117.51	121.00
83	Bx	73	ARG	NE-CZ-NH2	-5.81	117.39	120.30
85	AA	554	A	O4'-C1'-N9	5.81	112.85	108.20
85	AA	847	G	N9-C1'-C2'	-5.81	105.60	112.00
85	AA	1574	C	O5'-P-OP1	-5.81	100.47	105.70
4	A3	88	PHE	N-CA-CB	5.81	121.06	110.60
34	BA	230	A	C5-C6-N6	-5.81	119.05	123.70
34	BA	523	A	P-O3'-C3'	-5.81	112.73	119.70
34	BA	1039	G	N9-C1'-C2'	-5.81	105.61	112.00
34	BA	1156	U	C5'-C4'-C3'	5.81	125.30	116.00
35	BB	11	A	C1'-O4'-C4'	-5.81	105.25	109.90
35	BB	414	C	P-O3'-C3'	5.81	126.67	119.70
35	BB	912	C	C4'-C3'-C2'	-5.81	96.79	102.60
35	BB	961	G	P-O3'-C3'	-5.81	112.72	119.70
35	BB	977	G	N7-C8-N9	5.81	116.01	113.10
37	BD	75	G	N3-C4-N9	-5.81	122.51	126.00
40	BG	118	U	N1-C2-N3	5.81	118.39	114.90
42	BI	110	LEU	CB-CA-C	-5.81	99.16	110.20
44	BK	85	PHE	CB-CG-CD2	-5.81	116.73	120.80
50	BQ	179	ARG	CG-CD-NE	-5.81	99.59	111.80
77	Br	312	ARG	NE-CZ-NH2	-5.81	117.39	120.30
85	AA	246	C	C6-N1-C1'	5.81	127.77	120.80
85	AA	730	G	C2-N3-C4	5.81	114.81	111.90
85	AA	1003	G	O4'-C1'-N9	5.81	112.85	108.20
85	AA	1006	C	N1-C2-N3	5.81	123.27	119.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1679	U	C5'-C4'-C3'	-5.81	106.70	116.00
34	BA	56	G	N3-C2-N2	5.81	123.97	119.90
34	BA	579	U	OP2-P-O3'	5.81	117.98	105.20
34	BA	1208	U	O4'-C1'-C2'	5.81	112.83	107.60
34	BA	1519	G	C5'-C4'-O4'	5.81	116.07	109.10
34	BA	1731	A	O4'-C1'-N9	5.81	112.85	108.20
35	BB	59	U	O5'-P-OP1	-5.81	100.47	105.70
35	BB	103	C	N1-C2-O2	5.81	122.39	118.90
37	BD	2	G	O4'-C1'-N9	5.81	112.85	108.20
37	BD	31	U	C6-N1-C1'	5.81	129.34	121.20
39	BF	7	G	C5-C6-O6	-5.81	125.11	128.60
40	BG	175	G	O3'-P-O5'	5.81	115.04	104.00
53	BT	95	TRP	CB-CG-CD2	-5.81	119.05	126.60
85	AA	765	U	P-O3'-C3'	-5.81	112.73	119.70
34	BA	768	G	C1'-O4'-C4'	-5.81	105.25	109.90
34	BA	877	U	O4'-C4'-C3'	-5.81	98.19	104.00
34	BA	1269	C	C3'-C2'-C1'	-5.81	96.85	101.50
34	BA	1426	A	O5'-P-OP2	5.81	117.67	110.70
34	BA	1545	C	C6-N1-C1'	5.81	127.77	120.80
34	BA	1592	U	C5-C6-N1	-5.81	119.80	122.70
35	BB	566	A	C5'-C4'-C3'	-5.81	106.71	116.00
35	BB	1077	C	C6-N1-C2	-5.81	117.98	120.30
35	BB	1389	C	P-O3'-C3'	-5.81	112.73	119.70
35	BB	1531	G	C4-N9-C1'	-5.81	118.95	126.50
72	Bm	77	ALA	N-CA-CB	5.81	118.23	110.10
80	Bu	141	ASP	CB-CG-OD1	5.81	123.53	118.30
85	AA	78	A	C4'-C3'-C2'	-5.81	96.79	102.60
85	AA	586	G	C8-N9-C1'	5.81	134.55	127.00
34	BA	453	A	C1'-O4'-C4'	-5.81	105.25	109.90
34	BA	678	C	N3-C4-N4	5.81	122.07	118.00
34	BA	1211	G	OP1-P-OP2	-5.81	110.89	119.60
34	BA	1490	U	C5'-C4'-O4'	-5.81	102.13	109.10
35	BB	446	U	C1'-O4'-C4'	5.81	114.55	109.90
35	BB	449	C	N1-C2-N3	5.81	123.27	119.20
35	BB	698	C	C2-N3-C4	-5.81	117.00	119.90
35	BB	1336	G	O3'-P-O5'	-5.81	92.97	104.00
35	BB	1540	U	O4'-C4'-C3'	-5.81	98.19	104.00
38	BE	139	U	O4'-C1'-N1	5.81	112.84	108.20
40	BG	28	A	C8-N9-C4	5.81	108.12	105.80
85	AA	965	G	C4-N9-C1'	-5.81	118.95	126.50
85	AA	995	G	P-O5'-C5'	-5.81	111.61	120.90
85	AA	1235	G	C2-N3-C4	-5.81	109.00	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1717	G	C5-C6-O6	-5.81	125.12	128.60
85	AA	1938	G	O4'-C1'-N9	5.81	112.85	108.20
85	AA	2193	A	C5'-C4'-C3'	-5.81	106.71	116.00
86	AB	22	G	C8-N9-C1'	-5.81	119.45	127.00
34	BA	79	C	O4'-C1'-N1	5.81	112.84	108.20
34	BA	346	A	C1'-O4'-C4'	-5.81	105.25	109.90
34	BA	1491	U	C5'-C4'-O4'	5.81	116.07	109.10
35	BB	1032	U	P-O3'-C3'	5.81	126.67	119.70
35	BB	1098	G	O4'-C1'-C2'	5.81	112.83	107.60
35	BB	1388	A	C4-N9-C1'	-5.81	115.85	126.30
38	BE	11	A	C5-C6-N6	5.81	128.34	123.70
85	AA	278	C	P-O5'-C5'	-5.81	111.61	120.90
85	AA	870	U	O5'-C5'-C4'	-5.81	100.67	111.70
85	AA	1355	U	C5'-C4'-O4'	-5.81	102.13	109.10
85	AA	1922	A	C2'-C3'-O3'	5.81	122.99	113.70
85	AA	2138	G	C6-N1-C2	-5.81	121.62	125.10
34	BA	429	G	N9-C1'-C2'	-5.80	105.61	112.00
34	BA	1404	A	N9-C4-C5	5.80	108.12	105.80
34	BA	1540	C	N3-C4-N4	5.80	122.06	118.00
34	BA	1667	G	N3-C2-N2	5.80	123.96	119.90
35	BB	1056	A	C5'-C4'-C3'	-5.80	106.71	116.00
35	BB	1144	A	N1-C6-N6	-5.80	115.12	118.60
38	BE	10	G	N9-C4-C5	5.80	107.72	105.40
41	BH	34	G	O5'-P-OP2	-5.80	100.48	105.70
41	BH	46	C	O4'-C1'-N1	5.80	112.84	108.20
85	AA	118	C	C1'-O4'-C4'	-5.80	105.26	109.90
85	AA	260	A	P-O3'-C3'	5.80	126.67	119.70
85	AA	595	A	C5'-C4'-C3'	-5.80	106.71	116.00
85	AA	635	G	C1'-O4'-C4'	-5.80	105.26	109.90
85	AA	1361	A	OP2-P-O3'	5.80	117.97	105.20
85	AA	1729	C	C3'-C2'-C1'	-5.80	96.86	101.50
85	AA	1894	G	N3-C2-N2	5.80	123.96	119.90
85	AA	2225	G	P-O5'-C5'	-5.80	111.61	120.90
34	BA	215	C	P-O5'-C5'	5.80	130.19	120.90
34	BA	1278	A	C8-N9-C1'	5.80	138.15	127.70
35	BB	847	U	C4'-C3'-C2'	5.80	108.40	102.60
36	BC	99	U	O4'-C1'-N1	5.80	112.84	108.20
38	BE	31	A	C2'-C3'-O3'	5.80	122.98	113.70
69	Bj	71	HIS	CB-CA-C	5.80	122.01	110.40
85	AA	799	G	P-O5'-C5'	-5.80	111.62	120.90
85	AA	1242	A	N1-C6-N6	-5.80	115.12	118.60
85	AA	1589	G	P-O5'-C5'	-5.80	111.61	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	AT	25	ARG	CD-NE-CZ	-5.80	115.48	123.60
34	BA	221	G	C1'-O4'-C4'	-5.80	105.26	109.90
34	BA	679	U	C2-N3-C4	-5.80	123.52	127.00
34	BA	746	C	N3-C2-O2	-5.80	117.84	121.90
34	BA	1735	G	N3-C2-N2	5.80	123.96	119.90
35	BB	43	G	N9-C4-C5	-5.80	103.08	105.40
35	BB	1133	C	N3-C4-N4	-5.80	113.94	118.00
35	BB	1228	A	C4'-C3'-C2'	5.80	108.40	102.60
35	BB	1511	U	O4'-C1'-N1	5.80	112.84	108.20
38	BE	96	G	C6-C5-N7	-5.80	126.92	130.40
40	BG	172	C	C4'-C3'-C2'	-5.80	96.80	102.60
47	BN	102	VAL	N-CA-CB	-5.80	98.74	111.50
85	AA	14	C	N3-C4-N4	-5.80	113.94	118.00
85	AA	501	A	N1-C6-N6	-5.80	115.12	118.60
85	AA	593	U	O4'-C1'-N1	5.80	112.84	108.20
85	AA	1238	U	N1-C2-N3	5.80	118.38	114.90
85	AA	1495	G	C8-N9-C1'	5.80	134.54	127.00
85	AA	1628	U	O4'-C1'-N1	5.80	112.84	108.20
33	AZ	62	ARG	NE-CZ-NH1	5.80	123.20	120.30
34	BA	291	C	C1'-O4'-C4'	-5.80	105.26	109.90
34	BA	684	G	O4'-C1'-C2'	-5.80	100.00	105.80
34	BA	1184	A	C8-N9-C4	-5.80	103.48	105.80
34	BA	1316	G	C8-N9-C1'	-5.80	119.46	127.00
34	BA	1509	U	N1-C2-O2	5.80	126.86	122.80
34	BA	1642	A	C5-C6-N1	5.80	120.60	117.70
35	BB	1030	U	C5'-C4'-O4'	5.80	116.06	109.10
35	BB	1038	G	C4'-C3'-C2'	5.80	108.40	102.60
35	BB	1098	G	C1'-O4'-C4'	-5.80	105.26	109.90
35	BB	1358	A	P-O3'-C3'	-5.80	112.74	119.70
38	BE	208	G	C4'-C3'-C2'	-5.80	96.80	102.60
41	BH	109	G	C4-C5-C6	-5.80	115.32	118.80
42	BI	49	HIS	CB-CA-C	5.80	122.00	110.40
42	BI	94	LEU	CA-C-N	-5.80	104.44	117.20
85	AA	469	G	C2-N3-C4	-5.80	109.00	111.90
85	AA	484	G	O4'-C4'-C3'	-5.80	98.20	104.00
85	AA	571	G	C8-N9-C1'	5.80	134.54	127.00
85	AA	1047	G	C8-N9-C1'	5.80	134.54	127.00
85	AA	1535	C	O4'-C1'-C2'	-5.80	100.00	105.80
85	AA	1574	C	P-O3'-C3'	-5.80	112.74	119.70
85	AA	1692	U	C6-N1-C1'	5.80	129.32	121.20
85	AA	2216	A	N1-C6-N6	-5.80	115.12	118.60
6	A5	191	ARG	NE-CZ-NH1	5.80	123.20	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	527	C	C4'-C3'-C2'	-5.80	96.80	102.60
34	BA	1745	G	P-O5'-C5'	5.80	130.18	120.90
35	BB	1130	U	N1-C2-O2	5.80	126.86	122.80
35	BB	1226	G	N9-C1'-C2'	-5.80	105.62	112.00
37	BD	32	A	P-O3'-C3'	-5.80	112.74	119.70
84	By	10	THR	CA-CB-CG2	5.80	120.52	112.40
85	AA	976	G	O4'-C1'-C2'	5.80	112.82	107.60
85	AA	1061	C	C5'-C4'-C3'	5.80	125.28	116.00
85	AA	1270	C	C2-N3-C4	-5.80	117.00	119.90
85	AA	1884	A	C4'-C3'-C2'	-5.80	96.80	102.60
85	AA	2157	G	N3-C2-N2	-5.80	115.84	119.90
6	A5	60	LEU	CB-CA-C	-5.80	99.19	110.20
12	AD	39	GLY	N-CA-C	-5.80	98.61	113.10
34	BA	216	C	C6-N1-C1'	5.80	127.75	120.80
34	BA	430	A	O4'-C4'-C3'	-5.80	98.20	104.00
34	BA	962	U	O4'-C1'-N1	5.80	112.84	108.20
34	BA	1170	A	C5'-C4'-C3'	5.80	125.28	116.00
34	BA	1189	A	C8-N9-C1'	-5.80	117.27	127.70
34	BA	1540	C	C6-N1-C1'	-5.80	113.84	120.80
34	BA	1811	A	C2-N3-C4	-5.80	107.70	110.60
35	BB	643	G	C5-C6-O6	5.80	132.08	128.60
35	BB	842	G	O4'-C1'-N9	5.80	112.84	108.20
40	BG	182	G	C4-N9-C1'	5.80	134.04	126.50
85	AA	756	G	C5'-C4'-C3'	-5.80	106.73	116.00
85	AA	1096	G	N3-C4-C5	-5.80	125.70	128.60
85	AA	1097	G	O4'-C1'-N9	5.80	112.84	108.20
85	AA	1497	U	C4'-C3'-C2'	5.80	108.40	102.60
85	AA	1618	G	C5'-C4'-C3'	-5.80	106.72	116.00
85	AA	1902	C	C2-N3-C4	-5.80	117.00	119.90
34	BA	196	A	C5'-C4'-O4'	5.79	116.05	109.10
35	BB	26	C	N3-C4-C5	5.79	124.22	121.90
41	BH	85	C	OP1-P-O3'	5.79	117.95	105.20
47	BN	80	PHE	CB-CG-CD1	-5.79	116.74	120.80
85	AA	82	A	N9-C1'-C2'	-5.79	105.63	112.00
85	AA	1075	U	O4'-C1'-N1	5.79	112.84	108.20
85	AA	1765	G	N1-C6-O6	5.79	123.38	119.90
34	BA	75	U	C5-C6-N1	-5.79	119.80	122.70
34	BA	300	C	C2-N3-C4	-5.79	117.00	119.90
34	BA	661	C	C6-N1-C2	-5.79	117.98	120.30
34	BA	798	G	C5-N7-C8	-5.79	101.40	104.30
34	BA	1270	G	C1'-O4'-C4'	-5.79	105.27	109.90
38	BE	39	U	N3-C2-O2	-5.79	118.14	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	53	C	C5'-C4'-C3'	-5.79	106.73	116.00
40	BG	131	U	C5-C6-N1	-5.79	119.80	122.70
85	AA	388	G	O4'-C1'-N9	5.79	112.83	108.20
85	AA	859	G	C5'-C4'-C3'	-5.79	106.73	116.00
85	AA	1224	C	C2-N1-C1'	-5.79	112.43	118.80
85	AA	1490	A	N1-C6-N6	5.79	122.08	118.60
85	AA	2086	C	C4'-C3'-C2'	5.79	108.39	102.60
34	BA	134	U	O4'-C1'-N1	5.79	112.83	108.20
34	BA	329	G	C5'-C4'-C3'	5.79	125.27	116.00
34	BA	574	U	N1-C2-O2	5.79	126.85	122.80
34	BA	694	G	N1-C2-N2	-5.79	110.99	116.20
34	BA	953	G	O5'-P-OP1	-5.79	100.49	105.70
34	BA	1148	U	O5'-C5'-C4'	-5.79	100.69	111.70
35	BB	948	G	N1-C6-O6	5.79	123.38	119.90
35	BB	1005	A	C4-N9-C1'	-5.79	115.87	126.30
35	BB	1231	U	O5'-P-OP1	-5.79	100.49	105.70
35	BB	1289	G	C5-C6-N1	5.79	114.39	111.50
35	BB	1431	G	O5'-C5'-C4'	-5.79	100.70	111.70
36	BC	58	G	C5-C6-O6	-5.79	125.12	128.60
40	BG	41	U	O4'-C1'-N1	5.79	112.83	108.20
77	Br	297	ARG	NE-CZ-NH1	5.79	123.20	120.30
85	AA	557	G	C4-C5-N7	-5.79	108.48	110.80
85	AA	661	C	C5'-C4'-O4'	5.79	116.05	109.10
85	AA	816	A	P-O5'-C5'	5.79	130.17	120.90
85	AA	1413	G	C5-C6-O6	-5.79	125.12	128.60
85	AA	1522	U	P-O3'-C3'	-5.79	112.75	119.70
34	BA	500	C	O4'-C1'-C2'	5.79	112.81	107.60
34	BA	652	C	C1'-O4'-C4'	-5.79	105.27	109.90
34	BA	712	C	C5'-C4'-C3'	-5.79	106.73	116.00
34	BA	871	G	OP1-P-O3'	-5.79	92.46	105.20
34	BA	1620	U	C4'-C3'-O3'	-5.79	97.24	109.40
34	BA	1688	G	C4-N9-C1'	-5.79	118.97	126.50
35	BB	421	U	C1'-O4'-C4'	-5.79	105.27	109.90
35	BB	871	C	C5'-C4'-C3'	-5.79	106.74	116.00
35	BB	1358	A	C6-N1-C2	-5.79	115.13	118.60
41	BH	21	G	N1-C2-N2	-5.79	110.99	116.20
41	BH	122	U	P-O3'-C3'	-5.79	112.75	119.70
61	Bb	135	ARG	NE-CZ-NH1	5.79	123.19	120.30
65	Bf	352	ALA	N-CA-CB	5.79	118.21	110.10
85	AA	311	U	C5'-C4'-C3'	5.79	125.26	116.00
85	AA	2132	A	C2'-C3'-O3'	5.79	122.96	113.70
34	BA	477	C	N3-C4-N4	-5.79	113.95	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	771	A	C5'-C4'-O4'	5.79	116.05	109.10
34	BA	814	C	C4'-C3'-C2'	-5.79	96.81	102.60
34	BA	1522	G	C5-C6-O6	-5.79	125.13	128.60
34	BA	1809	G	N1-C6-O6	5.79	123.37	119.90
35	BB	588	A	N9-C1'-C2'	-5.79	105.63	112.00
35	BB	794	G	C8-N9-C1'	5.79	134.53	127.00
35	BB	879	G	O5'-C5'-C4'	5.79	122.70	111.70
35	BB	1187	G	C8-N9-C1'	5.79	134.52	127.00
38	BE	36	U	C6-N1-C1'	5.79	129.30	121.20
40	BG	105	A	P-O3'-C3'	5.79	126.65	119.70
85	AA	1185	G	O5'-P-OP2	5.79	117.65	110.70
85	AA	2196	G	C4-C5-N7	-5.79	108.48	110.80
85	AA	2236	U	P-O3'-C3'	-5.79	112.75	119.70
86	AB	68	C	O4'-C1'-N1	5.79	112.83	108.20
34	BA	706	C	C2-N3-C4	-5.79	117.01	119.90
34	BA	1034	U	N3-C4-O4	-5.79	115.35	119.40
34	BA	1376	U	C2-N1-C1'	-5.79	110.76	117.70
34	BA	1426	A	P-O3'-C3'	-5.79	112.75	119.70
35	BB	156	G	O4'-C1'-N9	5.79	112.83	108.20
35	BB	897	C	P-O3'-C3'	5.79	126.64	119.70
39	BF	52	A	N7-C8-N9	5.79	116.69	113.80
41	BH	20	A	C1'-O4'-C4'	-5.79	105.27	109.90
72	Bm	85	LEU	CB-CG-CD1	5.79	120.84	111.00
85	AA	369	A	C4-N9-C1'	-5.79	115.88	126.30
85	AA	1310	G	C2'-C3'-O3'	5.79	122.96	113.70
34	BA	146	G	N3-C2-N2	-5.79	115.85	119.90
34	BA	621	G	C5-C6-O6	-5.79	125.13	128.60
34	BA	1019	C	C3'-C2'-C1'	-5.79	96.87	101.50
34	BA	1471	U	N3-C2-O2	-5.79	118.15	122.20
34	BA	1794	A	O5'-C5'-C4'	5.79	122.69	111.70
35	BB	810	G	N9-C1'-C2'	-5.79	105.64	112.00
35	BB	971	A	C5-N7-C8	-5.79	101.01	103.90
35	BB	1351	G	C4-N9-C1'	-5.79	118.98	126.50
35	BB	1495	U	C3'-C2'-C1'	-5.79	96.87	101.50
36	BC	61	A	C4'-C3'-C2'	-5.79	96.81	102.60
38	BE	14	C	C5'-C4'-C3'	-5.79	106.74	116.00
39	BF	39	C	N3-C2-O2	-5.79	117.85	121.90
85	AA	361	U	P-O3'-C3'	-5.79	112.76	119.70
85	AA	474	C	C3'-C2'-C1'	-5.79	96.87	101.50
85	AA	893	G	N1-C2-N3	-5.79	120.43	123.90
85	AA	971	U	C6-N1-C2	-5.79	117.53	121.00
85	AA	1703	A	C4-N9-C1'	-5.79	115.89	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1866	A	C5'-C4'-C3'	-5.79	106.74	116.00
86	AB	63	G	N1-C6-O6	5.79	123.37	119.90
86	AB	71	G	C5'-C4'-O4'	5.79	116.04	109.10
33	AZ	60	ARG	NE-CZ-NH1	5.78	123.19	120.30
34	BA	711	C	N1-C2-O2	5.78	122.37	118.90
34	BA	1711	G	C8-N9-C4	5.78	108.71	106.40
35	BB	137	A	C5-C6-N1	5.78	120.59	117.70
35	BB	139	G	C1'-O4'-C4'	-5.78	105.27	109.90
35	BB	788	U	C5'-C4'-O4'	-5.78	102.16	109.10
35	BB	1250	A	C4-N9-C1'	-5.78	115.89	126.30
36	BC	6	G	C5-N7-C8	-5.78	101.41	104.30
36	BC	97	U	C5'-C4'-C3'	5.78	125.25	116.00
40	BG	117	C	C4'-C3'-C2'	-5.78	96.82	102.60
85	AA	85	U	O5'-C5'-C4'	-5.78	100.71	111.70
85	AA	727	U	C4'-C3'-C2'	-5.78	96.82	102.60
85	AA	1105	G	N1-C6-O6	-5.78	116.43	119.90
85	AA	1511	C	C1'-O4'-C4'	-5.78	105.27	109.90
85	AA	1616	U	C2-N1-C1'	-5.78	110.76	117.70
85	AA	1854	U	C5-C4-O4	-5.78	122.43	125.90
85	AA	2119	C	P-O3'-C3'	5.78	126.64	119.70
26	AS	119	ARG	CB-CA-C	5.78	121.96	110.40
34	BA	245	U	N1-C2-N3	5.78	118.37	114.90
34	BA	1252	G	C4-N9-C1'	-5.78	118.98	126.50
35	BB	69	A	C5'-C4'-C3'	-5.78	106.75	116.00
85	AA	253	C	C5'-C4'-O4'	5.78	116.04	109.10
85	AA	596	A	P-O3'-C3'	-5.78	112.76	119.70
85	AA	2029	G	C4'-C3'-C2'	-5.78	96.82	102.60
34	BA	213	A	C4-N9-C1'	-5.78	115.90	126.30
34	BA	1408	C	N3-C2-O2	-5.78	117.85	121.90
34	BA	1837	U	N3-C4-O4	-5.78	115.35	119.40
35	BB	1112	U	P-O3'-C3'	5.78	126.64	119.70
35	BB	1244	U	P-O3'-C3'	-5.78	112.76	119.70
39	BF	56	C	C6-N1-C1'	5.78	127.74	120.80
40	BG	3	G	C3'-C2'-C1'	-5.78	96.88	101.50
41	BH	20	A	C4'-C3'-C2'	-5.78	96.82	102.60
41	BH	76	G	P-O3'-C3'	5.78	126.64	119.70
62	Bc	108	ARG	NE-CZ-NH1	5.78	123.19	120.30
85	AA	1209	U	C2-N3-C4	-5.78	123.53	127.00
85	AA	1292	A	C5'-C4'-C3'	-5.78	106.75	116.00
85	AA	1585	A	C1'-O4'-C4'	-5.78	105.28	109.90
85	AA	1845	G	O4'-C1'-N9	5.78	112.82	108.20
85	AA	1970	A	C4-N9-C1'	-5.78	115.90	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
5	A4	158	PHE	CA-CB-CG	-5.78	100.03	113.90
34	BA	47	U	C5'-C4'-C3'	5.78	125.25	116.00
34	BA	717	U	C2-N3-C4	-5.78	123.53	127.00
34	BA	744	G	O4'-C1'-N9	5.78	112.82	108.20
34	BA	770	G	C8-N9-C4	-5.78	104.09	106.40
35	BB	794	G	C4-N9-C1'	-5.78	118.99	126.50
35	BB	1080	U	C2-N1-C1'	-5.78	110.77	117.70
35	BB	1380	G	OP2-P-O3'	5.78	117.91	105.20
36	BC	139	A	C3'-C2'-C1'	-5.78	96.88	101.50
41	BH	130	G	C3'-C2'-C1'	-5.78	96.88	101.50
85	AA	1034	U	C1'-O4'-C4'	-5.78	105.28	109.90
85	AA	1058	G	C5'-C4'-O4'	5.78	116.03	109.10
85	AA	1573	A	C5-C6-N1	-5.78	114.81	117.70
14	AF	41	ARG	NE-CZ-NH2	-5.78	117.41	120.30
34	BA	376	U	C5'-C4'-O4'	5.78	116.03	109.10
34	BA	662	U	C5-C6-N1	-5.78	119.81	122.70
34	BA	805	A	O4'-C4'-C3'	-5.78	98.22	104.00
35	BB	112	G	N1-C6-O6	5.78	123.37	119.90
35	BB	131	A	C1'-O4'-C4'	-5.78	105.28	109.90
35	BB	662	G	C4-N9-C1'	-5.78	118.99	126.50
35	BB	1164	U	P-O3'-C3'	-5.78	112.77	119.70
35	BB	1374	U	O4'-C1'-N1	5.78	112.82	108.20
37	BD	39	C	P-O5'-C5'	-5.78	111.66	120.90
38	BE	174	U	C1'-O4'-C4'	-5.78	105.28	109.90
82	Bw	238	GLU	CA-CB-CG	5.78	126.11	113.40
85	AA	56	U	C5'-C4'-C3'	5.78	125.24	116.00
85	AA	157	G	N9-C1'-C2'	-5.78	105.64	112.00
85	AA	486	G	O4'-C1'-C2'	5.78	112.80	107.60
85	AA	570	U	C4'-C3'-C2'	-5.78	96.82	102.60
85	AA	716	G	N9-C1'-C2'	-5.78	105.64	112.00
85	AA	1103	A	C2'-C3'-O3'	5.78	122.94	113.70
85	AA	1213	U	N3-C2-O2	-5.78	118.16	122.20
85	AA	1342	C	C4'-C3'-C2'	-5.78	96.82	102.60
85	AA	1441	G	P-O5'-C5'	-5.78	111.66	120.90
85	AA	1837	U	P-O3'-C3'	-5.78	112.77	119.70
85	AA	2120	C	N1-C1'-C2'	-5.78	105.64	112.00
11	AC	124	PHE	CB-CA-C	-5.78	98.85	110.40
18	AJ	78	ARG	NE-CZ-NH1	5.78	123.19	120.30
34	BA	395	G	C5'-C4'-O4'	5.78	116.03	109.10
34	BA	487	A	C5-C6-N1	5.78	120.59	117.70
34	BA	522	C	C4'-C3'-C2'	-5.78	96.83	102.60
35	BB	448	G	C3'-C2'-C1'	-5.78	96.88	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	836	U	N1-C2-N3	5.78	118.36	114.90
35	BB	1119	G	C4-N9-C1'	-5.78	118.99	126.50
35	BB	1366	C	O4'-C1'-N1	5.78	112.82	108.20
37	BD	3	G	C5'-C4'-C3'	-5.78	106.76	116.00
38	BE	206	G	O4'-C4'-C3'	-5.78	98.22	104.00
40	BG	139	U	C3'-C2'-C1'	-5.78	96.88	101.50
65	Bf	393	VAL	CB-CA-C	-5.78	100.43	111.40
85	AA	69	C	O4'-C1'-C2'	5.78	112.80	107.60
85	AA	378	A	O4'-C1'-N9	5.78	112.82	108.20
85	AA	718	C	P-O3'-C3'	-5.78	112.77	119.70
85	AA	824	C	N1-C1'-C2'	-5.78	105.65	112.00
85	AA	1066	U	O4'-C1'-N1	5.78	112.82	108.20
85	AA	1305	A	C3'-C2'-C1'	-5.78	96.88	101.50
85	AA	1713	A	C5'-C4'-C3'	-5.78	106.76	116.00
4	A3	231	HIS	CA-CB-CG	-5.77	103.78	113.60
24	AQ	75	ASP	N-CA-C	5.77	126.59	111.00
34	BA	1829	A	C5'-C4'-C3'	-5.77	106.76	116.00
35	BB	517	G	N1-C6-O6	-5.77	116.44	119.90
35	BB	566	A	C5-C6-N1	5.77	120.59	117.70
35	BB	1467	A	O4'-C4'-C3'	-5.77	98.23	104.00
36	BC	76	C	C1'-O4'-C4'	-5.77	105.28	109.90
77	Br	270	THR	CA-CB-OG1	5.77	121.12	109.00
85	AA	14	C	C3'-C2'-C1'	-5.77	96.88	101.50
85	AA	860	C	N1-C2-N3	5.77	123.24	119.20
85	AA	873	U	O3'-P-O5'	5.77	114.97	104.00
85	AA	1571	A	C4'-C3'-C2'	-5.77	96.83	102.60
85	AA	1732	G	C5-C6-N1	5.77	114.39	111.50
21	AM	119	ARG	N-CA-C	5.77	126.58	111.00
34	BA	554	A	N7-C8-N9	-5.77	110.91	113.80
34	BA	1616	A	N9-C4-C5	5.77	108.11	105.80
35	BB	849	A	O4'-C1'-N9	5.77	112.82	108.20
35	BB	1379	U	C4'-C3'-C2'	-5.77	96.83	102.60
35	BB	1487	G	N3-C4-C5	-5.77	125.71	128.60
36	BC	43	A	C3'-C2'-C1'	-5.77	96.88	101.50
36	BC	82	C	C5'-C4'-O4'	5.77	116.03	109.10
38	BE	119	U	C6-N1-C1'	5.77	129.28	121.20
47	BN	9	PRO	CA-N-CD	-5.77	103.42	111.50
85	AA	928	U	N3-C4-C5	5.77	118.06	114.60
86	AB	2	C	C6-N1-C2	-5.77	117.99	120.30
34	BA	85	C	O4'-C1'-N1	5.77	112.82	108.20
35	BB	444	U	C5-C6-N1	-5.77	119.81	122.70
37	BD	115	A	C1'-O4'-C4'	-5.77	105.28	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	135	U	C2-N1-C1'	-5.77	110.78	117.70
45	BL	170	PHE	CB-CG-CD2	-5.77	116.76	120.80
64	Be	71	ARG	NE-CZ-NH2	-5.77	117.41	120.30
68	Bi	47	TYR	CB-CG-CD1	-5.77	117.54	121.00
85	AA	957	A	P-O3'-C3'	-5.77	112.77	119.70
85	AA	1198	U	C2-N1-C1'	-5.77	110.78	117.70
16	AH	131	ASP	N-CA-C	5.77	126.58	111.00
34	BA	63	A	C3'-C2'-C1'	-5.77	96.88	101.50
34	BA	523	A	C8-N9-C1'	5.77	138.09	127.70
34	BA	868	C	N1-C2-O2	5.77	122.36	118.90
34	BA	1397	C	C5-C6-N1	5.77	123.89	121.00
34	BA	1504	A	C8-N9-C4	-5.77	103.49	105.80
34	BA	1682	A	C8-N9-C4	-5.77	103.49	105.80
35	BB	560	C	C6-N1-C1'	5.77	127.72	120.80
35	BB	775	U	C1'-O4'-C4'	-5.77	105.28	109.90
35	BB	991	C	C4'-C3'-C2'	-5.77	96.83	102.60
35	BB	1014	U	C3'-C2'-C1'	-5.77	96.88	101.50
38	BE	69	C	O4'-C1'-N1	5.77	112.81	108.20
62	Bc	13	ARG	NE-CZ-NH1	5.77	123.19	120.30
85	AA	570	U	C6-N1-C2	-5.77	117.54	121.00
85	AA	752	C	O4'-C1'-N1	5.77	112.81	108.20
85	AA	1168	C	C5'-C4'-C3'	5.77	125.23	116.00
85	AA	1645	G	N1-C6-O6	5.77	123.36	119.90
85	AA	1730	C	O4'-C1'-N1	5.77	112.81	108.20
85	AA	1815	U	O4'-C1'-N1	5.77	112.82	108.20
85	AA	1916	A	C5-N7-C8	-5.77	101.02	103.90
4	A3	209	ARG	NE-CZ-NH1	5.77	123.18	120.30
11	AC	75	HIS	N-CA-CB	5.77	120.98	110.60
34	BA	186	G	O4'-C1'-N9	5.77	112.81	108.20
34	BA	407	A	O5'-P-OP2	-5.77	100.51	105.70
34	BA	544	U	C1'-O4'-C4'	-5.77	105.29	109.90
34	BA	567	U	O4'-C1'-N1	5.77	112.81	108.20
34	BA	922	C	C2-N3-C4	-5.77	117.02	119.90
34	BA	1421	A	C4'-C3'-C2'	5.77	108.37	102.60
35	BB	135	C	C1'-O4'-C4'	-5.77	105.28	109.90
35	BB	143	G	C4-N9-C1'	-5.77	119.00	126.50
35	BB	389	G	C5'-C4'-C3'	-5.77	106.77	116.00
35	BB	637	G	N7-C8-N9	5.77	115.98	113.10
35	BB	812	G	C3'-C2'-C1'	-5.77	96.89	101.50
35	BB	1178	A	C1'-O4'-C4'	-5.77	105.29	109.90
36	BC	15	G	C1'-O4'-C4'	-5.77	105.29	109.90
36	BC	21	U	O4'-C1'-C2'	5.77	112.79	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	23	G	C3'-C2'-C1'	-5.77	96.89	101.50
40	BG	30	C	C2'-C3'-O3'	5.77	122.93	113.70
48	BO	28	ASP	CB-CG-OD1	5.77	123.49	118.30
59	BZ	111	ARG	NE-CZ-NH1	5.77	123.18	120.30
65	Bf	406	ARG	CD-NE-CZ	5.77	131.67	123.60
85	AA	324	U	O4'-C1'-N1	5.77	112.81	108.20
85	AA	534	A	C5'-C4'-C3'	5.77	125.23	116.00
85	AA	538	A	P-O5'-C5'	5.77	130.13	120.90
85	AA	764	U	C5'-C4'-C3'	-5.77	106.77	116.00
85	AA	922	A	C8-N9-C4	5.77	108.11	105.80
85	AA	1540	A	O3'-P-O5'	-5.77	93.04	104.00
85	AA	1708	A	C4-N9-C1'	-5.77	115.92	126.30
85	AA	1734	A	C6-N1-C2	-5.77	115.14	118.60
85	AA	1811	C	N3-C4-N4	5.77	122.04	118.00
16	AH	91	ARG	NE-CZ-NH1	5.77	123.18	120.30
34	BA	398	G	C4-N9-C1'	-5.77	119.00	126.50
34	BA	749	G	P-O5'-C5'	-5.77	111.67	120.90
34	BA	1001	G	P-O3'-C3'	-5.77	112.78	119.70
34	BA	1565	U	O4'-C1'-N1	5.77	112.81	108.20
34	BA	1591	G	O5'-C5'-C4'	-5.77	100.75	111.70
34	BA	1647	G	O5'-C5'-C4'	-5.77	100.75	111.70
34	BA	1727	A	C5-C6-N6	5.77	128.31	123.70
35	BB	436	G	C2-N3-C4	-5.77	109.02	111.90
35	BB	588	A	C4'-C3'-C2'	-5.77	96.83	102.60
35	BB	755	A	O4'-C1'-N9	5.77	112.81	108.20
35	BB	787	A	O4'-C1'-C2'	5.77	112.79	107.60
40	BG	108	G	P-O3'-C3'	-5.77	112.78	119.70
85	AA	1004	G	P-O5'-C5'	5.77	130.12	120.90
85	AA	2057	G	C5'-C4'-O4'	5.77	116.02	109.10
34	BA	11	U	C5-C6-N1	-5.76	119.82	122.70
34	BA	120	A	OP1-P-OP2	-5.76	110.95	119.60
35	BB	1053	G	N9-C4-C5	-5.76	103.09	105.40
35	BB	1306	G	N9-C1'-C2'	-5.76	105.66	112.00
36	BC	162	C	C4'-C3'-C2'	-5.76	96.83	102.60
37	BD	115	A	C5-C6-N6	-5.76	119.09	123.70
38	BE	36	U	C5'-C4'-O4'	-5.76	102.18	109.10
38	BE	56	U	C5'-C4'-C3'	5.76	125.22	116.00
39	BF	32	G	C8-N9-C1'	-5.76	119.51	127.00
39	BF	37	C	C5-C6-N1	5.76	123.88	121.00
40	BG	4	A	C5-C6-N6	5.76	128.31	123.70
41	BH	2	U	C2-N3-C4	-5.76	123.54	127.00
85	AA	169	G	C4-N9-C1'	-5.76	119.01	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	463	G	C8-N9-C4	5.76	108.71	106.40
85	AA	538	A	C8-N9-C1'	-5.76	117.33	127.70
85	AA	648	G	C1'-O4'-C4'	-5.76	105.29	109.90
85	AA	1272	G	O4'-C1'-N9	5.76	112.81	108.20
85	AA	1846	G	C4-N9-C1'	5.76	134.00	126.50
85	AA	1915	C	O4'-C1'-N1	5.76	112.81	108.20
85	AA	2019	G	C5-C6-O6	-5.76	125.14	128.60
85	AA	2089	G	O3'-P-O5'	5.76	114.95	104.00
34	BA	62	A	C8-N9-C4	5.76	108.11	105.80
34	BA	1105	A	O3'-P-O5'	-5.76	93.05	104.00
38	BE	58	U	N1-C2-N3	-5.76	111.44	114.90
39	BF	51	C	O4'-C4'-C3'	5.76	110.71	106.10
80	Bu	15	ARG	NE-CZ-NH2	-5.76	117.42	120.30
85	AA	596	A	O4'-C1'-N9	5.76	112.81	108.20
85	AA	767	A	O4'-C1'-C2'	5.76	112.79	107.60
85	AA	1480	C	C1'-O4'-C4'	-5.76	105.29	109.90
85	AA	1984	A	N9-C1'-C2'	-5.76	105.66	112.00
2	A1	206	ARG	NE-CZ-NH2	-5.76	117.42	120.30
34	BA	3	G	C5'-C4'-O4'	-5.76	102.19	109.10
34	BA	429	G	N1-C6-O6	5.76	123.36	119.90
34	BA	668	G	C4-N9-C1'	5.76	133.99	126.50
34	BA	903	C	C4'-C3'-C2'	5.76	108.36	102.60
34	BA	913	U	O4'-C1'-N1	5.76	112.81	108.20
34	BA	1241	U	C5'-C4'-C3'	5.76	125.22	116.00
34	BA	1559	C	P-O3'-C3'	-5.76	112.78	119.70
34	BA	1734	U	C6-N1-C2	-5.76	117.54	121.00
35	BB	566	A	C5'-C4'-O4'	5.76	116.02	109.10
35	BB	1066	G	P-O3'-C3'	-5.76	112.79	119.70
41	BH	63	G	N3-C4-N9	5.76	129.46	126.00
80	Bu	164	ARG	NE-CZ-NH1	5.76	123.18	120.30
85	AA	1	G	O4'-C4'-C3'	-5.76	98.24	104.00
85	AA	608	A	C2-N3-C4	5.76	113.48	110.60
85	AA	754	C	P-O3'-C3'	5.76	126.61	119.70
85	AA	755	G	C5-C6-O6	-5.76	125.14	128.60
85	AA	1631	C	O4'-C1'-N1	5.76	112.81	108.20
85	AA	1815	U	P-O3'-C3'	-5.76	112.79	119.70
85	AA	2208	G	N7-C8-N9	5.76	115.98	113.10
34	BA	114	U	C2-N3-C4	-5.76	123.54	127.00
34	BA	240	C	C3'-C2'-C1'	-5.76	96.89	101.50
34	BA	352	G	C5-C6-N1	5.76	114.38	111.50
34	BA	576	C	O5'-P-OP1	5.76	117.61	110.70
34	BA	633	G	OP1-P-O3'	5.76	117.87	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	680	C	C5'-C4'-O4'	5.76	116.01	109.10
34	BA	688	G	C6-N1-C2	-5.76	121.64	125.10
34	BA	1559	C	N1-C2-O2	5.76	122.36	118.90
35	BB	42	A	P-O5'-C5'	5.76	130.11	120.90
35	BB	88	U	O3'-P-O5'	5.76	114.94	104.00
35	BB	498	G	C1'-O4'-C4'	-5.76	105.29	109.90
35	BB	565	U	C2-N1-C1'	5.76	124.61	117.70
35	BB	687	C	O5'-C5'-C4'	-5.76	100.75	111.70
35	BB	711	C	O4'-C1'-N1	5.76	112.81	108.20
35	BB	799	A	C6-N1-C2	-5.76	115.14	118.60
35	BB	807	U	P-O5'-C5'	5.76	130.12	120.90
35	BB	854	G	C8-N9-C4	-5.76	104.10	106.40
36	BC	49	G	N3-C2-N2	5.76	123.93	119.90
39	BF	19	A	C5'-C4'-O4'	5.76	116.01	109.10
41	BH	133	U	O5'-C5'-C4'	-5.76	100.76	111.70
47	BN	44	ARG	CA-CB-CG	5.76	126.07	113.40
85	AA	119	G	N3-C2-N2	-5.76	115.87	119.90
85	AA	365	G	O4'-C4'-C3'	-5.76	98.24	104.00
85	AA	651	G	N1-C6-O6	5.76	123.36	119.90
85	AA	1858	G	O4'-C1'-N9	5.76	112.81	108.20
18	AJ	43	LYS	CB-CA-C	-5.76	98.89	110.40
34	BA	710	A	O4'-C1'-N9	5.76	112.81	108.20
34	BA	1366	C	C2-N1-C1'	5.76	125.13	118.80
34	BA	1489	U	C1'-O4'-C4'	-5.76	105.29	109.90
40	BG	168	A	C2'-C3'-O3'	5.76	122.91	113.70
85	AA	1570	A	O4'-C1'-N9	5.76	112.81	108.20
85	AA	2114	U	C2-N1-C1'	-5.76	110.79	117.70
85	AA	2163	G	O4'-C1'-N9	5.76	112.81	108.20
8	A7	81	PHE	CB-CG-CD2	-5.76	116.77	120.80
34	BA	39	C	P-O3'-C3'	-5.76	112.79	119.70
34	BA	233	U	C6-N1-C1'	5.76	129.26	121.20
34	BA	758	G	C5'-C4'-O4'	5.76	116.01	109.10
34	BA	878	G	C4-N9-C1'	-5.76	119.02	126.50
34	BA	908	G	C4-N9-C1'	-5.76	119.02	126.50
34	BA	955	G	P-O3'-C3'	-5.76	112.79	119.70
34	BA	1826	C	P-O5'-C5'	5.76	130.11	120.90
35	BB	824	C	C3'-C2'-C1'	-5.76	96.89	101.50
35	BB	1174	C	O4'-C1'-N1	5.76	112.81	108.20
41	BH	57	A	C8-N9-C1'	5.76	138.06	127.70
85	AA	10	G	P-O5'-C5'	-5.76	111.69	120.90
85	AA	453	G	C2'-C3'-O3'	5.76	122.91	113.70
85	AA	1026	U	C5'-C4'-O4'	5.76	116.01	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1392	C	O4'-C1'-N1	5.76	112.81	108.20
36	BC	126	G	C3'-C2'-C1'	-5.75	96.90	101.50
38	BE	116	U	C3'-C2'-C1'	-5.75	96.90	101.50
40	BG	13	A	C4'-C3'-C2'	-5.75	96.84	102.60
85	AA	1191	G	C8-N9-C4	5.75	108.70	106.40
85	AA	1248	U	N3-C2-O2	-5.75	118.17	122.20
34	BA	572	G	C5'-C4'-O4'	-5.75	102.19	109.10
34	BA	877	U	N1-C2-O2	5.75	126.83	122.80
34	BA	983	A	C4'-C3'-C2'	5.75	108.35	102.60
34	BA	1381	A	O4'-C1'-N9	5.75	112.80	108.20
34	BA	1450	G	C1'-O4'-C4'	-5.75	105.30	109.90
35	BB	535	U	C3'-C2'-C1'	-5.75	96.90	101.50
35	BB	1046	C	N3-C2-O2	-5.75	117.87	121.90
35	BB	1084	A	C5'-C4'-C3'	-5.75	106.80	116.00
35	BB	1462	G	C8-N9-C4	-5.75	104.10	106.40
53	BT	110	ARG	NE-CZ-NH2	-5.75	117.42	120.30
54	BU	34	PHE	CB-CG-CD1	-5.75	116.77	120.80
54	BU	74	VAL	CB-CA-C	-5.75	100.47	111.40
83	Bx	46	ASP	N-CA-C	5.75	126.53	111.00
85	AA	24	U	O4'-C1'-C2'	5.75	112.78	107.60
85	AA	343	U	O4'-C1'-N1	5.75	112.80	108.20
85	AA	486	G	C2-N3-C4	5.75	114.78	111.90
85	AA	522	A	O4'-C1'-N9	5.75	112.80	108.20
85	AA	716	G	C3'-C2'-C1'	-5.75	96.90	101.50
85	AA	766	G	N1-C6-O6	-5.75	116.45	119.90
85	AA	1204	A	P-O5'-C5'	-5.75	111.70	120.90
85	AA	1499	G	C8-N9-C4	5.75	108.70	106.40
85	AA	1681	G	P-O5'-C5'	-5.75	111.69	120.90
85	AA	2251	U	C6-N1-C2	-5.75	117.55	121.00
86	AB	24	G	C8-N9-C4	-5.75	104.10	106.40
34	BA	295	G	C5-C6-N1	5.75	114.38	111.50
34	BA	616	G	O3'-P-O5'	5.75	114.93	104.00
34	BA	1752	A	C4-N9-C1'	-5.75	115.95	126.30
35	BB	392	G	N1-C2-N2	-5.75	111.02	116.20
35	BB	450	A	N1-C2-N3	-5.75	126.42	129.30
35	BB	621	C	C6-N1-C1'	5.75	127.70	120.80
35	BB	870	C	C2-N3-C4	-5.75	117.03	119.90
35	BB	976	U	O5'-P-OP2	-5.75	100.52	105.70
36	BC	81	U	C2'-C3'-O3'	5.75	122.90	113.70
37	BD	50	A	O4'-C1'-C2'	5.75	112.78	107.60
40	BG	73	U	C5-C6-N1	-5.75	119.82	122.70
85	AA	164	G	P-O3'-C3'	-5.75	112.80	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	186	U	C2-N1-C1'	-5.75	110.80	117.70
85	AA	444	U	O5'-P-OP2	5.75	117.60	110.70
85	AA	585	G	N7-C8-N9	-5.75	110.22	113.10
85	AA	674	U	N3-C2-O2	-5.75	118.17	122.20
85	AA	888	A	O4'-C1'-N9	5.75	112.80	108.20
85	AA	1060	U	P-O3'-C3'	5.75	126.60	119.70
34	BA	374	U	C5'-C4'-C3'	-5.75	106.80	116.00
39	BF	63	U	C3'-C2'-C1'	5.75	106.10	101.50
42	BI	113	SER	N-CA-CB	5.75	119.12	110.50
8	A7	136	TRP	CB-CG-CD2	-5.75	119.13	126.60
34	BA	268	U	N1-C1'-C2'	-5.75	105.68	112.00
34	BA	483	A	P-O3'-C3'	5.75	126.60	119.70
34	BA	528	C	O4'-C1'-N1	5.75	112.80	108.20
34	BA	610	A	O3'-P-O5'	-5.75	93.08	104.00
34	BA	1279	U	C2-N3-C4	-5.75	123.55	127.00
34	BA	1538	G	N9-C4-C5	5.75	107.70	105.40
35	BB	36	U	O5'-C5'-C4'	-5.75	100.78	111.70
35	BB	518	G	C3'-C2'-C1'	-5.75	96.90	101.50
35	BB	809	U	O4'-C1'-N1	5.75	112.80	108.20
35	BB	994	A	C4-C5-C6	-5.75	114.12	117.00
35	BB	1240	A	N1-C6-N6	-5.75	115.15	118.60
35	BB	1390	U	P-O3'-C3'	5.75	126.60	119.70
35	BB	1488	G	C5'-C4'-C3'	-5.75	106.80	116.00
41	BH	117	U	O3'-P-O5'	5.75	114.92	104.00
41	BH	134	U	C3'-C2'-C1'	-5.75	96.90	101.50
69	Bj	8	TYR	CA-CB-CG	-5.75	102.48	113.40
85	AA	267	U	O4'-C1'-C2'	-5.75	100.05	105.80
85	AA	570	U	C2-N1-C1'	-5.75	110.80	117.70
85	AA	646	C	N1-C2-O2	5.75	122.35	118.90
85	AA	1416	U	O4'-C1'-N1	5.75	112.80	108.20
85	AA	1506	U	C6-N1-C2	-5.75	117.55	121.00
85	AA	1681	G	C4-N9-C1'	-5.75	119.03	126.50
85	AA	2020	C	C6-N1-C2	-5.75	118.00	120.30
34	BA	298	G	P-O5'-C5'	-5.75	111.71	120.90
34	BA	700	G	C5-C6-N1	5.75	114.37	111.50
34	BA	701	G	O4'-C1'-C2'	5.75	112.77	107.60
34	BA	851	C	C2-N3-C4	-5.75	117.03	119.90
34	BA	1249	G	N1-C2-N2	-5.75	111.03	116.20
34	BA	1269	C	C1'-O4'-C4'	-5.75	105.30	109.90
34	BA	1444	G	O4'-C1'-N9	5.75	112.80	108.20
34	BA	1523	U	N3-C2-O2	-5.75	118.18	122.20
35	BB	22	A	N9-C1'-C2'	-5.75	105.68	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	253	G	O4'-C1'-N9	5.75	112.80	108.20
35	BB	778	A	C8-N9-C4	5.75	108.10	105.80
35	BB	1068	G	C5-C6-O6	-5.75	125.15	128.60
35	BB	1097	U	P-O3'-C3'	-5.75	112.80	119.70
35	BB	1523	U	C6-N1-C1'	5.75	129.25	121.20
36	BC	23	G	N3-C4-C5	-5.75	125.73	128.60
38	BE	106	C	C5-C4-N4	-5.75	116.18	120.20
40	BG	109	C	N3-C2-O2	-5.75	117.88	121.90
80	Bu	107	ARG	NE-CZ-NH1	5.75	123.17	120.30
85	AA	654	A	C4-C5-C6	-5.75	114.13	117.00
85	AA	693	A	C8-N9-C4	-5.75	103.50	105.80
85	AA	697	G	P-O3'-C3'	5.75	126.59	119.70
85	AA	1280	U	C5'-C4'-C3'	5.75	125.19	116.00
85	AA	1991	C	C2'-C3'-O3'	5.75	122.90	113.70
85	AA	2121	G	O3'-P-O5'	5.75	114.92	104.00
5	A4	188	PHE	CA-CB-CG	-5.75	100.11	113.90
34	BA	554	A	C5-C6-N6	-5.75	119.10	123.70
34	BA	649	A	N1-C6-N6	5.75	122.05	118.60
34	BA	1352	G	C4-N9-C1'	-5.75	119.03	126.50
41	BH	63	G	C4-C5-C6	5.75	122.25	118.80
85	AA	1354	A	P-O3'-C3'	5.75	126.59	119.70
85	AA	1449	C	C6-N1-C1'	5.75	127.69	120.80
85	AA	1753	A	O4'-C1'-N9	5.75	112.80	108.20
1	A0	111	ARG	CG-CD-NE	-5.74	99.74	111.80
27	AT	69	THR	CA-CB-CG2	-5.74	104.36	112.40
31	AX	55	ARG	NE-CZ-NH1	5.74	123.17	120.30
34	BA	289	A	N1-C2-N3	-5.74	126.43	129.30
34	BA	741	A	N9-C1'-C2'	-5.74	105.68	112.00
34	BA	1193	A	P-O5'-C5'	5.74	130.09	120.90
34	BA	1376	U	O4'-C1'-N1	5.74	112.80	108.20
34	BA	1550	G	N9-C1'-C2'	-5.74	105.68	112.00
35	BB	153	G	N1-C6-O6	5.74	123.35	119.90
35	BB	746	A	C5'-C4'-C3'	-5.74	106.81	116.00
35	BB	773	G	P-O5'-C5'	-5.74	111.71	120.90
35	BB	878	G	C8-N9-C1'	5.74	134.47	127.00
36	BC	37	U	P-O3'-C3'	-5.74	112.81	119.70
37	BD	59	G	C8-N9-C1'	5.74	134.47	127.00
38	BE	177	U	C1'-O4'-C4'	-5.74	105.31	109.90
40	BG	111	C	N1-C2-O2	5.74	122.35	118.90
50	BQ	175	HIS	N-CA-CB	5.74	120.94	110.60
73	Bn	28	HIS	CB-CA-C	-5.74	98.91	110.40
85	AA	310	U	C6-N1-C1'	5.74	129.24	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	714	U	N1-C2-O2	5.74	126.82	122.80
85	AA	962	U	C3'-C2'-C1'	-5.74	96.91	101.50
85	AA	2042	G	P-O5'-C5'	-5.74	111.71	120.90
86	AB	57	G	C8-N9-C4	5.74	108.70	106.40
34	BA	30	A	O5'-C5'-C4'	-5.74	100.79	111.70
34	BA	689	C	C2-N3-C4	-5.74	117.03	119.90
34	BA	709	C	N3-C2-O2	-5.74	117.88	121.90
34	BA	955	G	C5'-C4'-C3'	-5.74	106.81	116.00
34	BA	1249	G	N3-C4-C5	-5.74	125.73	128.60
34	BA	1482	A	O4'-C1'-C2'	5.74	112.77	107.60
34	BA	1739	G	C5-N7-C8	-5.74	101.43	104.30
34	BA	1820	G	C5'-C4'-C3'	5.74	125.19	116.00
37	BD	19	C	C6-N1-C2	-5.74	118.00	120.30
40	BG	8	U	P-O5'-C5'	5.74	130.09	120.90
73	Bn	28	HIS	CA-CB-CG	-5.74	103.84	113.60
85	AA	1372	C	C5'-C4'-C3'	-5.74	106.81	116.00
11	AC	73	TYR	CB-CA-C	5.74	121.88	110.40
34	BA	262	A	C4-C5-C6	-5.74	114.13	117.00
34	BA	373	G	N9-C1'-C2'	-5.74	105.69	112.00
34	BA	762	A	C2'-C3'-O3'	5.74	122.88	113.70
34	BA	825	G	C3'-C2'-C1'	-5.74	96.91	101.50
34	BA	1823	A	P-O5'-C5'	5.74	130.09	120.90
35	BB	80	C	N3-C4-N4	-5.74	113.98	118.00
35	BB	397	C	N3-C4-N4	-5.74	113.98	118.00
35	BB	416	U	P-O3'-C3'	-5.74	112.81	119.70
35	BB	679	G	O4'-C1'-N9	5.74	112.79	108.20
35	BB	900	C	P-O5'-C5'	5.74	130.08	120.90
37	BD	18	G	C1'-O4'-C4'	-5.74	105.31	109.90
37	BD	45	U	C6-N1-C2	-5.74	117.56	121.00
47	BN	103	ASP	CB-CG-OD1	-5.74	113.13	118.30
62	Bc	75	LEU	N-CA-C	-5.74	95.50	111.00
81	Bv	47	ARG	NE-CZ-NH1	5.74	123.17	120.30
85	AA	1047	G	C4-N9-C1'	-5.74	119.04	126.50
85	AA	1295	G	O3'-P-O5'	5.74	114.91	104.00
85	AA	1299	A	C4-N9-C1'	-5.74	115.97	126.30
85	AA	2145	G	C5-C6-O6	-5.74	125.16	128.60
85	AA	2202	G	N1-C2-N2	-5.74	111.03	116.20
26	AS	105	PHE	CB-CA-C	-5.74	98.92	110.40
34	BA	6	C	O4'-C1'-C2'	5.74	112.77	107.60
34	BA	425	G	C4'-C3'-C2'	-5.74	96.86	102.60
34	BA	570	G	C6-C5-N7	-5.74	126.96	130.40
34	BA	854	A	C5-C6-N6	-5.74	119.11	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1481	U	O4'-C1'-N1	5.74	112.79	108.20
34	BA	1714	A	N1-C6-N6	-5.74	115.16	118.60
35	BB	399	A	C5-C6-N6	-5.74	119.11	123.70
35	BB	466	A	C5-C6-N6	5.74	128.29	123.70
35	BB	643	G	C5'-C4'-C3'	-5.74	106.82	116.00
35	BB	717	A	C6-N1-C2	-5.74	115.16	118.60
35	BB	880	G	N1-C6-O6	5.74	123.34	119.90
54	BU	56	TYR	CB-CG-CD2	-5.74	117.56	121.00
85	AA	486	G	C5-C6-N1	5.74	114.37	111.50
85	AA	508	C	O3'-P-O5'	-5.74	93.10	104.00
85	AA	766	G	C5'-C4'-O4'	5.74	115.99	109.10
85	AA	813	G	C3'-C2'-C1'	-5.74	96.91	101.50
85	AA	836	A	C5'-C4'-O4'	5.74	115.99	109.10
85	AA	1084	A	C5-C6-N6	5.74	128.29	123.70
85	AA	1620	G	O4'-C1'-N9	5.74	112.79	108.20
85	AA	1703	A	O5'-C5'-C4'	-5.74	100.80	111.70
85	AA	1883	C	C6-N1-C2	-5.74	118.00	120.30
85	AA	1886	U	O4'-C1'-N1	5.74	112.79	108.20
86	AB	17	C	C5'-C4'-O4'	5.74	115.99	109.10
86	AB	38	A	C5'-C4'-C3'	-5.74	106.82	116.00
34	BA	561	U	C5-C4-O4	-5.74	122.46	125.90
34	BA	628	U	C2-N3-C4	-5.74	123.56	127.00
35	BB	431	U	C4'-C3'-C2'	5.74	108.34	102.60
38	BE	89	G	C3'-C2'-C1'	-5.74	96.91	101.50
38	BE	206	G	N1-C6-O6	5.74	123.34	119.90
85	AA	123	A	C5-C6-N6	-5.74	119.11	123.70
85	AA	743	C	C5'-C4'-O4'	-5.74	102.22	109.10
85	AA	838	G	C5-C6-N1	5.74	114.37	111.50
85	AA	1378	U	O4'-C1'-N1	5.74	112.79	108.20
34	BA	102	G	C5-C6-O6	5.74	132.04	128.60
34	BA	591	G	OP2-P-O3'	5.74	117.82	105.20
34	BA	769	U	OP1-P-O3'	5.74	117.82	105.20
34	BA	791	A	O4'-C1'-N9	5.74	112.79	108.20
34	BA	1496	G	C6-C5-N7	-5.74	126.96	130.40
34	BA	1656	A	N7-C8-N9	-5.74	110.93	113.80
35	BB	463	C	N3-C2-O2	-5.74	117.89	121.90
35	BB	910	C	O4'-C1'-N1	5.74	112.79	108.20
35	BB	1002	G	O4'-C1'-N9	5.74	112.79	108.20
35	BB	1057	G	O5'-C5'-C4'	-5.74	100.80	111.70
35	BB	1207	C	N3-C4-C5	-5.74	119.61	121.90
35	BB	1217	C	C5'-C4'-C3'	-5.74	106.82	116.00
36	BC	42	G	C8-N9-C1'	5.74	134.46	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	139	U	N3-C2-O2	-5.74	118.19	122.20
83	Bx	198	ARG	NE-CZ-NH1	5.74	123.17	120.30
85	AA	496	C	C1'-O4'-C4'	-5.74	105.31	109.90
85	AA	1762	G	C4'-C3'-C2'	-5.74	96.86	102.60
85	AA	2195	A	C8-N9-C1'	5.74	138.03	127.70
34	BA	88	C	C6-N1-C1'	5.73	127.68	120.80
34	BA	940	C	C6-N1-C1'	5.73	127.68	120.80
34	BA	1470	G	P-O3'-C3'	5.73	126.58	119.70
34	BA	1742	G	N3-C2-N2	5.73	123.91	119.90
35	BB	834	U	C5-C6-N1	5.73	125.57	122.70
38	BE	112	G	C8-N9-C1'	5.73	134.45	127.00
39	BF	56	C	C5'-C4'-C3'	5.73	125.17	116.00
40	BG	22	G	C3'-C2'-C1'	-5.73	96.91	101.50
40	BG	43	U	O3'-P-O5'	5.73	114.89	104.00
47	BN	156	SER	N-CA-CB	5.73	119.10	110.50
85	AA	120	C	C6-N1-C2	-5.73	118.01	120.30
85	AA	1156	A	C5-C6-N6	-5.73	119.11	123.70
85	AA	1696	U	C4'-C3'-C2'	-5.73	96.87	102.60
85	AA	2216	A	O3'-P-O5'	5.73	114.89	104.00
23	AP	60	SER	N-CA-CB	5.73	119.10	110.50
25	AR	53	THR	N-CA-C	-5.73	95.52	111.00
34	BA	1	C	O4'-C1'-N1	5.73	112.79	108.20
34	BA	467	A	P-O3'-C3'	-5.73	112.82	119.70
34	BA	995	A	P-O3'-C3'	-5.73	112.82	119.70
34	BA	1573	C	P-O5'-C5'	5.73	130.07	120.90
34	BA	1606	A	C1'-O4'-C4'	-5.73	105.31	109.90
35	BB	53	C	C5'-C4'-C3'	5.73	125.17	116.00
35	BB	440	U	C3'-C2'-C1'	-5.73	96.91	101.50
35	BB	483	C	P-O5'-C5'	5.73	130.07	120.90
35	BB	1282	G	C6-N1-C2	-5.73	121.66	125.10
35	BB	1430	G	P-O3'-C3'	-5.73	112.82	119.70
38	BE	53	U	P-O3'-C3'	-5.73	112.82	119.70
40	BG	128	U	C2-N1-C1'	-5.73	110.82	117.70
59	BZ	24	ARG	NE-CZ-NH2	-5.73	117.43	120.30
84	By	4	LYS	C-N-CA	5.73	136.03	121.70
85	AA	274	A	C3'-C2'-C1'	-5.73	96.91	101.50
85	AA	381	A	C4-N9-C1'	-5.73	115.98	126.30
85	AA	451	G	C4-N9-C1'	-5.73	119.05	126.50
85	AA	543	A	C8-N9-C4	5.73	108.09	105.80
85	AA	764	U	C6-N1-C2	5.73	124.44	121.00
85	AA	936	C	C6-N1-C1'	5.73	127.68	120.80
85	AA	1163	G	C3'-C2'-C1'	-5.73	96.91	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2010	C	P-O3'-C3'	-5.73	112.82	119.70
85	AA	2046	G	C5'-C4'-C3'	-5.73	106.83	116.00
86	AB	63	G	C3'-C2'-C1'	-5.73	96.91	101.50
34	BA	863	G	C5-C6-O6	5.73	132.04	128.60
34	BA	1098	G	C8-N9-C4	5.73	108.69	106.40
34	BA	1211	G	C2'-C3'-O3'	5.73	122.87	113.70
34	BA	1738	G	C4-N9-C1'	-5.73	119.05	126.50
35	BB	512	C	O5'-P-OP2	-5.73	100.54	105.70
35	BB	1124	G	P-O3'-C3'	-5.73	112.82	119.70
35	BB	1220	A	N1-C6-N6	5.73	122.04	118.60
37	BD	21	G	N3-C2-N2	5.73	123.91	119.90
39	BF	48	G	C3'-C2'-C1'	-5.73	96.92	101.50
41	BH	111	U	N3-C4-O4	-5.73	115.39	119.40
41	BH	114	G	C4'-C3'-C2'	5.73	108.33	102.60
47	BN	9	PRO	N-CA-C	5.73	127.00	112.10
53	BT	88	ARG	CG-CD-NE	-5.73	99.77	111.80
85	AA	1496	U	C2-N1-C1'	-5.73	110.82	117.70
34	BA	837	U	C5-C4-O4	-5.73	122.46	125.90
34	BA	1198	U	O4'-C1'-N1	5.73	112.78	108.20
34	BA	1212	A	C6-N1-C2	-5.73	115.16	118.60
34	BA	1488	C	O4'-C1'-C2'	5.73	112.76	107.60
34	BA	1577	U	O5'-C5'-C4'	-5.73	100.81	111.70
35	BB	1268	C	C3'-C2'-C1'	-5.73	96.92	101.50
38	BE	40	C	N1-C1'-C2'	-5.73	105.70	112.00
48	BO	131	VAL	CG1-CB-CG2	5.73	120.07	110.90
85	AA	1334	C	O4'-C1'-N1	5.73	112.78	108.20
85	AA	1799	C	C5'-C4'-O4'	5.73	115.97	109.10
31	AX	98	MET	CB-CA-C	-5.73	98.94	110.40
34	BA	168	U	C3'-C2'-C1'	-5.73	96.92	101.50
34	BA	277	A	C8-N9-C4	5.73	108.09	105.80
34	BA	531	C	C1'-O4'-C4'	-5.73	105.32	109.90
34	BA	1099	U	P-O5'-C5'	-5.73	111.73	120.90
34	BA	1330	G	C5-C6-O6	-5.73	125.16	128.60
34	BA	1409	A	C4-N9-C1'	-5.73	115.99	126.30
34	BA	1728	G	N1-C2-N2	-5.73	111.05	116.20
35	BB	499	A	C8-N9-C4	5.73	108.09	105.80
35	BB	552	C	O5'-C5'-C4'	-5.73	100.82	111.70
35	BB	650	A	P-O5'-C5'	5.73	130.06	120.90
36	BC	134	G	N3-C4-C5	-5.73	125.74	128.60
41	BH	104	U	C2-N1-C1'	-5.73	110.83	117.70
85	AA	1328	U	C2-N1-C1'	5.73	124.57	117.70
34	BA	398	G	O4'-C1'-N9	5.73	112.78	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	872	U	O5'-P-OP2	5.73	117.57	110.70
34	BA	1710	C	OP1-P-O3'	5.73	117.80	105.20
35	BB	105	U	O4'-C1'-N1	5.73	112.78	108.20
35	BB	1038	G	O3'-P-O5'	-5.73	93.12	104.00
77	Br	206	PRO	N-CA-C	5.73	126.99	112.10
85	AA	496	C	C2-N3-C4	5.73	122.76	119.90
34	BA	265	A	P-O3'-C3'	5.72	126.57	119.70
34	BA	471	U	C2'-C3'-O3'	5.72	122.86	113.70
34	BA	799	A	C5'-C4'-C3'	5.72	125.16	116.00
34	BA	941	G	C5-C6-N1	5.72	114.36	111.50
34	BA	1033	G	C3'-C2'-C1'	-5.72	96.92	101.50
34	BA	1458	A	O5'-C5'-C4'	5.72	122.58	111.70
35	BB	416	U	N1-C1'-C2'	-5.72	105.70	112.00
35	BB	811	C	C3'-C2'-C1'	-5.72	96.92	101.50
36	BC	119	G	C4-N9-C1'	-5.72	119.06	126.50
76	Bq	30	ARG	CG-CD-NE	-5.72	99.78	111.80
85	AA	286	C	C1'-O4'-C4'	-5.72	105.32	109.90
85	AA	994	A	C5'-C4'-O4'	5.72	115.97	109.10
85	AA	1255	C	C5'-C4'-C3'	-5.72	106.84	116.00
34	BA	77	C	N1-C2-O2	5.72	122.33	118.90
34	BA	252	A	P-O3'-C3'	-5.72	112.83	119.70
34	BA	617	G	C8-N9-C1'	5.72	134.44	127.00
34	BA	1146	U	O4'-C1'-N1	5.72	112.78	108.20
34	BA	1381	A	C4-N9-C1'	5.72	136.60	126.30
34	BA	1803	A	C4'-C3'-C2'	5.72	108.32	102.60
35	BB	310	U	O4'-C1'-N1	5.72	112.78	108.20
35	BB	606	C	O4'-C1'-C2'	5.72	112.75	107.60
35	BB	697	G	C5'-C4'-C3'	-5.72	106.84	116.00
41	BH	134	U	C4'-C3'-C2'	-5.72	96.88	102.60
85	AA	158	C	O4'-C1'-N1	5.72	112.78	108.20
85	AA	560	C	O3'-P-O5'	-5.72	93.13	104.00
85	AA	893	G	C6-C5-N7	-5.72	126.97	130.40
85	AA	1236	G	C4-C5-C6	-5.72	115.37	118.80
34	BA	248	G	C4'-C3'-O3'	5.72	124.44	113.00
34	BA	1605	G	N3-C2-N2	5.72	123.91	119.90
34	BA	1724	G	P-O5'-C5'	5.72	130.05	120.90
35	BB	534	C	C5'-C4'-C3'	-5.72	106.85	116.00
85	AA	57	G	P-O3'-C3'	-5.72	112.83	119.70
85	AA	190	A	C5-C6-N1	5.72	120.56	117.70
85	AA	943	U	N3-C4-O4	5.72	123.40	119.40
86	AB	53	G	O4'-C1'-N9	5.72	112.78	108.20
34	BA	366	G	C5'-C4'-C3'	-5.72	106.85	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	955	G	C4-N9-C1'	-5.72	119.06	126.50
34	BA	1436	A	C4-N9-C1'	-5.72	116.00	126.30
34	BA	1558	C	C2-N3-C4	-5.72	117.04	119.90
34	BA	1656	A	C4-N9-C1'	-5.72	116.00	126.30
35	BB	108	G	C6-N1-C2	-5.72	121.67	125.10
35	BB	676	G	C5-C6-N1	5.72	114.36	111.50
35	BB	1269	A	P-O3'-C3'	-5.72	112.84	119.70
35	BB	1469	A	P-O3'-C3'	-5.72	112.84	119.70
36	BC	16	A	N7-C8-N9	-5.72	110.94	113.80
36	BC	159	U	O4'-C1'-C2'	-5.72	100.08	105.80
37	BD	79	G	O4'-C1'-C2'	5.72	112.75	107.60
38	BE	30	C	O3'-P-O5'	-5.72	93.13	104.00
38	BE	107	U	C5'-C4'-O4'	5.72	115.96	109.10
39	BF	58	U	C2-N3-C4	-5.72	123.57	127.00
65	Bf	265	PHE	CB-CG-CD1	-5.72	116.80	120.80
72	Bm	95	ARG	NE-CZ-NH2	-5.72	117.44	120.30
85	AA	241	U	C5'-C4'-O4'	5.72	115.96	109.10
85	AA	768	C	C6-N1-C2	-5.72	118.01	120.30
85	AA	803	C	N3-C4-N4	5.72	122.00	118.00
85	AA	993	G	O4'-C1'-N9	5.72	112.78	108.20
85	AA	1019	U	O4'-C1'-N1	5.72	112.78	108.20
85	AA	1286	C	N1-C2-O2	5.72	122.33	118.90
85	AA	1972	A	P-O5'-C5'	-5.72	111.75	120.90
85	AA	2013	A	N1-C6-N6	5.72	122.03	118.60
85	AA	2069	A	C5'-C4'-C3'	-5.72	106.85	116.00
34	BA	779	U	C2-N3-C4	-5.72	123.57	127.00
34	BA	1451	A	C1'-O4'-C4'	-5.72	105.33	109.90
34	BA	1566	G	C4'-C3'-C2'	-5.72	96.88	102.60
37	BD	92	G	C4-N9-C1'	-5.72	119.07	126.50
40	BG	60	A	C5-C6-N6	-5.72	119.12	123.70
40	BG	131	U	C2-N1-C1'	-5.72	110.84	117.70
52	BS	10	CYS	N-CA-CB	5.72	120.89	110.60
85	AA	312	G	P-O5'-C5'	-5.72	111.75	120.90
85	AA	430	G	N1-C6-O6	5.72	123.33	119.90
85	AA	674	U	C1'-O4'-C4'	-5.72	105.33	109.90
85	AA	1175	A	C5'-C4'-O4'	5.72	115.96	109.10
85	AA	1353	U	O4'-C1'-N1	5.72	112.77	108.20
4	A3	32	ARG	NE-CZ-NH2	5.72	123.16	120.30
34	BA	254	U	O3'-P-O5'	-5.72	93.14	104.00
34	BA	312	U	C2-N1-C1'	-5.72	110.84	117.70
34	BA	1022	C	C3'-C2'-C1'	-5.72	96.93	101.50
34	BA	1313	U	C5'-C4'-O4'	5.72	115.96	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1435	A	C3'-C2'-C1'	-5.72	96.93	101.50
35	BB	124	G	OP1-P-OP2	-5.72	111.03	119.60
35	BB	816	U	C6-N1-C2	-5.72	117.57	121.00
35	BB	854	G	P-O3'-C3'	-5.72	112.84	119.70
35	BB	975	G	P-O3'-C3'	-5.72	112.84	119.70
35	BB	1435	G	N1-C6-O6	5.72	123.33	119.90
38	BE	11	A	C2-N3-C4	-5.72	107.74	110.60
38	BE	122	G	N9-C1'-C2'	-5.72	105.71	112.00
38	BE	201	A	N7-C8-N9	-5.72	110.94	113.80
40	BG	147	U	C6-N1-C2	-5.72	117.57	121.00
47	BN	214	GLU	CB-CG-CD	-5.72	98.77	114.20
81	Bv	79	ARG	CD-NE-CZ	-5.72	115.60	123.60
85	AA	683	U	C3'-C2'-C1'	-5.72	96.93	101.50
85	AA	790	A	N9-C1'-C2'	-5.72	105.71	112.00
85	AA	881	C	C5-C4-N4	5.72	124.20	120.20
85	AA	983	A	C8-N9-C4	-5.72	103.51	105.80
85	AA	1359	U	C6-N1-C2	-5.72	117.57	121.00
85	AA	1464	G	N3-C2-N2	5.72	123.90	119.90
85	AA	1501	A	N9-C1'-C2'	-5.72	105.71	112.00
85	AA	2239	A	P-O5'-C5'	-5.72	111.75	120.90
86	AB	73	A	C1'-O4'-C4'	-5.72	105.33	109.90
34	BA	132	U	C2-N1-C1'	-5.71	110.84	117.70
34	BA	468	A	C4'-C3'-C2'	-5.71	96.89	102.60
34	BA	547	C	N1-C1'-C2'	-5.71	105.72	112.00
34	BA	696	A	C5-C6-N1	5.71	120.56	117.70
34	BA	821	G	P-O3'-C3'	-5.71	112.84	119.70
34	BA	1054	U	N1-C2-N3	5.71	118.33	114.90
34	BA	1537	G	C8-N9-C4	-5.71	104.11	106.40
34	BA	1561	C	O5'-C5'-C4'	-5.71	100.84	111.70
35	BB	420	U	C1'-O4'-C4'	-5.71	105.33	109.90
35	BB	522	A	O4'-C4'-C3'	-5.71	98.29	104.00
35	BB	670	G	N9-C1'-C2'	-5.71	105.71	112.00
35	BB	1317	U	C2-N1-C1'	5.71	124.56	117.70
35	BB	1458	U	C5-C4-O4	5.71	129.33	125.90
39	BF	42	G	N9-C1'-C2'	-5.71	105.71	112.00
40	BG	8	U	N1-C1'-C2'	-5.71	105.71	112.00
85	AA	108	C	C6-N1-C1'	-5.71	113.94	120.80
85	AA	395	G	P-O5'-C5'	-5.71	111.76	120.90
85	AA	425	G	P-O3'-C3'	-5.71	112.84	119.70
85	AA	1035	C	C5'-C4'-O4'	5.71	115.96	109.10
85	AA	1275	A	P-O3'-C3'	5.71	126.56	119.70
85	AA	2031	C	C2'-C3'-O3'	5.71	122.84	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	331	G	C3'-C2'-C1'	-5.71	96.93	101.50
34	BA	1146	U	O5'-C5'-C4'	-5.71	100.84	111.70
34	BA	1523	U	C2-N1-C1'	-5.71	110.84	117.70
34	BA	1803	A	N9-C4-C5	-5.71	103.52	105.80
35	BB	644	A	C8-N9-C4	-5.71	103.52	105.80
41	BH	118	U	P-O5'-C5'	5.71	130.04	120.90
84	By	16	THR	CA-CB-CG2	-5.71	104.40	112.40
85	AA	22	A	P-O3'-C3'	-5.71	112.84	119.70
85	AA	815	G	N1-C2-N2	-5.71	111.06	116.20
85	AA	845	A	P-O3'-C3'	5.71	126.56	119.70
85	AA	1682	U	O4'-C1'-N1	5.71	112.77	108.20
85	AA	1702	G	C5-C6-O6	-5.71	125.17	128.60
5	A4	4	GLN	N-CA-C	-5.71	95.58	111.00
19	AK	131	GLU	N-CA-C	5.71	126.42	111.00
34	BA	27	G	P-O5'-C5'	5.71	130.04	120.90
34	BA	496	G	N9-C1'-C2'	-5.71	105.72	112.00
34	BA	918	U	O4'-C1'-N1	5.71	112.77	108.20
34	BA	996	U	C2-N1-C1'	-5.71	110.84	117.70
34	BA	1223	C	O5'-C5'-C4'	-5.71	100.85	111.70
34	BA	1321	A	O5'-C5'-C4'	-5.71	100.85	111.70
34	BA	1563	G	N3-C4-C5	-5.71	125.75	128.60
34	BA	1587	C	O3'-P-O5'	-5.71	93.15	104.00
35	BB	525	U	C2-N1-C1'	-5.71	110.85	117.70
35	BB	1160	U	P-O3'-C3'	5.71	126.55	119.70
37	BD	34	C	O4'-C1'-N1	5.71	112.77	108.20
38	BE	85	G	N9-C1'-C2'	-5.71	105.72	112.00
39	BF	12	U	C4'-C3'-C2'	-5.71	96.89	102.60
40	BG	131	U	N1-C1'-C2'	-5.71	105.72	112.00
40	BG	177	U	P-O5'-C5'	-5.71	111.76	120.90
41	BH	53	C	P-O3'-C3'	-5.71	112.85	119.70
41	BH	87	U	O5'-P-OP1	5.71	117.55	110.70
41	BH	95	C	N3-C4-C5	-5.71	119.61	121.90
42	BI	78	LEU	N-CA-CB	5.71	121.83	110.40
52	BS	175	ARG	NE-CZ-NH1	5.71	123.16	120.30
80	Bu	23	ARG	CD-NE-CZ	-5.71	115.60	123.60
82	Bw	143	ARG	NE-CZ-NH1	5.71	123.16	120.30
85	AA	349	C	N1-C1'-C2'	-5.71	105.72	112.00
85	AA	1079	C	O4'-C1'-N1	5.71	112.77	108.20
85	AA	1362	A	O5'-C5'-C4'	5.71	122.55	111.70
85	AA	1377	C	O4'-C1'-N1	5.71	112.77	108.20
85	AA	1610	G	C4-N9-C1'	-5.71	119.08	126.50
85	AA	1839	G	O4'-C1'-N9	5.71	112.77	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1987	G	C5-C6-O6	-5.71	125.17	128.60
85	AA	2022	A	P-O3'-C3'	5.71	126.55	119.70
85	AA	2168	C	P-O5'-C5'	-5.71	111.76	120.90
34	BA	502	U	C5-C6-N1	-5.71	119.84	122.70
34	BA	768	G	C8-N9-C4	5.71	108.68	106.40
35	BB	768	A	C1'-O4'-C4'	-5.71	105.33	109.90
35	BB	981	A	P-O5'-C5'	-5.71	111.77	120.90
40	BG	85	C	C6-N1-C1'	-5.71	113.95	120.80
40	BG	95	U	C1'-O4'-C4'	-5.71	105.33	109.90
82	Bw	225	PRO	N-CA-CB	-5.71	96.32	102.60
85	AA	492	C	C1'-O4'-C4'	-5.71	105.33	109.90
85	AA	1720	C	C2-N3-C4	-5.71	117.05	119.90
85	AA	2215	C	P-O3'-C3'	5.71	126.55	119.70
2	A1	51	TYR	CA-CB-CG	5.71	124.24	113.40
16	AH	61	GLU	N-CA-C	5.71	126.41	111.00
34	BA	171	U	N1-C1'-C2'	-5.71	105.72	112.00
34	BA	234	A	C4-N9-C1'	-5.71	116.02	126.30
34	BA	259	C	N3-C2-O2	-5.71	117.90	121.90
34	BA	659	U	C2'-C3'-O3'	5.71	122.83	113.70
34	BA	1054	U	C2-N1-C1'	-5.71	110.85	117.70
34	BA	1261	G	P-O5'-C5'	-5.71	111.77	120.90
34	BA	1475	G	C4-N9-C1'	-5.71	119.08	126.50
35	BB	143	G	C8-N9-C1'	5.71	134.42	127.00
35	BB	1203	C	P-O5'-C5'	-5.71	111.77	120.90
35	BB	1282	G	N1-C6-O6	-5.71	116.47	119.90
35	BB	1416	A	O4'-C1'-N9	5.71	112.77	108.20
35	BB	1467	A	C3'-C2'-C1'	-5.71	96.93	101.50
36	BC	55	U	C5-C4-O4	5.71	129.33	125.90
37	BD	105	G	C5-C6-O6	-5.71	125.17	128.60
84	By	18	SER	CB-CA-C	-5.71	99.25	110.10
85	AA	805	A	C8-N9-C4	-5.71	103.52	105.80
85	AA	881	C	C5'-C4'-C3'	-5.71	106.87	116.00
85	AA	963	U	C5'-C4'-C3'	-5.71	106.87	116.00
85	AA	1267	A	C5'-C4'-C3'	-5.71	106.86	116.00
85	AA	1717	G	N1-C6-O6	5.71	123.33	119.90
85	AA	2080	U	C2'-C3'-O3'	5.71	122.83	113.70
25	AR	71	HIS	CA-CB-CG	-5.71	103.90	113.60
31	AX	89	ARG	NE-CZ-NH1	5.71	123.15	120.30
34	BA	42	A	O5'-P-OP2	-5.71	100.56	105.70
34	BA	401	A	C5'-C4'-C3'	-5.71	106.87	116.00
34	BA	554	A	C6-N1-C2	-5.71	115.18	118.60
34	BA	881	C	N3-C4-N4	-5.71	114.00	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1187	U	C5'-C4'-O4'	5.71	115.95	109.10
34	BA	1425	G	N1-C6-O6	5.71	123.32	119.90
34	BA	1438	C	O5'-C5'-C4'	-5.71	100.86	111.70
34	BA	1451	A	C5-C6-N1	5.71	120.55	117.70
34	BA	1535	G	N1-C6-O6	5.71	123.32	119.90
34	BA	1618	A	C8-N9-C4	-5.71	103.52	105.80
35	BB	837	A	C2'-C3'-O3'	5.71	122.83	113.70
35	BB	863	U	C5'-C4'-O4'	5.71	115.95	109.10
35	BB	1072	C	C3'-C2'-C1'	-5.71	96.93	101.50
35	BB	1193	G	C8-N9-C4	5.71	108.68	106.40
36	BC	21	U	C2-N1-C1'	-5.71	110.85	117.70
41	BH	26	C	N1-C2-O2	-5.71	115.48	118.90
43	BJ	212	ARG	NE-CZ-NH2	-5.71	117.45	120.30
85	AA	154	U	C3'-C2'-C1'	-5.71	96.94	101.50
85	AA	284	C	C1'-O4'-C4'	-5.71	105.33	109.90
85	AA	681	G	C5-C6-O6	-5.71	125.18	128.60
85	AA	684	G	N3-C4-C5	-5.71	125.75	128.60
85	AA	822	U	N3-C2-O2	-5.71	118.21	122.20
85	AA	1164	A	C1'-O4'-C4'	-5.71	105.33	109.90
85	AA	1187	G	C5-C6-O6	-5.71	125.18	128.60
85	AA	1705	G	C5-C6-O6	-5.71	125.18	128.60
85	AA	2151	U	C6-N1-C1'	5.71	129.19	121.20
85	AA	2183	U	N3-C2-O2	-5.71	118.20	122.20
85	AA	2228	G	O4'-C1'-N9	5.71	112.77	108.20
34	BA	1268	C	O4'-C1'-N1	5.71	112.76	108.20
34	BA	1579	G	N1-C2-N2	-5.71	111.06	116.20
85	AA	16	G	C5-C6-O6	-5.71	125.18	128.60
85	AA	156	G	P-O3'-C3'	-5.71	112.85	119.70
85	AA	649	C	C1'-O4'-C4'	-5.71	105.34	109.90
85	AA	2070	C	O4'-C1'-N1	5.71	112.76	108.20
21	AM	1	MET	CG-SD-CE	-5.70	91.08	100.20
34	BA	539	C	C6-N1-C1'	5.70	127.64	120.80
34	BA	733	G	C8-N9-C4	5.70	108.68	106.40
34	BA	1827	C	O4'-C1'-N1	5.70	112.76	108.20
35	BB	757	C	C1'-O4'-C4'	-5.70	105.34	109.90
35	BB	765	G	N1-C6-O6	5.70	123.32	119.90
35	BB	948	G	C5-C6-O6	-5.70	125.18	128.60
35	BB	1342	C	P-O3'-C3'	-5.70	112.86	119.70
56	BW	74	LYS	N-CA-CB	5.70	120.86	110.60
85	AA	80	G	O4'-C1'-N9	5.70	112.76	108.20
85	AA	392	G	N9-C1'-C2'	-5.70	105.73	112.00
85	AA	1116	G	C1'-O4'-C4'	-5.70	105.34	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1465	C	O4'-C1'-N1	5.70	112.76	108.20
85	AA	1652	A	N1-C6-N6	-5.70	115.18	118.60
85	AA	1916	A	N3-C4-N9	-5.70	122.84	127.40
34	BA	1067	G	N9-C1'-C2'	-5.70	105.73	112.00
34	BA	1613	G	N3-C2-N2	5.70	123.89	119.90
34	BA	1617	U	N1-C2-O2	5.70	126.79	122.80
35	BB	574	G	C5'-C4'-O4'	-5.70	102.26	109.10
35	BB	635	A	C8-N9-C4	-5.70	103.52	105.80
38	BE	67	A	C3'-C2'-C1'	-5.70	96.94	101.50
39	BF	38	C	C6-N1-C1'	5.70	127.64	120.80
85	AA	122	A	C5-C6-N1	5.70	120.55	117.70
85	AA	183	C	C1'-O4'-C4'	-5.70	105.34	109.90
85	AA	449	G	P-O3'-C3'	-5.70	112.86	119.70
85	AA	1154	A	C2'-C3'-O3'	5.70	122.82	113.70
85	AA	1346	C	C5'-C4'-C3'	-5.70	106.88	116.00
1	A0	228	GLN	N-CA-CB	-5.70	100.34	110.60
34	BA	512	U	P-O5'-C5'	5.70	130.02	120.90
34	BA	784	C	N3-C2-O2	-5.70	117.91	121.90
34	BA	847	U	C1'-O4'-C4'	-5.70	105.34	109.90
34	BA	1595	G	C4-N9-C1'	-5.70	119.09	126.50
35	BB	614	U	N3-C2-O2	-5.70	118.21	122.20
35	BB	876	G	N1-C6-O6	5.70	123.32	119.90
35	BB	1142	C	N1-C1'-C2'	-5.70	105.73	112.00
36	BC	59	A	C5-C6-N6	5.70	128.26	123.70
41	BH	63	G	OP2-P-O3'	5.70	117.74	105.20
41	BH	125	U	C5'-C4'-C3'	5.70	125.12	116.00
47	BN	96	ARG	NE-CZ-NH2	-5.70	117.45	120.30
52	BS	74	ARG	NE-CZ-NH1	5.70	123.15	120.30
67	Bh	90	ILE	C-N-CA	5.70	135.95	121.70
85	AA	70	U	N3-C2-O2	-5.70	118.21	122.20
85	AA	453	G	N3-C4-C5	-5.70	125.75	128.60
85	AA	715	G	C5'-C4'-C3'	5.70	125.12	116.00
85	AA	1110	A	C5-C6-N6	5.70	128.26	123.70
85	AA	1734	A	C1'-O4'-C4'	-5.70	105.34	109.90
8	A7	243	ILE	N-CA-CB	-5.70	97.69	110.80
23	AP	90	MET	CG-SD-CE	-5.70	91.08	100.20
34	BA	86	A	O4'-C1'-N9	5.70	112.76	108.20
34	BA	557	U	N3-C4-C5	-5.70	111.18	114.60
34	BA	560	U	C5-C4-O4	-5.70	122.48	125.90
34	BA	1070	G	C4-C5-N7	5.70	113.08	110.80
34	BA	1074	C	O4'-C1'-N1	5.70	112.76	108.20
34	BA	1197	U	O5'-C5'-C4'	-5.70	100.87	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1394	U	C6-N1-C2	-5.70	117.58	121.00
35	BB	314	A	N1-C6-N6	5.70	122.02	118.60
35	BB	503	G	C5'-C4'-O4'	5.70	115.94	109.10
35	BB	567	G	C6-N1-C2	-5.70	121.68	125.10
35	BB	1068	G	P-O3'-C3'	-5.70	112.86	119.70
35	BB	1437	U	P-O5'-C5'	-5.70	111.78	120.90
37	BD	54	A	N9-C1'-C2'	-5.70	105.73	112.00
42	BI	108	CYS	N-CA-CB	-5.70	100.34	110.60
44	BK	119	TYR	CB-CG-CD1	-5.70	117.58	121.00
85	AA	105	A	P-O3'-C3'	5.70	126.54	119.70
85	AA	832	U	P-O3'-C3'	5.70	126.54	119.70
85	AA	929	G	C8-N9-C4	-5.70	104.12	106.40
85	AA	1858	G	C5'-C4'-C3'	-5.70	106.88	116.00
85	AA	1908	A	C6-N1-C2	-5.70	115.18	118.60
85	AA	2144	C	O4'-C1'-N1	5.70	112.76	108.20
85	AA	2180	C	C2-N3-C4	-5.70	117.05	119.90
34	BA	627	U	P-O5'-C5'	5.70	130.02	120.90
34	BA	1189	A	O3'-P-O5'	-5.70	93.18	104.00
34	BA	1773	U	O4'-C1'-N1	5.70	112.76	108.20
35	BB	157	G	N1-C6-O6	5.70	123.32	119.90
35	BB	575	C	C2-N3-C4	-5.70	117.05	119.90
40	BG	44	G	P-O3'-C3'	-5.70	112.86	119.70
85	AA	1068	A	N1-C6-N6	-5.70	115.18	118.60
34	BA	1	C	N1-C1'-C2'	-5.70	105.73	112.00
34	BA	167	U	O5'-P-OP2	-5.70	100.57	105.70
34	BA	403	A	C8-N9-C1'	-5.70	117.45	127.70
34	BA	976	C	P-O3'-C3'	5.70	126.53	119.70
34	BA	1093	G	P-O5'-C5'	5.70	130.01	120.90
34	BA	1135	U	C3'-C2'-C1'	-5.70	96.94	101.50
34	BA	1145	U	C6-N1-C1'	5.70	129.18	121.20
34	BA	1282	G	N3-C2-N2	-5.70	115.91	119.90
35	BB	1268	C	P-O3'-C3'	-5.70	112.87	119.70
35	BB	1291	G	N1-C2-N2	-5.70	111.07	116.20
35	BB	1500	U	P-O3'-C3'	-5.70	112.86	119.70
36	BC	155	C	C6-N1-C1'	-5.70	113.97	120.80
38	BE	40	C	C4'-C3'-O3'	-5.70	97.44	109.40
38	BE	178	G	C8-N9-C1'	5.70	134.41	127.00
41	BH	50	A	C5'-C4'-C3'	5.70	125.11	116.00
85	AA	93	G	O4'-C1'-N9	5.70	112.76	108.20
85	AA	504	U	C4'-C3'-C2'	5.70	108.30	102.60
85	AA	507	C	N3-C4-N4	5.70	121.99	118.00
85	AA	791	C	C3'-C2'-C1'	-5.70	96.94	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	902	A	N9-C4-C5	5.70	108.08	105.80
85	AA	1185	G	O5'-C5'-C4'	5.70	122.52	111.70
85	AA	1415	G	C2'-C3'-O3'	5.70	122.81	113.70
85	AA	2033	C	C5'-C4'-O4'	5.70	115.94	109.10
85	AA	2165	C	N3-C4-N4	5.70	121.99	118.00
35	BB	339	C	O4'-C1'-N1	5.69	112.75	108.20
36	BC	137	C	O4'-C4'-C3'	-5.69	98.31	104.00
38	BE	146	U	N1-C1'-C2'	-5.69	105.74	112.00
40	BG	47	G	N1-C6-O6	5.69	123.32	119.90
41	BH	98	U	N3-C2-O2	-5.69	118.21	122.20
85	AA	452	A	C5'-C4'-C3'	-5.69	106.89	116.00
85	AA	1100	U	N3-C2-O2	-5.69	118.21	122.20
34	BA	304	G	O5'-P-OP2	-5.69	100.58	105.70
34	BA	454	G	C5'-C4'-C3'	-5.69	106.89	116.00
34	BA	821	G	C6-N1-C2	-5.69	121.68	125.10
34	BA	841	G	C3'-C2'-C1'	-5.69	96.95	101.50
34	BA	892	C	C4'-C3'-C2'	5.69	108.29	102.60
34	BA	1406	U	C6-N1-C1'	5.69	129.17	121.20
35	BB	25	A	C5'-C4'-O4'	-5.69	102.27	109.10
35	BB	1084	A	N1-C6-N6	5.69	122.02	118.60
37	BD	98	G	C8-N9-C1'	5.69	134.40	127.00
38	BE	113	C	O4'-C1'-N1	5.69	112.75	108.20
38	BE	132	U	N1-C2-N3	5.69	118.32	114.90
67	Bh	86	MET	CG-SD-CE	-5.69	91.09	100.20
85	AA	1137	C	C2-N3-C4	-5.69	117.05	119.90
85	AA	1621	U	O4'-C4'-C3'	-5.69	98.31	104.00
85	AA	1725	G	O4'-C1'-N9	5.69	112.75	108.20
85	AA	1955	U	O4'-C1'-N1	5.69	112.75	108.20
85	AA	2194	U	C6-N1-C1'	5.69	129.17	121.20
34	BA	366	G	N9-C1'-C2'	-5.69	105.74	112.00
34	BA	488	C	N3-C4-N4	5.69	121.98	118.00
34	BA	543	A	P-O5'-C5'	-5.69	111.80	120.90
34	BA	651	U	O3'-P-O5'	-5.69	93.19	104.00
34	BA	741	A	C8-N9-C4	5.69	108.08	105.80
34	BA	802	G	O5'-C5'-C4'	-5.69	100.89	111.70
34	BA	843	G	C5-C6-N1	5.69	114.34	111.50
34	BA	919	A	C5-C6-N6	5.69	128.25	123.70
34	BA	951	C	C2-N3-C4	-5.69	117.06	119.90
34	BA	1025	A	C5'-C4'-C3'	-5.69	106.90	116.00
34	BA	1355	G	P-O3'-C3'	-5.69	112.87	119.70
34	BA	1560	U	C3'-C2'-C1'	5.69	106.05	101.50
35	BB	130	G	C4'-C3'-C2'	5.69	108.29	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1093	C	C1'-O4'-C4'	-5.69	105.35	109.90
35	BB	1306	G	O4'-C1'-N9	5.69	112.75	108.20
35	BB	1307	C	P-O5'-C5'	5.69	130.00	120.90
35	BB	1405	G	C4'-C3'-C2'	5.69	108.29	102.60
37	BD	16	U	N1-C2-O2	5.69	126.78	122.80
37	BD	60	C	O4'-C1'-N1	5.69	112.75	108.20
41	BH	5	G	O4'-C1'-N9	5.69	112.75	108.20
44	BK	142	ASP	N-CA-CB	-5.69	100.36	110.60
54	BU	149	ARG	NE-CZ-NH2	-5.69	117.45	120.30
82	Bw	136	LYS	CA-C-N	5.69	133.03	117.10
85	AA	9	U	C2-N1-C1'	-5.69	110.87	117.70
85	AA	1285	C	O4'-C1'-N1	5.69	112.75	108.20
85	AA	1597	C	C5'-C4'-O4'	-5.69	102.27	109.10
85	AA	1684	U	C1'-O4'-C4'	-5.69	105.35	109.90
34	BA	1283	U	C1'-O4'-C4'	-5.69	105.35	109.90
34	BA	1505	G	C1'-O4'-C4'	-5.69	105.35	109.90
35	BB	1259	A	C3'-C2'-C1'	-5.69	96.95	101.50
38	BE	108	U	N1-C2-N3	5.69	118.31	114.90
85	AA	1891	U	O4'-C1'-N1	5.69	112.75	108.20
3	A2	34	TYR	CB-CG-CD1	-5.69	117.59	121.00
3	A2	38	SER	CB-CA-C	-5.69	99.29	110.10
34	BA	540	G	N1-C6-O6	5.69	123.31	119.90
34	BA	587	U	N3-C4-O4	5.69	123.38	119.40
34	BA	610	A	C5-C6-N1	5.69	120.54	117.70
34	BA	1351	G	C5-C6-O6	-5.69	125.19	128.60
34	BA	1597	G	C5-C6-N1	5.69	114.34	111.50
34	BA	1679	C	O4'-C1'-N1	5.69	112.75	108.20
35	BB	363	A	C8-N9-C4	-5.69	103.53	105.80
35	BB	405	U	C6-N1-C1'	-5.69	113.24	121.20
35	BB	432	C	O3'-P-O5'	5.69	114.81	104.00
35	BB	1221	G	C5-C6-O6	5.69	132.01	128.60
35	BB	1453	G	C5-C6-O6	-5.69	125.19	128.60
36	BC	106	G	N1-C2-N2	-5.69	111.08	116.20
38	BE	112	G	C6-N1-C2	-5.69	121.69	125.10
40	BG	33	G	C5-N7-C8	-5.69	101.46	104.30
40	BG	51	U	O3'-P-O5'	-5.69	93.19	104.00
59	BZ	50	ASP	C-N-CA	5.69	135.92	121.70
65	Bf	62	HIS	CA-CB-CG	-5.69	103.93	113.60
85	AA	144	A	O4'-C1'-N9	5.69	112.75	108.20
85	AA	848	C	P-O5'-C5'	5.69	130.00	120.90
85	AA	895	C	P-O5'-C5'	-5.69	111.80	120.90
85	AA	1645	G	C1'-O4'-C4'	-5.69	105.35	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1757	C	C3'-C2'-C1'	-5.69	96.95	101.50
85	AA	1894	G	C5'-C4'-C3'	5.69	125.10	116.00
85	AA	1969	A	C8-N9-C1'	5.69	137.94	127.70
85	AA	2196	G	C4'-C3'-C2'	-5.69	96.91	102.60
34	BA	97	A	C5'-C4'-O4'	5.69	115.92	109.10
34	BA	1626	U	C3'-C2'-C1'	5.69	106.05	101.50
34	BA	1686	G	P-O3'-C3'	-5.69	112.88	119.70
35	BB	792	G	N1-C6-O6	5.69	123.31	119.90
35	BB	998	G	N1-C6-O6	-5.69	116.49	119.90
35	BB	1185	G	O5'-C5'-C4'	-5.69	100.90	111.70
36	BC	134	G	N9-C1'-C2'	-5.69	105.75	112.00
85	AA	570	U	C5'-C4'-C3'	-5.69	106.90	116.00
85	AA	592	C	C6-N1-C2	-5.69	118.03	120.30
85	AA	665	A	P-O3'-C3'	5.69	126.52	119.70
85	AA	1347	C	C6-N1-C2	-5.69	118.03	120.30
85	AA	1802	U	C6-N1-C2	-5.69	117.59	121.00
85	AA	1850	G	P-O5'-C5'	5.69	130.00	120.90
85	AA	1989	A	C5-C6-N6	-5.69	119.15	123.70
85	AA	2084	U	C2-N3-C4	-5.69	123.59	127.00
85	AA	2148	C	C5'-C4'-C3'	-5.69	106.90	116.00
13	AE	99	SER	N-CA-C	-5.68	95.66	111.00
34	BA	107	C	O4'-C4'-C3'	-5.68	98.32	104.00
34	BA	279	U	C4'-C3'-C2'	-5.68	96.92	102.60
34	BA	608	G	OP1-P-OP2	-5.68	111.07	119.60
34	BA	1190	A	N9-C1'-C2'	-5.68	105.75	112.00
34	BA	1441	C	N3-C2-O2	-5.68	117.92	121.90
34	BA	1475	G	C1'-O4'-C4'	-5.68	105.35	109.90
34	BA	1582	C	O5'-P-OP1	-5.68	100.58	105.70
35	BB	1014	U	C1'-O4'-C4'	-5.68	105.35	109.90
35	BB	1084	A	C5-C6-N6	-5.68	119.15	123.70
35	BB	1128	U	C3'-C2'-C1'	-5.68	96.95	101.50
36	BC	41	A	C5-C6-N6	-5.68	119.15	123.70
37	BD	47	U	C5-C6-N1	-5.68	119.86	122.70
37	BD	104	C	C6-N1-C1'	5.68	127.62	120.80
38	BE	59	U	N3-C2-O2	-5.68	118.22	122.20
40	BG	164	U	N3-C4-O4	5.68	123.38	119.40
85	AA	109	G	C6-N1-C2	-5.68	121.69	125.10
85	AA	183	C	C4'-C3'-C2'	-5.68	96.92	102.60
85	AA	639	C	N1-C1'-C2'	5.68	121.39	114.00
85	AA	674	U	C6-N1-C2	-5.68	117.59	121.00
85	AA	930	G	C4'-C3'-C2'	-5.68	96.92	102.60
85	AA	1040	U	O4'-C1'-N1	5.68	112.75	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1128	G	C2'-C3'-O3'	5.68	122.79	113.70
85	AA	1697	C	O4'-C1'-N1	5.68	112.75	108.20
85	AA	1814	U	C2-N1-C1'	-5.68	110.88	117.70
85	AA	1976	G	O5'-C5'-C4'	-5.68	100.90	111.70
85	AA	2236	U	O4'-C4'-C3'	-5.68	98.31	104.00
34	BA	146	G	C4-N9-C1'	-5.68	119.11	126.50
34	BA	262	A	C6-N1-C2	-5.68	115.19	118.60
34	BA	668	G	C5-C6-N1	5.68	114.34	111.50
34	BA	1012	A	O3'-P-O5'	-5.68	93.20	104.00
35	BB	268	G	O4'-C1'-N9	5.68	112.75	108.20
35	BB	661	G	C4-N9-C1'	-5.68	119.11	126.50
35	BB	966	C	C6-N1-C2	-5.68	118.03	120.30
35	BB	980	G	C5-C6-O6	-5.68	125.19	128.60
35	BB	1008	U	O4'-C1'-N1	5.68	112.75	108.20
35	BB	1064	U	C6-N1-C2	-5.68	117.59	121.00
35	BB	1311	G	O4'-C1'-C2'	5.68	112.71	107.60
36	BC	24	G	C5'-C4'-C3'	-5.68	106.91	116.00
36	BC	88	A	C4-N9-C1'	-5.68	116.07	126.30
37	BD	106	G	C1'-O4'-C4'	-5.68	105.35	109.90
39	BF	59	U	C2-N3-C4	-5.68	123.59	127.00
41	BH	113	G	N1-C2-N3	-5.68	120.49	123.90
56	BW	127	LEU	N-CA-CB	-5.68	99.04	110.40
67	Bh	99	ARG	NE-CZ-NH1	5.68	123.14	120.30
74	Bo	44	LYS	N-CA-C	-5.68	95.66	111.00
81	Bv	27	ARG	NE-CZ-NH1	5.68	123.14	120.30
85	AA	107	A	N1-C6-N6	5.68	122.01	118.60
85	AA	310	U	N1-C2-N3	5.68	118.31	114.90
85	AA	414	C	O4'-C1'-N1	5.68	112.75	108.20
85	AA	680	U	C5-C4-O4	5.68	129.31	125.90
85	AA	763	U	C3'-C2'-C1'	-5.68	96.95	101.50
86	AB	43	C	P-O5'-C5'	5.68	129.99	120.90
34	BA	193	C	P-O3'-C3'	5.68	126.52	119.70
34	BA	1079	C	N1-C1'-C2'	-5.68	105.75	112.00
34	BA	1645	C	O4'-C1'-N1	5.68	112.75	108.20
35	BB	630	A	C1'-O4'-C4'	-5.68	105.36	109.90
35	BB	704	G	N1-C2-N2	-5.68	111.09	116.20
36	BC	128	U	C4'-C3'-C2'	-5.68	96.92	102.60
49	BP	102	PHE	CB-CG-CD1	5.68	124.78	120.80
85	AA	307	G	C8-N9-C1'	5.68	134.38	127.00
85	AA	790	A	C5'-C4'-C3'	5.68	125.09	116.00
85	AA	970	U	C6-N1-C1'	-5.68	113.25	121.20
85	AA	2248	A	P-O5'-C5'	5.68	129.99	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A0	114	TYR	CB-CG-CD1	5.68	124.41	121.00
34	BA	362	G	C4-N9-C1'	-5.68	119.12	126.50
34	BA	543	A	C1'-O4'-C4'	-5.68	105.36	109.90
34	BA	958	G	C8-N9-C4	-5.68	104.13	106.40
34	BA	1208	U	C6-N1-C2	-5.68	117.59	121.00
34	BA	1258	G	C6-C5-N7	-5.68	126.99	130.40
35	BB	66	G	N3-C2-N2	5.68	123.88	119.90
35	BB	96	A	C4-N9-C1'	5.68	136.52	126.30
35	BB	1368	A	N7-C8-N9	-5.68	110.96	113.80
35	BB	1378	U	P-O5'-C5'	-5.68	111.81	120.90
36	BC	138	C	C6-N1-C1'	5.68	127.62	120.80
38	BE	30	C	N3-C2-O2	-5.68	117.92	121.90
38	BE	129	G	C5-C6-N1	5.68	114.34	111.50
39	BF	47	C	C1'-O4'-C4'	-5.68	105.36	109.90
39	BF	64	U	C2-N1-C1'	5.68	124.52	117.70
41	BH	64	U	C4'-C3'-C2'	-5.68	96.92	102.60
85	AA	683	U	O4'-C1'-N1	5.68	112.74	108.20
85	AA	1235	G	P-O5'-C5'	5.68	129.99	120.90
7	A6	25	ARG	NE-CZ-NH1	5.68	123.14	120.30
34	BA	582	U	C5-C4-O4	5.68	129.31	125.90
34	BA	1435	A	N9-C4-C5	-5.68	103.53	105.80
34	BA	1618	A	O3'-P-O5'	-5.68	93.21	104.00
35	BB	127	U	O5'-P-OP2	-5.68	100.59	105.70
40	BG	73	U	O4'-C1'-C2'	5.68	112.71	107.60
40	BG	171	A	C4'-C3'-C2'	-5.68	96.92	102.60
48	BO	216	LEU	CB-CG-CD1	5.68	120.65	111.00
54	BU	83	ARG	NE-CZ-NH1	5.68	123.14	120.30
85	AA	132	G	C8-N9-C1'	5.68	134.38	127.00
85	AA	838	G	N1-C6-O6	5.68	123.31	119.90
85	AA	1093	C	P-O5'-C5'	5.68	129.99	120.90
26	AS	97	ASN	CA-CB-CG	-5.68	100.91	113.40
34	BA	115	U	O4'-C1'-N1	5.68	112.74	108.20
34	BA	530	A	N3-C4-N9	-5.68	122.86	127.40
34	BA	1208	U	N3-C2-O2	-5.68	118.23	122.20
35	BB	127	U	C2-N1-C1'	-5.68	110.89	117.70
35	BB	354	C	O4'-C1'-N1	5.68	112.74	108.20
35	BB	509	A	P-O3'-C3'	-5.68	112.89	119.70
35	BB	1191	G	C8-N9-C4	5.68	108.67	106.40
40	BG	82	U	C2-N1-C1'	-5.68	110.89	117.70
41	BH	48	G	N1-C6-O6	-5.68	116.49	119.90
41	BH	135	U	N3-C2-O2	-5.68	118.23	122.20
85	AA	264	A	C5'-C4'-C3'	5.68	125.08	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	455	G	C1'-O4'-C4'	-5.68	105.36	109.90
85	AA	1413	G	N1-C6-O6	5.68	123.31	119.90
85	AA	2047	U	P-O5'-C5'	-5.68	111.82	120.90
34	BA	140	C	O5'-C5'-C4'	5.67	122.48	111.70
34	BA	213	A	C4-C5-C6	-5.67	114.16	117.00
34	BA	494	A	C4'-C3'-C2'	5.67	108.28	102.60
34	BA	660	C	O4'-C1'-N1	5.67	112.74	108.20
34	BA	1261	G	C3'-C2'-C1'	-5.67	96.96	101.50
34	BA	1758	C	P-O3'-C3'	5.67	126.51	119.70
35	BB	369	A	N1-C6-N6	-5.67	115.19	118.60
35	BB	1128	U	P-O5'-C5'	-5.67	111.82	120.90
35	BB	1157	G	C4-N9-C1'	-5.67	119.12	126.50
35	BB	1218	G	C3'-C2'-C1'	-5.67	96.96	101.50
35	BB	1444	U	N3-C2-O2	-5.67	118.23	122.20
35	BB	1546	C	N3-C2-O2	-5.67	117.93	121.90
36	BC	70	C	C3'-C2'-C1'	-5.67	96.96	101.50
37	BD	109	U	C2-N3-C4	-5.67	123.59	127.00
38	BE	75	C	C3'-C2'-C1'	-5.67	96.96	101.50
38	BE	201	A	C5'-C4'-C3'	-5.67	106.92	116.00
59	BZ	24	ARG	CG-CD-NE	-5.67	99.89	111.80
85	AA	7	G	O3'-P-O5'	5.67	114.78	104.00
85	AA	542	G	C6-N1-C2	-5.67	121.69	125.10
85	AA	926	C	C2-N1-C1'	-5.67	112.56	118.80
85	AA	1104	G	O5'-C5'-C4'	-5.67	100.92	111.70
85	AA	2100	A	O4'-C1'-N9	5.67	112.74	108.20
34	BA	508	C	P-O5'-C5'	5.67	129.98	120.90
34	BA	580	U	P-O3'-C3'	5.67	126.51	119.70
35	BB	522	A	C6-N1-C2	-5.67	115.20	118.60
35	BB	1050	A	N9-C1'-C2'	-5.67	105.76	112.00
41	BH	16	A	N7-C8-N9	5.67	116.64	113.80
41	BH	123	G	C5'-C4'-C3'	5.67	125.08	116.00
85	AA	547	A	P-O3'-C3'	5.67	126.51	119.70
85	AA	1527	G	N1-C6-O6	5.67	123.30	119.90
34	BA	13	U	C5-C6-N1	-5.67	119.86	122.70
34	BA	168	U	C6-N1-C2	-5.67	117.60	121.00
34	BA	354	G	O4'-C1'-N9	5.67	112.74	108.20
34	BA	430	A	O5'-P-OP2	-5.67	100.60	105.70
34	BA	1074	C	N3-C2-O2	-5.67	117.93	121.90
34	BA	1444	G	O5'-P-OP1	-5.67	100.60	105.70
34	BA	1515	U	C1'-O4'-C4'	-5.67	105.36	109.90
34	BA	1844	U	C6-N1-C1'	5.67	129.14	121.20
35	BB	886	G	N1-C6-O6	5.67	123.30	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1214	U	C6-N1-C1'	5.67	129.14	121.20
35	BB	1254	G	C5-C6-O6	5.67	132.00	128.60
37	BD	73	U	C6-N1-C2	-5.67	117.60	121.00
38	BE	23	G	N3-C4-C5	-5.67	125.76	128.60
38	BE	163	A	C5-N7-C8	-5.67	101.06	103.90
39	BF	65	U	C4-C5-C6	-5.67	116.30	119.70
49	BP	82	LYS	CA-CB-CG	-5.67	100.92	113.40
50	BQ	127	TYR	CB-CG-CD2	-5.67	117.60	121.00
60	Ba	113	ARG	NE-CZ-NH1	5.67	123.14	120.30
62	Bc	13	ARG	CD-NE-CZ	5.67	131.54	123.60
72	Bm	46	ARG	NE-CZ-NH2	5.67	123.14	120.30
84	By	56	VAL	C-N-CA	5.67	135.88	121.70
85	AA	169	G	C5'-C4'-C3'	-5.67	106.93	116.00
85	AA	251	A	C1'-O4'-C4'	-5.67	105.36	109.90
34	BA	490	A	O4'-C1'-N9	5.67	112.74	108.20
34	BA	675	C	C6-N1-C2	-5.67	118.03	120.30
34	BA	1147	C	P-O5'-C5'	-5.67	111.83	120.90
34	BA	1735	G	C5'-C4'-O4'	-5.67	102.30	109.10
35	BB	688	U	N3-C2-O2	-5.67	118.23	122.20
35	BB	696	G	C5-C6-O6	-5.67	125.20	128.60
35	BB	868	C	N1-C2-N3	5.67	123.17	119.20
35	BB	1465	U	O4'-C1'-N1	5.67	112.74	108.20
38	BE	124	G	C8-N9-C4	5.67	108.67	106.40
85	AA	843	U	C5-C4-O4	-5.67	122.50	125.90
85	AA	887	A	C5-C6-N6	-5.67	119.16	123.70
85	AA	1011	G	C5-C6-O6	-5.67	125.20	128.60
85	AA	1180	C	C2'-C3'-O3'	5.67	122.77	113.70
85	AA	1469	G	N3-C4-C5	-5.67	125.77	128.60
6	A5	77	ARG	NE-CZ-NH2	-5.67	117.47	120.30
11	AC	160	VAL	CA-CB-CG1	5.67	119.40	110.90
34	BA	239	C	C5-C4-N4	-5.67	116.23	120.20
34	BA	481	A	C4'-C3'-C2'	-5.67	96.93	102.60
34	BA	1563	G	O5'-C5'-C4'	-5.67	100.93	111.70
35	BB	826	G	N3-C2-N2	5.67	123.87	119.90
35	BB	1134	G	C5'-C4'-C3'	-5.67	106.93	116.00
35	BB	1325	C	P-O5'-C5'	5.67	129.97	120.90
35	BB	1521	G	C8-N9-C1'	5.67	134.37	127.00
44	BK	170	TYR	CB-CG-CD2	-5.67	117.60	121.00
45	BL	170	PHE	CB-CG-CD1	5.67	124.77	120.80
47	BN	40	ARG	NE-CZ-NH2	-5.67	117.47	120.30
57	BX	60	TYR	CA-CB-CG	-5.67	102.63	113.40
71	Bl	47	TYR	CA-CB-CG	-5.67	102.63	113.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
72	Bm	90	ALA	CB-CA-C	-5.67	101.60	110.10
85	AA	178	U	C4'-C3'-C2'	-5.67	96.93	102.60
85	AA	188	G	O5'-P-OP1	-5.67	100.60	105.70
85	AA	663	C	O4'-C1'-N1	5.67	112.73	108.20
85	AA	936	C	C2-N1-C1'	-5.67	112.56	118.80
85	AA	1552	U	O4'-C1'-N1	5.67	112.73	108.20
85	AA	1921	G	C6-C5-N7	5.67	133.80	130.40
34	BA	431	A	P-O3'-C3'	-5.67	112.90	119.70
34	BA	585	G	O4'-C1'-N9	5.67	112.73	108.20
34	BA	745	A	C8-N9-C4	-5.67	103.53	105.80
34	BA	1293	A	C2'-C3'-O3'	5.67	122.77	113.70
35	BB	457	U	C4'-C3'-C2'	-5.67	96.93	102.60
35	BB	677	U	C2'-C3'-O3'	5.67	122.77	113.70
35	BB	814	A	C5'-C4'-C3'	-5.67	106.93	116.00
35	BB	1170	U	C5-C6-N1	-5.67	119.87	122.70
36	BC	108	A	P-O3'-C3'	-5.67	112.90	119.70
41	BH	118	U	C1'-O4'-C4'	-5.67	105.37	109.90
44	BK	144	TYR	CA-CB-CG	-5.67	102.63	113.40
65	Bf	217	THR	CA-CB-CG2	-5.67	104.47	112.40
80	Bu	179	VAL	CB-CA-C	-5.67	100.63	111.40
85	AA	889	G	C4'-C3'-C2'	-5.67	96.93	102.60
85	AA	943	U	P-O3'-C3'	-5.67	112.90	119.70
85	AA	1263	G	N3-C2-N2	5.67	123.87	119.90
85	AA	1312	G	O4'-C1'-N9	5.67	112.73	108.20
85	AA	1863	A	C5-C6-N6	5.67	128.23	123.70
85	AA	2070	C	N3-C2-O2	-5.67	117.93	121.90
34	BA	88	C	N3-C2-O2	-5.67	117.93	121.90
34	BA	353	U	N1-C1'-C2'	-5.67	105.77	112.00
34	BA	586	G	C3'-C2'-C1'	-5.67	96.97	101.50
34	BA	755	G	C4-N9-C1'	-5.67	119.14	126.50
34	BA	1692	U	N1-C2-N3	5.67	118.30	114.90
34	BA	1700	C	C2-N3-C4	-5.67	117.07	119.90
34	BA	1808	A	C3'-C2'-C1'	-5.67	96.97	101.50
35	BB	63	A	N1-C6-N6	5.67	122.00	118.60
35	BB	717	A	C5'-C4'-O4'	5.67	115.90	109.10
35	BB	1313	C	O5'-C5'-C4'	5.67	122.46	111.70
40	BG	1	G	C6-N1-C2	-5.67	121.70	125.10
41	BH	89	C	O5'-P-OP1	5.67	117.50	110.70
47	BN	42	ARG	NE-CZ-NH1	5.67	123.13	120.30
85	AA	835	C	C1'-O4'-C4'	-5.67	105.37	109.90
85	AA	919	U	O4'-C1'-N1	5.67	112.73	108.20
85	AA	945	A	C4'-C3'-C2'	5.67	108.27	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	963	U	C5-C6-N1	-5.67	119.87	122.70
85	AA	1937	G	C4-N9-C1'	-5.67	119.14	126.50
34	BA	470	C	P-O5'-C5'	-5.66	111.84	120.90
34	BA	555	C	O5'-P-OP1	-5.66	100.60	105.70
34	BA	557	U	C1'-C2'-O2'	-5.66	93.61	110.60
34	BA	613	A	P-O3'-C3'	-5.66	112.90	119.70
34	BA	1213	A	O5'-C5'-C4'	-5.66	100.94	111.70
35	BB	814	A	C2-N3-C4	5.66	113.43	110.60
37	BD	27	A	O5'-P-OP2	-5.66	100.60	105.70
38	BE	128	G	OP1-P-O3'	5.66	117.66	105.20
40	BG	167	C	C5'-C4'-C3'	-5.66	106.94	116.00
85	AA	332	A	C4'-C3'-C2'	-5.66	96.94	102.60
85	AA	1093	C	C6-N1-C1'	5.66	127.60	120.80
85	AA	1753	A	N1-C6-N6	-5.66	115.20	118.60
85	AA	1870	C	C6-N1-C2	-5.66	118.03	120.30
85	AA	2081	A	C4'-C3'-C2'	-5.66	96.94	102.60
85	AA	2105	G	C5-C6-N1	5.66	114.33	111.50
34	BA	244	A	N9-C1'-C2'	-5.66	105.77	112.00
34	BA	433	G	C1'-O4'-C4'	5.66	114.43	109.90
34	BA	1406	U	C2-N1-C1'	-5.66	110.91	117.70
34	BA	1475	G	N7-C8-N9	-5.66	110.27	113.10
38	BE	2	G	C6-N1-C2	-5.66	121.70	125.10
41	BH	83	U	O4'-C1'-N1	5.66	112.73	108.20
85	AA	121	C	P-O5'-C5'	-5.66	111.84	120.90
85	AA	472	A	N9-C4-C5	5.66	108.06	105.80
34	BA	98	A	C6-N1-C2	-5.66	115.20	118.60
34	BA	340	U	C5-C6-N1	-5.66	119.87	122.70
34	BA	619	U	C6-N1-C2	-5.66	117.60	121.00
34	BA	912	G	C6-N1-C2	-5.66	121.70	125.10
34	BA	1486	U	N1-C1'-C2'	-5.66	105.77	112.00
35	BB	477	U	P-O3'-C3'	5.66	126.49	119.70
35	BB	1303	A	O3'-P-O5'	-5.66	93.24	104.00
35	BB	1449	G	N1-C6-O6	5.66	123.30	119.90
36	BC	117	A	C5'-C4'-O4'	5.66	115.89	109.10
38	BE	74	U	N1-C1'-C2'	-5.66	105.77	112.00
38	BE	95	G	C2-N3-C4	5.66	114.73	111.90
38	BE	126	G	C5-C6-N1	5.66	114.33	111.50
40	BG	23	C	O3'-P-O5'	-5.66	93.25	104.00
40	BG	38	A	C5-C6-N6	-5.66	119.17	123.70
40	BG	129	G	C1'-O4'-C4'	-5.66	105.37	109.90
41	BH	100	A	N3-C4-C5	-5.66	122.84	126.80
47	BN	101	ARG	NE-CZ-NH1	5.66	123.13	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	187	C	C5-C4-N4	-5.66	116.24	120.20
85	AA	604	C	C6-N1-C2	-5.66	118.04	120.30
85	AA	1122	U	C2'-C3'-O3'	5.66	122.76	113.70
85	AA	1383	C	C2-N1-C1'	5.66	125.03	118.80
85	AA	1644	G	N3-C2-N2	5.66	123.86	119.90
85	AA	1883	C	C2-N1-C1'	5.66	125.03	118.80
85	AA	2172	A	P-O5'-C5'	5.66	129.96	120.90
5	A4	107	ARG	NE-CZ-NH1	5.66	123.13	120.30
34	BA	86	A	C4-C5-C6	-5.66	114.17	117.00
34	BA	174	A	C5'-C4'-O4'	5.66	115.89	109.10
34	BA	318	U	O3'-P-O5'	-5.66	93.25	104.00
34	BA	518	C	C2'-C3'-O3'	5.66	122.75	113.70
34	BA	687	G	C5'-C4'-O4'	-5.66	102.31	109.10
34	BA	936	A	C3'-C2'-C1'	-5.66	96.97	101.50
34	BA	1222	C	C2-N1-C1'	5.66	125.03	118.80
34	BA	1685	C	O5'-C5'-C4'	-5.66	100.95	111.70
35	BB	648	G	C5'-C4'-C3'	-5.66	106.94	116.00
35	BB	1337	C	C5'-C4'-C3'	-5.66	106.95	116.00
35	BB	1405	G	N9-C1'-C2'	-5.66	105.78	112.00
35	BB	1487	G	N1-C2-N2	-5.66	111.11	116.20
35	BB	1520	C	N3-C4-N4	-5.66	114.04	118.00
36	BC	26	U	P-O3'-C3'	5.66	126.49	119.70
40	BG	23	C	OP1-P-O3'	5.66	117.65	105.20
40	BG	102	G	C1'-O4'-C4'	-5.66	105.37	109.90
41	BH	23	G	C6-N1-C2	-5.66	121.70	125.10
41	BH	47	G	C3'-C2'-C1'	-5.66	96.97	101.50
59	BZ	9	ARG	NE-CZ-NH1	5.66	123.13	120.30
85	AA	660	G	O4'-C1'-N9	5.66	112.73	108.20
85	AA	1057	G	C4'-C3'-C2'	-5.66	96.94	102.60
85	AA	1120	G	C5-C6-N1	5.66	114.33	111.50
85	AA	1674	G	C5-N7-C8	-5.66	101.47	104.30
34	BA	228	A	N3-C4-C5	-5.66	122.84	126.80
34	BA	1043	C	C4'-C3'-C2'	5.66	108.26	102.60
35	BB	1230	A	C4'-C3'-C2'	-5.66	96.94	102.60
36	BC	155	C	O3'-P-O5'	-5.66	93.25	104.00
37	BD	9	C	C5-C6-N1	-5.66	118.17	121.00
38	BE	206	G	C8-N9-C1'	5.66	134.35	127.00
57	BX	62	ARG	N-CA-C	-5.66	95.73	111.00
71	Bl	126	PHE	CA-CB-CG	-5.66	100.32	113.90
83	Bx	244	ARG	NE-CZ-NH1	5.66	123.13	120.30
85	AA	22	A	N1-C6-N6	-5.66	115.21	118.60
85	AA	290	G	N9-C1'-C2'	-5.66	105.78	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	999	A	P-O3'-C3'	5.66	126.49	119.70
34	BA	4	A	O4'-C1'-N9	-5.66	103.67	108.20
34	BA	1413	G	C1'-O4'-C4'	-5.66	105.38	109.90
34	BA	1414	C	C2-N3-C4	-5.66	117.07	119.90
34	BA	1415	C	P-O3'-C3'	-5.66	112.91	119.70
34	BA	1500	G	C8-N9-C1'	5.66	134.35	127.00
34	BA	1651	C	O3'-P-O5'	-5.66	93.25	104.00
35	BB	50	A	N9-C4-C5	-5.66	103.54	105.80
53	BT	81	ARG	NE-CZ-NH1	-5.66	117.47	120.30
56	BW	120	VAL	O-C-N	-5.66	113.65	122.70
77	Br	215	GLY	C-N-CA	5.66	135.84	121.70
80	Bu	121	GLY	N-CA-C	5.66	127.24	113.10
85	AA	146	U	C2-N3-C4	5.66	130.39	127.00
85	AA	260	A	C5'-C4'-C3'	5.66	125.05	116.00
85	AA	544	A	N7-C8-N9	-5.66	110.97	113.80
85	AA	861	G	C5-C6-O6	-5.66	125.21	128.60
85	AA	995	G	N7-C8-N9	-5.66	110.27	113.10
85	AA	1128	G	C6-N1-C2	-5.66	121.71	125.10
85	AA	1571	A	O4'-C1'-N9	5.66	112.72	108.20
34	BA	91	C	C5-C4-N4	5.65	124.16	120.20
34	BA	744	G	C6-C5-N7	-5.65	127.01	130.40
34	BA	821	G	C4'-C3'-C2'	-5.65	96.95	102.60
34	BA	1037	C	O4'-C1'-N1	5.65	112.72	108.20
34	BA	1613	G	N9-C4-C5	-5.65	103.14	105.40
35	BB	4	C	OP2-P-O3'	5.65	117.64	105.20
35	BB	49	A	N1-C6-N6	5.65	121.99	118.60
35	BB	431	U	C5'-C4'-C3'	-5.65	106.95	116.00
35	BB	1513	U	N3-C2-O2	-5.65	118.24	122.20
44	BK	69	ARG	NE-CZ-NH2	5.65	123.13	120.30
85	AA	1799	C	C6-N1-C2	-5.65	118.04	120.30
85	AA	1921	G	C2-N3-C4	5.65	114.73	111.90
2	A1	70	ASP	CB-CG-OD1	5.65	123.39	118.30
3	A2	122	ARG	NE-CZ-NH1	5.65	123.13	120.30
34	BA	21	C	N1-C1'-C2'	-5.65	105.78	112.00
34	BA	165	C	O3'-P-O5'	-5.65	93.26	104.00
34	BA	230	A	C3'-C2'-C1'	5.65	106.02	101.50
34	BA	517	A	C5-C6-N1	5.65	120.53	117.70
34	BA	726	G	N3-C2-N2	5.65	123.86	119.90
34	BA	745	A	C5-C6-N1	5.65	120.53	117.70
34	BA	897	U	C2-N3-C4	-5.65	123.61	127.00
34	BA	1110	A	C8-N9-C4	5.65	108.06	105.80
34	BA	1613	G	C8-N9-C1'	5.65	134.35	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	491	A	C3'-C2'-C1'	-5.65	96.98	101.50
35	BB	596	C	N3-C2-O2	-5.65	117.94	121.90
35	BB	1024	G	C2'-C3'-O3'	5.65	122.75	113.70
36	BC	156	A	C8-N9-C4	5.65	108.06	105.80
70	Bk	123	TYR	CB-CG-CD1	-5.65	117.61	121.00
85	AA	354	C	C4'-C3'-C2'	-5.65	96.95	102.60
85	AA	404	A	C4'-C3'-C2'	5.65	108.25	102.60
85	AA	525	C	O4'-C1'-N1	5.65	112.72	108.20
85	AA	1221	G	C4-N9-C1'	-5.65	119.15	126.50
85	AA	1756	C	C5-C6-N1	5.65	123.83	121.00
85	AA	1812	C	N3-C2-O2	-5.65	117.94	121.90
85	AA	1825	A	C4-N9-C1'	5.65	136.47	126.30
85	AA	2151	U	O3'-P-O5'	-5.65	93.26	104.00
1	A0	114	TYR	CB-CG-CD2	-5.65	117.61	121.00
2	A1	71	GLY	N-CA-C	-5.65	98.97	113.10
34	BA	825	G	C5-C6-O6	-5.65	125.21	128.60
34	BA	831	U	C5-C6-N1	5.65	125.53	122.70
34	BA	1359	U	C1'-O4'-C4'	-5.65	105.38	109.90
35	BB	75	A	C1'-O4'-C4'	-5.65	105.38	109.90
35	BB	418	G	O4'-C1'-C2'	5.65	112.69	107.60
35	BB	1107	C	N3-C2-O2	-5.65	117.94	121.90
35	BB	1250	A	C4'-C3'-C2'	5.65	108.25	102.60
35	BB	1351	G	C4'-C3'-C2'	5.65	108.25	102.60
35	BB	1361	A	C5'-C4'-O4'	5.65	115.88	109.10
39	BF	62	U	C5'-C4'-C3'	-5.65	106.96	116.00
41	BH	67	G	C1'-O4'-C4'	-5.65	105.38	109.90
45	BL	67	ARG	NE-CZ-NH1	5.65	123.12	120.30
85	AA	1096	G	C3'-C2'-C1'	-5.65	96.98	101.50
85	AA	1253	G	C1'-O4'-C4'	-5.65	105.38	109.90
85	AA	1662	U	O4'-C1'-N1	5.65	112.72	108.20
85	AA	2063	C	C2'-C3'-O3'	5.65	122.74	113.70
86	AB	15	G	P-O5'-C5'	5.65	129.94	120.90
18	AJ	20	ARG	NE-CZ-NH1	5.65	123.12	120.30
34	BA	183	G	C8-N9-C1'	5.65	134.34	127.00
34	BA	489	A	C5'-C4'-O4'	5.65	115.88	109.10
35	BB	763	U	C6-N1-C1'	5.65	129.11	121.20
35	BB	1195	A	C8-N9-C1'	5.65	137.87	127.70
36	BC	42	G	O5'-C5'-C4'	-5.65	100.97	111.70
37	BD	119	U	C6-N1-C1'	-5.65	113.29	121.20
70	Bk	88	ARG	N-CA-CB	-5.65	100.43	110.60
81	Bv	174	TYR	CB-CG-CD2	-5.65	117.61	121.00
85	AA	495	G	C5'-C4'-C3'	-5.65	106.96	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1228	A	C3'-C2'-C1'	-5.65	96.98	101.50
85	AA	1371	C	O4'-C1'-N1	5.65	112.72	108.20
85	AA	1833	C	C3'-C2'-C1'	-5.65	96.98	101.50
23	AP	55	ASP	CB-CG-OD2	-5.65	113.22	118.30
34	BA	932	G	P-O3'-C3'	-5.65	112.92	119.70
34	BA	1505	G	N3-C4-N9	-5.65	122.61	126.00
35	BB	790	A	P-O3'-C3'	5.65	126.48	119.70
35	BB	1162	A	C5'-C4'-C3'	-5.65	106.97	116.00
35	BB	1274	G	C6-N1-C2	-5.65	121.71	125.10
38	BE	124	G	C5'-C4'-O4'	5.65	115.88	109.10
38	BE	141	A	C5'-C4'-C3'	-5.65	106.96	116.00
40	BG	31	G	N9-C4-C5	-5.65	103.14	105.40
41	BH	5	G	C8-N9-C1'	5.65	134.34	127.00
85	AA	628	C	OP1-P-O3'	5.65	117.63	105.20
34	BA	331	G	P-O3'-C3'	-5.65	112.92	119.70
34	BA	516	U	N3-C2-O2	-5.65	118.25	122.20
34	BA	1653	G	N7-C8-N9	-5.65	110.28	113.10
38	BE	104	G	OP1-P-OP2	-5.65	111.13	119.60
85	AA	194	U	C5'-C4'-C3'	-5.65	106.97	116.00
85	AA	1117	G	O3'-P-O5'	-5.65	93.27	104.00
85	AA	1959	G	O4'-C1'-N9	5.65	112.72	108.20
86	AB	5	G	P-O3'-C3'	-5.65	112.92	119.70
34	BA	495	A	C8-N9-C1'	5.64	137.86	127.70
34	BA	931	G	C4-N9-C1'	-5.64	119.16	126.50
34	BA	1374	G	O4'-C1'-N9	5.64	112.72	108.20
34	BA	1684	A	N1-C6-N6	-5.64	115.21	118.60
35	BB	487	A	O4'-C1'-C2'	5.64	112.68	107.60
35	BB	534	C	C5'-C4'-O4'	5.64	115.87	109.10
35	BB	751	A	N1-C2-N3	-5.64	126.48	129.30
35	BB	1470	G	N9-C1'-C2'	-5.64	105.79	112.00
38	BE	31	A	P-O3'-C3'	-5.64	112.93	119.70
38	BE	34	C	C2-N3-C4	-5.64	117.08	119.90
53	BT	5	LYS	CA-C-N	5.64	129.62	117.20
66	Bg	97	ARG	NE-CZ-NH1	5.64	123.12	120.30
75	Bp	48	ALA	C-N-CA	5.64	135.81	121.70
84	By	87	ARG	NE-CZ-NH1	5.64	123.12	120.30
85	AA	49	C	O5'-P-OP2	-5.64	100.62	105.70
85	AA	301	U	C6-N1-C1'	-5.64	113.30	121.20
85	AA	489	C	O4'-C4'-C3'	-5.64	98.36	104.00
85	AA	615	A	O4'-C1'-N9	5.64	112.72	108.20
85	AA	1142	G	C5-C6-O6	-5.64	125.21	128.60
85	AA	1283	C	C2-N1-C1'	-5.64	112.59	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1538	C	N3-C2-O2	-5.64	117.95	121.90
85	AA	2007	G	C4'-C3'-C2'	5.64	108.25	102.60
15	AG	18	TYR	CB-CG-CD1	-5.64	117.61	121.00
23	AP	101	SER	N-CA-CB	-5.64	102.03	110.50
23	AP	113	VAL	CA-CB-CG2	-5.64	102.44	110.90
34	BA	175	G	C3'-C2'-C1'	-5.64	96.99	101.50
34	BA	826	C	C3'-C2'-C1'	-5.64	96.98	101.50
35	BB	366	G	C5-C6-O6	-5.64	125.21	128.60
35	BB	824	C	C2-N3-C4	-5.64	117.08	119.90
35	BB	1001	G	O4'-C1'-N9	5.64	112.71	108.20
35	BB	1163	U	O4'-C1'-N1	5.64	112.71	108.20
35	BB	1234	G	C5-C6-O6	-5.64	125.22	128.60
38	BE	5	A	C6-N1-C2	-5.64	115.22	118.60
40	BG	54	G	P-O5'-C5'	-5.64	111.87	120.90
40	BG	76	C	P-O5'-C5'	-5.64	111.87	120.90
40	BG	79	U	C5-C6-N1	-5.64	119.88	122.70
41	BH	129	G	C8-N9-C4	5.64	108.66	106.40
85	AA	261	U	N1-C1'-C2'	-5.64	105.79	112.00
85	AA	484	G	C4-N9-C1'	-5.64	119.17	126.50
85	AA	611	G	N3-C4-N9	5.64	129.38	126.00
85	AA	648	G	C3'-C2'-C1'	-5.64	96.99	101.50
85	AA	726	U	C6-N1-C1'	-5.64	113.30	121.20
85	AA	1570	A	C8-N9-C1'	-5.64	117.54	127.70
30	AW	10	TYR	CA-C-N	-5.64	101.31	117.10
34	BA	86	A	C5-N7-C8	-5.64	101.08	103.90
34	BA	504	A	O3'-P-O5'	5.64	114.72	104.00
34	BA	719	G	N1-C6-O6	-5.64	116.52	119.90
34	BA	1054	U	O3'-P-O5'	5.64	114.72	104.00
35	BB	828	G	C1'-O4'-C4'	-5.64	105.39	109.90
36	BC	159	U	P-O3'-C3'	5.64	126.47	119.70
37	BD	14	C	O4'-C1'-C2'	5.64	112.68	107.60
37	BD	73	U	O5'-C5'-C4'	-5.64	100.98	111.70
85	AA	387	U	P-O5'-C5'	5.64	129.93	120.90
85	AA	438	G	C5-C6-N1	5.64	114.32	111.50
85	AA	536	C	O4'-C1'-N1	5.64	112.71	108.20
85	AA	2106	C	P-O5'-C5'	5.64	129.93	120.90
85	AA	2153	G	N1-C2-N3	-5.64	120.52	123.90
4	A3	118	LYS	N-CA-CB	5.64	120.75	110.60
34	BA	494	A	C4-N9-C1'	-5.64	116.15	126.30
34	BA	828	A	C8-N9-C1'	5.64	137.85	127.70
35	BB	3	C	O4'-C4'-C3'	5.64	110.61	106.10
35	BB	565	U	P-O5'-C5'	-5.64	111.88	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	632	U	O4'-C4'-C3'	-5.64	98.36	104.00
35	BB	688	U	C2-N1-C1'	-5.64	110.93	117.70
35	BB	1042	U	C4'-C3'-C2'	5.64	108.24	102.60
35	BB	1242	C	N1-C2-O2	5.64	122.28	118.90
35	BB	1322	A	C1'-O4'-C4'	-5.64	105.39	109.90
35	BB	1522	G	C5'-C4'-C3'	5.64	125.02	116.00
38	BE	73	A	C8-N9-C4	-5.64	103.54	105.80
41	BH	27	A	C5-C6-N6	5.64	128.21	123.70
41	BH	28	U	O5'-P-OP2	-5.64	100.62	105.70
44	BK	136	LEU	N-CA-C	-5.64	95.77	111.00
53	BT	88	ARG	NE-CZ-NH1	5.64	123.12	120.30
85	AA	424	A	C5'-C4'-O4'	-5.64	102.33	109.10
85	AA	537	G	C5'-C4'-C3'	-5.64	106.98	116.00
85	AA	703	U	N1-C2-N3	5.64	118.28	114.90
85	AA	1229	G	O4'-C1'-N9	5.64	112.71	108.20
85	AA	1281	G	N1-C2-N3	-5.64	120.52	123.90
85	AA	1537	A	O4'-C1'-N9	5.64	112.71	108.20
86	AB	11	C	C5'-C4'-C3'	-5.64	106.98	116.00
34	BA	46	C	P-O5'-C5'	-5.64	111.88	120.90
34	BA	181	G	P-O3'-C3'	-5.64	112.93	119.70
34	BA	921	G	P-O5'-C5'	-5.64	111.88	120.90
44	BK	99	ILE	CB-CA-C	-5.64	100.32	111.60
85	AA	34	G	O3'-P-O5'	-5.64	93.29	104.00
85	AA	982	G	N1-C2-N2	5.64	121.27	116.20
85	AA	1538	C	C6-N1-C2	-5.64	118.05	120.30
21	AM	89	ARG	NE-CZ-NH2	-5.64	117.48	120.30
34	BA	174	A	C4'-C3'-C2'	-5.64	96.96	102.60
34	BA	231	U	N1-C2-N3	5.64	118.28	114.90
34	BA	526	C	O4'-C1'-N1	5.64	112.71	108.20
35	BB	25	A	O5'-C5'-C4'	-5.64	100.99	111.70
35	BB	106	A	O4'-C1'-N9	5.64	112.71	108.20
35	BB	385	C	C1'-O4'-C4'	-5.64	105.39	109.90
35	BB	1480	G	O4'-C1'-C2'	5.64	112.67	107.60
38	BE	48	G	C8-N9-C4	-5.64	104.14	106.40
38	BE	75	C	O4'-C1'-N1	5.64	112.71	108.20
41	BH	5	G	C4-C5-C6	-5.64	115.42	118.80
60	Ba	74	VAL	CB-CA-C	-5.64	100.69	111.40
85	AA	473	C	P-O3'-C3'	5.64	126.46	119.70
85	AA	520	A	OP1-P-OP2	-5.64	111.14	119.60
85	AA	1320	G	C8-N9-C4	-5.64	104.14	106.40
85	AA	2153	G	C4-N9-C1'	-5.64	119.17	126.50
85	AA	2174	G	C1'-O4'-C4'	-5.64	105.39	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	138	C	C1'-O4'-C4'	-5.63	105.39	109.90
34	BA	588	C	O4'-C1'-N1	-5.63	103.69	108.20
34	BA	1039	G	C8-N9-C1'	5.63	134.32	127.00
34	BA	1259	C	O3'-P-O5'	-5.63	93.29	104.00
35	BB	74	U	O5'-C5'-C4'	-5.63	100.99	111.70
35	BB	666	A	C8-N9-C4	5.63	108.05	105.80
35	BB	1377	A	O3'-P-O5'	-5.63	93.29	104.00
35	BB	1462	G	N3-C2-N2	5.63	123.84	119.90
39	BF	5	U	O4'-C4'-C3'	-5.63	98.37	104.00
40	BG	66	C	O4'-C1'-C2'	5.63	112.67	107.60
41	BH	30	C	C6-N1-C1'	-5.63	114.04	120.80
54	BU	92	ARG	NE-CZ-NH2	-5.63	117.48	120.30
58	BY	20	ARG	NE-CZ-NH1	5.63	123.12	120.30
63	Bd	14	ARG	NE-CZ-NH2	-5.63	117.48	120.30
85	AA	84	C	O4'-C1'-N1	5.63	112.71	108.20
85	AA	633	C	C6-N1-C1'	-5.63	114.04	120.80
85	AA	879	G	C5'-C4'-O4'	5.63	115.86	109.10
85	AA	992	G	C3'-C2'-C1'	-5.63	96.99	101.50
85	AA	1501	A	O4'-C1'-N9	5.63	112.71	108.20
85	AA	1518	A	C4-C5-C6	-5.63	114.18	117.00
85	AA	1964	A	C5'-C4'-O4'	5.63	115.86	109.10
86	AB	67	C	C4'-C3'-C2'	-5.63	96.97	102.60
23	AP	200	PHE	CB-CG-CD1	5.63	124.74	120.80
34	BA	727	G	O4'-C1'-N9	5.63	112.71	108.20
34	BA	1077	G	C5-C6-N1	5.63	114.32	111.50
34	BA	1302	C	O4'-C1'-N1	5.63	112.71	108.20
35	BB	4	C	N3-C4-N4	-5.63	114.06	118.00
35	BB	583	G	O4'-C1'-N9	5.63	112.71	108.20
35	BB	1514	G	C3'-C2'-C1'	-5.63	96.99	101.50
36	BC	113	G	C6-C5-N7	-5.63	127.02	130.40
84	By	6	HIS	O-C-N	-5.63	113.69	122.70
7	A6	177	ASN	CA-CB-CG	-5.63	101.01	113.40
11	AC	121	ARG	NE-CZ-NH1	5.63	123.12	120.30
34	BA	253	U	P-O5'-C5'	-5.63	111.89	120.90
34	BA	923	C	C1'-O4'-C4'	-5.63	105.39	109.90
34	BA	1040	G	C3'-C2'-C1'	-5.63	96.99	101.50
34	BA	1103	G	C8-N9-C1'	5.63	134.32	127.00
34	BA	1455	C	O4'-C1'-N1	5.63	112.70	108.20
34	BA	1745	G	P-O3'-C3'	-5.63	112.94	119.70
35	BB	25	A	C5'-C4'-C3'	-5.63	106.99	116.00
35	BB	412	A	C5'-C4'-C3'	-5.63	106.99	116.00
35	BB	650	A	N1-C6-N6	-5.63	115.22	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1178	A	C2-N3-C4	-5.63	107.78	110.60
35	BB	1178	A	O4'-C4'-C3'	-5.63	98.37	104.00
35	BB	1364	C	C2-N1-C1'	-5.63	112.61	118.80
35	BB	1448	U	N3-C2-O2	-5.63	118.26	122.20
35	BB	1493	A	C4-N9-C1'	-5.63	116.16	126.30
38	BE	24	G	O4'-C1'-N9	5.63	112.71	108.20
57	BX	63	PRO	N-CA-C	5.63	126.74	112.10
85	AA	487	G	N3-C2-N2	5.63	123.84	119.90
85	AA	1276	A	O3'-P-O5'	5.63	114.70	104.00
85	AA	1486	G	N9-C1'-C2'	-5.63	105.80	112.00
85	AA	1975	G	C2'-C3'-O3'	5.63	122.71	113.70
34	BA	889	U	N1-C2-N3	5.63	118.28	114.90
34	BA	1162	U	N3-C2-O2	-5.63	118.26	122.20
34	BA	1415	C	C1'-O4'-C4'	-5.63	105.40	109.90
35	BB	420	U	C2-N3-C4	-5.63	123.62	127.00
35	BB	465	C	N3-C4-N4	-5.63	114.06	118.00
35	BB	567	G	C5'-C4'-O4'	5.63	115.86	109.10
35	BB	1029	U	O4'-C1'-C2'	5.63	112.67	107.60
38	BE	200	A	C8-N9-C4	5.63	108.05	105.80
40	BG	135	C	C4'-C3'-C2'	-5.63	96.97	102.60
40	BG	156	G	C5-N7-C8	-5.63	101.48	104.30
85	AA	1349	A	C5'-C4'-C3'	-5.63	106.99	116.00
85	AA	1650	G	C2'-C3'-O3'	5.63	122.71	113.70
85	AA	1829	C	C1'-O4'-C4'	-5.63	105.40	109.90
34	BA	962	U	C1'-O4'-C4'	-5.63	105.40	109.90
34	BA	1806	A	O4'-C1'-N9	5.63	112.70	108.20
35	BB	22	A	C8-N9-C1'	5.63	137.83	127.70
35	BB	35	G	N1-C6-O6	-5.63	116.52	119.90
35	BB	70	A	N1-C6-N6	-5.63	115.22	118.60
35	BB	606	C	N3-C2-O2	-5.63	117.96	121.90
35	BB	630	A	P-O3'-C3'	-5.63	112.94	119.70
35	BB	823	G	O4'-C1'-N9	5.63	112.70	108.20
35	BB	1425	A	O5'-P-OP2	5.63	117.45	110.70
35	BB	1450	G	C5-C6-N1	5.63	114.31	111.50
38	BE	208	G	N1-C6-O6	5.63	123.28	119.90
40	BG	176	G	C4'-C3'-C2'	5.63	108.23	102.60
40	BG	180	C	P-O3'-C3'	-5.63	112.95	119.70
41	BH	102	C	N3-C4-N4	-5.63	114.06	118.00
71	BI	116	HIS	N-CA-CB	5.63	120.73	110.60
85	AA	890	U	N1-C2-N3	5.63	118.28	114.90
85	AA	1566	A	C8-N9-C4	5.63	108.05	105.80
85	AA	1921	G	N3-C2-N2	-5.63	115.96	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	AC	164	ARG	N-CA-CB	-5.63	100.47	110.60
34	BA	241	U	C3'-C2'-C1'	-5.63	97.00	101.50
34	BA	376	U	C2-N1-C1'	-5.63	110.95	117.70
34	BA	435	U	N3-C2-O2	-5.63	118.26	122.20
34	BA	1333	G	C4-N9-C1'	-5.63	119.19	126.50
34	BA	1460	U	C6-N1-C1'	-5.63	113.32	121.20
34	BA	1483	U	N1-C2-O2	5.63	126.74	122.80
35	BB	345	U	O4'-C1'-N1	5.63	112.70	108.20
35	BB	433	C	P-O5'-C5'	5.63	129.90	120.90
35	BB	585	U	O4'-C1'-N1	5.63	112.70	108.20
36	BC	32	U	C4'-C3'-C2'	-5.63	96.97	102.60
36	BC	69	U	C5-C4-O4	-5.63	122.52	125.90
36	BC	72	A	P-O3'-C3'	-5.63	112.95	119.70
38	BE	200	A	O4'-C1'-N9	5.63	112.70	108.20
41	BH	127	A	O4'-C1'-N9	5.63	112.70	108.20
79	Bt	15	GLU	N-CA-CB	5.63	120.73	110.60
85	AA	17	C	C5-C6-N1	5.63	123.81	121.00
85	AA	77	C	C6-N1-C1'	-5.63	114.05	120.80
85	AA	1229	G	N7-C8-N9	5.63	115.91	113.10
85	AA	1480	C	C4'-C3'-C2'	-5.63	96.97	102.60
18	AJ	119	ARG	CB-CA-C	-5.62	99.15	110.40
34	BA	461	A	P-O5'-C5'	-5.62	111.90	120.90
34	BA	1093	G	C4-N9-C1'	-5.62	119.19	126.50
34	BA	1561	C	N3-C2-O2	-5.62	117.96	121.90
36	BC	124	A	C4-C5-C6	5.62	119.81	117.00
37	BD	65	G	C5-C6-N1	5.62	114.31	111.50
38	BE	94	U	P-O5'-C5'	-5.62	111.90	120.90
49	BP	11	ARG	N-CA-CB	5.62	120.73	110.60
85	AA	578	U	O3'-P-O5'	-5.62	93.31	104.00
85	AA	920	A	P-O3'-C3'	-5.62	112.95	119.70
6	A5	7	ARG	NE-CZ-NH2	-5.62	117.49	120.30
34	BA	12	G	N3-C4-N9	-5.62	122.63	126.00
34	BA	56	G	N1-C6-O6	5.62	123.27	119.90
34	BA	157	U	P-O5'-C5'	5.62	129.90	120.90
34	BA	449	G	C3'-C2'-C1'	-5.62	97.00	101.50
34	BA	484	A	C2'-C3'-O3'	5.62	122.70	113.70
35	BB	498	G	C8-N9-C1'	5.62	134.31	127.00
35	BB	1038	G	C5-C6-N1	5.62	114.31	111.50
53	BT	174	ARG	NE-CZ-NH1	5.62	123.11	120.30
72	Bm	76	ARG	N-CA-CB	5.62	120.72	110.60
85	AA	614	U	C5'-C4'-C3'	5.62	125.00	116.00
85	AA	738	C	N1-C2-N3	5.62	123.14	119.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1295	G	C8-N9-C4	-5.62	104.15	106.40
85	AA	2106	C	C6-N1-C2	-5.62	118.05	120.30
27	AT	67	PHE	CB-CG-CD1	-5.62	116.86	120.80
34	BA	96	G	C5'-C4'-C3'	-5.62	107.00	116.00
34	BA	330	A	C3'-C2'-C1'	-5.62	97.00	101.50
34	BA	423	G	C5-N7-C8	-5.62	101.49	104.30
34	BA	964	U	O4'-C1'-N1	5.62	112.70	108.20
34	BA	1487	U	N3-C2-O2	-5.62	118.27	122.20
35	BB	458	U	O4'-C1'-N1	5.62	112.70	108.20
35	BB	489	A	C2-N3-C4	-5.62	107.79	110.60
35	BB	529	A	N7-C8-N9	5.62	116.61	113.80
35	BB	827	U	C6-N1-C1'	5.62	129.07	121.20
35	BB	845	C	C4'-C3'-O3'	5.62	124.24	113.00
35	BB	1116	U	O5'-P-OP2	-5.62	100.64	105.70
35	BB	1384	A	O4'-C4'-C3'	-5.62	98.38	104.00
38	BE	61	A	C8-N9-C1'	5.62	137.82	127.70
39	BF	5	U	C5'-C4'-C3'	5.62	125.00	116.00
85	AA	54	C	C2-N1-C1'	-5.62	112.62	118.80
85	AA	353	G	N3-C2-N2	5.62	123.84	119.90
85	AA	560	C	P-O3'-C3'	5.62	126.44	119.70
85	AA	667	A	P-O3'-C3'	-5.62	112.95	119.70
85	AA	1323	G	C5-C6-O6	-5.62	125.23	128.60
85	AA	1353	U	O5'-C5'-C4'	5.62	122.38	111.70
85	AA	1884	A	C2'-C3'-O3'	5.62	122.69	113.70
85	AA	1931	C	C6-N1-C2	-5.62	118.05	120.30
34	BA	266	G	P-O5'-C5'	-5.62	111.91	120.90
34	BA	451	A	N9-C1'-C2'	-5.62	105.82	112.00
35	BB	347	G	O4'-C1'-N9	5.62	112.70	108.20
36	BC	52	A	O5'-C5'-C4'	-5.62	101.02	111.70
60	Ba	94	GLY	N-CA-C	-5.62	99.05	113.10
85	AA	125	A	C1'-O4'-C4'	-5.62	105.40	109.90
85	AA	586	G	N9-C1'-C2'	-5.62	105.82	112.00
85	AA	859	G	N9-C1'-C2'	-5.62	105.82	112.00
85	AA	2173	A	C5'-C4'-C3'	-5.62	107.01	116.00
5	A4	158	PHE	CB-CG-CD2	-5.62	116.87	120.80
34	BA	363	G	C5'-C4'-C3'	5.62	124.99	116.00
34	BA	465	A	N9-C1'-C2'	-5.62	105.82	112.00
34	BA	764	G	N3-C4-N9	5.62	129.37	126.00
34	BA	1079	C	C2-N3-C4	-5.62	117.09	119.90
34	BA	1460	U	C6-N1-C2	-5.62	117.63	121.00
34	BA	1771	U	O4'-C1'-N1	5.62	112.69	108.20
35	BB	68	G	P-O5'-C5'	-5.62	111.91	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	394	A	N1-C2-N3	-5.62	126.49	129.30
35	BB	568	A	C5-C6-N6	-5.62	119.20	123.70
35	BB	629	C	C4-C5-C6	-5.62	114.59	117.40
35	BB	1495	U	C2'-C3'-O3'	5.62	122.69	113.70
35	BB	1509	G	O3'-P-O5'	-5.62	93.32	104.00
36	BC	113	G	C5'-C4'-C3'	-5.62	107.01	116.00
38	BE	25	U	C2'-C3'-O3'	5.62	122.69	113.70
41	BH	113	G	C3'-C2'-C1'	-5.62	97.00	101.50
65	Bf	352	ALA	CA-C-N	-5.62	104.84	117.20
67	Bh	30	ARG	N-CA-CB	-5.62	100.49	110.60
85	AA	54	C	C3'-C2'-C1'	-5.62	97.00	101.50
85	AA	245	A	C4'-C3'-C2'	-5.62	96.98	102.60
85	AA	345	U	C4'-C3'-C2'	-5.62	96.98	102.60
85	AA	1147	A	P-O3'-C3'	5.62	126.44	119.70
85	AA	1275	A	C8-N9-C4	5.62	108.05	105.80
34	BA	125	G	N1-C6-O6	-5.62	116.53	119.90
35	BB	5	A	O4'-C4'-C3'	-5.62	98.38	104.00
38	BE	49	A	O4'-C1'-N9	5.62	112.69	108.20
48	BO	82	SER	CB-CA-C	-5.62	99.43	110.10
81	Bv	89	ASP	CA-CB-CG	-5.62	101.04	113.40
85	AA	547	A	C1'-O4'-C4'	-5.62	105.41	109.90
85	AA	793	C	O5'-P-OP2	5.62	117.44	110.70
85	AA	2225	G	C6-N1-C2	-5.62	121.73	125.10
4	A3	85	PHE	C-N-CA	5.62	135.74	121.70
31	AX	93	ARG	NE-CZ-NH1	5.62	123.11	120.30
34	BA	344	G	O4'-C1'-N9	5.62	112.69	108.20
34	BA	383	G	N1-C2-N2	5.62	121.25	116.20
34	BA	666	C	C4'-C3'-C2'	-5.62	96.98	102.60
34	BA	742	C	P-O3'-C3'	-5.62	112.96	119.70
34	BA	778	U	C4'-C3'-C2'	-5.62	96.98	102.60
34	BA	803	U	C2'-C3'-O3'	5.62	122.68	113.70
34	BA	1346	U	C5'-C4'-O4'	5.62	115.84	109.10
34	BA	1572	G	C5-C6-O6	5.62	131.97	128.60
34	BA	1628	A	C3'-C2'-C1'	-5.62	97.01	101.50
35	BB	1110	G	C8-N9-C1'	5.62	134.30	127.00
35	BB	1512	C	C5'-C4'-O4'	-5.62	102.36	109.10
36	BC	107	C	N1-C1'-C2'	-5.62	105.82	112.00
37	BD	38	U	C2-N3-C4	-5.62	123.63	127.00
41	BH	11	C	O5'-C5'-C4'	5.62	122.37	111.70
85	AA	674	U	C5'-C4'-C3'	-5.62	107.02	116.00
85	AA	820	G	C5-C6-O6	5.62	131.97	128.60
85	AA	1637	C	O3'-P-O5'	-5.62	93.33	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1664	G	P-O5'-C5'	-5.62	111.92	120.90
23	AP	111	PHE	CB-CG-CD1	5.61	124.73	120.80
34	BA	504	A	C8-N9-C4	5.61	108.05	105.80
34	BA	750	C	P-O5'-C5'	-5.61	111.92	120.90
35	BB	384	A	C1'-O4'-C4'	-5.61	105.41	109.90
35	BB	1331	U	C1'-O4'-C4'	-5.61	105.41	109.90
38	BE	55	C	O4'-C1'-N1	5.61	112.69	108.20
38	BE	79	G	O4'-C1'-N9	5.61	112.69	108.20
40	BG	10	U	C6-N1-C1'	5.61	129.06	121.20
85	AA	192	G	N3-C4-C5	-5.61	125.79	128.60
85	AA	324	U	C4'-C3'-C2'	5.61	108.21	102.60
85	AA	464	A	O5'-C5'-C4'	-5.61	101.03	111.70
85	AA	993	G	P-O3'-C3'	-5.61	112.96	119.70
85	AA	1223	A	C4-C5-C6	-5.61	114.19	117.00
85	AA	1456	A	C1'-O4'-C4'	-5.61	105.41	109.90
85	AA	2008	G	N1-C2-N3	-5.61	120.53	123.90
85	AA	2102	A	C5-C6-N6	-5.61	119.21	123.70
85	AA	2186	U	C5'-C4'-O4'	5.61	115.84	109.10
34	BA	423	G	C1'-O4'-C4'	-5.61	105.41	109.90
35	BB	545	C	O4'-C1'-N1	5.61	112.69	108.20
35	BB	823	G	N3-C4-C5	-5.61	125.79	128.60
35	BB	824	C	C5'-C4'-C3'	-5.61	107.02	116.00
67	Bh	28	GLU	C-N-CA	5.61	135.73	121.70
84	By	171	ARG	NE-CZ-NH2	-5.61	117.49	120.30
85	AA	2035	C	O4'-C1'-N1	5.61	112.69	108.20
85	AA	2125	A	C5-N7-C8	-5.61	101.09	103.90
34	BA	260	A	P-O5'-C5'	-5.61	111.92	120.90
34	BA	372	U	O5'-P-OP2	-5.61	100.65	105.70
34	BA	667	U	O4'-C1'-N1	5.61	112.69	108.20
34	BA	705	C	C4'-C3'-C2'	5.61	108.21	102.60
34	BA	883	C	O4'-C1'-N1	5.61	112.69	108.20
34	BA	901	C	N3-C2-O2	-5.61	117.97	121.90
34	BA	992	A	O4'-C1'-C2'	5.61	112.65	107.60
34	BA	1340	G	C6-N1-C2	-5.61	121.73	125.10
34	BA	1693	U	C2-N3-C4	-5.61	123.63	127.00
35	BB	147	C	C6-N1-C1'	-5.61	114.07	120.80
35	BB	638	G	C2'-C3'-O3'	5.61	122.68	113.70
35	BB	1147	G	C5-C6-O6	-5.61	125.23	128.60
35	BB	1336	G	N1-C6-O6	5.61	123.27	119.90
35	BB	1353	G	C8-N9-C4	-5.61	104.16	106.40
37	BD	79	G	C2-N3-C4	5.61	114.70	111.90
85	AA	4	C	C5-C4-N4	-5.61	116.27	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	124	A	C3'-C2'-C1'	5.61	105.99	101.50
85	AA	782	G	C3'-C2'-C1'	-5.61	97.01	101.50
85	AA	815	G	N9-C1'-C2'	-5.61	105.83	112.00
85	AA	999	A	O4'-C1'-N9	5.61	112.69	108.20
85	AA	1295	G	N3-C4-C5	-5.61	125.79	128.60
85	AA	1299	A	C5'-C4'-C3'	5.61	124.98	116.00
85	AA	1470	A	C8-N9-C4	5.61	108.04	105.80
85	AA	2058	C	N1-C2-N3	5.61	123.13	119.20
86	AB	28	G	C5-C6-O6	-5.61	125.23	128.60
26	AS	105	PHE	CB-CG-CD2	-5.61	116.87	120.80
34	BA	27	G	O4'-C1'-N9	5.61	112.69	108.20
34	BA	1318	G	O4'-C4'-C3'	-5.61	98.39	104.00
34	BA	1658	G	C3'-C2'-C1'	-5.61	97.01	101.50
35	BB	403	U	C2-N3-C4	-5.61	123.64	127.00
35	BB	852	G	OP2-P-O3'	5.61	117.54	105.20
35	BB	1443	C	O4'-C1'-N1	5.61	112.69	108.20
46	BM	67	ARG	NE-CZ-NH1	5.61	123.11	120.30
85	AA	66	U	C5-C4-O4	5.61	129.26	125.90
85	AA	138	C	C6-N1-C2	-5.61	118.06	120.30
85	AA	776	C	P-O5'-C5'	-5.61	111.92	120.90
85	AA	1266	C	C6-N1-C2	5.61	122.54	120.30
85	AA	1892	G	C1'-O4'-C4'	-5.61	105.41	109.90
14	AF	116	ARG	NE-CZ-NH2	-5.61	117.50	120.30
34	BA	10	G	C4-C5-C6	-5.61	115.44	118.80
34	BA	39	C	N3-C4-N4	5.61	121.93	118.00
34	BA	332	U	O4'-C1'-N1	5.61	112.69	108.20
34	BA	389	U	C3'-C2'-C1'	-5.61	97.02	101.50
34	BA	605	G	O5'-C5'-C4'	5.61	122.36	111.70
34	BA	794	G	C5'-C4'-C3'	-5.61	107.03	116.00
34	BA	796	G	C8-N9-C4	-5.61	104.16	106.40
34	BA	1395	C	C2-N3-C4	-5.61	117.10	119.90
34	BA	1415	C	O5'-P-OP2	-5.61	100.65	105.70
35	BB	1031	G	O4'-C1'-C2'	-5.61	100.19	105.80
36	BC	111	C	C5-C4-N4	-5.61	116.28	120.20
37	BD	105	G	C3'-C2'-C1'	-5.61	97.02	101.50
40	BG	119	A	N1-C6-N6	5.61	121.97	118.60
40	BG	172	C	C3'-C2'-C1'	-5.61	97.02	101.50
51	BR	141	ARG	NE-CZ-NH2	5.61	123.10	120.30
83	Bx	50	GLY	C-N-CA	5.61	135.72	121.70
85	AA	3	U	C4'-C3'-C2'	-5.61	96.99	102.60
85	AA	4	C	C5'-C4'-C3'	-5.61	107.03	116.00
85	AA	780	U	O4'-C1'-N1	5.61	112.69	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	851	G	N9-C4-C5	-5.61	103.16	105.40
85	AA	925	G	O5'-P-OP2	-5.61	100.65	105.70
85	AA	926	C	C6-N1-C1'	5.61	127.53	120.80
85	AA	1832	G	C4'-C3'-C2'	-5.61	96.99	102.60
85	AA	1850	G	C5'-C4'-O4'	5.61	115.83	109.10
85	AA	2061	C	C4'-C3'-C2'	-5.61	96.99	102.60
21	AM	109	ARG	NE-CZ-NH1	5.61	123.10	120.30
31	AX	9	MET	CA-CB-CG	5.61	122.83	113.30
34	BA	537	C	P-O5'-C5'	5.61	129.87	120.90
34	BA	771	A	C5'-C4'-C3'	5.61	124.97	116.00
34	BA	1563	G	C1'-O4'-C4'	5.61	114.38	109.90
35	BB	621	C	C5'-C4'-C3'	-5.61	107.03	116.00
35	BB	926	C	O4'-C1'-N1	5.61	112.68	108.20
38	BE	201	A	O4'-C1'-N9	5.61	112.69	108.20
85	AA	30	G	C1'-O4'-C4'	-5.61	105.42	109.90
85	AA	442	G	C5'-C4'-C3'	-5.61	107.03	116.00
85	AA	742	U	C5-C4-O4	5.61	129.26	125.90
85	AA	756	G	O5'-P-OP1	5.61	117.43	110.70
85	AA	1106	A	C5-N7-C8	-5.61	101.10	103.90
85	AA	1844	A	P-O3'-C3'	-5.61	112.97	119.70
85	AA	1892	G	N1-C6-O6	5.61	123.26	119.90
85	AA	2217	A	C5-C6-N6	-5.61	119.22	123.70
34	BA	83	G	N9-C1'-C2'	-5.60	105.84	112.00
85	AA	1879	U	C2'-C3'-O3'	5.60	122.67	113.70
85	AA	1990	U	P-O3'-C3'	-5.60	112.98	119.70
85	AA	2167	A	C4-N9-C1'	-5.60	116.21	126.30
34	BA	173	U	O4'-C1'-N1	5.60	112.68	108.20
34	BA	246	G	N3-C2-N2	5.60	123.82	119.90
34	BA	289	A	C5-C6-N6	-5.60	119.22	123.70
34	BA	515	U	C5'-C4'-C3'	5.60	124.97	116.00
34	BA	568	G	N9-C1'-C2'	-5.60	105.84	112.00
34	BA	799	A	P-O3'-C3'	5.60	126.42	119.70
34	BA	834	C	OP1-P-OP2	-5.60	111.20	119.60
34	BA	924	U	C6-N1-C2	-5.60	117.64	121.00
34	BA	1055	U	C4'-C3'-C2'	5.60	108.20	102.60
34	BA	1645	C	C3'-C2'-C1'	-5.60	97.02	101.50
35	BB	71	A	C1'-O4'-C4'	-5.60	105.42	109.90
35	BB	271	C	O4'-C1'-N1	5.60	112.68	108.20
35	BB	795	A	O4'-C1'-N9	5.60	112.68	108.20
35	BB	906	G	N1-C6-O6	5.60	123.26	119.90
36	BC	20	C	C6-N1-C2	-5.60	118.06	120.30
38	BE	112	G	O5'-C5'-C4'	5.60	122.34	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	154	A	O4'-C1'-N9	5.60	112.68	108.20
39	BF	61	A	O5'-C5'-C4'	5.60	122.35	111.70
42	BI	19	THR	C-N-CA	5.60	135.71	121.70
53	BT	189	ARG	NE-CZ-NH1	5.60	123.10	120.30
85	AA	249	C	C6-N1-C2	-5.60	118.06	120.30
85	AA	1302	A	N1-C6-N6	5.60	121.96	118.60
85	AA	1814	U	C3'-C2'-C1'	-5.60	97.02	101.50
85	AA	2150	G	C5-C6-O6	-5.60	125.24	128.60
1	A0	117	ARG	N-CA-C	-5.60	95.88	111.00
34	BA	223	U	C5'-C4'-O4'	-5.60	102.38	109.10
34	BA	297	A	C8-N9-C4	-5.60	103.56	105.80
34	BA	442	G	O4'-C1'-C2'	-5.60	100.20	105.80
34	BA	585	G	C5'-C4'-C3'	-5.60	107.04	116.00
34	BA	1410	C	C1'-O4'-C4'	-5.60	105.42	109.90
34	BA	1607	U	C1'-O4'-C4'	-5.60	105.42	109.90
35	BB	1515	C	C2-N3-C4	-5.60	117.10	119.90
41	BH	118	U	C3'-C2'-C1'	-5.60	97.02	101.50
59	BZ	55	VAL	CB-CA-C	-5.60	100.76	111.40
85	AA	750	A	C5-C6-N6	-5.60	119.22	123.70
34	BA	812	A	C4-N9-C1'	-5.60	116.22	126.30
34	BA	1166	A	C4-C5-C6	-5.60	114.20	117.00
34	BA	1496	G	N9-C4-C5	-5.60	103.16	105.40
34	BA	1559	C	N3-C2-O2	-5.60	117.98	121.90
34	BA	1617	U	C3'-C2'-C1'	-5.60	97.02	101.50
34	BA	1686	G	C8-N9-C4	5.60	108.64	106.40
34	BA	1697	U	OP1-P-OP2	-5.60	111.20	119.60
34	BA	1720	U	N3-C2-O2	-5.60	118.28	122.20
34	BA	1790	U	O4'-C1'-C2'	5.60	112.64	107.60
34	BA	1804	A	C4'-C3'-C2'	-5.60	97.00	102.60
37	BD	55	A	C5-C6-N6	5.60	128.18	123.70
38	BE	79	G	C5-C6-O6	-5.60	125.24	128.60
38	BE	131	C	C4'-C3'-C2'	5.60	108.20	102.60
53	BT	117	ARG	N-CA-CB	-5.60	100.52	110.60
85	AA	303	A	C5-C6-N6	5.60	128.18	123.70
85	AA	304	G	C3'-C2'-C1'	5.60	105.98	101.50
85	AA	422	G	C6-N1-C2	-5.60	121.74	125.10
85	AA	721	C	C4'-C3'-C2'	-5.60	97.00	102.60
85	AA	972	G	O5'-C5'-C4'	5.60	122.34	111.70
85	AA	1106	A	C5'-C4'-O4'	5.60	115.82	109.10
25	AR	14	VAL	N-CA-C	-5.60	95.89	111.00
34	BA	108	A	C5-C6-N1	5.60	120.50	117.70
34	BA	159	U	C3'-C2'-C1'	-5.60	97.02	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	499	C	C4'-C3'-C2'	5.60	108.20	102.60
34	BA	575	U	O3'-P-O5'	-5.60	93.37	104.00
34	BA	940	C	C4'-C3'-C2'	5.60	108.20	102.60
34	BA	1053	U	N1-C2-N3	5.60	118.26	114.90
34	BA	1424	G	C5'-C4'-C3'	-5.60	107.04	116.00
34	BA	1543	A	O4'-C1'-C2'	5.60	112.64	107.60
35	BB	531	U	O4'-C1'-N1	5.60	112.68	108.20
35	BB	829	C	C6-N1-C1'	5.60	127.52	120.80
35	BB	1046	C	C2-N3-C4	-5.60	117.10	119.90
35	BB	1314	G	P-O5'-C5'	-5.60	111.95	120.90
40	BG	104	A	C5'-C4'-C3'	-5.60	107.04	116.00
33	AZ	97	ARG	NE-CZ-NH2	-5.60	117.50	120.30
34	BA	1194	G	N1-C6-O6	-5.60	116.54	119.90
34	BA	1225	A	C1'-O4'-C4'	-5.60	105.42	109.90
34	BA	1249	G	N9-C1'-C2'	-5.60	105.84	112.00
34	BA	1337	A	C3'-C2'-C1'	-5.60	97.02	101.50
34	BA	1501	U	C5-C6-N1	-5.60	119.90	122.70
35	BB	1181	A	N9-C1'-C2'	-5.60	105.84	112.00
36	BC	154	A	N9-C4-C5	-5.60	103.56	105.80
51	BR	139	TYR	CB-CA-C	5.60	121.59	110.40
74	Bo	19	GLY	N-CA-C	5.60	127.09	113.10
82	Bw	46	ALA	CB-CA-C	-5.60	101.70	110.10
85	AA	438	G	P-O3'-C3'	-5.60	112.98	119.70
85	AA	988	C	C2-N1-C1'	5.60	124.96	118.80
85	AA	2171	A	OP2-P-O3'	5.60	117.51	105.20
34	BA	710	A	C5'-C4'-C3'	-5.59	107.05	116.00
34	BA	934	G	C8-N9-C4	5.59	108.64	106.40
34	BA	1300	G	C8-N9-C1'	5.59	134.27	127.00
34	BA	1634	A	O5'-C5'-C4'	5.59	122.33	111.70
34	BA	1728	G	C2-N3-C4	-5.59	109.10	111.90
35	BB	1498	G	C4-C5-C6	-5.59	115.44	118.80
35	BB	1522	G	C8-N9-C1'	5.59	134.27	127.00
38	BE	206	G	O4'-C1'-N9	5.59	112.67	108.20
39	BF	18	U	C6-N1-C1'	-5.59	113.37	121.20
40	BG	30	C	C6-N1-C2	-5.59	118.06	120.30
40	BG	91	U	O3'-P-O5'	5.59	114.63	104.00
53	BT	81	ARG	CB-CG-CD	5.59	126.15	111.60
64	Be	133	TYR	N-CA-C	-5.59	95.90	111.00
84	By	6	HIS	CA-C-N	5.59	129.51	117.20
85	AA	47	A	N3-C4-C5	-5.59	122.88	126.80
85	AA	257	U	O4'-C1'-N1	5.59	112.68	108.20
85	AA	455	G	N1-C2-N3	-5.59	120.54	123.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	873	U	O4'-C1'-N1	5.59	112.68	108.20
85	AA	1249	U	P-O5'-C5'	5.59	129.85	120.90
85	AA	1301	C	C6-N1-C2	-5.59	118.06	120.30
86	AB	43	C	C4'-C3'-C2'	-5.59	97.00	102.60
34	BA	78	U	C3'-C2'-C1'	-5.59	97.03	101.50
34	BA	976	C	C5-C4-N4	5.59	124.11	120.20
34	BA	1094	U	C2-N3-C4	-5.59	123.64	127.00
34	BA	1119	A	O4'-C1'-N9	5.59	112.67	108.20
34	BA	1485	U	N1-C1'-C2'	-5.59	105.85	112.00
35	BB	8	U	C2-N1-C1'	-5.59	110.99	117.70
35	BB	1068	G	N1-C6-O6	5.59	123.26	119.90
41	BH	4	U	C6-N1-C2	-5.59	117.64	121.00
41	BH	28	U	N1-C2-O2	5.59	126.72	122.80
47	BN	18	ASN	CA-CB-CG	-5.59	101.09	113.40
85	AA	329	G	C5-C6-O6	-5.59	125.24	128.60
85	AA	1016	G	C1'-O4'-C4'	-5.59	105.43	109.90
85	AA	1149	A	P-O3'-C3'	5.59	126.41	119.70
85	AA	1517	G	N1-C6-O6	5.59	123.26	119.90
23	AP	105	ARG	NE-CZ-NH1	5.59	123.09	120.30
34	BA	422	C	C5'-C4'-C3'	-5.59	107.06	116.00
34	BA	469	C	C6-N1-C2	-5.59	118.06	120.30
34	BA	709	C	C5'-C4'-C3'	-5.59	107.05	116.00
34	BA	941	G	N1-C2-N2	-5.59	111.17	116.20
34	BA	1090	A	P-O3'-C3'	-5.59	112.99	119.70
34	BA	1348	G	P-O3'-C3'	-5.59	112.99	119.70
34	BA	1695	G	C1'-O4'-C4'	-5.59	105.43	109.90
34	BA	1829	A	P-O5'-C5'	5.59	129.85	120.90
35	BB	673	C	P-O3'-C3'	-5.59	112.99	119.70
35	BB	1017	U	C3'-C2'-C1'	-5.59	97.03	101.50
35	BB	1503	U	C4'-C3'-C2'	-5.59	97.01	102.60
36	BC	3	C	N1-C2-N3	5.59	123.11	119.20
44	BK	49	CYS	N-CA-CB	5.59	120.66	110.60
59	BZ	106	LEU	C-N-CA	5.59	135.68	121.70
85	AA	373	G	C4'-C3'-C2'	5.59	108.19	102.60
85	AA	558	U	C3'-C2'-C1'	-5.59	97.03	101.50
85	AA	578	U	C5'-C4'-O4'	-5.59	102.39	109.10
85	AA	818	C	N3-C4-N4	5.59	121.91	118.00
85	AA	1457	C	C5-C6-N1	5.59	123.80	121.00
85	AA	2250	U	C3'-C2'-C1'	5.59	105.97	101.50
34	BA	52	G	C6-N1-C2	-5.59	121.75	125.10
34	BA	101	G	P-O5'-C5'	-5.59	111.96	120.90
34	BA	1415	C	C4'-C3'-C2'	5.59	108.19	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1729	G	N3-C2-N2	5.59	123.81	119.90
35	BB	719	G	P-O5'-C5'	5.59	129.84	120.90
35	BB	1015	U	P-O3'-C3'	-5.59	112.99	119.70
36	BC	156	A	O4'-C4'-C3'	-5.59	98.41	104.00
52	BS	116	HIS	N-CA-CB	5.59	120.66	110.60
85	AA	165	C	C4'-C3'-C2'	-5.59	97.01	102.60
85	AA	1324	G	C5'-C4'-C3'	-5.59	107.06	116.00
85	AA	1826	U	C2-N1-C1'	5.59	124.41	117.70
34	BA	548	G	N7-C8-N9	5.59	115.89	113.10
34	BA	702	G	O4'-C1'-N9	5.59	112.67	108.20
34	BA	1451	A	C5-C6-N6	-5.59	119.23	123.70
35	BB	391	G	O4'-C1'-N9	5.59	112.67	108.20
35	BB	456	A	N1-C6-N6	5.59	121.95	118.60
35	BB	1509	G	N9-C1'-C2'	-5.59	105.85	112.00
38	BE	182	U	P-O3'-C3'	-5.59	112.99	119.70
42	BI	108	CYS	CB-CA-C	5.59	121.58	110.40
85	AA	1661	U	O4'-C1'-C2'	5.59	112.63	107.60
85	AA	1700	C	C6-N1-C2	-5.59	118.06	120.30
15	AG	91	LEU	N-CA-CB	-5.59	99.23	110.40
34	BA	137	C	C1'-O4'-C4'	-5.59	105.43	109.90
34	BA	1700	C	P-O5'-C5'	-5.59	111.96	120.90
35	BB	776	U	O4'-C1'-N1	5.59	112.67	108.20
35	BB	1062	G	C5-C6-N1	5.59	114.29	111.50
35	BB	1467	A	N1-C6-N6	5.59	121.95	118.60
38	BE	184	G	C3'-C2'-C1'	-5.59	97.03	101.50
84	By	38	ARG	NE-CZ-NH2	5.59	123.09	120.30
85	AA	301	U	C2-N1-C1'	5.59	124.40	117.70
85	AA	340	G	C3'-C2'-C1'	-5.59	97.03	101.50
85	AA	443	A	P-O3'-C3'	5.59	126.40	119.70
85	AA	994	A	N1-C2-N3	-5.59	126.51	129.30
85	AA	1489	G	C4'-C3'-C2'	5.59	108.19	102.60
85	AA	1983	C	P-O5'-C5'	-5.59	111.96	120.90
30	AW	80	PHE	CB-CG-CD2	-5.58	116.89	120.80
34	BA	1053	U	O4'-C1'-N1	5.58	112.67	108.20
34	BA	1472	G	P-O3'-C3'	-5.58	113.00	119.70
34	BA	1755	U	O4'-C1'-N1	5.58	112.67	108.20
37	BD	25	G	C8-N9-C4	-5.58	104.17	106.40
47	BN	194	HIS	CA-CB-CG	5.58	123.09	113.60
59	BZ	7	ARG	N-CA-C	-5.58	95.92	111.00
77	Br	234	ASN	CA-CB-CG	-5.58	101.11	113.40
80	Bu	145	ARG	N-CA-CB	5.58	120.65	110.60
85	AA	1793	A	C5'-C4'-O4'	-5.58	102.40	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1797	U	O4'-C4'-C3'	-5.58	98.42	104.00
86	AB	57	G	C6-N1-C2	5.58	128.45	125.10
34	BA	111	U	N1-C1'-C2'	-5.58	105.86	112.00
34	BA	235	C	C2'-C3'-O3'	5.58	122.63	113.70
34	BA	861	C	C2-N3-C4	-5.58	117.11	119.90
34	BA	1023	G	N1-C6-O6	5.58	123.25	119.90
34	BA	1564	A	N1-C6-N6	5.58	121.95	118.60
35	BB	129	U	O4'-C1'-C2'	5.58	112.62	107.60
35	BB	383	U	C5'-C4'-O4'	5.58	115.80	109.10
35	BB	1356	G	C5-N7-C8	5.58	107.09	104.30
36	BC	99	U	O3'-P-O5'	-5.58	93.39	104.00
38	BE	97	G	P-O3'-C3'	-5.58	113.00	119.70
41	BH	22	A	N1-C6-N6	5.58	121.95	118.60
41	BH	102	C	C5'-C4'-O4'	-5.58	102.40	109.10
77	Br	277	LEU	N-CA-CB	5.58	121.57	110.40
85	AA	270	A	N1-C6-N6	-5.58	115.25	118.60
85	AA	584	G	C8-N9-C4	-5.58	104.17	106.40
85	AA	864	C	C1'-O4'-C4'	-5.58	105.43	109.90
85	AA	1493	A	N3-C4-N9	-5.58	122.93	127.40
34	BA	175	G	O4'-C4'-C3'	-5.58	98.42	104.00
34	BA	650	C	C3'-C2'-C1'	-5.58	97.03	101.50
34	BA	860	G	N9-C1'-C2'	-5.58	105.86	112.00
34	BA	1053	U	C2-N3-C4	-5.58	123.65	127.00
34	BA	1252	G	N9-C1'-C2'	-5.58	105.86	112.00
34	BA	1442	A	C5'-C4'-C3'	5.58	124.93	116.00
34	BA	1642	A	C3'-C2'-C1'	-5.58	97.03	101.50
35	BB	103	C	N3-C4-N4	-5.58	114.09	118.00
35	BB	1026	G	C5'-C4'-C3'	5.58	124.93	116.00
35	BB	1143	A	O4'-C1'-N9	5.58	112.67	108.20
45	BL	135	TYR	CB-CG-CD2	5.58	124.35	121.00
49	BP	48	ARG	NE-CZ-NH2	-5.58	117.51	120.30
85	AA	118	C	C2-N1-C1'	-5.58	112.66	118.80
85	AA	715	G	N1-C6-O6	-5.58	116.55	119.90
85	AA	855	G	N9-C1'-C2'	5.58	121.26	114.00
85	AA	1155	A	C4-N9-C1'	-5.58	116.25	126.30
85	AA	1268	C	C2-N1-C1'	-5.58	112.66	118.80
85	AA	1853	U	O4'-C1'-N1	5.58	112.67	108.20
85	AA	1945	A	O4'-C1'-N9	5.58	112.67	108.20
85	AA	2136	C	C2'-C3'-O3'	5.58	122.63	113.70
31	AX	116	ARG	NE-CZ-NH2	-5.58	117.51	120.30
34	BA	660	C	C2-N3-C4	-5.58	117.11	119.90
34	BA	1563	G	O4'-C1'-N9	-5.58	103.74	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	397	C	C5-C4-N4	5.58	124.11	120.20
85	AA	309	G	O4'-C1'-C2'	5.58	112.62	107.60
85	AA	365	G	C5-C6-O6	5.58	131.95	128.60
85	AA	400	G	C4-C5-C6	-5.58	115.45	118.80
85	AA	540	A	O4'-C1'-C2'	5.58	112.62	107.60
85	AA	1305	A	C5'-C4'-C3'	5.58	124.93	116.00
34	BA	54	A	C1'-O4'-C4'	-5.58	105.44	109.90
34	BA	89	G	N1-C2-N2	5.58	121.22	116.20
34	BA	967	C	P-O5'-C5'	-5.58	111.97	120.90
35	BB	122	U	C5-C6-N1	-5.58	119.91	122.70
35	BB	961	G	O4'-C1'-N9	5.58	112.66	108.20
35	BB	980	G	N1-C6-O6	5.58	123.25	119.90
35	BB	1076	U	N1-C2-N3	5.58	118.25	114.90
35	BB	1113	C	P-O5'-C5'	-5.58	111.97	120.90
35	BB	1202	G	N9-C1'-C2'	-5.58	105.86	112.00
35	BB	1213	U	N3-C4-C5	-5.58	111.25	114.60
35	BB	1421	C	C1'-O4'-C4'	-5.58	105.44	109.90
36	BC	123	G	O4'-C1'-N9	5.58	112.66	108.20
38	BE	69	C	C2-N3-C4	-5.58	117.11	119.90
85	AA	20	G	C5-C6-N1	5.58	114.29	111.50
85	AA	76	G	C8-N9-C1'	5.58	134.25	127.00
85	AA	82	A	C5'-C4'-C3'	-5.58	107.08	116.00
85	AA	290	G	N3-C2-N2	5.58	123.81	119.90
85	AA	465	A	N1-C2-N3	-5.58	126.51	129.30
85	AA	661	C	C2-N1-C1'	-5.58	112.66	118.80
85	AA	1191	G	C5-C6-N1	5.58	114.29	111.50
85	AA	1825	A	C8-N9-C4	-5.58	103.57	105.80
34	BA	811	C	P-O3'-C3'	-5.58	113.01	119.70
35	BB	919	U	O4'-C1'-N1	5.58	112.66	108.20
35	BB	1062	G	N1-C6-O6	5.58	123.25	119.90
41	BH	34	G	O4'-C1'-N9	5.58	112.66	108.20
47	BN	106	ARG	NE-CZ-NH1	5.58	123.09	120.30
48	BO	134	GLY	C-N-CA	5.58	134.01	122.30
65	Bf	101	LYS	CA-CB-CG	5.58	125.67	113.40
85	AA	132	G	C4-N9-C1'	-5.58	119.25	126.50
85	AA	1539	A	C8-N9-C4	5.58	108.03	105.80
85	AA	1667	C	N1-C2-N3	5.58	123.10	119.20
5	A4	81	ARG	NE-CZ-NH2	-5.58	117.51	120.30
34	BA	161	U	N3-C2-O2	5.58	126.10	122.20
34	BA	398	G	P-O5'-C5'	5.58	129.82	120.90
34	BA	526	C	N3-C4-N4	5.58	121.90	118.00
34	BA	661	C	O4'-C1'-C2'	5.58	112.62	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	854	A	P-O5'-C5'	-5.58	111.98	120.90
34	BA	1133	A	N1-C6-N6	5.58	121.94	118.60
34	BA	1612	C	N3-C4-N4	-5.58	114.10	118.00
35	BB	85	A	C5'-C4'-C3'	5.58	124.92	116.00
35	BB	568	A	P-O3'-C3'	-5.58	113.01	119.70
35	BB	573	C	C2-N1-C1'	5.58	124.93	118.80
35	BB	1369	A	O4'-C1'-C2'	5.58	112.62	107.60
41	BH	45	G	C5'-C4'-C3'	-5.58	107.08	116.00
48	BO	96	ASP	CB-CG-OD1	5.58	123.32	118.30
85	AA	474	C	N1-C2-O2	5.58	122.25	118.90
85	AA	836	A	N1-C6-N6	-5.58	115.25	118.60
85	AA	849	A	O4'-C4'-C3'	-5.58	98.42	104.00
85	AA	988	C	C1'-O4'-C4'	-5.58	105.44	109.90
85	AA	1018	G	C5'-C4'-C3'	-5.58	107.08	116.00
85	AA	1454	U	C2-N3-C4	-5.58	123.65	127.00
85	AA	1560	A	C4'-C3'-C2'	-5.58	97.03	102.60
85	AA	1854	U	N3-C2-O2	-5.58	118.30	122.20
85	AA	1996	A	C5'-C4'-C3'	5.58	124.92	116.00
34	BA	579	U	O3'-P-O5'	-5.57	93.41	104.00
34	BA	684	G	N3-C4-C5	-5.57	125.81	128.60
35	BB	32	C	C6-N1-C2	-5.57	118.07	120.30
35	BB	62	C	C4'-C3'-C2'	-5.57	97.03	102.60
36	BC	15	G	C4-N9-C1'	-5.57	119.25	126.50
38	BE	14	C	P-O5'-C5'	5.57	129.82	120.90
39	BF	28	C	C5'-C4'-C3'	-5.57	107.08	116.00
41	BH	16	A	C1'-O4'-C4'	-5.57	105.44	109.90
41	BH	54	U	C6-N1-C1'	5.57	129.00	121.20
41	BH	73	A	N1-C6-N6	5.57	121.94	118.60
50	BQ	40	GLN	N-CA-CB	-5.57	100.57	110.60
7	A6	139	ILE	CA-C-N	-5.57	104.94	117.20
34	BA	917	C	C1'-O4'-C4'	-5.57	105.44	109.90
34	BA	1275	G	C1'-O4'-C4'	-5.57	105.44	109.90
34	BA	1531	G	C3'-C2'-C1'	-5.57	97.04	101.50
35	BB	1405	G	N3-C4-C5	-5.57	125.81	128.60
35	BB	1508	G	P-O5'-C5'	-5.57	111.98	120.90
39	BF	40	U	OP1-P-O3'	5.57	117.46	105.20
41	BH	100	A	C5-N7-C8	-5.57	101.11	103.90
85	AA	1127	G	C8-N9-C1'	5.57	134.24	127.00
85	AA	1912	U	C4'-C3'-C2'	-5.57	97.03	102.60
34	BA	372	U	C1'-O4'-C4'	-5.57	105.44	109.90
34	BA	934	G	O4'-C1'-N9	5.57	112.66	108.20
34	BA	1080	U	O4'-C1'-N1	5.57	112.66	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1810	A	P-O5'-C5'	-5.57	111.99	120.90
35	BB	652	G	N3-C2-N2	5.57	123.80	119.90
35	BB	1220	A	C5-N7-C8	-5.57	101.11	103.90
36	BC	101	U	O5'-C5'-C4'	-5.57	101.12	111.70
36	BC	168	C	C2-N1-C1'	-5.57	112.67	118.80
38	BE	49	A	C5-C6-N6	-5.57	119.24	123.70
41	BH	72	G	C5'-C4'-O4'	-5.57	102.42	109.10
85	AA	169	G	N3-C2-N2	5.57	123.80	119.90
85	AA	268	A	P-O5'-C5'	5.57	129.81	120.90
85	AA	397	G	P-O3'-C3'	-5.57	113.02	119.70
85	AA	470	C	C4'-C3'-C2'	-5.57	97.03	102.60
85	AA	926	C	O4'-C1'-N1	5.57	112.66	108.20
85	AA	1385	C	C6-N1-C2	-5.57	118.07	120.30
85	AA	2007	G	N3-C4-N9	-5.57	122.66	126.00
23	AP	55	ASP	CB-CG-OD1	5.57	123.31	118.30
34	BA	322	U	C3'-C2'-C1'	-5.57	97.05	101.50
34	BA	800	G	N1-C2-N3	5.57	127.24	123.90
34	BA	1067	G	O4'-C1'-C2'	5.57	112.61	107.60
35	BB	805	G	C3'-C2'-C1'	-5.57	97.04	101.50
35	BB	1094	A	C8-N9-C1'	5.57	137.72	127.70
85	AA	161	A	C3'-C2'-C1'	-5.57	97.05	101.50
85	AA	195	C	C5'-C4'-C3'	-5.57	107.09	116.00
85	AA	696	G	C2'-C3'-O3'	5.57	122.61	113.70
85	AA	1503	G	N3-C2-N2	5.57	123.80	119.90
85	AA	1669	G	C5'-C4'-C3'	-5.57	107.09	116.00
2	A1	133	ILE	N-CA-C	-5.57	95.97	111.00
34	BA	195	G	N3-C4-C5	-5.57	125.82	128.60
34	BA	594	G	O4'-C1'-C2'	-5.57	100.23	105.80
34	BA	739	A	N7-C8-N9	5.57	116.58	113.80
34	BA	1068	C	C5-C4-N4	5.57	124.10	120.20
34	BA	1268	C	C5-C4-N4	-5.57	116.30	120.20
34	BA	1320	A	C3'-C2'-C1'	-5.57	97.05	101.50
34	BA	1444	G	C4'-C3'-C2'	-5.57	97.03	102.60
35	BB	444	U	O5'-P-OP1	-5.57	100.69	105.70
35	BB	899	C	C4'-C3'-C2'	-5.57	97.03	102.60
35	BB	1181	A	O4'-C1'-C2'	5.57	112.61	107.60
35	BB	1390	U	C4'-C3'-C2'	-5.57	97.03	102.60
36	BC	86	U	N3-C2-O2	-5.57	118.30	122.20
75	Bp	70	THR	C-N-CA	5.57	135.62	121.70
85	AA	313	A	C4-N9-C1'	-5.57	116.28	126.30
85	AA	657	C	C1'-O4'-C4'	-5.57	105.45	109.90
85	AA	1490	A	P-O5'-C5'	5.57	129.81	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1522	U	N1-C1'-C2'	-5.57	105.88	112.00
85	AA	1881	C	C6-N1-C1'	5.57	127.48	120.80
3	A2	76	VAL	CB-CA-C	5.57	121.97	111.40
16	AH	30	PHE	CA-CB-CG	-5.57	100.54	113.90
34	BA	118	C	C4'-C3'-C2'	-5.57	97.03	102.60
34	BA	423	G	C3'-C2'-C1'	-5.57	97.05	101.50
34	BA	538	G	C3'-C2'-C1'	-5.57	97.05	101.50
34	BA	574	U	C4'-C3'-C2'	5.57	108.17	102.60
34	BA	689	C	O3'-P-O5'	5.57	114.57	104.00
34	BA	1078	U	C1'-O4'-C4'	-5.57	105.45	109.90
34	BA	1650	G	P-O3'-C3'	-5.57	113.02	119.70
35	BB	555	G	C1'-O4'-C4'	-5.57	105.45	109.90
35	BB	571	C	C1'-O4'-C4'	-5.57	105.45	109.90
35	BB	1119	G	C5'-C4'-O4'	5.57	115.78	109.10
35	BB	1227	G	O5'-P-OP2	5.57	117.38	110.70
38	BE	45	G	O5'-C5'-C4'	-5.57	101.13	111.70
40	BG	156	G	N3-C4-N9	-5.57	122.66	126.00
85	AA	153	C	P-O5'-C5'	-5.57	112.00	120.90
85	AA	195	C	C1'-O4'-C4'	-5.57	105.45	109.90
85	AA	485	A	C1'-O4'-C4'	-5.57	105.45	109.90
85	AA	695	A	O3'-P-O5'	-5.57	93.43	104.00
85	AA	1116	G	C6-N1-C2	-5.57	121.76	125.10
85	AA	1221	G	C3'-C2'-C1'	-5.57	97.05	101.50
85	AA	1634	U	N3-C2-O2	-5.57	118.31	122.20
85	AA	1898	C	P-O5'-C5'	-5.57	111.99	120.90
85	AA	1962	U	C6-N1-C2	-5.57	117.66	121.00
85	AA	2016	A	P-O3'-C3'	-5.57	113.02	119.70
85	AA	2044	A	N1-C6-N6	5.57	121.94	118.60
85	AA	2071	U	O5'-C5'-C4'	-5.57	101.12	111.70
34	BA	2	A	C8-N9-C1'	5.56	137.72	127.70
34	BA	621	G	C8-N9-C4	-5.56	104.17	106.40
34	BA	718	U	C1'-O4'-C4'	-5.56	105.45	109.90
34	BA	749	G	C8-N9-C4	5.56	108.62	106.40
34	BA	1101	A	C3'-C2'-C1'	-5.56	97.05	101.50
34	BA	1459	U	C4'-C3'-C2'	-5.56	97.04	102.60
35	BB	505	G	N3-C2-N2	5.56	123.80	119.90
35	BB	1264	U	C1'-O4'-C4'	-5.56	105.45	109.90
47	BN	209	ARG	NE-CZ-NH1	5.56	123.08	120.30
85	AA	634	U	C6-N1-C1'	5.56	128.99	121.20
85	AA	2000	C	O3'-P-O5'	-5.56	93.43	104.00
85	AA	2230	U	C6-N1-C2	-5.56	117.66	121.00
15	AG	121	ARG	CG-CD-NE	-5.56	100.12	111.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	AM	90	ASP	C-N-CD	-5.56	108.36	120.60
24	AQ	77	TYR	CA-CB-CG	-5.56	102.83	113.40
34	BA	206	C	C5'-C4'-C3'	5.56	124.90	116.00
34	BA	719	G	P-O5'-C5'	-5.56	112.00	120.90
34	BA	1282	G	N1-C2-N2	5.56	121.21	116.20
34	BA	1291	A	C8-N9-C4	5.56	108.03	105.80
34	BA	1295	U	C2'-C3'-O3'	5.56	122.60	113.70
34	BA	1363	A	C8-N9-C4	5.56	108.03	105.80
34	BA	1573	C	O4'-C1'-C2'	5.56	112.61	107.60
35	BB	829	C	C2-N1-C1'	-5.56	112.68	118.80
35	BB	1288	G	C3'-C2'-C1'	-5.56	97.05	101.50
36	BC	169	G	C4'-C3'-C2'	-5.56	97.04	102.60
40	BG	99	A	C3'-C2'-C1'	-5.56	97.05	101.50
42	BI	71	MET	CB-CA-C	5.56	121.53	110.40
72	Bm	26	THR	C-N-CA	5.56	135.60	121.70
85	AA	138	C	O4'-C4'-C3'	-5.56	98.44	104.00
85	AA	164	G	C4'-C3'-C2'	-5.56	97.04	102.60
85	AA	174	U	C2'-C3'-O3'	5.56	122.60	113.70
85	AA	289	G	P-O3'-C3'	-5.56	113.03	119.70
85	AA	431	G	N1-C6-O6	5.56	123.24	119.90
85	AA	782	G	C4-N9-C1'	-5.56	119.27	126.50
85	AA	1467	U	C5'-C4'-C3'	-5.56	107.10	116.00
85	AA	1928	A	N7-C8-N9	-5.56	111.02	113.80
85	AA	2027	U	C6-N1-C2	-5.56	117.66	121.00
21	AM	109	ARG	NE-CZ-NH2	-5.56	117.52	120.30
34	BA	94	G	N9-C1'-C2'	-5.56	105.88	112.00
34	BA	1502	G	C1'-O4'-C4'	-5.56	105.45	109.90
35	BB	886	G	C8-N9-C4	-5.56	104.18	106.40
53	BT	121	ARG	NE-CZ-NH1	5.56	123.08	120.30
54	BU	52	MET	CG-SD-CE	-5.56	91.30	100.20
85	AA	161	A	N1-C6-N6	5.56	121.94	118.60
85	AA	830	A	O5'-C5'-C4'	5.56	122.27	111.70
85	AA	848	C	C5'-C4'-C3'	-5.56	107.10	116.00
85	AA	1224	C	C6-N1-C1'	5.56	127.47	120.80
85	AA	1283	C	C5-C4-N4	5.56	124.09	120.20
85	AA	1922	A	P-O5'-C5'	-5.56	112.00	120.90
85	AA	2147	A	C3'-C2'-C1'	-5.56	97.05	101.50
5	A4	68	TYR	CA-CB-CG	-5.56	102.83	113.40
22	AO	134	LYS	CB-CA-C	5.56	121.52	110.40
34	BA	167	U	C3'-C2'-C1'	-5.56	97.05	101.50
34	BA	619	U	O3'-P-O5'	-5.56	93.44	104.00
34	BA	801	U	C6-N1-C1'	5.56	128.98	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1194	G	N9-C1'-C2'	-5.56	105.88	112.00
35	BB	130	G	N3-C2-N2	5.56	123.79	119.90
35	BB	377	A	O5'-C5'-C4'	-5.56	101.14	111.70
35	BB	539	G	O4'-C1'-N9	5.56	112.65	108.20
35	BB	779	C	C5-C6-N1	-5.56	118.22	121.00
35	BB	1036	G	N9-C1'-C2'	-5.56	105.89	112.00
35	BB	1155	U	N1-C2-O2	5.56	126.69	122.80
35	BB	1169	A	P-O5'-C5'	5.56	129.80	120.90
35	BB	1178	A	C8-N9-C4	5.56	108.02	105.80
37	BD	58	G	C3'-C2'-C1'	-5.56	97.05	101.50
38	BE	88	G	O3'-P-O5'	5.56	114.56	104.00
40	BG	159	A	C5'-C4'-C3'	-5.56	107.10	116.00
60	Ba	51	LYS	N-CA-C	5.56	126.01	111.00
62	Bc	111	ARG	NE-CZ-NH2	5.56	123.08	120.30
64	Be	123	ARG	N-CA-CB	5.56	120.61	110.60
81	Bv	186	ALA	CB-CA-C	5.56	118.44	110.10
85	AA	198	U	N1-C2-N3	5.56	118.24	114.90
85	AA	269	G	C5'-C4'-C3'	-5.56	107.10	116.00
85	AA	332	A	N9-C4-C5	-5.56	103.58	105.80
85	AA	1459	C	O5'-C5'-C4'	-5.56	101.14	111.70
85	AA	1844	A	C5-C6-N1	5.56	120.48	117.70
85	AA	1938	G	N9-C1'-C2'	-5.56	105.89	112.00
86	AB	8	U	O4'-C4'-C3'	-5.56	98.44	104.00
1	A0	83	TYR	CA-CB-CG	5.56	123.96	113.40
15	AG	136	PRO	N-CA-C	-5.56	97.65	112.10
34	BA	458	G	C4'-C3'-C2'	-5.56	97.04	102.60
34	BA	970	U	P-O3'-C3'	-5.56	113.03	119.70
34	BA	1176	C	C2-N1-C1'	-5.56	112.69	118.80
35	BB	654	C	C4'-C3'-C2'	5.56	108.16	102.60
35	BB	791	A	O4'-C1'-C2'	-5.56	100.24	105.80
35	BB	1013	U	C3'-C2'-C1'	-5.56	97.06	101.50
35	BB	1306	G	N1-C2-N2	-5.56	111.20	116.20
35	BB	1451	C	C6-N1-C2	-5.56	118.08	120.30
35	BB	1462	G	C3'-C2'-C1'	5.56	105.95	101.50
36	BC	37	U	P-O5'-C5'	-5.56	112.01	120.90
37	BD	53	U	C1'-O4'-C4'	5.56	114.35	109.90
37	BD	73	U	O4'-C1'-C2'	5.56	112.60	107.60
85	AA	999	A	C5'-C4'-O4'	5.56	115.77	109.10
85	AA	1115	G	OP1-P-O3'	5.56	117.43	105.20
85	AA	1258	U	C2-N1-C1'	-5.56	111.03	117.70
35	BB	868	C	C2-N1-C1'	-5.56	112.69	118.80
35	BB	1048	A	O4'-C1'-N9	5.56	112.64	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1089	A	C4'-C3'-C2'	5.56	108.16	102.60
35	BB	1432	U	C5-C6-N1	-5.56	119.92	122.70
38	BE	148	C	C4'-C3'-C2'	-5.56	97.04	102.60
67	Bh	53	TYR	CB-CG-CD1	5.56	124.33	121.00
85	AA	101	C	P-O3'-C3'	5.56	126.37	119.70
85	AA	390	U	O3'-P-O5'	-5.56	93.44	104.00
85	AA	444	U	C1'-O4'-C4'	-5.56	105.45	109.90
85	AA	559	G	C5-C6-O6	-5.56	125.27	128.60
85	AA	638	G	C5'-C4'-C3'	-5.56	107.11	116.00
85	AA	744	C	C6-N1-C1'	-5.56	114.13	120.80
85	AA	917	A	P-O5'-C5'	-5.56	112.01	120.90
85	AA	1090	A	C4'-C3'-C2'	-5.56	97.04	102.60
5	A4	155	MET	CG-SD-CE	-5.55	91.31	100.20
23	AP	225	PHE	CB-CG-CD1	5.55	124.69	120.80
34	BA	2	A	N9-C1'-C2'	-5.55	105.89	112.00
34	BA	1082	U	C4'-C3'-C2'	-5.55	97.05	102.60
34	BA	1527	G	C8-N9-C4	5.55	108.62	106.40
34	BA	1831	A	C3'-C2'-C1'	5.55	105.94	101.50
35	BB	27	C	N1-C2-O2	5.55	122.23	118.90
35	BB	578	G	P-O3'-C3'	-5.55	113.03	119.70
35	BB	1438	U	O4'-C1'-N1	5.55	112.64	108.20
37	BD	11	A	O4'-C1'-N9	5.55	112.64	108.20
40	BG	152	G	C8-N9-C1'	5.55	134.22	127.00
41	BH	74	G	N7-C8-N9	5.55	115.88	113.10
85	AA	188	G	C1'-O4'-C4'	-5.55	105.46	109.90
85	AA	323	U	O5'-C5'-C4'	-5.55	101.14	111.70
85	AA	1549	G	C3'-C2'-C1'	-5.55	97.06	101.50
85	AA	1922	A	C4'-C3'-C2'	-5.55	97.05	102.60
34	BA	205	G	C5'-C4'-C3'	5.55	124.89	116.00
34	BA	217	C	O4'-C1'-N1	5.55	112.64	108.20
34	BA	285	C	C6-N1-C2	-5.55	118.08	120.30
34	BA	595	U	C2-N1-C1'	5.55	124.36	117.70
34	BA	759	A	C4-C5-C6	-5.55	114.22	117.00
34	BA	1006	G	C5-C6-O6	5.55	131.93	128.60
34	BA	1518	A	C6-C5-N7	-5.55	128.41	132.30
35	BB	524	C	O4'-C4'-C3'	-5.55	98.45	104.00
35	BB	1535	G	P-O3'-C3'	5.55	126.36	119.70
85	AA	293	A	C5-C6-N6	-5.55	119.26	123.70
85	AA	1432	C	C1'-O4'-C4'	5.55	114.34	109.90
85	AA	1521	U	C4'-C3'-O3'	-5.55	97.74	109.40
1	A0	49	ASN	CA-CB-CG	-5.55	101.19	113.40
34	BA	699	G	N1-C6-O6	5.55	123.23	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	856	G	N1-C6-O6	-5.55	116.57	119.90
34	BA	1063	G	N3-C2-N2	5.55	123.79	119.90
34	BA	1315	C	C1'-O4'-C4'	-5.55	105.46	109.90
34	BA	1592	U	C5'-C4'-O4'	5.55	115.76	109.10
35	BB	137	A	C4'-C3'-C2'	-5.55	97.05	102.60
35	BB	537	A	C5'-C4'-C3'	-5.55	107.12	116.00
35	BB	576	A	C5-C6-N1	5.55	120.48	117.70
35	BB	1005	A	O5'-C5'-C4'	-5.55	101.15	111.70
35	BB	1294	C	P-O5'-C5'	-5.55	112.02	120.90
38	BE	52	U	C1'-O4'-C4'	-5.55	105.46	109.90
53	BT	80	ARG	NE-CZ-NH2	-5.55	117.52	120.30
79	Bt	62	ALA	N-CA-C	5.55	125.99	111.00
85	AA	585	G	C8-N9-C4	5.55	108.62	106.40
85	AA	612	A	O4'-C1'-N9	5.55	112.64	108.20
85	AA	738	C	C2-N1-C1'	5.55	124.91	118.80
85	AA	889	G	O4'-C1'-N9	5.55	112.64	108.20
85	AA	1148	G	C5-C6-O6	-5.55	125.27	128.60
85	AA	1621	U	O4'-C1'-N1	5.55	112.64	108.20
85	AA	1865	C	C1'-O4'-C4'	-5.55	105.46	109.90
85	AA	1986	G	C5'-C4'-C3'	-5.55	107.12	116.00
85	AA	2064	A	C1'-O4'-C4'	-5.55	105.46	109.90
18	AJ	5	SER	N-CA-CB	5.55	118.82	110.50
34	BA	294	C	N3-C2-O2	-5.55	118.02	121.90
34	BA	366	G	N1-C6-O6	-5.55	116.57	119.90
34	BA	527	C	C5-C4-N4	-5.55	116.31	120.20
34	BA	693	G	C8-N9-C4	-5.55	104.18	106.40
34	BA	702	G	O4'-C1'-C2'	5.55	112.59	107.60
34	BA	815	C	OP1-P-OP2	-5.55	111.27	119.60
34	BA	1021	U	C2-N1-C1'	5.55	124.36	117.70
34	BA	1252	G	C5-C6-N1	5.55	114.27	111.50
34	BA	1337	A	O4'-C1'-N9	5.55	112.64	108.20
34	BA	1569	C	C2-N1-C1'	-5.55	112.70	118.80
34	BA	1604	A	C5-N7-C8	-5.55	101.12	103.90
34	BA	1718	C	C4'-C3'-C2'	5.55	108.15	102.60
35	BB	110	U	O5'-C5'-C4'	-5.55	101.16	111.70
35	BB	576	A	O3'-P-O5'	-5.55	93.45	104.00
35	BB	608	A	C8-N9-C4	5.55	108.02	105.80
35	BB	767	A	O3'-P-O5'	-5.55	93.46	104.00
35	BB	1033	U	O4'-C1'-N1	5.55	112.64	108.20
35	BB	1282	G	O5'-C5'-C4'	-5.55	101.16	111.70
35	BB	1546	C	C6-N1-C1'	5.55	127.46	120.80
36	BC	162	C	C1'-O4'-C4'	-5.55	105.46	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	83	U	OP1-P-OP2	-5.55	111.27	119.60
42	BI	17	HIS	CA-C-N	5.55	129.41	117.20
47	BN	70	TYR	CB-CG-CD2	5.55	124.33	121.00
85	AA	55	A	C1'-O4'-C4'	-5.55	105.46	109.90
85	AA	91	U	C5'-C4'-C3'	5.55	124.88	116.00
85	AA	655	U	P-O5'-C5'	5.55	129.78	120.90
85	AA	1222	A	C6-N1-C2	-5.55	115.27	118.60
85	AA	1591	U	C4'-C3'-C2'	-5.55	97.05	102.60
85	AA	1765	G	O3'-P-O5'	-5.55	93.45	104.00
85	AA	2090	C	C1'-O4'-C4'	-5.55	105.46	109.90
85	AA	2114	U	C1'-O4'-C4'	-5.55	105.46	109.90
34	BA	917	C	N3-C2-O2	-5.55	118.02	121.90
34	BA	968	G	C4-N9-C1'	-5.55	119.29	126.50
35	BB	83	G	N1-C6-O6	5.55	123.23	119.90
35	BB	145	G	C5'-C4'-C3'	5.55	124.88	116.00
35	BB	659	C	O5'-C5'-C4'	-5.55	101.16	111.70
44	BK	55	ARG	NE-CZ-NH1	5.55	123.07	120.30
85	AA	1235	G	C1'-O4'-C4'	-5.55	105.46	109.90
85	AA	1909	C	N3-C2-O2	-5.55	118.02	121.90
34	BA	182	U	O5'-C5'-C4'	-5.55	101.16	111.70
34	BA	483	A	C5'-C4'-O4'	5.55	115.75	109.10
34	BA	504	A	C4-N9-C1'	-5.55	116.31	126.30
34	BA	828	A	C4-N9-C1'	-5.55	116.32	126.30
35	BB	6	A	O4'-C1'-C2'	5.55	112.59	107.60
36	BC	122	A	C5'-C4'-C3'	-5.55	107.13	116.00
37	BD	69	U	C3'-C2'-C1'	-5.55	97.06	101.50
38	BE	121	G	P-O5'-C5'	-5.55	112.02	120.90
52	BS	134	HIS	CA-CB-CG	-5.55	104.17	113.60
56	BW	14	ARG	NE-CZ-NH1	5.55	123.07	120.30
85	AA	576	U	P-O3'-C3'	-5.55	113.04	119.70
85	AA	1855	U	C4'-C3'-O3'	-5.55	97.75	109.40
34	BA	306	G	N3-C4-C5	-5.54	125.83	128.60
34	BA	1419	A	C5'-C4'-O4'	-5.54	102.45	109.10
34	BA	1654	G	C8-N9-C1'	5.54	134.21	127.00
36	BC	149	A	C8-N9-C4	-5.54	103.58	105.80
38	BE	91	G	C3'-C2'-C1'	-5.54	97.06	101.50
40	BG	16	G	C5-N7-C8	5.54	107.07	104.30
65	Bf	233	HIS	C-N-CA	5.54	135.56	121.70
85	AA	329	G	C8-N9-C4	5.54	108.62	106.40
85	AA	849	A	N9-C1'-C2'	-5.54	105.90	112.00
85	AA	1624	U	O4'-C1'-C2'	5.54	112.59	107.60
34	BA	499	C	P-O5'-C5'	5.54	129.77	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1064	A	C5-C6-N1	5.54	120.47	117.70
34	BA	1708	A	C5-C6-N6	-5.54	119.27	123.70
35	BB	401	U	C6-N1-C2	5.54	124.33	121.00
35	BB	788	U	C5-C6-N1	-5.54	119.93	122.70
35	BB	805	G	C5'-C4'-C3'	5.54	124.87	116.00
35	BB	1013	U	O4'-C1'-N1	5.54	112.64	108.20
35	BB	1340	U	O4'-C1'-N1	5.54	112.64	108.20
40	BG	136	G	O4'-C1'-N9	5.54	112.64	108.20
73	Bn	66	TYR	CB-CG-CD1	-5.54	117.67	121.00
77	Br	189	ARG	NE-CZ-NH1	5.54	123.07	120.30
85	AA	657	C	N1-C2-O2	5.54	122.23	118.90
85	AA	703	U	C4'-C3'-C2'	-5.54	97.06	102.60
85	AA	930	G	N9-C1'-C2'	5.54	121.21	114.00
85	AA	1062	U	C5'-C4'-O4'	5.54	115.75	109.10
85	AA	1549	G	C4-N9-C1'	-5.54	119.29	126.50
85	AA	1854	U	C5'-C4'-C3'	-5.54	107.13	116.00
85	AA	2095	U	C4'-C3'-C2'	5.54	108.14	102.60
34	BA	77	C	P-O3'-C3'	-5.54	113.05	119.70
34	BA	1190	A	C8-N9-C1'	5.54	137.67	127.70
34	BA	1306	U	C1'-O4'-C4'	-5.54	105.47	109.90
35	BB	392	G	C8-N9-C1'	5.54	134.21	127.00
35	BB	978	C	C6-N1-C2	-5.54	118.08	120.30
35	BB	1160	U	N1-C2-O2	5.54	126.68	122.80
35	BB	1196	A	C5-N7-C8	-5.54	101.13	103.90
35	BB	1202	G	OP2-P-O3'	5.54	117.39	105.20
35	BB	1297	G	P-O5'-C5'	-5.54	112.03	120.90
35	BB	1347	C	N3-C2-O2	-5.54	118.02	121.90
36	BC	65	G	C8-N9-C4	5.54	108.62	106.40
36	BC	154	A	P-O3'-C3'	-5.54	113.05	119.70
39	BF	32	G	C8-N9-C4	-5.54	104.18	106.40
40	BG	46	G	C5'-C4'-C3'	-5.54	107.13	116.00
40	BG	122	G	C8-N9-C1'	5.54	134.20	127.00
40	BG	139	U	N3-C4-O4	-5.54	115.52	119.40
42	BI	72	ARG	N-CA-CB	-5.54	100.62	110.60
42	BI	74	ARG	NE-CZ-NH2	-5.54	117.53	120.30
77	Br	6	SER	CB-CA-C	5.54	120.63	110.10
85	AA	233	C	P-O5'-C5'	-5.54	112.03	120.90
85	AA	1107	A	P-O3'-C3'	5.54	126.35	119.70
85	AA	2142	A	OP1-P-O3'	5.54	117.39	105.20
34	BA	1785	G	C8-N9-C1'	5.54	134.20	127.00
35	BB	959	C	C4'-C3'-C2'	-5.54	97.06	102.60
40	BG	105	A	O4'-C1'-C2'	5.54	112.59	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1163	G	C5'-C4'-O4'	5.54	115.75	109.10
30	AW	80	PHE	CB-CG-CD1	-5.54	116.92	120.80
34	BA	158	U	C5-C4-O4	5.54	129.22	125.90
34	BA	235	C	C5-C6-N1	5.54	123.77	121.00
34	BA	260	A	C5'-C4'-C3'	-5.54	107.14	116.00
34	BA	301	U	O5'-P-OP1	5.54	117.35	110.70
34	BA	519	G	O4'-C4'-C3'	-5.54	98.46	104.00
34	BA	527	C	C4-C5-C6	-5.54	114.63	117.40
34	BA	593	G	N9-C4-C5	5.54	107.61	105.40
34	BA	1139	G	C3'-C2'-C1'	-5.54	97.07	101.50
34	BA	1697	U	C2-N1-C1'	-5.54	111.05	117.70
34	BA	1832	A	C4'-C3'-C2'	-5.54	97.06	102.60
35	BB	1116	U	C1'-O4'-C4'	-5.54	105.47	109.90
35	BB	1327	U	P-O5'-C5'	-5.54	112.04	120.90
35	BB	1475	U	C3'-C2'-C1'	-5.54	97.07	101.50
36	BC	32	U	C3'-C2'-C1'	-5.54	97.07	101.50
36	BC	127	C	P-O3'-C3'	-5.54	113.05	119.70
57	BX	54	PHE	CB-CG-CD2	-5.54	116.92	120.80
62	Bc	18	PHE	CB-CA-C	5.54	121.48	110.40
85	AA	138	C	P-O3'-C3'	-5.54	113.05	119.70
85	AA	479	C	C1'-O4'-C4'	-5.54	105.47	109.90
85	AA	575	G	C5'-C4'-C3'	-5.54	107.14	116.00
85	AA	981	A	P-O5'-C5'	-5.54	112.04	120.90
85	AA	1130	G	O5'-C5'-C4'	-5.54	101.17	111.70
85	AA	1178	A	C5'-C4'-C3'	5.54	124.86	116.00
85	AA	1359	U	N3-C2-O2	-5.54	118.32	122.20
85	AA	1456	A	C8-N9-C1'	-5.54	117.73	127.70
85	AA	1507	G	C4'-C3'-C2'	-5.54	97.06	102.60
85	AA	1540	A	C5-N7-C8	-5.54	101.13	103.90
85	AA	1580	A	C4-C5-C6	-5.54	114.23	117.00
85	AA	1794	U	O4'-C4'-C3'	5.54	110.53	106.10
12	AD	31	GLY	N-CA-C	5.54	126.94	113.10
20	AL	82	MET	C-N-CA	5.54	135.54	121.70
34	BA	652	C	C3'-C2'-C1'	-5.54	97.07	101.50
34	BA	1418	G	O5'-C5'-C4'	-5.54	101.18	111.70
35	BB	10	C	C6-N1-C1'	5.54	127.44	120.80
35	BB	370	A	C5'-C4'-C3'	-5.54	107.14	116.00
35	BB	660	G	C8-N9-C1'	5.54	134.20	127.00
35	BB	1149	A	P-O3'-C3'	-5.54	113.06	119.70
37	BD	29	C	N3-C4-C5	5.54	124.11	121.90
37	BD	75	G	N9-C1'-C2'	-5.54	105.91	112.00
38	BE	196	C	C1'-O4'-C4'	-5.54	105.47	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	54	G	C3'-C2'-C1'	-5.54	97.07	101.50
84	By	117	LYS	N-CA-CB	-5.54	100.63	110.60
85	AA	1457	C	O4'-C1'-C2'	-5.54	100.26	105.80
85	AA	1608	U	P-O3'-C3'	-5.54	113.06	119.70
34	BA	220	U	C2-N1-C1'	-5.54	111.06	117.70
34	BA	1443	U	C5'-C4'-O4'	-5.54	102.46	109.10
35	BB	163	G	O4'-C1'-N9	5.54	112.63	108.20
35	BB	1295	A	C5'-C4'-C3'	-5.54	107.14	116.00
38	BE	208	G	C5-C6-O6	-5.54	125.28	128.60
85	AA	67	C	OP1-P-OP2	-5.54	111.30	119.60
85	AA	88	G	C5-C6-N1	5.54	114.27	111.50
85	AA	406	U	P-O3'-C3'	-5.54	113.06	119.70
85	AA	483	G	C5'-C4'-C3'	-5.54	107.14	116.00
85	AA	804	A	C8-N9-C4	5.54	108.01	105.80
85	AA	1085	U	C6-N1-C1'	5.54	128.95	121.20
85	AA	1086	U	N3-C2-O2	-5.54	118.33	122.20
85	AA	1861	A	P-O3'-C3'	-5.54	113.06	119.70
86	AB	41	C	C2-N3-C4	5.54	122.67	119.90
86	AB	60	U	OP2-P-O3'	5.54	117.38	105.20
7	A6	4	TYR	N-CA-CB	-5.53	100.64	110.60
34	BA	187	G	N9-C1'-C2'	-5.53	105.91	112.00
34	BA	226	A	O4'-C1'-N9	5.53	112.63	108.20
34	BA	1055	U	O5'-C5'-C4'	-5.53	101.19	111.70
34	BA	1273	U	C2-N3-C4	-5.53	123.68	127.00
34	BA	1297	G	N3-C4-C5	-5.53	125.83	128.60
34	BA	1322	A	C5-C6-N6	-5.53	119.27	123.70
34	BA	1412	G	N1-C2-N2	-5.53	111.22	116.20
34	BA	1822	U	C2-N1-C1'	5.53	124.34	117.70
35	BB	410	A	C8-N9-C4	5.53	108.01	105.80
35	BB	663	G	C5'-C4'-O4'	5.53	115.74	109.10
35	BB	1327	U	N1-C1'-C2'	-5.53	105.91	112.00
39	BF	37	C	O3'-P-O5'	-5.53	93.49	104.00
69	Bj	4	PRO	O-C-N	-5.53	113.84	122.70
71	Bl	126	PHE	CB-CG-CD2	-5.53	116.93	120.80
76	Bq	6	PRO	N-CA-C	5.53	126.49	112.10
85	AA	867	G	N3-C2-N2	5.53	123.77	119.90
85	AA	1544	G	C5'-C4'-C3'	-5.53	107.15	116.00
34	BA	618	G	O4'-C1'-N9	5.53	112.63	108.20
34	BA	721	A	P-O5'-C5'	-5.53	112.05	120.90
34	BA	805	A	OP1-P-OP2	-5.53	111.30	119.60
34	BA	1215	U	C5'-C4'-O4'	5.53	115.74	109.10
35	BB	693	U	C4'-C3'-C2'	-5.53	97.07	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1276	U	C5'-C4'-O4'	-5.53	102.46	109.10
35	BB	1430	G	N3-C2-N2	5.53	123.77	119.90
41	BH	105	U	O5'-C5'-C4'	-5.53	101.19	111.70
65	Bf	303	ARG	CG-CD-NE	-5.53	100.18	111.80
77	Br	291	MET	CG-SD-CE	-5.53	91.35	100.20
85	AA	506	G	C5-C6-O6	-5.53	125.28	128.60
85	AA	1202	G	P-O3'-C3'	-5.53	113.06	119.70
85	AA	2234	C	C2-N1-C1'	-5.53	112.71	118.80
13	AE	58	PHE	CB-CG-CD1	-5.53	116.93	120.80
34	BA	99	G	N3-C4-C5	-5.53	125.83	128.60
34	BA	174	A	C6-N1-C2	-5.53	115.28	118.60
34	BA	383	G	C5-C6-N1	5.53	114.27	111.50
34	BA	543	A	O5'-C5'-C4'	5.53	122.21	111.70
34	BA	627	U	N1-C1'-C2'	-5.53	105.92	112.00
34	BA	717	U	OP2-P-O3'	5.53	117.37	105.20
34	BA	1664	C	O4'-C1'-N1	5.53	112.62	108.20
34	BA	1793	G	N1-C6-O6	5.53	123.22	119.90
35	BB	1060	U	C5'-C4'-C3'	-5.53	107.15	116.00
35	BB	1486	C	O3'-P-O5'	-5.53	93.49	104.00
39	BF	48	G	C5-C6-N1	5.53	114.27	111.50
85	AA	50	C	N3-C2-O2	-5.53	118.03	121.90
85	AA	104	C	P-O5'-C5'	-5.53	112.05	120.90
85	AA	530	A	N1-C6-N6	-5.53	115.28	118.60
85	AA	1150	G	N1-C6-O6	5.53	123.22	119.90
85	AA	1715	C	C5'-C4'-O4'	5.53	115.74	109.10
85	AA	2111	C	C3'-C2'-C1'	-5.53	97.08	101.50
22	AO	106	PRO	C-N-CA	5.53	135.52	121.70
34	BA	1284	G	C3'-C2'-C1'	-5.53	97.08	101.50
35	BB	1473	U	P-O3'-C3'	-5.53	113.06	119.70
36	BC	14	G	O4'-C1'-C2'	5.53	112.58	107.60
48	BO	100	ARG	NE-CZ-NH2	-5.53	117.54	120.30
85	AA	463	G	P-O5'-C5'	-5.53	112.05	120.90
85	AA	1968	A	O4'-C1'-N9	5.53	112.62	108.20
85	AA	2139	G	C2'-C3'-O3'	5.53	122.55	113.70
34	BA	398	G	N1-C2-N3	-5.53	120.58	123.90
34	BA	664	C	C5-C6-N1	5.53	123.76	121.00
34	BA	790	G	N1-C6-O6	5.53	123.22	119.90
34	BA	1569	C	O4'-C1'-N1	5.53	112.62	108.20
34	BA	1816	G	OP1-P-O3'	5.53	117.36	105.20
35	BB	558	U	C2-N3-C4	-5.53	123.68	127.00
35	BB	1528	U	C2-N1-C1'	-5.53	111.07	117.70
37	BD	62	A	N1-C2-N3	-5.53	126.54	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	BD	68	C	C1'-O4'-C4'	-5.53	105.48	109.90
37	BD	106	G	P-O3'-C3'	-5.53	113.07	119.70
64	Be	71	ARG	NE-CZ-NH1	5.53	123.06	120.30
80	Bu	193	SER	C-N-CA	5.53	135.52	121.70
85	AA	176	C	O4'-C1'-C2'	5.53	112.57	107.60
85	AA	206	U	P-O5'-C5'	5.53	129.74	120.90
85	AA	448	G	C8-N9-C1'	5.53	134.19	127.00
85	AA	737	G	C5-C6-O6	-5.53	125.28	128.60
85	AA	967	C	N3-C2-O2	-5.53	118.03	121.90
85	AA	1048	C	C6-N1-C2	-5.53	118.09	120.30
85	AA	1229	G	C5-C6-N1	5.53	114.26	111.50
85	AA	1369	U	C2-N1-C1'	-5.53	111.07	117.70
85	AA	2048	C	C3'-C2'-C1'	-5.53	97.08	101.50
4	A3	152	ARG	CG-CD-NE	-5.53	100.20	111.80
34	BA	281	C	C2'-C3'-O3'	5.53	122.54	113.70
34	BA	306	G	N1-C2-N2	-5.53	111.23	116.20
34	BA	557	U	N3-C2-O2	5.53	126.07	122.20
34	BA	825	G	C6-N1-C2	-5.53	121.78	125.10
34	BA	1125	G	C5-C6-O6	-5.53	125.28	128.60
34	BA	1497	A	O5'-P-OP1	-5.53	100.73	105.70
34	BA	1512	C	C5'-C4'-C3'	5.53	124.84	116.00
35	BB	471	U	C2-N1-C1'	-5.53	111.07	117.70
35	BB	1480	G	N3-C4-C5	-5.53	125.84	128.60
40	BG	11	G	O5'-P-OP1	-5.53	100.73	105.70
41	BH	124	C	P-O3'-C3'	-5.53	113.07	119.70
72	Bm	87	SER	N-CA-C	5.53	125.92	111.00
85	AA	1230	U	P-O5'-C5'	5.53	129.74	120.90
85	AA	1285	C	C6-N1-C2	-5.53	118.09	120.30
85	AA	1555	G	C8-N9-C1'	5.53	134.18	127.00
85	AA	1661	U	P-O5'-C5'	5.53	129.74	120.90
27	AT	121	ASN	N-CA-C	5.52	125.91	111.00
34	BA	492	G	C1'-O4'-C4'	-5.52	105.48	109.90
34	BA	803	U	C6-N1-C1'	5.52	128.93	121.20
34	BA	1141	C	C5'-C4'-C3'	-5.52	107.16	116.00
34	BA	1629	A	P-O3'-C3'	-5.52	113.07	119.70
35	BB	900	C	O4'-C1'-N1	5.52	112.62	108.20
35	BB	1108	G	O4'-C1'-N9	5.52	112.62	108.20
35	BB	1147	G	O5'-C5'-C4'	-5.52	101.20	111.70
35	BB	1166	A	N7-C8-N9	-5.52	111.04	113.80
35	BB	1218	G	N3-C4-N9	5.52	129.31	126.00
36	BC	63	G	C1'-O4'-C4'	-5.52	105.48	109.90
36	BC	71	A	OP1-P-OP2	-5.52	111.31	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	118	U	P-O3'-C3'	-5.52	113.07	119.70
49	BP	151	ALA	N-CA-C	-5.52	96.09	111.00
53	BT	163	ARG	NE-CZ-NH2	-5.52	117.54	120.30
54	BU	159	TYR	CB-CG-CD2	5.52	124.31	121.00
57	BX	68	PRO	C-N-CA	5.52	135.51	121.70
67	Bh	53	TYR	CA-CB-CG	5.52	123.90	113.40
80	Bu	204	ARG	NH1-CZ-NH2	-5.52	113.32	119.40
85	AA	1349	A	C3'-C2'-C1'	-5.52	97.08	101.50
85	AA	1477	A	N1-C6-N6	-5.52	115.28	118.60
34	BA	113	G	O4'-C4'-C3'	-5.52	98.48	104.00
34	BA	499	C	O3'-P-O5'	5.52	114.49	104.00
34	BA	603	U	C3'-C2'-C1'	5.52	105.92	101.50
34	BA	629	G	C5'-C4'-C3'	-5.52	107.17	116.00
34	BA	840	U	C5'-C4'-O4'	5.52	115.73	109.10
34	BA	934	G	C5-C6-N1	5.52	114.26	111.50
34	BA	1251	A	P-O5'-C5'	5.52	129.74	120.90
34	BA	1438	C	O3'-P-O5'	-5.52	93.51	104.00
34	BA	1535	G	P-O5'-C5'	-5.52	112.06	120.90
34	BA	1676	A	C5'-C4'-C3'	5.52	124.83	116.00
35	BB	909	U	C5'-C4'-C3'	5.52	124.84	116.00
35	BB	1147	G	C4-N9-C1'	-5.52	119.32	126.50
35	BB	1165	A	P-O3'-C3'	5.52	126.33	119.70
36	BC	7	U	C4-C5-C6	-5.52	116.39	119.70
38	BE	64	A	C1'-O4'-C4'	-5.52	105.48	109.90
41	BH	132	C	C6-N1-C2	-5.52	118.09	120.30
42	BI	189	ALA	N-CA-C	5.52	125.91	111.00
64	Be	80	GLU	C-N-CA	5.52	133.90	122.30
84	By	118	ARG	NE-CZ-NH1	5.52	123.06	120.30
85	AA	176	C	N3-C2-O2	-5.52	118.03	121.90
85	AA	348	G	P-O5'-C5'	5.52	129.74	120.90
85	AA	590	U	C5'-C4'-C3'	5.52	124.84	116.00
85	AA	1048	C	C2-N1-C1'	-5.52	112.72	118.80
85	AA	2030	U	C5'-C4'-O4'	5.52	115.73	109.10
85	AA	2170	G	N3-C2-N2	5.52	123.77	119.90
5	A4	90	ARG	NE-CZ-NH1	5.52	123.06	120.30
31	AX	63	ARG	C-N-CA	5.52	135.50	121.70
34	BA	282	A	C4'-C3'-C2'	-5.52	97.08	102.60
34	BA	584	A	O4'-C1'-N9	5.52	112.62	108.20
34	BA	1580	U	O3'-P-O5'	5.52	114.49	104.00
79	Bt	50	TYR	CB-CA-C	-5.52	99.36	110.40
85	AA	1448	A	C3'-C2'-C1'	5.52	105.92	101.50
8	A7	134	ARG	NE-CZ-NH2	-5.52	117.54	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	AC	46	MET	CG-SD-CE	-5.52	91.37	100.20
34	BA	290	G	C5-C6-O6	-5.52	125.29	128.60
34	BA	398	G	N3-C4-N9	-5.52	122.69	126.00
34	BA	553	A	C5-C6-N6	-5.52	119.28	123.70
34	BA	702	G	C5-C6-N1	5.52	114.26	111.50
34	BA	823	G	C1'-O4'-C4'	-5.52	105.48	109.90
34	BA	939	C	N1-C2-O2	5.52	122.21	118.90
34	BA	1445	U	O5'-P-OP2	-5.52	100.73	105.70
34	BA	1496	G	C4-C5-C6	-5.52	115.49	118.80
35	BB	27	C	N1-C1'-C2'	-5.52	105.93	112.00
35	BB	1245	A	C5-C6-N6	5.52	128.12	123.70
35	BB	1458	U	OP1-P-OP2	-5.52	111.32	119.60
36	BC	76	C	O3'-P-O5'	5.52	114.49	104.00
36	BC	149	A	C5'-C4'-O4'	5.52	115.72	109.10
38	BE	13	A	C6-N1-C2	-5.52	115.29	118.60
38	BE	38	C	C1'-O4'-C4'	-5.52	105.48	109.90
40	BG	89	A	C1'-O4'-C4'	-5.52	105.48	109.90
85	AA	104	C	N3-C2-O2	-5.52	118.04	121.90
85	AA	192	G	N3-C2-N2	5.52	123.76	119.90
85	AA	271	A	N9-C1'-C2'	-5.52	105.93	112.00
85	AA	1908	A	C5-C6-N1	5.52	120.46	117.70
34	BA	687	G	N3-C2-N2	5.52	123.76	119.90
34	BA	1542	A	C6-N1-C2	-5.52	115.29	118.60
35	BB	392	G	N7-C8-N9	-5.52	110.34	113.10
35	BB	550	G	C8-N9-C1'	5.52	134.17	127.00
35	BB	1251	G	O5'-P-OP2	-5.52	100.73	105.70
35	BB	1289	G	O4'-C1'-N9	5.52	112.61	108.20
35	BB	1424	G	P-O5'-C5'	-5.52	112.07	120.90
37	BD	34	C	C5'-C4'-C3'	-5.52	107.17	116.00
38	BE	187	G	O4'-C1'-N9	5.52	112.61	108.20
40	BG	58	G	O3'-P-O5'	-5.52	93.52	104.00
80	Bu	269	ARG	NE-CZ-NH2	-5.52	117.54	120.30
85	AA	12	U	O5'-P-OP2	-5.52	100.73	105.70
85	AA	87	C	N3-C2-O2	-5.52	118.04	121.90
85	AA	651	G	C4-N9-C1'	-5.52	119.33	126.50
85	AA	1054	U	O4'-C1'-N1	5.52	112.61	108.20
85	AA	1214	C	OP1-P-OP2	-5.52	111.32	119.60
85	AA	1452	C	P-O3'-C3'	5.52	126.32	119.70
85	AA	1736	U	O5'-P-OP2	-5.52	100.73	105.70
34	BA	83	G	N3-C2-N2	5.52	123.76	119.90
34	BA	350	C	O4'-C1'-N1	5.52	112.61	108.20
34	BA	1278	A	C4-N9-C1'	-5.52	116.37	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1807	G	C8-N9-C4	-5.52	104.19	106.40
35	BB	52	G	C2'-C3'-O3'	5.52	122.53	113.70
35	BB	62	C	N3-C4-N4	5.52	121.86	118.00
35	BB	381	C	C2'-C3'-O3'	5.52	122.53	113.70
35	BB	1308	G	N1-C6-O6	5.52	123.21	119.90
36	BC	4	G	P-O5'-C5'	-5.52	112.08	120.90
36	BC	106	G	N3-C2-N2	5.52	123.76	119.90
53	BT	73	GLY	N-CA-C	-5.52	99.31	113.10
85	AA	476	C	P-O3'-C3'	-5.52	113.08	119.70
85	AA	584	G	C4-C5-N7	5.52	113.01	110.80
85	AA	709	A	O4'-C4'-C3'	-5.52	98.48	104.00
85	AA	2039	G	C4-C5-C6	-5.52	115.49	118.80
17	AI	107	ARG	NE-CZ-NH1	5.51	123.06	120.30
34	BA	58	A	N1-C2-N3	-5.51	126.54	129.30
34	BA	650	C	P-O3'-C3'	5.51	126.32	119.70
34	BA	664	C	N3-C2-O2	-5.51	118.04	121.90
34	BA	951	C	O4'-C4'-C3'	-5.51	98.49	104.00
34	BA	1250	C	O4'-C1'-C2'	5.51	112.56	107.60
34	BA	1511	C	C5'-C4'-C3'	-5.51	107.18	116.00
35	BB	813	C	N1-C1'-C2'	-5.51	105.94	112.00
36	BC	59	A	N1-C2-N3	-5.51	126.54	129.30
36	BC	132	U	O4'-C1'-N1	5.51	112.61	108.20
40	BG	21	C	C5'-C4'-O4'	5.51	115.72	109.10
42	BI	11	LYS	C-N-CA	5.51	135.49	121.70
85	AA	2	A	O4'-C1'-N9	5.51	112.61	108.20
85	AA	77	C	C1'-O4'-C4'	-5.51	105.49	109.90
85	AA	618	A	O4'-C4'-C3'	-5.51	98.48	104.00
85	AA	625	G	N1-C6-O6	5.51	123.21	119.90
85	AA	969	U	O5'-C5'-C4'	5.51	122.18	111.70
85	AA	1807	A	P-O3'-C3'	5.51	126.32	119.70
85	AA	1818	C	N3-C4-C5	-5.51	119.69	121.90
34	BA	467	A	O3'-P-O5'	-5.51	93.53	104.00
35	BB	827	U	C2-N1-C1'	-5.51	111.08	117.70
35	BB	1023	G	P-O5'-C5'	5.51	129.72	120.90
35	BB	1291	G	C5-C6-N1	5.51	114.26	111.50
39	BF	9	C	C6-N1-C1'	-5.51	114.18	120.80
39	BF	18	U	C3'-C2'-C1'	5.51	105.91	101.50
70	Bk	60	THR	CA-CB-CG2	5.51	120.12	112.40
77	Br	286	ASP	CB-CA-C	5.51	121.43	110.40
85	AA	277	G	P-O3'-C3'	-5.51	113.08	119.70
85	AA	324	U	P-O3'-C3'	-5.51	113.08	119.70
85	AA	838	G	P-O5'-C5'	-5.51	112.08	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1110	A	C1'-O4'-C4'	-5.51	105.49	109.90
85	AA	1203	G	C1'-O4'-C4'	-5.51	105.49	109.90
6	A5	18	THR	CA-CB-CG2	-5.51	104.69	112.40
34	BA	86	A	O5'-P-OP1	5.51	117.31	110.70
34	BA	384	U	C1'-O4'-C4'	-5.51	105.49	109.90
34	BA	1095	G	P-O3'-C3'	-5.51	113.08	119.70
34	BA	1481	U	P-O3'-C3'	-5.51	113.09	119.70
34	BA	1619	U	O3'-P-O5'	-5.51	93.53	104.00
35	BB	10	C	C3'-C2'-C1'	-5.51	97.09	101.50
35	BB	659	C	O4'-C4'-C3'	5.51	110.51	106.10
35	BB	1001	G	N9-C1'-C2'	-5.51	105.94	112.00
35	BB	1392	A	OP2-P-O3'	5.51	117.32	105.20
35	BB	1521	G	C4-N9-C1'	-5.51	119.33	126.50
39	BF	7	G	N3-C4-N9	5.51	129.31	126.00
51	BR	135	ARG	NE-CZ-NH1	5.51	123.06	120.30
85	AA	648	G	C5'-C4'-C3'	-5.51	107.18	116.00
85	AA	735	G	N7-C8-N9	5.51	115.86	113.10
85	AA	1549	G	C1'-O4'-C4'	-5.51	105.49	109.90
85	AA	1891	U	N1-C2-N3	5.51	118.21	114.90
85	AA	2227	A	N1-C2-N3	-5.51	126.54	129.30
15	AG	71	ILE	CB-CA-C	-5.51	100.58	111.60
22	AO	153	PHE	CB-CG-CD2	-5.51	116.94	120.80
34	BA	656	U	C6-N1-C2	-5.51	117.69	121.00
34	BA	864	G	OP1-P-O3'	5.51	117.32	105.20
34	BA	1064	A	C3'-C2'-C1'	-5.51	97.09	101.50
34	BA	1121	U	OP2-P-O3'	5.51	117.32	105.20
34	BA	1463	U	C6-N1-C1'	5.51	128.91	121.20
35	BB	72	G	N1-C6-O6	-5.51	116.59	119.90
35	BB	88	U	C4'-C3'-C2'	-5.51	97.09	102.60
35	BB	781	U	O5'-P-OP1	-5.51	100.74	105.70
35	BB	1215	U	C5'-C4'-C3'	-5.51	107.19	116.00
35	BB	1374	U	N3-C2-O2	-5.51	118.34	122.20
35	BB	1480	G	C2-N3-C4	-5.51	109.15	111.90
39	BF	52	A	P-O5'-C5'	5.51	129.72	120.90
65	Bf	377	ASN	CB-CA-C	-5.51	99.38	110.40
83	Bx	265	ARG	NE-CZ-NH2	-5.51	117.55	120.30
85	AA	152	A	C6-C5-N7	-5.51	128.44	132.30
85	AA	154	U	C1'-O4'-C4'	-5.51	105.49	109.90
85	AA	351	C	C1'-O4'-C4'	-5.51	105.49	109.90
85	AA	433	U	C4'-C3'-C2'	-5.51	97.09	102.60
85	AA	441	C	P-O3'-C3'	5.51	126.31	119.70
85	AA	1355	U	C2-N1-C1'	-5.51	111.09	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2219	G	C5'-C4'-O4'	5.51	115.71	109.10
85	AA	2237	G	N3-C2-N2	5.51	123.76	119.90
34	BA	513	U	O3'-P-O5'	-5.51	93.53	104.00
34	BA	770	G	N3-C4-C5	-5.51	125.85	128.60
34	BA	877	U	N3-C2-O2	-5.51	118.34	122.20
34	BA	1816	G	C8-N9-C4	5.51	108.60	106.40
35	BB	1053	G	N9-C1'-C2'	-5.51	105.94	112.00
35	BB	1205	A	C5-C6-N6	-5.51	119.29	123.70
39	BF	16	C	OP2-P-O3'	5.51	117.32	105.20
85	AA	754	C	C4'-C3'-C2'	5.51	108.11	102.60
85	AA	1539	A	C5-C6-N6	-5.51	119.29	123.70
85	AA	1612	C	O3'-P-O5'	-5.51	93.53	104.00
85	AA	1762	G	O4'-C1'-N9	5.51	112.61	108.20
85	AA	2135	A	C4-N9-C1'	-5.51	116.39	126.30
7	A6	68	ARG	N-CA-CB	-5.51	100.69	110.60
34	BA	7	U	O5'-P-OP1	5.51	117.31	110.70
34	BA	160	G	OP1-P-OP2	-5.51	111.34	119.60
34	BA	983	A	N9-C1'-C2'	-5.51	105.94	112.00
34	BA	1321	A	N9-C1'-C2'	-5.51	105.94	112.00
35	BB	268	G	P-O5'-C5'	5.51	129.71	120.90
35	BB	364	U	O4'-C1'-N1	5.51	112.61	108.20
35	BB	376	A	P-O5'-C5'	-5.51	112.09	120.90
35	BB	668	A	C4'-C3'-C2'	-5.51	97.09	102.60
35	BB	738	G	C4-N9-C1'	-5.51	119.34	126.50
35	BB	1039	A	C2'-C3'-O3'	5.51	122.51	113.70
35	BB	1077	C	P-O5'-C5'	-5.51	112.09	120.90
35	BB	1464	G	C8-N9-C4	-5.51	104.20	106.40
36	BC	100	U	O4'-C1'-N1	5.51	112.61	108.20
37	BD	53	U	C2-N1-C1'	-5.51	111.09	117.70
39	BF	11	C	OP1-P-OP2	-5.51	111.34	119.60
50	BQ	146	TRP	CA-CB-CG	-5.51	103.24	113.70
84	By	45	ARG	CG-CD-NE	-5.51	100.23	111.80
85	AA	506	G	C8-N9-C4	5.51	108.60	106.40
85	AA	2172	A	C4-N9-C1'	5.51	136.21	126.30
26	AS	61	GLN	N-CA-C	-5.50	96.14	111.00
34	BA	92	G	C3'-C2'-C1'	-5.50	97.10	101.50
34	BA	290	G	O4'-C1'-N9	5.50	112.60	108.20
34	BA	739	A	P-O5'-C5'	-5.50	112.09	120.90
34	BA	759	A	P-O3'-C3'	-5.50	113.09	119.70
34	BA	811	C	O4'-C1'-N1	5.50	112.60	108.20
36	BC	104	A	C5'-C4'-O4'	5.50	115.71	109.10
36	BC	149	A	C5'-C4'-C3'	-5.50	107.19	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	151	C	C2'-C3'-O3'	5.50	122.51	113.70
49	BP	76	GLU	N-CA-CB	-5.50	100.69	110.60
58	BY	5	ASP	CA-CB-CG	-5.50	101.29	113.40
85	AA	250	C	C1'-O4'-C4'	-5.50	105.50	109.90
85	AA	402	G	O3'-P-O5'	5.50	114.46	104.00
85	AA	636	G	C1'-O4'-C4'	-5.50	105.50	109.90
85	AA	1474	U	O4'-C1'-C2'	-5.50	100.30	105.80
85	AA	1549	G	P-O3'-C3'	-5.50	113.09	119.70
34	BA	193	C	N1-C1'-C2'	-5.50	105.95	112.00
34	BA	243	C	C3'-C2'-C1'	-5.50	97.10	101.50
34	BA	587	U	O3'-P-O5'	-5.50	93.54	104.00
34	BA	676	G	C8-N9-C4	-5.50	104.20	106.40
34	BA	762	A	P-O5'-C5'	5.50	129.71	120.90
34	BA	1221	A	N7-C8-N9	-5.50	111.05	113.80
34	BA	1498	A	N9-C1'-C2'	-5.50	105.95	112.00
34	BA	1545	C	O5'-C5'-C4'	-5.50	101.25	111.70
34	BA	1585	A	N1-C6-N6	-5.50	115.30	118.60
35	BB	603	U	N1-C1'-C2'	-5.50	105.95	112.00
35	BB	686	A	C4'-C3'-C2'	5.50	108.10	102.60
35	BB	770	G	C8-N9-C1'	5.50	134.15	127.00
35	BB	1070	G	C4-N9-C1'	-5.50	119.35	126.50
85	AA	4	C	C3'-C2'-C1'	-5.50	97.10	101.50
85	AA	1356	U	C5-C4-O4	-5.50	122.60	125.90
85	AA	2079	U	O4'-C1'-N1	5.50	112.60	108.20
85	AA	2204	A	O4'-C1'-N9	5.50	112.60	108.20
34	BA	72	U	C4'-C3'-C2'	5.50	108.10	102.60
34	BA	691	A	N9-C1'-C2'	-5.50	105.95	112.00
34	BA	811	C	C3'-C2'-C1'	-5.50	97.10	101.50
34	BA	902	C	C5'-C4'-C3'	-5.50	107.20	116.00
34	BA	1295	U	OP2-P-O3'	5.50	117.30	105.20
35	BB	12	G	N1-C6-O6	-5.50	116.60	119.90
35	BB	50	A	N7-C8-N9	-5.50	111.05	113.80
35	BB	156	G	C5-C6-O6	-5.50	125.30	128.60
35	BB	471	U	C6-N1-C2	-5.50	117.70	121.00
35	BB	484	G	O5'-C5'-C4'	5.50	122.15	111.70
35	BB	845	C	C6-N1-C2	-5.50	118.10	120.30
35	BB	878	G	C4'-C3'-C2'	5.50	108.10	102.60
35	BB	1049	G	C4-N9-C1'	-5.50	119.35	126.50
35	BB	1200	A	N9-C1'-C2'	-5.50	105.95	112.00
68	Bi	106	ARG	CD-NE-CZ	-5.50	115.90	123.60
85	AA	16	G	N7-C8-N9	-5.50	110.35	113.10
85	AA	364	C	C5'-C4'-C3'	-5.50	107.20	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1195	U	O4'-C1'-N1	5.50	112.60	108.20
85	AA	1337	A	C1'-O4'-C4'	-5.50	105.50	109.90
85	AA	1433	C	O4'-C1'-N1	5.50	112.60	108.20
85	AA	1802	U	C2-N3-C4	-5.50	123.70	127.00
85	AA	2034	G	N1-C6-O6	5.50	123.20	119.90
85	AA	2149	C	C5'-C4'-O4'	5.50	115.70	109.10
35	BB	688	U	C2'-C3'-O3'	5.50	122.50	113.70
35	BB	1283	C	C1'-O4'-C4'	-5.50	105.50	109.90
85	AA	863	C	C3'-C2'-C1'	-5.50	97.10	101.50
85	AA	1115	G	N1-C2-N3	5.50	127.20	123.90
86	AB	20	U	N1-C1'-C2'	-5.50	105.95	112.00
34	BA	520	G	C8-N9-C1'	5.50	134.15	127.00
34	BA	996	U	O4'-C1'-C2'	5.50	112.55	107.60
35	BB	520	G	N3-C2-N2	5.50	123.75	119.90
35	BB	802	G	C5'-C4'-O4'	5.50	115.70	109.10
35	BB	1146	C	C5'-C4'-O4'	5.50	115.70	109.10
35	BB	1371	G	O4'-C1'-C2'	5.50	112.55	107.60
35	BB	1393	C	C2-N1-C1'	5.50	124.85	118.80
35	BB	1395	G	N9-C1'-C2'	-5.50	105.95	112.00
35	BB	1462	G	C4'-C3'-C2'	-5.50	97.10	102.60
36	BC	8	C	O5'-C5'-C4'	-5.50	101.25	111.70
36	BC	68	A	C1'-O4'-C4'	-5.50	105.50	109.90
36	BC	102	G	C1'-O4'-C4'	-5.50	105.50	109.90
38	BE	76	U	C5'-C4'-C3'	-5.50	107.20	116.00
42	BI	56	ARG	NE-CZ-NH2	-5.50	117.55	120.30
85	AA	447	C	N3-C4-N4	-5.50	114.15	118.00
85	AA	644	A	C5-C6-N6	-5.50	119.30	123.70
85	AA	676	U	C4'-C3'-C2'	5.50	108.10	102.60
85	AA	760	U	C6-N1-C1'	-5.50	113.50	121.20
85	AA	779	G	P-O5'-C5'	5.50	129.70	120.90
85	AA	929	G	C1'-O4'-C4'	5.50	114.30	109.90
85	AA	1082	U	C6-N1-C1'	-5.50	113.50	121.20
85	AA	1240	A	C8-N9-C4	5.50	108.00	105.80
85	AA	1558	U	N3-C4-O4	-5.50	115.55	119.40
86	AB	70	G	C5'-C4'-C3'	-5.50	107.20	116.00
7	A6	138	GLN	CB-CA-C	5.50	121.39	110.40
34	BA	49	A	C3'-C2'-C1'	-5.50	97.10	101.50
34	BA	229	C	O4'-C1'-N1	5.50	112.60	108.20
34	BA	476	U	P-O5'-C5'	-5.50	112.11	120.90
34	BA	800	G	N1-C2-N2	-5.50	111.25	116.20
34	BA	819	G	C1'-O4'-C4'	-5.50	105.50	109.90
34	BA	1299	G	N1-C2-N2	5.50	121.15	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1657	A	N9-C1'-C2'	-5.50	105.95	112.00
35	BB	803	U	O4'-C1'-N1	5.50	112.60	108.20
35	BB	838	G	O5'-P-OP2	5.50	117.30	110.70
35	BB	971	A	C5-C6-N6	-5.50	119.30	123.70
35	BB	1267	C	C5'-C4'-O4'	5.50	115.69	109.10
35	BB	1370	G	C4-N9-C1'	5.50	133.65	126.50
38	BE	34	C	N1-C2-N3	5.50	123.05	119.20
38	BE	153	C	N3-C4-N4	5.50	121.85	118.00
41	BH	10	U	C1'-O4'-C4'	-5.50	105.50	109.90
41	BH	104	U	N3-C4-C5	5.50	117.90	114.60
53	BT	166	ASP	N-CA-CB	-5.50	100.71	110.60
85	AA	670	C	O4'-C1'-N1	5.50	112.60	108.20
85	AA	1108	U	O4'-C1'-N1	5.50	112.60	108.20
85	AA	1484	G	N3-C2-N2	5.50	123.75	119.90
85	AA	1683	U	O3'-P-O5'	-5.50	93.56	104.00
85	AA	1982	C	C3'-C2'-C1'	5.50	105.90	101.50
86	AB	66	U	O5'-C5'-C4'	-5.50	101.26	111.70
34	BA	55	G	C8-N9-C4	-5.50	104.20	106.40
34	BA	174	A	C1'-O4'-C4'	-5.50	105.50	109.90
34	BA	516	U	N3-C4-C5	5.50	117.90	114.60
34	BA	840	U	P-O3'-C3'	-5.50	113.11	119.70
34	BA	1557	G	O5'-C5'-C4'	-5.50	101.26	111.70
35	BB	475	A	C4-C5-C6	-5.50	114.25	117.00
35	BB	1070	G	C8-N9-C1'	5.50	134.14	127.00
53	BT	97	ARG	NE-CZ-NH2	-5.50	117.55	120.30
85	AA	11	A	N1-C6-N6	-5.50	115.30	118.60
85	AA	416	U	N3-C4-C5	-5.50	111.30	114.60
85	AA	657	C	C5'-C4'-O4'	5.50	115.69	109.10
26	AS	134	TYR	CB-CA-C	5.49	121.39	110.40
34	BA	531	C	C5'-C4'-O4'	5.49	115.69	109.10
34	BA	941	G	C5'-C4'-O4'	5.49	115.69	109.10
34	BA	969	A	C4'-C3'-C2'	-5.49	97.11	102.60
34	BA	1031	U	O4'-C1'-N1	5.49	112.59	108.20
34	BA	1144	A	C8-N9-C1'	5.49	137.59	127.70
34	BA	1821	A	O5'-C5'-C4'	-5.49	101.26	111.70
35	BB	99	G	O5'-C5'-C4'	-5.49	101.26	111.70
35	BB	161	G	C5-C6-O6	-5.49	125.30	128.60
35	BB	381	C	C2-N1-C1'	5.49	124.84	118.80
35	BB	628	A	C1'-O4'-C4'	-5.49	105.50	109.90
35	BB	661	G	C5-C6-O6	5.49	131.90	128.60
35	BB	661	G	N1-C6-O6	-5.49	116.60	119.90
35	BB	808	U	C2-N1-C1'	-5.49	111.11	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	895	U	N3-C2-O2	-5.49	118.35	122.20
35	BB	1147	G	N3-C2-N2	-5.49	116.06	119.90
35	BB	1518	U	C6-N1-C2	-5.49	117.70	121.00
37	BD	118	C	C2-N3-C4	-5.49	117.15	119.90
38	BE	162	U	C2-N1-C1'	-5.49	111.11	117.70
39	BF	4	A	C2-N3-C4	-5.49	107.85	110.60
40	BG	2	U	N1-C2-O2	-5.49	118.95	122.80
49	BP	61	PHE	N-CA-C	-5.49	96.17	111.00
85	AA	23	G	O4'-C1'-N9	5.49	112.59	108.20
85	AA	241	U	P-O5'-C5'	5.49	129.69	120.90
85	AA	346	U	O4'-C1'-N1	5.49	112.59	108.20
85	AA	475	A	C4'-C3'-C2'	5.49	108.09	102.60
85	AA	830	A	C8-N9-C4	5.49	108.00	105.80
85	AA	913	U	N3-C2-O2	5.49	126.05	122.20
85	AA	1685	G	C5-C6-O6	-5.49	125.30	128.60
85	AA	1730	C	OP1-P-O3'	5.49	117.29	105.20
1	A0	210	ARG	CG-CD-NE	-5.49	100.27	111.80
34	BA	684	G	C6-C5-N7	-5.49	127.11	130.40
34	BA	1785	G	C5'-C4'-C3'	5.49	124.79	116.00
35	BB	856	U	P-O3'-C3'	-5.49	113.11	119.70
36	BC	145	G	P-O3'-C3'	-5.49	113.11	119.70
84	By	184	ASN	CA-CB-CG	-5.49	101.32	113.40
85	AA	91	U	P-O5'-C5'	-5.49	112.11	120.90
85	AA	1509	A	C3'-C2'-C1'	-5.49	97.11	101.50
85	AA	2184	A	C8-N9-C4	5.49	108.00	105.80
34	BA	651	U	O4'-C4'-C3'	-5.49	98.51	104.00
34	BA	1218	G	C5-C6-N1	5.49	114.25	111.50
34	BA	1523	U	C6-N1-C2	-5.49	117.71	121.00
35	BB	317	C	O4'-C1'-N1	5.49	112.59	108.20
35	BB	506	G	C1'-O4'-C4'	-5.49	105.51	109.90
35	BB	558	U	C3'-C2'-C1'	-5.49	97.11	101.50
35	BB	807	U	C2'-C3'-O3'	5.49	122.48	113.70
35	BB	1493	A	P-O5'-C5'	-5.49	112.12	120.90
36	BC	60	U	OP2-P-O3'	5.49	117.28	105.20
38	BE	193	A	P-O3'-C3'	-5.49	113.11	119.70
41	BH	31	A	P-O5'-C5'	-5.49	112.12	120.90
53	BT	170	ARG	N-CA-CB	-5.49	100.72	110.60
85	AA	54	C	N3-C2-O2	-5.49	118.06	121.90
85	AA	500	C	P-O5'-C5'	-5.49	112.11	120.90
85	AA	809	A	P-O3'-C3'	5.49	126.29	119.70
85	AA	891	G	O5'-C5'-C4'	5.49	122.13	111.70
85	AA	1721	A	C2'-C3'-O3'	5.49	122.48	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1904	C	N3-C4-C5	-5.49	119.70	121.90
85	AA	1914	U	O4'-C1'-N1	5.49	112.59	108.20
86	AB	55	U	C2-N1-C1'	5.49	124.29	117.70
34	BA	108	A	C6-N1-C2	-5.49	115.31	118.60
34	BA	934	G	C3'-C2'-C1'	-5.49	97.11	101.50
34	BA	960	C	C2-N1-C1'	-5.49	112.76	118.80
34	BA	993	C	C6-N1-C1'	-5.49	114.21	120.80
34	BA	1414	C	N3-C2-O2	-5.49	118.06	121.90
34	BA	1457	C	P-O5'-C5'	5.49	129.68	120.90
35	BB	636	G	C3'-C2'-C1'	-5.49	97.11	101.50
35	BB	1019	C	N3-C2-O2	-5.49	118.06	121.90
35	BB	1510	G	C4-N9-C1'	-5.49	119.36	126.50
40	BG	124	A	N1-C6-N6	-5.49	115.31	118.60
40	BG	130	G	N9-C1'-C2'	-5.49	105.96	112.00
64	Be	72	VAL	CB-CA-C	5.49	121.83	111.40
80	Bu	13	TYR	CB-CG-CD2	5.49	124.29	121.00
85	AA	244	G	O4'-C1'-N9	5.49	112.59	108.20
85	AA	531	G	O4'-C1'-N9	5.49	112.59	108.20
85	AA	549	A	O4'-C1'-N9	5.49	112.59	108.20
85	AA	854	A	O5'-P-OP1	5.49	117.28	110.70
85	AA	965	G	C8-N9-C1'	5.49	134.13	127.00
85	AA	1462	A	O5'-P-OP2	5.49	117.28	110.70
85	AA	1679	U	P-O3'-C3'	5.49	126.29	119.70
85	AA	1795	C	C1'-O4'-C4'	-5.49	105.51	109.90
85	AA	1829	C	N3-C4-C5	5.49	124.09	121.90
85	AA	1996	A	C5-C6-N6	5.49	128.09	123.70
85	AA	2121	G	N7-C8-N9	5.49	115.84	113.10
34	BA	11	U	O5'-C5'-C4'	-5.49	101.27	111.70
34	BA	1095	G	O5'-C5'-C4'	-5.49	101.28	111.70
34	BA	1108	U	P-O3'-C3'	-5.49	113.11	119.70
34	BA	1475	G	C4'-C3'-C2'	-5.49	97.11	102.60
35	BB	778	A	C5-C6-N6	-5.49	119.31	123.70
35	BB	953	G	O4'-C1'-N9	5.49	112.59	108.20
35	BB	1062	G	C6-N1-C2	-5.49	121.81	125.10
44	BK	49	CYS	CA-CB-SG	-5.49	104.12	114.00
52	BS	83	TYR	CB-CG-CD1	-5.49	117.71	121.00
85	AA	634	U	C5'-C4'-C3'	-5.49	107.22	116.00
85	AA	1202	G	C4'-C3'-C2'	-5.49	97.11	102.60
85	AA	1680	U	N3-C2-O2	-5.49	118.36	122.20
4	A3	152	ARG	C-N-CA	5.49	135.41	121.70
7	A6	106	ARG	NE-CZ-NH1	5.49	123.04	120.30
34	BA	83	G	C5-C6-O6	-5.49	125.31	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	472	G	C5'-C4'-O4'	5.49	115.68	109.10
34	BA	711	C	C5'-C4'-C3'	5.49	124.78	116.00
34	BA	875	G	C4'-C3'-C2'	5.49	108.08	102.60
34	BA	1057	C	C3'-C2'-C1'	5.49	105.89	101.50
34	BA	1234	U	C6-N1-C1'	5.49	128.88	121.20
34	BA	1443	U	O4'-C4'-C3'	5.49	110.49	106.10
34	BA	1722	U	N3-C4-O4	-5.49	115.56	119.40
35	BB	26	C	N3-C2-O2	-5.49	118.06	121.90
35	BB	590	G	N9-C1'-C2'	-5.49	105.97	112.00
38	BE	189	A	C3'-C2'-C1'	-5.49	97.11	101.50
40	BG	24	A	C4'-C3'-O3'	-5.49	97.88	109.40
41	BH	110	C	P-O3'-C3'	5.49	126.28	119.70
51	BR	75	GLU	N-CA-CB	-5.49	100.73	110.60
63	Bd	6	ASN	CA-CB-CG	-5.49	101.33	113.40
80	Bu	153	ASP	N-CA-CB	5.49	120.47	110.60
82	Bw	223	ALA	CB-CA-C	-5.49	101.87	110.10
85	AA	128	U	C6-N1-C1'	-5.49	113.52	121.20
85	AA	235	U	O4'-C1'-N1	5.49	112.59	108.20
85	AA	678	A	C4'-C3'-C2'	5.49	108.08	102.60
85	AA	1714	G	N3-C4-N9	-5.49	122.71	126.00
85	AA	2151	U	O4'-C1'-N1	5.49	112.59	108.20
5	A4	101	GLN	N-CA-CB	5.48	120.47	110.60
8	A7	43	TRP	CB-CG-CD2	-5.48	119.47	126.60
34	BA	583	G	C5-C6-N1	5.48	114.24	111.50
34	BA	1107	A	C3'-C2'-C1'	-5.48	97.11	101.50
34	BA	1108	U	C6-N1-C1'	5.48	128.88	121.20
35	BB	802	G	C5-C6-O6	5.48	131.89	128.60
35	BB	816	U	O4'-C1'-N1	5.48	112.59	108.20
35	BB	879	G	C3'-C2'-C1'	-5.48	97.11	101.50
35	BB	1348	C	P-O5'-C5'	-5.48	112.13	120.90
35	BB	1359	G	C5'-C4'-C3'	-5.48	107.22	116.00
41	BH	48	G	O3'-P-O5'	-5.48	93.58	104.00
42	BI	87	ALA	N-CA-CB	-5.48	102.42	110.10
59	BZ	6	CYS	O-C-N	-5.48	113.92	122.70
85	AA	1428	A	P-O5'-C5'	-5.48	112.13	120.90
85	AA	1514	A	C5'-C4'-C3'	-5.48	107.22	116.00
85	AA	1658	G	C5-C6-N1	5.48	114.24	111.50
85	AA	1792	C	P-O3'-C3'	5.48	126.28	119.70
2	A1	155	ASP	CB-CG-OD2	-5.48	113.37	118.30
30	AW	50	HIS	CA-CB-CG	-5.48	104.28	113.60
34	BA	114	U	C6-N1-C1'	5.48	128.88	121.20
34	BA	125	G	N9-C1'-C2'	-5.48	105.97	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	279	U	C1'-O4'-C4'	-5.48	105.51	109.90
34	BA	719	G	O4'-C1'-N9	5.48	112.59	108.20
34	BA	1468	U	O5'-C5'-C4'	-5.48	101.28	111.70
35	BB	4	C	N3-C2-O2	-5.48	118.06	121.90
35	BB	1199	A	C4-C5-C6	-5.48	114.26	117.00
35	BB	1249	G	C1'-O4'-C4'	-5.48	105.51	109.90
35	BB	1370	G	P-O5'-C5'	-5.48	112.13	120.90
38	BE	108	U	C2-N1-C1'	-5.48	111.12	117.70
38	BE	144	A	C1'-O4'-C4'	-5.48	105.51	109.90
40	BG	87	G	C5'-C4'-C3'	-5.48	107.23	116.00
85	AA	334	A	O3'-P-O5'	5.48	114.42	104.00
85	AA	420	C	C5'-C4'-C3'	-5.48	107.23	116.00
85	AA	491	G	N3-C2-N2	5.48	123.74	119.90
85	AA	623	G	C5'-C4'-C3'	-5.48	107.23	116.00
85	AA	653	A	C3'-C2'-C1'	-5.48	97.11	101.50
85	AA	1141	U	P-O3'-C3'	-5.48	113.12	119.70
85	AA	1367	C	P-O5'-C5'	-5.48	112.13	120.90
85	AA	1923	A	N9-C1'-C2'	5.48	121.13	114.00
86	AB	3	C	C5-C4-N4	-5.48	116.36	120.20
29	AV	91	ARG	NE-CZ-NH2	-5.48	117.56	120.30
34	BA	315	U	C5'-C4'-O4'	5.48	115.68	109.10
34	BA	328	A	O4'-C1'-N9	5.48	112.58	108.20
34	BA	829	U	C5'-C4'-O4'	5.48	115.68	109.10
34	BA	1379	G	N9-C4-C5	5.48	107.59	105.40
35	BB	1012	G	O4'-C1'-N9	5.48	112.58	108.20
35	BB	1427	A	P-O3'-C3'	-5.48	113.12	119.70
38	BE	80	G	O4'-C1'-N9	5.48	112.58	108.20
40	BG	33	G	C4'-C3'-C2'	5.48	108.08	102.60
65	Bf	353	THR	N-CA-C	-5.48	96.20	111.00
67	Bh	60	ALA	N-CA-CB	5.48	117.77	110.10
85	AA	24	U	C4'-C3'-C2'	5.48	108.08	102.60
85	AA	364	C	O4'-C1'-C2'	5.48	112.53	107.60
85	AA	774	C	C3'-C2'-C1'	-5.48	97.11	101.50
85	AA	788	G	OP1-P-OP2	-5.48	111.38	119.60
85	AA	1116	G	C8-N9-C1'	5.48	134.12	127.00
85	AA	1212	C	P-O5'-C5'	5.48	129.67	120.90
34	BA	736	G	C3'-C2'-C1'	-5.48	97.12	101.50
34	BA	1151	A	P-O3'-C3'	5.48	126.28	119.70
34	BA	1205	A	C5-C6-N1	5.48	120.44	117.70
34	BA	1217	A	C5'-C4'-C3'	-5.48	107.23	116.00
35	BB	990	G	C4-N9-C1'	5.48	133.62	126.50
35	BB	1229	A	P-O3'-C3'	-5.48	113.13	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1446	C	N3-C2-O2	-5.48	118.06	121.90
38	BE	205	G	C8-N9-C1'	5.48	134.12	127.00
41	BH	130	G	N3-C2-N2	5.48	123.73	119.90
52	BS	76	TYR	CB-CG-CD2	-5.48	117.71	121.00
85	AA	819	G	C5-N7-C8	5.48	107.04	104.30
85	AA	1092	G	N3-C4-C5	-5.48	125.86	128.60
34	BA	174	A	C5-C6-N6	-5.48	119.32	123.70
34	BA	429	G	C5-C6-N1	5.48	114.24	111.50
34	BA	612	U	C1'-O4'-C4'	-5.48	105.52	109.90
34	BA	1246	G	C1'-O4'-C4'	-5.48	105.52	109.90
34	BA	1413	G	C6-N1-C2	-5.48	121.81	125.10
34	BA	1500	G	C4-N9-C1'	-5.48	119.38	126.50
35	BB	789	G	C4-C5-C6	-5.48	115.51	118.80
35	BB	1176	G	N1-C2-N2	-5.48	111.27	116.20
35	BB	1176	G	O4'-C1'-C2'	-5.48	100.32	105.80
36	BC	84	U	O4'-C1'-N1	5.48	112.58	108.20
37	BD	97	U	P-O3'-C3'	-5.48	113.13	119.70
38	BE	21	C	C5'-C4'-C3'	-5.48	107.24	116.00
38	BE	23	G	N3-C2-N2	5.48	123.73	119.90
40	BG	175	G	O4'-C4'-C3'	-5.48	98.52	104.00
62	Bc	21	LYS	CA-C-O	-5.48	108.60	120.10
65	Bf	160	HIS	N-CA-C	-5.48	96.21	111.00
70	Bk	47	ARG	NE-CZ-NH1	5.48	123.04	120.30
85	AA	17	C	C5'-C4'-O4'	5.48	115.67	109.10
85	AA	601	A	O4'-C1'-N9	5.48	112.58	108.20
34	BA	170	U	C5'-C4'-C3'	-5.48	107.24	116.00
34	BA	803	U	C5'-C4'-C3'	-5.48	107.24	116.00
34	BA	872	U	N1-C2-O2	-5.48	118.97	122.80
34	BA	1303	U	P-O3'-C3'	-5.48	113.13	119.70
35	BB	85	A	C5-C6-N6	-5.48	119.32	123.70
35	BB	775	U	O5'-P-OP2	-5.48	100.77	105.70
35	BB	1072	C	O4'-C1'-N1	5.48	112.58	108.20
35	BB	1426	G	N3-C4-N9	5.48	129.28	126.00
85	AA	485	A	C4-N9-C1'	-5.48	116.44	126.30
85	AA	648	G	N1-C6-O6	-5.48	116.61	119.90
85	AA	735	G	C4-N9-C1'	-5.48	119.38	126.50
34	BA	125	G	C8-N9-C1'	5.47	134.12	127.00
34	BA	184	C	O4'-C1'-N1	5.47	112.58	108.20
34	BA	569	C	C1'-O4'-C4'	-5.47	105.52	109.90
34	BA	928	C	P-O5'-C5'	-5.47	112.14	120.90
34	BA	1653	G	P-O3'-C3'	-5.47	113.13	119.70
34	BA	1815	G	N3-C2-N2	5.47	123.73	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	475	A	C5-C6-N6	5.47	128.08	123.70
35	BB	997	G	C5-C6-N1	5.47	114.24	111.50
35	BB	1202	G	C4-N9-C1'	5.47	133.62	126.50
53	BT	93	ASP	N-CA-CB	-5.47	100.75	110.60
73	Bn	49	TRP	CB-CG-CD1	5.47	134.12	127.00
82	Bw	175	ARG	NE-CZ-NH1	5.47	123.04	120.30
85	AA	10	G	C8-N9-C1'	5.47	134.12	127.00
85	AA	353	G	N1-C6-O6	5.47	123.19	119.90
85	AA	678	A	C6-N1-C2	-5.47	115.32	118.60
85	AA	897	A	P-O3'-C3'	-5.47	113.13	119.70
85	AA	970	U	C3'-C2'-C1'	5.47	105.88	101.50
85	AA	1823	G	C5-C6-O6	-5.47	125.32	128.60
85	AA	2240	G	P-O3'-C3'	-5.47	113.13	119.70
86	AB	57	G	C4-N9-C1'	-5.47	119.38	126.50
34	BA	148	G	O4'-C1'-N9	5.47	112.58	108.20
34	BA	266	G	C5-C6-N1	5.47	114.24	111.50
34	BA	909	G	C6-N1-C2	-5.47	121.82	125.10
34	BA	936	A	C1'-O4'-C4'	-5.47	105.52	109.90
35	BB	18	A	OP2-P-O3'	5.47	117.24	105.20
35	BB	355	A	P-O5'-C5'	5.47	129.66	120.90
35	BB	424	U	C5-C6-N1	-5.47	119.96	122.70
35	BB	1467	A	C4'-C3'-C2'	-5.47	97.13	102.60
35	BB	1510	G	O4'-C1'-N9	5.47	112.58	108.20
42	BI	135	MET	CA-CB-CG	5.47	122.60	113.30
71	Bl	130	SER	N-CA-CB	5.47	118.71	110.50
85	AA	38	C	O3'-P-O5'	5.47	114.40	104.00
85	AA	111	A	C5'-C4'-C3'	-5.47	107.25	116.00
85	AA	409	C	O4'-C1'-N1	5.47	112.58	108.20
85	AA	556	C	C5'-C4'-O4'	-5.47	102.53	109.10
85	AA	557	G	C5'-C4'-C3'	5.47	124.75	116.00
1	A0	22	GLU	CB-CA-C	5.47	121.34	110.40
34	BA	241	U	C5'-C4'-O4'	5.47	115.67	109.10
34	BA	1377	A	P-O3'-C3'	-5.47	113.14	119.70
34	BA	1771	U	C6-N1-C2	-5.47	117.72	121.00
35	BB	562	A	OP1-P-OP2	-5.47	111.39	119.60
37	BD	13	A	P-O5'-C5'	-5.47	112.15	120.90
37	BD	56	G	N1-C6-O6	5.47	123.18	119.90
38	BE	95	G	C6-C5-N7	5.47	133.68	130.40
47	BN	175	ARG	N-CA-CB	-5.47	100.75	110.60
57	BX	81	ALA	N-CA-CB	-5.47	102.44	110.10
69	Bj	85	HIS	CA-CB-CG	5.47	122.90	113.60
85	AA	1294	U	C5-C6-N1	-5.47	119.96	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2155	U	C3'-C2'-C1'	-5.47	97.12	101.50
13	AE	43	ASN	N-CA-CB	5.47	120.44	110.60
34	BA	138	C	C3'-C2'-C1'	-5.47	97.12	101.50
34	BA	537	C	C3'-C2'-C1'	-5.47	97.12	101.50
34	BA	584	A	C3'-C2'-C1'	5.47	105.88	101.50
34	BA	953	G	C5'-C4'-C3'	-5.47	107.25	116.00
34	BA	1205	A	O4'-C1'-N9	-5.47	103.82	108.20
34	BA	1258	G	O3'-P-O5'	-5.47	93.61	104.00
34	BA	1488	C	C2-N1-C1'	-5.47	112.78	118.80
34	BA	1537	G	C6-N1-C2	-5.47	121.82	125.10
35	BB	445	G	C4-N9-C1'	5.47	133.61	126.50
35	BB	457	U	N1-C1'-C2'	-5.47	105.98	112.00
35	BB	900	C	C1'-O4'-C4'	-5.47	105.53	109.90
37	BD	9	C	N1-C2-O2	5.47	122.18	118.90
38	BE	32	U	N3-C4-C5	-5.47	111.32	114.60
38	BE	92	C	N3-C2-O2	-5.47	118.07	121.90
85	AA	539	A	N9-C1'-C2'	-5.47	105.98	112.00
85	AA	544	A	C4-N9-C1'	-5.47	116.45	126.30
85	AA	849	A	C5-C6-N6	-5.47	119.32	123.70
85	AA	1160	U	C2-N1-C1'	-5.47	111.14	117.70
85	AA	1180	C	C5'-C4'-C3'	5.47	124.75	116.00
85	AA	1283	C	N3-C4-N4	-5.47	114.17	118.00
85	AA	1932	C	O4'-C1'-N1	5.47	112.58	108.20
34	BA	507	U	C5'-C4'-O4'	-5.47	102.54	109.10
34	BA	1401	C	O4'-C1'-N1	5.47	112.57	108.20
34	BA	1439	C	N1-C1'-C2'	-5.47	105.98	112.00
34	BA	1501	U	O4'-C4'-C3'	-5.47	98.53	104.00
35	BB	1017	U	O4'-C1'-N1	5.47	112.57	108.20
51	BR	76	TRP	CA-CB-CG	-5.47	103.31	113.70
84	By	9	ILE	N-CA-C	5.47	125.76	111.00
85	AA	146	U	N1-C1'-C2'	-5.47	105.99	112.00
85	AA	1101	C	P-O3'-C3'	-5.47	113.14	119.70
28	AU	96	ARG	NE-CZ-NH1	5.47	123.03	120.30
34	BA	266	G	P-O3'-C3'	-5.47	113.14	119.70
34	BA	774	A	N9-C1'-C2'	-5.47	105.99	112.00
34	BA	865	C	P-O5'-C5'	-5.47	112.15	120.90
34	BA	881	C	C1'-O4'-C4'	-5.47	105.53	109.90
34	BA	994	G	C5-N7-C8	-5.47	101.57	104.30
34	BA	1033	G	C5-N7-C8	-5.47	101.57	104.30
34	BA	1407	C	O4'-C1'-N1	5.47	112.57	108.20
34	BA	1563	G	P-O5'-C5'	5.47	129.65	120.90
34	BA	1833	G	P-O3'-C3'	-5.47	113.14	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	139	G	N7-C8-N9	-5.47	110.37	113.10
35	BB	435	A	C4-N9-C1'	-5.47	116.46	126.30
35	BB	648	G	O4'-C1'-N9	5.47	112.57	108.20
35	BB	852	G	O4'-C1'-N9	5.47	112.57	108.20
35	BB	1239	A	C4-N9-C1'	-5.47	116.46	126.30
35	BB	1509	G	C4'-C3'-C2'	-5.47	97.13	102.60
37	BD	61	C	C6-N1-C2	-5.47	118.11	120.30
38	BE	152	U	C5-C4-O4	-5.47	122.62	125.90
40	BG	173	C	C4'-C3'-C2'	-5.47	97.13	102.60
57	BX	122	ARG	NE-CZ-NH2	5.47	123.03	120.30
85	AA	327	G	N3-C4-C5	-5.47	125.87	128.60
85	AA	381	A	O5'-C5'-C4'	-5.47	101.31	111.70
85	AA	521	A	C4'-C3'-C2'	5.47	108.07	102.60
85	AA	2053	A	N7-C8-N9	5.47	116.53	113.80
34	BA	137	C	N3-C2-O2	-5.46	118.08	121.90
34	BA	288	U	N3-C2-O2	-5.46	118.38	122.20
34	BA	525	A	C4-C5-C6	-5.46	114.27	117.00
34	BA	692	U	O5'-P-OP2	-5.46	100.78	105.70
34	BA	697	A	C5-C6-N1	5.46	120.43	117.70
34	BA	896	U	P-O5'-C5'	-5.46	112.16	120.90
34	BA	1411	C	C4'-C3'-O3'	-5.46	97.92	109.40
34	BA	1526	C	C5'-C4'-O4'	5.46	115.66	109.10
35	BB	260	A	O4'-C1'-N9	5.46	112.57	108.20
35	BB	1007	U	P-O5'-C5'	-5.46	112.16	120.90
35	BB	1133	C	N3-C2-O2	-5.46	118.08	121.90
35	BB	1209	A	C5-N7-C8	-5.46	101.17	103.90
37	BD	91	U	N3-C2-O2	-5.46	118.38	122.20
39	BF	57	C	O4'-C1'-N1	5.46	112.57	108.20
40	BG	30	C	N1-C2-O2	5.46	122.18	118.90
40	BG	37	G	C1'-O4'-C4'	-5.46	105.53	109.90
40	BG	162	A	C3'-C2'-C1'	5.46	105.87	101.50
41	BH	65	G	C4-N9-C1'	-5.46	119.40	126.50
85	AA	111	A	C3'-C2'-C1'	5.46	105.87	101.50
85	AA	1302	A	O4'-C1'-N9	5.46	112.57	108.20
85	AA	2057	G	C5-C6-O6	-5.46	125.32	128.60
85	AA	2112	G	C8-N9-C1'	5.46	134.10	127.00
34	BA	493	G	N1-C6-O6	-5.46	116.62	119.90
34	BA	919	A	N7-C8-N9	-5.46	111.07	113.80
34	BA	1499	A	C2-N3-C4	5.46	113.33	110.60
35	BB	10	C	C4'-C3'-C2'	-5.46	97.14	102.60
35	BB	54	U	C4'-C3'-C2'	5.46	108.06	102.60
35	BB	114	A	O4'-C1'-N9	5.46	112.57	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	977	G	C3'-C2'-C1'	-5.46	97.13	101.50
35	BB	977	G	OP1-P-OP2	-5.46	111.41	119.60
37	BD	93	G	N3-C4-C5	-5.46	125.87	128.60
65	Bf	392	ARG	N-CA-C	5.46	125.75	111.00
85	AA	577	U	C2'-C3'-O3'	5.46	122.44	113.70
85	AA	1294	U	C6-N1-C1'	5.46	128.85	121.20
85	AA	2215	C	C3'-C2'-C1'	-5.46	97.13	101.50
34	BA	381	A	C5-C6-N6	-5.46	119.33	123.70
34	BA	427	G	N3-C2-N2	5.46	123.72	119.90
34	BA	635	G	O4'-C1'-N9	5.46	112.57	108.20
34	BA	1019	C	C5'-C4'-C3'	-5.46	107.26	116.00
35	BB	83	G	P-O5'-C5'	5.46	129.64	120.90
35	BB	950	G	N1-C6-O6	5.46	123.18	119.90
35	BB	1233	U	C5-C6-N1	-5.46	119.97	122.70
35	BB	1306	G	C5'-C4'-C3'	-5.46	107.26	116.00
35	BB	1501	U	O4'-C1'-N1	5.46	112.57	108.20
37	BD	11	A	C5'-C4'-C3'	-5.46	107.26	116.00
40	BG	111	C	P-O3'-C3'	-5.46	113.15	119.70
40	BG	169	A	C2-N3-C4	-5.46	107.87	110.60
77	Br	121	PHE	CB-CG-CD2	-5.46	116.98	120.80
85	AA	461	G	N9-C4-C5	-5.46	103.22	105.40
85	AA	727	U	C3'-C2'-C1'	-5.46	97.13	101.50
85	AA	813	G	C5-C6-O6	-5.46	125.32	128.60
85	AA	858	G	N9-C1'-C2'	-5.46	105.99	112.00
85	AA	1202	G	O4'-C1'-N9	5.46	112.57	108.20
85	AA	1247	A	N9-C1'-C2'	-5.46	105.99	112.00
34	BA	346	A	C4'-C3'-C2'	-5.46	97.14	102.60
34	BA	679	U	C1'-O4'-C4'	5.46	114.27	109.90
34	BA	1233	U	N3-C2-O2	-5.46	118.38	122.20
35	BB	37	C	N3-C2-O2	-5.46	118.08	121.90
35	BB	121	A	C3'-C2'-C1'	-5.46	97.13	101.50
40	BG	152	G	C5'-C4'-C3'	-5.46	107.26	116.00
67	Bh	139	GLY	N-CA-C	-5.46	99.45	113.10
85	AA	477	U	C5'-C4'-C3'	5.46	124.74	116.00
85	AA	792	A	O4'-C1'-N9	5.46	112.57	108.20
85	AA	984	A	C8-N9-C4	-5.46	103.62	105.80
34	BA	223	U	C4'-C3'-C2'	-5.46	97.14	102.60
34	BA	387	A	C5'-C4'-C3'	-5.46	107.27	116.00
34	BA	389	U	O4'-C4'-C3'	-5.46	98.54	104.00
34	BA	484	A	C5'-C4'-O4'	-5.46	102.55	109.10
34	BA	513	U	C5'-C4'-C3'	-5.46	107.27	116.00
34	BA	674	G	N1-C6-O6	5.46	123.17	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	786	U	P-O3'-C3'	5.46	126.25	119.70
34	BA	868	C	OP1-P-OP2	5.46	127.79	119.60
34	BA	1269	C	N3-C2-O2	-5.46	118.08	121.90
34	BA	1496	G	OP1-P-OP2	-5.46	111.41	119.60
34	BA	1608	C	O4'-C1'-N1	5.46	112.57	108.20
34	BA	1779	U	C6-N1-C1'	5.46	128.84	121.20
35	BB	429	C	N1-C2-N3	5.46	123.02	119.20
35	BB	445	G	C8-N9-C4	-5.46	104.22	106.40
36	BC	18	G	C4-C5-C6	-5.46	115.53	118.80
37	BD	49	A	C5-N7-C8	-5.46	101.17	103.90
38	BE	107	U	C4'-C3'-C2'	-5.46	97.14	102.60
64	Be	115	ASN	CA-CB-CG	-5.46	101.39	113.40
85	AA	439	U	P-O5'-C5'	-5.46	112.17	120.90
85	AA	757	A	P-O3'-C3'	-5.46	113.15	119.70
85	AA	889	G	C5'-C4'-O4'	5.46	115.65	109.10
85	AA	1217	U	C5-C4-O4	5.46	129.18	125.90
85	AA	2163	G	C5-C6-O6	5.46	131.88	128.60
21	AM	84	PHE	N-CA-C	-5.46	96.27	111.00
34	BA	335	C	C6-N1-C2	-5.46	118.12	120.30
34	BA	528	C	C3'-C2'-C1'	-5.46	97.14	101.50
34	BA	798	G	C6-C5-N7	-5.46	127.13	130.40
34	BA	1247	G	C4-N9-C1'	-5.46	119.41	126.50
34	BA	1786	C	N3-C2-O2	-5.46	118.08	121.90
34	BA	1839	G	C4-N9-C1'	-5.46	119.41	126.50
35	BB	86	A	O3'-P-O5'	-5.46	93.63	104.00
35	BB	618	U	N1-C2-O2	5.46	126.62	122.80
35	BB	1453	G	O4'-C4'-C3'	-5.46	98.54	104.00
37	BD	25	G	O3'-P-O5'	-5.46	93.63	104.00
38	BE	107	U	O4'-C1'-N1	5.46	112.57	108.20
40	BG	142	A	P-O3'-C3'	5.46	126.25	119.70
77	Br	217	ARG	CD-NE-CZ	5.46	131.24	123.60
80	Bu	99	TYR	N-CA-CB	-5.46	100.78	110.60
85	AA	36	U	O4'-C1'-N1	5.46	112.57	108.20
85	AA	760	U	N1-C1'-C2'	-5.46	106.00	112.00
85	AA	1114	A	P-O3'-C3'	-5.46	113.15	119.70
85	AA	1704	C	N3-C2-O2	-5.46	118.08	121.90
85	AA	1821	C	C5'-C4'-C3'	-5.46	107.27	116.00
85	AA	1877	G	N7-C8-N9	5.46	115.83	113.10
85	AA	2073	U	C6-N1-C1'	5.46	128.84	121.20
85	AA	2190	U	C2-N1-C1'	-5.46	111.15	117.70
12	AD	11	ASP	N-CA-CB	-5.46	100.78	110.60
34	BA	52	G	C1'-O4'-C4'	-5.46	105.54	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1287	G	P-O3'-C3'	-5.46	113.15	119.70
35	BB	101	U	O3'-P-O5'	-5.46	93.64	104.00
35	BB	606	C	O4'-C1'-N1	5.46	112.56	108.20
35	BB	786	A	P-O3'-C3'	-5.46	113.15	119.70
35	BB	1449	G	C5-C6-N1	5.46	114.23	111.50
39	BF	24	G	C4'-C3'-C2'	-5.46	97.14	102.60
40	BG	166	C	P-O5'-C5'	5.46	129.63	120.90
62	Bc	13	ARG	CB-CG-CD	5.46	125.78	111.60
85	AA	1134	G	P-O3'-C3'	-5.46	113.15	119.70
22	AO	117	LEU	CB-CA-C	-5.45	99.84	110.20
34	BA	78	U	O3'-P-O5'	-5.45	93.64	104.00
34	BA	661	C	N1-C2-O2	5.45	122.17	118.90
34	BA	704	G	C5-C6-O6	-5.45	125.33	128.60
34	BA	987	C	N3-C2-O2	-5.45	118.08	121.90
34	BA	1739	G	C2-N3-C4	-5.45	109.17	111.90
35	BB	121	A	C1'-O4'-C4'	-5.45	105.54	109.90
35	BB	1050	A	P-O5'-C5'	-5.45	112.17	120.90
35	BB	1146	C	C6-N1-C1'	-5.45	114.26	120.80
35	BB	1356	G	C5-C6-N1	5.45	114.23	111.50
35	BB	1506	C	C2-N3-C4	-5.45	117.17	119.90
35	BB	1546	C	C2-N3-C4	-5.45	117.17	119.90
41	BH	10	U	N1-C2-N3	5.45	118.17	114.90
85	AA	266	U	O3'-P-O5'	5.45	114.36	104.00
85	AA	809	A	C2'-C3'-O3'	5.45	122.43	113.70
85	AA	2089	G	C8-N9-C4	5.45	108.58	106.40
2	A1	51	TYR	N-CA-C	-5.45	96.28	111.00
34	BA	557	U	C4-C5-C6	5.45	122.97	119.70
34	BA	602	G	N1-C2-N2	5.45	121.11	116.20
34	BA	736	G	N3-C2-N2	5.45	123.72	119.90
34	BA	747	G	N1-C6-O6	-5.45	116.63	119.90
34	BA	941	G	P-O5'-C5'	-5.45	112.18	120.90
34	BA	1232	C	C3'-C2'-C1'	-5.45	97.14	101.50
34	BA	1329	U	C2-N3-C4	-5.45	123.73	127.00
35	BB	40	C	N3-C4-N4	-5.45	114.18	118.00
41	BH	127	A	C5-C6-N1	5.45	120.43	117.70
77	Br	329	LEU	O-C-N	-5.45	113.98	122.70
85	AA	880	A	C4-N9-C1'	-5.45	116.49	126.30
16	AH	13	SER	C-N-CA	5.45	135.33	121.70
34	BA	362	G	C1'-O4'-C4'	-5.45	105.54	109.90
34	BA	530	A	P-O5'-C5'	-5.45	112.18	120.90
34	BA	697	A	C6-N1-C2	-5.45	115.33	118.60
34	BA	701	G	P-O3'-C3'	5.45	126.24	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	748	C	C3'-C2'-C1'	-5.45	97.14	101.50
34	BA	1177	C	O3'-P-O5'	-5.45	93.64	104.00
34	BA	1338	G	P-O3'-C3'	-5.45	113.16	119.70
35	BB	107	A	O4'-C1'-N9	5.45	112.56	108.20
35	BB	968	C	C6-N1-C2	-5.45	118.12	120.30
35	BB	1201	G	C5'-C4'-O4'	-5.45	102.56	109.10
35	BB	1284	U	C2-N1-C1'	-5.45	111.16	117.70
40	BG	161	C	N3-C2-O2	-5.45	118.08	121.90
41	BH	51	C	N3-C2-O2	-5.45	118.08	121.90
69	Bj	5	ARG	NE-CZ-NH2	-5.45	117.58	120.30
80	Bu	56	THR	CA-CB-OG1	5.45	120.45	109.00
85	AA	100	A	O3'-P-O5'	5.45	114.36	104.00
85	AA	386	G	N9-C1'-C2'	-5.45	106.00	112.00
85	AA	815	G	C3'-C2'-C1'	5.45	105.86	101.50
85	AA	1538	C	C2-N3-C4	-5.45	117.17	119.90
85	AA	1605	G	P-O5'-C5'	-5.45	112.18	120.90
85	AA	1753	A	C3'-C2'-C1'	5.45	105.86	101.50
86	AB	50	U	O3'-P-O5'	-5.45	93.64	104.00
34	BA	113	G	C4'-C3'-O3'	5.45	123.89	113.00
34	BA	375	C	P-O5'-C5'	5.45	129.62	120.90
34	BA	887	U	N3-C2-O2	-5.45	118.39	122.20
34	BA	1187	U	C4'-C3'-C2'	-5.45	97.15	102.60
34	BA	1813	C	P-O5'-C5'	-5.45	112.18	120.90
35	BB	107	A	C3'-C2'-C1'	-5.45	97.14	101.50
35	BB	619	A	C8-N9-C1'	5.45	137.51	127.70
35	BB	1053	G	N1-C2-N2	-5.45	111.30	116.20
37	BD	30	A	N9-C4-C5	-5.45	103.62	105.80
39	BF	62	U	C1'-O4'-C4'	-5.45	105.54	109.90
40	BG	29	U	C3'-C2'-C1'	-5.45	97.14	101.50
40	BG	75	C	C4'-C3'-C2'	5.45	108.05	102.60
40	BG	113	G	C4-N9-C1'	-5.45	119.42	126.50
41	BH	72	G	C8-N9-C4	5.45	108.58	106.40
41	BH	102	C	C2-N3-C4	-5.45	117.17	119.90
55	BV	73	ARG	NE-CZ-NH1	5.45	123.03	120.30
63	Bd	55	ALA	O-C-N	-5.45	113.98	122.70
65	Bf	391	ARG	NE-CZ-NH2	-5.45	117.58	120.30
76	Bq	46	ARG	NE-CZ-NH1	5.45	123.02	120.30
85	AA	425	G	C3'-C2'-C1'	-5.45	97.14	101.50
85	AA	788	G	N3-C4-C5	-5.45	125.88	128.60
6	A5	25	MET	N-CA-C	5.45	125.71	111.00
34	BA	348	U	N1-C2-N3	5.45	118.17	114.90
34	BA	1240	G	C8-N9-C1'	-5.45	119.92	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1296	U	N3-C4-C5	-5.45	111.33	114.60
38	BE	60	C	N3-C2-O2	-5.45	118.09	121.90
39	BF	55	A	C6-N1-C2	-5.45	115.33	118.60
66	Bg	19	ASP	CB-CG-OD1	-5.45	113.40	118.30
85	AA	286	C	N3-C4-N4	5.45	121.81	118.00
85	AA	370	A	C5-C6-N1	5.45	120.42	117.70
85	AA	523	U	C2-N1-C1'	-5.45	111.16	117.70
85	AA	1483	A	C4-C5-C6	-5.45	114.28	117.00
34	BA	95	C	O4'-C1'-N1	5.45	112.56	108.20
34	BA	110	C	O4'-C4'-C3'	-5.45	98.56	104.00
34	BA	339	G	C2'-C3'-O3'	5.45	122.41	113.70
34	BA	953	G	P-O5'-C5'	-5.45	112.19	120.90
34	BA	1090	A	C4'-C3'-C2'	-5.45	97.15	102.60
34	BA	1685	C	N3-C2-O2	-5.45	118.09	121.90
35	BB	760	C	C6-N1-C2	-5.45	118.12	120.30
35	BB	850	U	C2-N1-C1'	5.45	124.24	117.70
35	BB	1359	G	C3'-C2'-C1'	-5.45	97.14	101.50
35	BB	1361	A	O4'-C1'-N9	5.45	112.56	108.20
35	BB	1466	A	P-O3'-C3'	-5.45	113.17	119.70
37	BD	41	G	C5'-C4'-C3'	5.45	124.71	116.00
37	BD	91	U	C3'-C2'-C1'	-5.45	97.14	101.50
38	BE	168	C	C5-C4-N4	5.45	124.01	120.20
52	BS	36	PHE	CB-CG-CD2	-5.45	116.99	120.80
85	AA	97	A	C4'-C3'-C2'	-5.45	97.15	102.60
85	AA	423	G	C6-N1-C2	-5.45	121.83	125.10
85	AA	765	U	O4'-C4'-C3'	-5.45	98.55	104.00
85	AA	1578	G	C5'-C4'-C3'	5.45	124.71	116.00
13	AE	69	TYR	CB-CG-CD1	-5.44	117.73	121.00
34	BA	290	G	C8-N9-C1'	5.44	134.08	127.00
34	BA	538	G	C5'-C4'-C3'	5.44	124.71	116.00
34	BA	1227	U	OP1-P-O3'	5.44	117.18	105.20
34	BA	1505	G	N3-C4-C5	-5.44	125.88	128.60
35	BB	10	C	O4'-C1'-N1	5.44	112.56	108.20
35	BB	883	G	C5'-C4'-O4'	5.44	115.63	109.10
35	BB	1071	G	C3'-C2'-C1'	-5.44	97.14	101.50
35	BB	1367	U	C2-N1-C1'	-5.44	111.17	117.70
35	BB	1546	C	N1-C2-N3	5.44	123.01	119.20
37	BD	66	G	C1'-O4'-C4'	-5.44	105.55	109.90
34	BA	347	A	P-O5'-C5'	5.44	129.61	120.90
34	BA	353	U	C5'-C4'-O4'	5.44	115.63	109.10
34	BA	471	U	O3'-P-O5'	-5.44	93.66	104.00
34	BA	592	G	C5'-C4'-C3'	5.44	124.71	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	651	U	P-O3'-C3'	-5.44	113.17	119.70
34	BA	1087	A	C6-C5-N7	-5.44	128.49	132.30
34	BA	1557	G	C4-N9-C1'	-5.44	119.42	126.50
35	BB	790	A	O4'-C1'-N9	5.44	112.55	108.20
35	BB	850	U	N1-C1'-C2'	5.44	121.08	114.00
35	BB	1129	C	P-O3'-C3'	-5.44	113.17	119.70
35	BB	1218	G	N1-C6-O6	5.44	123.17	119.90
35	BB	1508	G	C5-C6-O6	5.44	131.87	128.60
36	BC	14	G	C8-N9-C1'	5.44	134.08	127.00
38	BE	142	A	C5-C6-N6	5.44	128.06	123.70
60	Ba	63	ARG	NE-CZ-NH2	-5.44	117.58	120.30
77	Br	338	ALA	N-CA-CB	-5.44	102.48	110.10
80	Bu	184	ASN	CA-CB-CG	-5.44	101.42	113.40
82	Bw	234	ARG	NE-CZ-NH1	5.44	123.02	120.30
85	AA	99	U	C1'-O4'-C4'	-5.44	105.55	109.90
85	AA	474	C	C1'-O4'-C4'	-5.44	105.55	109.90
85	AA	639	C	O3'-P-O5'	-5.44	93.66	104.00
85	AA	670	C	C5-C4-N4	5.44	124.01	120.20
85	AA	803	C	P-O3'-C3'	-5.44	113.17	119.70
85	AA	1276	A	N1-C6-N6	-5.44	115.33	118.60
85	AA	1442	U	C6-N1-C2	5.44	124.27	121.00
16	AH	14	ALA	N-CA-C	-5.44	96.31	111.00
31	AX	155	TYR	N-CA-C	-5.44	96.31	111.00
34	BA	205	G	O4'-C1'-N9	5.44	112.55	108.20
34	BA	575	U	P-O5'-C5'	5.44	129.61	120.90
34	BA	607	C	C2-N1-C1'	5.44	124.78	118.80
34	BA	633	G	C3'-C2'-C1'	-5.44	97.15	101.50
34	BA	1071	G	O4'-C1'-N9	5.44	112.55	108.20
34	BA	1091	U	C6-N1-C1'	5.44	128.82	121.20
34	BA	1348	G	C5'-C4'-C3'	-5.44	107.30	116.00
35	BB	37	C	P-O5'-C5'	-5.44	112.19	120.90
35	BB	404	A	O4'-C1'-N9	5.44	112.55	108.20
35	BB	416	U	C1'-O4'-C4'	-5.44	105.55	109.90
35	BB	810	G	P-O3'-C3'	-5.44	113.17	119.70
35	BB	1124	G	N9-C1'-C2'	-5.44	106.02	112.00
35	BB	1363	A	C8-N9-C4	5.44	107.98	105.80
35	BB	1511	U	C6-N1-C1'	5.44	128.82	121.20
36	BC	86	U	C2-N1-C1'	-5.44	111.17	117.70
37	BD	94	C	O3'-P-O5'	-5.44	93.66	104.00
38	BE	123	A	C5-C6-N1	-5.44	114.98	117.70
38	BE	162	U	P-O3'-C3'	-5.44	113.17	119.70
41	BH	89	C	C2-N1-C1'	5.44	124.78	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
62	Bc	109	ARG	NE-CZ-NH1	5.44	123.02	120.30
85	AA	474	C	O5'-C5'-C4'	-5.44	101.36	111.70
85	AA	549	A	N9-C1'-C2'	-5.44	106.02	112.00
85	AA	898	A	C8-N9-C1'	-5.44	117.91	127.70
85	AA	1057	G	C5'-C4'-C3'	-5.44	107.30	116.00
85	AA	1096	G	P-O3'-C3'	-5.44	113.17	119.70
5	A4	114	LEU	N-CA-C	5.44	125.69	111.00
34	BA	52	G	P-O5'-C5'	-5.44	112.20	120.90
34	BA	284	U	OP1-P-OP2	-5.44	111.44	119.60
34	BA	661	C	C1'-O4'-C4'	-5.44	105.55	109.90
34	BA	820	C	O4'-C1'-N1	5.44	112.55	108.20
34	BA	1613	G	C4-N9-C1'	-5.44	119.43	126.50
34	BA	1617	U	C5-C6-N1	-5.44	119.98	122.70
35	BB	997	G	P-O3'-C3'	-5.44	113.17	119.70
35	BB	1033	U	N1-C1'-C2'	-5.44	106.02	112.00
35	BB	1467	A	P-O5'-C5'	5.44	129.60	120.90
71	Bl	124	VAL	C-N-CA	5.44	135.30	121.70
85	AA	675	A	C8-N9-C1'	5.44	137.49	127.70
1	A0	192	ARG	N-CA-CB	-5.44	100.81	110.60
34	BA	680	C	N3-C4-N4	-5.44	114.19	118.00
34	BA	955	G	O3'-P-O5'	-5.44	93.67	104.00
34	BA	1473	A	C5-C6-N1	5.44	120.42	117.70
34	BA	1489	U	C5-C6-N1	5.44	125.42	122.70
35	BB	598	C	C2-N1-C1'	-5.44	112.82	118.80
35	BB	681	G	C5'-C4'-C3'	-5.44	107.30	116.00
35	BB	1374	U	P-O3'-C3'	-5.44	113.17	119.70
36	BC	73	U	P-O3'-C3'	-5.44	113.17	119.70
36	BC	75	G	C8-N9-C4	-5.44	104.22	106.40
40	BG	137	G	C4-N9-C1'	-5.44	119.43	126.50
47	BN	23	GLN	CB-CA-C	-5.44	99.53	110.40
59	BZ	10	ARG	NE-CZ-NH2	-5.44	117.58	120.30
65	Bf	220	ARG	NE-CZ-NH1	5.44	123.02	120.30
85	AA	478	U	N3-C4-C5	-5.44	111.34	114.60
85	AA	492	C	N3-C2-O2	-5.44	118.09	121.90
85	AA	1132	A	O4'-C4'-C3'	-5.44	98.56	104.00
85	AA	1759	U	O3'-P-O5'	-5.44	93.67	104.00
85	AA	1848	G	C4-N9-C1'	-5.44	119.43	126.50
85	AA	2039	G	O4'-C4'-C3'	-5.44	98.56	104.00
85	AA	2105	G	N9-C1'-C2'	-5.44	106.02	112.00
34	BA	1164	C	N3-C2-O2	-5.44	118.09	121.90
34	BA	1609	U	O4'-C1'-C2'	5.44	112.49	107.60
35	BB	25	A	O4'-C1'-C2'	-5.44	100.36	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1156	U	O4'-C1'-N1	5.44	112.55	108.20
51	BR	30	PHE	CB-CG-CD2	-5.44	117.00	120.80
85	AA	504	U	C3'-C2'-C1'	-5.44	97.15	101.50
85	AA	1811	C	P-O3'-C3'	5.44	126.22	119.70
21	AM	85	LEU	N-CA-C	-5.43	96.33	111.00
34	BA	294	C	C2'-C3'-O3'	5.43	122.39	113.70
34	BA	579	U	C5'-C4'-C3'	-5.43	107.31	116.00
34	BA	598	G	OP1-P-O3'	5.43	117.16	105.20
34	BA	607	C	O4'-C1'-N1	5.43	112.55	108.20
34	BA	859	G	C8-N9-C4	5.43	108.57	106.40
34	BA	1331	G	C4'-C3'-C2'	5.43	108.03	102.60
34	BA	1381	A	C8-N9-C1'	-5.43	117.92	127.70
34	BA	1551	G	C8-N9-C4	5.43	108.57	106.40
35	BB	981	A	O4'-C1'-N9	5.43	112.55	108.20
35	BB	1506	C	O5'-C5'-C4'	5.43	122.03	111.70
36	BC	33	U	N3-C4-O4	5.43	123.20	119.40
38	BE	143	A	N1-C2-N3	-5.43	126.58	129.30
39	BF	60	C	N3-C4-C5	-5.43	119.73	121.90
40	BG	47	G	C4-N9-C1'	-5.43	119.44	126.50
41	BH	101	A	C6-N1-C2	-5.43	115.34	118.60
50	BQ	182	THR	CA-CB-CG2	-5.43	104.79	112.40
54	BU	133	ARG	O-C-N	-5.43	114.00	122.70
85	AA	260	A	C3'-C2'-C1'	5.43	105.85	101.50
85	AA	336	C	N3-C4-N4	5.43	121.80	118.00
85	AA	1547	G	P-O3'-C3'	-5.43	113.18	119.70
85	AA	2177	C	C1'-O4'-C4'	-5.43	105.55	109.90
7	A6	36	ARG	NE-CZ-NH2	-5.43	117.58	120.30
34	BA	6	C	N3-C4-N4	-5.43	114.20	118.00
34	BA	159	U	O4'-C4'-C3'	-5.43	98.57	104.00
34	BA	800	G	C8-N9-C1'	5.43	134.06	127.00
34	BA	876	C	P-O5'-C5'	-5.43	112.21	120.90
34	BA	1598	U	P-O5'-C5'	5.43	129.59	120.90
34	BA	1763	U	O4'-C1'-N1	5.43	112.55	108.20
35	BB	57	G	N9-C4-C5	5.43	107.57	105.40
35	BB	592	G	C5-C6-O6	-5.43	125.34	128.60
36	BC	119	G	N3-C2-N2	5.43	123.70	119.90
40	BG	4	A	C1'-O4'-C4'	-5.43	105.56	109.90
51	BR	23	ARG	NE-CZ-NH2	-5.43	117.58	120.30
61	Bb	114	HIS	N-CA-CB	5.43	120.38	110.60
85	AA	91	U	O3'-P-O5'	5.43	114.32	104.00
85	AA	159	G	O4'-C1'-N9	5.43	112.55	108.20
85	AA	190	A	C8-N9-C4	5.43	107.97	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	559	G	C1'-O4'-C4'	-5.43	105.55	109.90
85	AA	1224	C	N3-C2-O2	-5.43	118.10	121.90
85	AA	1230	U	C4'-C3'-C2'	-5.43	97.17	102.60
85	AA	1338	C	P-O5'-C5'	-5.43	112.21	120.90
85	AA	1458	G	N3-C2-N2	5.43	123.70	119.90
85	AA	1607	A	N7-C8-N9	5.43	116.52	113.80
34	BA	280	A	C5-N7-C8	-5.43	101.18	103.90
34	BA	1468	U	C6-N1-C2	-5.43	117.74	121.00
34	BA	1546	C	P-O5'-C5'	-5.43	112.21	120.90
34	BA	1779	U	C5'-C4'-C3'	-5.43	107.31	116.00
37	BD	77	A	N7-C8-N9	-5.43	111.08	113.80
37	BD	98	G	C4'-C3'-C2'	-5.43	97.17	102.60
38	BE	134	A	C6-C5-N7	5.43	136.10	132.30
40	BG	24	A	O5'-C5'-C4'	5.43	122.02	111.70
40	BG	88	G	C5'-C4'-C3'	-5.43	107.31	116.00
65	Bf	199	ARG	N-CA-CB	-5.43	100.82	110.60
85	AA	464	A	C5-C6-N6	-5.43	119.36	123.70
85	AA	962	U	C5-C6-N1	-5.43	119.98	122.70
85	AA	1211	C	N3-C2-O2	-5.43	118.10	121.90
85	AA	1690	A	N9-C1'-C2'	-5.43	106.03	112.00
33	AZ	38	ALA	N-CA-C	-5.43	96.34	111.00
34	BA	276	C	N1-C2-O2	5.43	122.16	118.90
34	BA	305	C	P-O3'-C3'	-5.43	113.18	119.70
34	BA	415	C	C5-C4-N4	5.43	124.00	120.20
34	BA	1055	U	O4'-C1'-N1	5.43	112.54	108.20
34	BA	1588	U	C5-C6-N1	-5.43	119.99	122.70
35	BB	532	C	N3-C2-O2	-5.43	118.10	121.90
35	BB	624	A	C4'-C3'-C2'	5.43	108.03	102.60
35	BB	1331	U	C6-N1-C1'	-5.43	113.60	121.20
35	BB	1356	G	N3-C4-C5	-5.43	125.89	128.60
35	BB	1514	G	O3'-P-O5'	-5.43	93.69	104.00
36	BC	67	U	N3-C2-O2	-5.43	118.40	122.20
37	BD	88	U	O4'-C1'-C2'	-5.43	100.37	105.80
37	BD	114	U	O5'-P-OP2	-5.43	100.81	105.70
38	BE	163	A	C8-N9-C4	5.43	107.97	105.80
38	BE	185	G	C1'-O4'-C4'	-5.43	105.56	109.90
59	BZ	43	ARG	N-CA-CB	-5.43	100.83	110.60
73	Bn	52	LYS	N-CA-CB	-5.43	100.83	110.60
85	AA	886	A	N1-C6-N6	-5.43	115.34	118.60
85	AA	1448	A	P-O3'-C3'	-5.43	113.18	119.70
85	AA	1677	A	C8-N9-C4	5.43	107.97	105.80
85	AA	1700	C	C2-N1-C1'	-5.43	112.83	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1904	C	O5'-C5'-C4'	-5.43	101.38	111.70
85	AA	2036	A	C5-N7-C8	5.43	106.61	103.90
85	AA	2073	U	O4'-C1'-N1	5.43	112.54	108.20
34	BA	1557	G	C8-N9-C1'	5.43	134.06	127.00
35	BB	76	C	N3-C2-O2	-5.43	118.10	121.90
35	BB	139	G	O5'-C5'-C4'	5.43	122.01	111.70
35	BB	394	A	P-O3'-C3'	-5.43	113.19	119.70
35	BB	1289	G	P-O5'-C5'	5.43	129.59	120.90
35	BB	1487	G	C8-N9-C1'	5.43	134.06	127.00
85	AA	1274	A	P-O5'-C5'	-5.43	112.22	120.90
85	AA	1491	G	C6-C5-N7	-5.43	127.14	130.40
85	AA	2039	G	N1-C2-N3	-5.43	120.64	123.90
34	BA	8	G	C8-N9-C4	5.43	108.57	106.40
34	BA	308	C	C2-N3-C4	-5.43	117.19	119.90
34	BA	315	U	N3-C4-O4	-5.43	115.60	119.40
34	BA	326	A	O4'-C4'-C3'	-5.43	98.57	104.00
34	BA	812	A	C8-N9-C1'	5.43	137.47	127.70
34	BA	965	A	P-O3'-C3'	-5.43	113.19	119.70
34	BA	1192	A	N9-C1'-C2'	-5.43	106.03	112.00
34	BA	1789	A	O3'-P-O5'	5.43	114.31	104.00
35	BB	342	U	O4'-C1'-N1	5.43	112.54	108.20
35	BB	424	U	N3-C2-O2	-5.43	118.40	122.20
35	BB	1100	C	C5-C6-N1	-5.43	118.29	121.00
38	BE	65	U	C6-N1-C2	-5.43	117.74	121.00
50	BQ	111	LEU	N-CA-CB	5.43	121.25	110.40
63	Bd	20	GLY	C-N-CA	5.43	135.27	121.70
85	AA	35	U	C5-C6-N1	5.43	125.41	122.70
85	AA	586	G	C5-C6-N1	5.43	114.21	111.50
85	AA	1522	U	C2-N3-C4	-5.43	123.74	127.00
85	AA	1831	U	O3'-P-O5'	-5.43	93.69	104.00
85	AA	2076	C	N3-C4-N4	-5.43	114.20	118.00
25	AR	11	GLY	N-CA-C	-5.42	99.54	113.10
34	BA	51	C	N3-C4-N4	-5.42	114.20	118.00
34	BA	160	G	C8-N9-C4	5.42	108.57	106.40
34	BA	730	C	P-O3'-C3'	-5.42	113.19	119.70
34	BA	822	U	O3'-P-O5'	5.42	114.31	104.00
34	BA	1032	A	P-O3'-C3'	-5.42	113.19	119.70
34	BA	1154	U	N3-C4-O4	-5.42	115.60	119.40
34	BA	1371	U	C6-N1-C2	-5.42	117.75	121.00
35	BB	115	A	P-O5'-C5'	-5.42	112.22	120.90
35	BB	1108	G	N3-C2-N2	5.42	123.70	119.90
35	BB	1192	C	C6-N1-C2	-5.42	118.13	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1498	G	C5'-C4'-O4'	-5.42	102.59	109.10
38	BE	67	A	C5-C6-N1	5.42	120.41	117.70
38	BE	108	U	C6-N1-C1'	5.42	128.79	121.20
38	BE	125	C	P-O3'-C3'	-5.42	113.19	119.70
39	BF	2	G	C4'-C3'-C2'	-5.42	97.18	102.60
39	BF	56	C	C5'-C4'-O4'	-5.42	102.59	109.10
64	Be	216	HIS	C-N-CA	5.42	135.26	121.70
83	Bx	51	ARG	NE-CZ-NH1	-5.42	117.59	120.30
85	AA	163	C	C5-C4-N4	-5.42	116.40	120.20
85	AA	285	C	C5'-C4'-O4'	5.42	115.61	109.10
85	AA	586	G	N1-C6-O6	5.42	123.15	119.90
85	AA	762	U	C5-C4-O4	-5.42	122.65	125.90
85	AA	1717	G	O4'-C1'-N9	5.42	112.54	108.20
34	BA	16	C	C6-N1-C1'	5.42	127.31	120.80
34	BA	53	G	N9-C4-C5	-5.42	103.23	105.40
35	BB	660	G	C4-N9-C1'	-5.42	119.45	126.50
35	BB	1230	A	O5'-P-OP2	-5.42	100.82	105.70
35	BB	1364	C	C6-N1-C2	-5.42	118.13	120.30
35	BB	1471	A	C8-N9-C1'	-5.42	117.94	127.70
85	AA	190	A	N7-C8-N9	-5.42	111.09	113.80
85	AA	1712	A	C4-N9-C1'	-5.42	116.54	126.30
34	BA	93	A	C8-N9-C1'	5.42	137.46	127.70
34	BA	309	U	C3'-C2'-C1'	-5.42	97.16	101.50
34	BA	687	G	O4'-C1'-N9	5.42	112.54	108.20
34	BA	769	U	C5'-C4'-O4'	5.42	115.61	109.10
34	BA	818	G	O4'-C1'-N9	5.42	112.54	108.20
34	BA	824	C	C6-N1-C2	-5.42	118.13	120.30
34	BA	898	G	C8-N9-C4	-5.42	104.23	106.40
34	BA	1667	G	C5-C6-N1	5.42	114.21	111.50
35	BB	1533	U	O4'-C1'-N1	5.42	112.54	108.20
36	BC	11	G	P-O5'-C5'	-5.42	112.22	120.90
38	BE	103	C	O4'-C1'-N1	5.42	112.54	108.20
48	BO	183	SER	CB-CA-C	5.42	120.40	110.10
66	Bg	69	TYR	CB-CG-CD1	5.42	124.25	121.00
68	Bi	14	ARG	CG-CD-NE	-5.42	100.42	111.80
81	Bv	169	LYS	N-CA-C	-5.42	96.36	111.00
83	Bx	38	SER	N-CA-C	-5.42	96.36	111.00
85	AA	185	A	O4'-C4'-C3'	-5.42	98.58	104.00
85	AA	267	U	C4'-C3'-O3'	5.42	123.84	113.00
85	AA	715	G	N3-C4-C5	-5.42	125.89	128.60
85	AA	779	G	N3-C2-N2	5.42	123.69	119.90
85	AA	811	A	P-O3'-C3'	-5.42	113.19	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1559	U	C5'-C4'-O4'	5.42	115.61	109.10
85	AA	1816	C	N1-C2-N3	-5.42	115.40	119.20
85	AA	1826	U	C6-N1-C2	-5.42	117.75	121.00
85	AA	1991	C	O5'-P-OP2	-5.42	100.82	105.70
34	BA	1145	U	P-O5'-C5'	-5.42	112.23	120.90
34	BA	1534	U	N1-C2-O2	5.42	126.59	122.80
34	BA	1628	A	N7-C8-N9	5.42	116.51	113.80
34	BA	1653	G	C8-N9-C4	5.42	108.57	106.40
35	BB	1119	G	C8-N9-C1'	5.42	134.04	127.00
35	BB	1483	A	C4'-C3'-C2'	5.42	108.02	102.60
45	BL	160	ARG	NE-CZ-NH2	-5.42	117.59	120.30
64	Be	73	LYS	N-CA-C	5.42	125.63	111.00
85	AA	706	U	C2-N1-C1'	-5.42	111.20	117.70
85	AA	879	G	P-O3'-C3'	-5.42	113.20	119.70
85	AA	1205	U	O4'-C1'-N1	5.42	112.54	108.20
85	AA	1291	A	C4'-C3'-C2'	-5.42	97.18	102.60
85	AA	1676	G	C4'-C3'-O3'	-5.42	98.02	109.40
1	A0	108	THR	C-N-CA	5.42	135.25	121.70
15	AG	20	ARG	NE-CZ-NH2	-5.42	117.59	120.30
16	AH	39	ASP	N-CA-CB	-5.42	100.85	110.60
29	AV	44	ARG	NE-CZ-NH1	5.42	123.01	120.30
34	BA	260	A	N9-C4-C5	-5.42	103.63	105.80
34	BA	320	G	C5-C6-N1	5.42	114.21	111.50
34	BA	329	G	N3-C2-N2	5.42	123.69	119.90
34	BA	784	C	N3-C4-C5	5.42	124.07	121.90
34	BA	1001	G	N9-C1'-C2'	-5.42	106.04	112.00
35	BB	127	U	O5'-C5'-C4'	-5.42	101.40	111.70
35	BB	1199	A	P-O5'-C5'	5.42	129.57	120.90
36	BC	26	U	C2'-C3'-O3'	5.42	122.37	113.70
36	BC	55	U	C2-N3-C4	5.42	130.25	127.00
36	BC	84	U	C2-N1-C1'	5.42	124.20	117.70
37	BD	4	U	C1'-O4'-C4'	-5.42	105.56	109.90
40	BG	109	C	C6-N1-C1'	5.42	127.30	120.80
41	BH	5	G	O5'-P-OP2	-5.42	100.82	105.70
67	Bh	146	ARG	NE-CZ-NH1	5.42	123.01	120.30
68	Bi	14	ARG	NE-CZ-NH2	-5.42	117.59	120.30
85	AA	273	C	N1-C2-O2	5.42	122.15	118.90
85	AA	334	A	N1-C6-N6	-5.42	115.35	118.60
85	AA	489	C	C2-N1-C1'	5.42	124.76	118.80
85	AA	496	C	C5'-C4'-C3'	5.42	124.67	116.00
34	BA	210	G	O4'-C1'-N9	5.42	112.53	108.20
34	BA	456	G	C5-C6-O6	-5.42	125.35	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	467	A	N9-C1'-C2'	-5.42	106.04	112.00
34	BA	1299	G	C2'-C3'-O3'	5.42	122.37	113.70
34	BA	1318	G	C4'-C3'-C2'	-5.42	97.18	102.60
34	BA	1323	G	C1'-O4'-C4'	-5.42	105.57	109.90
34	BA	1325	G	N1-C6-O6	5.42	123.15	119.90
34	BA	1840	C	N1-C2-O2	5.42	122.15	118.90
35	BB	78	C	P-O3'-C3'	-5.42	113.20	119.70
35	BB	384	A	C4-N9-C1'	5.42	136.05	126.30
35	BB	465	C	O5'-C5'-C4'	-5.42	101.41	111.70
35	BB	1248	A	C6-N1-C2	-5.42	115.35	118.60
35	BB	1446	C	P-O3'-C3'	5.42	126.20	119.70
38	BE	148	C	N3-C2-O2	-5.42	118.11	121.90
38	BE	162	U	N3-C2-O2	-5.42	118.41	122.20
40	BG	148	C	P-O3'-C3'	-5.42	113.20	119.70
77	Br	250	TRP	CB-CG-CD1	5.42	134.04	127.00
85	AA	467	U	P-O3'-C3'	-5.42	113.20	119.70
85	AA	550	G	N9-C1'-C2'	-5.42	106.04	112.00
85	AA	604	C	O4'-C1'-C2'	-5.42	100.38	105.80
85	AA	660	G	N9-C4-C5	-5.42	103.23	105.40
85	AA	797	C	O4'-C1'-N1	5.42	112.53	108.20
85	AA	1119	A	P-O5'-C5'	-5.42	112.23	120.90
85	AA	1672	G	C3'-C2'-C1'	-5.42	97.17	101.50
85	AA	1816	C	N3-C4-N4	5.42	121.79	118.00
85	AA	2200	A	C4-N9-C1'	-5.42	116.55	126.30
34	BA	175	G	C5-C6-O6	-5.42	125.35	128.60
34	BA	176	G	N9-C1'-C2'	-5.42	106.04	112.00
34	BA	1180	A	O5'-C5'-C4'	-5.42	101.41	111.70
35	BB	387	G	C5'-C4'-C3'	-5.42	107.33	116.00
35	BB	853	U	O3'-P-O5'	5.42	114.29	104.00
35	BB	1167	C	O5'-C5'-C4'	-5.42	101.41	111.70
37	BD	103	C	O4'-C1'-N1	5.42	112.53	108.20
38	BE	197	A	N9-C1'-C2'	-5.42	106.04	112.00
62	Bc	22	ARG	N-CA-C	5.42	125.62	111.00
62	Bc	33	ASN	CA-CB-CG	-5.42	101.49	113.40
85	AA	828	U	C6-N1-C2	-5.42	117.75	121.00
85	AA	1754	G	N1-C6-O6	5.42	123.15	119.90
85	AA	1894	G	C8-N9-C4	5.42	108.57	106.40
3	A2	30	ARG	NE-CZ-NH2	-5.41	117.59	120.30
34	BA	41	U	O4'-C1'-N1	5.41	112.53	108.20
34	BA	122	U	O4'-C1'-C2'	5.41	112.47	107.60
34	BA	454	G	C8-N9-C1'	5.41	134.04	127.00
34	BA	529	A	N1-C2-N3	-5.41	126.59	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	931	G	C8-N9-C4	5.41	108.56	106.40
34	BA	1816	G	N9-C4-C5	-5.41	103.23	105.40
35	BB	343	U	P-O3'-C3'	5.41	126.20	119.70
35	BB	466	A	P-O5'-C5'	-5.41	112.24	120.90
35	BB	505	G	C6-N1-C2	-5.41	121.85	125.10
35	BB	716	G	C5'-C4'-O4'	5.41	115.60	109.10
35	BB	790	A	C5'-C4'-C3'	-5.41	107.34	116.00
35	BB	1183	U	C3'-C2'-C1'	-5.41	97.17	101.50
35	BB	1232	A	C5-N7-C8	-5.41	101.19	103.90
38	BE	39	U	OP1-P-O3'	5.41	117.11	105.20
40	BG	157	A	C5-N7-C8	5.41	106.61	103.90
66	Bg	104	THR	CA-CB-CG2	-5.41	104.82	112.40
85	AA	33	U	C1'-O4'-C4'	-5.41	105.57	109.90
85	AA	116	G	C8-N9-C4	5.41	108.56	106.40
85	AA	588	G	O5'-C5'-C4'	-5.41	101.41	111.70
85	AA	674	U	C5-C6-N1	-5.41	119.99	122.70
85	AA	713	G	P-O5'-C5'	-5.41	112.24	120.90
85	AA	2139	G	O5'-C5'-C4'	-5.41	101.41	111.70
2	A1	51	TYR	CB-CG-CD1	5.41	124.25	121.00
34	BA	96	G	C1'-O4'-C4'	-5.41	105.57	109.90
34	BA	99	G	C1'-O4'-C4'	-5.41	105.57	109.90
34	BA	289	A	C5'-C4'-O4'	-5.41	102.61	109.10
34	BA	614	A	C5'-C4'-C3'	-5.41	107.34	116.00
34	BA	692	U	C4'-C3'-C2'	-5.41	97.19	102.60
34	BA	780	U	C5'-C4'-C3'	5.41	124.66	116.00
34	BA	908	G	P-O3'-C3'	-5.41	113.20	119.70
34	BA	1699	A	C4-C5-C6	-5.41	114.29	117.00
35	BB	740	A	O4'-C1'-N9	5.41	112.53	108.20
35	BB	933	U	O4'-C1'-N1	5.41	112.53	108.20
35	BB	1452	U	O5'-C5'-C4'	-5.41	101.42	111.70
35	BB	1453	G	C8-N9-C1'	5.41	134.04	127.00
38	BE	170	U	C5-C6-N1	-5.41	119.99	122.70
52	BS	115	ARG	NE-CZ-NH1	5.41	123.01	120.30
8	A7	245	PHE	CB-CG-CD1	5.41	124.59	120.80
34	BA	204	U	C6-N1-C2	-5.41	117.75	121.00
34	BA	329	G	C2-N3-C4	5.41	114.61	111.90
34	BA	1049	G	O4'-C1'-N9	5.41	112.53	108.20
34	BA	1432	C	N3-C2-O2	-5.41	118.11	121.90
34	BA	1510	C	O4'-C1'-C2'	5.41	112.47	107.60
35	BB	70	A	N9-C1'-C2'	-5.41	106.05	112.00
35	BB	638	G	O4'-C1'-C2'	5.41	112.47	107.60
35	BB	873	C	C6-N1-C2	-5.41	118.14	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1292	G	P-O3'-C3'	-5.41	113.21	119.70
36	BC	9	G	O5'-C5'-C4'	-5.41	101.42	111.70
40	BG	132	U	P-O5'-C5'	5.41	129.56	120.90
60	Ba	127	PHE	N-CA-CB	-5.41	100.86	110.60
80	Bu	78	ALA	N-CA-CB	5.41	117.67	110.10
85	AA	66	U	OP1-P-O3'	5.41	117.10	105.20
85	AA	388	G	C6-N1-C2	-5.41	121.85	125.10
85	AA	487	G	C5-C6-O6	5.41	131.85	128.60
85	AA	556	C	C5'-C4'-C3'	5.41	124.66	116.00
85	AA	695	A	C4'-C3'-C2'	5.41	108.01	102.60
85	AA	764	U	N1-C2-N3	-5.41	111.65	114.90
85	AA	988	C	C6-N1-C2	-5.41	118.14	120.30
85	AA	1295	G	O4'-C1'-N9	5.41	112.53	108.20
85	AA	2123	U	C5'-C4'-O4'	-5.41	102.61	109.10
85	AA	2246	U	O5'-P-OP2	-5.41	100.83	105.70
15	AG	135	LEU	N-CA-C	-5.41	96.40	111.00
34	BA	236	A	N9-C4-C5	-5.41	103.64	105.80
34	BA	1017	C	N3-C4-N4	-5.41	114.21	118.00
34	BA	1118	C	C4'-C3'-C2'	-5.41	97.19	102.60
34	BA	1633	C	C3'-C2'-C1'	5.41	105.83	101.50
34	BA	1817	G	C6-N1-C2	-5.41	121.86	125.10
35	BB	641	C	C2-N3-C4	-5.41	117.20	119.90
35	BB	1024	G	C6-N1-C2	-5.41	121.86	125.10
35	BB	1041	A	C4'-C3'-O3'	5.41	123.82	113.00
35	BB	1134	G	N3-C2-N2	5.41	123.69	119.90
35	BB	1199	A	C1'-O4'-C4'	-5.41	105.57	109.90
35	BB	1286	G	C4'-C3'-C2'	5.41	108.01	102.60
35	BB	1470	G	C5'-C4'-C3'	5.41	124.65	116.00
65	Bf	425	HIS	C-N-CA	5.41	133.66	122.30
85	AA	179	G	P-O3'-C3'	5.41	126.19	119.70
85	AA	213	G	O4'-C1'-N9	5.41	112.53	108.20
85	AA	453	G	N3-C2-N2	5.41	123.69	119.90
85	AA	852	C	N3-C4-N4	5.41	121.79	118.00
85	AA	929	G	C3'-C2'-C1'	5.41	105.83	101.50
85	AA	2199	G	P-O5'-C5'	-5.41	112.25	120.90
34	BA	401	A	O3'-P-O5'	-5.41	93.73	104.00
34	BA	458	G	C5'-C4'-C3'	-5.41	107.35	116.00
34	BA	532	C	N3-C4-C5	-5.41	119.74	121.90
34	BA	1067	G	C1'-O4'-C4'	-5.41	105.57	109.90
34	BA	1568	A	C6-N1-C2	-5.41	115.36	118.60
85	AA	338	G	C8-N9-C1'	5.41	134.03	127.00
85	AA	426	C	C2-N3-C4	5.41	122.60	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	830	A	C4-C5-C6	-5.41	114.30	117.00
85	AA	1281	G	N9-C1'-C2'	-5.41	106.05	112.00
10	A9	150	TYR	CB-CG-CD1	5.41	124.24	121.00
12	AD	33	TRP	CA-C-N	-5.41	105.31	117.20
34	BA	128	C	P-O3'-C3'	-5.41	113.21	119.70
34	BA	942	G	O4'-C1'-N9	5.41	112.53	108.20
34	BA	1561	C	N3-C4-N4	5.41	121.78	118.00
34	BA	1661	U	C3'-C2'-C1'	-5.41	97.18	101.50
35	BB	280	C	P-O3'-C3'	5.41	126.19	119.70
35	BB	611	U	O5'-C5'-C4'	-5.41	101.43	111.70
35	BB	859	U	O4'-C1'-N1	5.41	112.53	108.20
35	BB	1031	G	C5-C6-N1	5.41	114.20	111.50
35	BB	1035	C	P-O3'-C3'	-5.41	113.21	119.70
35	BB	1208	G	N3-C2-N2	5.41	123.68	119.90
35	BB	1215	U	N3-C2-O2	-5.41	118.42	122.20
36	BC	20	C	N3-C4-C5	-5.41	119.74	121.90
77	Br	307	PRO	C-N-CA	5.41	135.21	121.70
85	AA	794	A	C5'-C4'-O4'	5.41	115.59	109.10
85	AA	2042	G	N7-C8-N9	-5.41	110.40	113.10
34	BA	79	C	N1-C1'-C2'	-5.40	106.06	112.00
34	BA	635	G	O4'-C4'-C3'	-5.40	98.60	104.00
34	BA	1275	G	C3'-C2'-C1'	-5.40	97.18	101.50
35	BB	131	A	C8-N9-C1'	5.40	137.43	127.70
35	BB	496	C	C6-N1-C2	-5.40	118.14	120.30
35	BB	1254	G	C8-N9-C1'	5.40	134.03	127.00
37	BD	81	C	C5-C6-N1	-5.40	118.30	121.00
74	Bo	67	GLY	N-CA-C	5.40	126.61	113.10
85	AA	388	G	C5'-C4'-O4'	5.40	115.58	109.10
34	BA	148	G	C3'-C2'-C1'	5.40	105.82	101.50
34	BA	422	C	C6-N1-C2	-5.40	118.14	120.30
34	BA	1064	A	C4-N9-C1'	-5.40	116.58	126.30
34	BA	1800	G	O3'-P-O5'	5.40	114.27	104.00
34	BA	1807	G	O4'-C1'-N9	5.40	112.52	108.20
34	BA	1814	U	N1-C2-N3	5.40	118.14	114.90
35	BB	133	G	C4-N9-C1'	-5.40	119.48	126.50
35	BB	139	G	C4'-C3'-C2'	-5.40	97.20	102.60
35	BB	499	A	C4-N9-C1'	-5.40	116.58	126.30
35	BB	657	A	OP1-P-O3'	5.40	117.09	105.20
35	BB	792	G	C4-C5-N7	5.40	112.96	110.80
35	BB	828	G	O4'-C1'-N9	5.40	112.52	108.20
35	BB	956	G	N7-C8-N9	5.40	115.80	113.10
35	BB	1210	U	P-O3'-C3'	-5.40	113.22	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1310	C	C5'-C4'-O4'	5.40	115.58	109.10
35	BB	1467	A	C4'-C3'-O3'	5.40	123.81	113.00
37	BD	110	G	N7-C8-N9	-5.40	110.40	113.10
40	BG	119	A	C5-C6-N6	-5.40	119.38	123.70
40	BG	139	U	N1-C1'-C2'	-5.40	106.06	112.00
73	Bn	11	ARG	CD-NE-CZ	-5.40	116.04	123.60
77	Br	198	ARG	NE-CZ-NH2	5.40	123.00	120.30
84	By	88	PHE	CB-CG-CD2	-5.40	117.02	120.80
85	AA	418	G	C4-C5-C6	-5.40	115.56	118.80
85	AA	1087	G	P-O5'-C5'	-5.40	112.26	120.90
85	AA	1257	A	P-O5'-C5'	-5.40	112.26	120.90
85	AA	2083	G	N1-C6-O6	-5.40	116.66	119.90
3	A2	157	GLU	N-CA-CB	-5.40	100.88	110.60
26	AS	22	TRP	CB-CG-CD2	-5.40	119.58	126.60
34	BA	974	G	C6-N1-C2	-5.40	121.86	125.10
34	BA	1704	G	C5'-C4'-O4'	5.40	115.58	109.10
35	BB	361	A	O4'-C1'-N9	5.40	112.52	108.20
35	BB	623	A	C5-C6-N6	-5.40	119.38	123.70
35	BB	1104	A	N1-C6-N6	-5.40	115.36	118.60
35	BB	1126	A	C5'-C4'-O4'	5.40	115.58	109.10
35	BB	1165	A	C8-N9-C1'	5.40	137.42	127.70
35	BB	1338	U	C5'-C4'-C3'	-5.40	107.36	116.00
35	BB	1401	G	C1'-O4'-C4'	-5.40	105.58	109.90
37	BD	52	U	P-O5'-C5'	5.40	129.54	120.90
38	BE	170	U	N1-C2-O2	5.40	126.58	122.80
49	BP	17	ARG	NE-CZ-NH1	5.40	123.00	120.30
65	Bf	274	CYS	N-CA-CB	5.40	120.32	110.60
85	AA	103	U	C5'-C4'-O4'	5.40	115.58	109.10
85	AA	395	G	O3'-P-O5'	-5.40	93.74	104.00
85	AA	486	G	C8-N9-C4	5.40	108.56	106.40
85	AA	830	A	C5'-C4'-O4'	-5.40	102.62	109.10
85	AA	1524	A	P-O3'-C3'	-5.40	113.22	119.70
85	AA	1786	G	C1'-O4'-C4'	-5.40	105.58	109.90
85	AA	1885	A	C4'-C3'-C2'	5.40	108.00	102.60
31	AX	119	MET	CA-CB-CG	5.40	122.48	113.30
34	BA	142	A	O3'-P-O5'	-5.40	93.74	104.00
34	BA	590	U	P-O5'-C5'	5.40	129.54	120.90
34	BA	666	C	C1'-O4'-C4'	-5.40	105.58	109.90
34	BA	1472	G	N9-C4-C5	-5.40	103.24	105.40
35	BB	421	U	C3'-C2'-C1'	-5.40	97.18	101.50
35	BB	1230	A	OP1-P-OP2	-5.40	111.50	119.60
35	BB	1464	G	C3'-C2'-C1'	5.40	105.82	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1529	G	P-O5'-C5'	-5.40	112.26	120.90
36	BC	25	C	P-O5'-C5'	5.40	129.54	120.90
85	AA	815	G	C5'-C4'-O4'	5.40	115.58	109.10
85	AA	1491	G	C4-N9-C1'	-5.40	119.48	126.50
85	AA	2104	C	C5'-C4'-C3'	5.40	124.64	116.00
85	AA	2246	U	C3'-C2'-C1'	5.40	105.82	101.50
34	BA	218	G	N3-C2-N2	5.40	123.68	119.90
34	BA	289	A	N9-C4-C5	-5.40	103.64	105.80
34	BA	350	C	N3-C2-O2	-5.40	118.12	121.90
34	BA	521	C	O4'-C4'-C3'	-5.40	98.60	104.00
34	BA	708	C	C4'-C3'-C2'	5.40	108.00	102.60
34	BA	709	C	O4'-C1'-N1	5.40	112.52	108.20
34	BA	761	U	C1'-O4'-C4'	-5.40	105.58	109.90
34	BA	997	U	C1'-O4'-C4'	-5.40	105.58	109.90
34	BA	1311	G	C8-N9-C4	5.40	108.56	106.40
34	BA	1426	A	C3'-C2'-C1'	5.40	105.82	101.50
34	BA	1463	U	O5'-C5'-C4'	-5.40	101.44	111.70
35	BB	132	G	C6-N1-C2	-5.40	121.86	125.10
35	BB	474	G	C3'-C2'-C1'	-5.40	97.18	101.50
35	BB	975	G	O5'-C5'-C4'	-5.40	101.44	111.70
35	BB	1362	G	C6-N1-C2	-5.40	121.86	125.10
38	BE	88	G	C2'-C3'-O3'	5.40	122.34	113.70
38	BE	127	G	N3-C2-N2	5.40	123.68	119.90
39	BF	28	C	P-O3'-C3'	-5.40	113.22	119.70
40	BG	102	G	C4'-C3'-C2'	-5.40	97.20	102.60
41	BH	26	C	C2-N1-C1'	5.40	124.74	118.80
85	AA	192	G	N3-C4-N9	5.40	129.24	126.00
85	AA	965	G	C2'-C3'-O3'	5.40	122.34	113.70
85	AA	1010	U	N3-C2-O2	-5.40	118.42	122.20
85	AA	1211	C	P-O5'-C5'	-5.40	112.26	120.90
85	AA	1448	A	C5'-C4'-O4'	-5.40	102.62	109.10
85	AA	1998	A	O4'-C1'-N9	5.40	112.52	108.20
85	AA	2087	C	O4'-C1'-C2'	5.40	112.46	107.60
85	AA	2092	A	C2'-C3'-O3'	5.40	122.34	113.70
85	AA	2234	C	O4'-C1'-C2'	5.40	112.46	107.60
34	BA	754	G	N3-C2-N2	-5.40	116.12	119.90
34	BA	1516	G	C5-C6-O6	-5.40	125.36	128.60
34	BA	1803	A	N7-C8-N9	-5.40	111.10	113.80
36	BC	95	A	C4'-C3'-C2'	5.40	108.00	102.60
77	Br	186	ARG	NE-CZ-NH1	5.40	123.00	120.30
85	AA	148	G	C5'-C4'-C3'	-5.40	107.37	116.00
85	AA	1670	U	P-O5'-C5'	-5.40	112.27	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	AR	83	VAL	N-CA-C	-5.39	96.43	111.00
34	BA	734	G	C8-N9-C4	5.39	108.56	106.40
34	BA	757	G	N9-C4-C5	-5.39	103.24	105.40
34	BA	1241	U	C3'-C2'-C1'	-5.39	97.19	101.50
34	BA	1711	G	C4-C5-C6	-5.39	115.56	118.80
35	BB	688	U	C4'-C3'-C2'	5.39	107.99	102.60
35	BB	999	G	C1'-O4'-C4'	-5.39	105.58	109.90
35	BB	1049	G	N3-C2-N2	5.39	123.68	119.90
35	BB	1086	G	N9-C1'-C2'	-5.39	106.07	112.00
35	BB	1169	A	O5'-C5'-C4'	-5.39	101.45	111.70
35	BB	1487	G	C8-N9-C4	-5.39	104.24	106.40
37	BD	60	C	N3-C2-O2	-5.39	118.12	121.90
37	BD	93	G	C4'-C3'-C2'	-5.39	97.20	102.60
41	BH	17	A	P-O5'-C5'	-5.39	112.27	120.90
85	AA	386	G	C5-C6-O6	-5.39	125.36	128.60
85	AA	455	G	C4-C5-C6	-5.39	115.56	118.80
85	AA	455	G	O4'-C1'-C2'	5.39	112.45	107.60
85	AA	486	G	N7-C8-N9	-5.39	110.40	113.10
85	AA	2001	C	C5'-C4'-O4'	5.39	115.57	109.10
34	BA	157	U	OP1-P-OP2	-5.39	111.51	119.60
34	BA	364	C	N3-C2-O2	-5.39	118.12	121.90
34	BA	420	A	N1-C6-N6	5.39	121.84	118.60
34	BA	518	C	O4'-C1'-N1	5.39	112.51	108.20
34	BA	1428	G	C5-C6-O6	-5.39	125.36	128.60
34	BA	1719	G	N3-C4-C5	-5.39	125.90	128.60
34	BA	1818	A	N3-C4-N9	-5.39	123.09	127.40
34	BA	1832	A	C8-N9-C1'	5.39	137.41	127.70
35	BB	781	U	C2-N3-C4	-5.39	123.77	127.00
35	BB	797	C	C5'-C4'-O4'	-5.39	102.63	109.10
35	BB	1047	C	N1-C2-O2	5.39	122.14	118.90
35	BB	1502	U	O4'-C1'-N1	5.39	112.51	108.20
36	BC	22	U	C5'-C4'-C3'	-5.39	107.37	116.00
36	BC	78	G	C6-N1-C2	-5.39	121.86	125.10
37	BD	16	U	P-O3'-C3'	-5.39	113.23	119.70
40	BG	163	G	C4-C5-N7	5.39	112.96	110.80
85	AA	355	G	C8-N9-C4	5.39	108.56	106.40
85	AA	1001	G	N3-C2-N2	5.39	123.67	119.90
85	AA	1444	U	N3-C2-O2	-5.39	118.43	122.20
85	AA	1899	A	C5'-C4'-O4'	-5.39	102.63	109.10
85	AA	1927	G	OP2-P-O3'	5.39	117.06	105.20
85	AA	2214	A	O4'-C1'-N9	5.39	112.51	108.20
34	BA	212	A	O5'-C5'-C4'	-5.39	101.46	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	573	U	C2-N3-C4	-5.39	123.77	127.00
34	BA	795	G	N1-C6-O6	5.39	123.14	119.90
34	BA	871	G	O3'-P-O5'	-5.39	93.76	104.00
34	BA	1119	A	C4-C5-C6	-5.39	114.30	117.00
34	BA	1356	C	P-O3'-C3'	-5.39	113.23	119.70
34	BA	1666	U	N1-C1'-C2'	-5.39	106.07	112.00
34	BA	1777	U	O4'-C1'-N1	5.39	112.51	108.20
34	BA	1844	U	C2-N1-C1'	-5.39	111.23	117.70
35	BB	1360	A	C1'-O4'-C4'	-5.39	105.59	109.90
77	Br	54	ASN	CB-CA-C	5.39	121.18	110.40
85	AA	886	A	C5'-C4'-O4'	5.39	115.57	109.10
86	AB	66	U	C5'-C4'-C3'	5.39	124.62	116.00
2	A1	127	TYR	CB-CG-CD1	-5.39	117.77	121.00
34	BA	430	A	P-O3'-C3'	-5.39	113.23	119.70
34	BA	1175	G	N9-C1'-C2'	-5.39	106.07	112.00
34	BA	1470	G	C8-N9-C1'	5.39	134.01	127.00
35	BB	430	A	N9-C1'-C2'	-5.39	106.07	112.00
35	BB	1164	U	O5'-C5'-C4'	-5.39	101.46	111.70
35	BB	1415	G	O4'-C1'-N9	5.39	112.51	108.20
40	BG	64	C	C5'-C4'-C3'	-5.39	107.38	116.00
40	BG	80	G	C8-N9-C4	-5.39	104.24	106.40
41	BH	30	C	C4'-C3'-C2'	-5.39	97.21	102.60
64	Be	132	ASP	CA-CB-CG	-5.39	101.54	113.40
85	AA	96	C	N3-C2-O2	-5.39	118.13	121.90
85	AA	1091	C	P-O5'-C5'	5.39	129.52	120.90
85	AA	1499	G	C5-C6-N1	5.39	114.19	111.50
85	AA	1732	G	C6-N1-C2	-5.39	121.87	125.10
85	AA	2120	C	O4'-C1'-N1	5.39	112.51	108.20
85	AA	2200	A	N7-C8-N9	-5.39	111.11	113.80
5	A4	190	PHE	C-N-CA	5.39	135.17	121.70
34	BA	151	A	C8-N9-C4	-5.39	103.64	105.80
34	BA	251	U	N1-C1'-C2'	-5.39	106.07	112.00
34	BA	647	U	OP1-P-OP2	-5.39	111.52	119.60
35	BB	366	G	C3'-C2'-C1'	-5.39	97.19	101.50
35	BB	562	A	N1-C6-N6	-5.39	115.37	118.60
35	BB	677	U	O5'-P-OP2	-5.39	100.85	105.70
35	BB	1334	C	C5'-C4'-O4'	5.39	115.57	109.10
35	BB	1373	U	N1-C2-O2	5.39	126.57	122.80
38	BE	72	C	N3-C4-N4	5.39	121.77	118.00
38	BE	104	G	O4'-C4'-C3'	-5.39	98.61	104.00
39	BF	6	C	C3'-C2'-C1'	-5.39	97.19	101.50
40	BG	95	U	C5'-C4'-C3'	-5.39	107.38	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	BN	149	LYS	N-CA-C	-5.39	96.45	111.00
64	Be	196	TRP	CB-CG-CD2	-5.39	119.60	126.60
85	AA	357	C	N3-C4-C5	-5.39	119.75	121.90
85	AA	1126	G	N1-C2-N2	-5.39	111.35	116.20
24	AQ	41	ARG	CG-CD-NE	-5.39	100.49	111.80
34	BA	292	C	O5'-C5'-C4'	-5.39	101.47	111.70
34	BA	630	U	C4'-C3'-C2'	-5.39	97.21	102.60
34	BA	1201	G	O4'-C1'-N9	5.39	112.51	108.20
35	BB	278	U	O3'-P-O5'	-5.39	93.77	104.00
35	BB	506	G	C5-C6-O6	5.39	131.83	128.60
35	BB	511	A	C8-N9-C4	-5.39	103.64	105.80
35	BB	994	A	C5'-C4'-C3'	-5.39	107.38	116.00
39	BF	35	C	O4'-C1'-N1	5.39	112.51	108.20
42	BI	176	ARG	NE-CZ-NH1	5.39	122.99	120.30
48	BO	208	MET	N-CA-C	-5.39	96.46	111.00
85	AA	110	U	C2'-C3'-O3'	5.39	122.32	113.70
85	AA	216	U	C6-N1-C1'	-5.39	113.66	121.20
85	AA	724	A	O4'-C1'-N9	5.39	112.51	108.20
85	AA	1356	U	C2-N1-C1'	5.39	124.16	117.70
85	AA	1472	G	C8-N9-C1'	5.39	134.00	127.00
85	AA	1491	G	C8-N9-C4	5.39	108.56	106.40
85	AA	1668	G	N1-C6-O6	-5.39	116.67	119.90
85	AA	1975	G	N7-C8-N9	5.39	115.79	113.10
85	AA	2031	C	O4'-C1'-N1	5.39	112.51	108.20
5	A4	40	ARG	NE-CZ-NH1	5.38	122.99	120.30
20	AL	111	ARG	NE-CZ-NH1	5.38	122.99	120.30
34	BA	310	C	N1-C1'-C2'	-5.38	106.08	112.00
34	BA	653	U	C1'-O4'-C4'	-5.38	105.59	109.90
34	BA	679	U	C4'-C3'-C2'	5.38	107.98	102.60
34	BA	758	G	C5'-C4'-C3'	-5.38	107.39	116.00
34	BA	809	U	O3'-P-O5'	-5.38	93.77	104.00
34	BA	917	C	C3'-C2'-C1'	-5.38	97.19	101.50
34	BA	1176	C	C4'-C3'-O3'	-5.38	98.09	109.40
34	BA	1269	C	P-O5'-C5'	-5.38	112.28	120.90
34	BA	1299	G	C5'-C4'-C3'	5.38	124.62	116.00
34	BA	1665	G	C5-C6-N1	5.38	114.19	111.50
34	BA	1691	G	O4'-C1'-N9	5.38	112.51	108.20
35	BB	423	G	C4-N9-C1'	-5.38	119.50	126.50
35	BB	657	A	O5'-C5'-C4'	-5.38	101.47	111.70
35	BB	1172	U	C4'-C3'-C2'	5.38	107.98	102.60
36	BC	129	C	C5-C6-N1	5.38	123.69	121.00
40	BG	28	A	N7-C8-N9	-5.38	111.11	113.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	81	G	O5'-C5'-C4'	-5.38	101.47	111.70
68	Bi	50	GLN	N-CA-C	5.38	125.54	111.00
85	AA	159	G	N3-C2-N2	-5.38	116.13	119.90
85	AA	283	A	C4'-C3'-C2'	-5.38	97.22	102.60
85	AA	313	A	P-O5'-C5'	-5.38	112.28	120.90
85	AA	355	G	C1'-O4'-C4'	-5.38	105.59	109.90
85	AA	480	U	C6-N1-C1'	5.38	128.74	121.20
85	AA	769	C	O5'-C5'-C4'	-5.38	101.47	111.70
85	AA	882	C	P-O3'-C3'	5.38	126.16	119.70
85	AA	1897	A	P-O3'-C3'	5.38	126.16	119.70
85	AA	2125	A	C4'-C3'-O3'	-5.38	98.09	109.40
19	AK	144	PHE	CA-CB-CG	5.38	126.82	113.90
34	BA	664	C	C2-N1-C1'	-5.38	112.88	118.80
34	BA	1651	C	C2'-C3'-O3'	5.38	122.31	113.70
34	BA	1806	A	C4'-C3'-C2'	-5.38	97.22	102.60
40	BG	111	C	O4'-C1'-N1	5.38	112.51	108.20
40	BG	153	C	C6-N1-C1'	5.38	127.26	120.80
56	BW	26	ASN	CA-CB-CG	-5.38	101.56	113.40
85	AA	781	G	O4'-C1'-N9	5.38	112.51	108.20
85	AA	835	C	C6-N1-C1'	5.38	127.26	120.80
85	AA	1532	G	C5-C6-O6	-5.38	125.37	128.60
15	AG	69	ARG	CG-CD-NE	-5.38	100.50	111.80
33	AZ	41	GLY	N-CA-C	-5.38	99.64	113.10
34	BA	372	U	C2-N3-C4	-5.38	123.77	127.00
34	BA	575	U	O4'-C4'-C3'	-5.38	98.62	104.00
34	BA	734	G	C6-N1-C2	-5.38	121.87	125.10
34	BA	1048	C	O4'-C1'-N1	5.38	112.50	108.20
34	BA	1642	A	P-O3'-C3'	-5.38	113.24	119.70
34	BA	1808	A	N9-C4-C5	5.38	107.95	105.80
35	BB	445	G	P-O3'-C3'	-5.38	113.24	119.70
35	BB	600	C	C5-C4-N4	5.38	123.97	120.20
35	BB	671	A	C3'-C2'-C1'	5.38	105.81	101.50
35	BB	737	C	C6-N1-C2	-5.38	118.15	120.30
40	BG	86	U	O4'-C4'-C3'	-5.38	98.62	104.00
77	Br	99	ARG	N-CA-CB	5.38	120.29	110.60
85	AA	258	G	P-O3'-C3'	-5.38	113.24	119.70
85	AA	300	C	C2'-C3'-O3'	5.38	122.31	113.70
85	AA	409	C	N3-C2-O2	-5.38	118.13	121.90
85	AA	526	G	C8-N9-C4	-5.38	104.25	106.40
85	AA	1523	G	O3'-P-O5'	5.38	114.23	104.00
85	AA	1703	A	P-O3'-C3'	5.38	126.16	119.70
85	AA	2027	U	N3-C2-O2	-5.38	118.43	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	370	U	N3-C2-O2	-5.38	118.43	122.20
34	BA	1461	A	C1'-O4'-C4'	-5.38	105.60	109.90
35	BB	1293	C	C5'-C4'-C3'	-5.38	107.39	116.00
41	BH	74	G	N3-C2-N2	5.38	123.67	119.90
6	A5	72	SER	N-CA-C	-5.38	96.48	111.00
13	AE	105	ASN	CA-CB-CG	-5.38	101.57	113.40
34	BA	149	G	P-O3'-C3'	-5.38	113.25	119.70
34	BA	254	U	N1-C1'-C2'	-5.38	106.08	112.00
34	BA	432	A	C2-N3-C4	-5.38	107.91	110.60
34	BA	1540	C	P-O5'-C5'	-5.38	112.30	120.90
34	BA	1600	G	C6-C5-N7	5.38	133.63	130.40
34	BA	1627	U	C4'-C3'-C2'	5.38	107.98	102.60
35	BB	42	A	P-O3'-C3'	-5.38	113.25	119.70
35	BB	144	G	C5-C6-O6	-5.38	125.37	128.60
35	BB	545	C	C1'-O4'-C4'	5.38	114.20	109.90
35	BB	1045	G	C4-N9-C1'	-5.38	119.51	126.50
35	BB	1187	G	C3'-C2'-C1'	-5.38	97.20	101.50
35	BB	1357	C	O4'-C1'-N1	5.38	112.50	108.20
35	BB	1456	G	N1-C6-O6	5.38	123.13	119.90
36	BC	18	G	C8-N9-C1'	5.38	133.99	127.00
38	BE	48	G	O5'-C5'-C4'	-5.38	101.48	111.70
39	BF	45	G	C3'-C2'-C1'	-5.38	97.20	101.50
40	BG	38	A	N7-C8-N9	-5.38	111.11	113.80
40	BG	52	A	C3'-C2'-C1'	-5.38	97.20	101.50
40	BG	88	G	N3-C4-C5	-5.38	125.91	128.60
41	BH	59	G	C8-N9-C4	-5.38	104.25	106.40
42	BI	105	LEU	N-CA-C	5.38	125.52	111.00
56	BW	17	LEU	C-N-CA	5.38	135.14	121.70
67	Bh	29	ALA	N-CA-C	-5.38	96.48	111.00
85	AA	329	G	C5'-C4'-O4'	5.38	115.56	109.10
85	AA	470	C	O4'-C1'-N1	5.38	112.50	108.20
85	AA	787	U	O4'-C4'-C3'	5.38	110.40	106.10
85	AA	904	U	P-O3'-C3'	-5.38	113.25	119.70
85	AA	1464	G	C4-N9-C1'	-5.38	119.51	126.50
14	AF	116	ARG	CG-CD-NE	-5.38	100.51	111.80
34	BA	351	A	O4'-C1'-N9	5.38	112.50	108.20
34	BA	386	A	O5'-C5'-C4'	5.38	121.92	111.70
34	BA	429	G	C5'-C4'-C3'	-5.38	107.40	116.00
34	BA	805	A	OP1-P-O3'	5.38	117.03	105.20
34	BA	999	G	C3'-C2'-C1'	-5.38	97.20	101.50
34	BA	1325	G	C1'-O4'-C4'	-5.38	105.60	109.90
34	BA	1496	G	C5-C6-N1	5.38	114.19	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1509	U	O3'-P-O5'	5.38	114.22	104.00
35	BB	14	C	C1'-O4'-C4'	-5.38	105.60	109.90
35	BB	141	G	O4'-C1'-N9	5.38	112.50	108.20
35	BB	835	C	C2-N3-C4	-5.38	117.21	119.90
35	BB	857	G	P-O3'-C3'	-5.38	113.25	119.70
37	BD	31	U	C3'-C2'-C1'	-5.38	97.20	101.50
41	BH	13	C	P-O5'-C5'	-5.38	112.30	120.90
42	BI	27	LYS	N-CA-CB	-5.38	100.92	110.60
78	Bs	20	CYS	CB-CA-C	-5.38	99.65	110.40
85	AA	490	A	C5'-C4'-O4'	5.38	115.55	109.10
85	AA	660	G	N3-C2-N2	5.38	123.66	119.90
85	AA	800	A	N9-C4-C5	5.38	107.95	105.80
85	AA	859	G	P-O3'-C3'	-5.38	113.25	119.70
85	AA	1197	U	C5-C4-O4	5.38	129.13	125.90
85	AA	1557	U	C6-N1-C2	-5.38	117.77	121.00
85	AA	2093	U	C4'-C3'-C2'	-5.38	97.22	102.60
85	AA	2199	G	O4'-C4'-C3'	-5.38	98.62	104.00
86	AB	69	G	C5-C6-O6	-5.38	125.37	128.60
35	BB	382	U	C4'-C3'-C2'	-5.38	97.22	102.60
35	BB	1144	A	P-O3'-C3'	-5.38	113.25	119.70
40	BG	87	G	C1'-O4'-C4'	-5.38	105.60	109.90
41	BH	85	C	C2'-C3'-O3'	5.38	122.30	113.70
61	Bb	45	PHE	CA-CB-CG	-5.38	101.00	113.90
85	AA	1483	A	C5-C6-N1	5.38	120.39	117.70
7	A6	158	ALA	N-CA-C	-5.37	96.49	111.00
34	BA	135	G	O4'-C4'-C3'	-5.37	98.63	104.00
34	BA	486	G	N1-C6-O6	5.37	123.12	119.90
34	BA	487	A	OP2-P-O3'	5.37	117.02	105.20
34	BA	575	U	N1-C2-O2	5.37	126.56	122.80
34	BA	1173	C	O4'-C1'-N1	5.37	112.50	108.20
34	BA	1177	C	C2-N3-C4	5.37	122.59	119.90
34	BA	1228	G	C5-C6-O6	-5.37	125.38	128.60
34	BA	1707	C	O4'-C1'-C2'	5.37	112.44	107.60
34	BA	1784	G	O4'-C1'-N9	5.37	112.50	108.20
35	BB	528	G	N9-C1'-C2'	-5.37	106.09	112.00
35	BB	684	U	O4'-C1'-N1	5.37	112.50	108.20
35	BB	852	G	N1-C2-N2	-5.37	111.36	116.20
35	BB	977	G	C2-N3-C4	-5.37	109.21	111.90
35	BB	1051	U	N1-C2-O2	5.37	126.56	122.80
38	BE	183	C	C5-C6-N1	5.37	123.69	121.00
39	BF	56	C	N3-C4-N4	5.37	121.76	118.00
41	BH	13	C	C5-C4-N4	-5.37	116.44	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	94	C	C5'-C4'-C3'	-5.37	107.40	116.00
85	AA	251	A	C8-N9-C1'	-5.37	118.03	127.70
85	AA	327	G	N3-C4-N9	5.37	129.22	126.00
85	AA	1450	U	C2-N3-C4	-5.37	123.78	127.00
85	AA	1456	A	O4'-C4'-C3'	-5.37	98.63	104.00
85	AA	1845	G	N9-C1'-C2'	-5.37	106.09	112.00
85	AA	1953	G	C5-C6-O6	-5.37	125.38	128.60
85	AA	1979	A	P-O3'-C3'	5.37	126.15	119.70
34	BA	306	G	N3-C2-N2	5.37	123.66	119.90
34	BA	500	C	N3-C2-O2	-5.37	118.14	121.90
34	BA	517	A	N9-C4-C5	-5.37	103.65	105.80
34	BA	776	U	C2-N1-C1'	-5.37	111.25	117.70
34	BA	961	C	C5'-C4'-O4'	5.37	115.55	109.10
34	BA	1019	C	C5'-C4'-O4'	5.37	115.55	109.10
34	BA	1184	A	N9-C4-C5	5.37	107.95	105.80
34	BA	1278	A	C3'-C2'-C1'	-5.37	97.20	101.50
34	BA	1566	G	C5-C6-N1	5.37	114.19	111.50
34	BA	1739	G	C8-N9-C4	-5.37	104.25	106.40
35	BB	65	A	C5-C6-N1	5.37	120.39	117.70
35	BB	495	A	C5'-C4'-O4'	5.37	115.55	109.10
35	BB	978	C	C3'-C2'-C1'	-5.37	97.20	101.50
35	BB	1405	G	P-O3'-C3'	-5.37	113.25	119.70
35	BB	1464	G	N9-C1'-C2'	5.37	120.98	114.00
35	BB	1512	C	N3-C2-O2	-5.37	118.14	121.90
38	BE	32	U	C4-C5-C6	-5.37	116.48	119.70
38	BE	118	C	C3'-C2'-C1'	-5.37	97.20	101.50
85	AA	421	G	C5-C6-O6	5.37	131.82	128.60
85	AA	589	A	O5'-C5'-C4'	5.37	121.91	111.70
85	AA	731	U	O5'-C5'-C4'	-5.37	101.50	111.70
85	AA	1274	A	N1-C6-N6	-5.37	115.38	118.60
85	AA	1465	C	C3'-C2'-C1'	-5.37	97.20	101.50
85	AA	1799	C	C4'-C3'-C2'	-5.37	97.23	102.60
85	AA	2083	G	C6-N1-C2	-5.37	121.88	125.10
85	AA	2193	A	N1-C6-N6	5.37	121.82	118.60
85	AA	2204	A	C4-N9-C1'	-5.37	116.63	126.30
34	BA	223	U	N3-C4-C5	5.37	117.82	114.60
35	BB	527	U	C5-C6-N1	5.37	125.39	122.70
35	BB	583	G	P-O3'-C3'	-5.37	113.26	119.70
35	BB	677	U	P-O5'-C5'	-5.37	112.31	120.90
65	Bf	320	VAL	N-CA-C	-5.37	96.50	111.00
65	Bf	346	SER	CA-C-O	-5.37	108.82	120.10
74	Bo	50	LYS	N-CA-C	-5.37	96.50	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	488	G	O3'-P-O5'	5.37	114.20	104.00
85	AA	1846	G	N3-C2-N2	5.37	123.66	119.90
85	AA	1880	C	P-O3'-C3'	5.37	126.14	119.70
16	AH	59	ARG	NE-CZ-NH2	-5.37	117.62	120.30
34	BA	464	U	P-O3'-C3'	-5.37	113.26	119.70
34	BA	1084	A	C8-N9-C4	-5.37	103.65	105.80
34	BA	1240	G	C4-N9-C1'	5.37	133.48	126.50
35	BB	413	A	P-O5'-C5'	-5.37	112.31	120.90
35	BB	850	U	O5'-C5'-C4'	5.37	121.90	111.70
35	BB	1113	C	C5-C4-N4	5.37	123.96	120.20
35	BB	1205	A	C8-N9-C4	5.37	107.95	105.80
35	BB	1230	A	C3'-C2'-C1'	-5.37	97.20	101.50
35	BB	1505	U	N1-C1'-C2'	-5.37	106.09	112.00
36	BC	104	A	P-O5'-C5'	-5.37	112.31	120.90
38	BE	1	U	N1-C2-N3	5.37	118.12	114.90
40	BG	4	A	O3'-P-O5'	-5.37	93.80	104.00
40	BG	16	G	C4-N9-C1'	-5.37	119.52	126.50
40	BG	17	A	C5-C6-N6	-5.37	119.41	123.70
47	BN	112	GLU	N-CA-CB	-5.37	100.94	110.60
47	BN	120	ARG	CB-CA-C	-5.37	99.66	110.40
85	AA	369	A	N7-C8-N9	-5.37	111.11	113.80
85	AA	1164	A	C3'-C2'-C1'	-5.37	97.20	101.50
85	AA	2108	C	N1-C1'-C2'	-5.37	106.09	112.00
34	BA	380	A	C5'-C4'-O4'	5.37	115.54	109.10
34	BA	771	A	N9-C1'-C2'	5.37	120.98	114.00
34	BA	1137	U	C3'-C2'-C1'	-5.37	97.21	101.50
34	BA	1181	G	O3'-P-O5'	5.37	114.20	104.00
34	BA	1744	C	C6-N1-C1'	5.37	127.24	120.80
35	BB	993	A	OP2-P-O3'	5.37	117.01	105.20
35	BB	1155	U	N1-C1'-C2'	-5.37	106.10	112.00
35	BB	1419	G	N1-C6-O6	5.37	123.12	119.90
85	AA	642	G	P-O3'-C3'	-5.37	113.26	119.70
85	AA	2059	A	C3'-C2'-C1'	-5.37	97.21	101.50
85	AA	2235	C	N1-C2-N3	5.37	122.96	119.20
1	A0	153	SER	C-N-CA	5.37	135.11	121.70
34	BA	38	G	N1-C2-N2	-5.37	111.37	116.20
34	BA	406	G	C4-N9-C1'	-5.37	119.52	126.50
34	BA	619	U	OP2-P-O3'	5.37	117.00	105.20
34	BA	1213	A	C1'-O4'-C4'	-5.37	105.61	109.90
34	BA	1320	A	C6-N1-C2	-5.37	115.38	118.60
34	BA	1369	C	C5-C4-N4	-5.37	116.44	120.20
35	BB	606	C	C4'-C3'-C2'	5.37	107.97	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	805	G	N1-C6-O6	5.37	123.12	119.90
35	BB	1448	U	C1'-O4'-C4'	-5.37	105.61	109.90
36	BC	153	C	C6-N1-C1'	-5.37	114.36	120.80
37	BD	19	C	C5-C6-N1	5.37	123.68	121.00
37	BD	53	U	C2-N3-C4	-5.37	123.78	127.00
41	BH	14	C	P-O3'-C3'	5.37	126.14	119.70
41	BH	43	G	C5'-C4'-O4'	-5.37	102.66	109.10
80	Bu	126	ASP	CA-CB-CG	-5.37	101.59	113.40
85	AA	98	U	P-O5'-C5'	5.37	129.48	120.90
85	AA	161	A	C1'-O4'-C4'	-5.37	105.61	109.90
85	AA	415	G	O3'-P-O5'	5.37	114.19	104.00
85	AA	472	A	C3'-C2'-C1'	-5.37	97.21	101.50
85	AA	863	C	P-O3'-C3'	-5.37	113.26	119.70
85	AA	2101	C	O4'-C1'-N1	5.37	112.49	108.20
85	AA	2149	C	C3'-C2'-C1'	-5.37	97.21	101.50
85	AA	2203	C	O4'-C1'-C2'	5.37	112.43	107.60
86	AB	59	U	C6-N1-C1'	-5.37	113.69	121.20
5	A4	122	THR	N-CA-CB	5.36	120.49	110.30
34	BA	232	U	C2-N3-C4	-5.36	123.78	127.00
34	BA	403	A	O4'-C1'-C2'	-5.36	100.44	105.80
34	BA	734	G	C4-N9-C1'	-5.36	119.53	126.50
34	BA	799	A	N7-C8-N9	5.36	116.48	113.80
34	BA	941	G	O3'-P-O5'	-5.36	93.81	104.00
35	BB	463	C	C5'-C4'-C3'	-5.36	107.42	116.00
35	BB	561	C	C5'-C4'-O4'	5.36	115.54	109.10
35	BB	1108	G	N1-C2-N2	-5.36	111.37	116.20
35	BB	1365	G	C4'-C3'-C2'	-5.36	97.24	102.60
35	BB	1521	G	C5'-C4'-C3'	-5.36	107.42	116.00
35	BB	1539	C	C5'-C4'-C3'	5.36	124.58	116.00
38	BE	206	G	C5-C6-O6	-5.36	125.38	128.60
40	BG	12	A	C8-N9-C4	-5.36	103.65	105.80
41	BH	43	G	C6-N1-C2	-5.36	121.88	125.10
72	Bm	97	LYS	N-CA-CB	-5.36	100.95	110.60
85	AA	638	G	C5-C6-O6	-5.36	125.38	128.60
85	AA	745	C	N3-C2-O2	-5.36	118.15	121.90
85	AA	1142	G	C4-N9-C1'	-5.36	119.53	126.50
85	AA	1286	C	N3-C2-O2	-5.36	118.15	121.90
85	AA	1389	G	O4'-C1'-N9	5.36	112.49	108.20
86	AB	34	G	C5-C6-O6	-5.36	125.38	128.60
34	BA	95	C	C2-N1-C1'	-5.36	112.90	118.80
34	BA	1357	C	C6-N1-C2	-5.36	118.16	120.30
35	BB	403	U	C6-N1-C2	-5.36	117.78	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	493	U	C2-N3-C4	-5.36	123.78	127.00
35	BB	1110	G	C8-N9-C4	-5.36	104.25	106.40
35	BB	1416	A	O3'-P-O5'	5.36	114.19	104.00
36	BC	138	C	N3-C2-O2	-5.36	118.15	121.90
40	BG	77	U	C4'-C3'-C2'	-5.36	97.24	102.60
41	BH	32	U	P-O3'-C3'	-5.36	113.27	119.70
65	Bf	425	HIS	CA-CB-CG	-5.36	104.48	113.60
85	AA	149	A	C4-N9-C1'	-5.36	116.65	126.30
85	AA	450	A	N9-C1'-C2'	-5.36	106.10	112.00
85	AA	685	U	C5'-C4'-O4'	5.36	115.53	109.10
34	BA	260	A	O3'-P-O5'	5.36	114.18	104.00
34	BA	341	U	C5-C6-N1	-5.36	120.02	122.70
34	BA	875	G	C4-C5-C6	-5.36	115.58	118.80
34	BA	1309	U	C4'-C3'-C2'	-5.36	97.24	102.60
34	BA	1414	C	O4'-C1'-N1	5.36	112.49	108.20
35	BB	47	C	N3-C4-N4	-5.36	114.25	118.00
35	BB	416	U	C4-C5-C6	-5.36	116.48	119.70
35	BB	904	C	O4'-C1'-N1	5.36	112.49	108.20
35	BB	1251	G	C1'-O4'-C4'	-5.36	105.61	109.90
37	BD	68	C	C5'-C4'-C3'	-5.36	107.42	116.00
38	BE	186	C	C6-N1-C1'	-5.36	114.37	120.80
50	BQ	72	THR	N-CA-C	5.36	125.47	111.00
82	Bw	251	ARG	CD-NE-CZ	-5.36	116.09	123.60
85	AA	239	G	C4-N9-C1'	5.36	133.47	126.50
85	AA	296	A	P-O3'-C3'	-5.36	113.27	119.70
85	AA	764	U	C5-C6-N1	-5.36	120.02	122.70
85	AA	1540	A	C4'-C3'-C2'	-5.36	97.24	102.60
85	AA	1582	U	C2-N3-C4	-5.36	123.78	127.00
85	AA	1654	G	O4'-C1'-N9	5.36	112.49	108.20
86	AB	49	C	C6-N1-C2	-5.36	118.16	120.30
4	A3	102	GLY	C-N-CA	5.36	135.10	121.70
29	AV	5	ARG	NE-CZ-NH2	-5.36	117.62	120.30
34	BA	1644	A	O5'-P-OP1	-5.36	100.88	105.70
34	BA	1830	A	P-O5'-C5'	-5.36	112.33	120.90
35	BB	873	C	C1'-O4'-C4'	-5.36	105.61	109.90
35	BB	874	G	P-O3'-C3'	-5.36	113.27	119.70
38	BE	127	G	C5'-C4'-O4'	-5.36	102.67	109.10
40	BG	82	U	P-O3'-C3'	-5.36	113.27	119.70
85	AA	400	G	C5-N7-C8	-5.36	101.62	104.30
85	AA	737	G	O4'-C1'-N9	5.36	112.49	108.20
85	AA	820	G	O5'-P-OP2	5.36	117.13	110.70
85	AA	1440	C	OP1-P-O3'	5.36	116.99	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1801	U	O4'-C1'-N1	5.36	112.49	108.20
34	BA	78	U	P-O5'-C5'	-5.36	112.33	120.90
34	BA	440	A	C5-C6-N6	5.36	127.99	123.70
34	BA	458	G	O5'-C5'-C4'	-5.36	101.52	111.70
34	BA	571	G	N9-C1'-C2'	5.36	120.96	114.00
34	BA	753	G	P-O3'-C3'	-5.36	113.27	119.70
34	BA	859	G	C6-N1-C2	-5.36	121.89	125.10
34	BA	1067	G	N7-C8-N9	-5.36	110.42	113.10
34	BA	1092	U	P-O5'-C5'	-5.36	112.33	120.90
34	BA	1177	C	O4'-C1'-C2'	5.36	112.42	107.60
34	BA	1277	G	O4'-C1'-N9	5.36	112.49	108.20
34	BA	1321	A	C4'-C3'-C2'	-5.36	97.24	102.60
35	BB	72	G	O4'-C1'-C2'	5.36	112.42	107.60
35	BB	474	G	N1-C6-O6	5.36	123.11	119.90
35	BB	561	C	C5'-C4'-C3'	5.36	124.57	116.00
35	BB	734	A	C4-N9-C1'	-5.36	116.66	126.30
35	BB	787	A	P-O3'-C3'	-5.36	113.27	119.70
35	BB	1185	G	P-O3'-C3'	-5.36	113.27	119.70
35	BB	1210	U	C5-C4-O4	-5.36	122.69	125.90
38	BE	24	G	N1-C6-O6	5.36	123.11	119.90
38	BE	127	G	N1-C2-N3	-5.36	120.69	123.90
39	BF	10	A	C6-C5-N7	-5.36	128.55	132.30
39	BF	13	U	O5'-C5'-C4'	-5.36	101.52	111.70
40	BG	128	U	N1-C2-O2	5.36	126.55	122.80
41	BH	130	G	C1'-O4'-C4'	-5.36	105.61	109.90
50	BQ	173	HIS	CA-CB-CG	-5.36	104.49	113.60
58	BY	37	ARG	CB-CG-CD	5.36	125.53	111.60
85	AA	30	G	C8-N9-C1'	5.36	133.96	127.00
85	AA	464	A	N1-C6-N6	5.36	121.81	118.60
85	AA	902	A	C8-N9-C1'	5.36	137.34	127.70
85	AA	1192	C	O4'-C1'-N1	5.36	112.49	108.20
85	AA	1997	G	P-O5'-C5'	5.36	129.47	120.90
85	AA	2096	G	N1-C6-O6	5.36	123.11	119.90
4	A3	219	ARG	NE-CZ-NH1	5.36	122.98	120.30
4	A3	222	ALA	N-CA-C	5.36	125.46	111.00
34	BA	455	A	C1'-O4'-C4'	-5.36	105.61	109.90
34	BA	1300	G	C2'-C3'-O3'	5.36	122.27	113.70
34	BA	1311	G	O4'-C1'-C2'	5.36	112.42	107.60
34	BA	1632	G	C3'-C2'-C1'	5.36	105.78	101.50
35	BB	312	U	O4'-C1'-N1	5.36	112.48	108.20
35	BB	463	C	C3'-C2'-C1'	-5.36	97.22	101.50
35	BB	813	C	C6-N1-C2	-5.36	118.16	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	998	G	C6-N1-C2	-5.36	121.89	125.10
35	BB	1055	G	OP2-P-O3'	5.36	116.98	105.20
38	BE	117	A	C4'-C3'-C2'	-5.36	97.25	102.60
48	BO	166	TRP	CA-C-N	-5.36	105.42	117.20
77	Br	176	ILE	CB-CA-C	-5.36	100.89	111.60
77	Br	196	ARG	NE-CZ-NH1	-5.36	117.62	120.30
85	AA	207	G	N1-C6-O6	5.36	123.11	119.90
85	AA	368	C	P-O5'-C5'	-5.36	112.33	120.90
85	AA	413	G	N1-C6-O6	-5.36	116.69	119.90
85	AA	674	U	O5'-P-OP1	-5.36	100.88	105.70
85	AA	880	A	C1'-O4'-C4'	-5.36	105.62	109.90
1	A0	144	PHE	CB-CA-C	-5.35	99.69	110.40
34	BA	5	C	C1'-O4'-C4'	-5.35	105.62	109.90
34	BA	1033	G	O4'-C1'-N9	5.35	112.48	108.20
34	BA	1537	G	C2'-C3'-O3'	5.35	122.27	113.70
34	BA	1831	A	N9-C1'-C2'	5.35	120.96	114.00
35	BB	611	U	C6-N1-C1'	5.35	128.69	121.20
36	BC	134	G	N7-C8-N9	5.35	115.78	113.10
42	BI	49	HIS	N-CA-CB	-5.35	100.96	110.60
85	AA	20	G	O4'-C1'-N9	5.35	112.48	108.20
85	AA	63	G	O4'-C1'-N9	-5.35	103.92	108.20
85	AA	771	A	N9-C4-C5	5.35	107.94	105.80
85	AA	1437	G	O4'-C1'-C2'	5.35	112.42	107.60
85	AA	1806	C	O4'-C1'-N1	5.35	112.48	108.20
34	BA	1463	U	N3-C2-O2	-5.35	118.45	122.20
34	BA	1835	A	P-O3'-C3'	-5.35	113.28	119.70
35	BB	148	C	O4'-C1'-N1	5.35	112.48	108.20
35	BB	769	C	P-O5'-C5'	5.35	129.47	120.90
35	BB	772	U	C6-N1-C2	-5.35	117.79	121.00
35	BB	1144	A	C5'-C4'-O4'	5.35	115.52	109.10
35	BB	1476	C	C5'-C4'-C3'	5.35	124.56	116.00
38	BE	7	U	C5'-C4'-O4'	5.35	115.52	109.10
38	BE	197	A	N1-C6-N6	-5.35	115.39	118.60
45	BL	104	SER	N-CA-CB	5.35	118.53	110.50
70	Bk	79	LEU	C-N-CA	5.35	135.08	121.70
77	Br	246	ARG	CG-CD-NE	-5.35	100.56	111.80
85	AA	68	A	C5'-C4'-O4'	5.35	115.52	109.10
85	AA	177	A	P-O3'-C3'	-5.35	113.28	119.70
85	AA	452	A	C4-N9-C1'	-5.35	116.67	126.30
85	AA	708	G	N3-C4-N9	-5.35	122.79	126.00
85	AA	935	A	C1'-O4'-C4'	-5.35	105.62	109.90
85	AA	976	G	N1-C6-O6	-5.35	116.69	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1482	C	O4'-C1'-N1	5.35	112.48	108.20
85	AA	1633	A	C5-C6-N6	-5.35	119.42	123.70
85	AA	1755	U	C4'-C3'-C2'	5.35	107.95	102.60
85	AA	1919	G	C5'-C4'-O4'	5.35	115.52	109.10
85	AA	2018	U	N1-C2-N3	5.35	118.11	114.90
3	A2	42	TRP	C-N-CA	5.35	135.08	121.70
6	A5	36	THR	CB-CA-C	-5.35	97.15	111.60
34	BA	20	A	N1-C6-N6	5.35	121.81	118.60
34	BA	62	A	O4'-C1'-N9	5.35	112.48	108.20
34	BA	399	G	P-O3'-C3'	-5.35	113.28	119.70
34	BA	616	G	C4'-C3'-C2'	-5.35	97.25	102.60
34	BA	1542	A	C4'-C3'-C2'	5.35	107.95	102.60
34	BA	1668	C	O4'-C1'-C2'	5.35	112.42	107.60
35	BB	452	A	C5-C6-N1	5.35	120.38	117.70
35	BB	777	C	P-O5'-C5'	5.35	129.46	120.90
35	BB	1062	G	C3'-C2'-C1'	5.35	105.78	101.50
38	BE	74	U	C5-C4-O4	-5.35	122.69	125.90
38	BE	91	G	C5-C6-N1	5.35	114.18	111.50
41	BH	73	A	P-O3'-C3'	5.35	126.12	119.70
85	AA	337	C	OP2-P-O3'	5.35	116.97	105.20
85	AA	1059	C	O4'-C1'-N1	5.35	112.48	108.20
85	AA	1564	U	C2-N1-C1'	-5.35	111.28	117.70
24	AQ	112	THR	N-CA-CB	5.35	120.47	110.30
34	BA	34	U	O4'-C1'-N1	5.35	112.48	108.20
34	BA	460	G	N3-C2-N2	5.35	123.64	119.90
34	BA	685	C	OP1-P-O3'	5.35	116.97	105.20
34	BA	741	A	N7-C8-N9	-5.35	111.13	113.80
34	BA	849	G	N9-C4-C5	-5.35	103.26	105.40
34	BA	919	A	O4'-C1'-N9	5.35	112.48	108.20
35	BB	600	C	N1-C2-N3	-5.35	115.46	119.20
35	BB	977	G	N3-C4-C5	-5.35	125.92	128.60
35	BB	1476	C	C6-N1-C1'	5.35	127.22	120.80
44	BK	209	TYR	CB-CG-CD1	-5.35	117.79	121.00
62	Bc	111	ARG	NH1-CZ-NH2	-5.35	113.52	119.40
65	Bf	223	ARG	NE-CZ-NH1	5.35	122.97	120.30
85	AA	97	A	N1-C6-N6	5.35	121.81	118.60
85	AA	164	G	OP1-P-OP2	-5.35	111.58	119.60
85	AA	1085	U	C2-N1-C1'	-5.35	111.28	117.70
85	AA	1660	U	C5-C6-N1	-5.35	120.03	122.70
85	AA	2016	A	O4'-C1'-C2'	5.35	112.42	107.60
85	AA	2047	U	N1-C2-N3	5.35	118.11	114.90
86	AB	21	A	C4-N9-C1'	-5.35	116.67	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	AR	83	VAL	CA-CB-CG2	-5.35	102.88	110.90
34	BA	6	C	C5'-C4'-O4'	5.35	115.52	109.10
34	BA	56	G	C4-N9-C1'	-5.35	119.55	126.50
34	BA	222	C	C3'-C2'-C1'	-5.35	97.22	101.50
34	BA	276	C	P-O3'-C3'	-5.35	113.28	119.70
34	BA	1262	A	C1'-O4'-C4'	-5.35	105.62	109.90
34	BA	1479	G	P-O5'-C5'	5.35	129.46	120.90
35	BB	329	U	C6-N1-C1'	-5.35	113.71	121.20
35	BB	793	A	C3'-C2'-C1'	-5.35	97.22	101.50
35	BB	1327	U	C5-C6-N1	-5.35	120.03	122.70
38	BE	35	A	C5'-C4'-C3'	-5.35	107.44	116.00
41	BH	32	U	P-O5'-C5'	-5.35	112.34	120.90
50	BQ	167	TRP	CB-CG-CD2	-5.35	119.65	126.60
52	BS	126	LEU	C-N-CA	-5.35	111.07	122.30
85	AA	36	U	C5-C6-N1	-5.35	120.03	122.70
85	AA	123	A	OP1-P-O3'	5.35	116.96	105.20
85	AA	927	A	N3-C4-N9	5.35	131.68	127.40
85	AA	1052	C	P-O3'-C3'	5.35	126.12	119.70
85	AA	1596	A	C1'-O4'-C4'	-5.35	105.62	109.90
85	AA	1978	G	N3-C4-C5	5.35	131.27	128.60
85	AA	2078	A	C4-C5-C6	-5.35	114.33	117.00
15	AG	104	ARG	CB-CA-C	-5.35	99.71	110.40
19	AK	104	ASN	CA-CB-CG	-5.35	101.64	113.40
34	BA	296	G	OP2-P-O3'	5.35	116.96	105.20
34	BA	567	U	P-O3'-C3'	-5.35	113.28	119.70
34	BA	704	G	C4-N9-C1'	-5.35	119.55	126.50
34	BA	1249	G	N1-C6-O6	-5.35	116.69	119.90
34	BA	1475	G	C5-C6-O6	-5.35	125.39	128.60
34	BA	1641	G	O5'-C5'-C4'	-5.35	101.54	111.70
35	BB	1511	U	C6-N1-C2	-5.35	117.79	121.00
38	BE	31	A	C5-N7-C8	-5.35	101.23	103.90
39	BF	1	C	C1'-O4'-C4'	-5.35	105.62	109.90
69	Bj	94	ARG	NE-CZ-NH2	-5.35	117.63	120.30
85	AA	488	G	N3-C4-C5	-5.35	125.93	128.60
85	AA	997	U	N3-C4-C5	-5.35	111.39	114.60
85	AA	1033	C	P-O5'-C5'	5.35	129.45	120.90
85	AA	1105	G	N7-C8-N9	5.35	115.77	113.10
85	AA	1678	U	P-O3'-C3'	-5.35	113.28	119.70
85	AA	1810	C	N3-C2-O2	-5.35	118.16	121.90
34	BA	694	G	C4-N9-C1'	5.34	133.45	126.50
34	BA	1078	U	C3'-C2'-C1'	-5.34	97.22	101.50
34	BA	1409	A	C8-N9-C1'	5.34	137.32	127.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1601	C	O4'-C1'-C2'	5.34	112.41	107.60
35	BB	456	A	P-O5'-C5'	-5.34	112.35	120.90
35	BB	1458	U	N1-C2-O2	-5.34	119.06	122.80
35	BB	1544	A	C2-N3-C4	5.34	113.27	110.60
40	BG	138	C	C5'-C4'-C3'	-5.34	107.45	116.00
73	Bn	20	ARG	NE-CZ-NH1	5.34	122.97	120.30
73	Bn	73	ARG	N-CA-CB	-5.34	100.98	110.60
85	AA	36	U	N3-C4-O4	-5.34	115.66	119.40
85	AA	314	C	O4'-C1'-N1	5.34	112.47	108.20
85	AA	427	G	O4'-C1'-N9	5.34	112.48	108.20
85	AA	538	A	C4-N9-C1'	5.34	135.92	126.30
85	AA	641	A	P-O5'-C5'	5.34	129.45	120.90
85	AA	726	U	O4'-C1'-N1	5.34	112.48	108.20
85	AA	930	G	C4-N9-C1'	5.34	133.45	126.50
85	AA	1229	G	N1-C2-N2	-5.34	111.39	116.20
85	AA	1309	G	C8-N9-C1'	5.34	133.95	127.00
34	BA	1028	A	P-O3'-C3'	-5.34	113.29	119.70
34	BA	1409	A	O4'-C1'-N9	5.34	112.47	108.20
35	BB	762	C	C5-C6-N1	5.34	123.67	121.00
85	AA	1262	A	C3'-C2'-C1'	-5.34	97.23	101.50
2	A1	10	TYR	CB-CG-CD2	-5.34	117.80	121.00
11	AC	154	GLN	N-CA-CB	-5.34	100.99	110.60
34	BA	1330	G	C2-N3-C4	5.34	114.57	111.90
34	BA	1482	A	N7-C8-N9	-5.34	111.13	113.80
34	BA	1636	C	N1-C2-N3	-5.34	115.46	119.20
34	BA	1690	U	C5-C4-O4	5.34	129.10	125.90
35	BB	412	A	C5'-C4'-O4'	5.34	115.51	109.10
35	BB	1216	G	C5-C6-O6	-5.34	125.39	128.60
35	BB	1322	A	C4-N9-C1'	-5.34	116.68	126.30
35	BB	1375	G	C8-N9-C1'	5.34	133.94	127.00
80	Bu	77	ALA	N-CA-CB	-5.34	102.62	110.10
85	AA	105	A	N1-C6-N6	-5.34	115.39	118.60
85	AA	260	A	O5'-C5'-C4'	-5.34	101.55	111.70
85	AA	545	A	O5'-C5'-C4'	-5.34	101.55	111.70
85	AA	1194	U	P-O3'-C3'	-5.34	113.29	119.70
85	AA	2206	A	C1'-O4'-C4'	-5.34	105.63	109.90
5	A4	79	ILE	C-N-CA	5.34	135.05	121.70
34	BA	52	G	N9-C1'-C2'	-5.34	106.13	112.00
34	BA	196	A	O4'-C4'-C3'	-5.34	98.66	104.00
34	BA	1012	A	P-O3'-C3'	5.34	126.11	119.70
34	BA	1134	A	C3'-C2'-C1'	-5.34	97.23	101.50
34	BA	1230	G	N1-C6-O6	-5.34	116.70	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1257	U	O5'-P-OP2	-5.34	100.89	105.70
34	BA	1425	G	C5'-C4'-C3'	-5.34	107.46	116.00
35	BB	4	C	C2-N1-C1'	5.34	124.67	118.80
35	BB	832	C	C1'-O4'-C4'	-5.34	105.63	109.90
35	BB	921	U	O4'-C1'-N1	5.34	112.47	108.20
35	BB	1216	G	C3'-C2'-C1'	-5.34	97.23	101.50
35	BB	1456	G	C5-C6-O6	-5.34	125.40	128.60
36	BC	135	A	C3'-C2'-C1'	-5.34	97.23	101.50
37	BD	51	G	C5-C6-O6	-5.34	125.40	128.60
38	BE	114	G	C8-N9-C4	5.34	108.54	106.40
41	BH	54	U	N1-C2-O2	5.34	126.54	122.80
72	Bm	71	VAL	C-N-CA	5.34	133.51	122.30
85	AA	402	G	C8-N9-C1'	5.34	133.94	127.00
85	AA	1097	G	C3'-C2'-C1'	-5.34	97.23	101.50
85	AA	1321	G	O4'-C1'-N9	5.34	112.47	108.20
85	AA	2198	G	P-O5'-C5'	-5.34	112.36	120.90
85	AA	2231	G	O4'-C4'-C3'	-5.34	98.66	104.00
34	BA	721	A	C4'-C3'-C2'	-5.34	97.26	102.60
34	BA	1153	C	N3-C4-N4	5.34	121.74	118.00
34	BA	1235	C	C5'-C4'-C3'	-5.34	107.46	116.00
35	BB	108	G	C5-C6-O6	-5.34	125.40	128.60
35	BB	1507	U	O4'-C4'-C3'	-5.34	98.66	104.00
36	BC	115	G	P-O5'-C5'	5.34	129.44	120.90
38	BE	160	C	C2-N1-C1'	-5.34	112.93	118.80
41	BH	21	G	O4'-C1'-N9	5.34	112.47	108.20
54	BU	35	LYS	N-CA-C	-5.34	96.59	111.00
85	AA	1469	G	C6-N1-C2	-5.34	121.90	125.10
85	AA	1565	G	P-O3'-C3'	-5.34	113.30	119.70
85	AA	2218	G	C5-N7-C8	5.34	106.97	104.30
34	BA	553	A	O4'-C1'-N9	5.34	112.47	108.20
34	BA	667	U	O3'-P-O5'	5.34	114.14	104.00
34	BA	774	A	C5-C6-N6	5.34	127.97	123.70
34	BA	1065	U	C2-N3-C4	-5.34	123.80	127.00
34	BA	1117	G	C4'-C3'-C2'	5.34	107.94	102.60
34	BA	1156	U	C6-N1-C1'	5.34	128.67	121.20
34	BA	1251	A	C8-N9-C1'	5.34	137.30	127.70
34	BA	1799	G	C5-C6-N1	5.34	114.17	111.50
35	BB	14	C	O4'-C4'-C3'	-5.34	98.66	104.00
35	BB	412	A	C2-N3-C4	-5.34	107.93	110.60
35	BB	760	C	N3-C4-N4	-5.34	114.27	118.00
35	BB	1052	G	O3'-P-O5'	-5.34	93.86	104.00
35	BB	1093	C	C3'-C2'-C1'	-5.34	97.23	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1197	G	C4-N9-C1'	-5.34	119.56	126.50
35	BB	1477	C	N1-C2-O2	5.34	122.10	118.90
36	BC	148	C	P-O3'-C3'	-5.34	113.30	119.70
85	AA	114	C	C1'-O4'-C4'	-5.34	105.63	109.90
85	AA	437	G	C8-N9-C4	5.34	108.53	106.40
85	AA	455	G	O5'-C5'-C4'	-5.34	101.56	111.70
85	AA	753	U	O5'-P-OP2	5.34	117.10	110.70
85	AA	1732	G	C5'-C4'-C3'	-5.34	107.46	116.00
85	AA	1900	C	C5-C6-N1	-5.34	118.33	121.00
85	AA	1973	G	C5-C6-O6	-5.34	125.40	128.60
85	AA	2032	G	O3'-P-O5'	-5.34	93.86	104.00
34	BA	296	G	O5'-P-OP1	-5.33	100.90	105.70
34	BA	711	C	N3-C4-C5	-5.33	119.77	121.90
34	BA	728	A	C5'-C4'-C3'	5.33	124.54	116.00
34	BA	1224	A	C1'-O4'-C4'	5.33	114.17	109.90
35	BB	430	A	C1'-O4'-C4'	-5.33	105.63	109.90
35	BB	1014	U	N1-C2-N3	5.33	118.10	114.90
35	BB	1044	U	O3'-P-O5'	5.33	114.14	104.00
35	BB	1370	G	O4'-C1'-N9	5.33	112.47	108.20
39	BF	11	C	O3'-P-O5'	-5.33	93.86	104.00
39	BF	41	U	O5'-C5'-C4'	-5.33	101.56	111.70
40	BG	31	G	C4'-C3'-C2'	5.33	107.94	102.60
41	BH	27	A	C3'-C2'-C1'	5.33	105.77	101.50
54	BU	132	THR	C-N-CA	5.33	135.04	121.70
85	AA	119	G	C6-N1-C2	-5.33	121.90	125.10
85	AA	882	C	C6-N1-C1'	5.33	127.20	120.80
85	AA	1848	G	C4-C5-C6	-5.33	115.60	118.80
85	AA	2122	A	P-O5'-C5'	5.33	129.44	120.90
34	BA	132	U	C6-N1-C1'	5.33	128.67	121.20
34	BA	184	C	C6-N1-C1'	5.33	127.20	120.80
34	BA	258	C	C5-C6-N1	5.33	123.67	121.00
34	BA	730	C	O4'-C1'-N1	5.33	112.47	108.20
34	BA	777	C	C4'-C3'-C2'	-5.33	97.27	102.60
34	BA	889	U	C6-N1-C1'	5.33	128.67	121.20
34	BA	1619	U	C4'-C3'-C2'	5.33	107.93	102.60
35	BB	148	C	C6-N1-C2	-5.33	118.17	120.30
35	BB	678	U	C4'-C3'-C2'	5.33	107.93	102.60
35	BB	1289	G	C6-N1-C2	-5.33	121.90	125.10
35	BB	1422	G	P-O3'-C3'	-5.33	113.30	119.70
37	BD	114	U	O5'-C5'-C4'	-5.33	101.56	111.70
41	BH	56	C	P-O3'-C3'	-5.33	113.30	119.70
42	BI	44	PHE	N-CA-CB	-5.33	101.00	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	116	G	P-O3'-C3'	-5.33	113.30	119.70
85	AA	467	U	C3'-C2'-C1'	5.33	105.77	101.50
85	AA	490	A	P-O3'-C3'	-5.33	113.30	119.70
85	AA	978	U	O4'-C1'-C2'	5.33	112.40	107.60
85	AA	1063	U	P-O5'-C5'	5.33	129.44	120.90
85	AA	1251	G	C6-N1-C2	-5.33	121.90	125.10
85	AA	1490	A	C5-C6-N6	-5.33	119.43	123.70
85	AA	1509	A	C5-N7-C8	-5.33	101.23	103.90
34	BA	415	C	N3-C4-N4	-5.33	114.27	118.00
34	BA	648	C	C3'-C2'-C1'	-5.33	97.23	101.50
34	BA	900	A	C8-N9-C4	5.33	107.93	105.80
34	BA	1203	G	C3'-C2'-C1'	-5.33	97.23	101.50
34	BA	1205	A	C4-N9-C1'	-5.33	116.70	126.30
34	BA	1644	A	C1'-O4'-C4'	-5.33	105.64	109.90
35	BB	89	C	O4'-C1'-N1	5.33	112.47	108.20
35	BB	552	C	C1'-O4'-C4'	-5.33	105.64	109.90
35	BB	835	C	N3-C2-O2	-5.33	118.17	121.90
35	BB	1017	U	C2-N3-C4	-5.33	123.80	127.00
35	BB	1391	G	P-O5'-C5'	-5.33	112.37	120.90
36	BC	4	G	O4'-C1'-N9	5.33	112.47	108.20
36	BC	14	G	C1'-O4'-C4'	-5.33	105.64	109.90
36	BC	58	G	C8-N9-C4	5.33	108.53	106.40
38	BE	96	G	O4'-C1'-N9	5.33	112.47	108.20
64	Be	74	GLU	N-CA-C	5.33	125.39	111.00
65	Bf	362	THR	N-CA-C	-5.33	96.61	111.00
85	AA	46	U	O4'-C1'-N1	5.33	112.47	108.20
85	AA	52	U	C5'-C4'-O4'	5.33	115.50	109.10
85	AA	352	G	N7-C8-N9	-5.33	110.43	113.10
85	AA	935	A	C2'-C3'-O3'	5.33	122.23	113.70
85	AA	1531	G	O5'-C5'-C4'	-5.33	101.57	111.70
85	AA	1702	G	O4'-C4'-C3'	5.33	110.36	106.10
85	AA	1950	G	C4-N9-C1'	-5.33	119.57	126.50
23	AP	65	PHE	CB-CG-CD1	5.33	124.53	120.80
34	BA	136	A	N9-C1'-C2'	-5.33	106.14	112.00
34	BA	210	G	P-O5'-C5'	5.33	129.43	120.90
34	BA	595	U	C4-C5-C6	-5.33	116.50	119.70
34	BA	596	G	O3'-P-O5'	5.33	114.13	104.00
34	BA	1070	G	N1-C2-N2	-5.33	111.40	116.20
34	BA	1331	G	N1-C6-O6	5.33	123.10	119.90
34	BA	1334	G	O4'-C1'-N9	5.33	112.46	108.20
34	BA	1745	G	C6-N1-C2	-5.33	121.90	125.10
35	BB	54	U	C5'-C4'-O4'	5.33	115.50	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	617	C	C2-N1-C1'	5.33	124.66	118.80
42	BI	31	LYS	CB-CA-C	5.33	121.06	110.40
58	BY	63	HIS	N-CA-C	5.33	125.39	111.00
86	AB	18	G	C4-N9-C1'	-5.33	119.57	126.50
22	AO	157	LEU	N-CA-C	5.33	125.39	111.00
34	BA	63	A	P-O3'-C3'	-5.33	113.31	119.70
34	BA	584	A	P-O5'-C5'	5.33	129.43	120.90
34	BA	604	G	O5'-C5'-C4'	-5.33	101.57	111.70
34	BA	945	A	O4'-C1'-N9	5.33	112.46	108.20
34	BA	1119	A	P-O3'-C3'	-5.33	113.31	119.70
34	BA	1148	U	C2'-C3'-O3'	5.33	122.23	113.70
34	BA	1173	C	C3'-C2'-C1'	-5.33	97.24	101.50
34	BA	1351	G	N1-C6-O6	5.33	123.10	119.90
35	BB	61	A	O4'-C1'-N9	5.33	112.46	108.20
35	BB	427	U	C4'-C3'-C2'	-5.33	97.27	102.60
37	BD	3	G	N3-C2-N2	5.33	123.63	119.90
38	BE	128	G	O5'-C5'-C4'	-5.33	101.58	111.70
38	BE	138	U	C4'-C3'-C2'	-5.33	97.27	102.60
41	BH	110	C	C5'-C4'-C3'	-5.33	107.47	116.00
57	BX	58	HIS	CB-CA-C	-5.33	99.74	110.40
62	Bc	46	LEU	N-CA-CB	-5.33	99.75	110.40
85	AA	100	A	C5-N7-C8	-5.33	101.24	103.90
85	AA	401	U	C2-N3-C4	-5.33	123.80	127.00
85	AA	449	G	C3'-C2'-C1'	-5.33	97.24	101.50
85	AA	662	U	O4'-C1'-N1	5.33	112.46	108.20
85	AA	750	A	C8-N9-C1'	5.33	137.29	127.70
85	AA	1543	C	P-O3'-C3'	-5.33	113.31	119.70
85	AA	1579	A	C8-N9-C4	5.33	107.93	105.80
85	AA	1729	C	C5-C4-N4	-5.33	116.47	120.20
85	AA	2024	U	C2-N3-C4	-5.33	123.80	127.00
85	AA	2060	G	O5'-C5'-C4'	-5.33	101.58	111.70
85	AA	2068	A	O5'-C5'-C4'	5.33	121.82	111.70
24	AQ	24	ARG	NE-CZ-NH2	-5.33	117.64	120.30
35	BB	991	C	O4'-C4'-C3'	-5.33	98.67	104.00
35	BB	1167	C	C5'-C4'-O4'	5.33	115.49	109.10
35	BB	1464	G	O3'-P-O5'	5.33	114.12	104.00
35	BB	1540	U	C3'-C2'-C1'	-5.33	97.24	101.50
38	BE	73	A	N1-C6-N6	5.33	121.80	118.60
39	BF	67	A	C1'-O4'-C4'	-5.33	105.64	109.90
40	BG	159	A	P-O5'-C5'	-5.33	112.38	120.90
41	BH	59	G	N1-C6-O6	5.33	123.10	119.90
85	AA	385	A	O4'-C1'-N9	5.33	112.46	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2066	C	O4'-C4'-C3'	-5.33	98.67	104.00
11	AC	130	VAL	CA-CB-CG2	-5.33	102.91	110.90
34	BA	171	U	C3'-C2'-C1'	-5.33	97.24	101.50
34	BA	348	U	O4'-C1'-N1	5.33	112.46	108.20
34	BA	960	C	C6-N1-C2	-5.33	118.17	120.30
34	BA	1430	C	N3-C2-O2	-5.33	118.17	121.90
34	BA	1462	U	N3-C2-O2	-5.33	118.47	122.20
34	BA	1507	C	O4'-C1'-N1	5.33	112.46	108.20
35	BB	107	A	O3'-P-O5'	5.33	114.12	104.00
35	BB	128	C	C6-N1-C1'	-5.33	114.41	120.80
35	BB	398	A	N1-C6-N6	-5.33	115.41	118.60
35	BB	616	U	C2-N3-C4	-5.33	123.81	127.00
35	BB	1024	G	C5-C6-N1	5.33	114.16	111.50
35	BB	1108	G	C6-N1-C2	-5.33	121.91	125.10
35	BB	1221	G	N9-C1'-C2'	-5.33	106.14	112.00
35	BB	1301	U	C4'-C3'-C2'	-5.33	97.27	102.60
37	BD	33	U	C5'-C4'-O4'	5.33	115.49	109.10
40	BG	156	G	C4'-C3'-C2'	5.33	107.92	102.60
55	BV	69	ARG	NE-CZ-NH1	5.33	122.96	120.30
59	BZ	17	PHE	CB-CA-C	-5.33	99.75	110.40
75	Bp	2	PRO	N-CA-C	-5.33	98.25	112.10
85	AA	327	G	C5-C6-O6	-5.33	125.41	128.60
85	AA	469	G	P-O5'-C5'	5.33	129.42	120.90
85	AA	898	A	C1'-O4'-C4'	-5.33	105.64	109.90
85	AA	1793	A	C2-N3-C4	5.33	113.26	110.60
85	AA	1812	C	C1'-O4'-C4'	-5.33	105.64	109.90
85	AA	2086	C	C2-N1-C1'	5.33	124.66	118.80
85	AA	2251	U	C5'-C4'-C3'	5.33	124.52	116.00
1	A0	66	TYR	CA-CB-CG	-5.32	103.28	113.40
19	AK	105	GLU	CB-CA-C	-5.32	99.75	110.40
34	BA	17	A	N9-C1'-C2'	-5.32	106.14	112.00
34	BA	841	G	C8-N9-C4	-5.32	104.27	106.40
34	BA	1078	U	N3-C2-O2	5.32	125.93	122.20
34	BA	1160	U	C3'-C2'-C1'	-5.32	97.24	101.50
34	BA	1612	C	C3'-C2'-C1'	-5.32	97.24	101.50
34	BA	1831	A	C4-N9-C1'	5.32	135.88	126.30
35	BB	755	A	C5-C6-N6	5.32	127.96	123.70
35	BB	766	G	C5'-C4'-C3'	-5.32	107.48	116.00
35	BB	1047	C	O4'-C1'-N1	5.32	112.46	108.20
35	BB	1209	A	C5'-C4'-C3'	5.32	124.52	116.00
36	BC	10	C	C5'-C4'-O4'	5.32	115.49	109.10
38	BE	147	G	C8-N9-C4	5.32	108.53	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	175	G	N3-C4-C5	-5.32	125.94	128.60
41	BH	33	G	C5-N7-C8	-5.32	101.64	104.30
61	Bb	21	ARG	NE-CZ-NH1	5.32	122.96	120.30
77	Br	359	ARG	NE-CZ-NH1	5.32	122.96	120.30
85	AA	11	A	C1'-O4'-C4'	-5.32	105.64	109.90
85	AA	140	C	P-O3'-C3'	-5.32	113.31	119.70
85	AA	290	G	C1'-O4'-C4'	-5.32	105.64	109.90
85	AA	785	C	C6-N1-C2	-5.32	118.17	120.30
85	AA	830	A	C4-N9-C1'	-5.32	116.72	126.30
85	AA	1103	A	C4'-C3'-C2'	5.32	107.92	102.60
85	AA	1347	C	C4'-C3'-C2'	-5.32	97.28	102.60
85	AA	2061	C	C5'-C4'-C3'	5.32	124.52	116.00
6	A5	199	ASP	C-N-CA	-5.32	111.12	122.30
34	BA	382	G	N9-C4-C5	5.32	107.53	105.40
34	BA	912	G	C5-C6-N1	5.32	114.16	111.50
34	BA	1515	U	C4'-C3'-C2'	-5.32	97.28	102.60
35	BB	553	U	C5-C6-N1	-5.32	120.04	122.70
35	BB	626	C	C5'-C4'-O4'	5.32	115.49	109.10
38	BE	16	C	C5'-C4'-C3'	5.32	124.52	116.00
41	BH	57	A	C4-N9-C1'	-5.32	116.72	126.30
65	Bf	296	ARG	N-CA-CB	-5.32	101.02	110.60
84	By	131	TYR	CG-CD1-CE1	-5.32	117.04	121.30
85	AA	469	G	C4-C5-N7	5.32	112.93	110.80
85	AA	1116	G	P-O5'-C5'	5.32	129.41	120.90
6	A5	102	VAL	C-N-CA	5.32	135.00	121.70
23	AP	238	ASP	N-CA-CB	-5.32	101.02	110.60
34	BA	52	G	C5-C6-O6	-5.32	125.41	128.60
34	BA	422	C	N3-C2-O2	-5.32	118.18	121.90
34	BA	1029	C	C5'-C4'-C3'	-5.32	107.49	116.00
34	BA	1249	G	N3-C4-N9	5.32	129.19	126.00
35	BB	481	A	C8-N9-C4	-5.32	103.67	105.80
35	BB	831	C	P-O5'-C5'	-5.32	112.39	120.90
35	BB	1092	G	O4'-C4'-C3'	-5.32	98.68	104.00
35	BB	1272	G	N3-C2-N2	5.32	123.62	119.90
35	BB	1372	G	N3-C2-N2	5.32	123.62	119.90
35	BB	1491	G	C5-C6-O6	-5.32	125.41	128.60
35	BB	1515	C	O5'-C5'-C4'	-5.32	101.59	111.70
42	BI	175	SER	N-CA-CB	5.32	118.48	110.50
57	BX	87	TYR	CD1-CG-CD2	-5.32	112.05	117.90
85	AA	7	G	C8-N9-C1'	5.32	133.92	127.00
85	AA	312	G	N1-C2-N2	-5.32	111.41	116.20
85	AA	358	U	C4'-C3'-C2'	5.32	107.92	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1177	G	C4-N9-C1'	5.32	133.42	126.50
85	AA	1527	G	C5-C6-O6	-5.32	125.41	128.60
85	AA	1550	C	C2-N3-C4	-5.32	117.24	119.90
85	AA	2110	U	P-O3'-C3'	5.32	126.09	119.70
15	AG	94	ARG	NE-CZ-NH2	-5.32	117.64	120.30
18	AJ	34	VAL	CB-CA-C	5.32	121.51	111.40
34	BA	303	C	N3-C2-O2	-5.32	118.18	121.90
34	BA	428	C	N3-C2-O2	-5.32	118.18	121.90
34	BA	485	C	C6-N1-C2	-5.32	118.17	120.30
34	BA	750	C	C6-N1-C2	-5.32	118.17	120.30
34	BA	1341	A	P-O3'-C3'	-5.32	113.32	119.70
34	BA	1809	G	O5'-C5'-C4'	-5.32	101.59	111.70
35	BB	417	A	C5-N7-C8	-5.32	101.24	103.90
35	BB	890	U	O4'-C1'-N1	5.32	112.45	108.20
67	Bh	6	PHE	CB-CG-CD1	5.32	124.52	120.80
85	AA	623	G	C5-C6-N1	5.32	114.16	111.50
85	AA	1394	C	O4'-C1'-N1	5.32	112.46	108.20
85	AA	1462	A	P-O5'-C5'	-5.32	112.39	120.90
85	AA	2152	C	P-O3'-C3'	-5.32	113.32	119.70
34	BA	103	G	C5-C6-N1	5.32	114.16	111.50
34	BA	552	C	O4'-C1'-N1	5.32	112.45	108.20
34	BA	668	G	P-O5'-C5'	5.32	129.41	120.90
34	BA	800	G	N9-C4-C5	5.32	107.53	105.40
34	BA	1501	U	C2-N1-C1'	-5.32	111.32	117.70
35	BB	323	C	C5'-C4'-O4'	5.32	115.48	109.10
35	BB	567	G	O4'-C1'-N9	5.32	112.45	108.20
35	BB	1018	U	C3'-C2'-C1'	-5.32	97.25	101.50
35	BB	1060	U	N1-C2-O2	5.32	126.52	122.80
35	BB	1289	G	C5'-C4'-C3'	-5.32	107.49	116.00
36	BC	88	A	N1-C6-N6	-5.32	115.41	118.60
36	BC	147	G	N3-C4-N9	5.32	129.19	126.00
40	BG	43	U	P-O3'-C3'	-5.32	113.32	119.70
41	BH	109	G	C5-C6-N1	5.32	114.16	111.50
41	BH	117	U	C2-N3-C4	-5.32	123.81	127.00
61	Bb	46	ARG	N-CA-CB	-5.32	101.03	110.60
80	Bu	79	TYR	CB-CG-CD2	-5.32	117.81	121.00
85	AA	119	G	P-O5'-C5'	-5.32	112.39	120.90
85	AA	367	A	O3'-P-O5'	5.32	114.10	104.00
85	AA	472	A	C5-C6-N6	5.32	127.95	123.70
85	AA	766	G	C1'-O4'-C4'	-5.32	105.64	109.90
85	AA	978	U	C5'-C4'-O4'	5.32	115.48	109.10
85	AA	1284	A	C5'-C4'-O4'	5.32	115.48	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1897	A	OP1-P-OP2	-5.32	111.62	119.60
85	AA	2121	G	C1'-O4'-C4'	-5.32	105.65	109.90
20	AL	37	ASP	N-CA-CB	-5.32	101.03	110.60
33	AZ	65	ALA	N-CA-C	-5.32	96.65	111.00
34	BA	66	C	O4'-C4'-C3'	-5.32	98.68	104.00
34	BA	327	G	N9-C1'-C2'	-5.32	106.15	112.00
34	BA	487	A	C4-C5-C6	-5.32	114.34	117.00
34	BA	578	C	N3-C2-O2	-5.32	118.18	121.90
34	BA	882	G	N1-C6-O6	5.32	123.09	119.90
34	BA	1399	A	P-O3'-C3'	-5.32	113.32	119.70
34	BA	1412	G	N3-C4-C5	-5.32	125.94	128.60
34	BA	1551	G	C6-N1-C2	-5.32	121.91	125.10
35	BB	322	G	C5-C6-O6	-5.32	125.41	128.60
35	BB	734	A	C8-N9-C1'	5.32	137.27	127.70
35	BB	1197	G	C3'-C2'-C1'	-5.32	97.25	101.50
35	BB	1240	A	P-O5'-C5'	-5.32	112.40	120.90
35	BB	1521	G	O4'-C1'-N9	5.32	112.45	108.20
35	BB	1526	C	O4'-C1'-N1	5.32	112.45	108.20
35	BB	1540	U	N3-C2-O2	-5.32	118.48	122.20
36	BC	131	C	C1'-O4'-C4'	-5.32	105.65	109.90
38	BE	139	U	C6-N1-C1'	5.32	128.64	121.20
40	BG	75	C	C5-C6-N1	-5.32	118.34	121.00
54	BU	130	VAL	N-CA-C	-5.32	96.65	111.00
85	AA	160	A	C3'-C2'-C1'	5.32	105.75	101.50
85	AA	217	G	C5-C6-O6	-5.32	125.41	128.60
85	AA	738	C	P-O5'-C5'	-5.32	112.39	120.90
85	AA	768	C	C4'-C3'-O3'	-5.32	98.24	109.40
85	AA	803	C	C5-C6-N1	5.32	123.66	121.00
85	AA	867	G	C5'-C4'-O4'	5.32	115.48	109.10
85	AA	902	A	C4-N9-C1'	-5.32	116.73	126.30
85	AA	1011	G	N1-C6-O6	5.32	123.09	119.90
85	AA	1108	U	C3'-C2'-C1'	5.32	105.75	101.50
85	AA	1495	G	P-O3'-C3'	-5.32	113.32	119.70
85	AA	1610	G	C8-N9-C1'	5.32	133.91	127.00
85	AA	1848	G	N9-C1'-C2'	-5.32	106.15	112.00
3	A2	42	TRP	CB-CG-CD1	5.31	133.91	127.00
26	AS	51	VAL	CB-CA-C	-5.31	101.30	111.40
34	BA	90	G	C1'-O4'-C4'	-5.31	105.65	109.90
34	BA	162	G	C4-N9-C1'	-5.31	119.59	126.50
34	BA	171	U	C1'-O4'-C4'	-5.31	105.65	109.90
34	BA	213	A	C8-N9-C4	5.31	107.93	105.80
34	BA	1180	A	C2'-C3'-O3'	5.31	122.20	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1733	G	C5-C6-O6	-5.31	125.41	128.60
35	BB	490	G	C1'-O4'-C4'	-5.31	105.65	109.90
35	BB	994	A	N1-C6-N6	-5.31	115.41	118.60
35	BB	1147	G	C8-N9-C1'	5.31	133.91	127.00
36	BC	73	U	C3'-C2'-C1'	-5.31	97.25	101.50
38	BE	75	C	C6-N1-C2	-5.31	118.17	120.30
85	AA	85	U	C5-C4-O4	-5.31	122.71	125.90
85	AA	226	C	P-O3'-C3'	5.31	126.08	119.70
85	AA	1176	C	O4'-C1'-N1	5.31	112.45	108.20
85	AA	1219	A	P-O5'-C5'	-5.31	112.40	120.90
85	AA	1494	C	C6-N1-C2	-5.31	118.17	120.30
85	AA	1955	U	C6-N1-C1'	5.31	128.64	121.20
34	BA	543	A	O5'-P-OP2	5.31	117.07	110.70
34	BA	1011	G	C1'-O4'-C4'	-5.31	105.65	109.90
34	BA	1072	U	C2-N3-C4	-5.31	123.81	127.00
34	BA	1223	C	P-O5'-C5'	-5.31	112.40	120.90
34	BA	1398	C	N3-C2-O2	-5.31	118.18	121.90
34	BA	1442	A	N3-C4-N9	5.31	131.65	127.40
34	BA	1559	C	C3'-C2'-C1'	-5.31	97.25	101.50
35	BB	51	U	N3-C4-O4	5.31	123.12	119.40
35	BB	784	C	C4'-C3'-C2'	5.31	107.91	102.60
35	BB	1507	U	N3-C2-O2	-5.31	118.48	122.20
36	BC	9	G	C5-C6-O6	-5.31	125.41	128.60
36	BC	111	C	C4-C5-C6	5.31	120.06	117.40
38	BE	30	C	N1-C2-N3	5.31	122.92	119.20
65	Bf	188	ALA	CB-CA-C	-5.31	102.13	110.10
85	AA	307	G	C3'-C2'-C1'	-5.31	97.25	101.50
85	AA	391	G	C4-N9-C1'	-5.31	119.59	126.50
85	AA	399	A	C4'-C3'-C2'	-5.31	97.29	102.60
85	AA	685	U	OP1-P-OP2	-5.31	111.63	119.60
85	AA	1243	G	N1-C6-O6	-5.31	116.71	119.90
85	AA	1484	G	P-O3'-C3'	-5.31	113.33	119.70
85	AA	1765	G	C5-C6-O6	-5.31	125.41	128.60
85	AA	2065	U	P-O3'-C3'	-5.31	113.33	119.70
34	BA	665	C	C5-C6-N1	5.31	123.66	121.00
34	BA	744	G	N1-C2-N2	-5.31	111.42	116.20
34	BA	1197	U	C4'-C3'-O3'	-5.31	98.25	109.40
35	BB	530	C	C6-N1-C2	-5.31	118.18	120.30
35	BB	1539	C	C5-C6-N1	-5.31	118.34	121.00
37	BD	84	U	C4'-C3'-C2'	-5.31	97.29	102.60
38	BE	47	U	C5'-C4'-C3'	5.31	124.50	116.00
85	AA	634	U	C1'-O4'-C4'	-5.31	105.65	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	196	A	C5-N7-C8	5.31	106.55	103.90
34	BA	444	A	C5-C6-N6	-5.31	119.45	123.70
34	BA	593	G	N3-C4-C5	-5.31	125.94	128.60
34	BA	902	C	C1'-O4'-C4'	-5.31	105.65	109.90
34	BA	972	C	N1-C1'-C2'	-5.31	106.16	112.00
34	BA	1538	G	C5-C6-N1	5.31	114.16	111.50
34	BA	1808	A	C1'-O4'-C4'	-5.31	105.65	109.90
35	BB	358	U	O3'-P-O5'	-5.31	93.91	104.00
35	BB	616	U	O3'-P-O5'	-5.31	93.91	104.00
35	BB	707	G	N9-C1'-C2'	-5.31	106.16	112.00
35	BB	1063	C	C5-C6-N1	-5.31	118.34	121.00
35	BB	1152	U	O4'-C1'-N1	5.31	112.45	108.20
35	BB	1195	A	C8-N9-C4	-5.31	103.68	105.80
35	BB	1288	G	P-O5'-C5'	-5.31	112.40	120.90
35	BB	1320	U	O5'-C5'-C4'	-5.31	101.61	111.70
35	BB	1397	G	C4-N9-C1'	-5.31	119.60	126.50
35	BB	1545	U	C3'-C2'-C1'	-5.31	97.25	101.50
37	BD	116	C	N3-C2-O2	-5.31	118.18	121.90
54	BU	32	THR	N-CA-C	5.31	125.34	111.00
71	Bl	111	ARG	NE-CZ-NH1	5.31	122.95	120.30
85	AA	453	G	C1'-O4'-C4'	-5.31	105.65	109.90
85	AA	466	A	C2-N3-C4	-5.31	107.95	110.60
85	AA	483	G	C4-N9-C1'	5.31	133.40	126.50
85	AA	1307	U	O4'-C1'-N1	5.31	112.45	108.20
85	AA	2080	U	O3'-P-O5'	-5.31	93.91	104.00
85	AA	2105	G	C8-N9-C1'	5.31	133.90	127.00
34	BA	499	C	N1-C1'-C2'	-5.31	106.16	112.00
34	BA	744	G	O3'-P-O5'	5.31	114.08	104.00
34	BA	764	G	P-O3'-C3'	-5.31	113.33	119.70
34	BA	914	G	O4'-C1'-N9	5.31	112.45	108.20
34	BA	1548	A	O3'-P-O5'	-5.31	93.91	104.00
34	BA	1579	G	C8-N9-C1'	-5.31	120.10	127.00
34	BA	1692	U	C2-N3-C4	-5.31	123.81	127.00
35	BB	575	C	P-O5'-C5'	5.31	129.39	120.90
35	BB	1381	U	C4'-C3'-C2'	-5.31	97.29	102.60
36	BC	130	U	N1-C2-O2	5.31	126.52	122.80
36	BC	138	C	C1'-O4'-C4'	-5.31	105.65	109.90
40	BG	52	A	C1'-O4'-C4'	-5.31	105.65	109.90
45	BL	144	ARG	NE-CZ-NH1	5.31	122.95	120.30
85	AA	1066	U	C6-N1-C1'	5.31	128.63	121.20
85	AA	1155	A	N9-C1'-C2'	-5.31	106.16	112.00
85	AA	1217	U	N1-C2-N3	5.31	118.08	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	803	U	C4'-C3'-O3'	-5.31	98.26	109.40
34	BA	867	C	C6-N1-C1'	-5.31	114.43	120.80
34	BA	1256	A	O4'-C1'-C2'	-5.31	100.49	105.80
34	BA	1312	A	C8-N9-C1'	5.31	137.25	127.70
35	BB	8	U	N3-C4-O4	5.31	123.11	119.40
35	BB	1313	C	N1-C2-O2	5.31	122.08	118.90
36	BC	133	C	O4'-C1'-N1	5.31	112.44	108.20
36	BC	169	G	N9-C1'-C2'	-5.31	106.16	112.00
40	BG	29	U	C4-C5-C6	-5.31	116.52	119.70
55	BV	87	LYS	C-N-CA	5.31	134.97	121.70
31	AX	50	ARG	NE-CZ-NH2	-5.30	117.65	120.30
34	BA	204	U	C2-N3-C4	-5.30	123.82	127.00
34	BA	562	C	O4'-C1'-N1	5.30	112.44	108.20
34	BA	575	U	O4'-C1'-C2'	5.30	112.38	107.60
34	BA	636	G	OP1-P-OP2	-5.30	111.64	119.60
34	BA	909	G	C5-C6-O6	-5.30	125.42	128.60
34	BA	1067	G	C8-N9-C4	5.30	108.52	106.40
34	BA	1315	C	O4'-C1'-N1	5.30	112.44	108.20
35	BB	1071	G	C6-N1-C2	-5.30	121.92	125.10
35	BB	1335	G	O4'-C1'-N9	5.30	112.44	108.20
35	BB	1468	A	C5'-C4'-C3'	-5.30	107.51	116.00
36	BC	67	U	C2-N1-C1'	-5.30	111.33	117.70
38	BE	65	U	P-O3'-C3'	-5.30	113.33	119.70
39	BF	13	U	C5-C4-O4	5.30	129.08	125.90
40	BG	4	A	C4-C5-N7	-5.30	108.05	110.70
48	BO	128	VAL	N-CA-C	5.30	125.32	111.00
54	BU	4	SER	N-CA-CB	-5.30	102.54	110.50
68	Bi	22	ARG	NE-CZ-NH2	-5.30	117.65	120.30
83	Bx	40	PHE	CB-CG-CD2	-5.30	117.09	120.80
84	By	57	ARG	N-CA-C	5.30	125.32	111.00
85	AA	374	C	C4'-C3'-C2'	-5.30	97.30	102.60
85	AA	531	G	N3-C2-N2	5.30	123.61	119.90
85	AA	602	U	N3-C2-O2	5.30	125.91	122.20
18	AJ	96	SER	N-CA-CB	5.30	118.45	110.50
34	BA	465	A	O4'-C4'-C3'	-5.30	98.70	104.00
34	BA	764	G	O5'-P-OP2	5.30	117.06	110.70
35	BB	597	C	O4'-C1'-N1	5.30	112.44	108.20
35	BB	675	U	C5-C4-O4	5.30	129.08	125.90
68	Bi	44	ARG	NE-CZ-NH2	-5.30	117.65	120.30
85	AA	745	C	C2-N3-C4	-5.30	117.25	119.90
85	AA	1428	A	O4'-C1'-N9	5.30	112.44	108.20
85	AA	1459	C	C6-N1-C1'	5.30	127.16	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1715	C	C2-N3-C4	5.30	122.55	119.90
85	AA	2049	U	O4'-C1'-N1	5.30	112.44	108.20
23	AP	240	TRP	CB-CA-C	5.30	121.00	110.40
28	AU	54	ASP	N-CA-CB	-5.30	101.06	110.60
34	BA	30	A	O4'-C1'-N9	5.30	112.44	108.20
34	BA	71	G	N9-C4-C5	5.30	107.52	105.40
34	BA	375	C	O4'-C1'-N1	5.30	112.44	108.20
34	BA	417	A	C4-N9-C1'	-5.30	116.76	126.30
34	BA	541	C	C5'-C4'-C3'	-5.30	107.52	116.00
34	BA	1307	U	N1-C2-N3	5.30	118.08	114.90
34	BA	1477	C	C5'-C4'-C3'	-5.30	107.52	116.00
34	BA	1641	G	C5-C6-N1	5.30	114.15	111.50
34	BA	1682	A	P-O3'-C3'	-5.30	113.34	119.70
35	BB	481	A	O4'-C1'-C2'	5.30	112.37	107.60
35	BB	554	C	P-O3'-C3'	-5.30	113.34	119.70
35	BB	760	C	C5-C4-N4	5.30	123.91	120.20
35	BB	837	A	C2-N3-C4	5.30	113.25	110.60
35	BB	897	C	C5'-C4'-C3'	-5.30	107.52	116.00
35	BB	1015	U	C3'-C2'-C1'	-5.30	97.26	101.50
35	BB	1229	A	C2-N3-C4	-5.30	107.95	110.60
85	AA	483	G	P-O3'-C3'	-5.30	113.34	119.70
85	AA	597	A	N1-C2-N3	-5.30	126.65	129.30
85	AA	787	U	C3'-C2'-C1'	5.30	105.74	101.50
85	AA	1278	C	C5-C4-N4	-5.30	116.49	120.20
85	AA	1786	G	C3'-C2'-C1'	-5.30	97.26	101.50
1	A0	84	ARG	C-N-CA	5.30	134.95	121.70
11	AC	129	LEU	N-CA-CB	-5.30	99.80	110.40
34	BA	130	U	N1-C2-N3	5.30	118.08	114.90
34	BA	1577	U	C4-C5-C6	-5.30	116.52	119.70
35	BB	452	A	C2'-C3'-O3'	5.30	122.18	113.70
35	BB	586	U	C5'-C4'-C3'	-5.30	107.52	116.00
35	BB	657	A	C5-C6-N1	5.30	120.35	117.70
35	BB	1239	A	P-O5'-C5'	5.30	129.38	120.90
35	BB	1541	G	C8-N9-C4	-5.30	104.28	106.40
38	BE	102	U	N1-C1'-C2'	5.30	120.89	114.00
38	BE	164	C	C5'-C4'-C3'	-5.30	107.52	116.00
38	BE	177	U	O4'-C1'-N1	5.30	112.44	108.20
40	BG	72	G	C5-C6-N1	5.30	114.15	111.50
41	BH	78	C	O4'-C1'-N1	5.30	112.44	108.20
59	BZ	86	LYS	N-CA-CB	-5.30	101.06	110.60
85	AA	277	G	N1-C6-O6	5.30	123.08	119.90
85	AA	690	G	C4-N9-C1'	-5.30	119.61	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1507	G	C1'-O4'-C4'	-5.30	105.66	109.90
85	AA	1534	A	C8-N9-C4	-5.30	103.68	105.80
86	AB	54	U	O4'-C1'-N1	5.30	112.44	108.20
34	BA	20	A	C4-C5-C6	5.30	119.65	117.00
34	BA	316	G	N9-C1'-C2'	-5.30	106.17	112.00
34	BA	860	G	C8-N9-C4	5.30	108.52	106.40
34	BA	1458	A	O4'-C4'-C3'	-5.30	98.70	104.00
34	BA	1504	A	C3'-C2'-C1'	-5.30	97.26	101.50
35	BB	1	U	C3'-C2'-C1'	5.30	105.74	101.50
35	BB	790	A	C8-N9-C4	5.30	107.92	105.80
35	BB	1509	G	C5'-C4'-O4'	5.30	115.46	109.10
85	AA	284	C	C4'-C3'-C2'	-5.30	97.30	102.60
85	AA	870	U	C2-N1-C1'	-5.30	111.34	117.70
85	AA	1686	G	C3'-C2'-C1'	-5.30	97.26	101.50
13	AE	153	TYR	CA-CB-CG	-5.30	103.33	113.40
32	AY	58	ASN	CA-CB-CG	-5.30	101.75	113.40
34	BA	386	A	C5'-C4'-C3'	5.30	124.47	116.00
34	BA	627	U	P-O3'-C3'	5.30	126.06	119.70
34	BA	726	G	N1-C6-O6	-5.30	116.72	119.90
34	BA	908	G	C8-N9-C4	5.30	108.52	106.40
34	BA	1006	G	C5'-C4'-C3'	-5.30	107.53	116.00
34	BA	1152	A	P-O5'-C5'	-5.30	112.43	120.90
34	BA	1166	A	N1-C6-N6	-5.30	115.42	118.60
34	BA	1500	G	C6-N1-C2	-5.30	121.92	125.10
34	BA	1738	G	C5'-C4'-C3'	-5.30	107.52	116.00
34	BA	1777	U	C2-N1-C1'	5.30	124.06	117.70
35	BB	62	C	C4'-C3'-O3'	5.30	123.59	113.00
35	BB	651	G	C4-N9-C1'	5.30	133.39	126.50
35	BB	861	C	P-O3'-C3'	-5.30	113.34	119.70
35	BB	1425	A	C3'-C2'-C1'	-5.30	97.26	101.50
40	BG	70	C	O5'-C5'-C4'	-5.30	101.64	111.70
61	Bb	114	HIS	CB-CA-C	-5.30	99.81	110.40
65	Bf	331	ARG	N-CA-C	-5.30	96.70	111.00
84	By	39	HIS	CA-CB-CG	5.30	122.61	113.60
85	AA	112	A	N1-C6-N6	-5.30	115.42	118.60
85	AA	152	A	C8-N9-C4	-5.30	103.68	105.80
85	AA	250	C	C2-N3-C4	-5.30	117.25	119.90
85	AA	508	C	P-O3'-C3'	-5.30	113.34	119.70
85	AA	1522	U	C5'-C4'-C3'	-5.30	107.53	116.00
85	AA	1537	A	P-O5'-C5'	5.30	129.37	120.90
85	AA	1539	A	C1'-O4'-C4'	-5.30	105.66	109.90
85	AA	1724	A	P-O5'-C5'	5.30	129.37	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1979	A	C1'-O4'-C4'	-5.30	105.66	109.90
85	AA	2115	G	N1-C6-O6	5.30	123.08	119.90
34	BA	105	U	C3'-C2'-C1'	-5.29	97.26	101.50
34	BA	1229	G	P-O3'-C3'	-5.29	113.35	119.70
34	BA	1612	C	C2-N1-C1'	-5.29	112.98	118.80
34	BA	1687	A	C5'-C4'-C3'	5.29	124.47	116.00
35	BB	281	U	O4'-C1'-N1	5.29	112.44	108.20
35	BB	708	C	C5'-C4'-O4'	5.29	115.45	109.10
36	BC	151	G	C5-C6-O6	-5.29	125.42	128.60
37	BD	107	G	O4'-C1'-N9	5.29	112.44	108.20
38	BE	116	U	C5'-C4'-O4'	5.29	115.45	109.10
38	BE	121	G	C8-N9-C4	5.29	108.52	106.40
40	BG	24	A	C5-N7-C8	-5.29	101.25	103.90
40	BG	73	U	N3-C4-O4	-5.29	115.69	119.40
40	BG	106	G	C8-N9-C4	5.29	108.52	106.40
85	AA	244	G	C8-N9-C1'	5.29	133.88	127.00
85	AA	462	A	P-O5'-C5'	-5.29	112.43	120.90
85	AA	577	U	N3-C2-O2	-5.29	118.49	122.20
85	AA	910	G	C3'-C2'-C1'	-5.29	97.26	101.50
85	AA	1927	G	OP1-P-OP2	-5.29	111.66	119.60
85	AA	2020	C	O4'-C1'-N1	5.29	112.44	108.20
10	A9	150	TYR	CB-CG-CD2	-5.29	117.82	121.00
20	AL	27	ASP	CB-CA-C	-5.29	99.81	110.40
34	BA	81	C	C6-N1-C2	5.29	122.42	120.30
34	BA	440	A	C4'-C3'-C2'	-5.29	97.31	102.60
34	BA	857	C	N3-C2-O2	-5.29	118.19	121.90
34	BA	1410	C	C3'-C2'-C1'	-5.29	97.26	101.50
34	BA	1429	A	O4'-C1'-N9	5.29	112.43	108.20
35	BB	970	C	C4'-C3'-C2'	-5.29	97.31	102.60
35	BB	1042	U	P-O5'-C5'	5.29	129.37	120.90
35	BB	1230	A	N1-C6-N6	5.29	121.78	118.60
35	BB	1547	U	P-O3'-C3'	-5.29	113.35	119.70
41	BH	93	G	P-O5'-C5'	5.29	129.37	120.90
44	BK	168	SER	CB-CA-C	-5.29	100.04	110.10
56	BW	106	ASN	CB-CA-C	5.29	120.98	110.40
61	Bb	75	LEU	N-CA-C	5.29	125.29	111.00
85	AA	203	C	P-O3'-C3'	5.29	126.05	119.70
85	AA	493	A	C5'-C4'-C3'	5.29	124.47	116.00
85	AA	504	U	N1-C2-N3	-5.29	111.72	114.90
85	AA	710	A	N1-C6-N6	5.29	121.78	118.60
85	AA	1275	A	C1'-O4'-C4'	-5.29	105.67	109.90
85	AA	1435	C	O4'-C1'-C2'	-5.29	100.51	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1931	C	P-O5'-C5'	5.29	129.37	120.90
85	AA	2203	C	O4'-C4'-C3'	-5.29	98.71	104.00
27	AT	29	VAL	N-CA-C	5.29	125.29	111.00
27	AT	128	ALA	CB-CA-C	5.29	118.04	110.10
34	BA	189	G	C1'-O4'-C4'	-5.29	105.67	109.90
34	BA	298	G	C8-N9-C1'	-5.29	120.12	127.00
34	BA	1640	G	C8-N9-C1'	5.29	133.88	127.00
34	BA	1787	U	C6-N1-C1'	5.29	128.61	121.20
35	BB	132	G	N1-C6-O6	5.29	123.08	119.90
35	BB	541	U	C5'-C4'-O4'	5.29	115.45	109.10
35	BB	836	U	C1'-O4'-C4'	-5.29	105.67	109.90
35	BB	1132	A	C2'-C3'-O3'	5.29	122.17	113.70
35	BB	1494	G	C1'-O4'-C4'	-5.29	105.67	109.90
35	BB	1512	C	C5'-C4'-C3'	-5.29	107.53	116.00
38	BE	168	C	C5'-C4'-C3'	5.29	124.47	116.00
44	BK	170	TYR	CB-CG-CD1	5.29	124.17	121.00
49	BP	20	ARG	NH1-CZ-NH2	-5.29	113.58	119.40
59	BZ	63	ARG	NE-CZ-NH1	5.29	122.94	120.30
63	Bd	23	PRO	CA-C-N	5.29	131.92	117.10
77	Br	20	THR	N-CA-C	-5.29	96.71	111.00
85	AA	474	C	C6-N1-C1'	5.29	127.15	120.80
85	AA	490	A	C2'-C3'-O3'	5.29	122.17	113.70
85	AA	593	U	P-O5'-C5'	-5.29	112.43	120.90
85	AA	1190	G	N1-C6-O6	5.29	123.07	119.90
85	AA	1519	A	P-O3'-C3'	-5.29	113.35	119.70
85	AA	1686	G	C5'-C4'-C3'	-5.29	107.53	116.00
34	BA	297	A	O5'-P-OP2	-5.29	100.94	105.70
34	BA	484	A	N9-C1'-C2'	-5.29	106.18	112.00
34	BA	684	G	C5'-C4'-O4'	5.29	115.45	109.10
34	BA	702	G	P-O5'-C5'	5.29	129.37	120.90
34	BA	1637	G	C6-N1-C2	-5.29	121.93	125.10
34	BA	1706	A	P-O5'-C5'	-5.29	112.44	120.90
34	BA	1779	U	N3-C2-O2	-5.29	118.50	122.20
35	BB	3	C	N3-C4-C5	-5.29	119.78	121.90
35	BB	51	U	C2-N3-C4	5.29	130.17	127.00
35	BB	86	A	C8-N9-C1'	-5.29	118.18	127.70
35	BB	767	A	O5'-C5'-C4'	-5.29	101.65	111.70
36	BC	140	U	N3-C4-O4	5.29	123.10	119.40
64	Be	95	PRO	C-N-CA	5.29	134.93	121.70
85	AA	748	C	N1-C1'-C2'	-5.29	106.18	112.00
85	AA	927	A	C1'-O4'-C4'	-5.29	105.67	109.90
85	AA	1787	G	C4-N9-C1'	-5.29	119.62	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2066	C	C1'-O4'-C4'	-5.29	105.67	109.90
11	AC	193	VAL	N-CA-CB	5.29	123.14	111.50
20	AL	20	TYR	CB-CA-C	5.29	120.97	110.40
34	BA	41	U	P-O3'-C3'	-5.29	113.35	119.70
34	BA	47	U	C5'-C4'-O4'	-5.29	102.75	109.10
34	BA	449	G	C1'-O4'-C4'	-5.29	105.67	109.90
34	BA	523	A	O4'-C1'-C2'	5.29	112.36	107.60
34	BA	647	U	O4'-C1'-N1	5.29	112.43	108.20
34	BA	1825	U	C2-N3-C4	-5.29	123.83	127.00
35	BB	109	U	C6-N1-C1'	5.29	128.60	121.20
35	BB	1130	U	N3-C4-C5	5.29	117.77	114.60
35	BB	1483	A	O5'-C5'-C4'	-5.29	101.65	111.70
38	BE	107	U	OP2-P-O3'	5.29	116.83	105.20
38	BE	200	A	P-O3'-C3'	-5.29	113.35	119.70
40	BG	105	A	C4-C5-C6	-5.29	114.36	117.00
55	BV	89	TYR	CB-CG-CD1	-5.29	117.83	121.00
85	AA	386	G	C5'-C4'-C3'	-5.29	107.54	116.00
85	AA	661	C	N1-C2-O2	5.29	122.07	118.90
85	AA	821	U	C6-N1-C1'	5.29	128.60	121.20
85	AA	1017	G	P-O5'-C5'	-5.29	112.44	120.90
85	AA	1515	A	C1'-O4'-C4'	-5.29	105.67	109.90
85	AA	1578	G	OP1-P-OP2	-5.29	111.67	119.60
85	AA	1846	G	N3-C4-C5	-5.29	125.96	128.60
85	AA	1897	A	C5-C6-N6	5.29	127.93	123.70
34	BA	876	C	C3'-C2'-C1'	-5.29	97.27	101.50
34	BA	1043	C	O3'-P-O5'	5.29	114.05	104.00
35	BB	6	A	C5'-C4'-O4'	5.29	115.44	109.10
35	BB	727	U	P-O5'-C5'	5.29	129.36	120.90
35	BB	1226	G	N7-C8-N9	-5.29	110.46	113.10
38	BE	110	U	O3'-P-O5'	-5.29	93.95	104.00
85	AA	120	C	N1-C2-O2	5.29	122.07	118.90
85	AA	1461	A	C4'-C3'-C2'	-5.29	97.31	102.60
4	A3	37	VAL	CA-CB-CG2	-5.29	102.97	110.90
5	A4	107	ARG	N-CA-C	5.29	125.27	111.00
31	AX	39	ARG	NE-CZ-NH1	5.29	122.94	120.30
34	BA	95	C	N3-C4-N4	-5.29	114.30	118.00
34	BA	181	G	O3'-P-O5'	5.29	114.04	104.00
34	BA	494	A	O4'-C1'-C2'	5.29	112.36	107.60
34	BA	807	U	C2-N3-C4	-5.29	123.83	127.00
34	BA	1740	U	C2-N1-C1'	-5.29	111.36	117.70
35	BB	52	G	C8-N9-C1'	5.29	133.87	127.00
35	BB	129	U	N3-C2-O2	-5.29	118.50	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1078	U	N3-C2-O2	-5.29	118.50	122.20
35	BB	1099	U	O5'-C5'-C4'	-5.29	101.66	111.70
35	BB	1221	G	OP1-P-O3'	5.29	116.83	105.20
35	BB	1256	C	O4'-C1'-N1	5.29	112.43	108.20
35	BB	1397	G	C5-C6-O6	5.29	131.77	128.60
39	BF	13	U	C2-N1-C1'	-5.29	111.36	117.70
62	Bc	113	MET	CG-SD-CE	-5.29	91.74	100.20
67	Bh	83	ASN	CA-CB-CG	-5.29	101.77	113.40
85	AA	310	U	C6-N1-C2	-5.29	117.83	121.00
85	AA	487	G	C5'-C4'-O4'	5.29	115.44	109.10
85	AA	504	U	C4'-C3'-O3'	-5.29	98.30	109.40
85	AA	678	A	N9-C1'-C2'	-5.29	106.19	112.00
85	AA	1105	G	C1'-O4'-C4'	-5.29	105.67	109.90
85	AA	1247	A	C5-C6-N1	5.29	120.34	117.70
85	AA	1407	C	C6-N1-C2	-5.29	118.19	120.30
85	AA	1542	A	C5'-C4'-C3'	-5.29	107.54	116.00
85	AA	1801	U	C5'-C4'-C3'	-5.29	107.54	116.00
86	AB	34	G	O4'-C1'-N9	5.29	112.43	108.20
34	BA	341	U	N1-C2-O2	5.28	126.50	122.80
34	BA	409	A	P-O3'-C3'	-5.28	113.36	119.70
34	BA	948	C	C2-N1-C1'	-5.28	112.99	118.80
34	BA	986	G	C8-N9-C1'	5.28	133.87	127.00
34	BA	1420	A	C5'-C4'-C3'	5.28	124.45	116.00
34	BA	1511	C	C1'-O4'-C4'	-5.28	105.67	109.90
34	BA	1555	G	C6-N1-C2	-5.28	121.93	125.10
34	BA	1599	A	O4'-C1'-N9	5.28	112.43	108.20
35	BB	353	G	C4'-C3'-C2'	-5.28	97.32	102.60
35	BB	635	A	O4'-C1'-N9	5.28	112.43	108.20
35	BB	1292	G	C5'-C4'-C3'	-5.28	107.55	116.00
35	BB	1473	U	C6-N1-C1'	-5.28	113.80	121.20
38	BE	111	C	N3-C4-N4	5.28	121.70	118.00
39	BF	34	C	N1-C2-O2	5.28	122.07	118.90
61	Bb	110	LEU	CB-CA-C	5.28	120.24	110.20
85	AA	626	G	N3-C2-N2	5.28	123.60	119.90
85	AA	643	C	C1'-O4'-C4'	-5.28	105.67	109.90
85	AA	790	A	O5'-P-OP1	5.28	117.04	110.70
85	AA	848	C	N3-C4-N4	5.28	121.70	118.00
85	AA	864	C	P-O3'-C3'	-5.28	113.36	119.70
85	AA	1663	U	C4'-C3'-C2'	-5.28	97.32	102.60
85	AA	1786	G	N7-C8-N9	5.28	115.74	113.10
34	BA	923	C	N3-C2-O2	-5.28	118.20	121.90
34	BA	1226	G	C5-C6-N1	5.28	114.14	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1308	C	O4'-C1'-N1	5.28	112.42	108.20
35	BB	1463	A	N3-C4-C5	-5.28	123.10	126.80
36	BC	42	G	N9-C1'-C2'	-5.28	106.19	112.00
36	BC	163	A	C8-N9-C1'	5.28	137.21	127.70
40	BG	3	G	N1-C6-O6	-5.28	116.73	119.90
41	BH	46	C	N3-C2-O2	-5.28	118.20	121.90
85	AA	59	C	C5-C4-N4	-5.28	116.50	120.20
85	AA	78	A	C5'-C4'-O4'	5.28	115.44	109.10
85	AA	238	C	C5'-C4'-O4'	5.28	115.44	109.10
6	A5	116	HIS	N-CA-CB	-5.28	101.09	110.60
34	BA	29	U	O4'-C1'-N1	5.28	112.42	108.20
34	BA	136	A	C5-N7-C8	5.28	106.54	103.90
34	BA	439	A	O3'-P-O5'	-5.28	93.97	104.00
34	BA	572	G	P-O5'-C5'	5.28	129.35	120.90
34	BA	731	A	C5'-C4'-C3'	5.28	124.45	116.00
34	BA	893	U	P-O3'-C3'	-5.28	113.36	119.70
34	BA	1120	U	C5'-C4'-O4'	5.28	115.44	109.10
34	BA	1221	A	C8-N9-C1'	5.28	137.21	127.70
34	BA	1612	C	O4'-C1'-N1	5.28	112.42	108.20
35	BB	486	G	C4-N9-C1'	-5.28	119.64	126.50
35	BB	532	C	P-O5'-C5'	-5.28	112.45	120.90
35	BB	967	G	C4-N9-C1'	-5.28	119.64	126.50
35	BB	1134	G	C4-C5-N7	5.28	112.91	110.80
35	BB	1170	U	C5'-C4'-C3'	-5.28	107.55	116.00
37	BD	72	U	N1-C1'-C2'	-5.28	106.19	112.00
40	BG	29	U	C5-C4-O4	-5.28	122.73	125.90
49	BP	123	TRP	CB-CG-CD2	-5.28	119.73	126.60
51	BR	155	GLU	CB-CA-C	-5.28	99.84	110.40
78	Bs	24	TYR	CB-CG-CD2	-5.28	117.83	121.00
85	AA	117	C	C6-N1-C1'	-5.28	114.46	120.80
85	AA	233	C	C3'-C2'-C1'	-5.28	97.28	101.50
85	AA	1086	U	C6-N1-C2	-5.28	117.83	121.00
85	AA	1120	G	C6-N1-C2	-5.28	121.93	125.10
85	AA	1219	A	P-O3'-C3'	-5.28	113.36	119.70
85	AA	1626	U	O4'-C1'-N1	5.28	112.42	108.20
85	AA	1901	G	C4'-C3'-C2'	-5.28	97.32	102.60
34	BA	460	G	C5'-C4'-C3'	5.28	124.45	116.00
34	BA	758	G	C4-C5-C6	-5.28	115.63	118.80
34	BA	1417	C	C1'-O4'-C4'	-5.28	105.68	109.90
35	BB	962	U	C1'-O4'-C4'	-5.28	105.68	109.90
36	BC	132	U	P-O5'-C5'	5.28	129.35	120.90
40	BG	84	U	N3-C4-C5	-5.28	111.43	114.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
78	Bs	16	GLU	N-CA-C	5.28	125.25	111.00
85	AA	286	C	C5-C6-N1	5.28	123.64	121.00
85	AA	354	C	P-O3'-C3'	-5.28	113.36	119.70
85	AA	1457	C	C3'-C2'-C1'	-5.28	97.28	101.50
85	AA	1638	C	N3-C2-O2	-5.28	118.20	121.90
34	BA	494	A	P-O5'-C5'	5.28	129.34	120.90
34	BA	612	U	O4'-C1'-N1	5.28	112.42	108.20
34	BA	651	U	C2-N1-C1'	-5.28	111.37	117.70
34	BA	743	A	N3-C4-C5	5.28	130.49	126.80
34	BA	1177	C	O4'-C1'-N1	5.28	112.42	108.20
34	BA	1537	G	O4'-C4'-C3'	5.28	110.32	106.10
34	BA	1697	U	O5'-P-OP1	-5.28	100.95	105.70
34	BA	1726	U	O4'-C1'-N1	5.28	112.42	108.20
35	BB	23	U	C1'-O4'-C4'	-5.28	105.68	109.90
35	BB	169	U	P-O3'-C3'	5.28	126.03	119.70
35	BB	435	A	O4'-C1'-C2'	5.28	112.35	107.60
35	BB	1339	C	O4'-C1'-N1	5.28	112.42	108.20
38	BE	162	U	O5'-C5'-C4'	-5.28	101.67	111.70
39	BF	60	C	O4'-C1'-C2'	-5.28	100.52	105.80
40	BG	38	A	C8-N9-C1'	5.28	137.20	127.70
40	BG	169	A	O5'-C5'-C4'	5.28	121.73	111.70
42	BI	92	ASP	N-CA-CB	5.28	120.10	110.60
48	BO	187	HIS	N-CA-CB	-5.28	101.10	110.60
54	BU	148	ARG	C-N-CA	5.28	134.89	121.70
80	Bu	249	TYR	CB-CG-CD2	-5.28	117.83	121.00
85	AA	307	G	N9-C1'-C2'	-5.28	106.19	112.00
85	AA	943	U	N3-C4-C5	-5.28	111.43	114.60
85	AA	1470	A	N7-C8-N9	-5.28	111.16	113.80
85	AA	1854	U	C6-N1-C2	-5.28	117.83	121.00
86	AB	39	U	C5'-C4'-C3'	5.28	124.44	116.00
21	AM	27	VAL	N-CA-CB	5.28	123.11	111.50
34	BA	230	A	C4'-C3'-C2'	5.28	107.88	102.60
34	BA	238	C	C5'-C4'-C3'	-5.28	107.56	116.00
34	BA	507	U	P-O5'-C5'	5.28	129.34	120.90
34	BA	576	C	OP1-P-OP2	-5.28	111.69	119.60
34	BA	633	G	OP2-P-O3'	5.28	116.81	105.20
34	BA	769	U	C4-C5-C6	5.28	122.86	119.70
34	BA	785	G	N9-C1'-C2'	-5.28	106.20	112.00
34	BA	1116	G	O3'-P-O5'	5.28	114.02	104.00
34	BA	1470	G	C5-C6-N1	5.28	114.14	111.50
34	BA	1612	C	N3-C2-O2	-5.28	118.21	121.90
34	BA	1663	U	C6-N1-C2	-5.28	117.83	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1752	A	C3'-C2'-C1'	-5.28	97.28	101.50
35	BB	391	G	C5-C6-N1	5.28	114.14	111.50
35	BB	1113	C	O3'-P-O5'	-5.28	93.98	104.00
35	BB	1520	C	C2-N3-C4	-5.28	117.26	119.90
39	BF	69	A	N1-C6-N6	5.28	121.77	118.60
40	BG	171	A	C8-N9-C1'	5.28	137.20	127.70
47	BN	15	LYS	C-N-CA	5.28	134.89	121.70
65	Bf	372	TYR	CB-CG-CD2	-5.28	117.83	121.00
79	Bt	81	ILE	CB-CA-C	-5.28	101.05	111.60
85	AA	436	G	C6-N1-C2	-5.28	121.94	125.10
85	AA	719	C	C1'-O4'-C4'	-5.28	105.68	109.90
85	AA	892	C	C5'-C4'-O4'	5.28	115.43	109.10
85	AA	1036	A	P-O5'-C5'	-5.28	112.46	120.90
85	AA	1463	A	C5-C6-N6	-5.28	119.48	123.70
85	AA	1553	G	P-O3'-C3'	-5.28	113.37	119.70
85	AA	1976	G	C3'-C2'-C1'	-5.28	97.28	101.50
28	AU	108	THR	N-CA-C	-5.27	96.76	111.00
34	BA	236	A	O4'-C1'-N9	-5.27	103.98	108.20
34	BA	619	U	N3-C2-O2	-5.27	118.51	122.20
34	BA	768	G	O4'-C1'-N9	5.27	112.42	108.20
34	BA	1546	C	N1-C2-O2	5.27	122.06	118.90
35	BB	1331	U	C3'-C2'-C1'	-5.27	97.28	101.50
35	BB	1373	U	O3'-P-O5'	5.27	114.02	104.00
36	BC	144	C	C5-C6-N1	-5.27	118.36	121.00
40	BG	63	U	C5'-C4'-C3'	-5.27	107.56	116.00
85	AA	796	U	P-O5'-C5'	-5.27	112.46	120.90
85	AA	862	U	N1-C2-O2	5.27	126.49	122.80
85	AA	1841	G	C5'-C4'-C3'	-5.27	107.56	116.00
86	AB	71	G	N1-C6-O6	5.27	123.06	119.90
3	A2	138	TYR	CB-CG-CD2	-5.27	117.84	121.00
16	AH	121	ARG	NE-CZ-NH1	5.27	122.94	120.30
22	AO	45	MET	CG-SD-CE	-5.27	91.77	100.20
22	AO	149	ASP	CA-CB-CG	5.27	125.00	113.40
34	BA	17	A	O4'-C1'-N9	5.27	112.42	108.20
34	BA	462	C	C2-N1-C1'	5.27	124.60	118.80
34	BA	574	U	C2-N1-C1'	5.27	124.03	117.70
34	BA	719	G	C6-N1-C2	-5.27	121.94	125.10
34	BA	947	A	C1'-O4'-C4'	-5.27	105.68	109.90
34	BA	1314	A	OP2-P-O3'	5.27	116.80	105.20
34	BA	1390	C	C5'-C4'-O4'	5.27	115.43	109.10
35	BB	434	A	C1'-O4'-C4'	-5.27	105.68	109.90
35	BB	852	G	N3-C4-C5	-5.27	125.96	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1027	U	O5'-C5'-C4'	-5.27	101.68	111.70
35	BB	1096	G	C4-N9-C1'	-5.27	119.65	126.50
35	BB	1373	U	N3-C2-O2	-5.27	118.51	122.20
35	BB	1533	U	C2-N3-C4	-5.27	123.84	127.00
36	BC	102	G	C5'-C4'-C3'	-5.27	107.56	116.00
40	BG	177	U	N1-C1'-C2'	-5.27	106.20	112.00
41	BH	18	C	C4'-C3'-C2'	-5.27	97.33	102.60
41	BH	29	G	C5-C6-O6	5.27	131.76	128.60
69	Bj	103	ILE	CB-CA-C	5.27	122.14	111.60
85	AA	85	U	P-O3'-C3'	5.27	126.03	119.70
85	AA	86	G	C8-N9-C1'	5.27	133.85	127.00
85	AA	424	A	C6-N1-C2	-5.27	115.44	118.60
85	AA	479	C	C3'-C2'-C1'	-5.27	97.28	101.50
85	AA	559	G	O4'-C1'-N9	5.27	112.42	108.20
85	AA	620	U	C5'-C4'-C3'	-5.27	107.56	116.00
85	AA	873	U	C3'-C2'-C1'	5.27	105.72	101.50
85	AA	882	C	O4'-C1'-N1	5.27	112.42	108.20
85	AA	1189	A	C5-C6-N6	-5.27	119.48	123.70
85	AA	1670	U	C1'-O4'-C4'	-5.27	105.68	109.90
85	AA	1922	A	OP1-P-O3'	5.27	116.80	105.20
85	AA	2140	U	N3-C2-O2	-5.27	118.51	122.20
86	AB	70	G	C2'-C3'-O3'	5.27	122.14	113.70
30	AW	18	MET	CG-SD-CE	-5.27	91.77	100.20
34	BA	236	A	C5-C6-N1	5.27	120.33	117.70
34	BA	1314	A	C8-N9-C1'	-5.27	118.21	127.70
35	BB	74	U	N1-C2-N3	5.27	118.06	114.90
35	BB	431	U	O5'-C5'-C4'	-5.27	101.69	111.70
35	BB	1359	G	C4-N9-C1'	-5.27	119.65	126.50
36	BC	9	G	C6-N1-C2	-5.27	121.94	125.10
50	BQ	70	TYR	CA-CB-CG	-5.27	103.39	113.40
85	AA	1574	C	C6-N1-C2	-5.27	118.19	120.30
34	BA	10	G	C4'-C3'-C2'	5.27	107.87	102.60
34	BA	745	A	O4'-C1'-N9	5.27	112.42	108.20
34	BA	1267	A	C8-N9-C1'	-5.27	118.21	127.70
34	BA	1619	U	C6-N1-C1'	5.27	128.58	121.20
34	BA	1780	U	C5'-C4'-C3'	5.27	124.43	116.00
34	BA	1797	A	OP1-P-O3'	5.27	116.79	105.20
35	BB	317	C	P-O5'-C5'	5.27	129.33	120.90
35	BB	500	C	P-O3'-C3'	-5.27	113.38	119.70
35	BB	790	A	P-O5'-C5'	-5.27	112.47	120.90
35	BB	801	G	C8-N9-C4	-5.27	104.29	106.40
35	BB	803	U	OP1-P-OP2	-5.27	111.69	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1311	G	C1'-O4'-C4'	-5.27	105.68	109.90
35	BB	1416	A	C5'-C4'-O4'	5.27	115.42	109.10
35	BB	1422	G	C4-N9-C1'	-5.27	119.65	126.50
35	BB	1468	A	C2'-C3'-O3'	5.27	122.13	113.70
36	BC	72	A	C4'-C3'-C2'	-5.27	97.33	102.60
37	BD	27	A	C1'-O4'-C4'	-5.27	105.68	109.90
41	BH	20	A	C3'-C2'-C1'	-5.27	97.28	101.50
41	BH	133	U	C6-N1-C1'	5.27	128.58	121.20
47	BN	59	LEU	CB-CG-CD1	5.27	119.96	111.00
53	BT	78	THR	CA-CB-CG2	-5.27	105.02	112.40
83	Bx	213	PHE	CA-CB-CG	-5.27	101.25	113.90
84	By	35	LYS	C-N-CA	5.27	134.88	121.70
85	AA	109	G	C8-N9-C1'	5.27	133.85	127.00
85	AA	540	A	C4-C5-C6	-5.27	114.36	117.00
85	AA	1474	U	C1'-O4'-C4'	-5.27	105.68	109.90
85	AA	1511	C	O5'-C5'-C4'	-5.27	101.69	111.70
85	AA	1833	C	C4-C5-C6	-5.27	114.77	117.40
2	A1	61	ILE	CB-CA-C	-5.27	101.07	111.60
8	A7	238	ALA	N-CA-CB	-5.27	102.73	110.10
34	BA	567	U	O4'-C4'-C3'	-5.27	98.73	104.00
34	BA	800	G	N7-C8-N9	5.27	115.73	113.10
34	BA	849	G	C4-C5-C6	-5.27	115.64	118.80
34	BA	1022	C	C5'-C4'-C3'	5.27	124.43	116.00
34	BA	1703	A	O4'-C1'-N9	5.27	112.41	108.20
35	BB	665	A	C5-C6-N6	-5.27	119.48	123.70
35	BB	1198	C	C3'-C2'-C1'	-5.27	97.29	101.50
35	BB	1461	C	C4'-C3'-C2'	5.27	107.87	102.60
37	BD	32	A	C1'-O4'-C4'	-5.27	105.69	109.90
38	BE	14	C	OP1-P-O3'	5.27	116.79	105.20
38	BE	91	G	C1'-O4'-C4'	-5.27	105.69	109.90
40	BG	19	C	P-O3'-C3'	-5.27	113.38	119.70
40	BG	76	C	P-O3'-C3'	-5.27	113.38	119.70
41	BH	76	G	N3-C4-C5	-5.27	125.97	128.60
52	BS	118	ALA	N-CA-CB	-5.27	102.73	110.10
65	Bf	451	ARG	NE-CZ-NH1	5.27	122.93	120.30
85	AA	86	G	C8-N9-C4	5.27	108.51	106.40
85	AA	99	U	C2-N1-C1'	-5.27	111.38	117.70
85	AA	156	G	P-O5'-C5'	5.27	129.33	120.90
85	AA	943	U	O5'-P-OP2	-5.27	100.96	105.70
85	AA	1056	C	C6-N1-C2	-5.27	118.19	120.30
85	AA	1292	A	C5'-C4'-O4'	5.27	115.42	109.10
85	AA	2235	C	C2-N3-C4	-5.27	117.27	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	841	G	C4-N9-C1'	-5.27	119.66	126.50
34	BA	1542	A	O5'-P-OP2	5.27	117.02	110.70
34	BA	1603	A	C6-N1-C2	-5.27	115.44	118.60
35	BB	19	C	N1-C1'-C2'	-5.27	106.21	112.00
35	BB	506	G	N1-C2-N2	-5.27	111.46	116.20
35	BB	671	A	P-O5'-C5'	5.27	129.33	120.90
35	BB	1485	G	N3-C4-C5	-5.27	125.97	128.60
40	BG	53	C	N3-C4-C5	-5.27	119.79	121.90
40	BG	54	G	C8-N9-C1'	5.27	133.85	127.00
85	AA	59	C	O4'-C1'-N1	5.27	112.41	108.20
85	AA	60	U	C2-N3-C4	-5.27	123.84	127.00
85	AA	932	A	C5'-C4'-C3'	5.27	124.43	116.00
85	AA	1461	A	OP2-P-O3'	5.27	116.78	105.20
85	AA	1856	G	O5'-C5'-C4'	-5.27	101.69	111.70
22	AO	153	PHE	CB-CG-CD1	5.26	124.48	120.80
34	BA	465	A	N1-C6-N6	-5.26	115.44	118.60
34	BA	472	G	N7-C8-N9	5.26	115.73	113.10
34	BA	803	U	O4'-C1'-N1	5.26	112.41	108.20
34	BA	881	C	C2-N3-C4	-5.26	117.27	119.90
34	BA	1313	U	P-O5'-C5'	-5.26	112.48	120.90
34	BA	1709	A	C4'-C3'-C2'	5.26	107.86	102.60
35	BB	962	U	C3'-C2'-C1'	-5.26	97.29	101.50
35	BB	967	G	C8-N9-C4	-5.26	104.30	106.40
35	BB	1142	C	C1'-O4'-C4'	-5.26	105.69	109.90
35	BB	1386	C	N3-C2-O2	-5.26	118.22	121.90
35	BB	1436	U	C3'-C2'-C1'	5.26	105.71	101.50
35	BB	1481	C	C6-N1-C2	-5.26	118.19	120.30
37	BD	69	U	C6-N1-C1'	5.26	128.57	121.20
38	BE	6	A	P-O5'-C5'	-5.26	112.48	120.90
38	BE	57	U	O4'-C1'-N1	5.26	112.41	108.20
40	BG	86	U	C6-N1-C1'	-5.26	113.83	121.20
56	BW	49	ASN	CA-CB-CG	-5.26	101.82	113.40
60	Ba	116	GLN	CA-CB-CG	-5.26	101.82	113.40
69	Bj	77	ARG	N-CA-C	-5.26	96.79	111.00
85	AA	4	C	P-O3'-C3'	5.26	126.02	119.70
85	AA	415	G	O4'-C1'-N9	5.26	112.41	108.20
85	AA	800	A	C4-C5-C6	-5.26	114.37	117.00
85	AA	1644	G	N3-C4-N9	5.26	129.16	126.00
85	AA	2133	A	P-O3'-C3'	-5.26	113.38	119.70
34	BA	398	G	C5-N7-C8	-5.26	101.67	104.30
34	BA	1110	A	N7-C8-N9	-5.26	111.17	113.80
34	BA	1201	G	O5'-C5'-C4'	5.26	121.70	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1016	C	O4'-C4'-C3'	-5.26	98.74	104.00
35	BB	1093	C	C5'-C4'-C3'	-5.26	107.58	116.00
35	BB	1338	U	C1'-O4'-C4'	-5.26	105.69	109.90
77	Br	217	ARG	O-C-N	-5.26	114.28	122.70
85	AA	1098	C	C3'-C2'-C1'	-5.26	97.29	101.50
85	AA	1256	C	O5'-C5'-C4'	-5.26	101.70	111.70
85	AA	2080	U	P-O3'-C3'	5.26	126.02	119.70
85	AA	2200	A	C1'-O4'-C4'	-5.26	105.69	109.90
8	A7	81	PHE	CB-CG-CD1	5.26	124.48	120.80
34	BA	79	C	N1-C2-O2	5.26	122.06	118.90
34	BA	96	G	C5-N7-C8	-5.26	101.67	104.30
34	BA	117	C	O5'-P-OP1	5.26	117.01	110.70
34	BA	400	A	C1'-O4'-C4'	-5.26	105.69	109.90
34	BA	498	A	C5'-C4'-C3'	5.26	124.42	116.00
34	BA	569	C	O4'-C1'-N1	5.26	112.41	108.20
34	BA	758	G	C6-N1-C2	-5.26	121.94	125.10
34	BA	1030	C	P-O5'-C5'	-5.26	112.48	120.90
34	BA	1088	G	C5-C6-N1	5.26	114.13	111.50
34	BA	1321	A	O4'-C1'-N9	5.26	112.41	108.20
34	BA	1363	A	O5'-C5'-C4'	-5.26	101.70	111.70
34	BA	1372	C	C2'-C3'-O3'	5.26	122.12	113.70
35	BB	130	G	C1'-O4'-C4'	-5.26	105.69	109.90
35	BB	514	G	C3'-C2'-C1'	-5.26	97.29	101.50
35	BB	817	C	C3'-C2'-C1'	-5.26	97.29	101.50
35	BB	1114	A	C1'-O4'-C4'	-5.26	105.69	109.90
35	BB	1200	A	O4'-C1'-N9	5.26	112.41	108.20
36	BC	158	U	C3'-C2'-C1'	-5.26	97.29	101.50
38	BE	114	G	C5-C6-N1	5.26	114.13	111.50
38	BE	180	G	N3-C4-N9	-5.26	122.84	126.00
39	BF	24	G	O3'-P-O5'	5.26	114.00	104.00
40	BG	9	G	N3-C4-C5	-5.26	125.97	128.60
40	BG	147	U	C2-N3-C4	-5.26	123.84	127.00
41	BH	40	C	O3'-P-O5'	5.26	114.00	104.00
41	BH	127	A	O3'-P-O5'	-5.26	94.00	104.00
59	BZ	78	HIS	CA-CB-CG	-5.26	104.66	113.60
85	AA	130	G	N3-C2-N2	5.26	123.58	119.90
85	AA	856	G	O5'-C5'-C4'	5.26	121.70	111.70
13	AE	83	ARG	CD-NE-CZ	-5.26	116.24	123.60
34	BA	102	G	C5'-C4'-C3'	-5.26	107.58	116.00
34	BA	196	A	N7-C8-N9	-5.26	111.17	113.80
34	BA	458	G	N3-C2-N2	5.26	123.58	119.90
34	BA	535	G	C5-N7-C8	5.26	106.93	104.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	558	C	C5'-C4'-O4'	5.26	115.41	109.10
34	BA	670	U	N1-C1'-C2'	-5.26	106.22	112.00
34	BA	768	G	C6-N1-C2	5.26	128.25	125.10
35	BB	804	U	O5'-P-OP1	5.26	117.01	110.70
38	BE	64	A	C3'-C2'-C1'	-5.26	97.29	101.50
39	BF	18	U	N1-C1'-C2'	-5.26	106.22	112.00
40	BG	150	A	C4'-C3'-C2'	5.26	107.86	102.60
40	BG	171	A	P-O3'-C3'	-5.26	113.39	119.70
47	BN	104	ARG	NE-CZ-NH2	5.26	122.93	120.30
71	Bl	101	ARG	NE-CZ-NH2	-5.26	117.67	120.30
85	AA	29	U	C2-N1-C1'	-5.26	111.39	117.70
85	AA	202	U	N3-C4-O4	5.26	123.08	119.40
85	AA	279	C	O4'-C1'-N1	5.26	112.41	108.20
85	AA	579	U	C3'-C2'-C1'	-5.26	97.29	101.50
85	AA	849	A	C4'-C3'-C2'	-5.26	97.34	102.60
85	AA	851	G	N1-C6-O6	5.26	123.06	119.90
85	AA	982	G	O4'-C1'-N9	5.26	112.41	108.20
85	AA	1441	G	O4'-C1'-N9	5.26	112.41	108.20
85	AA	2208	G	C2'-C3'-O3'	5.26	122.11	113.70
85	AA	2227	A	O4'-C1'-N9	5.26	112.41	108.20
27	AT	35	PRO	O-C-N	-5.26	114.26	123.20
34	BA	139	U	C5'-C4'-O4'	5.26	115.41	109.10
34	BA	1334	G	N1-C6-O6	5.26	123.06	119.90
34	BA	1591	G	O4'-C1'-N9	5.26	112.41	108.20
34	BA	1630	A	C6-N1-C2	-5.26	115.44	118.60
34	BA	1832	A	C4'-C3'-O3'	-5.26	98.36	109.40
71	Bl	118	THR	CA-CB-CG2	5.26	119.76	112.40
83	Bx	97	ARG	NE-CZ-NH1	5.26	122.93	120.30
34	BA	121	A	C6-N1-C2	5.26	121.75	118.60
34	BA	333	A	C5'-C4'-C3'	-5.26	107.59	116.00
34	BA	436	U	P-O3'-C3'	-5.26	113.39	119.70
34	BA	614	A	C4'-C3'-C2'	-5.26	97.34	102.60
34	BA	687	G	C4-N9-C1'	-5.26	119.67	126.50
34	BA	816	G	C6-C5-N7	-5.26	127.25	130.40
34	BA	1092	U	O4'-C1'-N1	5.26	112.41	108.20
34	BA	1566	G	C5'-C4'-C3'	-5.26	107.59	116.00
34	BA	1716	A	C5-C6-N6	5.26	127.91	123.70
35	BB	54	U	O5'-P-OP2	5.26	117.01	110.70
35	BB	813	C	C1'-O4'-C4'	-5.26	105.69	109.90
35	BB	999	G	C8-N9-C1'	5.26	133.83	127.00
35	BB	1119	G	C1'-O4'-C4'	-5.26	105.69	109.90
35	BB	1379	U	P-O5'-C5'	-5.26	112.49	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	63	C	O5'-C5'-C4'	-5.26	101.71	111.70
65	Bf	298	THR	CA-CB-CG2	-5.26	105.04	112.40
85	AA	4	C	N3-C4-N4	5.26	121.68	118.00
85	AA	47	A	N3-C4-N9	5.26	131.61	127.40
85	AA	102	A	C8-N9-C4	5.26	107.90	105.80
85	AA	387	U	N3-C2-O2	-5.26	118.52	122.20
85	AA	483	G	C2-N3-C4	5.26	114.53	111.90
85	AA	606	A	C1'-O4'-C4'	-5.26	105.69	109.90
85	AA	751	C	O4'-C1'-C2'	-5.26	100.54	105.80
85	AA	1103	A	P-O3'-C3'	5.26	126.01	119.70
85	AA	1299	A	C8-N9-C1'	5.26	137.16	127.70
85	AA	1647	G	C5-C6-O6	5.26	131.75	128.60
25	AR	7	TYR	CA-CB-CG	-5.25	103.42	113.40
34	BA	495	A	N9-C1'-C2'	-5.25	106.22	112.00
34	BA	1706	A	O4'-C4'-C3'	-5.25	98.75	104.00
35	BB	524	C	P-O3'-C3'	-5.25	113.39	119.70
40	BG	44	G	N3-C2-N2	5.25	123.58	119.90
67	Bh	177	SER	N-CA-CB	5.25	118.38	110.50
79	Bt	38	ARG	NE-CZ-NH1	5.25	122.93	120.30
85	AA	118	C	P-O3'-C3'	-5.25	113.39	119.70
85	AA	303	A	N1-C6-N6	-5.25	115.45	118.60
85	AA	1463	A	O4'-C1'-N9	5.25	112.40	108.20
85	AA	1735	U	O4'-C1'-N1	5.25	112.40	108.20
85	AA	2109	G	C3'-C2'-C1'	-5.25	97.30	101.50
34	BA	68	A	N3-C4-N9	-5.25	123.20	127.40
34	BA	467	A	P-O5'-C5'	5.25	129.31	120.90
34	BA	540	G	O5'-C5'-C4'	-5.25	101.72	111.70
34	BA	1014	A	C2'-C3'-O3'	5.25	122.11	113.70
34	BA	1239	G	C5'-C4'-C3'	-5.25	107.59	116.00
34	BA	1346	U	C5'-C4'-C3'	-5.25	107.59	116.00
34	BA	1409	A	C5'-C4'-C3'	-5.25	107.59	116.00
34	BA	1721	U	N1-C2-N3	-5.25	111.75	114.90
34	BA	1834	A	P-O3'-C3'	-5.25	113.39	119.70
35	BB	367	C	C4'-C3'-C2'	-5.25	97.35	102.60
35	BB	650	A	C4-C5-C6	-5.25	114.37	117.00
35	BB	1046	C	P-O3'-C3'	-5.25	113.39	119.70
35	BB	1229	A	O4'-C1'-N9	5.25	112.40	108.20
36	BC	82	C	C4'-C3'-C2'	-5.25	97.35	102.60
37	BD	96	C	N3-C2-O2	-5.25	118.22	121.90
41	BH	113	G	O4'-C1'-C2'	5.25	112.33	107.60
53	BT	81	ARG	NE-CZ-NH2	5.25	122.93	120.30
85	AA	637	U	C6-N1-C1'	5.25	128.56	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1184	A	O4'-C1'-N9	5.25	112.40	108.20
34	BA	676	G	C1'-O4'-C4'	-5.25	105.70	109.90
34	BA	760	G	C8-N9-C1'	5.25	133.83	127.00
34	BA	1069	U	N1-C2-O2	5.25	126.48	122.80
34	BA	1083	A	C8-N9-C4	5.25	107.90	105.80
34	BA	1622	U	C4'-C3'-C2'	-5.25	97.35	102.60
35	BB	970	C	C6-N1-C2	-5.25	118.20	120.30
37	BD	77	A	C5'-C4'-O4'	5.25	115.40	109.10
49	BP	144	LEU	N-CA-CB	-5.25	99.90	110.40
65	Bf	322	ARG	N-CA-CB	5.25	120.05	110.60
84	By	59	PHE	CB-CG-CD2	-5.25	117.12	120.80
85	AA	109	G	C4'-C3'-C2'	5.25	107.85	102.60
85	AA	114	C	N1-C1'-C2'	-5.25	106.22	112.00
85	AA	196	U	C1'-O4'-C4'	-5.25	105.70	109.90
85	AA	1021	G	O4'-C1'-N9	5.25	112.40	108.20
85	AA	1292	A	C4-C5-C6	-5.25	114.37	117.00
85	AA	1518	A	C4-N9-C1'	-5.25	116.85	126.30
28	AU	100	CYS	CA-CB-SG	-5.25	104.55	114.00
34	BA	842	U	C5-C6-N1	-5.25	120.08	122.70
34	BA	985	C	N3-C2-O2	-5.25	118.22	121.90
34	BA	1305	A	N1-C2-N3	5.25	131.93	129.30
34	BA	1575	U	N3-C2-O2	-5.25	118.53	122.20
35	BB	399	A	C1'-O4'-C4'	-5.25	105.70	109.90
35	BB	1312	U	O5'-C5'-C4'	-5.25	101.72	111.70
85	AA	565	G	C1'-O4'-C4'	-5.25	105.70	109.90
85	AA	741	G	N1-C6-O6	-5.25	116.75	119.90
85	AA	966	G	O4'-C1'-N9	5.25	112.40	108.20
85	AA	1441	G	O5'-P-OP2	-5.25	100.97	105.70
85	AA	1996	A	N3-C4-N9	-5.25	123.20	127.40
86	AB	34	G	N1-C6-O6	5.25	123.05	119.90
3	A2	145	ARG	CD-NE-CZ	-5.25	116.25	123.60
4	A3	145	ARG	NE-CZ-NH1	5.25	122.92	120.30
34	BA	102	G	C8-N9-C4	-5.25	104.30	106.40
34	BA	357	A	N9-C1'-C2'	-5.25	106.23	112.00
34	BA	398	G	C2-N3-C4	5.25	114.52	111.90
34	BA	688	G	C5'-C4'-O4'	5.25	115.40	109.10
34	BA	785	G	C4-N9-C1'	-5.25	119.68	126.50
34	BA	1259	C	N3-C2-O2	-5.25	118.23	121.90
34	BA	1585	A	C2-N3-C4	-5.25	107.98	110.60
34	BA	1621	U	P-O3'-C3'	-5.25	113.40	119.70
35	BB	109	U	O5'-C5'-C4'	-5.25	101.73	111.70
35	BB	696	G	C8-N9-C1'	5.25	133.82	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1128	U	C6-N1-C1'	5.25	128.55	121.20
35	BB	1375	G	C4-N9-C1'	-5.25	119.68	126.50
37	BD	94	C	C2-N1-C1'	5.25	124.57	118.80
38	BE	92	C	O3'-P-O5'	5.25	113.97	104.00
51	BR	4	TYR	N-CA-C	5.25	125.17	111.00
77	Br	269	SER	N-CA-CB	5.25	118.37	110.50
85	AA	418	G	C5-C6-N1	5.25	114.12	111.50
85	AA	462	A	N1-C6-N6	-5.25	115.45	118.60
85	AA	517	A	C3'-C2'-C1'	5.25	105.70	101.50
85	AA	521	A	N9-C1'-C2'	-5.25	106.23	112.00
85	AA	1517	G	C3'-C2'-C1'	-5.25	97.30	101.50
85	AA	1780	A	O4'-C1'-N9	5.25	112.40	108.20
85	AA	1905	A	P-O5'-C5'	5.25	129.30	120.90
85	AA	2083	G	N3-C4-C5	-5.25	125.98	128.60
34	BA	579	U	C5-C6-N1	-5.25	120.08	122.70
34	BA	600	G	C4'-C3'-C2'	-5.25	97.35	102.60
34	BA	689	C	C6-N1-C2	-5.25	118.20	120.30
34	BA	851	C	C2-N1-C1'	-5.25	113.03	118.80
34	BA	993	C	C5'-C4'-O4'	5.25	115.39	109.10
34	BA	1201	G	P-O3'-C3'	-5.25	113.40	119.70
34	BA	1260	G	N9-C1'-C2'	-5.25	106.23	112.00
35	BB	782	A	C8-N9-C4	-5.25	103.70	105.80
35	BB	1255	U	N1-C2-N3	5.25	118.05	114.90
35	BB	1312	U	C2-N1-C1'	-5.25	111.41	117.70
35	BB	1476	C	N1-C2-O2	5.25	122.05	118.90
37	BD	44	U	O4'-C1'-N1	5.25	112.40	108.20
38	BE	144	A	C4-N9-C1'	-5.25	116.86	126.30
39	BF	16	C	O5'-P-OP1	5.25	117.00	110.70
41	BH	128	G	N7-C8-N9	-5.25	110.48	113.10
62	Bc	84	PRO	CA-N-CD	-5.25	104.16	111.50
85	AA	527	A	C5'-C4'-C3'	-5.25	107.61	116.00
85	AA	1451	U	C6-N1-C2	-5.25	117.85	121.00
85	AA	1538	C	C2-N1-C1'	-5.25	113.03	118.80
85	AA	1542	A	C5-C6-N1	5.25	120.32	117.70
85	AA	2210	C	C5'-C4'-O4'	-5.25	102.80	109.10
34	BA	25	C	N3-C2-O2	-5.25	118.23	121.90
34	BA	182	U	C6-N1-C1'	5.25	128.54	121.20
35	BB	746	A	P-O5'-C5'	-5.25	112.51	120.90
36	BC	32	U	C5'-C4'-O4'	5.25	115.39	109.10
38	BE	170	U	C6-N1-C2	-5.25	117.85	121.00
71	Bl	52	HIS	N-CA-CB	5.25	120.04	110.60
85	AA	5	U	O5'-C5'-C4'	-5.25	101.73	111.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	246	C	C2-N1-C1'	-5.25	113.03	118.80
85	AA	451	G	C8-N9-C1'	5.25	133.82	127.00
85	AA	746	G	C1'-O4'-C4'	-5.25	105.70	109.90
85	AA	1189	A	N1-C6-N6	5.25	121.75	118.60
85	AA	2036	A	C4-C5-C6	5.25	119.62	117.00
85	AA	2155	U	C5-C6-N1	5.25	125.32	122.70
85	AA	2219	G	O4'-C1'-N9	5.25	112.40	108.20
25	AR	75	HIS	N-CA-CB	5.24	120.04	110.60
34	BA	66	C	O4'-C1'-N1	5.24	112.40	108.20
34	BA	79	C	C2-N1-C1'	-5.24	113.03	118.80
34	BA	859	G	C4'-C3'-C2'	-5.24	97.36	102.60
34	BA	1119	A	N9-C1'-C2'	-5.24	106.23	112.00
34	BA	1629	A	C8-N9-C4	-5.24	103.70	105.80
34	BA	1647	G	N7-C8-N9	-5.24	110.48	113.10
34	BA	1738	G	C1'-O4'-C4'	-5.24	105.71	109.90
35	BB	149	A	O4'-C1'-N9	5.24	112.39	108.20
35	BB	425	G	C4'-C3'-C2'	-5.24	97.36	102.60
35	BB	620	G	P-O5'-C5'	-5.24	112.51	120.90
35	BB	1157	G	C1'-O4'-C4'	-5.24	105.71	109.90
35	BB	1204	C	C5-C4-N4	-5.24	116.53	120.20
35	BB	1491	G	O3'-P-O5'	-5.24	94.04	104.00
35	BB	1519	U	N1-C1'-C2'	-5.24	106.23	112.00
36	BC	24	G	O5'-C5'-C4'	-5.24	101.74	111.70
37	BD	36	C	N3-C2-O2	-5.24	118.23	121.90
39	BF	31	U	P-O5'-C5'	-5.24	112.51	120.90
62	Bc	142	ARG	NE-CZ-NH1	5.24	122.92	120.30
85	AA	352	G	O4'-C1'-C2'	5.24	112.32	107.60
85	AA	818	C	C4'-C3'-C2'	-5.24	97.36	102.60
85	AA	1031	G	O4'-C1'-N9	5.24	112.39	108.20
85	AA	1199	C	O4'-C1'-N1	5.24	112.39	108.20
85	AA	1801	U	P-O5'-C5'	5.24	129.29	120.90
86	AB	21	A	C8-N9-C1'	5.24	137.14	127.70
7	A6	56	ARG	NH1-CZ-NH2	-5.24	113.63	119.40
23	AP	225	PHE	CB-CG-CD2	-5.24	117.13	120.80
34	BA	64	A	O3'-P-O5'	5.24	113.96	104.00
34	BA	166	G	C5-C6-O6	-5.24	125.45	128.60
35	BB	382	U	C6-N1-C1'	5.24	128.54	121.20
35	BB	817	C	C6-N1-C2	-5.24	118.20	120.30
35	BB	1399	A	N9-C4-C5	-5.24	103.70	105.80
37	BD	16	U	P-O5'-C5'	5.24	129.29	120.90
37	BD	25	G	C2'-C3'-O3'	5.24	122.09	113.70
38	BE	62	C	C1'-O4'-C4'	-5.24	105.71	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	34	C	C5'-C4'-O4'	-5.24	102.81	109.10
41	BH	25	A	C4-C5-C6	-5.24	114.38	117.00
48	BO	23	ASP	N-CA-CB	-5.24	101.17	110.60
85	AA	150	U	N3-C4-C5	-5.24	111.45	114.60
85	AA	2228	G	O5'-C5'-C4'	-5.24	101.74	111.70
1	A0	42	GLN	CB-CG-CD	5.24	125.22	111.60
33	AZ	49	ARG	NE-CZ-NH1	5.24	122.92	120.30
34	BA	61	G	C8-N9-C4	-5.24	104.30	106.40
34	BA	99	G	O4'-C1'-N9	5.24	112.39	108.20
34	BA	133	A	C5-N7-C8	-5.24	101.28	103.90
34	BA	217	C	P-O3'-C3'	-5.24	113.41	119.70
34	BA	660	C	C5'-C4'-C3'	5.24	124.38	116.00
34	BA	854	A	O4'-C4'-C3'	-5.24	98.76	104.00
34	BA	867	C	C4'-C3'-C2'	5.24	107.84	102.60
34	BA	1493	U	C6-N1-C2	-5.24	117.86	121.00
34	BA	1605	G	P-O5'-C5'	5.24	129.29	120.90
35	BB	649	A	O4'-C1'-N9	5.24	112.39	108.20
35	BB	651	G	C5-N7-C8	-5.24	101.68	104.30
38	BE	85	G	N1-C6-O6	5.24	123.04	119.90
38	BE	138	U	O3'-P-O5'	-5.24	94.04	104.00
49	BP	48	ARG	N-CA-C	-5.24	96.85	111.00
61	Bb	52	TYR	CB-CG-CD1	5.24	124.14	121.00
69	Bj	5	ARG	N-CA-C	-5.24	96.85	111.00
34	BA	228	A	C4'-C3'-O3'	-5.24	98.40	109.40
34	BA	263	G	O4'-C1'-C2'	-5.24	100.56	105.80
34	BA	749	G	N1-C6-O6	5.24	123.04	119.90
34	BA	1611	A	C6-N1-C2	-5.24	115.46	118.60
34	BA	1802	C	N3-C2-O2	-5.24	118.23	121.90
35	BB	534	C	C5-C6-N1	5.24	123.62	121.00
35	BB	835	C	C2-N1-C1'	5.24	124.56	118.80
35	BB	971	A	O3'-P-O5'	-5.24	94.05	104.00
35	BB	1504	U	O5'-C5'-C4'	5.24	121.65	111.70
36	BC	46	G	C5'-C4'-C3'	-5.24	107.62	116.00
36	BC	120	G	O4'-C1'-C2'	5.24	112.31	107.60
37	BD	76	U	N3-C4-O4	5.24	123.07	119.40
39	BF	27	G	C5-C6-O6	-5.24	125.46	128.60
40	BG	14	G	O4'-C4'-C3'	-5.24	98.76	104.00
40	BG	14	G	P-O5'-C5'	-5.24	112.52	120.90
85	AA	408	C	N3-C4-N4	-5.24	114.33	118.00
85	AA	443	A	C4'-C3'-O3'	-5.24	98.40	109.40
85	AA	537	G	N9-C4-C5	-5.24	103.31	105.40
85	AA	1680	U	O4'-C1'-N1	5.24	112.39	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2096	G	N3-C4-C5	-5.24	125.98	128.60
34	BA	281	C	N1-C2-O2	5.24	122.04	118.90
34	BA	1297	G	C5'-C4'-C3'	-5.24	107.62	116.00
34	BA	1433	U	N1-C1'-C2'	-5.24	106.24	112.00
35	BB	485	U	O4'-C1'-C2'	5.24	112.31	107.60
35	BB	541	U	N3-C4-O4	5.24	123.07	119.40
40	BG	41	U	N1-C2-O2	5.24	126.47	122.80
63	Bd	8	THR	N-CA-CB	5.24	120.25	110.30
85	AA	1903	G	N9-C1'-C2'	-5.24	106.24	112.00
2	A1	45	ILE	CA-CB-CG1	5.24	120.95	111.00
34	BA	246	G	N1-C6-O6	-5.24	116.76	119.90
34	BA	648	C	P-O3'-C3'	-5.24	113.42	119.70
34	BA	764	G	N1-C6-O6	-5.24	116.76	119.90
34	BA	875	G	C8-N9-C1'	5.24	133.81	127.00
34	BA	1015	G	O3'-P-O5'	-5.24	94.05	104.00
34	BA	1528	U	C2-N1-C1'	-5.24	111.42	117.70
34	BA	1596	C	C5-C6-N1	-5.24	118.38	121.00
34	BA	1740	U	O4'-C1'-N1	5.24	112.39	108.20
34	BA	1817	G	OP1-P-OP2	-5.24	111.75	119.60
36	BC	113	G	O3'-P-O5'	-5.24	94.05	104.00
40	BG	147	U	C5'-C4'-C3'	-5.24	107.62	116.00
41	BH	103	C	C6-N1-C2	-5.24	118.21	120.30
69	Bj	53	ARG	NE-CZ-NH1	5.24	122.92	120.30
85	AA	492	C	C3'-C2'-C1'	-5.24	97.31	101.50
85	AA	492	C	C4'-C3'-C2'	-5.24	97.36	102.60
85	AA	574	U	C5'-C4'-O4'	5.24	115.38	109.10
85	AA	583	U	OP2-P-O3'	5.24	116.72	105.20
85	AA	619	A	N9-C1'-C2'	-5.24	106.24	112.00
85	AA	640	C	C6-N1-C2	-5.24	118.21	120.30
85	AA	768	C	C2-N3-C4	-5.24	117.28	119.90
85	AA	1009	G	C5-C6-O6	-5.24	125.46	128.60
85	AA	2130	G	N9-C1'-C2'	-5.24	106.24	112.00
85	AA	2154	C	P-O3'-C3'	5.24	125.98	119.70
34	BA	551	U	N3-C4-O4	-5.23	115.74	119.40
34	BA	1065	U	N3-C2-O2	-5.23	118.54	122.20
34	BA	1150	A	C8-N9-C4	-5.23	103.71	105.80
35	BB	60	A	N9-C1'-C2'	-5.23	106.24	112.00
35	BB	472	C	N1-C2-N3	5.23	122.86	119.20
36	BC	114	C	C5'-C4'-O4'	5.23	115.38	109.10
38	BE	62	C	O5'-C5'-C4'	-5.23	101.76	111.70
38	BE	63	C	C5-C6-N1	-5.23	118.38	121.00
38	BE	170	U	N1-C2-N3	-5.23	111.76	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	111	C	C5'-C4'-C3'	5.23	124.38	116.00
40	BG	177	U	C5'-C4'-C3'	5.23	124.37	116.00
54	BU	12	ARG	NH1-CZ-NH2	-5.23	113.64	119.40
85	AA	537	G	O4'-C1'-C2'	5.23	112.31	107.60
85	AA	659	A	P-O3'-C3'	-5.23	113.42	119.70
85	AA	1150	G	O4'-C1'-C2'	5.23	112.31	107.60
85	AA	1170	C	N3-C4-N4	-5.23	114.34	118.00
85	AA	2058	C	P-O3'-C3'	-5.23	113.42	119.70
34	BA	21	C	C4'-C3'-C2'	-5.23	97.37	102.60
34	BA	117	C	C6-N1-C2	5.23	122.39	120.30
34	BA	191	G	O4'-C1'-N9	5.23	112.39	108.20
34	BA	232	U	N1-C1'-C2'	-5.23	106.24	112.00
34	BA	765	U	N1-C2-N3	5.23	118.04	114.90
34	BA	1093	G	P-O3'-C3'	-5.23	113.42	119.70
34	BA	1702	G	C5'-C4'-C3'	-5.23	107.63	116.00
35	BB	404	A	O5'-C5'-C4'	-5.23	101.76	111.70
35	BB	645	C	N3-C2-O2	-5.23	118.24	121.90
40	BG	62	C	N1-C2-N3	5.23	122.86	119.20
40	BG	152	G	N1-C6-O6	-5.23	116.76	119.90
41	BH	34	G	C6-N1-C2	-5.23	121.96	125.10
47	BN	174	PRO	N-CA-C	5.23	125.70	112.10
85	AA	657	C	C5'-C4'-C3'	-5.23	107.63	116.00
85	AA	1211	C	O4'-C1'-N1	5.23	112.39	108.20
85	AA	1543	C	N3-C2-O2	-5.23	118.24	121.90
8	A7	74	ALA	N-CA-C	-5.23	96.88	111.00
24	AQ	59	LYS	CB-CA-C	-5.23	99.94	110.40
34	BA	128	C	O4'-C1'-C2'	5.23	112.31	107.60
34	BA	501	U	C2-N3-C4	-5.23	123.86	127.00
34	BA	1103	G	N7-C8-N9	-5.23	110.48	113.10
34	BA	1248	A	N1-C6-N6	-5.23	115.46	118.60
35	BB	25	A	C6-C5-N7	5.23	135.96	132.30
35	BB	68	G	C2'-C3'-O3'	5.23	122.07	113.70
35	BB	815	G	N7-C8-N9	-5.23	110.48	113.10
35	BB	842	G	C3'-C2'-C1'	-5.23	97.31	101.50
35	BB	870	C	C6-N1-C1'	-5.23	114.52	120.80
35	BB	1098	G	C4-C5-C6	-5.23	115.66	118.80
37	BD	70	C	N3-C2-O2	-5.23	118.24	121.90
37	BD	118	C	C6-N1-C1'	5.23	127.08	120.80
38	BE	117	A	O4'-C1'-N9	5.23	112.39	108.20
38	BE	123	A	C5-N7-C8	-5.23	101.28	103.90
39	BF	11	C	OP1-P-O3'	5.23	116.71	105.20
40	BG	16	G	C4-C5-N7	-5.23	108.71	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
50	BQ	66	ARG	NE-CZ-NH2	5.23	122.92	120.30
67	Bh	121	LYS	CA-C-N	-5.23	105.69	117.20
71	Bl	53	GLY	C-N-CA	5.23	134.78	121.70
85	AA	483	G	N1-C6-O6	-5.23	116.76	119.90
85	AA	567	G	O4'-C1'-N9	5.23	112.38	108.20
85	AA	584	G	N9-C1'-C2'	-5.23	106.25	112.00
85	AA	655	U	C4-C5-C6	-5.23	116.56	119.70
85	AA	655	U	C5-C6-N1	-5.23	120.08	122.70
85	AA	674	U	O5'-C5'-C4'	-5.23	101.76	111.70
85	AA	839	C	C5'-C4'-C3'	-5.23	107.63	116.00
85	AA	942	A	C5'-C4'-O4'	5.23	115.38	109.10
85	AA	1117	G	C5-C6-N1	5.23	114.12	111.50
85	AA	1258	U	C2-N3-C4	-5.23	123.86	127.00
85	AA	2050	C	C3'-C2'-C1'	-5.23	97.32	101.50
85	AA	2101	C	P-O3'-C3'	-5.23	113.42	119.70
86	AB	61	C	C4'-C3'-C2'	-5.23	97.37	102.60
34	BA	1456	C	O4'-C1'-C2'	5.23	112.31	107.60
34	BA	1481	U	C1'-O4'-C4'	-5.23	105.72	109.90
35	BB	1175	A	N1-C6-N6	-5.23	115.46	118.60
37	BD	16	U	O4'-C1'-C2'	5.23	112.31	107.60
85	AA	1547	G	C5-C6-O6	5.23	131.74	128.60
85	AA	2215	C	N1-C2-N3	5.23	122.86	119.20
34	BA	145	U	C4-C5-C6	-5.23	116.56	119.70
34	BA	531	C	N1-C1'-C2'	-5.23	106.25	112.00
34	BA	533	U	C5-C6-N1	-5.23	120.09	122.70
34	BA	788	C	O4'-C1'-C2'	5.23	112.30	107.60
34	BA	1138	C	O4'-C1'-N1	5.23	112.38	108.20
34	BA	1219	G	P-O3'-C3'	-5.23	113.43	119.70
34	BA	1434	U	C3'-C2'-C1'	5.23	105.68	101.50
35	BB	265	C	P-O3'-C3'	-5.23	113.43	119.70
35	BB	697	G	C5-C6-O6	-5.23	125.46	128.60
35	BB	1424	G	C8-N9-C4	5.23	108.49	106.40
37	BD	47	U	O4'-C4'-C3'	-5.23	98.77	104.00
40	BG	71	C	N1-C1'-C2'	-5.23	106.25	112.00
85	AA	36	U	C4'-C3'-C2'	5.23	107.83	102.60
85	AA	165	C	N1-C2-O2	5.23	122.04	118.90
85	AA	1018	G	C5-C6-O6	-5.23	125.46	128.60
85	AA	1220	A	N1-C6-N6	-5.23	115.46	118.60
85	AA	1292	A	N9-C4-C5	-5.23	103.71	105.80
85	AA	2048	C	N1-C2-O2	5.23	122.04	118.90
11	AC	90	THR	CA-CB-CG2	-5.23	105.08	112.40
34	BA	182	U	O4'-C1'-N1	5.23	112.38	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	670	U	C5-C6-N1	-5.23	120.09	122.70
34	BA	848	U	OP1-P-O3'	5.23	116.70	105.20
34	BA	1824	U	O4'-C1'-N1	5.23	112.38	108.20
35	BB	586	U	C5'-C4'-O4'	5.23	115.37	109.10
85	AA	259	A	C8-N9-C1'	5.23	137.11	127.70
85	AA	279	C	C6-N1-C2	-5.23	118.21	120.30
85	AA	1098	C	O4'-C1'-N1	5.23	112.38	108.20
85	AA	2195	A	C4-N9-C1'	-5.23	116.89	126.30
34	BA	184	C	P-O3'-C3'	-5.22	113.43	119.70
34	BA	421	G	C4-N9-C1'	-5.22	119.71	126.50
34	BA	1610	A	C8-N9-C1'	5.22	137.10	127.70
34	BA	1819	U	O4'-C1'-N1	5.22	112.38	108.20
35	BB	443	A	P-O3'-C3'	5.22	125.97	119.70
35	BB	586	U	C6-N1-C1'	5.22	128.51	121.20
35	BB	688	U	O3'-P-O5'	5.22	113.93	104.00
35	BB	873	C	N3-C2-O2	-5.22	118.24	121.90
35	BB	1341	U	C4'-C3'-C2'	-5.22	97.38	102.60
37	BD	24	U	C2'-C3'-O3'	5.22	122.06	113.70
38	BE	119	U	O4'-C1'-N1	5.22	112.38	108.20
39	BF	31	U	C5'-C4'-O4'	5.22	115.37	109.10
40	BG	23	C	O4'-C1'-N1	5.22	112.38	108.20
50	BQ	57	ARG	NE-CZ-NH2	-5.22	117.69	120.30
55	BV	50	TYR	CB-CG-CD1	5.22	124.13	121.00
85	AA	109	G	N9-C4-C5	5.22	107.49	105.40
85	AA	806	G	P-O3'-C3'	-5.22	113.43	119.70
85	AA	964	C	O5'-C5'-C4'	5.22	121.63	111.70
85	AA	1846	G	N1-C2-N2	-5.22	111.50	116.20
85	AA	2124	G	C5-C6-N1	5.22	114.11	111.50
85	AA	2124	G	N9-C1'-C2'	-5.22	106.25	112.00
85	AA	2198	G	O4'-C1'-N9	5.22	112.38	108.20
85	AA	2237	G	P-O5'-C5'	5.22	129.26	120.90
28	AU	78	SER	N-CA-CB	5.22	118.33	110.50
34	BA	240	C	P-O3'-C3'	-5.22	113.43	119.70
34	BA	1615	A	O4'-C1'-N9	5.22	112.38	108.20
35	BB	528	G	C2-N3-C4	-5.22	109.29	111.90
35	BB	778	A	N1-C6-N6	5.22	121.73	118.60
35	BB	1293	C	O4'-C1'-N1	5.22	112.38	108.20
61	Bb	10	HIS	CA-CB-CG	-5.22	104.72	113.60
85	AA	38	C	P-O3'-C3'	-5.22	113.43	119.70
85	AA	126	U	O4'-C4'-C3'	-5.22	98.78	104.00
85	AA	370	A	N1-C6-N6	-5.22	115.47	118.60
85	AA	413	G	C4'-C3'-C2'	-5.22	97.38	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	835	C	C5-C4-N4	5.22	123.86	120.20
85	AA	878	U	N1-C1'-C2'	-5.22	106.25	112.00
85	AA	1544	G	C5'-C4'-O4'	5.22	115.37	109.10
85	AA	1617	G	N1-C6-O6	5.22	123.03	119.90
85	AA	1685	G	C5'-C4'-C3'	-5.22	107.64	116.00
85	AA	1930	U	C4'-C3'-C2'	-5.22	97.38	102.60
34	BA	607	C	C5'-C4'-O4'	5.22	115.36	109.10
34	BA	963	G	N9-C1'-C2'	-5.22	106.26	112.00
34	BA	1284	G	C4-N9-C1'	-5.22	119.71	126.50
34	BA	1729	G	C3'-C2'-C1'	5.22	105.68	101.50
35	BB	813	C	N3-C2-O2	-5.22	118.25	121.90
35	BB	1516	C	P-O5'-C5'	-5.22	112.55	120.90
38	BE	10	G	OP1-P-OP2	-5.22	111.77	119.60
38	BE	97	G	C1'-O4'-C4'	-5.22	105.72	109.90
40	BG	60	A	N1-C6-N6	5.22	121.73	118.60
40	BG	127	G	N1-C2-N2	-5.22	111.50	116.20
85	AA	436	G	N7-C8-N9	-5.22	110.49	113.10
85	AA	456	A	C8-N9-C1'	5.22	137.10	127.70
85	AA	787	U	O5'-C5'-C4'	-5.22	101.78	111.70
17	AI	47	ARG	NE-CZ-NH1	5.22	122.91	120.30
34	BA	448	U	C2-N1-C1'	-5.22	111.44	117.70
34	BA	979	G	C3'-C2'-C1'	-5.22	97.33	101.50
34	BA	1223	C	C2-N1-C1'	5.22	124.54	118.80
34	BA	1263	A	C5'-C4'-O4'	5.22	115.36	109.10
34	BA	1412	G	N3-C4-N9	5.22	129.13	126.00
35	BB	106	A	N9-C1'-C2'	-5.22	106.26	112.00
35	BB	752	A	C4-C5-C6	-5.22	114.39	117.00
35	BB	1109	A	P-O5'-C5'	5.22	129.25	120.90
35	BB	1221	G	C4-N9-C1'	-5.22	119.71	126.50
35	BB	1410	G	C3'-C2'-C1'	-5.22	97.32	101.50
37	BD	31	U	C2-N1-C1'	-5.22	111.44	117.70
37	BD	78	C	N3-C2-O2	-5.22	118.25	121.90
38	BE	1	U	C1'-O4'-C4'	-5.22	105.72	109.90
40	BG	21	C	C4-C5-C6	5.22	120.01	117.40
40	BG	123	C	C5'-C4'-C3'	-5.22	107.65	116.00
41	BH	67	G	C4'-C3'-C2'	-5.22	97.38	102.60
47	BN	12	HIS	CA-CB-CG	-5.22	104.73	113.60
56	BW	13	PHE	N-CA-C	-5.22	96.91	111.00
60	Ba	101	ALA	N-CA-C	-5.22	96.91	111.00
65	Bf	127	VAL	CB-CA-C	-5.22	101.48	111.40
66	Bg	81	TYR	CB-CG-CD1	-5.22	117.87	121.00
85	AA	93	G	P-O3'-C3'	-5.22	113.44	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	102	A	N9-C1'-C2'	-5.22	106.26	112.00
85	AA	362	G	O4'-C1'-C2'	5.22	112.30	107.60
85	AA	1978	G	C4-C5-N7	5.22	112.89	110.80
85	AA	2001	C	C2'-C3'-O3'	5.22	122.05	113.70
23	AP	69	MET	CG-SD-CE	-5.22	91.85	100.20
34	BA	12	G	C5-C6-N1	5.22	114.11	111.50
34	BA	886	G	C3'-C2'-C1'	-5.22	97.33	101.50
34	BA	1182	U	O4'-C4'-C3'	-5.22	98.78	104.00
34	BA	1542	A	O4'-C1'-C2'	5.22	112.30	107.60
34	BA	1826	C	N3-C2-O2	-5.22	118.25	121.90
35	BB	22	A	O5'-P-OP1	5.22	116.96	110.70
41	BH	51	C	C6-N1-C1'	5.22	127.06	120.80
47	BN	63	VAL	CA-CB-CG2	5.22	118.73	110.90
62	Bc	28	SER	N-CA-C	5.22	125.09	111.00
85	AA	197	C	N3-C4-N4	5.22	121.65	118.00
85	AA	2229	G	N9-C1'-C2'	5.22	120.78	114.00
15	AG	130	ARG	NE-CZ-NH1	5.22	122.91	120.30
34	BA	355	U	O4'-C1'-N1	5.22	112.37	108.20
34	BA	697	A	C8-N9-C4	-5.22	103.71	105.80
34	BA	1247	G	C5-C6-O6	-5.22	125.47	128.60
34	BA	1323	G	P-O5'-C5'	5.22	129.25	120.90
35	BB	75	A	C4-N9-C1'	-5.22	116.91	126.30
35	BB	608	A	C5-C6-N1	5.22	120.31	117.70
35	BB	685	G	P-O5'-C5'	-5.22	112.56	120.90
35	BB	692	G	C1'-O4'-C4'	-5.22	105.73	109.90
35	BB	1035	C	N3-C2-O2	-5.22	118.25	121.90
35	BB	1357	C	C2-N3-C4	5.22	122.51	119.90
35	BB	1493	A	C5'-C4'-C3'	5.22	124.34	116.00
37	BD	109	U	O5'-C5'-C4'	-5.22	101.79	111.70
39	BF	57	C	C2-N3-C4	-5.22	117.29	119.90
40	BG	109	C	C5-C6-N1	-5.22	118.39	121.00
41	BH	18	C	C5'-C4'-C3'	-5.22	107.66	116.00
49	BP	134	LYS	N-CA-CB	5.22	119.99	110.60
63	Bd	50	ASN	N-CA-C	5.22	125.08	111.00
67	Bh	9	LYS	N-CA-CB	-5.22	101.21	110.60
84	By	127	ASP	CA-C-N	-5.22	105.72	117.20
84	By	138	VAL	CB-CA-C	5.22	121.31	111.40
85	AA	244	G	P-O5'-C5'	5.22	129.25	120.90
85	AA	394	C	N3-C4-C5	5.22	123.99	121.90
85	AA	743	C	C6-N1-C1'	5.22	127.06	120.80
85	AA	812	C	P-O5'-C5'	-5.22	112.55	120.90
85	AA	1248	U	C2-N1-C1'	-5.22	111.44	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1854	U	O4'-C1'-N1	5.22	112.37	108.20
85	AA	2095	U	N1-C2-O2	5.22	126.45	122.80
34	BA	171	U	C5-C6-N1	-5.21	120.09	122.70
34	BA	695	A	C6-N1-C2	-5.21	115.47	118.60
34	BA	1027	C	C3'-C2'-C1'	-5.21	97.33	101.50
34	BA	1099	U	C5'-C4'-O4'	5.21	115.36	109.10
34	BA	1277	G	O5'-C5'-C4'	-5.21	101.79	111.70
34	BA	1386	G	C5-C6-O6	-5.21	125.47	128.60
34	BA	1469	G	N3-C2-N2	5.21	123.55	119.90
34	BA	1640	G	C5-C6-N1	5.21	114.11	111.50
34	BA	1692	U	C5'-C4'-C3'	-5.21	107.66	116.00
34	BA	1800	G	C8-N9-C1'	-5.21	120.22	127.00
34	BA	1824	U	P-O3'-C3'	-5.21	113.44	119.70
34	BA	1842	U	C5'-C4'-O4'	5.21	115.36	109.10
35	BB	599	U	C1'-O4'-C4'	-5.21	105.73	109.90
35	BB	718	G	P-O3'-C3'	-5.21	113.44	119.70
35	BB	1133	C	C2'-C3'-O3'	5.21	122.04	113.70
35	BB	1509	G	C4-N9-C1'	-5.21	119.72	126.50
36	BC	116	C	C5'-C4'-C3'	-5.21	107.66	116.00
36	BC	120	G	C5-N7-C8	-5.21	101.69	104.30
40	BG	176	G	O4'-C1'-C2'	5.21	112.29	107.60
85	AA	97	A	C5-C6-N6	-5.21	119.53	123.70
85	AA	516	G	C5-C6-N1	5.21	114.11	111.50
85	AA	730	G	C4'-C3'-C2'	-5.21	97.39	102.60
85	AA	884	A	N1-C2-N3	-5.21	126.69	129.30
85	AA	1463	A	C8-N9-C4	-5.21	103.71	105.80
34	BA	177	G	C5-C6-N1	5.21	114.11	111.50
34	BA	191	G	N1-C2-N2	-5.21	111.51	116.20
34	BA	212	A	N7-C8-N9	-5.21	111.19	113.80
34	BA	589	A	P-O5'-C5'	-5.21	112.56	120.90
35	BB	60	A	O3'-P-O5'	5.21	113.90	104.00
35	BB	285	C	O4'-C1'-N1	5.21	112.37	108.20
35	BB	301	G	O4'-C1'-N9	5.21	112.37	108.20
35	BB	1495	U	C5-C4-O4	-5.21	122.77	125.90
40	BG	104	A	C3'-C2'-C1'	-5.21	97.33	101.50
40	BG	166	C	O5'-P-OP2	-5.21	101.01	105.70
85	AA	285	C	C3'-C2'-C1'	-5.21	97.33	101.50
85	AA	682	C	C2'-C3'-O3'	5.21	122.04	113.70
85	AA	1017	G	O4'-C1'-N9	5.21	112.37	108.20
20	AL	91	VAL	N-CA-C	-5.21	96.93	111.00
34	BA	248	G	O5'-C5'-C4'	5.21	121.60	111.70
34	BA	617	G	C4-N9-C1'	-5.21	119.72	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	745	A	C6-N1-C2	-5.21	115.47	118.60
34	BA	751	A	C3'-C2'-C1'	-5.21	97.33	101.50
34	BA	887	U	C6-N1-C1'	5.21	128.50	121.20
34	BA	943	G	C3'-C2'-C1'	-5.21	97.33	101.50
34	BA	1520	A	N9-C1'-C2'	-5.21	106.27	112.00
34	BA	1540	C	N1-C2-N3	5.21	122.85	119.20
34	BA	1549	U	P-O5'-C5'	-5.21	112.56	120.90
34	BA	1706	A	O5'-C5'-C4'	5.21	121.60	111.70
34	BA	1732	A	C4-C5-N7	-5.21	108.09	110.70
35	BB	1313	C	C3'-C2'-C1'	-5.21	97.33	101.50
36	BC	114	C	O4'-C1'-N1	5.21	112.37	108.20
38	BE	205	G	O3'-P-O5'	-5.21	94.10	104.00
40	BG	138	C	N3-C2-O2	-5.21	118.25	121.90
51	BR	76	TRP	CB-CG-CD2	-5.21	119.83	126.60
84	By	78	MET	CG-SD-CE	-5.21	91.86	100.20
85	AA	253	C	P-O3'-C3'	-5.21	113.45	119.70
85	AA	496	C	C5-C4-N4	5.21	123.85	120.20
85	AA	594	C	C6-N1-C1'	5.21	127.05	120.80
85	AA	811	A	C5-C6-N1	5.21	120.31	117.70
85	AA	1221	G	C5-N7-C8	-5.21	101.69	104.30
85	AA	1523	G	C5-C6-N1	5.21	114.11	111.50
85	AA	1912	U	P-O5'-C5'	5.21	129.24	120.90
85	AA	1916	A	C5-C6-N1	-5.21	115.09	117.70
85	AA	2014	G	C5'-C4'-C3'	-5.21	107.66	116.00
11	AC	77	ARG	NE-CZ-NH2	-5.21	117.69	120.30
34	BA	1618	A	C5'-C4'-C3'	-5.21	107.66	116.00
35	BB	77	A	N7-C8-N9	-5.21	111.19	113.80
35	BB	403	U	C1'-O4'-C4'	-5.21	105.73	109.90
35	BB	1424	G	C5'-C4'-O4'	5.21	115.35	109.10
40	BG	67	A	C5-C6-N6	-5.21	119.53	123.70
40	BG	107	U	N1-C1'-C2'	-5.21	106.27	112.00
63	Bd	29	MET	CA-CB-CG	5.21	122.16	113.30
85	AA	824	C	C2-N1-C1'	-5.21	113.07	118.80
85	AA	1736	U	C3'-C2'-C1'	5.21	105.67	101.50
34	BA	273	G	C5-C6-N1	5.21	114.11	111.50
34	BA	300	C	C5-C6-N1	-5.21	118.40	121.00
34	BA	819	G	C3'-C2'-C1'	-5.21	97.33	101.50
34	BA	1045	C	C6-N1-C2	-5.21	118.22	120.30
34	BA	1650	G	C5-C6-O6	5.21	131.72	128.60
35	BB	501	G	N9-C1'-C2'	-5.21	106.27	112.00
35	BB	640	A	C5'-C4'-O4'	-5.21	102.85	109.10
35	BB	1351	G	C5'-C4'-C3'	5.21	124.33	116.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1471	A	C4-N9-C1'	5.21	135.68	126.30
83	Bx	270	ASP	C-N-CA	5.21	134.72	121.70
84	By	117	LYS	CB-CA-C	5.21	120.82	110.40
85	AA	206	U	C2'-C3'-O3'	5.21	122.03	113.70
85	AA	419	A	C3'-C2'-C1'	-5.21	97.33	101.50
85	AA	794	A	C5-C6-N6	-5.21	119.53	123.70
85	AA	1674	G	N9-C1'-C2'	-5.21	106.27	112.00
85	AA	1879	U	C2-N1-C1'	-5.21	111.45	117.70
85	AA	1896	G	O4'-C1'-N9	-5.21	104.03	108.20
85	AA	2037	A	C5-C6-N6	-5.21	119.53	123.70
34	BA	366	G	N3-C4-N9	5.21	129.12	126.00
34	BA	699	G	C5-C6-O6	-5.21	125.48	128.60
34	BA	1202	G	O3'-P-O5'	-5.21	94.11	104.00
34	BA	1435	A	N1-C6-N6	5.21	121.72	118.60
35	BB	829	C	O4'-C1'-N1	5.21	112.37	108.20
35	BB	1108	G	N3-C4-N9	5.21	129.12	126.00
35	BB	1339	C	C1'-O4'-C4'	-5.21	105.73	109.90
35	BB	1344	U	N1-C2-N3	5.21	118.02	114.90
35	BB	1431	G	C5-C6-N1	5.21	114.10	111.50
35	BB	1513	U	C6-N1-C1'	5.21	128.49	121.20
36	BC	76	C	C3'-C2'-C1'	-5.21	97.33	101.50
36	BC	113	G	C4-C5-N7	5.21	112.88	110.80
40	BG	28	A	C4-N9-C1'	-5.21	116.93	126.30
40	BG	146	C	C5'-C4'-C3'	-5.21	107.67	116.00
41	BH	119	U	C2-N3-C4	-5.21	123.88	127.00
77	Br	79	HIS	CA-CB-CG	-5.21	104.75	113.60
85	AA	878	U	N1-C2-N3	5.21	118.02	114.90
85	AA	1116	G	C2'-C3'-O3'	5.21	122.03	113.70
85	AA	1693	C	O4'-C1'-N1	5.21	112.36	108.20
34	BA	809	U	C5'-C4'-O4'	5.21	115.35	109.10
34	BA	1545	C	C2'-C3'-O3'	5.21	122.03	113.70
35	BB	1084	A	O5'-P-OP1	5.21	116.95	110.70
35	BB	1393	C	C5'-C4'-C3'	-5.21	107.67	116.00
35	BB	1482	A	C2'-C3'-O3'	5.21	122.03	113.70
85	AA	381	A	C2'-C3'-O3'	5.21	122.03	113.70
85	AA	842	G	O4'-C1'-N9	5.21	112.36	108.20
3	A2	189	ASN	CA-CB-CG	-5.20	101.95	113.40
23	AP	248	ASP	CB-CA-C	-5.20	99.99	110.40
34	BA	352	G	C5-C6-O6	-5.20	125.48	128.60
34	BA	751	A	C4'-C3'-C2'	-5.20	97.40	102.60
34	BA	959	G	C8-N9-C4	5.20	108.48	106.40
34	BA	1120	U	C2-N1-C1'	-5.20	111.46	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1263	A	P-O3'-C3'	-5.20	113.46	119.70
34	BA	1333	G	C3'-C2'-C1'	-5.20	97.34	101.50
35	BB	612	A	O4'-C1'-C2'	5.20	112.28	107.60
35	BB	648	G	C4-C5-N7	5.20	112.88	110.80
35	BB	687	C	O4'-C1'-C2'	5.20	112.28	107.60
35	BB	770	G	C4-N9-C1'	-5.20	119.74	126.50
35	BB	915	U	O4'-C1'-N1	5.20	112.36	108.20
35	BB	1340	U	P-O3'-C3'	-5.20	113.46	119.70
35	BB	1377	A	P-O3'-C3'	-5.20	113.46	119.70
37	BD	118	C	O4'-C1'-N1	5.20	112.36	108.20
38	BE	182	U	O3'-P-O5'	-5.20	94.11	104.00
38	BE	208	G	C8-N9-C1'	-5.20	120.23	127.00
40	BG	130	G	N7-C8-N9	-5.20	110.50	113.10
41	BH	28	U	O4'-C1'-N1	5.20	112.36	108.20
59	BZ	79	ILE	CA-C-N	-5.20	105.75	117.20
62	Bc	25	ILE	O-C-N	-5.20	114.37	122.70
67	Bh	6	PHE	CB-CG-CD2	-5.20	117.16	120.80
85	AA	378	A	N1-C2-N3	-5.20	126.70	129.30
85	AA	1554	C	C6-N1-C2	5.20	122.38	120.30
85	AA	1671	G	C1'-O4'-C4'	-5.20	105.74	109.90
86	AB	5	G	C8-N9-C4	-5.20	104.32	106.40
34	BA	953	G	N1-C6-O6	-5.20	116.78	119.90
34	BA	1515	U	C2-N1-C1'	-5.20	111.46	117.70
35	BB	376	A	C4'-C3'-C2'	-5.20	97.40	102.60
35	BB	1021	C	N3-C2-O2	-5.20	118.26	121.90
35	BB	1318	U	P-O3'-C3'	-5.20	113.46	119.70
35	BB	1522	G	O4'-C1'-N9	5.20	112.36	108.20
38	BE	8	G	N3-C2-N2	5.20	123.54	119.90
40	BG	1	G	N9-C1'-C2'	-5.20	106.28	112.00
65	Bf	318	TYR	CB-CG-CD1	5.20	124.12	121.00
85	AA	118	C	C5-C6-N1	-5.20	118.40	121.00
85	AA	131	C	C4'-C3'-C2'	-5.20	97.40	102.60
85	AA	423	G	C5-C6-O6	-5.20	125.48	128.60
85	AA	898	A	C5-C6-N6	-5.20	119.54	123.70
85	AA	1480	C	O4'-C1'-N1	5.20	112.36	108.20
85	AA	1729	C	P-O5'-C5'	-5.20	112.58	120.90
85	AA	2152	C	C2-N1-C1'	-5.20	113.08	118.80
8	A7	136	TRP	CB-CG-CD1	5.20	133.76	127.00
26	AS	33	PHE	CB-CG-CD2	-5.20	117.16	120.80
34	BA	131	A	O3'-P-O5'	-5.20	94.12	104.00
34	BA	255	G	C5'-C4'-O4'	5.20	115.34	109.10
34	BA	715	U	N3-C2-O2	-5.20	118.56	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	746	C	C5'-C4'-C3'	-5.20	107.68	116.00
34	BA	932	G	C6-C5-N7	-5.20	127.28	130.40
34	BA	967	C	C2-N1-C1'	-5.20	113.08	118.80
34	BA	1833	G	O4'-C1'-N9	5.20	112.36	108.20
35	BB	692	G	C4-N9-C1'	-5.20	119.74	126.50
35	BB	832	C	P-O5'-C5'	5.20	129.22	120.90
35	BB	1410	G	C6-N1-C2	-5.20	121.98	125.10
38	BE	73	A	O4'-C1'-N9	5.20	112.36	108.20
41	BH	74	G	O5'-C5'-C4'	5.20	121.58	111.70
60	Ba	52	VAL	N-CA-C	5.20	125.04	111.00
62	Bc	64	THR	N-CA-CB	5.20	120.18	110.30
85	AA	84	C	O3'-P-O5'	5.20	113.88	104.00
85	AA	307	G	C4-N9-C1'	-5.20	119.74	126.50
85	AA	482	C	O4'-C1'-N1	5.20	112.36	108.20
85	AA	1309	G	C4-N9-C1'	-5.20	119.74	126.50
85	AA	1447	U	P-O5'-C5'	-5.20	112.58	120.90
85	AA	1589	G	C1'-O4'-C4'	-5.20	105.74	109.90
85	AA	1977	G	C4-N9-C1'	-5.20	119.74	126.50
85	AA	2141	G	O3'-P-O5'	5.20	113.88	104.00
5	A4	163	ASP	CB-CA-C	5.20	120.80	110.40
34	BA	976	C	P-O5'-C5'	-5.20	112.58	120.90
34	BA	1060	C	C3'-C2'-C1'	-5.20	97.34	101.50
35	BB	363	A	C5-C6-N6	5.20	127.86	123.70
35	BB	557	C	C2-N1-C1'	-5.20	113.08	118.80
35	BB	557	C	O5'-C5'-C4'	-5.20	101.82	111.70
37	BD	118	C	C5'-C4'-C3'	-5.20	107.68	116.00
38	BE	25	U	C4'-C3'-C2'	5.20	107.80	102.60
38	BE	39	U	O5'-C5'-C4'	-5.20	101.82	111.70
38	BE	176	G	P-O5'-C5'	5.20	129.22	120.90
40	BG	66	C	N1-C2-O2	5.20	122.02	118.90
68	Bi	128	ARG	N-CA-C	5.20	125.04	111.00
70	Bk	57	ARG	O-C-N	-5.20	114.38	122.70
78	Bs	26	ARG	NE-CZ-NH2	-5.20	117.70	120.30
79	Bt	38	ARG	N-CA-CB	-5.20	101.24	110.60
85	AA	227	A	O4'-C4'-C3'	-5.20	98.80	104.00
85	AA	622	G	P-O5'-C5'	-5.20	112.58	120.90
85	AA	983	A	C5-N7-C8	-5.20	101.30	103.90
85	AA	1519	A	C3'-C2'-C1'	-5.20	97.34	101.50
85	AA	1790	G	C5-C6-O6	-5.20	125.48	128.60
85	AA	1894	G	C4-N9-C1'	-5.20	119.74	126.50
85	AA	2204	A	O4'-C1'-C2'	5.20	112.28	107.60
85	AA	2225	G	C8-N9-C4	5.20	108.48	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	90	G	C4'-C3'-C2'	-5.20	97.40	102.60
34	BA	237	A	N9-C4-C5	-5.20	103.72	105.80
34	BA	335	C	O5'-P-OP2	-5.20	101.02	105.70
34	BA	813	C	N3-C4-N4	-5.20	114.36	118.00
35	BB	1185	G	C3'-C2'-C1'	-5.20	97.34	101.50
37	BD	110	G	C5-C6-N1	5.20	114.10	111.50
61	Bb	16	PHE	CB-CA-C	-5.20	100.00	110.40
65	Bf	356	TYR	CA-CB-CG	-5.20	103.53	113.40
72	Bm	42	HIS	N-CA-CB	5.20	119.95	110.60
85	AA	36	U	N1-C2-N3	5.20	118.02	114.90
85	AA	179	G	C5'-C4'-C3'	-5.20	107.68	116.00
85	AA	286	C	C5-C4-N4	-5.20	116.56	120.20
85	AA	542	G	O4'-C4'-C3'	-5.20	98.80	104.00
85	AA	735	G	N3-C2-N2	5.20	123.54	119.90
25	AR	38	GLN	CB-CA-C	-5.20	100.01	110.40
34	BA	112	C	O4'-C1'-N1	5.20	112.36	108.20
34	BA	531	C	C2-N3-C4	-5.20	117.30	119.90
34	BA	606	G	C5-C6-N1	5.20	114.10	111.50
34	BA	891	C	C6-N1-C2	5.20	122.38	120.30
34	BA	919	A	C4-N9-C1'	-5.20	116.95	126.30
34	BA	1181	G	C3'-C2'-C1'	-5.20	97.34	101.50
34	BA	1347	G	C1'-O4'-C4'	-5.20	105.74	109.90
34	BA	1596	C	C5-C4-N4	5.20	123.84	120.20
34	BA	1611	A	N9-C4-C5	5.20	107.88	105.80
34	BA	1646	U	C4'-C3'-C2'	5.20	107.80	102.60
35	BB	65	A	N9-C1'-C2'	-5.20	106.28	112.00
35	BB	1234	G	O4'-C1'-N9	5.20	112.36	108.20
36	BC	154	A	C4-C5-C6	-5.20	114.40	117.00
37	BD	87	G	C5-C6-O6	-5.20	125.48	128.60
40	BG	7	U	P-O5'-C5'	5.20	129.21	120.90
40	BG	56	G	C5'-C4'-O4'	5.20	115.34	109.10
41	BH	28	U	C4-C5-C6	-5.20	116.58	119.70
51	BR	123	PRO	C-N-CA	5.20	134.69	121.70
60	Ba	14	SER	C-N-CA	5.20	133.21	122.30
85	AA	300	C	P-O5'-C5'	5.20	129.21	120.90
85	AA	672	U	P-O5'-C5'	-5.20	112.59	120.90
85	AA	814	G	O4'-C1'-N9	5.20	112.36	108.20
85	AA	1439	A	C3'-C2'-C1'	-5.20	97.34	101.50
7	A6	18	PHE	CB-CG-CD1	5.19	124.44	120.80
30	AW	55	VAL	CB-CA-C	-5.19	101.53	111.40
34	BA	265	A	O4'-C1'-N9	5.19	112.36	108.20
34	BA	578	C	N3-C4-N4	5.19	121.64	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	820	C	C4'-C3'-C2'	-5.19	97.41	102.60
34	BA	1171	C	C5'-C4'-C3'	-5.19	107.69	116.00
34	BA	1583	A	C5'-C4'-O4'	5.19	115.33	109.10
34	BA	1656	A	C5'-C4'-O4'	5.19	115.33	109.10
34	BA	1761	U	O4'-C1'-N1	5.19	112.36	108.20
35	BB	802	G	C8-N9-C4	-5.19	104.32	106.40
35	BB	1278	A	N9-C1'-C2'	-5.19	106.29	112.00
36	BC	108	A	C3'-C2'-C1'	5.19	105.66	101.50
40	BG	116	G	O4'-C1'-N9	5.19	112.36	108.20
68	Bi	15	THR	N-CA-CB	5.19	120.17	110.30
85	AA	971	U	C5'-C4'-C3'	5.19	124.31	116.00
85	AA	1070	G	O4'-C1'-N9	5.19	112.36	108.20
3	A2	130	MET	N-CA-CB	5.19	119.95	110.60
34	BA	623	U	O4'-C1'-N1	5.19	112.35	108.20
34	BA	746	C	N1-C1'-C2'	-5.19	106.29	112.00
34	BA	1245	C	C6-N1-C1'	5.19	127.03	120.80
34	BA	1336	U	O4'-C1'-N1	5.19	112.35	108.20
35	BB	380	G	C5-C6-N1	5.19	114.10	111.50
35	BB	474	G	C4-C5-C6	-5.19	115.68	118.80
35	BB	989	C	N3-C2-O2	-5.19	118.27	121.90
35	BB	1220	A	C4'-C3'-C2'	-5.19	97.41	102.60
35	BB	1385	C	C2'-C3'-O3'	5.19	122.01	113.70
35	BB	1516	C	O4'-C1'-N1	5.19	112.35	108.20
36	BC	164	G	C5-C6-O6	-5.19	125.48	128.60
37	BD	13	A	O4'-C1'-N9	5.19	112.35	108.20
39	BF	57	C	C5-C6-N1	-5.19	118.40	121.00
40	BG	109	C	O4'-C1'-C2'	5.19	112.27	107.60
85	AA	530	A	C4-N9-C1'	5.19	135.65	126.30
85	AA	1447	U	C2-N3-C4	-5.19	123.88	127.00
85	AA	1462	A	P-O3'-C3'	-5.19	113.47	119.70
85	AA	1495	G	N1-C6-O6	-5.19	116.78	119.90
85	AA	1593	C	C5-C4-N4	5.19	123.83	120.20
85	AA	1957	C	N1-C1'-C2'	-5.19	106.29	112.00
22	AO	163	GLU	CA-C-N	5.19	128.62	117.20
34	BA	80	U	C4-C5-C6	-5.19	116.58	119.70
34	BA	664	C	C6-N1-C1'	5.19	127.03	120.80
34	BA	920	U	C5'-C4'-C3'	5.19	124.30	116.00
34	BA	1053	U	N3-C2-O2	-5.19	118.57	122.20
35	BB	367	C	C5'-C4'-O4'	5.19	115.33	109.10
35	BB	404	A	N3-C4-N9	-5.19	123.25	127.40
35	BB	453	C	N3-C2-O2	-5.19	118.27	121.90
35	BB	1219	A	P-O5'-C5'	-5.19	112.59	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1237	C	O4'-C1'-N1	5.19	112.35	108.20
36	BC	30	U	N3-C4-O4	-5.19	115.77	119.40
37	BD	82	G	C3'-C2'-C1'	-5.19	97.35	101.50
38	BE	76	U	C4'-C3'-C2'	5.19	107.79	102.60
39	BF	11	C	C4'-C3'-C2'	5.19	107.79	102.60
40	BG	38	A	C4'-C3'-C2'	5.19	107.79	102.60
85	AA	305	A	C3'-C2'-C1'	-5.19	97.35	101.50
85	AA	604	C	OP2-P-O3'	5.19	116.62	105.20
85	AA	1525	C	O5'-C5'-C4'	-5.19	101.84	111.70
85	AA	2074	G	N9-C1'-C2'	-5.19	106.29	112.00
85	AA	2208	G	N9-C4-C5	5.19	107.48	105.40
86	AB	4	C	C5'-C4'-O4'	5.19	115.33	109.10
2	A1	159	TYR	CA-CB-CG	-5.19	103.54	113.40
34	BA	1306	U	N3-C2-O2	-5.19	118.57	122.20
35	BB	518	G	C5-C6-N1	5.19	114.09	111.50
35	BB	1113	C	C1'-O4'-C4'	-5.19	105.75	109.90
35	BB	1128	U	C2-N1-C1'	-5.19	111.47	117.70
35	BB	1163	U	C4'-C3'-C2'	-5.19	97.41	102.60
35	BB	1378	U	C5-C6-N1	-5.19	120.11	122.70
35	BB	1534	U	N3-C2-O2	-5.19	118.57	122.20
36	BC	16	A	C4'-C3'-C2'	-5.19	97.41	102.60
62	Bc	98	ILE	CA-C-N	5.19	128.62	117.20
85	AA	48	G	C4'-C3'-C2'	-5.19	97.41	102.60
85	AA	463	G	O4'-C1'-C2'	5.19	112.27	107.60
85	AA	588	G	C6-N1-C2	-5.19	121.99	125.10
85	AA	2183	U	C5'-C4'-O4'	5.19	115.33	109.10
1	A0	213	ARG	CB-CA-C	-5.19	100.02	110.40
34	BA	48	C	O4'-C1'-N1	5.19	112.35	108.20
34	BA	167	U	N1-C2-N3	5.19	118.01	114.90
34	BA	353	U	O4'-C1'-N1	5.19	112.35	108.20
34	BA	419	U	C2-N3-C4	-5.19	123.89	127.00
34	BA	717	U	P-O3'-C3'	5.19	125.92	119.70
34	BA	845	U	C1'-O4'-C4'	-5.19	105.75	109.90
34	BA	1192	A	C1'-O4'-C4'	-5.19	105.75	109.90
34	BA	1463	U	C2-N1-C1'	-5.19	111.47	117.70
34	BA	1562	G	O4'-C1'-C2'	5.19	112.27	107.60
34	BA	1588	U	N1-C2-O2	5.19	126.43	122.80
35	BB	639	A	P-O3'-C3'	5.19	125.92	119.70
35	BB	1111	C	O4'-C1'-N1	5.19	112.35	108.20
35	BB	1229	A	C8-N9-C4	5.19	107.88	105.80
36	BC	114	C	N1-C2-O2	5.19	122.01	118.90
40	BG	109	C	N1-C1'-C2'	-5.19	106.29	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
42	BI	104	ALA	N-CA-CB	-5.19	102.84	110.10
42	BI	111	ARG	CA-CB-CG	5.19	124.81	113.40
52	BS	1	MET	N-CA-CB	-5.19	101.26	110.60
69	Bj	11	ARG	NE-CZ-NH2	-5.19	117.71	120.30
85	AA	999	A	O5'-C5'-C4'	5.19	121.56	111.70
85	AA	1298	G	C8-N9-C4	-5.19	104.33	106.40
85	AA	1363	U	C4'-C3'-O3'	-5.19	98.51	109.40
85	AA	1403	G	O4'-C1'-N9	5.19	112.35	108.20
85	AA	1948	A	O3'-P-O5'	-5.19	94.14	104.00
85	AA	2122	A	C4-N9-C1'	-5.19	116.96	126.30
34	BA	12	G	O3'-P-O5'	-5.19	94.15	104.00
34	BA	626	G	N3-C2-N2	5.19	123.53	119.90
34	BA	770	G	O5'-P-OP1	-5.19	101.03	105.70
34	BA	1096	C	C6-N1-C1'	-5.19	114.58	120.80
34	BA	1245	C	C2-N1-C1'	-5.19	113.10	118.80
34	BA	1288	U	C1'-O4'-C4'	-5.19	105.75	109.90
34	BA	1671	A	C5-C6-N6	5.19	127.85	123.70
35	BB	517	G	N9-C1'-C2'	-5.19	106.30	112.00
35	BB	656	A	OP1-P-O3'	5.19	116.61	105.20
40	BG	71	C	N3-C4-C5	5.19	123.97	121.90
65	Bf	209	ASP	CA-CB-CG	-5.19	101.99	113.40
85	AA	590	U	C1'-O4'-C4'	-5.19	105.75	109.90
85	AA	696	G	C3'-C2'-C1'	-5.19	97.35	101.50
85	AA	901	C	O3'-P-O5'	5.19	113.85	104.00
85	AA	1170	C	C2-N3-C4	5.19	122.49	119.90
1	A0	192	ARG	NE-CZ-NH1	5.18	122.89	120.30
16	AH	44	GLU	N-CA-CB	5.18	119.93	110.60
34	BA	682	A	N1-C6-N6	5.18	121.71	118.60
34	BA	718	U	O3'-P-O5'	-5.18	94.15	104.00
34	BA	1141	C	C3'-C2'-C1'	-5.18	97.35	101.50
34	BA	1344	G	C3'-C2'-C1'	-5.18	97.35	101.50
34	BA	1502	G	N1-C2-N2	-5.18	111.53	116.20
34	BA	1563	G	N7-C8-N9	5.18	115.69	113.10
34	BA	1647	G	C4-N9-C1'	-5.18	119.76	126.50
34	BA	1667	G	C8-N9-C1'	5.18	133.74	127.00
35	BB	643	G	N1-C2-N2	-5.18	111.53	116.20
35	BB	1417	C	N3-C2-O2	-5.18	118.27	121.90
35	BB	1483	A	C5-N7-C8	-5.18	101.31	103.90
36	BC	107	C	C5-C6-N1	-5.18	118.41	121.00
37	BD	54	A	OP1-P-OP2	-5.18	111.82	119.60
40	BG	39	A	C1'-O4'-C4'	-5.18	105.75	109.90
80	Bu	264	ARG	NE-CZ-NH1	5.18	122.89	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	64	A	P-O3'-C3'	-5.18	113.48	119.70
85	AA	156	G	C4-N9-C1'	-5.18	119.76	126.50
85	AA	1200	A	C4-N9-C1'	-5.18	116.97	126.30
85	AA	1666	U	O3'-P-O5'	-5.18	94.15	104.00
34	BA	179	U	C6-N1-C1'	5.18	128.46	121.20
34	BA	202	A	C5'-C4'-C3'	-5.18	107.71	116.00
34	BA	235	C	C5-C4-N4	-5.18	116.57	120.20
34	BA	879	C	C4'-C3'-C2'	5.18	107.78	102.60
34	BA	890	G	C5-C6-O6	-5.18	125.49	128.60
34	BA	1353	U	C6-N1-C1'	-5.18	113.94	121.20
34	BA	1414	C	C1'-O4'-C4'	-5.18	105.75	109.90
34	BA	1844	U	C6-N1-C2	-5.18	117.89	121.00
35	BB	618	U	N3-C2-O2	-5.18	118.57	122.20
35	BB	925	U	O4'-C1'-N1	5.18	112.35	108.20
35	BB	1045	G	O4'-C1'-N9	5.18	112.35	108.20
35	BB	1064	U	O4'-C1'-N1	5.18	112.35	108.20
35	BB	1091	C	O5'-P-OP2	-5.18	101.04	105.70
35	BB	1396	G	C1'-O4'-C4'	-5.18	105.75	109.90
35	BB	1419	G	C4'-C3'-C2'	-5.18	97.42	102.60
36	BC	124	A	N7-C8-N9	5.18	116.39	113.80
37	BD	62	A	C1'-O4'-C4'	-5.18	105.75	109.90
38	BE	125	C	C5'-C4'-C3'	-5.18	107.71	116.00
39	BF	62	U	O4'-C1'-C2'	-5.18	100.62	105.80
41	BH	24	U	C5'-C4'-O4'	5.18	115.32	109.10
51	BR	4	TYR	CB-CG-CD2	-5.18	117.89	121.00
60	Ba	126	TRP	CA-CB-CG	-5.18	103.85	113.70
74	Bo	34	HIS	CB-CA-C	-5.18	100.03	110.40
85	AA	279	C	C1'-O4'-C4'	-5.18	105.75	109.90
85	AA	384	C	O4'-C1'-C2'	5.18	112.26	107.60
85	AA	703	U	C1'-O4'-C4'	-5.18	105.75	109.90
85	AA	760	U	O5'-C5'-C4'	5.18	121.55	111.70
85	AA	1442	U	C2-N1-C1'	-5.18	111.48	117.70
85	AA	1520	A	C5'-C4'-C3'	-5.18	107.71	116.00
85	AA	1576	G	C2'-C3'-O3'	5.18	121.99	113.70
85	AA	1673	A	N1-C2-N3	-5.18	126.71	129.30
85	AA	2231	G	C8-N9-C1'	-5.18	120.26	127.00
1	A0	105	MET	CA-CB-CG	5.18	122.11	113.30
5	A4	2	SER	C-N-CA	5.18	134.65	121.70
8	A7	273	GLU	C-N-CA	5.18	134.65	121.70
13	AE	40	LYS	CB-CA-C	5.18	120.76	110.40
34	BA	45	A	N1-C6-N6	-5.18	115.49	118.60
34	BA	352	G	C6-N1-C2	-5.18	121.99	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	428	C	O5'-C5'-C4'	-5.18	101.86	111.70
34	BA	930	A	C4-N9-C1'	-5.18	116.97	126.30
35	BB	844	G	C5-C6-O6	-5.18	125.49	128.60
85	AA	190	A	N1-C2-N3	-5.18	126.71	129.30
85	AA	1194	U	C3'-C2'-C1'	-5.18	97.36	101.50
85	AA	1735	U	N3-C2-O2	-5.18	118.57	122.20
1	A0	36	LYS	C-N-CA	5.18	134.65	121.70
34	BA	163	G	C4-N9-C1'	5.18	133.23	126.50
34	BA	164	C	N3-C2-O2	-5.18	118.28	121.90
34	BA	436	U	C2'-C3'-O3'	5.18	121.99	113.70
34	BA	541	C	C1'-O4'-C4'	-5.18	105.76	109.90
34	BA	563	A	O4'-C4'-C3'	-5.18	98.82	104.00
34	BA	1571	C	C2-N1-C1'	-5.18	113.10	118.80
34	BA	1698	C	O4'-C1'-N1	5.18	112.34	108.20
34	BA	1775	U	O3'-P-O5'	5.18	113.84	104.00
35	BB	58	G	N9-C1'-C2'	-5.18	106.30	112.00
35	BB	123	U	O3'-P-O5'	5.18	113.84	104.00
35	BB	330	U	O4'-C1'-N1	5.18	112.34	108.20
35	BB	448	G	N1-C6-O6	-5.18	116.79	119.90
35	BB	772	U	C2'-C3'-O3'	5.18	121.99	113.70
35	BB	1001	G	C6-N1-C2	-5.18	121.99	125.10
38	BE	158	U	C3'-C2'-C1'	-5.18	97.36	101.50
40	BG	89	A	C4-N9-C1'	-5.18	116.98	126.30
40	BG	163	G	O4'-C1'-N9	-5.18	104.06	108.20
76	Bq	45	ARG	NE-CZ-NH1	5.18	122.89	120.30
85	AA	1194	U	O4'-C1'-N1	5.18	112.34	108.20
85	AA	1491	G	N3-C2-N2	-5.18	116.27	119.90
85	AA	1661	U	C1'-O4'-C4'	-5.18	105.76	109.90
85	AA	2039	G	C1'-O4'-C4'	-5.18	105.76	109.90
85	AA	2169	C	N3-C4-N4	-5.18	114.37	118.00
85	AA	2179	C	O5'-C5'-C4'	-5.18	101.86	111.70
34	BA	332	U	C4'-C3'-C2'	-5.18	97.42	102.60
34	BA	905	A	N9-C1'-C2'	-5.18	106.30	112.00
35	BB	582	G	C6-N1-C2	-5.18	121.99	125.10
35	BB	830	G	N9-C1'-C2'	-5.18	106.30	112.00
35	BB	1423	U	O4'-C4'-C3'	-5.18	98.82	104.00
37	BD	108	G	C2'-C3'-O3'	5.18	121.98	113.70
41	BH	4	U	C5-C6-N1	-5.18	120.11	122.70
41	BH	39	G	N3-C4-N9	5.18	129.11	126.00
67	Bh	1	MET	N-CA-CB	-5.18	101.28	110.60
77	Br	308	LYS	N-CA-C	5.18	124.98	111.00
85	AA	87	C	P-O5'-C5'	-5.18	112.61	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	853	G	P-O5'-C5'	-5.18	112.61	120.90
85	AA	874	A	N9-C4-C5	-5.18	103.73	105.80
85	AA	1856	G	N9-C1'-C2'	-5.18	106.30	112.00
11	AC	118	PHE	CA-CB-CG	-5.18	101.48	113.90
13	AE	35	ARG	NE-CZ-NH1	5.18	122.89	120.30
17	AI	130	TYR	N-CA-C	5.18	124.98	111.00
22	AO	73	ARG	NE-CZ-NH1	5.18	122.89	120.30
31	AX	122	ILE	CB-CA-C	5.18	121.95	111.60
34	BA	187	G	C8-N9-C1'	5.18	133.73	127.00
34	BA	683	C	N3-C4-C5	-5.18	119.83	121.90
34	BA	732	A	O5'-C5'-C4'	-5.18	101.86	111.70
34	BA	961	C	C5-C6-N1	5.18	123.59	121.00
35	BB	35	G	C8-N9-C1'	5.18	133.73	127.00
35	BB	669	A	O4'-C1'-C2'	-5.18	100.62	105.80
35	BB	1260	A	C8-N9-C4	5.18	107.87	105.80
35	BB	1357	C	C6-N1-C2	5.18	122.37	120.30
38	BE	116	U	C1'-O4'-C4'	-5.18	105.76	109.90
41	BH	66	G	C5-C6-O6	-5.18	125.49	128.60
41	BH	111	U	C5-C6-N1	-5.18	120.11	122.70
45	BL	23	LYS	N-CA-CB	5.18	119.92	110.60
67	Bh	69	THR	CA-CB-CG2	-5.18	105.15	112.40
85	AA	644	A	O4'-C1'-C2'	5.18	112.26	107.60
85	AA	893	G	C6-N1-C2	-5.18	121.99	125.10
85	AA	1106	A	C2-N3-C4	-5.18	108.01	110.60
85	AA	1576	G	C4-N9-C1'	-5.18	119.77	126.50
86	AB	52	G	N1-C6-O6	5.18	123.00	119.90
2	A1	86	VAL	CA-CB-CG2	-5.17	103.14	110.90
8	A7	13	ARG	NE-CZ-NH2	-5.17	117.71	120.30
20	AL	124	VAL	CA-C-N	5.17	128.58	117.20
34	BA	177	G	N7-C8-N9	-5.17	110.51	113.10
34	BA	295	G	C3'-C2'-C1'	-5.17	97.36	101.50
34	BA	371	U	C5-C4-O4	5.17	129.00	125.90
34	BA	500	C	C6-N1-C1'	5.17	127.01	120.80
34	BA	542	A	C3'-C2'-C1'	-5.17	97.36	101.50
34	BA	1068	C	C6-N1-C1'	5.17	127.01	120.80
34	BA	1243	A	O4'-C1'-N9	5.17	112.34	108.20
35	BB	137	A	O4'-C1'-N9	5.17	112.34	108.20
35	BB	470	C	C6-N1-C1'	5.17	127.01	120.80
35	BB	538	A	OP2-P-O3'	5.17	116.58	105.20
35	BB	723	A	N9-C1'-C2'	-5.17	106.31	112.00
35	BB	1353	G	C2'-C3'-O3'	5.17	121.98	113.70
36	BC	24	G	C5'-C4'-O4'	5.17	115.31	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	18	U	O4'-C1'-C2'	5.17	112.26	107.60
40	BG	21	C	P-O3'-C3'	5.17	125.91	119.70
40	BG	94	G	O5'-C5'-C4'	-5.17	101.87	111.70
47	BN	52	PHE	CB-CA-C	5.17	120.75	110.40
60	Ba	90	ARG	CB-CA-C	-5.17	100.05	110.40
75	Bp	34	PHE	CA-CB-CG	-5.17	101.48	113.90
85	AA	636	G	P-O5'-C5'	-5.17	112.62	120.90
85	AA	1096	G	C6-N1-C2	-5.17	122.00	125.10
85	AA	1485	G	C5-C6-N1	5.17	114.09	111.50
85	AA	1520	A	C1'-O4'-C4'	-5.17	105.76	109.90
85	AA	1596	A	C3'-C2'-C1'	-5.17	97.36	101.50
85	AA	1731	G	O4'-C1'-N9	5.17	112.34	108.20
85	AA	1913	G	O4'-C1'-N9	5.17	112.34	108.20
85	AA	1928	A	C3'-C2'-C1'	-5.17	97.36	101.50
85	AA	1953	G	C8-N9-C1'	5.17	133.73	127.00
34	BA	419	U	C5'-C4'-C3'	5.17	124.28	116.00
34	BA	1433	U	O3'-P-O5'	-5.17	94.17	104.00
35	BB	1201	G	P-O5'-C5'	-5.17	112.62	120.90
51	BR	124	ARG	N-CA-C	5.17	124.97	111.00
70	Bk	67	ARG	N-CA-CB	-5.17	101.29	110.60
85	AA	438	G	O3'-P-O5'	5.17	113.83	104.00
85	AA	930	G	C5-C6-N1	5.17	114.09	111.50
85	AA	1652	A	P-O5'-C5'	5.17	129.18	120.90
20	AL	75	GLU	N-CA-CB	-5.17	101.29	110.60
34	BA	5	C	C3'-C2'-C1'	-5.17	97.36	101.50
34	BA	215	C	C2-N3-C4	-5.17	117.31	119.90
34	BA	492	G	N3-C2-N2	5.17	123.52	119.90
34	BA	493	G	P-O3'-C3'	-5.17	113.49	119.70
34	BA	649	A	C1'-O4'-C4'	-5.17	105.76	109.90
34	BA	1705	C	N1-C1'-C2'	-5.17	106.31	112.00
34	BA	1836	A	C4-N9-C1'	-5.17	116.99	126.30
35	BB	119	G	N3-C2-N2	5.17	123.52	119.90
35	BB	696	G	C4-N9-C1'	-5.17	119.78	126.50
35	BB	971	A	OP2-P-O3'	5.17	116.58	105.20
35	BB	1255	U	C1'-O4'-C4'	-5.17	105.76	109.90
39	BF	14	C	OP1-P-OP2	-5.17	111.84	119.60
45	BL	58	ARG	NE-CZ-NH2	-5.17	117.71	120.30
65	Bf	426	GLY	C-N-CA	5.17	134.63	121.70
71	Bl	123	ARG	NE-CZ-NH1	5.17	122.89	120.30
85	AA	153	C	C3'-C2'-C1'	-5.17	97.36	101.50
85	AA	363	A	N9-C4-C5	5.17	107.87	105.80
85	AA	409	C	C5-C4-N4	5.17	123.82	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	427	G	C4-N9-C1'	-5.17	119.78	126.50
85	AA	463	G	N9-C1'-C2'	-5.17	106.31	112.00
85	AA	804	A	C4-N9-C1'	-5.17	116.99	126.30
85	AA	1153	G	C6-C5-N7	-5.17	127.30	130.40
85	AA	1359	U	C5'-C4'-O4'	5.17	115.31	109.10
34	BA	566	G	O4'-C4'-C3'	-5.17	98.83	104.00
34	BA	1218	G	C5-N7-C8	5.17	106.89	104.30
34	BA	1226	G	C1'-O4'-C4'	-5.17	105.76	109.90
34	BA	1747	C	N3-C2-O2	-5.17	118.28	121.90
35	BB	2	C	C5-C6-N1	-5.17	118.42	121.00
35	BB	27	C	N3-C2-O2	-5.17	118.28	121.90
35	BB	125	G	N3-C2-N2	5.17	123.52	119.90
35	BB	315	C	C2-N3-C4	-5.17	117.31	119.90
41	BH	28	U	C6-N1-C1'	-5.17	113.96	121.20
52	BS	130	SER	N-CA-CB	5.17	118.25	110.50
85	AA	57	G	N3-C4-N9	5.17	129.10	126.00
85	AA	429	G	C5-C6-N1	5.17	114.08	111.50
85	AA	1477	A	O4'-C1'-N9	5.17	112.34	108.20
85	AA	2077	G	C4-N9-C1'	-5.17	119.78	126.50
1	A0	102	PHE	CB-CG-CD2	-5.17	117.18	120.80
34	BA	306	G	N9-C1'-C2'	-5.17	106.31	112.00
34	BA	500	C	N1-C1'-C2'	-5.17	106.31	112.00
34	BA	563	A	C5'-C4'-O4'	-5.17	102.90	109.10
34	BA	592	G	O5'-C5'-C4'	5.17	121.52	111.70
34	BA	991	U	N1-C2-N3	5.17	118.00	114.90
34	BA	1104	C	P-O3'-C3'	5.17	125.90	119.70
34	BA	1243	A	P-O3'-C3'	-5.17	113.50	119.70
34	BA	1311	G	C4-N9-C1'	-5.17	119.78	126.50
34	BA	1547	G	C1'-O4'-C4'	-5.17	105.77	109.90
35	BB	71	A	C3'-C2'-C1'	-5.17	97.36	101.50
35	BB	611	U	N3-C2-O2	-5.17	118.58	122.20
35	BB	1078	U	O4'-C1'-N1	5.17	112.33	108.20
36	BC	149	A	O4'-C1'-N9	5.17	112.33	108.20
36	BC	154	A	C4-N9-C1'	-5.17	117.00	126.30
39	BF	33	C	C2-N3-C4	-5.17	117.31	119.90
40	BG	33	G	C5-C6-N1	-5.17	108.92	111.50
40	BG	112	C	N3-C2-O2	-5.17	118.28	121.90
40	BG	139	U	P-O5'-C5'	-5.17	112.63	120.90
42	BI	191	ARG	C-N-CA	5.17	134.62	121.70
56	BW	58	ASP	N-CA-CB	-5.17	101.30	110.60
68	Bi	55	ASN	CB-CA-C	-5.17	100.06	110.40
85	AA	485	A	C5-C6-N6	-5.17	119.56	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	691	U	N1-C2-N3	5.17	118.00	114.90
85	AA	706	U	O4'-C1'-N1	5.17	112.34	108.20
85	AA	807	A	C4'-C3'-C2'	-5.17	97.43	102.60
85	AA	1191	G	N7-C8-N9	-5.17	110.52	113.10
85	AA	1459	C	P-O3'-C3'	-5.17	113.50	119.70
85	AA	1652	A	O4'-C1'-N9	5.17	112.33	108.20
85	AA	1694	C	P-O3'-C3'	-5.17	113.50	119.70
85	AA	1726	G	P-O3'-C3'	-5.17	113.50	119.70
85	AA	1731	G	C2'-C3'-O3'	5.17	121.97	113.70
85	AA	1916	A	N1-C2-N3	5.17	131.88	129.30
85	AA	2011	C	C5'-C4'-C3'	5.17	124.27	116.00
85	AA	2170	G	C5-C6-O6	-5.17	125.50	128.60
85	AA	2203	C	C1'-O4'-C4'	-5.17	105.77	109.90
4	A3	62	PHE	N-CA-CB	5.17	119.90	110.60
6	A5	180	ARG	NE-CZ-NH2	-5.17	117.72	120.30
14	AF	123	CYS	N-CA-C	-5.17	97.05	111.00
23	AP	191	ALA	N-CA-C	5.17	124.95	111.00
34	BA	528	C	C5'-C4'-O4'	-5.17	102.90	109.10
34	BA	1087	A	N9-C4-C5	-5.17	103.73	105.80
34	BA	1106	A	P-O3'-C3'	-5.17	113.50	119.70
34	BA	1247	G	C4'-C3'-C2'	-5.17	97.43	102.60
34	BA	1346	U	C3'-C2'-C1'	-5.17	97.37	101.50
35	BB	135	C	N1-C1'-C2'	-5.17	106.32	112.00
35	BB	853	U	C4'-C3'-O3'	5.17	123.33	113.00
35	BB	1061	G	N1-C2-N2	-5.17	111.55	116.20
35	BB	1226	G	C4-N9-C1'	-5.17	119.78	126.50
35	BB	1386	C	O4'-C1'-N1	5.17	112.33	108.20
36	BC	61	A	C1'-O4'-C4'	-5.17	105.77	109.90
39	BF	24	G	C5-C6-O6	-5.17	125.50	128.60
40	BG	16	G	C8-N9-C1'	5.17	133.72	127.00
40	BG	27	C	N3-C4-N4	-5.17	114.38	118.00
62	Bc	25	ILE	CB-CA-C	5.17	121.93	111.60
74	Bo	71	LEU	C-N-CA	5.17	134.61	121.70
77	Br	118	GLN	N-CA-CB	-5.17	101.30	110.60
81	Bv	69	SER	N-CA-CB	-5.17	102.75	110.50
85	AA	15	U	C6-N1-C1'	5.17	128.43	121.20
85	AA	832	U	P-O5'-C5'	5.17	129.17	120.90
85	AA	1125	G	N3-C2-N2	-5.17	116.28	119.90
85	AA	1497	U	C2-N1-C1'	-5.17	111.50	117.70
85	AA	1599	G	N1-C6-O6	5.17	123.00	119.90
85	AA	1927	G	O4'-C1'-C2'	5.17	112.25	107.60
85	AA	2216	A	C2'-C3'-O3'	5.17	121.97	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	64	A	P-O3'-C3'	-5.17	113.50	119.70
34	BA	325	A	C3'-C2'-C1'	-5.17	97.37	101.50
34	BA	478	G	N1-C2-N2	-5.17	111.55	116.20
34	BA	536	C	C2-N1-C1'	-5.17	113.12	118.80
34	BA	1427	U	C5'-C4'-C3'	-5.17	107.73	116.00
35	BB	276	U	O4'-C1'-N1	5.17	112.33	108.20
35	BB	339	C	C6-N1-C2	-5.17	118.23	120.30
35	BB	1075	A	P-O3'-C3'	-5.17	113.50	119.70
44	BK	64	ALA	N-CA-CB	-5.17	102.87	110.10
85	AA	153	C	P-O3'-C3'	-5.17	113.50	119.70
85	AA	1261	U	P-O3'-C3'	-5.17	113.50	119.70
85	AA	1519	A	N9-C4-C5	5.17	107.87	105.80
5	A4	121	ARG	CB-CA-C	5.16	120.73	110.40
34	BA	12	G	C4-N9-C1'	-5.16	119.79	126.50
34	BA	127	U	O5'-C5'-C4'	-5.16	101.89	111.70
34	BA	161	U	C5-C6-N1	5.16	125.28	122.70
34	BA	393	G	O5'-C5'-C4'	5.16	121.51	111.70
34	BA	788	C	N1-C1'-C2'	-5.16	106.32	112.00
34	BA	809	U	C6-N1-C1'	5.16	128.43	121.20
34	BA	921	G	N3-C2-N2	5.16	123.51	119.90
34	BA	1601	C	C2'-C3'-O3'	5.16	121.96	113.70
35	BB	486	G	C8-N9-C1'	5.16	133.71	127.00
35	BB	732	G	C3'-C2'-C1'	-5.16	97.37	101.50
35	BB	764	C	C5'-C4'-C3'	5.16	124.26	116.00
35	BB	795	A	O5'-C5'-C4'	-5.16	101.89	111.70
38	BE	21	C	C5'-C4'-O4'	5.16	115.30	109.10
38	BE	52	U	O5'-C5'-C4'	-5.16	101.89	111.70
38	BE	58	U	N3-C4-O4	5.16	123.01	119.40
38	BE	154	A	C5'-C4'-O4'	5.16	115.30	109.10
40	BG	104	A	O4'-C1'-N9	5.16	112.33	108.20
40	BG	112	C	C3'-C2'-C1'	-5.16	97.37	101.50
42	BI	98	ARG	CG-CD-NE	-5.16	100.95	111.80
61	Bb	68	THR	N-CA-CB	-5.16	100.49	110.30
76	Bq	47	THR	N-CA-CB	5.16	120.11	110.30
80	Bu	217	VAL	CA-CB-CG2	-5.16	103.16	110.90
85	AA	197	C	C6-N1-C2	-5.16	118.23	120.30
85	AA	403	G	C5-C6-O6	-5.16	125.50	128.60
85	AA	573	U	C4'-C3'-C2'	5.16	107.76	102.60
85	AA	780	U	C1'-O4'-C4'	-5.16	105.77	109.90
85	AA	1210	U	C2'-C3'-O3'	5.16	121.96	113.70
85	AA	1897	A	C3'-C2'-C1'	5.16	105.63	101.50
85	AA	1918	U	O4'-C1'-N1	5.16	112.33	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2144	C	N3-C4-C5	-5.16	119.83	121.90
85	AA	2248	A	C5-C6-N6	5.16	127.83	123.70
86	AB	6	G	P-O5'-C5'	5.16	129.16	120.90
15	AG	33	VAL	CA-CB-CG2	5.16	118.64	110.90
31	AX	72	LEU	CB-CG-CD1	5.16	119.78	111.00
34	BA	515	U	C2-N1-C1'	-5.16	111.50	117.70
34	BA	804	G	C2'-C3'-O3'	5.16	121.96	113.70
34	BA	954	U	O5'-C5'-C4'	-5.16	101.89	111.70
34	BA	1035	A	C4'-C3'-C2'	5.16	107.76	102.60
34	BA	1590	G	C5'-C4'-O4'	5.16	115.29	109.10
35	BB	364	U	C5'-C4'-C3'	-5.16	107.74	116.00
35	BB	580	A	C3'-C2'-C1'	-5.16	97.37	101.50
35	BB	1231	U	C6-N1-C2	-5.16	117.90	121.00
35	BB	1257	A	P-O3'-C3'	-5.16	113.51	119.70
36	BC	160	C	N1-C2-O2	5.16	122.00	118.90
47	BN	99	GLY	N-CA-C	-5.16	100.20	113.10
85	AA	1124	G	C8-N9-C4	-5.16	104.33	106.40
85	AA	1465	C	N1-C2-O2	5.16	122.00	118.90
85	AA	1790	G	O4'-C1'-N9	-5.16	104.07	108.20
85	AA	1881	C	C2-N1-C1'	-5.16	113.12	118.80
31	AX	202	ASP	N-CA-CB	-5.16	101.31	110.60
34	BA	92	G	N3-C2-N2	5.16	123.51	119.90
34	BA	149	G	C5'-C4'-O4'	5.16	115.29	109.10
34	BA	249	A	C3'-C2'-C1'	5.16	105.63	101.50
34	BA	560	U	N3-C4-O4	5.16	123.01	119.40
34	BA	758	G	N9-C4-C5	-5.16	103.34	105.40
34	BA	988	U	C6-N1-C1'	5.16	128.43	121.20
34	BA	1564	A	O3'-P-O5'	5.16	113.81	104.00
35	BB	128	C	C3'-C2'-C1'	-5.16	97.37	101.50
35	BB	895	U	C2-N1-C1'	-5.16	111.51	117.70
35	BB	1001	G	C5-C6-N1	5.16	114.08	111.50
35	BB	1106	G	N3-C4-C5	-5.16	126.02	128.60
36	BC	140	U	C5-C4-O4	-5.16	122.80	125.90
38	BE	138	U	OP1-P-O3'	5.16	116.55	105.20
40	BG	132	U	N1-C1'-C2'	-5.16	106.32	112.00
47	BN	66	PRO	C-N-CA	5.16	134.60	121.70
85	AA	373	G	O4'-C4'-C3'	-5.16	98.84	104.00
85	AA	392	G	C1'-O4'-C4'	-5.16	105.77	109.90
85	AA	453	G	O4'-C1'-N9	5.16	112.33	108.20
85	AA	585	G	C4-C5-N7	-5.16	108.74	110.80
85	AA	681	G	C1'-O4'-C4'	5.16	114.03	109.90
85	AA	846	U	C2-N1-C1'	5.16	123.89	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2111	C	C2-N1-C1'	-5.16	113.12	118.80
85	AA	2162	G	C4-N9-C1'	-5.16	119.79	126.50
85	AA	2221	A	O4'-C4'-C3'	-5.16	98.84	104.00
85	AA	2246	U	O4'-C1'-N1	5.16	112.33	108.20
86	AB	28	G	C8-N9-C1'	5.16	133.71	127.00
34	BA	147	U	N3-C4-C5	-5.16	111.50	114.60
34	BA	287	U	C2-N1-C1'	5.16	123.89	117.70
34	BA	355	U	O5'-C5'-C4'	-5.16	101.90	111.70
34	BA	425	G	P-O5'-C5'	-5.16	112.65	120.90
34	BA	616	G	N9-C1'-C2'	-5.16	106.33	112.00
34	BA	1153	C	P-O3'-C3'	5.16	125.89	119.70
34	BA	1218	G	O4'-C1'-N9	5.16	112.33	108.20
34	BA	1432	C	P-O5'-C5'	5.16	129.16	120.90
34	BA	1584	G	C5'-C4'-O4'	5.16	115.29	109.10
34	BA	1606	A	O5'-P-OP2	5.16	116.89	110.70
36	BC	39	G	C6-N1-C2	-5.16	122.00	125.10
37	BD	15	U	O4'-C1'-N1	5.16	112.33	108.20
38	BE	45	G	O4'-C1'-N9	5.16	112.33	108.20
38	BE	72	C	N1-C1'-C2'	-5.16	106.33	112.00
40	BG	18	U	P-O3'-C3'	-5.16	113.51	119.70
62	Bc	12	VAL	CA-C-N	5.16	128.55	117.20
85	AA	273	C	P-O5'-C5'	5.16	129.15	120.90
85	AA	438	G	C4'-C3'-C2'	-5.16	97.44	102.60
85	AA	685	U	C5'-C4'-C3'	-5.16	107.75	116.00
85	AA	1776	C	C6-N1-C2	-5.16	118.24	120.30
85	AA	1933	G	O3'-P-O5'	5.16	113.80	104.00
85	AA	2223	C	O4'-C1'-N1	5.16	112.33	108.20
6	A5	83	TYR	CB-CG-CD2	-5.16	117.91	121.00
34	BA	583	G	C4-N9-C1'	-5.16	119.80	126.50
34	BA	1095	G	C5'-C4'-C3'	-5.16	107.75	116.00
34	BA	1577	U	C6-N1-C2	-5.16	117.91	121.00
34	BA	1614	G	P-O5'-C5'	-5.16	112.65	120.90
34	BA	1837	U	C3'-C2'-C1'	-5.16	97.37	101.50
35	BB	625	A	C8-N9-C4	-5.16	103.74	105.80
37	BD	74	A	C3'-C2'-C1'	-5.16	97.37	101.50
38	BE	162	U	O4'-C1'-N1	5.16	112.33	108.20
40	BG	27	C	C5'-C4'-O4'	5.16	115.29	109.10
85	AA	899	A	O5'-C5'-C4'	5.16	121.50	111.70
85	AA	1302	A	C5'-C4'-C3'	5.16	124.25	116.00
85	AA	2130	G	P-O3'-C3'	-5.16	113.51	119.70
2	A1	145	ARG	NE-CZ-NH1	5.16	122.88	120.30
29	AV	6	ARG	NE-CZ-NH1	5.16	122.88	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	458	G	C5-C6-O6	-5.16	125.51	128.60
34	BA	535	G	N1-C2-N2	-5.16	111.56	116.20
34	BA	852	C	O4'-C1'-N1	5.16	112.32	108.20
34	BA	1348	G	N9-C1'-C2'	-5.16	106.33	112.00
34	BA	1556	A	N9-C1'-C2'	-5.16	106.33	112.00
35	BB	96	A	C8-N9-C1'	-5.16	118.42	127.70
35	BB	501	G	C8-N9-C4	-5.16	104.34	106.40
35	BB	1025	A	P-O5'-C5'	-5.16	112.65	120.90
37	BD	93	G	O5'-C5'-C4'	-5.16	101.90	111.70
38	BE	100	U	N3-C2-O2	-5.16	118.59	122.20
39	BF	56	C	C1'-O4'-C4'	5.16	114.02	109.90
40	BG	95	U	O4'-C1'-C2'	5.16	112.24	107.60
47	BN	171	ARG	NE-CZ-NH1	5.16	122.88	120.30
51	BR	139	TYR	CA-CB-CG	5.16	123.19	113.40
85	AA	271	A	C4-N9-C1'	-5.16	117.02	126.30
85	AA	506	G	C1'-O4'-C4'	-5.16	105.78	109.90
85	AA	725	G	C5'-C4'-C3'	-5.16	107.75	116.00
85	AA	915	G	N9-C1'-C2'	-5.16	106.33	112.00
85	AA	983	A	C8-N9-C1'	5.16	136.98	127.70
85	AA	2163	G	C1'-O4'-C4'	-5.16	105.78	109.90
15	AG	127	ARG	NE-CZ-NH2	-5.15	117.72	120.30
34	BA	774	A	P-O3'-C3'	-5.15	113.52	119.70
34	BA	1131	G	N1-C6-O6	5.15	122.99	119.90
34	BA	1437	G	N3-C2-N2	5.15	123.51	119.90
34	BA	1670	A	O4'-C1'-N9	5.15	112.32	108.20
35	BB	716	G	P-O3'-C3'	-5.15	113.52	119.70
35	BB	1054	G	C1'-O4'-C4'	-5.15	105.78	109.90
35	BB	1123	A	C8-N9-C4	5.15	107.86	105.80
35	BB	1397	G	C4'-C3'-C2'	-5.15	97.45	102.60
79	Bt	91	PHE	N-CA-CB	-5.15	101.32	110.60
85	AA	136	U	C4'-C3'-C2'	-5.15	97.45	102.60
85	AA	931	G	C1'-O4'-C4'	-5.15	105.78	109.90
85	AA	993	G	N1-C6-O6	5.15	122.99	119.90
85	AA	1127	G	C5-C6-O6	-5.15	125.51	128.60
85	AA	1215	A	OP1-P-O3'	5.15	116.54	105.20
8	A7	112	VAL	CA-CB-CG2	-5.15	103.17	110.90
34	BA	266	G	N1-C6-O6	-5.15	116.81	119.90
34	BA	387	A	C5-C6-N6	5.15	127.82	123.70
34	BA	624	G	C1'-O4'-C4'	-5.15	105.78	109.90
34	BA	1103	G	N3-C2-N2	5.15	123.51	119.90
34	BA	1176	C	O3'-P-O5'	-5.15	94.21	104.00
34	BA	1202	G	OP1-P-OP2	-5.15	111.87	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	101	U	C4'-C3'-C2'	-5.15	97.45	102.60
35	BB	309	G	O4'-C1'-N9	5.15	112.32	108.20
35	BB	658	G	OP1-P-OP2	-5.15	111.87	119.60
35	BB	669	A	C3'-C2'-C1'	-5.15	97.38	101.50
35	BB	1109	A	C5-C6-N1	5.15	120.28	117.70
35	BB	1237	C	P-O5'-C5'	-5.15	112.66	120.90
35	BB	1307	C	C5'-C4'-C3'	-5.15	107.76	116.00
37	BD	108	G	C1'-O4'-C4'	-5.15	105.78	109.90
38	BE	136	G	C4-N9-C1'	-5.15	119.80	126.50
41	BH	109	G	N3-C4-N9	-5.15	122.91	126.00
48	BO	23	ASP	CB-CG-OD1	5.15	122.94	118.30
70	Bk	88	ARG	NE-CZ-NH1	5.15	122.88	120.30
85	AA	431	G	N3-C2-N2	-5.15	116.29	119.90
85	AA	868	A	C1'-O4'-C4'	-5.15	105.78	109.90
85	AA	1156	A	N9-C1'-C2'	-5.15	106.33	112.00
85	AA	1174	G	N9-C1'-C2'	-5.15	106.33	112.00
85	AA	1323	G	O4'-C1'-N9	5.15	112.32	108.20
85	AA	1469	G	O5'-C5'-C4'	-5.15	101.91	111.70
85	AA	2072	G	O5'-C5'-C4'	-5.15	101.91	111.70
31	AX	168	ASP	N-CA-CB	-5.15	101.33	110.60
34	BA	112	C	C5'-C4'-C3'	-5.15	107.76	116.00
34	BA	236	A	C2'-C3'-O3'	5.15	121.94	113.70
34	BA	504	A	O5'-P-OP1	-5.15	101.06	105.70
34	BA	565	U	P-O3'-C3'	-5.15	113.52	119.70
34	BA	667	U	O5'-C5'-C4'	-5.15	101.91	111.70
34	BA	1336	U	P-O5'-C5'	5.15	129.14	120.90
34	BA	1471	U	N1-C2-O2	5.15	126.41	122.80
34	BA	1524	G	C6-N1-C2	-5.15	122.01	125.10
34	BA	1597	G	C4-N9-C1'	-5.15	119.81	126.50
34	BA	1776	G	O4'-C1'-C2'	5.15	112.23	107.60
35	BB	822	G	C5-N7-C8	5.15	106.88	104.30
40	BG	123	C	N3-C2-O2	-5.15	118.29	121.90
45	BL	127	THR	CA-CB-CG2	-5.15	105.19	112.40
49	BP	119	SER	CB-CA-C	-5.15	100.31	110.10
60	Ba	86	MET	CG-SD-CE	-5.15	91.96	100.20
85	AA	90	A	N1-C6-N6	5.15	121.69	118.60
85	AA	250	C	C6-N1-C1'	5.15	126.98	120.80
85	AA	492	C	P-O3'-C3'	-5.15	113.52	119.70
85	AA	713	G	C4-N9-C1'	5.15	133.20	126.50
85	AA	719	C	P-O3'-C3'	-5.15	113.52	119.70
85	AA	750	A	P-O5'-C5'	5.15	129.14	120.90
85	AA	1445	C	C4'-C3'-C2'	5.15	107.75	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1829	C	C2-N1-C1'	-5.15	113.13	118.80
85	AA	1969	A	C4-N9-C1'	-5.15	117.03	126.30
85	AA	2091	C	O4'-C1'-N1	5.15	112.32	108.20
34	BA	799	A	C6-C5-N7	-5.15	128.70	132.30
34	BA	835	U	C5'-C4'-C3'	-5.15	107.76	116.00
34	BA	1172	C	O4'-C1'-N1	5.15	112.32	108.20
34	BA	1563	G	O3'-P-O5'	5.15	113.78	104.00
35	BB	527	U	C6-N1-C2	-5.15	117.91	121.00
57	BX	55	ARG	N-CA-CB	5.15	119.87	110.60
65	Bf	262	ASP	CB-CG-OD2	5.15	122.93	118.30
80	Bu	99	TYR	CB-CG-CD1	-5.15	117.91	121.00
85	AA	855	G	O3'-P-O5'	-5.15	94.22	104.00
85	AA	1016	G	C8-N9-C1'	5.15	133.69	127.00
85	AA	1458	G	C5-C6-N1	5.15	114.08	111.50
85	AA	1798	U	O3'-P-O5'	5.15	113.78	104.00
86	AB	16	U	C2-N1-C1'	5.15	123.88	117.70
4	A3	45	ALA	N-CA-CB	-5.15	102.89	110.10
34	BA	266	G	N3-C4-C5	-5.15	126.03	128.60
34	BA	805	A	O4'-C1'-N9	5.15	112.32	108.20
34	BA	1247	G	C1'-O4'-C4'	-5.15	105.78	109.90
34	BA	1452	U	C5'-C4'-O4'	5.15	115.28	109.10
34	BA	1614	G	C4-C5-N7	5.15	112.86	110.80
34	BA	1699	A	C8-N9-C4	-5.15	103.74	105.80
35	BB	627	G	O3'-P-O5'	5.15	113.78	104.00
35	BB	857	G	C4-N9-C1'	-5.15	119.81	126.50
35	BB	1005	A	C5'-C4'-C3'	-5.15	107.76	116.00
35	BB	1535	G	C8-N9-C1'	-5.15	120.31	127.00
36	BC	161	U	P-O3'-C3'	-5.15	113.52	119.70
40	BG	94	G	N9-C1'-C2'	-5.15	106.34	112.00
40	BG	135	C	C5'-C4'-C3'	-5.15	107.77	116.00
50	BQ	22	MET	N-CA-CB	-5.15	101.33	110.60
66	Bg	78	ILE	N-CA-C	-5.15	97.10	111.00
85	AA	682	C	P-O5'-C5'	-5.15	112.66	120.90
85	AA	831	C	O4'-C4'-C3'	-5.15	98.85	104.00
85	AA	1110	A	O5'-C5'-C4'	-5.15	101.92	111.70
85	AA	2016	A	N9-C1'-C2'	-5.15	106.34	112.00
85	AA	2180	C	C6-N1-C2	-5.15	118.24	120.30
85	AA	2180	C	N1-C2-N3	5.15	122.80	119.20
34	BA	323	C	C4'-C3'-O3'	5.15	123.29	113.00
34	BA	667	U	C6-N1-C1'	5.15	128.40	121.20
34	BA	735	A	C5-N7-C8	-5.15	101.33	103.90
34	BA	786	U	C4'-C3'-C2'	-5.15	97.45	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1591	G	P-O3'-C3'	-5.15	113.53	119.70
34	BA	1630	A	P-O5'-C5'	-5.15	112.67	120.90
35	BB	694	C	O4'-C1'-N1	5.15	112.32	108.20
35	BB	830	G	N3-C2-N2	5.15	123.50	119.90
35	BB	990	G	C8-N9-C1'	-5.15	120.31	127.00
62	Bc	22	ARG	CG-CD-NE	5.15	122.61	111.80
84	By	66	SER	CB-CA-C	-5.15	100.32	110.10
85	AA	97	A	C4-N9-C1'	-5.15	117.04	126.30
85	AA	1096	G	O4'-C1'-N9	5.15	112.32	108.20
85	AA	1570	A	C4-N9-C1'	5.15	135.56	126.30
85	AA	1619	A	P-O3'-C3'	-5.15	113.53	119.70
16	AH	130	THR	C-N-CA	5.14	134.56	121.70
34	BA	315	U	OP2-P-O3'	5.14	116.52	105.20
34	BA	603	U	P-O5'-C5'	-5.14	112.67	120.90
34	BA	1162	U	O3'-P-O5'	5.14	113.77	104.00
34	BA	1189	A	C5-C6-N6	5.14	127.82	123.70
34	BA	1220	C	N1-C2-O2	5.14	121.99	118.90
34	BA	1283	U	C6-N1-C1'	5.14	128.40	121.20
34	BA	1485	U	C4'-C3'-C2'	5.14	107.74	102.60
34	BA	1673	G	N3-C4-C5	-5.14	126.03	128.60
34	BA	1719	G	C5'-C4'-C3'	-5.14	107.77	116.00
34	BA	1752	A	O4'-C1'-N9	5.14	112.31	108.20
35	BB	386	G	P-O5'-C5'	5.14	129.13	120.90
35	BB	973	G	C5'-C4'-O4'	5.14	115.27	109.10
35	BB	1033	U	C2-N1-C1'	-5.14	111.53	117.70
35	BB	1081	U	C5'-C4'-C3'	5.14	124.23	116.00
35	BB	1473	U	C2-N1-C1'	5.14	123.87	117.70
35	BB	1545	U	C4'-C3'-C2'	-5.14	97.45	102.60
39	BF	55	A	C8-N9-C4	-5.14	103.74	105.80
52	BS	74	ARG	CB-CA-C	5.14	120.69	110.40
79	Bt	93	LEU	N-CA-C	-5.14	97.11	111.00
85	AA	11	A	C5-C6-N1	5.14	120.27	117.70
85	AA	301	U	C1'-O4'-C4'	-5.14	105.78	109.90
85	AA	720	A	O4'-C1'-N9	5.14	112.32	108.20
85	AA	978	U	C5'-C4'-C3'	-5.14	107.77	116.00
85	AA	1421	U	O4'-C1'-C2'	-5.14	100.66	105.80
85	AA	1449	C	C5'-C4'-O4'	5.14	115.27	109.10
85	AA	1873	U	N1-C2-N3	5.14	117.99	114.90
11	AC	202	ARG	C-N-CA	5.14	133.10	122.30
34	BA	24	C	C2-N1-C1'	-5.14	113.14	118.80
34	BA	291	C	C5-C4-N4	-5.14	116.60	120.20
34	BA	470	C	OP1-P-OP2	-5.14	111.89	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	531	C	C3'-C2'-C1'	-5.14	97.39	101.50
34	BA	539	C	C2-N1-C1'	-5.14	113.14	118.80
34	BA	856	G	C6-N1-C2	-5.14	122.02	125.10
34	BA	1137	U	C6-N1-C1'	5.14	128.40	121.20
34	BA	1161	G	C3'-C2'-C1'	-5.14	97.39	101.50
34	BA	1191	C	O3'-P-O5'	-5.14	94.23	104.00
34	BA	1588	U	O4'-C1'-N1	5.14	112.31	108.20
34	BA	1689	U	O4'-C1'-N1	5.14	112.31	108.20
34	BA	1750	A	C8-N9-C1'	5.14	136.96	127.70
35	BB	410	A	N9-C1'-C2'	-5.14	106.34	112.00
35	BB	499	A	C5'-C4'-C3'	-5.14	107.77	116.00
35	BB	579	A	C4-C5-N7	5.14	113.27	110.70
35	BB	743	C	N1-C2-N3	5.14	122.80	119.20
35	BB	1246	C	C5-C4-N4	5.14	123.80	120.20
35	BB	1437	U	C2-N3-C4	-5.14	123.91	127.00
41	BH	54	U	C2-N1-C1'	-5.14	111.53	117.70
42	BI	119	ARG	N-CA-CB	-5.14	101.34	110.60
85	AA	66	U	O5'-C5'-C4'	5.14	121.47	111.70
85	AA	1022	G	N1-C6-O6	5.14	122.99	119.90
85	AA	1241	A	O4'-C1'-N9	5.14	112.31	108.20
85	AA	1864	G	C6-N1-C2	-5.14	122.01	125.10
86	AB	21	A	C4'-C3'-C2'	-5.14	97.46	102.60
34	BA	704	G	C5-C6-N1	5.14	114.07	111.50
34	BA	1065	U	O4'-C1'-N1	5.14	112.31	108.20
35	BB	750	G	P-O5'-C5'	5.14	129.12	120.90
35	BB	824	C	N3-C2-O2	-5.14	118.30	121.90
35	BB	1003	G	N1-C6-O6	-5.14	116.82	119.90
36	BC	73	U	P-O5'-C5'	-5.14	112.67	120.90
40	BG	109	C	C5'-C4'-C3'	-5.14	107.78	116.00
42	BI	84	SER	C-N-CD	-5.14	109.29	120.60
57	BX	139	ASP	CB-CG-OD2	-5.14	113.67	118.30
65	Bf	372	TYR	CB-CG-CD1	5.14	124.08	121.00
77	Br	326	ARG	NE-CZ-NH1	5.14	122.87	120.30
85	AA	486	G	C1'-O4'-C4'	-5.14	105.79	109.90
85	AA	740	A	O4'-C4'-C3'	-5.14	98.86	104.00
85	AA	1226	A	N1-C6-N6	-5.14	115.52	118.60
85	AA	1491	G	O5'-P-OP2	-5.14	101.07	105.70
85	AA	1703	A	C1'-O4'-C4'	-5.14	105.79	109.90
6	A5	117	TYR	CB-CG-CD1	5.14	124.08	121.00
8	A7	244	CYS	N-CA-CB	5.14	119.85	110.60
34	BA	103	G	O4'-C1'-N9	5.14	112.31	108.20
34	BA	144	C	C1'-O4'-C4'	-5.14	105.79	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	161	U	C5'-C4'-O4'	5.14	115.27	109.10
34	BA	235	C	N3-C4-N4	5.14	121.60	118.00
34	BA	327	G	P-O5'-C5'	-5.14	112.68	120.90
34	BA	643	U	C5-C4-O4	5.14	128.98	125.90
34	BA	1025	A	C8-N9-C4	5.14	107.86	105.80
34	BA	1364	G	O3'-P-O5'	-5.14	94.23	104.00
34	BA	1691	G	C5-C6-N1	5.14	114.07	111.50
35	BB	10	C	O3'-P-O5'	5.14	113.77	104.00
35	BB	79	U	C6-N1-C1'	5.14	128.40	121.20
35	BB	589	U	O3'-P-O5'	5.14	113.76	104.00
35	BB	779	C	N1-C2-O2	5.14	121.98	118.90
35	BB	825	U	N3-C2-O2	-5.14	118.60	122.20
35	BB	1305	A	P-O3'-C3'	-5.14	113.53	119.70
35	BB	1522	G	C1'-O4'-C4'	-5.14	105.79	109.90
41	BH	29	G	O4'-C1'-C2'	5.14	112.23	107.60
58	BY	37	ARG	NE-CZ-NH1	5.14	122.87	120.30
65	Bf	456	ARG	NE-CZ-NH1	5.14	122.87	120.30
75	Bp	37	ARG	CD-NE-CZ	-5.14	116.41	123.60
82	Bw	183	MET	N-CA-CB	5.14	119.85	110.60
85	AA	459	C	C6-N1-C2	-5.14	118.24	120.30
85	AA	1871	U	C5'-C4'-C3'	-5.14	107.78	116.00
85	AA	2105	G	C2'-C3'-O3'	5.14	121.92	113.70
19	AK	35	ILE	CB-CA-C	-5.14	101.33	111.60
34	BA	135	G	N9-C1'-C2'	-5.14	106.35	112.00
34	BA	1505	G	C6-N1-C2	-5.14	122.02	125.10
34	BA	1525	G	C8-N9-C4	-5.14	104.34	106.40
35	BB	1503	U	C2-N3-C4	-5.14	123.92	127.00
36	BC	24	G	C4-N9-C1'	-5.14	119.82	126.50
37	BD	83	A	C5'-C4'-O4'	5.14	115.27	109.10
39	BF	41	U	C5-C6-N1	-5.14	120.13	122.70
85	AA	431	G	C3'-C2'-C1'	-5.14	97.39	101.50
85	AA	2074	G	O4'-C1'-C2'	5.14	112.22	107.60
21	AM	116	ARG	CB-CA-C	5.14	120.67	110.40
34	BA	71	G	O3'-P-O5'	5.14	113.76	104.00
34	BA	192	G	C5'-C4'-O4'	-5.14	102.94	109.10
34	BA	417	A	P-O3'-C3'	-5.14	113.54	119.70
34	BA	513	U	O5'-C5'-C4'	-5.14	101.94	111.70
34	BA	889	U	C2'-C3'-O3'	5.14	121.92	113.70
34	BA	1204	U	C2-N3-C4	-5.14	123.92	127.00
34	BA	1260	G	N7-C8-N9	-5.14	110.53	113.10
34	BA	1490	U	C5'-C4'-C3'	5.14	124.22	116.00
34	BA	1514	A	N9-C1'-C2'	-5.14	106.35	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1531	G	C6-N1-C2	-5.14	122.02	125.10
34	BA	1692	U	OP1-P-OP2	-5.14	111.89	119.60
35	BB	778	A	P-O5'-C5'	-5.14	112.68	120.90
35	BB	781	U	O4'-C4'-C3'	-5.14	98.86	104.00
35	BB	1139	A	N9-C1'-C2'	-5.14	106.35	112.00
35	BB	1241	U	C1'-O4'-C4'	-5.14	105.79	109.90
36	BC	30	U	C5-C4-O4	5.14	128.98	125.90
38	BE	200	A	C6-N1-C2	-5.14	115.52	118.60
39	BF	63	U	OP1-P-O3'	5.14	116.50	105.20
40	BG	61	A	C1'-O4'-C4'	-5.14	105.79	109.90
40	BG	159	A	P-O3'-C3'	5.14	125.86	119.70
41	BH	112	U	C6-N1-C1'	5.14	128.39	121.20
50	BQ	66	ARG	CD-NE-CZ	-5.14	116.41	123.60
65	Bf	184	ARG	N-CA-CB	-5.14	101.35	110.60
72	Bm	37	ARG	NE-CZ-NH2	-5.14	117.73	120.30
77	Br	151	LEU	CB-CA-C	5.14	119.96	110.20
85	AA	191	C	O3'-P-O5'	-5.14	94.24	104.00
85	AA	292	C	O4'-C4'-C3'	-5.14	98.86	104.00
85	AA	340	G	C8-N9-C1'	5.14	133.68	127.00
85	AA	1177	G	C6-C5-N7	-5.14	127.32	130.40
85	AA	1770	U	C2-N1-C1'	-5.14	111.54	117.70
85	AA	1867	G	C5-C6-N1	5.14	114.07	111.50
85	AA	1899	A	C4'-C3'-C2'	-5.14	97.46	102.60
85	AA	2119	C	N3-C4-C5	5.14	123.95	121.90
29	AV	62	TYR	CB-CG-CD1	5.13	124.08	121.00
34	BA	337	C	P-O3'-C3'	-5.13	113.54	119.70
34	BA	734	G	O5'-C5'-C4'	-5.13	101.95	111.70
34	BA	1034	U	C1'-O4'-C4'	-5.13	105.79	109.90
34	BA	1267	A	C4-N9-C1'	5.13	135.54	126.30
35	BB	502	C	N3-C4-N4	-5.13	114.41	118.00
35	BB	511	A	O4'-C1'-N9	5.13	112.31	108.20
35	BB	1185	G	C2-N3-C4	5.13	114.47	111.90
36	BC	25	C	C2-N3-C4	-5.13	117.33	119.90
36	BC	71	A	O4'-C1'-C2'	-5.13	100.67	105.80
40	BG	29	U	C6-N1-C1'	5.13	128.39	121.20
45	BL	27	ASN	N-CA-CB	5.13	119.84	110.60
69	Bj	47	TRP	CA-CB-CG	5.13	123.45	113.70
85	AA	543	A	N7-C8-N9	-5.13	111.23	113.80
85	AA	1309	G	O4'-C1'-N9	5.13	112.31	108.20
2	A1	159	TYR	CB-CG-CD1	-5.13	117.92	121.00
2	A1	185	ASN	CA-CB-CG	-5.13	102.11	113.40
34	BA	213	A	C5'-C4'-O4'	-5.13	102.94	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1435	G	P-O3'-C3'	-5.13	113.54	119.70
35	BB	1510	G	O3'-P-O5'	-5.13	94.25	104.00
37	BD	93	G	P-O3'-C3'	-5.13	113.54	119.70
39	BF	35	C	C4'-C3'-C2'	-5.13	97.47	102.60
40	BG	122	G	N7-C8-N9	-5.13	110.53	113.10
52	BS	36	PHE	N-CA-CB	-5.13	101.36	110.60
85	AA	112	A	C5'-C4'-O4'	5.13	115.26	109.10
85	AA	799	G	C6-C5-N7	-5.13	127.32	130.40
85	AA	971	U	O4'-C4'-C3'	-5.13	98.87	104.00
85	AA	1506	U	N1-C1'-C2'	-5.13	106.35	112.00
85	AA	2188	C	N3-C2-O2	-5.13	118.31	121.90
15	AG	25	TRP	N-CA-CB	5.13	119.83	110.60
23	AP	217	ARG	NH1-CZ-NH2	-5.13	113.76	119.40
34	BA	74	A	O4'-C1'-N9	5.13	112.31	108.20
34	BA	408	U	O4'-C1'-N1	5.13	112.31	108.20
34	BA	462	C	O5'-C5'-C4'	-5.13	101.95	111.70
34	BA	596	G	N9-C4-C5	-5.13	103.35	105.40
34	BA	681	G	P-O3'-C3'	5.13	125.86	119.70
34	BA	750	C	C4'-C3'-C2'	-5.13	97.47	102.60
34	BA	919	A	P-O5'-C5'	5.13	129.11	120.90
34	BA	1540	C	C5'-C4'-O4'	5.13	115.26	109.10
34	BA	1568	A	C3'-C2'-C1'	-5.13	97.40	101.50
34	BA	1812	C	C5-C4-N4	-5.13	116.61	120.20
35	BB	29	C	C6-N1-C2	5.13	122.35	120.30
35	BB	474	G	P-O5'-C5'	-5.13	112.69	120.90
35	BB	964	G	N7-C8-N9	-5.13	110.53	113.10
35	BB	1017	U	C1'-O4'-C4'	-5.13	105.80	109.90
35	BB	1195	A	C4-N9-C1'	-5.13	117.06	126.30
38	BE	108	U	N3-C2-O2	-5.13	118.61	122.20
42	BI	185	LYS	N-CA-CB	5.13	119.83	110.60
44	BK	22	PHE	CB-CG-CD1	-5.13	117.21	120.80
47	BN	120	ARG	NE-CZ-NH2	-5.13	117.73	120.30
67	Bh	134	PHE	CB-CG-CD1	-5.13	117.21	120.80
85	AA	445	U	C5'-C4'-O4'	5.13	115.26	109.10
85	AA	2139	G	C4'-C3'-C2'	5.13	107.73	102.60
86	AB	68	C	C2-N3-C4	-5.13	117.33	119.90
20	AL	42	ARG	NE-CZ-NH1	5.13	122.86	120.30
34	BA	330	A	C5'-C4'-C3'	5.13	124.21	116.00
34	BA	502	U	O4'-C1'-C2'	5.13	112.22	107.60
34	BA	557	U	N1-C2-N3	-5.13	111.82	114.90
35	BB	1153	G	N3-C2-N2	5.13	123.49	119.90
36	BC	113	G	C5-N7-C8	-5.13	101.73	104.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	86	G	O4'-C1'-N9	5.13	112.30	108.20
85	AA	97	A	N9-C4-C5	-5.13	103.75	105.80
85	AA	1280	U	C5-C6-N1	-5.13	120.14	122.70
85	AA	1963	G	C5-C6-O6	-5.13	125.52	128.60
7	A6	4	TYR	C-N-CA	5.13	134.52	121.70
34	BA	183	G	C4-N9-C1'	-5.13	119.83	126.50
34	BA	580	U	C5'-C4'-C3'	-5.13	107.79	116.00
34	BA	852	C	N3-C2-O2	-5.13	118.31	121.90
34	BA	1360	G	P-O3'-C3'	5.13	125.86	119.70
35	BB	15	C	C2-N3-C4	-5.13	117.34	119.90
35	BB	427	U	O4'-C1'-N1	5.13	112.30	108.20
35	BB	754	U	P-O3'-C3'	5.13	125.85	119.70
35	BB	1154	C	C4'-C3'-C2'	5.13	107.73	102.60
35	BB	1169	A	C3'-C2'-C1'	-5.13	97.40	101.50
37	BD	110	G	C8-N9-C4	5.13	108.45	106.40
38	BE	185	G	O5'-C5'-C4'	-5.13	101.96	111.70
47	BN	10	HIS	CA-CB-CG	5.13	122.32	113.60
72	Bm	6	ARG	NH1-CZ-NH2	-5.13	113.76	119.40
85	AA	321	C	N3-C4-N4	5.13	121.59	118.00
85	AA	627	A	C4'-C3'-C2'	-5.13	97.47	102.60
85	AA	893	G	C8-N9-C4	5.13	108.45	106.40
85	AA	1003	G	P-O3'-C3'	-5.13	113.55	119.70
85	AA	1189	A	O4'-C1'-N9	5.13	112.30	108.20
85	AA	1469	G	N1-C6-O6	5.13	122.98	119.90
85	AA	1870	C	P-O3'-C3'	-5.13	113.55	119.70
85	AA	1873	U	O4'-C1'-N1	5.13	112.30	108.20
85	AA	1903	G	C1'-O4'-C4'	-5.13	105.80	109.90
85	AA	2187	G	N7-C8-N9	-5.13	110.54	113.10
34	BA	6	C	C5'-C4'-C3'	-5.13	107.80	116.00
34	BA	68	A	P-O3'-C3'	5.13	125.85	119.70
34	BA	81	C	C2-N1-C1'	-5.13	113.16	118.80
34	BA	417	A	C3'-C2'-C1'	-5.13	97.40	101.50
34	BA	503	C	C5'-C4'-C3'	-5.13	107.80	116.00
34	BA	545	U	C6-N1-C2	-5.13	117.92	121.00
34	BA	829	U	C1'-O4'-C4'	-5.13	105.80	109.90
34	BA	1802	C	P-O5'-C5'	-5.13	112.70	120.90
34	BA	1827	C	N1-C1'-C2'	-5.13	106.36	112.00
35	BB	642	G	C1'-O4'-C4'	-5.13	105.80	109.90
35	BB	689	C	C1'-O4'-C4'	-5.13	105.80	109.90
35	BB	1130	U	C2'-C3'-O3'	5.13	121.90	113.70
35	BB	1186	A	N1-C6-N6	-5.13	115.52	118.60
35	BB	1448	U	C4'-C3'-C2'	-5.13	97.47	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1481	C	C5-C6-N1	5.13	123.56	121.00
35	BB	1526	C	O3'-P-O5'	-5.13	94.26	104.00
39	BF	46	G	N1-C6-O6	-5.13	116.82	119.90
40	BG	54	G	C6-N1-C2	-5.13	122.02	125.10
40	BG	75	C	C6-N1-C2	5.13	122.35	120.30
43	BJ	38	TYR	CB-CG-CD2	-5.13	117.92	121.00
48	BO	36	ARG	NE-CZ-NH1	5.13	122.86	120.30
65	Bf	216	HIS	CA-CB-CG	-5.13	104.89	113.60
77	Br	203	ARG	NE-CZ-NH1	5.13	122.86	120.30
85	AA	367	A	N3-C4-N9	-5.13	123.30	127.40
85	AA	441	C	C3'-C2'-C1'	-5.13	97.40	101.50
85	AA	765	U	C1'-O4'-C4'	-5.13	105.80	109.90
85	AA	795	C	P-O5'-C5'	-5.13	112.70	120.90
85	AA	974	U	C5'-C4'-C3'	5.13	124.20	116.00
85	AA	1348	C	C4'-C3'-C2'	-5.13	97.47	102.60
85	AA	1977	G	C3'-C2'-C1'	-5.13	97.40	101.50
34	BA	102	G	N1-C2-N2	-5.12	111.59	116.20
34	BA	1003	A	N1-C6-N6	-5.12	115.53	118.60
34	BA	1398	C	P-O5'-C5'	-5.12	112.70	120.90
35	BB	1406	C	N3-C2-O2	-5.12	118.31	121.90
40	BG	118	U	O4'-C1'-N1	5.12	112.30	108.20
44	BK	171	TRP	CB-CG-CD1	5.12	133.66	127.00
79	Bt	41	ARG	NE-CZ-NH1	5.12	122.86	120.30
85	AA	903	G	C6-N1-C2	-5.12	122.03	125.10
85	AA	1093	C	P-O3'-C3'	-5.12	113.55	119.70
85	AA	1155	A	O5'-C5'-C4'	-5.12	101.96	111.70
34	BA	575	U	C2-N1-C1'	5.12	123.85	117.70
34	BA	1137	U	C5'-C4'-O4'	5.12	115.25	109.10
34	BA	1244	G	N1-C6-O6	5.12	122.97	119.90
34	BA	1457	C	P-O3'-C3'	5.12	125.85	119.70
34	BA	1519	G	C4-N9-C1'	-5.12	119.84	126.50
34	BA	1801	G	C5'-C4'-O4'	-5.12	102.95	109.10
35	BB	462	G	C4-N9-C1'	-5.12	119.84	126.50
35	BB	784	C	N1-C2-O2	5.12	121.97	118.90
35	BB	849	A	C4-N9-C1'	5.12	135.52	126.30
35	BB	969	C	C3'-C2'-C1'	-5.12	97.40	101.50
35	BB	971	A	P-O5'-C5'	5.12	129.10	120.90
35	BB	1047	C	P-O5'-C5'	5.12	129.10	120.90
35	BB	1164	U	C6-N1-C2	-5.12	117.93	121.00
35	BB	1383	C	C3'-C2'-C1'	-5.12	97.40	101.50
35	BB	1397	G	C1'-O4'-C4'	-5.12	105.80	109.90
40	BG	70	C	O4'-C1'-C2'	5.12	112.21	107.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	33	G	N3-C4-C5	5.12	131.16	128.60
44	BK	201	THR	N-CA-CB	5.12	120.03	110.30
55	BV	121	TYR	CB-CG-CD2	-5.12	117.93	121.00
65	Bf	422	LYS	N-CA-C	-5.12	97.16	111.00
77	Br	177	ASP	CB-CA-C	-5.12	100.15	110.40
85	AA	99	U	N3-C4-O4	5.12	122.99	119.40
85	AA	246	C	C4'-C3'-C2'	-5.12	97.48	102.60
85	AA	280	U	O4'-C1'-N1	5.12	112.30	108.20
85	AA	354	C	N1-C2-O2	5.12	121.97	118.90
85	AA	705	G	C8-N9-C1'	5.12	133.66	127.00
85	AA	1102	C	C5-C6-N1	5.12	123.56	121.00
85	AA	2164	G	O4'-C1'-N9	5.12	112.30	108.20
85	AA	2209	U	O4'-C1'-N1	5.12	112.30	108.20
34	BA	228	A	O3'-P-O5'	-5.12	94.27	104.00
34	BA	588	C	C4'-C3'-C2'	-5.12	97.48	102.60
34	BA	757	G	O4'-C4'-C3'	-5.12	98.88	104.00
34	BA	1699	A	C8-N9-C1'	5.12	136.92	127.70
34	BA	1723	U	O5'-P-OP2	-5.12	101.09	105.70
34	BA	1744	C	P-O3'-C3'	-5.12	113.55	119.70
35	BB	34	G	C5'-C4'-O4'	5.12	115.25	109.10
35	BB	404	A	C5'-C4'-C3'	5.12	124.19	116.00
35	BB	411	A	N9-C4-C5	-5.12	103.75	105.80
35	BB	1410	G	C1'-O4'-C4'	-5.12	105.80	109.90
37	BD	20	C	N3-C2-O2	-5.12	118.31	121.90
38	BE	173	G	C5'-C4'-O4'	5.12	115.25	109.10
41	BH	92	A	N7-C8-N9	-5.12	111.24	113.80
41	BH	123	G	N7-C8-N9	-5.12	110.54	113.10
67	Bh	38	TRP	CB-CG-CD1	5.12	133.66	127.00
84	By	154	ARG	NE-CZ-NH1	5.12	122.86	120.30
85	AA	72	C	P-O3'-C3'	-5.12	113.56	119.70
85	AA	204	U	O4'-C4'-C3'	-5.12	98.88	104.00
85	AA	586	G	C4'-C3'-C2'	-5.12	97.48	102.60
85	AA	719	C	C6-N1-C2	-5.12	118.25	120.30
85	AA	1531	G	N1-C6-O6	5.12	122.97	119.90
27	AT	38	CYS	CB-CA-C	5.12	120.64	110.40
34	BA	34	U	C3'-C2'-C1'	-5.12	97.40	101.50
34	BA	207	A	P-O5'-C5'	-5.12	112.71	120.90
34	BA	894	G	N1-C6-O6	-5.12	116.83	119.90
34	BA	1481	U	O5'-C5'-C4'	-5.12	101.97	111.70
35	BB	977	G	OP2-P-O3'	5.12	116.47	105.20
35	BB	1250	A	C8-N9-C1'	5.12	136.92	127.70
40	BG	57	A	C1'-O4'-C4'	-5.12	105.80	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
40	BG	73	U	N1-C2-N3	5.12	117.97	114.90
65	Bf	409	THR	N-CA-CB	5.12	120.03	110.30
85	AA	8	U	C5'-C4'-C3'	5.12	124.19	116.00
85	AA	517	A	C4'-C3'-O3'	5.12	123.24	113.00
85	AA	1135	U	N3-C2-O2	-5.12	118.62	122.20
85	AA	1495	G	C1'-O4'-C4'	-5.12	105.80	109.90
85	AA	2036	A	O4'-C1'-N9	5.12	112.30	108.20
9	A8	35	TYR	CB-CG-CD1	5.12	124.07	121.00
34	BA	161	U	N3-C4-O4	5.12	122.98	119.40
34	BA	166	G	O3'-P-O5'	-5.12	94.28	104.00
34	BA	232	U	C6-N1-C2	-5.12	117.93	121.00
34	BA	1068	C	N3-C2-O2	-5.12	118.32	121.90
34	BA	1347	G	C5-C6-O6	-5.12	125.53	128.60
34	BA	1352	G	C8-N9-C4	-5.12	104.35	106.40
34	BA	1527	G	C4-N9-C1'	-5.12	119.84	126.50
34	BA	1812	C	N3-C4-C5	5.12	123.95	121.90
35	BB	365	U	C6-N1-C2	-5.12	117.93	121.00
35	BB	451	A	N7-C8-N9	-5.12	111.24	113.80
35	BB	475	A	C5-N7-C8	-5.12	101.34	103.90
35	BB	866	A	C5-C6-N6	-5.12	119.61	123.70
35	BB	964	G	C6-N1-C2	-5.12	122.03	125.10
41	BH	8	C	C5'-C4'-O4'	5.12	115.24	109.10
41	BH	52	G	C6-N1-C2	-5.12	122.03	125.10
51	BR	57	CYS	N-CA-CB	5.12	119.81	110.60
59	BZ	91	THR	CA-CB-CG2	-5.12	105.23	112.40
85	AA	106	G	P-O3'-C3'	-5.12	113.56	119.70
85	AA	172	A	C5'-C4'-C3'	5.12	124.19	116.00
85	AA	494	G	N1-C2-N2	-5.12	111.59	116.20
85	AA	675	A	C4-N9-C1'	-5.12	117.09	126.30
85	AA	1160	U	C5'-C4'-C3'	-5.12	107.81	116.00
85	AA	1235	G	C5-N7-C8	-5.12	101.74	104.30
85	AA	1255	C	P-O3'-C3'	-5.12	113.56	119.70
85	AA	1368	G	C8-N9-C1'	-5.12	120.34	127.00
85	AA	1547	G	C5-C6-N1	5.12	114.06	111.50
85	AA	1644	G	P-O3'-C3'	-5.12	113.56	119.70
85	AA	1712	A	C1'-O4'-C4'	-5.12	105.80	109.90
31	AX	19	GLU	N-CA-CB	-5.12	101.39	110.60
34	BA	111	U	C5-C6-N1	-5.12	120.14	122.70
34	BA	701	G	N3-C2-N2	5.12	123.48	119.90
34	BA	702	G	C5'-C4'-O4'	5.12	115.24	109.10
34	BA	1025	A	C5-C6-N6	-5.12	119.61	123.70
38	BE	16	C	C2-N1-C1'	5.12	124.43	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	BF	56	C	P-O5'-C5'	-5.12	112.71	120.90
40	BG	34	A	P-O5'-C5'	-5.12	112.71	120.90
50	BQ	116	ARG	NE-CZ-NH2	-5.12	117.74	120.30
85	AA	821	U	C1'-O4'-C4'	-5.12	105.81	109.90
85	AA	1844	A	O4'-C1'-N9	5.12	112.29	108.20
85	AA	1883	C	N3-C4-N4	5.12	121.58	118.00
15	AG	80	MET	CG-SD-CE	-5.12	92.02	100.20
34	BA	712	C	C5-C4-N4	-5.12	116.62	120.20
34	BA	1104	C	O4'-C1'-C2'	5.12	112.20	107.60
34	BA	1342	C	C2'-C3'-O3'	5.12	121.88	113.70
34	BA	1357	C	N3-C4-N4	-5.12	114.42	118.00
34	BA	1669	C	P-O5'-C5'	5.12	129.09	120.90
35	BB	390	G	N3-C4-C5	-5.12	126.04	128.60
35	BB	1016	C	C6-N1-C1'	5.12	126.94	120.80
35	BB	1094	A	N7-C8-N9	-5.12	111.24	113.80
35	BB	1165	A	P-O5'-C5'	-5.12	112.72	120.90
35	BB	1186	A	C8-N9-C1'	5.12	136.91	127.70
36	BC	96	A	C8-N9-C4	5.12	107.85	105.80
38	BE	14	C	N1-C2-O2	5.12	121.97	118.90
38	BE	35	A	C4-N9-C1'	-5.12	117.09	126.30
38	BE	159	A	O4'-C4'-C3'	-5.12	98.89	104.00
41	BH	25	A	N9-C1'-C2'	5.12	120.65	114.00
41	BH	45	G	N9-C4-C5	-5.12	103.35	105.40
65	Bf	364	THR	CB-CA-C	-5.12	97.79	111.60
74	Bo	58	ASP	C-N-CA	-5.12	111.56	122.30
77	Br	154	GLU	CB-CA-C	-5.12	100.17	110.40
79	Bt	93	LEU	CB-CA-C	-5.12	100.48	110.20
84	By	87	ARG	NE-CZ-NH2	-5.12	117.74	120.30
85	AA	319	U	C5-C6-N1	-5.12	120.14	122.70
85	AA	526	G	O5'-C5'-C4'	-5.12	101.98	111.70
85	AA	687	G	N3-C4-C5	-5.12	126.04	128.60
34	BA	276	C	C3'-C2'-C1'	-5.11	97.41	101.50
34	BA	429	G	C4-C5-N7	-5.11	108.75	110.80
34	BA	469	C	C2-N3-C4	-5.11	117.34	119.90
34	BA	764	G	C5-C6-O6	5.11	131.67	128.60
35	BB	430	A	C4'-C3'-C2'	-5.11	97.49	102.60
35	BB	474	G	N9-C1'-C2'	-5.11	106.38	112.00
35	BB	522	A	N1-C6-N6	-5.11	115.53	118.60
35	BB	541	U	O4'-C1'-N1	5.11	112.29	108.20
35	BB	663	G	N1-C6-O6	-5.11	116.83	119.90
35	BB	802	G	N3-C4-N9	5.11	129.07	126.00
35	BB	856	U	C3'-C2'-C1'	-5.11	97.41	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	976	U	OP1-P-O3'	5.11	116.45	105.20
35	BB	1031	G	N3-C2-N2	5.11	123.48	119.90
35	BB	1051	U	C6-N1-C1'	5.11	128.36	121.20
35	BB	1101	C	C2-N3-C4	-5.11	117.34	119.90
35	BB	1227	G	C1'-O4'-C4'	-5.11	105.81	109.90
36	BC	7	U	O4'-C1'-C2'	-5.11	100.69	105.80
38	BE	152	U	C5-C6-N1	5.11	125.26	122.70
39	BF	2	G	C5'-C4'-C3'	-5.11	107.82	116.00
40	BG	9	G	C2'-C3'-O3'	5.11	121.88	113.70
41	BH	121	A	C5'-C4'-O4'	5.11	115.24	109.10
62	Bc	141	VAL	C-N-CA	5.11	134.49	121.70
85	AA	334	A	C4'-C3'-C2'	5.11	107.71	102.60
85	AA	701	C	N1-C2-N3	5.11	122.78	119.20
85	AA	717	G	C1'-O4'-C4'	-5.11	105.81	109.90
85	AA	777	U	C5-C4-O4	-5.11	122.83	125.90
85	AA	863	C	N3-C4-C5	5.11	123.94	121.90
85	AA	983	A	C2'-C3'-O3'	5.11	121.88	113.70
85	AA	1953	G	C4-N9-C1'	-5.11	119.85	126.50
85	AA	2055	G	P-O5'-C5'	-5.11	112.72	120.90
85	AA	2133	A	O3'-P-O5'	-5.11	94.28	104.00
85	AA	2171	A	C3'-C2'-C1'	-5.11	97.41	101.50
86	AB	63	G	O4'-C1'-C2'	5.11	112.20	107.60
34	BA	505	U	C2-N3-C4	-5.11	123.93	127.00
34	BA	561	U	O4'-C1'-N1	5.11	112.29	108.20
34	BA	1109	G	N3-C2-N2	5.11	123.48	119.90
34	BA	1682	A	C5'-C4'-C3'	-5.11	107.82	116.00
36	BC	96	A	P-O3'-C3'	-5.11	113.57	119.70
1	A0	121	THR	C-N-CA	5.11	134.48	121.70
7	A6	118	ALA	N-CA-CB	5.11	117.26	110.10
11	AC	156	ARG	NE-CZ-NH2	5.11	122.86	120.30
34	BA	26	C	O3'-P-O5'	-5.11	94.29	104.00
34	BA	205	G	O5'-C5'-C4'	-5.11	101.99	111.70
34	BA	290	G	O4'-C1'-C2'	5.11	112.20	107.60
34	BA	335	C	O4'-C1'-N1	5.11	112.29	108.20
34	BA	488	C	C5-C4-N4	-5.11	116.62	120.20
34	BA	544	U	C4-C5-C6	-5.11	116.63	119.70
34	BA	671	C	C6-N1-C2	-5.11	118.26	120.30
34	BA	800	G	N3-C4-N9	5.11	129.07	126.00
34	BA	935	A	C5-C6-N6	-5.11	119.61	123.70
34	BA	965	A	P-O5'-C5'	5.11	129.08	120.90
34	BA	1294	C	C5-C6-N1	-5.11	118.44	121.00
34	BA	1484	A	C3'-C2'-C1'	-5.11	97.41	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1641	G	C4-C5-C6	-5.11	115.73	118.80
34	BA	1655	G	C5-C6-O6	-5.11	125.53	128.60
34	BA	1837	U	N1-C2-O2	5.11	126.38	122.80
35	BB	543	G	C5'-C4'-C3'	-5.11	107.82	116.00
35	BB	858	U	C1'-O4'-C4'	-5.11	105.81	109.90
35	BB	1124	G	C5-C6-N1	5.11	114.06	111.50
35	BB	1223	A	O3'-P-O5'	5.11	113.71	104.00
35	BB	1260	A	N9-C1'-C2'	-5.11	106.38	112.00
35	BB	1365	G	C4-N9-C1'	-5.11	119.86	126.50
35	BB	1487	G	O3'-P-O5'	-5.11	94.29	104.00
37	BD	65	G	C5-C6-O6	-5.11	125.53	128.60
40	BG	150	A	C5'-C4'-C3'	-5.11	107.82	116.00
44	BK	162	ARG	NE-CZ-NH1	5.11	122.86	120.30
49	BP	122	TYR	CA-CB-CG	-5.11	103.69	113.40
85	AA	527	A	O4'-C4'-C3'	-5.11	98.89	104.00
85	AA	586	G	C6-N1-C2	-5.11	122.03	125.10
85	AA	633	C	N3-C2-O2	-5.11	118.32	121.90
85	AA	1092	G	C5'-C4'-C3'	-5.11	107.82	116.00
85	AA	1110	A	N9-C1'-C2'	-5.11	106.38	112.00
85	AA	1114	A	N9-C1'-C2'	-5.11	106.38	112.00
85	AA	1667	C	N3-C2-O2	-5.11	118.32	121.90
85	AA	2208	G	O4'-C1'-N9	5.11	112.29	108.20
26	AS	119	ARG	NE-CZ-NH1	5.11	122.86	120.30
34	BA	12	G	C6-N1-C2	-5.11	122.03	125.10
34	BA	429	G	O4'-C1'-N9	5.11	112.29	108.20
34	BA	955	G	C6-N1-C2	-5.11	122.03	125.10
34	BA	973	U	O5'-P-OP2	-5.11	101.10	105.70
35	BB	157	G	C5-C6-O6	-5.11	125.53	128.60
35	BB	1466	A	N7-C8-N9	5.11	116.36	113.80
47	BN	80	PHE	CB-CG-CD2	-5.11	117.22	120.80
47	BN	204	ASN	C-N-CA	5.11	134.47	121.70
85	AA	565	G	C5-C6-O6	-5.11	125.53	128.60
85	AA	1888	U	O4'-C1'-N1	5.11	112.29	108.20
85	AA	2179	C	N1-C2-O2	5.11	121.97	118.90
34	BA	813	C	O4'-C1'-N1	5.11	112.29	108.20
34	BA	1204	U	C3'-C2'-C1'	-5.11	97.41	101.50
34	BA	1642	A	C4'-C3'-C2'	-5.11	97.49	102.60
34	BA	1797	A	O4'-C1'-N9	5.11	112.29	108.20
34	BA	1809	G	N9-C1'-C2'	-5.11	106.38	112.00
35	BB	148	C	C2'-C3'-O3'	5.11	121.87	113.70
35	BB	401	U	C3'-C2'-C1'	-5.11	97.41	101.50
35	BB	470	C	C2-N1-C1'	-5.11	113.18	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	502	C	N3-C2-O2	-5.11	118.32	121.90
35	BB	682	U	C4-C5-C6	-5.11	116.64	119.70
35	BB	700	C	O4'-C1'-N1	5.11	112.29	108.20
35	BB	1226	G	N1-C2-N2	5.11	120.80	116.20
35	BB	1261	U	P-O5'-C5'	-5.11	112.73	120.90
35	BB	1432	U	C1'-O4'-C4'	-5.11	105.81	109.90
38	BE	73	A	C5'-C4'-O4'	-5.11	102.97	109.10
38	BE	195	G	P-O5'-C5'	5.11	129.07	120.90
70	Bk	83	LYS	N-CA-C	-5.11	97.21	111.00
85	AA	173	A	OP1-P-O3'	5.11	116.44	105.20
85	AA	333	A	C8-N9-C4	5.11	107.84	105.80
85	AA	1115	G	O4'-C4'-C3'	-5.11	98.89	104.00
85	AA	1184	A	C5-C6-N1	5.11	120.25	117.70
85	AA	1970	A	N9-C1'-C2'	-5.11	106.38	112.00
2	A1	192	ILE	CB-CA-C	-5.11	101.39	111.60
7	A6	7	PHE	CB-CG-CD1	5.11	124.37	120.80
34	BA	109	A	C4-N9-C1'	-5.11	117.11	126.30
34	BA	359	G	C4'-C3'-C2'	5.11	107.71	102.60
34	BA	468	A	OP1-P-O3'	5.11	116.43	105.20
34	BA	622	G	C8-N9-C1'	5.11	133.64	127.00
34	BA	668	G	O4'-C1'-N9	5.11	112.28	108.20
34	BA	1276	G	C4-N9-C1'	-5.11	119.86	126.50
34	BA	1517	U	C2-N1-C1'	-5.11	111.57	117.70
34	BA	1601	C	C6-N1-C1'	5.11	126.93	120.80
34	BA	1728	G	C8-N9-C4	5.11	108.44	106.40
35	BB	86	A	OP2-P-O3'	5.11	116.43	105.20
35	BB	967	G	C8-N9-C1'	5.11	133.64	127.00
35	BB	1287	U	C5'-C4'-O4'	5.11	115.23	109.10
35	BB	1479	C	C6-N1-C1'	5.11	126.93	120.80
36	BC	143	C	N1-C2-O2	5.11	121.96	118.90
38	BE	173	G	C5-N7-C8	-5.11	101.75	104.30
38	BE	175	U	C5'-C4'-C3'	5.11	124.17	116.00
47	BN	175	ARG	C-N-CA	5.11	134.46	121.70
63	Bd	58	ALA	O-C-N	-5.11	114.53	122.70
79	Bt	6	LYS	CB-CA-C	-5.11	100.19	110.40
85	AA	4	C	C4'-C3'-C2'	5.11	107.71	102.60
85	AA	39	A	C3'-C2'-C1'	-5.11	97.42	101.50
85	AA	889	G	N9-C4-C5	5.11	107.44	105.40
85	AA	1149	A	C8-N9-C4	5.11	107.84	105.80
85	AA	1186	C	N3-C2-O2	-5.11	118.33	121.90
85	AA	1288	A	N7-C8-N9	5.11	116.35	113.80
85	AA	1292	A	C4'-C3'-C2'	5.11	107.70	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1561	A	C4-C5-C6	-5.11	114.45	117.00
34	BA	190	U	O5'-C5'-C4'	-5.10	102.00	111.70
34	BA	300	C	O4'-C4'-C3'	-5.10	98.90	104.00
34	BA	488	C	N1-C1'-C2'	5.10	120.64	114.00
34	BA	765	U	C2-N3-C4	-5.10	123.94	127.00
34	BA	1104	C	C3'-C2'-C1'	-5.10	97.42	101.50
34	BA	1277	G	C6-N1-C2	-5.10	122.04	125.10
34	BA	1800	G	C5-C6-O6	-5.10	125.54	128.60
35	BB	590	G	C5-C6-O6	5.10	131.66	128.60
35	BB	971	A	C4-N9-C1'	-5.10	117.11	126.30
35	BB	1070	G	O5'-C5'-C4'	-5.10	102.00	111.70
38	BE	36	U	O4'-C1'-N1	5.10	112.28	108.20
50	BQ	144	PHE	CB-CA-C	-5.10	100.19	110.40
85	AA	693	A	P-O5'-C5'	5.10	129.07	120.90
85	AA	1161	U	C1'-O4'-C4'	-5.10	105.82	109.90
85	AA	1409	U	P-O5'-C5'	5.10	129.07	120.90
85	AA	1646	U	O4'-C1'-N1	5.10	112.28	108.20
34	BA	442	G	C3'-C2'-C1'	-5.10	97.42	101.50
34	BA	567	U	C1'-O4'-C4'	-5.10	105.82	109.90
34	BA	859	G	P-O3'-C3'	-5.10	113.58	119.70
34	BA	1499	A	N9-C4-C5	5.10	107.84	105.80
34	BA	1621	U	O4'-C1'-N1	5.10	112.28	108.20
34	BA	1827	C	C5'-C4'-C3'	-5.10	107.84	116.00
35	BB	50	A	P-O3'-C3'	-5.10	113.58	119.70
35	BB	397	C	C1'-O4'-C4'	-5.10	105.82	109.90
35	BB	569	G	O4'-C1'-N9	5.10	112.28	108.20
35	BB	977	G	O5'-P-OP1	-5.10	101.11	105.70
35	BB	1028	C	C2-N1-C1'	-5.10	113.19	118.80
35	BB	1444	U	C5'-C4'-O4'	5.10	115.22	109.10
35	BB	1511	U	N3-C2-O2	-5.10	118.63	122.20
36	BC	87	C	C2'-C3'-O3'	5.10	121.86	113.70
51	BR	139	TYR	N-CA-C	-5.10	97.22	111.00
52	BS	93	VAL	CA-CB-CG2	-5.10	103.25	110.90
67	Bh	1	MET	CB-CG-SD	5.10	127.71	112.40
85	AA	104	C	C1'-O4'-C4'	-5.10	105.82	109.90
85	AA	800	A	N3-C4-N9	-5.10	123.32	127.40
85	AA	851	G	N9-C1'-C2'	-5.10	106.39	112.00
85	AA	1246	G	N3-C4-C5	-5.10	126.05	128.60
85	AA	1593	C	N3-C4-N4	-5.10	114.43	118.00
85	AA	2109	G	O4'-C4'-C3'	-5.10	98.90	104.00
85	AA	2132	A	C3'-C2'-C1'	-5.10	97.42	101.50
5	A4	111	VAL	N-CA-C	5.10	124.77	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	474	A	P-O3'-C3'	-5.10	113.58	119.70
34	BA	957	A	C4'-C3'-O3'	-5.10	98.69	109.40
34	BA	1100	A	O4'-C1'-N9	5.10	112.28	108.20
34	BA	1656	A	C5'-C4'-C3'	-5.10	107.84	116.00
34	BA	1708	A	C4-N9-C1'	-5.10	117.12	126.30
34	BA	1815	G	N1-C2-N2	-5.10	111.61	116.20
35	BB	1168	G	O5'-C5'-C4'	-5.10	102.01	111.70
35	BB	1485	G	C6-N1-C2	-5.10	122.04	125.10
85	AA	384	C	N1-C1'-C2'	-5.10	106.39	112.00
85	AA	1263	G	C1'-O4'-C4'	-5.10	105.82	109.90
85	AA	2095	U	C5'-C4'-O4'	-5.10	102.98	109.10
86	AB	18	G	O4'-C1'-C2'	-5.10	100.70	105.80
34	BA	558	C	C1'-O4'-C4'	-5.10	105.82	109.90
34	BA	799	A	C8-N9-C4	-5.10	103.76	105.80
34	BA	1055	U	C6-N1-C2	-5.10	117.94	121.00
34	BA	1101	A	C5-C6-N6	5.10	127.78	123.70
34	BA	1167	A	C3'-C2'-C1'	-5.10	97.42	101.50
34	BA	1581	G	C4'-C3'-C2'	-5.10	97.50	102.60
34	BA	1776	G	C3'-C2'-C1'	-5.10	97.42	101.50
35	BB	130	G	C5-C6-N1	5.10	114.05	111.50
35	BB	519	A	C1'-O4'-C4'	-5.10	105.82	109.90
35	BB	578	G	C3'-C2'-C1'	-5.10	97.42	101.50
35	BB	845	C	N3-C2-O2	-5.10	118.33	121.90
35	BB	1220	A	C3'-C2'-C1'	-5.10	97.42	101.50
35	BB	1263	A	C5'-C4'-C3'	-5.10	107.84	116.00
35	BB	1390	U	O3'-P-O5'	5.10	113.69	104.00
35	BB	1457	A	N1-C6-N6	5.10	121.66	118.60
37	BD	66	G	C5'-C4'-C3'	-5.10	107.84	116.00
37	BD	76	U	OP2-P-O3'	5.10	116.42	105.20
39	BF	47	C	N3-C2-O2	-5.10	118.33	121.90
40	BG	49	A	O3'-P-O5'	-5.10	94.31	104.00
40	BG	175	G	C8-N9-C4	-5.10	104.36	106.40
53	BT	100	ARG	N-CA-CB	-5.10	101.42	110.60
61	Bb	29	PRO	N-CA-C	5.10	125.36	112.10
85	AA	402	G	C4-N9-C1'	-5.10	119.87	126.50
85	AA	485	A	C4'-C3'-C2'	-5.10	97.50	102.60
85	AA	534	A	N1-C2-N3	-5.10	126.75	129.30
85	AA	576	U	C3'-C2'-C1'	-5.10	97.42	101.50
85	AA	1230	U	O5'-C5'-C4'	-5.10	102.01	111.70
85	AA	1287	C	N3-C2-O2	-5.10	118.33	121.90
85	AA	1554	C	C1'-O4'-C4'	-5.10	105.82	109.90
85	AA	1788	U	C6-N1-C2	-5.10	117.94	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
86	AB	57	G	P-O5'-C5'	-5.10	112.74	120.90
9	A8	20	ARG	NE-CZ-NH1	5.10	122.85	120.30
34	BA	481	A	C5-C6-N1	-5.10	115.15	117.70
34	BA	544	U	O5'-P-OP2	-5.10	101.11	105.70
35	BB	837	A	N1-C2-N3	-5.10	126.75	129.30
35	BB	966	C	C1'-O4'-C4'	-5.10	105.82	109.90
35	BB	1000	U	N1-C1'-C2'	-5.10	106.39	112.00
35	BB	1535	G	C4'-C3'-C2'	-5.10	97.50	102.60
40	BG	48	U	C6-N1-C1'	5.10	128.34	121.20
40	BG	123	C	P-O3'-C3'	-5.10	113.58	119.70
48	BO	24	ILE	N-CA-CB	5.10	122.52	110.80
65	Bf	459	ARG	NE-CZ-NH2	-5.10	117.75	120.30
85	AA	189	G	N3-C4-C5	-5.10	126.05	128.60
85	AA	745	C	C6-N1-C1'	5.10	126.92	120.80
85	AA	762	U	C2-N1-C1'	5.10	123.82	117.70
85	AA	917	A	C4-C5-C6	-5.10	114.45	117.00
85	AA	1197	U	C1'-O4'-C4'	-5.10	105.82	109.90
85	AA	1230	U	C2-N3-C4	-5.10	123.94	127.00
85	AA	1514	A	O4'-C1'-N9	5.10	112.28	108.20
85	AA	1789	C	O4'-C1'-N1	5.10	112.28	108.20
85	AA	2250	U	C2-N3-C4	-5.10	123.94	127.00
34	BA	842	U	C1'-O4'-C4'	-5.10	105.82	109.90
34	BA	1067	G	C5-C6-O6	-5.10	125.54	128.60
34	BA	1706	A	P-O3'-C3'	-5.10	113.58	119.70
40	BG	25	G	O3'-P-O5'	5.10	113.68	104.00
40	BG	41	U	C5'-C4'-O4'	5.10	115.22	109.10
40	BG	181	C	P-O5'-C5'	-5.10	112.75	120.90
67	Bh	73	LEU	C-N-CA	-5.10	108.96	121.70
75	Bp	31	VAL	CA-CB-CG2	5.10	118.54	110.90
85	AA	107	A	C5'-C4'-C3'	-5.10	107.85	116.00
85	AA	791	C	C5'-C4'-C3'	-5.10	107.85	116.00
85	AA	983	A	C5-C6-N6	-5.10	119.62	123.70
85	AA	1379	A	P-O3'-C3'	5.10	125.81	119.70
85	AA	1405	U	P-O3'-C3'	-5.10	113.58	119.70
85	AA	1622	G	N1-C2-N2	-5.10	111.61	116.20
85	AA	1833	C	C2'-C3'-O3'	5.10	121.85	113.70
4	A3	176	ALA	N-CA-CB	-5.09	102.97	110.10
7	A6	149	VAL	N-CA-C	5.09	124.75	111.00
31	AX	88	GLU	N-CA-CB	5.09	119.77	110.60
34	BA	263	G	O4'-C1'-N9	5.09	112.28	108.20
34	BA	384	U	N1-C2-N3	5.09	117.96	114.90
34	BA	525	A	O3'-P-O5'	5.09	113.68	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	618	G	O5'-C5'-C4'	-5.09	102.02	111.70
34	BA	676	G	P-O3'-C3'	-5.09	113.59	119.70
34	BA	764	G	C8-N9-C1'	-5.09	120.38	127.00
34	BA	786	U	O4'-C1'-N1	5.09	112.28	108.20
34	BA	1036	G	C1'-O4'-C4'	-5.09	105.82	109.90
34	BA	1606	A	N1-C6-N6	-5.09	115.54	118.60
34	BA	1665	G	C6-N1-C2	-5.09	122.04	125.10
34	BA	1734	U	N1-C2-N3	5.09	117.96	114.90
35	BB	65	A	C8-N9-C4	5.09	107.84	105.80
35	BB	775	U	C2-N1-C1'	-5.09	111.59	117.70
35	BB	1342	C	N1-C2-O2	5.09	121.96	118.90
39	BF	9	C	N3-C2-O2	-5.09	118.33	121.90
41	BH	13	C	N3-C4-N4	5.09	121.57	118.00
42	BI	187	ARG	N-CA-CB	5.09	119.77	110.60
44	BK	7	ARG	NE-CZ-NH2	-5.09	117.75	120.30
78	Bs	2	MET	CB-CA-C	5.09	120.59	110.40
80	Bu	221	ALA	C-N-CA	5.09	134.44	121.70
85	AA	55	A	C3'-C2'-C1'	-5.09	97.42	101.50
85	AA	980	U	P-O3'-C3'	-5.09	113.59	119.70
85	AA	1024	G	O4'-C1'-N9	5.09	112.28	108.20
85	AA	1772	U	O4'-C1'-N1	5.09	112.28	108.20
85	AA	2011	C	C6-N1-C2	5.09	122.34	120.30
85	AA	2096	G	N3-C2-N2	5.09	123.47	119.90
85	AA	2187	G	C5-C6-N1	5.09	114.05	111.50
5	A4	78	LYS	N-CA-CB	-5.09	101.43	110.60
25	AR	29	ILE	N-CA-CB	5.09	122.51	110.80
34	BA	559	C	C4'-C3'-C2'	-5.09	97.51	102.60
35	BB	439	G	C5'-C4'-O4'	5.09	115.21	109.10
35	BB	1188	A	C5'-C4'-C3'	-5.09	107.85	116.00
35	BB	1442	C	N3-C2-O2	-5.09	118.33	121.90
40	BG	2	U	C6-N1-C2	-5.09	117.94	121.00
40	BG	16	G	N3-C2-N2	5.09	123.47	119.90
15	AG	89	TYR	CB-CG-CD1	-5.09	117.95	121.00
25	AR	13	ASN	CA-C-N	-5.09	106.00	117.20
34	BA	418	G	C6-N1-C2	-5.09	122.05	125.10
34	BA	449	G	C2-N3-C4	5.09	114.45	111.90
34	BA	478	G	N1-C6-O6	-5.09	116.84	119.90
34	BA	683	C	O3'-P-O5'	5.09	113.68	104.00
34	BA	705	C	C2-N1-C1'	-5.09	113.20	118.80
34	BA	932	G	N1-C6-O6	5.09	122.95	119.90
34	BA	1161	G	C5-C6-O6	-5.09	125.55	128.60
34	BA	1220	C	N3-C4-N4	-5.09	114.44	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1537	G	C5'-C4'-O4'	5.09	115.21	109.10
35	BB	330	U	P-O3'-C3'	5.09	125.81	119.70
35	BB	835	C	P-O5'-C5'	5.09	129.05	120.90
35	BB	927	U	P-O3'-C3'	5.09	125.81	119.70
35	BB	999	G	O4'-C4'-C3'	-5.09	98.91	104.00
35	BB	1072	C	P-O5'-C5'	-5.09	112.75	120.90
35	BB	1404	A	N9-C4-C5	-5.09	103.76	105.80
35	BB	1513	U	C1'-O4'-C4'	-5.09	105.83	109.90
36	BC	154	A	C5-C6-N1	5.09	120.25	117.70
39	BF	62	U	C5'-C4'-O4'	5.09	115.21	109.10
52	BS	135	GLU	O-C-N	-5.09	114.55	122.70
85	AA	285	C	N1-C1'-C2'	-5.09	106.40	112.00
85	AA	512	U	P-O5'-C5'	5.09	129.05	120.90
85	AA	518	A	O5'-C5'-C4'	5.09	121.37	111.70
85	AA	1371	C	C4'-C3'-O3'	5.09	123.18	113.00
85	AA	1955	U	C4'-C3'-O3'	5.09	123.18	113.00
2	A1	155	ASP	CB-CG-OD1	5.09	122.88	118.30
26	AS	65	ALA	N-CA-CB	5.09	117.22	110.10
33	AZ	88	ASP	CB-CA-C	5.09	120.58	110.40
34	BA	465	A	C1'-O4'-C4'	-5.09	105.83	109.90
34	BA	487	A	C5'-C4'-O4'	5.09	115.21	109.10
34	BA	523	A	C5'-C4'-C3'	5.09	124.14	116.00
34	BA	784	C	O4'-C1'-C2'	5.09	112.18	107.60
34	BA	1044	A	C5-C6-N1	5.09	120.25	117.70
35	BB	673	C	C2-N3-C4	-5.09	117.36	119.90
35	BB	1109	A	O5'-P-OP2	-5.09	101.12	105.70
35	BB	1123	A	P-O3'-C3'	-5.09	113.59	119.70
35	BB	1304	U	C2-N1-C1'	5.09	123.81	117.70
36	BC	124	A	C5-C6-N1	5.09	120.25	117.70
37	BD	101	A	C1'-O4'-C4'	-5.09	105.83	109.90
38	BE	7	U	C2-N1-C1'	-5.09	111.59	117.70
38	BE	195	G	C2-N3-C4	5.09	114.44	111.90
40	BG	44	G	C5'-C4'-C3'	-5.09	107.86	116.00
65	Bf	233	HIS	CA-C-N	-5.09	106.00	117.20
66	Bg	78	ILE	CB-CA-C	-5.09	101.42	111.60
81	Bv	46	ARG	NE-CZ-NH2	5.09	122.84	120.30
84	By	30	ARG	NE-CZ-NH1	5.09	122.84	120.30
85	AA	514	U	C5'-C4'-C3'	-5.09	107.86	116.00
85	AA	579	U	O3'-P-O5'	-5.09	94.33	104.00
85	AA	609	U	C4'-C3'-C2'	-5.09	97.51	102.60
85	AA	866	U	C2-N1-C1'	-5.09	111.59	117.70
85	AA	887	A	N1-C2-N3	-5.09	126.75	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1191	G	C5'-C4'-C3'	-5.09	107.86	116.00
34	BA	215	C	O4'-C1'-N1	5.09	112.27	108.20
34	BA	868	C	P-O3'-C3'	-5.09	113.59	119.70
35	BB	45	A	C8-N9-C4	5.09	107.83	105.80
35	BB	325	G	O4'-C1'-N9	5.09	112.27	108.20
36	BC	121	G	N1-C6-O6	5.09	122.95	119.90
37	BD	75	G	O4'-C1'-C2'	-5.09	100.71	105.80
38	BE	158	U	O4'-C1'-N1	5.09	112.27	108.20
40	BG	105	A	C4-N9-C1'	-5.09	117.14	126.30
40	BG	114	A	P-O3'-C3'	-5.09	113.59	119.70
85	AA	259	A	C8-N9-C4	5.09	107.83	105.80
85	AA	513	G	C6-N1-C2	-5.09	122.05	125.10
85	AA	601	A	C5'-C4'-C3'	5.09	124.14	116.00
85	AA	2182	A	C4-C5-N7	5.09	113.24	110.70
34	BA	826	C	C6-N1-C1'	5.09	126.90	120.80
34	BA	1082	U	C5'-C4'-O4'	-5.09	103.00	109.10
34	BA	1115	A	O3'-P-O5'	5.09	113.67	104.00
34	BA	1143	U	N1-C2-O2	5.09	126.36	122.80
34	BA	1352	G	C3'-C2'-C1'	-5.09	97.43	101.50
34	BA	1484	A	OP1-P-O3'	5.09	116.39	105.20
34	BA	1615	A	C5'-C4'-C3'	5.09	124.14	116.00
35	BB	1047	C	P-O3'-C3'	-5.09	113.60	119.70
35	BB	1427	A	O4'-C4'-C3'	-5.09	98.91	104.00
35	BB	1455	A	C4-N9-C1'	5.09	135.46	126.30
35	BB	1488	G	N1-C6-O6	-5.09	116.85	119.90
36	BC	8	C	C1'-O4'-C4'	5.09	113.97	109.90
40	BG	175	G	C3'-C2'-C1'	-5.09	97.43	101.50
41	BH	60	A	C3'-C2'-C1'	5.09	105.57	101.50
44	BK	85	PHE	CB-CG-CD1	-5.09	117.24	120.80
85	AA	161	A	O4'-C1'-C2'	5.09	112.18	107.60
85	AA	800	A	C5-C6-N6	-5.09	119.63	123.70
85	AA	872	U	C2-N3-C4	-5.09	123.95	127.00
85	AA	999	A	P-O5'-C5'	5.09	129.04	120.90
85	AA	1710	C	O4'-C1'-N1	5.09	112.27	108.20
85	AA	2019	G	P-O3'-C3'	-5.09	113.60	119.70
11	AC	109	ASP	CB-CG-OD1	5.08	122.88	118.30
20	AL	81	ARG	NE-CZ-NH1	5.08	122.84	120.30
21	AM	4	THR	N-CA-CB	5.08	119.96	110.30
27	AT	100	ARG	NE-CZ-NH2	5.08	122.84	120.30
34	BA	566	G	C5'-C4'-C3'	5.08	124.14	116.00
34	BA	581	U	P-O3'-C3'	-5.08	113.60	119.70
34	BA	742	C	C2-N3-C4	5.08	122.44	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	387	G	P-O3'-C3'	-5.08	113.60	119.70
35	BB	598	C	P-O3'-C3'	-5.08	113.60	119.70
35	BB	850	U	O4'-C1'-N1	5.08	112.27	108.20
35	BB	1273	G	C5'-C4'-C3'	5.08	124.14	116.00
39	BF	70	A	N9-C1'-C2'	-5.08	106.41	112.00
84	By	118	ARG	CG-CD-NE	-5.08	101.12	111.80
85	AA	39	A	O4'-C4'-C3'	-5.08	98.92	104.00
85	AA	595	A	C6-N1-C2	-5.08	115.55	118.60
4	A3	224	ARG	N-CA-CB	-5.08	101.45	110.60
34	BA	98	A	C5'-C4'-C3'	5.08	124.13	116.00
34	BA	361	C	N3-C4-N4	5.08	121.56	118.00
34	BA	369	A	P-O3'-C3'	-5.08	113.60	119.70
34	BA	711	C	P-O5'-C5'	5.08	129.03	120.90
34	BA	827	A	C3'-C2'-C1'	-5.08	97.43	101.50
34	BA	1555	G	C5'-C4'-C3'	-5.08	107.86	116.00
35	BB	129	U	C5-C6-N1	-5.08	120.16	122.70
35	BB	505	G	C1'-O4'-C4'	-5.08	105.83	109.90
35	BB	715	G	C1'-O4'-C4'	-5.08	105.83	109.90
35	BB	733	G	P-O5'-C5'	-5.08	112.77	120.90
35	BB	1302	C	O4'-C1'-N1	5.08	112.27	108.20
35	BB	1369	A	C5-C6-N6	-5.08	119.63	123.70
35	BB	1429	A	C3'-C2'-C1'	-5.08	97.43	101.50
38	BE	17	U	C5'-C4'-O4'	5.08	115.20	109.10
38	BE	163	A	C2-N3-C4	-5.08	108.06	110.60
41	BH	135	U	C4'-C3'-C2'	5.08	107.68	102.60
42	BI	162	ALA	N-CA-CB	5.08	117.22	110.10
47	BN	142	ASP	CB-CA-C	5.08	120.57	110.40
62	Bc	53	ILE	C-N-CA	5.08	134.41	121.70
67	Bh	70	ARG	NE-CZ-NH1	5.08	122.84	120.30
68	Bi	50	GLN	C-N-CA	5.08	134.41	121.70
79	Bt	87	ARG	CD-NE-CZ	-5.08	116.48	123.60
85	AA	233	C	N3-C4-N4	-5.08	114.44	118.00
85	AA	585	G	O3'-P-O5'	-5.08	94.34	104.00
85	AA	1147	A	O3'-P-O5'	5.08	113.66	104.00
85	AA	1257	A	C5-C6-N1	5.08	120.24	117.70
85	AA	1689	G	P-O3'-C3'	-5.08	113.60	119.70
85	AA	1895	C	C2-N1-C1'	5.08	124.39	118.80
85	AA	1916	A	C4'-C3'-C2'	-5.08	97.52	102.60
85	AA	2185	U	N1-C2-N3	-5.08	111.85	114.90
86	AB	39	U	O4'-C4'-C3'	-5.08	98.92	104.00
15	AG	60	ILE	N-CA-C	-5.08	97.28	111.00
31	AX	16	PHE	CB-CG-CD2	-5.08	117.24	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	152	C	O5'-C5'-C4'	5.08	121.36	111.70
34	BA	167	U	N1-C1'-C2'	-5.08	106.41	112.00
34	BA	370	U	N3-C4-C5	5.08	117.65	114.60
34	BA	370	U	N3-C4-O4	-5.08	115.84	119.40
34	BA	733	G	N1-C6-O6	5.08	122.95	119.90
34	BA	839	U	P-O3'-C3'	-5.08	113.60	119.70
34	BA	930	A	C4-C5-C6	-5.08	114.46	117.00
34	BA	936	A	O3'-P-O5'	5.08	113.66	104.00
34	BA	959	G	P-O3'-C3'	5.08	125.80	119.70
34	BA	999	G	O4'-C1'-N9	5.08	112.27	108.20
34	BA	1086	A	P-O5'-C5'	-5.08	112.77	120.90
34	BA	1165	A	C1'-O4'-C4'	-5.08	105.83	109.90
34	BA	1285	G	C5-C6-N1	5.08	114.04	111.50
34	BA	1679	C	OP1-P-OP2	-5.08	111.98	119.60
34	BA	1783	C	C6-N1-C2	-5.08	118.27	120.30
35	BB	5	A	C5-N7-C8	-5.08	101.36	103.90
35	BB	87	G	P-O5'-C5'	5.08	129.03	120.90
35	BB	849	A	N7-C8-N9	5.08	116.34	113.80
35	BB	866	A	O4'-C1'-N9	5.08	112.27	108.20
35	BB	1115	G	C4'-C3'-C2'	-5.08	97.52	102.60
35	BB	1249	G	C5-C6-N1	5.08	114.04	111.50
35	BB	1416	A	C5-C6-N6	5.08	127.77	123.70
36	BC	41	A	C5-N7-C8	-5.08	101.36	103.90
36	BC	89	U	P-O3'-C3'	-5.08	113.60	119.70
37	BD	92	G	N3-C2-N2	5.08	123.46	119.90
38	BE	60	C	C4'-C3'-C2'	-5.08	97.52	102.60
38	BE	164	C	O4'-C1'-N1	5.08	112.26	108.20
61	Bb	52	TYR	CB-CG-CD2	-5.08	117.95	121.00
81	Bv	75	ARG	CA-CB-CG	5.08	124.58	113.40
85	AA	339	A	N9-C1'-C2'	-5.08	106.41	112.00
85	AA	610	C	C1'-O4'-C4'	-5.08	105.83	109.90
85	AA	1201	A	P-O3'-C3'	-5.08	113.60	119.70
85	AA	1512	U	P-O5'-C5'	-5.08	112.77	120.90
85	AA	1525	C	N1-C2-O2	5.08	121.95	118.90
85	AA	1579	A	P-O5'-C5'	-5.08	112.77	120.90
85	AA	1833	C	C5'-C4'-C3'	-5.08	107.87	116.00
86	AB	5	G	C5-C6-O6	-5.08	125.55	128.60
34	BA	343	G	OP1-P-OP2	-5.08	111.98	119.60
34	BA	619	U	O5'-C5'-C4'	-5.08	102.05	111.70
34	BA	1122	G	N9-C1'-C2'	-5.08	106.41	112.00
35	BB	53	C	C5'-C4'-O4'	-5.08	103.00	109.10
35	BB	887	G	N7-C8-N9	5.08	115.64	113.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	1407	U	C5-C6-N1	-5.08	120.16	122.70
40	BG	26	G	N9-C1'-C2'	-5.08	106.41	112.00
44	BK	79	ARG	CG-CD-NE	-5.08	101.13	111.80
57	BX	129	THR	CA-CB-CG2	-5.08	105.29	112.40
65	Bf	413	THR	N-CA-C	-5.08	97.28	111.00
85	AA	763	U	C6-N1-C1'	5.08	128.31	121.20
17	AI	42	ALA	C-N-CA	5.08	134.40	121.70
26	AS	22	TRP	CB-CG-CD1	5.08	133.60	127.00
34	BA	961	C	P-O3'-C3'	-5.08	113.61	119.70
34	BA	1088	G	N7-C8-N9	-5.08	110.56	113.10
34	BA	1256	A	O3'-P-O5'	5.08	113.65	104.00
34	BA	1345	U	P-O3'-C3'	5.08	125.80	119.70
34	BA	1533	G	C5-C6-N1	5.08	114.04	111.50
34	BA	1617	U	OP2-P-O3'	5.08	116.37	105.20
35	BB	108	G	C5-C6-N1	5.08	114.04	111.50
35	BB	152	G	P-O3'-C3'	5.08	125.80	119.70
35	BB	324	A	O4'-C1'-N9	5.08	112.26	108.20
35	BB	455	G	C4'-C3'-C2'	-5.08	97.52	102.60
35	BB	507	G	N1-C2-N2	-5.08	111.63	116.20
35	BB	723	A	C8-N9-C4	-5.08	103.77	105.80
35	BB	1052	G	C1'-O4'-C4'	-5.08	105.84	109.90
35	BB	1210	U	O4'-C1'-N1	5.08	112.26	108.20
35	BB	1342	C	C5'-C4'-O4'	5.08	115.19	109.10
38	BE	124	G	N9-C1'-C2'	-5.08	106.42	112.00
40	BG	75	C	O4'-C1'-C2'	5.08	112.17	107.60
41	BH	1	U	C5'-C4'-O4'	5.08	115.19	109.10
50	BQ	102	ASN	N-CA-C	5.08	124.71	111.00
77	Br	155	ASP	CB-CG-OD2	-5.08	113.73	118.30
81	Bv	171	MET	N-CA-CB	5.08	119.74	110.60
85	AA	36	U	O5'-C5'-C4'	-5.08	102.05	111.70
85	AA	308	U	C6-N1-C1'	-5.08	114.09	121.20
85	AA	329	G	O4'-C1'-N9	5.08	112.26	108.20
85	AA	928	U	N1-C2-N3	5.08	117.95	114.90
85	AA	1146	C	P-O3'-C3'	-5.08	113.61	119.70
85	AA	1360	C	O4'-C1'-N1	5.08	112.26	108.20
85	AA	1647	G	P-O3'-C3'	-5.08	113.61	119.70
85	AA	1825	A	N1-C6-N6	-5.08	115.55	118.60
85	AA	2157	G	P-O3'-C3'	-5.08	113.61	119.70
85	AA	2175	U	C5'-C4'-C3'	-5.08	107.88	116.00
86	AB	28	G	C4-N9-C1'	-5.08	119.90	126.50
26	AS	30	SER	N-CA-CB	-5.08	102.88	110.50
34	BA	876	C	C4'-C3'-C2'	5.08	107.68	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1039	G	C5'-C4'-O4'	5.08	115.19	109.10
34	BA	1052	G	C1'-O4'-C4'	-5.08	105.84	109.90
34	BA	1670	A	C5'-C4'-O4'	5.08	115.19	109.10
35	BB	70	A	C5-C6-N6	5.08	127.76	123.70
35	BB	365	U	C5'-C4'-O4'	5.08	115.19	109.10
39	BF	39	C	P-O5'-C5'	5.08	129.02	120.90
85	AA	807	A	C3'-C2'-C1'	-5.08	97.44	101.50
85	AA	1225	C	C5'-C4'-C3'	-5.08	107.88	116.00
85	AA	1999	C	P-O3'-C3'	-5.08	113.61	119.70
85	AA	2059	A	C1'-O4'-C4'	-5.08	105.84	109.90
23	AP	226	TYR	CB-CG-CD2	-5.08	117.95	121.00
34	BA	548	G	C1'-O4'-C4'	5.08	113.96	109.90
34	BA	636	G	OP2-P-O3'	5.08	116.36	105.20
34	BA	1553	G	O4'-C1'-N9	5.08	112.26	108.20
35	BB	609	G	C1'-O4'-C4'	-5.08	105.84	109.90
35	BB	694	C	C6-N1-C2	-5.08	118.27	120.30
35	BB	716	G	N3-C4-C5	-5.08	126.06	128.60
35	BB	1076	U	P-O5'-C5'	-5.08	112.78	120.90
35	BB	1263	A	C6-N1-C2	-5.08	115.56	118.60
35	BB	1272	G	N1-C2-N2	-5.08	111.63	116.20
35	BB	1314	G	O5'-C5'-C4'	-5.08	102.06	111.70
35	BB	1347	C	N1-C2-N3	5.08	122.75	119.20
36	BC	9	G	C5-C6-N1	5.08	114.04	111.50
40	BG	161	C	C6-N1-C2	-5.08	118.27	120.30
44	BK	69	ARG	CD-NE-CZ	-5.08	116.49	123.60
53	BT	120	TYR	CA-CB-CG	5.08	123.04	113.40
53	BT	190	ARG	NE-CZ-NH1	5.08	122.84	120.30
77	Br	368	ALA	CB-CA-C	5.08	117.71	110.10
85	AA	336	C	O4'-C1'-N1	5.08	112.26	108.20
85	AA	1291	A	C5-C6-N6	5.08	127.76	123.70
85	AA	1684	U	C4'-C3'-C2'	-5.08	97.53	102.60
85	AA	1692	U	C2-N1-C1'	-5.08	111.61	117.70
85	AA	1967	A	P-O5'-C5'	5.08	129.02	120.90
85	AA	2082	C	C4-C5-C6	5.08	119.94	117.40
85	AA	2162	G	P-O3'-C3'	-5.08	113.61	119.70
85	AA	2193	A	P-O5'-C5'	-5.08	112.78	120.90
85	AA	2195	A	C3'-C2'-C1'	-5.08	97.44	101.50
15	AG	10	GLY	N-CA-C	-5.07	100.42	113.10
15	AG	123	HIS	CB-CA-C	-5.07	100.25	110.40
34	BA	644	C	O5'-P-OP2	5.07	116.79	110.70
34	BA	803	U	P-O5'-C5'	-5.07	112.78	120.90
34	BA	1039	G	C1'-O4'-C4'	-5.07	105.84	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1145	U	C2-N3-C4	-5.07	123.95	127.00
34	BA	1634	A	N9-C1'-C2'	5.07	120.60	114.00
34	BA	1713	U	C5'-C4'-C3'	5.07	124.12	116.00
35	BB	8	U	O5'-C5'-C4'	-5.07	102.06	111.70
35	BB	428	G	P-O5'-C5'	5.07	129.02	120.90
35	BB	486	G	O4'-C1'-C2'	5.07	112.17	107.60
35	BB	653	G	N9-C4-C5	5.07	107.43	105.40
35	BB	682	U	C3'-C2'-C1'	-5.07	97.44	101.50
35	BB	1074	U	N3-C2-O2	-5.07	118.65	122.20
35	BB	1266	A	C1'-O4'-C4'	-5.07	105.84	109.90
36	BC	100	U	N3-C2-O2	-5.07	118.65	122.20
37	BD	1	G	C2-N3-C4	5.07	114.44	111.90
37	BD	94	C	N3-C4-N4	5.07	121.55	118.00
38	BE	6	A	OP2-P-O3'	5.07	116.36	105.20
38	BE	10	G	C5-C6-N1	-5.07	108.96	111.50
39	BF	1	C	C6-N1-C2	-5.07	118.27	120.30
41	BH	28	U	C4'-C3'-C2'	5.07	107.67	102.60
80	Bu	299	VAL	N-CA-C	-5.07	97.30	111.00
81	Bv	80	TYR	CA-CB-CG	-5.07	103.76	113.40
85	AA	12	U	C2-N1-C1'	-5.07	111.61	117.70
85	AA	24	U	C6-N1-C1'	5.07	128.30	121.20
85	AA	146	U	C5-C6-N1	-5.07	120.16	122.70
85	AA	168	A	C3'-C2'-C1'	-5.07	97.44	101.50
85	AA	1065	G	O4'-C1'-N9	5.07	112.26	108.20
85	AA	1143	C	O5'-C5'-C4'	-5.07	102.06	111.70
85	AA	1207	C	O4'-C4'-C3'	5.07	110.16	106.10
85	AA	1668	G	O4'-C1'-N9	5.07	112.26	108.20
85	AA	2188	C	C5'-C4'-O4'	5.07	115.19	109.10
85	AA	2211	G	O4'-C1'-N9	5.07	112.26	108.20
5	A4	94	ASN	CA-CB-CG	-5.07	102.24	113.40
34	BA	771	A	OP1-P-OP2	-5.07	111.99	119.60
34	BA	1054	U	C6-N1-C1'	5.07	128.30	121.20
35	BB	718	G	C4-N9-C1'	-5.07	119.91	126.50
35	BB	1490	G	P-O5'-C5'	5.07	129.02	120.90
57	BX	51	ARG	NE-CZ-NH1	5.07	122.84	120.30
66	Bg	107	GLY	C-N-CA	5.07	134.38	121.70
85	AA	35	U	O5'-C5'-C4'	-5.07	102.06	111.70
85	AA	2045	U	C5'-C4'-C3'	-5.07	107.88	116.00
86	AB	48	C	C5'-C4'-O4'	5.07	115.19	109.10
22	AO	111	ARG	CB-CA-C	-5.07	100.26	110.40
34	BA	229	C	C5'-C4'-C3'	-5.07	107.89	116.00
34	BA	351	A	C6-N1-C2	-5.07	115.56	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	397	A	P-O5'-C5'	-5.07	112.79	120.90
34	BA	483	A	O4'-C1'-C2'	-5.07	100.73	105.80
34	BA	571	G	C3'-C2'-C1'	5.07	105.56	101.50
34	BA	670	U	P-O3'-C3'	-5.07	113.62	119.70
34	BA	812	A	C1'-O4'-C4'	-5.07	105.84	109.90
34	BA	870	C	P-O3'-C3'	-5.07	113.61	119.70
34	BA	909	G	C4-C5-C6	-5.07	115.76	118.80
34	BA	1240	G	O3'-P-O5'	-5.07	94.36	104.00
34	BA	1717	C	C5'-C4'-C3'	-5.07	107.89	116.00
35	BB	561	C	O4'-C4'-C3'	-5.07	98.93	104.00
35	BB	753	A	C1'-O4'-C4'	-5.07	105.84	109.90
35	BB	835	C	C4'-C3'-O3'	-5.07	98.75	109.40
35	BB	1387	C	N1-C2-N3	5.07	122.75	119.20
36	BC	155	C	C6-N1-C2	5.07	122.33	120.30
37	BD	70	C	C3'-C2'-C1'	-5.07	97.44	101.50
41	BH	48	G	N7-C8-N9	5.07	115.64	113.10
41	BH	121	A	C5-N7-C8	-5.07	101.36	103.90
47	BN	124	TYR	CB-CG-CD2	-5.07	117.96	121.00
50	BQ	89	LYS	CB-CA-C	-5.07	100.26	110.40
67	Bh	126	ARG	NE-CZ-NH1	5.07	122.83	120.30
85	AA	308	U	O4'-C4'-C3'	-5.07	98.93	104.00
85	AA	863	C	C6-N1-C2	-5.07	118.27	120.30
85	AA	1483	A	C4-N9-C1'	-5.07	117.17	126.30
85	AA	1973	G	C8-N9-C1'	5.07	133.59	127.00
8	A7	206	PRO	C-N-CA	5.07	134.37	121.70
25	AR	47	ASN	C-N-CA	-5.07	111.66	122.30
34	BA	550	U	N3-C2-O2	-5.07	118.65	122.20
34	BA	910	U	C5-C6-N1	-5.07	120.17	122.70
35	BB	366	G	P-O3'-C3'	-5.07	113.62	119.70
35	BB	703	U	C2-N1-C1'	-5.07	111.62	117.70
35	BB	1154	C	N1-C2-O2	5.07	121.94	118.90
36	BC	88	A	OP1-P-O3'	5.07	116.35	105.20
40	BG	137	G	C6-N1-C2	-5.07	122.06	125.10
41	BH	69	C	N3-C2-O2	-5.07	118.35	121.90
62	Bc	18	PHE	CB-CG-CD1	5.07	124.35	120.80
85	AA	74	U	O4'-C1'-N1	5.07	112.26	108.20
85	AA	735	G	O4'-C1'-N9	5.07	112.26	108.20
85	AA	1627	U	O4'-C1'-N1	5.07	112.25	108.20
11	AC	240	TYR	CB-CG-CD1	-5.07	117.96	121.00
34	BA	136	A	C2'-C3'-O3'	5.07	121.81	113.70
34	BA	1426	A	OP2-P-O3'	5.07	116.35	105.20
35	BB	522	A	P-O3'-C3'	-5.07	113.62	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	778	A	C4-N9-C1'	-5.07	117.18	126.30
35	BB	1003	G	N1-C2-N3	5.07	126.94	123.90
35	BB	1156	U	C5'-C4'-C3'	-5.07	107.89	116.00
35	BB	1452	U	C2-N3-C4	-5.07	123.96	127.00
38	BE	48	G	OP1-P-O3'	5.07	116.35	105.20
39	BF	56	C	N1-C2-N3	-5.07	115.65	119.20
85	AA	121	C	O3'-P-O5'	5.07	113.63	104.00
85	AA	166	C	C5'-C4'-C3'	5.07	124.11	116.00
85	AA	686	U	C1'-O4'-C4'	-5.07	105.85	109.90
85	AA	1372	C	C2-N1-C1'	-5.07	113.23	118.80
85	AA	1648	G	O3'-P-O5'	-5.07	94.37	104.00
85	AA	1674	G	P-O3'-C3'	-5.07	113.62	119.70
85	AA	1841	G	N3-C4-C5	-5.07	126.07	128.60
85	AA	1882	U	O5'-C5'-C4'	-5.07	102.07	111.70
2	A1	206	ARG	CG-CD-NE	-5.07	101.16	111.80
12	AD	15	ARG	NE-CZ-NH1	5.07	122.83	120.30
34	BA	281	C	OP2-P-O3'	5.07	116.35	105.20
34	BA	329	G	N3-C4-C5	-5.07	126.07	128.60
34	BA	768	G	C4'-C3'-C2'	-5.07	97.53	102.60
34	BA	1042	U	OP1-P-OP2	-5.07	112.00	119.60
34	BA	1307	U	OP2-P-O3'	5.07	116.34	105.20
34	BA	1693	U	C4'-C3'-C2'	5.07	107.67	102.60
35	BB	129	U	C5-C4-O4	-5.07	122.86	125.90
35	BB	130	G	N1-C2-N2	-5.07	111.64	116.20
35	BB	497	C	C4'-C3'-C2'	5.07	107.67	102.60
35	BB	1193	G	N9-C1'-C2'	-5.07	106.43	112.00
35	BB	1312	U	C6-N1-C2	5.07	124.04	121.00
39	BF	7	G	C3'-C2'-C1'	-5.07	97.45	101.50
39	BF	65	U	O4'-C1'-N1	-5.07	104.15	108.20
39	BF	70	A	O4'-C4'-C3'	-5.07	98.93	104.00
41	BH	64	U	O3'-P-O5'	-5.07	94.38	104.00
77	Br	209	VAL	CA-CB-CG2	-5.07	103.30	110.90
82	Bw	177	LYS	N-CA-CB	5.07	119.72	110.60
85	AA	127	U	O4'-C4'-C3'	-5.07	98.94	104.00
85	AA	369	A	C4-C5-C6	-5.07	114.47	117.00
85	AA	424	A	C5'-C4'-C3'	5.07	124.10	116.00
85	AA	534	A	P-O5'-C5'	-5.07	112.79	120.90
85	AA	862	U	C4'-C3'-O3'	-5.07	98.76	109.40
85	AA	926	C	O3'-P-O5'	5.07	113.62	104.00
85	AA	1067	G	N1-C6-O6	-5.07	116.86	119.90
85	AA	1552	U	P-O3'-C3'	-5.07	113.62	119.70
34	BA	513	U	C5'-C4'-O4'	5.06	115.18	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	762	A	C8-N9-C4	5.06	107.83	105.80
35	BB	558	U	P-O3'-C3'	-5.06	113.62	119.70
35	BB	805	G	C6-N1-C2	-5.06	122.06	125.10
35	BB	1205	A	C4'-C3'-C2'	-5.06	97.54	102.60
38	BE	10	G	C1'-O4'-C4'	-5.06	105.85	109.90
85	AA	492	C	N1-C2-O2	5.06	121.94	118.90
85	AA	788	G	N1-C6-O6	-5.06	116.86	119.90
85	AA	1810	C	O4'-C4'-C3'	-5.06	98.94	104.00
85	AA	1828	C	O4'-C1'-N1	5.06	112.25	108.20
85	AA	1887	G	C4-N9-C1'	-5.06	119.92	126.50
85	AA	2084	U	C5'-C4'-O4'	5.06	115.18	109.10
9	A8	26	SER	C-N-CA	5.06	134.35	121.70
34	BA	4	A	C3'-C2'-C1'	-5.06	97.45	101.50
34	BA	72	U	C5'-C4'-O4'	5.06	115.17	109.10
34	BA	194	G	C8-N9-C1'	5.06	133.58	127.00
34	BA	374	U	O5'-C5'-C4'	-5.06	102.08	111.70
34	BA	383	G	C5'-C4'-O4'	5.06	115.18	109.10
34	BA	403	A	C8-N9-C4	5.06	107.83	105.80
34	BA	696	A	C6-N1-C2	-5.06	115.56	118.60
34	BA	740	A	C5'-C4'-C3'	5.06	124.10	116.00
34	BA	1215	U	O4'-C1'-C2'	5.06	112.16	107.60
34	BA	1221	A	C5'-C4'-O4'	-5.06	103.03	109.10
34	BA	1270	G	O4'-C1'-C2'	5.06	112.16	107.60
35	BB	130	G	C8-N9-C1'	5.06	133.58	127.00
35	BB	404	A	C5'-C4'-O4'	-5.06	103.03	109.10
35	BB	418	G	C4-N9-C1'	-5.06	119.92	126.50
35	BB	503	G	O4'-C1'-N9	5.06	112.25	108.20
35	BB	557	C	N3-C4-N4	-5.06	114.46	118.00
35	BB	614	U	N1-C2-O2	5.06	126.34	122.80
35	BB	1019	C	C2-N3-C4	-5.06	117.37	119.90
35	BB	1235	A	O5'-P-OP1	-5.06	101.14	105.70
35	BB	1313	C	C4'-C3'-C2'	5.06	107.66	102.60
35	BB	1463	A	O4'-C1'-N9	5.06	112.25	108.20
35	BB	1543	C	P-O3'-C3'	-5.06	113.62	119.70
36	BC	38	U	C5-C6-N1	-5.06	120.17	122.70
38	BE	196	C	P-O3'-C3'	5.06	125.78	119.70
38	BE	198	A	N7-C8-N9	-5.06	111.27	113.80
40	BG	94	G	C6-N1-C2	5.06	128.14	125.10
77	Br	308	LYS	C-N-CA	5.06	134.35	121.70
85	AA	106	G	C1'-O4'-C4'	-5.06	105.85	109.90
85	AA	361	U	N3-C2-O2	-5.06	118.66	122.20
85	AA	740	A	N1-C2-N3	5.06	131.83	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1270	C	C4'-C3'-C2'	-5.06	97.54	102.60
85	AA	1295	G	C3'-C2'-C1'	-5.06	97.45	101.50
85	AA	1388	G	O4'-C1'-N9	5.06	112.25	108.20
85	AA	1463	A	C5-C6-N1	5.06	120.23	117.70
85	AA	1954	C	C5-C6-N1	5.06	123.53	121.00
86	AB	4	C	N1-C2-O2	-5.06	115.86	118.90
34	BA	13	U	O5'-C5'-C4'	-5.06	102.08	111.70
34	BA	202	A	C6-N1-C2	5.06	121.64	118.60
34	BA	339	G	C5-C6-N1	5.06	114.03	111.50
34	BA	341	U	P-O3'-C3'	-5.06	113.63	119.70
34	BA	348	U	C5-C6-N1	-5.06	120.17	122.70
34	BA	936	A	C8-N9-C4	5.06	107.82	105.80
34	BA	1622	U	O4'-C4'-C3'	-5.06	98.94	104.00
34	BA	1729	G	C5'-C4'-C3'	5.06	124.10	116.00
35	BB	1065	G	C4-N9-C1'	5.06	133.08	126.50
35	BB	1486	C	C6-N1-C1'	5.06	126.87	120.80
35	BB	1542	C	N3-C2-O2	-5.06	118.36	121.90
49	BP	168	ARG	NE-CZ-NH2	5.06	122.83	120.30
50	BQ	88	ARG	CG-CD-NE	-5.06	101.17	111.80
85	AA	396	U	C4'-C3'-C2'	5.06	107.66	102.60
85	AA	1302	A	C8-N9-C4	-5.06	103.78	105.80
85	AA	1469	G	C2'-C3'-O3'	5.06	121.80	113.70
85	AA	1766	G	P-O3'-C3'	-5.06	113.63	119.70
85	AA	1962	U	O4'-C1'-N1	5.06	112.25	108.20
1	A0	88	PHE	CA-CB-CG	-5.06	101.76	113.90
34	BA	283	U	C5-C4-O4	5.06	128.94	125.90
34	BA	711	C	O5'-C5'-C4'	5.06	121.31	111.70
34	BA	866	C	C6-N1-C2	-5.06	118.28	120.30
34	BA	892	C	O4'-C1'-C2'	5.06	112.15	107.60
34	BA	1562	G	N3-C2-N2	5.06	123.44	119.90
36	BC	146	U	C2-N1-C1'	-5.06	111.63	117.70
52	BS	114	SER	CB-CA-C	-5.06	100.49	110.10
85	AA	112	A	C4'-C3'-O3'	-5.06	98.77	109.40
85	AA	167	A	C4'-C3'-C2'	-5.06	97.54	102.60
85	AA	889	G	O3'-P-O5'	5.06	113.61	104.00
85	AA	906	U	P-O3'-C3'	5.06	125.77	119.70
85	AA	926	C	C1'-O4'-C4'	-5.06	105.85	109.90
85	AA	1228	A	C1'-O4'-C4'	-5.06	105.85	109.90
85	AA	1240	A	N9-C4-C5	-5.06	103.78	105.80
85	AA	2156	C	O4'-C1'-N1	5.06	112.25	108.20
85	AA	2210	C	N1-C2-N3	5.06	122.74	119.20
3	A2	54	ILE	N-CA-CB	-5.06	99.17	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
7	A6	15	ARG	N-CA-C	5.06	124.65	111.00
13	AE	51	SER	N-CA-C	-5.06	97.34	111.00
14	AF	114	SER	N-CA-CB	5.06	118.09	110.50
34	BA	39	C	C4'-C3'-C2'	5.06	107.66	102.60
34	BA	104	A	P-O5'-C5'	-5.06	112.81	120.90
34	BA	156	U	C2'-C3'-O3'	5.06	121.79	113.70
34	BA	196	A	N1-C6-N6	5.06	121.63	118.60
34	BA	591	G	C4-C5-C6	-5.06	115.77	118.80
34	BA	661	C	N3-C2-O2	-5.06	118.36	121.90
34	BA	810	A	C8-N9-C1'	5.06	136.80	127.70
34	BA	847	U	P-O3'-C3'	5.06	125.77	119.70
34	BA	1612	C	C2'-C3'-O3'	5.06	121.79	113.70
35	BB	7	C	C6-N1-C1'	5.06	126.87	120.80
35	BB	822	G	C4'-C3'-O3'	5.06	123.11	113.00
35	BB	998	G	N1-C2-N2	-5.06	111.65	116.20
35	BB	1434	G	N1-C6-O6	5.06	122.93	119.90
35	BB	1454	G	O4'-C4'-C3'	-5.06	98.94	104.00
37	BD	65	G	C1'-O4'-C4'	-5.06	105.86	109.90
41	BH	30	C	C3'-C2'-C1'	5.06	105.55	101.50
50	BQ	79	PHE	CB-CG-CD1	-5.06	117.26	120.80
54	BU	65	TRP	CB-CG-CD1	5.06	133.57	127.00
58	BY	70	ARG	NE-CZ-NH1	5.06	122.83	120.30
60	Ba	35	ARG	N-CA-C	-5.06	97.34	111.00
69	Bj	145	ALA	C-N-CA	5.06	134.34	121.70
71	Bl	62	ARG	N-CA-CB	5.06	119.70	110.60
85	AA	5	U	P-O5'-C5'	5.06	128.99	120.90
85	AA	79	G	C5-C6-O6	-5.06	125.57	128.60
85	AA	702	G	N3-C2-N2	5.06	123.44	119.90
85	AA	716	G	C1'-O4'-C4'	-5.06	105.86	109.90
85	AA	1272	G	C4'-C3'-C2'	-5.06	97.54	102.60
85	AA	1325	C	O4'-C1'-N1	5.06	112.25	108.20
34	BA	56	G	C5-C6-N1	5.06	114.03	111.50
34	BA	575	U	O5'-C5'-C4'	5.06	121.31	111.70
34	BA	578	C	C2'-C3'-O3'	5.06	121.79	113.70
34	BA	591	G	C8-N9-C4	5.06	108.42	106.40
34	BA	626	G	N9-C1'-C2'	-5.06	106.44	112.00
34	BA	757	G	O4'-C1'-N9	5.06	112.25	108.20
34	BA	1776	G	N9-C1'-C2'	-5.06	106.44	112.00
35	BB	1485	G	C3'-C2'-C1'	-5.06	97.45	101.50
54	BU	18	LYS	N-CA-C	5.06	124.65	111.00
65	Bf	382	LEU	CB-CA-C	-5.06	100.59	110.20
85	AA	35	U	O4'-C1'-N1	5.06	112.25	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	487	G	O3'-P-O5'	5.06	113.61	104.00
85	AA	1782	C	O4'-C1'-N1	5.06	112.25	108.20
85	AA	2043	A	N1-C6-N6	5.06	121.63	118.60
86	AB	36	A	O3'-P-O5'	-5.06	94.39	104.00
34	BA	292	C	C5'-C4'-C3'	-5.05	107.91	116.00
34	BA	400	A	C5'-C4'-O4'	5.05	115.17	109.10
34	BA	462	C	C5'-C4'-C3'	-5.05	107.91	116.00
34	BA	663	U	P-O5'-C5'	5.05	128.99	120.90
34	BA	690	G	O3'-P-O5'	5.05	113.60	104.00
34	BA	1026	C	C2-N3-C4	-5.05	117.37	119.90
34	BA	1074	C	OP1-P-OP2	-5.05	112.02	119.60
34	BA	1122	G	C1'-O4'-C4'	-5.05	105.86	109.90
34	BA	1305	A	C8-N9-C4	-5.05	103.78	105.80
35	BB	1356	G	N3-C4-N9	5.05	129.03	126.00
36	BC	15	G	C8-N9-C4	5.05	108.42	106.40
38	BE	204	U	P-O3'-C3'	-5.05	113.64	119.70
40	BG	73	U	C4'-C3'-C2'	5.05	107.65	102.60
41	BH	47	G	P-O5'-C5'	-5.05	112.81	120.90
77	Br	315	GLN	CB-CG-CD	-5.05	98.46	111.60
82	Bw	103	ARG	NE-CZ-NH1	5.05	122.83	120.30
85	AA	88	G	C4'-C3'-C2'	-5.05	97.55	102.60
85	AA	890	U	O3'-P-O5'	5.05	113.60	104.00
85	AA	1458	G	C1'-O4'-C4'	-5.05	105.86	109.90
85	AA	1848	G	N1-C6-O6	5.05	122.93	119.90
85	AA	2073	U	C5-C4-O4	5.05	128.93	125.90
34	BA	1428	G	C1'-O4'-C4'	-5.05	105.86	109.90
35	BB	38	C	C5'-C4'-O4'	5.05	115.16	109.10
35	BB	851	U	OP2-P-O3'	5.05	116.32	105.20
35	BB	1042	U	N1-C1'-C2'	-5.05	106.44	112.00
35	BB	1522	G	C4-N9-C1'	-5.05	119.93	126.50
85	AA	344	U	O5'-C5'-C4'	5.05	121.30	111.70
85	AA	627	A	C8-N9-C4	5.05	107.82	105.80
85	AA	788	G	O5'-C5'-C4'	-5.05	102.10	111.70
85	AA	884	A	C4'-C3'-C2'	-5.05	97.55	102.60
85	AA	1594	G	C3'-C2'-C1'	-5.05	97.46	101.50
85	AA	1621	U	P-O5'-C5'	5.05	128.99	120.90
85	AA	1865	C	N3-C2-O2	-5.05	118.36	121.90
85	AA	2154	C	C6-N1-C2	-5.05	118.28	120.30
1	A0	106	ASP	CB-CA-C	5.05	120.50	110.40
34	BA	74	A	C5'-C4'-O4'	5.05	115.16	109.10
34	BA	201	A	C5'-C4'-O4'	5.05	115.16	109.10
34	BA	470	C	O5'-P-OP1	5.05	116.76	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	498	A	C4-N9-C1'	-5.05	117.21	126.30
34	BA	997	U	N3-C4-O4	-5.05	115.86	119.40
34	BA	1119	A	C6-N1-C2	-5.05	115.57	118.60
34	BA	1133	A	C5-C6-N6	-5.05	119.66	123.70
34	BA	1626	U	O4'-C1'-N1	5.05	112.24	108.20
34	BA	1811	A	OP1-P-O3'	5.05	116.31	105.20
35	BB	683	U	N3-C2-O2	-5.05	118.66	122.20
35	BB	1040	C	C5'-C4'-C3'	5.05	124.08	116.00
35	BB	1051	U	C2-N1-C1'	-5.05	111.64	117.70
35	BB	1254	G	C8-N9-C4	5.05	108.42	106.40
35	BB	1372	G	C5-C6-N1	5.05	114.03	111.50
35	BB	1493	A	P-O3'-C3'	-5.05	113.64	119.70
38	BE	105	A	P-O5'-C5'	-5.05	112.82	120.90
40	BG	34	A	C1'-O4'-C4'	-5.05	105.86	109.90
41	BH	18	C	C5-C6-N1	-5.05	118.47	121.00
41	BH	97	C	O3'-P-O5'	-5.05	94.40	104.00
41	BH	98	U	C3'-C2'-C1'	-5.05	97.46	101.50
48	BO	143	GLN	CB-CA-C	5.05	120.50	110.40
48	BO	166	TRP	CA-CB-CG	-5.05	104.10	113.70
62	Bc	66	SER	N-CA-CB	5.05	118.08	110.50
75	Bp	72	ARG	NE-CZ-NH1	5.05	122.83	120.30
85	AA	367	A	C8-N9-C4	-5.05	103.78	105.80
85	AA	740	A	N7-C8-N9	5.05	116.33	113.80
85	AA	819	G	C6-C5-N7	-5.05	127.37	130.40
85	AA	1199	C	N3-C2-O2	-5.05	118.36	121.90
85	AA	1228	A	C5-N7-C8	-5.05	101.37	103.90
85	AA	1256	C	O4'-C1'-N1	5.05	112.24	108.20
85	AA	1600	G	O4'-C1'-N9	5.05	112.24	108.20
85	AA	1673	A	C4-N9-C1'	-5.05	117.21	126.30
85	AA	1906	C	C3'-C2'-C1'	-5.05	97.46	101.50
2	A1	54	ASN	CA-CB-CG	5.05	124.51	113.40
34	BA	149	G	C5'-C4'-C3'	-5.05	107.92	116.00
34	BA	809	U	C5-C6-N1	-5.05	120.17	122.70
34	BA	1123	G	C8-N9-C4	-5.05	104.38	106.40
34	BA	1488	C	P-O3'-C3'	-5.05	113.64	119.70
35	BB	387	G	C3'-C2'-C1'	-5.05	97.46	101.50
35	BB	830	G	C4-N9-C1'	-5.05	119.94	126.50
35	BB	1543	C	N3-C2-O2	-5.05	118.37	121.90
37	BD	3	G	C1'-O4'-C4'	-5.05	105.86	109.90
38	BE	89	G	N1-C2-N2	-5.05	111.66	116.20
79	Bt	80	ILE	CA-CB-CG1	-5.05	101.41	111.00
85	AA	47	A	C6-C5-N7	-5.05	128.77	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	67	C	N3-C4-C5	-5.05	119.88	121.90
85	AA	70	U	C1'-O4'-C4'	-5.05	105.86	109.90
85	AA	141	A	N1-C2-N3	-5.05	126.78	129.30
85	AA	455	G	C5'-C4'-O4'	5.05	115.16	109.10
85	AA	511	A	C5'-C4'-C3'	-5.05	107.92	116.00
85	AA	864	C	C5'-C4'-C3'	-5.05	107.92	116.00
85	AA	880	A	N1-C6-N6	-5.05	115.57	118.60
85	AA	1091	C	N3-C4-N4	5.05	121.53	118.00
85	AA	1355	U	C1'-O4'-C4'	5.05	113.94	109.90
85	AA	1884	A	C1'-O4'-C4'	-5.05	105.86	109.90
85	AA	2174	G	C4-N9-C1'	-5.05	119.94	126.50
4	A3	163	ARG	CG-CD-NE	-5.05	101.20	111.80
34	BA	429	G	C2'-C3'-O3'	5.05	121.78	113.70
34	BA	1233	U	C6-N1-C1'	5.05	128.27	121.20
35	BB	364	U	P-O3'-C3'	-5.05	113.64	119.70
35	BB	538	A	O5'-C5'-C4'	-5.05	102.11	111.70
35	BB	829	C	N1-C2-O2	5.05	121.93	118.90
35	BB	838	G	N3-C2-N2	5.05	123.43	119.90
35	BB	889	U	C5'-C4'-C3'	-5.05	107.92	116.00
35	BB	916	U	C2-N1-C1'	5.05	123.76	117.70
35	BB	1414	A	C8-N9-C4	5.05	107.82	105.80
38	BE	18	U	N3-C2-O2	-5.05	118.67	122.20
38	BE	169	C	C2'-C3'-O3'	5.05	121.78	113.70
39	BF	45	G	N9-C1'-C2'	-5.05	106.45	112.00
63	Bd	58	ALA	CA-C-O	5.05	130.70	120.10
85	AA	30	G	O4'-C1'-N9	5.05	112.24	108.20
85	AA	525	C	C3'-C2'-C1'	-5.05	97.46	101.50
85	AA	1016	G	O4'-C1'-N9	5.05	112.24	108.20
85	AA	2208	G	C1'-O4'-C4'	-5.05	105.86	109.90
34	BA	709	C	C5'-C4'-O4'	5.05	115.16	109.10
34	BA	1113	A	C6-N1-C2	-5.05	115.57	118.60
34	BA	1602	A	N1-C6-N6	5.05	121.63	118.60
34	BA	1718	C	C5'-C4'-C3'	5.05	124.08	116.00
34	BA	1735	G	N1-C2-N3	-5.05	120.87	123.90
35	BB	47	C	P-O3'-C3'	5.05	125.76	119.70
35	BB	579	A	O4'-C1'-N9	5.05	112.24	108.20
35	BB	822	G	C4'-C3'-C2'	5.05	107.65	102.60
35	BB	1048	A	N7-C8-N9	-5.05	111.28	113.80
35	BB	1157	G	C5-C6-N1	5.05	114.02	111.50
35	BB	1209	A	P-O5'-C5'	5.05	128.98	120.90
35	BB	1532	C	C6-N1-C1'	5.05	126.86	120.80
36	BC	28	C	C2-N1-C1'	5.05	124.35	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	BC	58	G	C8-N9-C1'	5.05	133.56	127.00
36	BC	113	G	C3'-C2'-C1'	-5.05	97.46	101.50
36	BC	127	C	N1-C2-O2	5.05	121.93	118.90
38	BE	162	U	C6-N1-C1'	5.05	128.26	121.20
65	Bf	265	PHE	CA-CB-CG	-5.05	101.79	113.90
81	Bv	179	PHE	N-CA-C	-5.05	97.37	111.00
85	AA	63	G	C5'-C4'-O4'	5.05	115.16	109.10
85	AA	226	C	O4'-C1'-N1	5.05	112.24	108.20
85	AA	261	U	O4'-C1'-N1	5.05	112.24	108.20
85	AA	455	G	N3-C4-N9	-5.05	122.97	126.00
85	AA	844	C	C2-N1-C1'	5.05	124.35	118.80
85	AA	937	G	C3'-C2'-C1'	-5.05	97.46	101.50
85	AA	1612	C	N3-C4-N4	5.05	121.53	118.00
85	AA	1666	U	O5'-C5'-C4'	-5.05	102.11	111.70
85	AA	1876	U	C6-N1-C2	-5.05	117.97	121.00
85	AA	2066	C	C5'-C4'-C3'	5.05	124.07	116.00
85	AA	2135	A	P-O3'-C3'	5.05	125.75	119.70
31	AX	66	ARG	N-CA-CB	-5.04	101.52	110.60
34	BA	649	A	C5-C6-N6	-5.04	119.66	123.70
34	BA	1315	C	N1-C1'-C2'	-5.04	106.45	112.00
35	BB	458	U	C3'-C2'-C1'	-5.04	97.46	101.50
35	BB	533	U	C2-N3-C4	-5.04	123.97	127.00
35	BB	901	U	P-O3'-C3'	-5.04	113.65	119.70
35	BB	1364	C	O4'-C1'-N1	5.04	112.24	108.20
80	Bu	118	LYS	CA-CB-CG	5.04	124.50	113.40
85	AA	25	C	C2'-C3'-O3'	5.04	121.77	113.70
85	AA	34	G	O4'-C1'-N9	5.04	112.24	108.20
85	AA	243	A	P-O5'-C5'	-5.04	112.83	120.90
85	AA	694	A	C4'-C3'-C2'	5.04	107.64	102.60
85	AA	869	A	O4'-C1'-N9	-5.04	104.16	108.20
85	AA	1471	G	N1-C6-O6	5.04	122.93	119.90
85	AA	1820	G	N1-C6-O6	5.04	122.93	119.90
85	AA	1981	A	C4'-C3'-C2'	-5.04	97.56	102.60
21	AM	27	VAL	CA-CB-CG1	5.04	118.47	110.90
34	BA	191	G	C5-C6-N1	5.04	114.02	111.50
34	BA	630	U	P-O5'-C5'	5.04	128.97	120.90
34	BA	1050	A	C5'-C4'-C3'	5.04	124.07	116.00
34	BA	1130	U	C4'-C3'-C2'	-5.04	97.56	102.60
34	BA	1433	U	OP1-P-O3'	5.04	116.29	105.20
34	BA	1498	A	O3'-P-O5'	-5.04	94.42	104.00
34	BA	1500	G	N1-C6-O6	-5.04	116.87	119.90
35	BB	25	A	C4-C5-C6	-5.04	114.48	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	123	U	C2-N3-C4	-5.04	123.97	127.00
35	BB	314	A	P-O5'-C5'	-5.04	112.83	120.90
35	BB	409	U	O5'-C5'-C4'	-5.04	102.12	111.70
35	BB	1446	C	C2-N3-C4	-5.04	117.38	119.90
36	BC	9	G	N9-C1'-C2'	-5.04	106.45	112.00
37	BD	98	G	P-O5'-C5'	-5.04	112.83	120.90
38	BE	113	C	C2'-C3'-O3'	5.04	121.77	113.70
40	BG	132	U	C4'-C3'-C2'	-5.04	97.56	102.60
40	BG	139	U	C6-N1-C2	-5.04	117.97	121.00
45	BL	67	ARG	NE-CZ-NH2	-5.04	117.78	120.30
51	BR	56	ARG	CD-NE-CZ	5.04	130.66	123.60
73	Bn	34	CYS	N-CA-C	-5.04	97.39	111.00
85	AA	87	C	C2'-C3'-O3'	5.04	121.77	113.70
85	AA	247	G	C8-N9-C1'	5.04	133.56	127.00
85	AA	308	U	O5'-C5'-C4'	5.04	121.28	111.70
85	AA	636	G	N3-C2-N2	5.04	123.43	119.90
85	AA	773	G	N3-C2-N2	-5.04	116.37	119.90
85	AA	1341	U	O4'-C1'-N1	5.04	112.23	108.20
85	AA	1376	U	C6-N1-C2	-5.04	117.97	121.00
7	A6	17	PRO	N-CA-C	5.04	125.21	112.10
17	AI	121	HIS	CA-CB-CG	-5.04	105.03	113.60
34	BA	338	U	O4'-C1'-N1	5.04	112.23	108.20
34	BA	527	C	N1-C1'-C2'	-5.04	106.45	112.00
34	BA	697	A	O3'-P-O5'	5.04	113.58	104.00
34	BA	1294	C	C5'-C4'-C3'	-5.04	107.94	116.00
34	BA	1428	G	N9-C1'-C2'	-5.04	106.45	112.00
34	BA	1591	G	C2'-C3'-O3'	5.04	121.77	113.70
34	BA	1615	A	C3'-C2'-C1'	-5.04	97.47	101.50
34	BA	1617	U	N3-C2-O2	-5.04	118.67	122.20
35	BB	38	C	C5-C4-N4	5.04	123.73	120.20
35	BB	393	A	C4-N9-C1'	-5.04	117.23	126.30
35	BB	565	U	C6-N1-C1'	-5.04	114.14	121.20
35	BB	853	U	P-O5'-C5'	-5.04	112.83	120.90
37	BD	91	U	C4'-C3'-C2'	5.04	107.64	102.60
38	BE	196	C	C5'-C4'-C3'	5.04	124.07	116.00
45	BL	62	ARG	N-CA-CB	5.04	119.67	110.60
59	BZ	51	ASP	N-CA-C	5.04	124.61	111.00
80	Bu	133	ARG	CG-CD-NE	-5.04	101.22	111.80
85	AA	267	U	N3-C4-O4	5.04	122.93	119.40
85	AA	694	A	O5'-P-OP2	5.04	116.75	110.70
85	AA	1043	U	P-O3'-C3'	-5.04	113.65	119.70
85	AA	1140	G	C6-N1-C2	5.04	128.12	125.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1198	U	N1-C2-O2	5.04	126.33	122.80
85	AA	1517	G	C5-C6-N1	5.04	114.02	111.50
85	AA	1650	G	C4-N9-C1'	-5.04	119.95	126.50
85	AA	1913	G	O5'-C5'-C4'	-5.04	102.12	111.70
85	AA	2009	A	C8-N9-C4	-5.04	103.78	105.80
85	AA	2076	C	C5'-C4'-O4'	-5.04	103.05	109.10
85	AA	2140	U	C6-N1-C1'	5.04	128.26	121.20
34	BA	589	A	C8-N9-C1'	-5.04	118.63	127.70
34	BA	1596	C	P-O3'-C3'	5.04	125.75	119.70
35	BB	1060	U	P-O3'-C3'	5.04	125.75	119.70
37	BD	47	U	O3'-P-O5'	5.04	113.58	104.00
40	BG	70	C	P-O3'-C3'	-5.04	113.65	119.70
41	BH	113	G	P-O5'-C5'	-5.04	112.84	120.90
73	Bn	80	THR	CA-CB-CG2	-5.04	105.34	112.40
77	Br	256	GLU	N-CA-CB	-5.04	101.53	110.60
85	AA	526	G	C1'-O4'-C4'	-5.04	105.87	109.90
24	AQ	89	ARG	N-CA-C	-5.04	97.39	111.00
34	BA	95	C	C5-C4-N4	5.04	123.73	120.20
34	BA	816	G	O4'-C4'-C3'	-5.04	98.96	104.00
34	BA	954	U	C6-N1-C2	-5.04	117.98	121.00
34	BA	1287	G	C5-C6-O6	-5.04	125.58	128.60
34	BA	1840	C	C5'-C4'-O4'	5.04	115.15	109.10
35	BB	25	A	P-O5'-C5'	-5.04	112.84	120.90
35	BB	379	U	C4-C5-C6	-5.04	116.68	119.70
35	BB	640	A	O5'-P-OP1	5.04	116.75	110.70
35	BB	1054	G	N1-C6-O6	5.04	122.92	119.90
35	BB	1517	G	N3-C4-N9	-5.04	122.98	126.00
36	BC	132	U	P-O3'-C3'	-5.04	113.66	119.70
40	BG	14	G	P-O3'-C3'	5.04	125.75	119.70
41	BH	22	A	C2'-C3'-O3'	5.04	121.76	113.70
41	BH	108	U	O3'-P-O5'	-5.04	94.43	104.00
65	Bf	364	THR	CA-CB-CG2	-5.04	105.35	112.40
85	AA	37	U	P-O5'-C5'	-5.04	112.84	120.90
85	AA	147	G	N9-C1'-C2'	-5.04	106.46	112.00
85	AA	420	C	C3'-C2'-C1'	-5.04	97.47	101.50
85	AA	830	A	O3'-P-O5'	5.04	113.57	104.00
85	AA	1028	C	C1'-O4'-C4'	-5.04	105.87	109.90
85	AA	1567	C	P-O3'-C3'	-5.04	113.65	119.70
85	AA	1758	C	O4'-C1'-N1	5.04	112.23	108.20
85	AA	2184	A	O5'-C5'-C4'	-5.04	102.13	111.70
85	AA	2241	C	C6-N1-C2	-5.04	118.28	120.30
34	BA	633	G	O4'-C1'-N9	5.04	112.23	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	803	U	N3-C2-O2	-5.04	118.67	122.20
34	BA	1038	U	C3'-C2'-C1'	-5.04	97.47	101.50
34	BA	1131	G	O4'-C1'-N9	5.04	112.23	108.20
34	BA	1165	A	C3'-C2'-C1'	-5.04	97.47	101.50
34	BA	1436	A	C8-N9-C1'	5.04	136.77	127.70
34	BA	1507	C	P-O3'-C3'	5.04	125.74	119.70
34	BA	1746	G	C4'-C3'-C2'	-5.04	97.56	102.60
35	BB	77	A	N1-C6-N6	-5.04	115.58	118.60
35	BB	387	G	O4'-C1'-N9	5.04	112.23	108.20
35	BB	1091	C	N3-C2-O2	-5.04	118.37	121.90
35	BB	1542	C	O4'-C4'-C3'	-5.04	98.96	104.00
41	BH	16	A	N3-C4-N9	-5.04	123.37	127.40
41	BH	17	A	O5'-C5'-C4'	-5.04	102.13	111.70
85	AA	290	G	O4'-C1'-N9	5.04	112.23	108.20
85	AA	1371	C	C1'-O4'-C4'	-5.04	105.87	109.90
85	AA	2227	A	O4'-C1'-C2'	5.04	112.13	107.60
34	BA	216	C	C3'-C2'-C1'	-5.04	97.47	101.50
34	BA	406	G	N1-C6-O6	5.04	122.92	119.90
34	BA	422	C	O5'-C5'-C4'	-5.04	102.13	111.70
34	BA	490	A	O3'-P-O5'	5.04	113.57	104.00
34	BA	493	G	C5-C6-N1	5.04	114.02	111.50
34	BA	618	G	N3-C4-C5	-5.04	126.08	128.60
34	BA	622	G	C4-N9-C1'	-5.04	119.95	126.50
34	BA	691	A	O5'-C5'-C4'	5.04	121.27	111.70
34	BA	1034	U	N1-C2-N3	5.04	117.92	114.90
34	BA	1167	A	C2-N3-C4	-5.04	108.08	110.60
34	BA	1531	G	N1-C2-N2	-5.04	111.67	116.20
35	BB	268	G	C4'-C3'-C2'	-5.04	97.56	102.60
35	BB	504	C	N3-C2-O2	5.04	125.42	121.90
35	BB	576	A	P-O3'-C3'	-5.04	113.66	119.70
35	BB	613	C	N3-C2-O2	-5.04	118.38	121.90
35	BB	806	U	N1-C2-N3	5.04	117.92	114.90
35	BB	956	G	C5-C6-O6	-5.04	125.58	128.60
35	BB	1047	C	C4'-C3'-C2'	5.04	107.64	102.60
35	BB	1195	A	C3'-C2'-C1'	-5.04	97.47	101.50
35	BB	1202	G	O5'-C5'-C4'	5.04	121.27	111.70
35	BB	1410	G	C8-N9-C1'	-5.04	120.45	127.00
35	BB	1512	C	C5-C4-N4	-5.04	116.67	120.20
37	BD	54	A	C5-C6-N6	-5.04	119.67	123.70
38	BE	131	C	O5'-C5'-C4'	-5.04	102.13	111.70
39	BF	37	C	C1'-O4'-C4'	-5.04	105.87	109.90
40	BG	179	C	C2-N3-C4	-5.04	117.38	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	BH	103	C	N1-C1'-C2'	-5.04	106.46	112.00
47	BN	158	TYR	CB-CG-CD1	-5.04	117.98	121.00
57	BX	72	VAL	N-CA-C	-5.04	97.40	111.00
65	Bf	454	LYS	CB-CG-CD	5.04	124.69	111.60
69	Bj	22	ARG	NE-CZ-NH1	5.04	122.82	120.30
85	AA	33	U	O5'-C5'-C4'	-5.04	102.13	111.70
85	AA	315	U	O4'-C4'-C3'	-5.04	98.97	104.00
85	AA	436	G	C8-N9-C1'	5.04	133.55	127.00
85	AA	617	C	C2-N1-C1'	5.04	124.34	118.80
85	AA	671	G	O4'-C1'-N9	5.04	112.23	108.20
85	AA	797	C	P-O5'-C5'	5.04	128.96	120.90
85	AA	986	U	P-O5'-C5'	5.04	128.96	120.90
85	AA	999	A	N1-C6-N6	-5.04	115.58	118.60
85	AA	1063	U	C5'-C4'-C3'	-5.04	107.94	116.00
85	AA	1155	A	C6-N1-C2	-5.04	115.58	118.60
85	AA	1538	C	P-O3'-C3'	-5.04	113.66	119.70
85	AA	2007	G	C4-C5-C6	-5.04	115.78	118.80
85	AA	2131	C	C5'-C4'-O4'	5.04	115.14	109.10
34	BA	27	G	C1'-O4'-C4'	-5.03	105.87	109.90
34	BA	345	G	N1-C6-O6	-5.03	116.88	119.90
34	BA	494	A	N9-C1'-C2'	-5.03	106.46	112.00
34	BA	754	G	OP1-P-O3'	5.03	116.27	105.20
34	BA	1251	A	C4-N9-C1'	-5.03	117.24	126.30
34	BA	1452	U	C5'-C4'-C3'	-5.03	107.95	116.00
35	BB	45	A	P-O5'-C5'	-5.03	112.84	120.90
35	BB	1038	G	C8-N9-C4	5.03	108.41	106.40
35	BB	1536	G	C4-N9-C1'	5.03	133.04	126.50
36	BC	113	G	O4'-C1'-N9	5.03	112.23	108.20
38	BE	107	U	N1-C2-N3	-5.03	111.88	114.90
40	BG	76	C	C2-N1-C1'	5.03	124.34	118.80
61	Bb	76	ASP	C-N-CA	5.03	134.29	121.70
82	Bw	26	SER	C-N-CA	5.03	134.28	121.70
85	AA	191	C	C5-C4-N4	-5.03	116.68	120.20
85	AA	244	G	O5'-C5'-C4'	-5.03	102.14	111.70
85	AA	336	C	C1'-O4'-C4'	-5.03	105.87	109.90
85	AA	427	G	N9-C1'-C2'	-5.03	106.46	112.00
85	AA	450	A	P-O3'-C3'	-5.03	113.66	119.70
85	AA	641	A	O4'-C1'-N9	5.03	112.23	108.20
85	AA	736	U	C5-C6-N1	5.03	125.22	122.70
85	AA	1429	U	C2-N1-C1'	-5.03	111.66	117.70
85	AA	1586	C	C5-C6-N1	5.03	123.52	121.00
85	AA	1887	G	N3-C2-N2	5.03	123.42	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	2030	U	O4'-C1'-N1	5.03	112.23	108.20
85	AA	2032	G	C5-C6-O6	-5.03	125.58	128.60
85	AA	2222	G	C5'-C4'-C3'	-5.03	107.94	116.00
16	AH	117	MET	CG-SD-CE	-5.03	92.15	100.20
34	BA	376	U	N1-C1'-C2'	-5.03	106.47	112.00
34	BA	554	A	N1-C2-N3	5.03	131.82	129.30
34	BA	1223	C	N1-C2-N3	-5.03	115.68	119.20
34	BA	1423	U	OP2-P-O3'	5.03	116.27	105.20
34	BA	1593	U	P-O5'-C5'	5.03	128.95	120.90
35	BB	306	U	C2-N3-C4	5.03	130.02	127.00
35	BB	1094	A	C8-N9-C4	5.03	107.81	105.80
85	AA	892	C	C6-N1-C2	-5.03	118.29	120.30
85	AA	1964	A	O4'-C4'-C3'	-5.03	98.97	104.00
2	A1	129	GLY	N-CA-C	-5.03	100.52	113.10
34	BA	34	U	C6-N1-C2	-5.03	117.98	121.00
34	BA	91	C	C4'-C3'-C2'	-5.03	97.57	102.60
34	BA	202	A	N1-C6-N6	5.03	121.62	118.60
34	BA	412	G	P-O3'-C3'	5.03	125.74	119.70
34	BA	754	G	P-O3'-C3'	-5.03	113.66	119.70
34	BA	807	U	O5'-C5'-C4'	5.03	121.26	111.70
34	BA	951	C	N1-C2-N3	5.03	122.72	119.20
34	BA	1078	U	C2'-C3'-O3'	5.03	121.75	113.70
34	BA	1306	U	C6-N1-C2	-5.03	117.98	121.00
35	BB	56	U	P-O3'-C3'	-5.03	113.66	119.70
35	BB	613	C	O5'-C5'-C4'	-5.03	102.14	111.70
35	BB	960	C	P-O5'-C5'	-5.03	112.85	120.90
35	BB	1063	C	O5'-C5'-C4'	-5.03	102.14	111.70
35	BB	1197	G	N1-C2-N3	-5.03	120.88	123.90
35	BB	1352	C	C5-C6-N1	5.03	123.52	121.00
35	BB	1376	G	C5-C6-N1	5.03	114.02	111.50
37	BD	14	C	C5'-C4'-C3'	5.03	124.05	116.00
38	BE	2	G	C8-N9-C4	-5.03	104.39	106.40
38	BE	171	U	C2-N3-C4	-5.03	123.98	127.00
62	Bc	36	ALA	C-N-CA	5.03	134.28	121.70
64	Be	230	PRO	N-CA-CB	5.03	109.34	103.30
65	Bf	204	PHE	CB-CG-CD1	5.03	124.32	120.80
70	Bk	71	ARG	NE-CZ-NH1	5.03	122.82	120.30
74	Bo	33	GLN	N-CA-CB	-5.03	101.55	110.60
85	AA	445	U	O4'-C4'-C3'	-5.03	98.97	104.00
85	AA	506	G	C5'-C4'-C3'	-5.03	107.95	116.00
85	AA	687	G	C8-N9-C4	-5.03	104.39	106.40
85	AA	783	C	C3'-C2'-C1'	-5.03	97.47	101.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	960	G	C3'-C2'-C1'	-5.03	97.48	101.50
85	AA	1673	A	C8-N9-C4	5.03	107.81	105.80
85	AA	1733	G	N1-C6-O6	-5.03	116.88	119.90
85	AA	1901	G	C5-C6-O6	-5.03	125.58	128.60
85	AA	2004	U	C2'-C3'-O3'	5.03	121.75	113.70
85	AA	2099	C	P-O3'-C3'	5.03	125.74	119.70
34	BA	289	A	N7-C8-N9	-5.03	111.28	113.80
34	BA	549	G	C5-C6-O6	5.03	131.62	128.60
34	BA	550	U	OP2-P-O3'	5.03	116.26	105.20
34	BA	1602	A	C8-N9-C4	5.03	107.81	105.80
35	BB	806	U	C5'-C4'-C3'	5.03	124.05	116.00
35	BB	1021	C	C2-N3-C4	-5.03	117.39	119.90
35	BB	1372	G	C5'-C4'-O4'	5.03	115.14	109.10
38	BE	39	U	N1-C1'-C2'	-5.03	106.47	112.00
38	BE	172	U	C5'-C4'-O4'	5.03	115.14	109.10
38	BE	202	C	N3-C4-N4	-5.03	114.48	118.00
39	BF	15	U	C1'-O4'-C4'	-5.03	105.88	109.90
39	BF	34	C	OP1-P-OP2	-5.03	112.06	119.60
79	Bt	100	GLY	N-CA-C	5.03	125.67	113.10
84	By	104	GLU	C-N-CA	5.03	134.27	121.70
85	AA	773	G	C2-N3-C4	5.03	114.42	111.90
85	AA	1961	U	N3-C2-O2	-5.03	118.68	122.20
34	BA	20	A	C6-C5-N7	-5.03	128.78	132.30
34	BA	480	G	O3'-P-O5'	-5.03	94.45	104.00
34	BA	543	A	N1-C2-N3	5.03	131.81	129.30
34	BA	646	C	O5'-P-OP1	5.03	116.73	110.70
34	BA	777	C	N3-C4-N4	5.03	121.52	118.00
35	BB	45	A	P-O3'-C3'	5.03	125.73	119.70
35	BB	66	G	C4-C5-N7	5.03	112.81	110.80
35	BB	379	U	P-O3'-C3'	-5.03	113.67	119.70
35	BB	461	U	N3-C2-O2	-5.03	118.68	122.20
35	BB	755	A	C2-N3-C4	5.03	113.11	110.60
35	BB	1221	G	N3-C4-N9	-5.03	122.98	126.00
65	Bf	72	ARG	N-CA-C	-5.03	97.43	111.00
70	Bk	46	GLY	C-N-CA	5.03	134.27	121.70
70	Bk	92	THR	N-CA-CB	5.03	119.85	110.30
85	AA	874	A	N9-C1'-C2'	-5.03	106.47	112.00
85	AA	1191	G	N1-C2-N3	-5.03	120.88	123.90
85	AA	1735	U	O4'-C4'-C3'	-5.03	98.97	104.00
85	AA	1812	C	C6-N1-C1'	5.03	126.83	120.80
85	AA	1818	C	P-O3'-C3'	5.03	125.73	119.70
86	AB	68	C	C4'-C3'-C2'	5.03	107.63	102.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	175	G	C5'-C4'-C3'	5.03	124.04	116.00
34	BA	201	A	C1'-O4'-C4'	-5.03	105.88	109.90
34	BA	308	C	C6-N1-C1'	5.03	126.83	120.80
34	BA	543	A	C5'-C4'-O4'	5.03	115.13	109.10
34	BA	1203	G	N3-C2-N2	5.03	123.42	119.90
34	BA	1446	G	C5'-C4'-C3'	-5.03	107.96	116.00
35	BB	938	G	C5-C6-O6	-5.03	125.58	128.60
35	BB	1147	G	C1'-O4'-C4'	-5.03	105.88	109.90
35	BB	1386	C	P-O3'-C3'	-5.03	113.67	119.70
36	BC	42	G	C8-N9-C4	5.03	108.41	106.40
37	BD	29	C	C5'-C4'-O4'	5.03	115.13	109.10
38	BE	32	U	C5-C6-N1	-5.03	120.19	122.70
40	BG	61	A	N9-C1'-C2'	-5.03	106.47	112.00
40	BG	88	G	N9-C1'-C2'	-5.03	106.47	112.00
65	Bf	339	TYR	N-CA-CB	-5.03	101.55	110.60
70	Bk	120	ARG	NE-CZ-NH2	-5.03	117.79	120.30
85	AA	58	C	C4-C5-C6	-5.03	114.89	117.40
85	AA	114	C	N3-C2-O2	-5.03	118.38	121.90
85	AA	180	A	C5'-C4'-C3'	-5.03	107.96	116.00
85	AA	247	G	N1-C6-O6	5.03	122.92	119.90
85	AA	933	U	C5-C4-O4	-5.03	122.89	125.90
85	AA	2204	A	C4'-C3'-C2'	5.03	107.63	102.60
25	AR	45	ASP	N-CA-C	-5.02	97.44	111.00
34	BA	61	G	N9-C4-C5	5.02	107.41	105.40
34	BA	550	U	O5'-P-OP1	-5.02	101.18	105.70
34	BA	1064	A	P-O3'-C3'	-5.02	113.67	119.70
34	BA	1072	U	O4'-C1'-N1	5.02	112.22	108.20
36	BC	119	G	C8-N9-C4	5.02	108.41	106.40
39	BF	52	A	O4'-C4'-C3'	-5.02	98.98	104.00
41	BH	125	U	C1'-O4'-C4'	-5.02	105.88	109.90
41	BH	135	U	N3-C4-O4	-5.02	115.88	119.40
59	BZ	91	THR	N-CA-CB	5.02	119.84	110.30
77	Br	107	PHE	N-CA-CB	5.02	119.64	110.60
85	AA	372	U	C5-C4-O4	5.02	128.91	125.90
85	AA	2213	A	P-O5'-C5'	-5.02	112.86	120.90
34	BA	107	C	C5'-C4'-O4'	5.02	115.13	109.10
34	BA	274	C	C2-N3-C4	5.02	122.41	119.90
34	BA	375	C	C5-C6-N1	5.02	123.51	121.00
34	BA	793	A	C5-C6-N6	-5.02	119.68	123.70
34	BA	878	G	N3-C2-N2	5.02	123.42	119.90
34	BA	963	G	C6-N1-C2	-5.02	122.09	125.10
34	BA	1297	G	O4'-C1'-N9	5.02	112.22	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	1460	U	C2-N1-C1'	5.02	123.73	117.70
34	BA	1818	A	N9-C1'-C2'	5.02	120.53	114.00
35	BB	68	G	O4'-C4'-C3'	5.02	110.12	106.10
35	BB	423	G	C5-C6-O6	5.02	131.61	128.60
35	BB	603	U	OP1-P-OP2	-5.02	112.07	119.60
35	BB	856	U	O4'-C1'-N1	5.02	112.22	108.20
35	BB	923	U	O4'-C1'-N1	5.02	112.22	108.20
35	BB	1232	A	C8-N9-C4	-5.02	103.79	105.80
35	BB	1252	G	O4'-C1'-N9	-5.02	104.18	108.20
37	BD	41	G	O4'-C1'-C2'	5.02	112.12	107.60
41	BH	46	C	C3'-C2'-C1'	-5.02	97.48	101.50
68	Bi	128	ARG	CA-C-N	5.02	128.25	117.20
85	AA	363	A	O5'-C5'-C4'	-5.02	102.16	111.70
85	AA	725	G	N3-C2-N2	5.02	123.42	119.90
85	AA	902	A	O5'-C5'-C4'	-5.02	102.16	111.70
85	AA	1093	C	C2-N1-C1'	-5.02	113.28	118.80
85	AA	1226	A	C2-N3-C4	5.02	113.11	110.60
85	AA	1253	G	C5-C6-O6	-5.02	125.59	128.60
85	AA	1270	C	C1'-O4'-C4'	-5.02	105.88	109.90
85	AA	1510	A	N1-C6-N6	-5.02	115.59	118.60
85	AA	1574	C	OP1-P-O3'	5.02	116.25	105.20
85	AA	1668	G	N3-C2-N2	-5.02	116.39	119.90
85	AA	1975	G	P-O5'-C5'	5.02	128.94	120.90
85	AA	1992	A	C4'-C3'-C2'	-5.02	97.58	102.60
23	AP	136	ARG	CG-CD-NE	-5.02	101.25	111.80
34	BA	533	U	C1'-O4'-C4'	-5.02	105.88	109.90
34	BA	1574	C	N1-C2-O2	5.02	121.91	118.90
35	BB	83	G	C8-N9-C1'	5.02	133.53	127.00
35	BB	634	A	C1'-O4'-C4'	-5.02	105.88	109.90
36	BC	122	A	N9-C1'-C2'	-5.02	106.48	112.00
85	AA	807	A	C8-N9-C4	-5.02	103.79	105.80
86	AB	67	C	C5-C4-N4	5.02	123.72	120.20
34	BA	200	C	C3'-C2'-C1'	-5.02	97.48	101.50
34	BA	437	G	N3-C4-N9	5.02	129.01	126.00
34	BA	469	C	C2'-C3'-O3'	5.02	121.73	113.70
34	BA	636	G	O4'-C1'-N9	5.02	112.22	108.20
34	BA	1194	G	C8-N9-C4	5.02	108.41	106.40
34	BA	1468	U	C1'-O4'-C4'	-5.02	105.89	109.90
34	BA	1611	A	C4-N9-C1'	-5.02	117.26	126.30
35	BB	475	A	OP1-P-OP2	-5.02	112.07	119.60
35	BB	708	C	C5'-C4'-C3'	-5.02	107.97	116.00
35	BB	811	C	N3-C2-O2	-5.02	118.39	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	866	A	N9-C1'-C2'	-5.02	106.48	112.00
35	BB	1242	C	C5-C4-N4	-5.02	116.69	120.20
35	BB	1406	C	C2-N3-C4	-5.02	117.39	119.90
35	BB	1421	C	O5'-C5'-C4'	-5.02	102.17	111.70
40	BG	126	G	C6-N1-C2	-5.02	122.09	125.10
40	BG	168	A	N1-C2-N3	-5.02	126.79	129.30
41	BH	18	C	P-O5'-C5'	-5.02	112.87	120.90
41	BH	30	C	C2-N3-C4	-5.02	117.39	119.90
52	BS	110	ASN	CA-CB-CG	-5.02	102.36	113.40
53	BT	117	ARG	NE-CZ-NH2	-5.02	117.79	120.30
85	AA	270	A	C8-N9-C4	-5.02	103.79	105.80
85	AA	553	G	N3-C4-C5	-5.02	126.09	128.60
85	AA	630	A	C5'-C4'-O4'	-5.02	103.08	109.10
85	AA	737	G	O5'-P-OP2	5.02	116.72	110.70
85	AA	869	A	O4'-C4'-C3'	5.02	110.12	106.10
85	AA	1442	U	O4'-C1'-N1	5.02	112.22	108.20
85	AA	1835	U	C5'-C4'-O4'	5.02	115.12	109.10
85	AA	1836	U	C4'-C3'-C2'	-5.02	97.58	102.60
85	AA	2100	A	C2'-C3'-O3'	5.02	121.73	113.70
11	AC	66	SER	N-CA-C	-5.02	97.45	111.00
34	BA	19	G	C1'-O4'-C4'	-5.02	105.89	109.90
34	BA	1174	A	O3'-P-O5'	-5.02	94.47	104.00
34	BA	1216	G	C5'-C4'-O4'	5.02	115.12	109.10
34	BA	1627	U	C5'-C4'-O4'	5.02	115.12	109.10
34	BA	1695	G	C5'-C4'-O4'	5.02	115.12	109.10
35	BB	347	G	P-O5'-C5'	5.02	128.93	120.90
35	BB	690	C	C2-N3-C4	5.02	122.41	119.90
35	BB	864	U	O4'-C1'-N1	5.02	112.21	108.20
36	BC	114	C	N1-C1'-C2'	-5.02	106.48	112.00
36	BC	122	A	C3'-C2'-C1'	-5.02	97.49	101.50
37	BD	31	U	C2-N3-C4	-5.02	123.99	127.00
38	BE	30	C	C5'-C4'-O4'	-5.02	103.08	109.10
38	BE	99	C	O4'-C1'-N1	5.02	112.21	108.20
64	Be	193	ARG	NE-CZ-NH2	-5.02	117.79	120.30
66	Bg	52	LEU	CB-CG-CD1	5.02	119.53	111.00
77	Br	295	GLU	CA-C-N	-5.02	106.16	117.20
85	AA	329	G	OP1-P-OP2	-5.02	112.08	119.60
85	AA	373	G	C4-N9-C1'	-5.02	119.98	126.50
85	AA	385	A	P-O3'-C3'	-5.02	113.68	119.70
85	AA	766	G	C8-N9-C4	-5.02	104.39	106.40
85	AA	883	A	O5'-P-OP1	5.02	116.72	110.70
85	AA	986	U	C2'-C3'-O3'	5.02	121.73	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
85	AA	1185	G	C4'-C3'-C2'	5.02	107.62	102.60
85	AA	1466	U	C2'-C3'-O3'	5.02	121.73	113.70
85	AA	1685	G	N9-C1'-C2'	-5.02	106.48	112.00
85	AA	1695	G	P-O5'-C5'	-5.02	112.87	120.90
85	AA	2090	C	C3'-C2'-C1'	-5.02	97.48	101.50
2	A1	197	ARG	NE-CZ-NH1	5.02	122.81	120.30
12	AD	24	SER	N-CA-CB	5.02	118.03	110.50
34	BA	115	U	P-O5'-C5'	5.02	128.93	120.90
34	BA	125	G	C3'-C2'-C1'	-5.02	97.49	101.50
34	BA	408	U	C1'-O4'-C4'	-5.02	105.89	109.90
34	BA	680	C	C2'-C3'-O3'	5.02	121.73	113.70
35	BB	434	A	P-O3'-C3'	-5.02	113.68	119.70
35	BB	703	U	O4'-C1'-C2'	5.02	112.11	107.60
35	BB	1005	A	C8-N9-C1'	5.02	136.73	127.70
37	BD	92	G	C1'-O4'-C4'	-5.02	105.89	109.90
85	AA	420	C	P-O3'-C3'	-5.02	113.68	119.70
85	AA	1782	C	C6-N1-C2	-5.02	118.29	120.30
85	AA	2149	C	O3'-P-O5'	-5.02	94.47	104.00
10	A9	103	TYR	CB-CG-CD2	-5.01	117.99	121.00
34	BA	588	C	C5'-C4'-O4'	5.01	115.12	109.10
34	BA	599	U	C1'-O4'-C4'	-5.01	105.89	109.90
34	BA	880	G	C8-N9-C4	-5.01	104.39	106.40
34	BA	1384	G	N3-C2-N2	5.01	123.41	119.90
34	BA	1645	C	C1'-O4'-C4'	-5.01	105.89	109.90
35	BB	3	C	C4'-C3'-C2'	-5.01	97.59	102.60
35	BB	465	C	P-O5'-C5'	5.01	128.92	120.90
35	BB	712	U	O3'-P-O5'	-5.01	94.47	104.00
35	BB	1269	A	C5-C6-N1	5.01	120.21	117.70
35	BB	1287	U	P-O5'-C5'	5.01	128.93	120.90
35	BB	1545	U	O3'-P-O5'	-5.01	94.47	104.00
36	BC	69	U	N3-C4-C5	5.01	117.61	114.60
37	BD	106	G	C4'-C3'-C2'	-5.01	97.58	102.60
38	BE	173	G	C6-C5-N7	-5.01	127.39	130.40
40	BG	168	A	C5-C6-N1	5.01	120.21	117.70
51	BR	118	GLN	N-CA-C	-5.01	97.46	111.00
65	Bf	272	ASP	CB-CG-OD1	5.01	122.81	118.30
65	Bf	315	ARG	CD-NE-CZ	-5.01	116.58	123.60
77	Br	350	GLU	N-CA-CB	-5.01	101.58	110.60
83	Bx	120	ASP	N-CA-CB	-5.01	101.57	110.60
85	AA	778	C	N1-C2-N3	5.01	122.71	119.20
85	AA	1775	U	C6-N1-C2	-5.01	117.99	121.00
85	AA	1920	A	C2'-C3'-O3'	5.01	121.72	113.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	A5	121	LEU	C-N-CA	-5.01	111.77	122.30
34	BA	11	U	C5'-C4'-O4'	-5.01	103.08	109.10
34	BA	1634	A	O5'-P-OP1	5.01	116.72	110.70
35	BB	77	A	C4-C5-C6	-5.01	114.49	117.00
40	BG	135	C	N1-C1'-C2'	-5.01	106.48	112.00
50	BQ	29	LYS	CA-CB-CG	-5.01	102.37	113.40
60	Ba	84	TYR	CB-CG-CD1	-5.01	117.99	121.00
74	Bo	59	GLY	CA-C-N	-5.01	106.17	117.20
85	AA	239	G	C5-C6-O6	-5.01	125.59	128.60
85	AA	449	G	C1'-O4'-C4'	-5.01	105.89	109.90
85	AA	1196	C	C6-N1-C1'	5.01	126.82	120.80
85	AA	1469	G	OP1-P-OP2	-5.01	112.08	119.60
85	AA	2243	G	C8-N9-C4	-5.01	104.39	106.40
86	AB	49	C	P-O3'-C3'	5.01	125.72	119.70
34	BA	69	C	C4'-C3'-C2'	5.01	107.61	102.60
34	BA	421	G	C5'-C4'-C3'	-5.01	107.98	116.00
34	BA	432	A	C8-N9-C4	5.01	107.81	105.80
34	BA	919	A	C8-N9-C4	5.01	107.80	105.80
35	BB	16	G	O3'-P-O5'	5.01	113.52	104.00
35	BB	34	G	N1-C2-N2	-5.01	111.69	116.20
35	BB	63	A	O5'-P-OP2	-5.01	101.19	105.70
35	BB	89	C	C3'-C2'-C1'	-5.01	97.49	101.50
35	BB	645	C	C4'-C3'-C2'	5.01	107.61	102.60
35	BB	813	C	C6-N1-C1'	5.01	126.81	120.80
35	BB	979	G	P-O3'-C3'	-5.01	113.69	119.70
36	BC	7	U	O3'-P-O5'	-5.01	94.48	104.00
37	BD	95	G	N9-C4-C5	5.01	107.40	105.40
41	BH	20	A	P-O5'-C5'	-5.01	112.88	120.90
42	BI	103	PRO	C-N-CA	5.01	134.23	121.70
64	Be	155	LYS	N-CA-CB	5.01	119.62	110.60
65	Bf	294	LEU	CB-CA-C	5.01	119.72	110.20
74	Bo	85	ARG	CD-NE-CZ	-5.01	116.58	123.60
85	AA	334	A	N1-C2-N3	-5.01	126.79	129.30
85	AA	743	C	C5-C6-N1	-5.01	118.49	121.00
85	AA	815	G	C5-C6-N1	5.01	114.01	111.50
85	AA	1157	U	O3'-P-O5'	5.01	113.52	104.00
85	AA	1199	C	C6-N1-C2	-5.01	118.30	120.30
85	AA	1376	U	P-O3'-C3'	5.01	125.71	119.70
85	AA	1501	A	C6-N1-C2	-5.01	115.59	118.60
85	AA	2170	G	N9-C1'-C2'	-5.01	106.49	112.00
86	AB	67	C	C1'-O4'-C4'	-5.01	105.89	109.90
34	BA	290	G	N1-C2-N2	-5.01	111.69	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	BA	359	G	C3'-C2'-C1'	-5.01	97.49	101.50
34	BA	661	C	N1-C1'-C2'	-5.01	106.49	112.00
34	BA	901	C	C2-N3-C4	-5.01	117.40	119.90
34	BA	1724	G	C5'-C4'-O4'	-5.01	103.09	109.10
35	BB	612	A	C1'-O4'-C4'	-5.01	105.89	109.90
35	BB	1154	C	N3-C2-O2	-5.01	118.39	121.90
35	BB	1169	A	C1'-O4'-C4'	-5.01	105.89	109.90
35	BB	1331	U	C2-N1-C1'	5.01	123.71	117.70
36	BC	117	A	C5'-C4'-C3'	-5.01	107.98	116.00
37	BD	119	U	C5'-C4'-O4'	5.01	115.11	109.10
39	BF	12	U	C2-N3-C4	5.01	130.01	127.00
39	BF	54	U	N1-C2-N3	5.01	117.91	114.90
41	BH	100	A	C6-N1-C2	-5.01	115.59	118.60
41	BH	126	C	C1'-O4'-C4'	-5.01	105.89	109.90
85	AA	18	C	N3-C2-O2	-5.01	118.39	121.90
85	AA	29	U	N1-C1'-C2'	-5.01	106.49	112.00
85	AA	520	A	C1'-O4'-C4'	-5.01	105.89	109.90
85	AA	1026	U	P-O5'-C5'	-5.01	112.88	120.90
85	AA	1105	G	C2-N3-C4	-5.01	109.39	111.90
85	AA	1207	C	P-O5'-C5'	-5.01	112.89	120.90
85	AA	1998	A	OP1-P-OP2	-5.01	112.09	119.60
85	AA	2221	A	C5'-C4'-C3'	5.01	124.01	116.00
19	AK	31	PRO	CA-N-CD	-5.01	104.49	111.50
34	BA	101	G	C1'-O4'-C4'	-5.01	105.89	109.90
34	BA	196	A	C3'-C2'-C1'	-5.01	97.49	101.50
34	BA	501	U	O4'-C1'-C2'	-5.01	100.79	105.80
34	BA	622	G	C1'-O4'-C4'	-5.01	105.89	109.90
34	BA	1103	G	C5-C6-N1	5.01	114.00	111.50
34	BA	1151	A	OP2-P-O3'	5.01	116.22	105.20
35	BB	499	A	P-O5'-C5'	-5.01	112.89	120.90
35	BB	640	A	C5'-C4'-C3'	5.01	124.01	116.00
40	BG	173	C	C2-N3-C4	-5.01	117.40	119.90
84	By	7	ASP	N-CA-CB	-5.01	101.59	110.60
85	AA	354	C	C5-C4-N4	5.01	123.71	120.20
85	AA	1120	G	C1'-O4'-C4'	-5.01	105.89	109.90
85	AA	1146	C	N3-C4-N4	-5.01	114.49	118.00
85	AA	1484	G	O4'-C1'-N9	5.01	112.21	108.20
21	AM	25	ARG	N-CA-CB	5.01	119.61	110.60
34	BA	308	C	C6-N1-C2	-5.01	118.30	120.30
34	BA	643	U	N3-C4-O4	-5.01	115.90	119.40
34	BA	929	A	C2-N3-C4	-5.01	108.10	110.60
34	BA	997	U	C5-C4-O4	5.01	128.90	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	BB	323	C	C1'-O4'-C4'	-5.01	105.89	109.90
35	BB	804	U	N3-C2-O2	-5.01	118.69	122.20
35	BB	1254	G	C5'-C4'-C3'	-5.01	107.99	116.00
35	BB	1412	U	C4'-C3'-C2'	-5.01	97.59	102.60
35	BB	1434	G	C5-C6-N1	5.01	114.00	111.50
35	BB	1439	U	O4'-C1'-N1	5.01	112.21	108.20
37	BD	5	A	C4'-C3'-C2'	5.01	107.61	102.60
40	BG	165	C	C5-C6-N1	5.01	123.50	121.00
71	Bl	95	TRP	N-CA-CB	-5.01	101.59	110.60
85	AA	139	G	C5-C6-O6	-5.01	125.60	128.60
85	AA	334	A	C5'-C4'-O4'	5.01	115.11	109.10
85	AA	364	C	C5'-C4'-O4'	5.01	115.11	109.10
85	AA	1299	A	O4'-C1'-C2'	5.01	112.11	107.60
85	AA	1490	A	N3-C4-N9	-5.01	123.39	127.40
85	AA	1491	G	C5-C6-O6	-5.01	125.60	128.60
85	AA	1679	U	O5'-C5'-C4'	-5.01	102.19	111.70
85	AA	1998	A	N9-C1'-C2'	-5.01	106.49	112.00
34	BA	160	G	C5-C6-N1	5.00	114.00	111.50
34	BA	321	G	C3'-C2'-C1'	-5.00	97.50	101.50
34	BA	433	G	C4'-C3'-C2'	5.00	107.61	102.60
34	BA	1073	G	C8-N9-C4	-5.00	104.40	106.40
34	BA	1291	A	C1'-O4'-C4'	-5.00	105.90	109.90
34	BA	1560	U	C5-C6-N1	-5.00	120.20	122.70
35	BB	738	G	C5'-C4'-C3'	-5.00	107.99	116.00
35	BB	1156	U	N3-C2-O2	-5.00	118.70	122.20
41	BH	112	U	N3-C4-C5	5.00	117.60	114.60
48	BO	23	ASP	O-C-N	-5.00	114.69	122.70
85	AA	1056	C	N1-C1'-C2'	-5.00	106.49	112.00
85	AA	1574	C	N1-C2-N3	5.00	122.70	119.20
85	AA	1877	G	C5'-C4'-O4'	5.00	115.11	109.10
27	AT	68	LYS	N-CA-C	-5.00	97.49	111.00
34	BA	155	U	O5'-C5'-C4'	5.00	121.21	111.70
34	BA	313	C	C2-N1-C1'	-5.00	113.30	118.80
34	BA	999	G	N9-C4-C5	5.00	107.40	105.40
34	BA	1152	A	C2-N3-C4	-5.00	108.10	110.60
34	BA	1325	G	N3-C2-N2	5.00	123.40	119.90
34	BA	1556	A	O4'-C1'-C2'	5.00	112.10	107.60
34	BA	1794	A	O4'-C4'-C3'	-5.00	99.00	104.00
35	BB	574	G	O5'-C5'-C4'	-5.00	102.19	111.70
35	BB	754	U	C2-N3-C4	-5.00	124.00	127.00
37	BD	48	G	C4-C5-C6	-5.00	115.80	118.80
38	BE	89	G	O4'-C1'-N9	5.00	112.20	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	BE	179	A	P-O5'-C5'	5.00	128.91	120.90
40	BG	15	G	O4'-C4'-C3'	-5.00	99.00	104.00
40	BG	61	A	C8-N9-C4	5.00	107.80	105.80
85	AA	71	G	C5'-C4'-C3'	-5.00	107.99	116.00
85	AA	619	A	N1-C6-N6	-5.00	115.60	118.60
85	AA	932	A	N9-C4-C5	-5.00	103.80	105.80
85	AA	1652	A	N9-C1'-C2'	-5.00	106.50	112.00
85	AA	1871	U	O5'-C5'-C4'	-5.00	102.19	111.70
85	AA	1895	C	P-O3'-C3'	-5.00	113.69	119.70
85	AA	2059	A	P-O3'-C3'	5.00	125.70	119.70
85	AA	2151	U	C3'-C2'-C1'	-5.00	97.50	101.50
10	A9	125	ASN	CA-C-N	5.00	131.10	117.10
34	BA	478	G	C5'-C4'-O4'	5.00	115.10	109.10
34	BA	843	G	C3'-C2'-C1'	-5.00	97.50	101.50
34	BA	1015	G	C5-C6-O6	-5.00	125.60	128.60
34	BA	1122	G	P-O3'-C3'	-5.00	113.70	119.70
34	BA	1342	C	C3'-C2'-C1'	-5.00	97.50	101.50
34	BA	1551	G	N1-C6-O6	-5.00	116.90	119.90
35	BB	6	A	C4-N9-C1'	-5.00	117.30	126.30
35	BB	899	C	N1-C2-N3	5.00	122.70	119.20
35	BB	967	G	O4'-C1'-N9	5.00	112.20	108.20
35	BB	1336	G	N3-C2-N2	5.00	123.40	119.90
36	BC	58	G	N9-C1'-C2'	-5.00	106.50	112.00
36	BC	120	G	C2-N3-C4	-5.00	109.40	111.90
37	BD	17	G	N1-C6-O6	5.00	122.90	119.90
39	BF	24	G	N9-C1'-C2'	-5.00	106.50	112.00
39	BF	58	U	C2-N1-C1'	-5.00	111.70	117.70
44	BK	195	LEU	C-N-CA	5.00	134.20	121.70
63	Bd	28	TYR	C-N-CA	5.00	134.21	121.70
81	Bv	178	VAL	N-CA-C	-5.00	97.50	111.00
85	AA	1267	A	C5'-C4'-O4'	5.00	115.10	109.10
85	AA	1446	U	C6-N1-C2	-5.00	118.00	121.00
85	AA	2191	C	P-O3'-C3'	-5.00	113.70	119.70
85	AA	2192	A	C4-N9-C1'	-5.00	117.30	126.30
85	AA	2198	G	N1-C2-N3	5.00	126.90	123.90

All (3) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
47	BN	66	PRO	CA
69	Bj	5	ARG	CA
84	By	7	ASP	CA

All (5934) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A0	103	HIS	Sidechain
1	A0	111	ARG	Sidechain
1	A0	114	TYR	Sidechain
1	A0	138	ARG	Sidechain
1	A0	155	ASN	Mainchain
1	A0	157	TYR	Sidechain
1	A0	167	ARG	Sidechain
1	A0	169	ARG	Sidechain
1	A0	176	ARG	Sidechain
1	A0	192	ARG	Sidechain
1	A0	202	ARG	Sidechain
1	A0	210	ARG	Sidechain
1	A0	211	ASP	Peptide
1	A0	213	ARG	Sidechain
1	A0	221	ARG	Sidechain
1	A0	41	ARG	Sidechain
1	A0	43	PHE	Sidechain
1	A0	64	ARG	Sidechain
1	A0	83	TYR	Sidechain
2	A1	10	TYR	Sidechain
2	A1	100	TYR	Sidechain
2	A1	127	TYR	Sidechain
2	A1	132	ARG	Sidechain
2	A1	139	HIS	Sidechain
2	A1	153	ARG	Sidechain
2	A1	159	TYR	Sidechain
2	A1	186	ARG	Sidechain
2	A1	188	ARG	Sidechain
2	A1	206	ARG	Sidechain
2	A1	218	ARG	Sidechain
2	A1	27	ARG	Sidechain
2	A1	29	ARG	Sidechain
2	A1	33	HIS	Sidechain
2	A1	51	TYR	Sidechain
2	A1	87	VAL	Mainchain
2	A1	96	PHE	Sidechain
2	A1	97	ARG	Sidechain
3	A2	122	ARG	Sidechain
3	A2	131	ARG	Sidechain
3	A2	145	ARG	Sidechain
3	A2	190	ARG	Sidechain
3	A2	27	TYR	Sidechain

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Mol	Chain	Res	Type	Group
3	A2	46	ARG	Sidechain
3	A2	48	ARG	Sidechain
3	A2	57	ARG	Sidechain
3	A2	64	PHE	Sidechain
3	A2	80	ARG	Sidechain
3	A2	9	PHE	Sidechain
4	A3	101	ARG	Sidechain
4	A3	104	ILE	Peptide
4	A3	124	ILE	Peptide
4	A3	134	ARG	Sidechain
4	A3	152	ARG	Sidechain
4	A3	162	ARG	Sidechain
4	A3	170	LYS	Peptide
4	A3	173	ARG	Sidechain
4	A3	190	ARG	Sidechain
4	A3	209	ARG	Sidechain
4	A3	212	TYR	Sidechain
4	A3	224	ARG	Sidechain
4	A3	226	ARG	Sidechain
4	A3	25	ARG	Sidechain
4	A3	29	GLY	Peptide
4	A3	31	TYR	Sidechain
4	A3	32	ARG	Sidechain
4	A3	47	ARG	Sidechain
4	A3	59	LYS	Mainchain
4	A3	73	ARG	Sidechain
4	A3	85	PHE	Sidechain
4	A3	89	ARG	Sidechain
4	A3	91	TYR	Sidechain
5	A4	11	ARG	Sidechain
5	A4	114	LEU	Mainchain
5	A4	12	LYS	Mainchain
5	A4	135	ILE	Peptide
5	A4	143	ARG	Sidechain
5	A4	144	ARG	Sidechain
5	A4	146	ARG	Sidechain
5	A4	15	ARG	Sidechain
5	A4	163	ASP	Peptide,Mainchain
5	A4	166	ARG	Sidechain
5	A4	178	TYR	Sidechain
5	A4	184	ARG	Sidechain
5	A4	188	PHE	Sidechain

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Mol	Chain	Res	Type	Group
5	A4	27	ARG	Sidechain
5	A4	35	SER	Mainchain
5	A4	40	ARG	Sidechain
5	A4	45	ARG	Sidechain
5	A4	46	PHE	Sidechain
5	A4	52	ARG	Sidechain
5	A4	60	LYS	Peptide
5	A4	66	ILE	Mainchain
5	A4	68	TYR	Sidechain
5	A4	81	ARG	Sidechain
5	A4	9	LYS	Mainchain
5	A4	90	ARG	Sidechain
5	A4	91	PHE	Sidechain
6	A5	11	ARG	Sidechain
6	A5	110	ARG	Sidechain
6	A5	113	TYR	Sidechain
6	A5	12	LYS	Mainchain
6	A5	13	ILE	Mainchain
6	A5	157	LYS	Peptide,Mainchain
6	A5	160	ARG	Sidechain
6	A5	164	TYR	Sidechain
6	A5	165	ARG	Sidechain
6	A5	172	GLU	Peptide
6	A5	183	ARG	Sidechain
6	A5	187	ARG	Sidechain
6	A5	197	ARG	Sidechain
6	A5	21	HIS	Sidechain
6	A5	22	ARG	Sidechain
6	A5	24	ARG	Sidechain
6	A5	31	ARG	Sidechain
6	A5	42	ARG	Sidechain
6	A5	49	ARG	Sidechain
6	A5	5	ARG	Sidechain
6	A5	7	ARG	Sidechain
6	A5	77	ARG	Sidechain
7	A6	106	ARG	Sidechain
7	A6	113	PHE	Sidechain
7	A6	122	HIS	Sidechain
7	A6	125	ARG	Sidechain
7	A6	131	ARG	Sidechain
7	A6	132	HIS	Sidechain
7	A6	139	ILE	Mainchain

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Mol	Chain	Res	Type	Group
7	A6	15	ARG	Sidechain
7	A6	157	PHE	Peptide,Mainchain,Sidechain
7	A6	170	ARG	Sidechain
7	A6	176	ARG	Sidechain
7	A6	33	TYR	Sidechain
7	A6	36	ARG	Sidechain
7	A6	39	ARG	Sidechain
7	A6	4	TYR	Sidechain
7	A6	52	ARG	Sidechain
7	A6	65	HIS	Sidechain
7	A6	68	ARG	Sidechain
7	A6	77	ARG	Sidechain
7	A6	78	ARG	Sidechain
7	A6	82	TYR	Sidechain
7	A6	84	PHE	Sidechain
7	A6	9	ARG	Sidechain
7	A6	94	TYR	Sidechain
8	A7	103	TYR	Sidechain
8	A7	117	PHE	Sidechain
8	A7	134	ARG	Sidechain
8	A7	159	ARG	Sidechain
8	A7	187	ARG	Sidechain
8	A7	205	SER	Mainchain
8	A7	249	ARG	Sidechain
8	A7	261	ARG	Sidechain
8	A7	284	ILE	Peptide
8	A7	301	TYR	Sidechain
8	A7	310	ARG	Sidechain
8	A7	46	ASN	Peptide
8	A7	49	ARG	Sidechain
8	A7	50	HIS	Sidechain
8	A7	56	TYR	Sidechain
8	A7	62	ARG	Sidechain
8	A7	92	ARG	Sidechain
9	A8	20	ARG	Sidechain
9	A8	33	ARG	Sidechain
9	A8	41	ARG	Sidechain
9	A8	44	PHE	Sidechain
10	A9	103	TYR	Sidechain
10	A9	121	ASP	Peptide
10	A9	150	TYR	Sidechain
10	A9	99	ARG	Peptide

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Mol	Chain	Res	Type	Group
85	AA	1	G	Sidechain
85	AA	10	G	Sidechain
85	AA	100	A	Sidechain
85	AA	1000	U	Sidechain
85	AA	1001	G	Sidechain
85	AA	1002	G	Sidechain
85	AA	1003	G	Sidechain
85	AA	1007	G	Sidechain
85	AA	101	C	Sidechain
85	AA	1010	U	Sidechain
85	AA	1011	G	Sidechain
85	AA	1015	U	Sidechain
85	AA	1016	G	Sidechain
85	AA	1018	G	Sidechain
85	AA	1019	U	Sidechain
85	AA	102	A	Sidechain
85	AA	1021	G	Sidechain
85	AA	1022	G	Sidechain
85	AA	1023	U	Sidechain
85	AA	1024	G	Sidechain
85	AA	1027	U	Sidechain
85	AA	1028	C	Sidechain
85	AA	1030	U	Sidechain
85	AA	1033	C	Sidechain
85	AA	1034	U	Sidechain
85	AA	1037	U	Sidechain
85	AA	104	C	Sidechain
85	AA	1042	G	Sidechain
85	AA	1043	U	Sidechain
85	AA	1044	G	Sidechain
85	AA	105	A	Sidechain
85	AA	1050	C	Sidechain
85	AA	1051	A	Sidechain
85	AA	1053	A	Sidechain
85	AA	1055	U	Sidechain
85	AA	1056	C	Sidechain
85	AA	1058	G	Sidechain
85	AA	106	G	Sidechain
85	AA	1060	U	Sidechain
85	AA	1063	U	Sidechain
85	AA	1064	C	Sidechain
85	AA	1065	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1066	U	Sidechain
85	AA	1067	G	Sidechain
85	AA	1070	G	Sidechain
85	AA	1071	U	Sidechain
85	AA	1072	U	Sidechain
85	AA	1077	U	Sidechain
85	AA	108	C	Sidechain
85	AA	1081	U	Sidechain
85	AA	1082	U	Sidechain
85	AA	1083	C	Sidechain
85	AA	1086	U	Sidechain
85	AA	1089	G	Sidechain
85	AA	109	G	Sidechain
85	AA	1090	A	Sidechain
85	AA	1091	C	Sidechain
85	AA	1092	G	Sidechain
85	AA	1093	C	Sidechain
85	AA	1094	G	Sidechain
85	AA	1096	G	Sidechain
85	AA	1097	G	Sidechain
85	AA	1098	C	Sidechain
85	AA	1099	U	Sidechain
85	AA	110	U	Sidechain
85	AA	1100	U	Sidechain
85	AA	1101	C	Sidechain
85	AA	1102	C	Sidechain
85	AA	1103	A	Sidechain
85	AA	1104	G	Sidechain
85	AA	1105	G	Sidechain
85	AA	1106	A	Sidechain
85	AA	1107	A	Sidechain
85	AA	1108	U	Sidechain
85	AA	1109	G	Sidechain
85	AA	1110	A	Sidechain
85	AA	1111	A	Sidechain
85	AA	1112	G	Sidechain
85	AA	1114	A	Sidechain
85	AA	1116	G	Sidechain
85	AA	1117	G	Sidechain
85	AA	1118	U	Sidechain
85	AA	1119	A	Sidechain
85	AA	1120	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1121	U	Sidechain
85	AA	1122	U	Sidechain
85	AA	1123	C	Sidechain
85	AA	1124	G	Sidechain
85	AA	1125	G	Sidechain
85	AA	1127	G	Sidechain
85	AA	1128	G	Sidechain
85	AA	1129	A	Sidechain
85	AA	113	U	Sidechain
85	AA	1130	G	Sidechain
85	AA	1131	A	Sidechain
85	AA	1132	A	Sidechain
85	AA	1133	C	Sidechain
85	AA	1134	G	Sidechain
85	AA	1135	U	Sidechain
85	AA	1136	A	Sidechain
85	AA	1137	C	Sidechain
85	AA	1139	G	Sidechain
85	AA	114	C	Sidechain
85	AA	1140	G	Sidechain
85	AA	1141	U	Sidechain
85	AA	1143	C	Sidechain
85	AA	1145	U	Sidechain
85	AA	1146	C	Sidechain
85	AA	1147	A	Sidechain
85	AA	1148	G	Sidechain
85	AA	1149	A	Sidechain
85	AA	115	U	Sidechain
85	AA	1151	G	Sidechain
85	AA	1152	U	Sidechain
85	AA	1153	G	Sidechain
85	AA	1155	A	Sidechain
85	AA	1157	U	Sidechain
85	AA	1158	U	Sidechain
85	AA	1159	C	Sidechain
85	AA	116	G	Sidechain
85	AA	1160	U	Sidechain
85	AA	1161	U	Sidechain
85	AA	1162	A	Sidechain
85	AA	1163	G	Sidechain
85	AA	1165	C	Sidechain
85	AA	1166	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1167	G	Sidechain
85	AA	1168	C	Sidechain
85	AA	117	C	Sidechain
85	AA	1170	C	Sidechain
85	AA	1171	C	Sidechain
85	AA	1172	A	Sidechain
85	AA	1173	A	Sidechain
85	AA	1174	G	Sidechain
85	AA	1175	A	Sidechain
85	AA	1176	C	Sidechain
85	AA	1177	G	Sidechain
85	AA	118	C	Sidechain
85	AA	1180	C	Sidechain
85	AA	1181	U	Sidechain
85	AA	1182	A	Sidechain
85	AA	1184	A	Sidechain
85	AA	1185	G	Sidechain
85	AA	1187	G	Sidechain
85	AA	1188	A	Sidechain
85	AA	1189	A	Sidechain
85	AA	119	G	Sidechain
85	AA	1190	G	Sidechain
85	AA	1191	G	Sidechain
85	AA	1193	A	Sidechain
85	AA	1194	U	Sidechain
85	AA	1195	U	Sidechain
85	AA	1196	C	Sidechain
85	AA	1197	U	Sidechain
85	AA	1199	C	Sidechain
85	AA	12	U	Sidechain
85	AA	120	C	Sidechain
85	AA	1200	A	Sidechain
85	AA	1201	A	Sidechain
85	AA	1202	G	Sidechain
85	AA	1205	U	Sidechain
85	AA	1206	A	Sidechain
85	AA	1207	C	Sidechain
85	AA	1209	U	Sidechain
85	AA	121	C	Sidechain
85	AA	1210	U	Sidechain
85	AA	1211	C	Sidechain
85	AA	1212	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1213	U	Sidechain
85	AA	1214	C	Sidechain
85	AA	1215	A	Sidechain
85	AA	1218	C	Sidechain
85	AA	1219	A	Sidechain
85	AA	122	A	Sidechain
85	AA	1220	A	Sidechain
85	AA	1221	G	Sidechain
85	AA	1224	C	Sidechain
85	AA	1225	C	Sidechain
85	AA	1226	A	Sidechain
85	AA	1227	A	Sidechain
85	AA	1228	A	Sidechain
85	AA	1229	G	Sidechain
85	AA	1230	U	Sidechain
85	AA	1231	G	Sidechain
85	AA	1233	G	Sidechain
85	AA	1234	G	Sidechain
85	AA	1235	G	Sidechain
85	AA	1236	G	Sidechain
85	AA	1237	A	Sidechain
85	AA	1238	U	Sidechain
85	AA	1239	C	Sidechain
85	AA	124	A	Sidechain
85	AA	1240	A	Sidechain
85	AA	1241	A	Sidechain
85	AA	1242	A	Sidechain
85	AA	1243	G	Sidechain
85	AA	1244	A	Sidechain
85	AA	1245	U	Sidechain
85	AA	1246	G	Sidechain
85	AA	1248	U	Sidechain
85	AA	1251	G	Sidechain
85	AA	1253	G	Sidechain
85	AA	1257	A	Sidechain
85	AA	1258	U	Sidechain
85	AA	1259	U	Sidechain
85	AA	126	U	Sidechain
85	AA	1260	G	Sidechain
85	AA	1261	U	Sidechain
85	AA	1265	C	Sidechain
85	AA	1268	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	127	U	Sidechain
85	AA	1270	C	Sidechain
85	AA	1272	G	Sidechain
85	AA	1273	C	Sidechain
85	AA	1275	A	Sidechain
85	AA	1277	C	Sidechain
85	AA	1278	C	Sidechain
85	AA	1279	A	Sidechain
85	AA	128	U	Sidechain
85	AA	1280	U	Sidechain
85	AA	1281	G	Sidechain
85	AA	1282	A	Sidechain
85	AA	1283	C	Sidechain
85	AA	1284	A	Sidechain
85	AA	1285	C	Sidechain
85	AA	1287	C	Sidechain
85	AA	1288	A	Sidechain
85	AA	1289	U	Sidechain
85	AA	129	U	Sidechain
85	AA	1290	G	Sidechain
85	AA	1291	A	Sidechain
85	AA	1292	A	Sidechain
85	AA	1293	U	Sidechain
85	AA	1294	U	Sidechain
85	AA	1295	G	Sidechain
85	AA	1297	G	Sidechain
85	AA	1299	A	Sidechain
85	AA	13	U	Sidechain
85	AA	130	G	Sidechain
85	AA	1301	C	Sidechain
85	AA	1302	A	Sidechain
85	AA	1304	C	Sidechain
85	AA	1305	A	Sidechain
85	AA	1306	U	Sidechain
85	AA	1310	G	Sidechain
85	AA	1317	U	Sidechain
85	AA	1318	G	Sidechain
85	AA	132	G	Sidechain
85	AA	1324	G	Sidechain
85	AA	1328	U	Sidechain
85	AA	133	G	Sidechain
85	AA	1338	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1341	U	Sidechain
85	AA	1346	C	Sidechain
85	AA	135	C	Sidechain
85	AA	1351	U	Sidechain
85	AA	1352	U	Sidechain
85	AA	1355	U	Sidechain
85	AA	1356	U	Sidechain
85	AA	1357	U	Sidechain
85	AA	1359	U	Sidechain
85	AA	136	U	Sidechain
85	AA	1361	A	Sidechain
85	AA	1363	U	Sidechain
85	AA	1365	U	Sidechain
85	AA	1366	A	Sidechain
85	AA	1367	C	Sidechain
85	AA	1368	G	Sidechain
85	AA	1369	U	Sidechain
85	AA	137	C	Sidechain
85	AA	1371	C	Sidechain
85	AA	1373	U	Sidechain
85	AA	1374	A	Sidechain
85	AA	1375	U	Sidechain
85	AA	1378	U	Sidechain
85	AA	138	C	Sidechain
85	AA	1380	U	Sidechain
85	AA	1383	C	Sidechain
85	AA	139	G	Sidechain
85	AA	1390	U	Sidechain
85	AA	14	C	Sidechain
85	AA	1409	U	Sidechain
85	AA	1413	G	Sidechain
85	AA	1419	U	Sidechain
85	AA	142	U	Sidechain
85	AA	1421	U	Sidechain
85	AA	1422	A	Sidechain
85	AA	1423	C	Sidechain
85	AA	1427	A	Sidechain
85	AA	1429	U	Sidechain
85	AA	143	U	Sidechain
85	AA	1430	A	Sidechain
85	AA	1431	U	Sidechain
85	AA	1432	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1433	C	Sidechain
85	AA	1434	U	Sidechain
85	AA	1435	C	Sidechain
85	AA	1438	C	Sidechain
85	AA	1441	G	Sidechain
85	AA	1442	U	Sidechain
85	AA	1443	U	Sidechain
85	AA	1444	U	Sidechain
85	AA	1445	C	Sidechain
85	AA	1446	U	Sidechain
85	AA	1447	U	Sidechain
85	AA	1448	A	Sidechain
85	AA	145	C	Sidechain
85	AA	1450	U	Sidechain
85	AA	1451	U	Sidechain
85	AA	1452	C	Sidechain
85	AA	1453	U	Sidechain
85	AA	1455	C	Sidechain
85	AA	1456	A	Sidechain
85	AA	1458	G	Sidechain
85	AA	1459	C	Sidechain
85	AA	146	U	Sidechain
85	AA	1460	G	Sidechain
85	AA	1462	A	Sidechain
85	AA	1464	G	Sidechain
85	AA	1465	C	Sidechain
85	AA	1466	U	Sidechain
85	AA	1467	U	Sidechain
85	AA	1468	G	Sidechain
85	AA	1469	G	Sidechain
85	AA	147	G	Sidechain
85	AA	1470	A	Sidechain
85	AA	1471	G	Sidechain
85	AA	1473	U	Sidechain
85	AA	1474	U	Sidechain
85	AA	1476	C	Sidechain
85	AA	1477	A	Sidechain
85	AA	1478	G	Sidechain
85	AA	1479	U	Sidechain
85	AA	148	G	Sidechain
85	AA	1480	C	Sidechain
85	AA	1481	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1482	C	Sidechain
85	AA	1483	A	Sidechain
85	AA	1484	G	Sidechain
85	AA	1485	G	Sidechain
85	AA	1487	G	Sidechain
85	AA	1488	G	Sidechain
85	AA	1489	G	Sidechain
85	AA	1490	A	Sidechain
85	AA	1491	G	Sidechain
85	AA	1492	U	Sidechain
85	AA	1493	A	Sidechain
85	AA	1495	G	Sidechain
85	AA	1497	U	Sidechain
85	AA	1498	C	Sidechain
85	AA	1499	G	Sidechain
85	AA	15	U	Sidechain
85	AA	150	U	Sidechain
85	AA	1501	A	Sidechain
85	AA	1502	A	Sidechain
85	AA	1503	G	Sidechain
85	AA	1505	G	Sidechain
85	AA	1506	U	Sidechain
85	AA	1507	G	Sidechain
85	AA	1508	A	Sidechain
85	AA	1509	A	Sidechain
85	AA	151	A	Sidechain
85	AA	1510	A	Sidechain
85	AA	1511	C	Sidechain
85	AA	1512	U	Sidechain
85	AA	1513	U	Sidechain
85	AA	1515	A	Sidechain
85	AA	1516	A	Sidechain
85	AA	1517	G	Sidechain
85	AA	1518	A	Sidechain
85	AA	1519	A	Sidechain
85	AA	152	A	Sidechain
85	AA	1521	U	Sidechain
85	AA	1522	U	Sidechain
85	AA	1523	G	Sidechain
85	AA	1524	A	Sidechain
85	AA	1525	C	Sidechain
85	AA	1526	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1529	A	Sidechain
85	AA	153	C	Sidechain
85	AA	1530	U	Sidechain
85	AA	1531	G	Sidechain
85	AA	1532	G	Sidechain
85	AA	1533	C	Sidechain
85	AA	1535	C	Sidechain
85	AA	1536	C	Sidechain
85	AA	1538	C	Sidechain
85	AA	1539	A	Sidechain
85	AA	154	U	Sidechain
85	AA	1540	A	Sidechain
85	AA	1541	G	Sidechain
85	AA	1542	A	Sidechain
85	AA	1543	C	Sidechain
85	AA	1544	G	Sidechain
85	AA	1545	U	Sidechain
85	AA	1546	G	Sidechain
85	AA	1548	A	Sidechain
85	AA	1549	G	Sidechain
85	AA	155	U	Sidechain
85	AA	1551	G	Sidechain
85	AA	1552	U	Sidechain
85	AA	1553	G	Sidechain
85	AA	1554	C	Sidechain
85	AA	1555	G	Sidechain
85	AA	1556	G	Sidechain
85	AA	1557	U	Sidechain
85	AA	1559	U	Sidechain
85	AA	156	G	Sidechain
85	AA	1560	A	Sidechain
85	AA	1561	A	Sidechain
85	AA	1562	U	Sidechain
85	AA	1563	U	Sidechain
85	AA	1564	U	Sidechain
85	AA	1565	G	Sidechain
85	AA	1566	A	Sidechain
85	AA	1569	C	Sidechain
85	AA	157	G	Sidechain
85	AA	1572	C	Sidechain
85	AA	1573	A	Sidechain
85	AA	1574	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1575	G	Sidechain
85	AA	1576	G	Sidechain
85	AA	1577	G	Sidechain
85	AA	1578	G	Sidechain
85	AA	1579	A	Sidechain
85	AA	158	C	Sidechain
85	AA	1580	A	Sidechain
85	AA	1581	C	Sidechain
85	AA	1582	U	Sidechain
85	AA	1583	U	Sidechain
85	AA	1584	U	Sidechain
85	AA	1585	A	Sidechain
85	AA	1589	G	Sidechain
85	AA	159	G	Sidechain
85	AA	1590	A	Sidechain
85	AA	1591	U	Sidechain
85	AA	1592	C	Sidechain
85	AA	1593	C	Sidechain
85	AA	1594	G	Sidechain
85	AA	1595	G	Sidechain
85	AA	1597	C	Sidechain
85	AA	1598	A	Sidechain
85	AA	1599	G	Sidechain
85	AA	160	A	Sidechain
85	AA	1600	G	Sidechain
85	AA	1601	G	Sidechain
85	AA	1603	G	Sidechain
85	AA	1604	A	Sidechain
85	AA	1605	G	Sidechain
85	AA	1606	G	Sidechain
85	AA	1608	U	Sidechain
85	AA	1609	U	Sidechain
85	AA	161	A	Sidechain
85	AA	1612	C	Sidechain
85	AA	1613	A	Sidechain
85	AA	1614	G	Sidechain
85	AA	1617	G	Sidechain
85	AA	1618	G	Sidechain
85	AA	1619	A	Sidechain
85	AA	162	A	Sidechain
85	AA	1620	G	Sidechain
85	AA	1621	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1622	G	Sidechain
85	AA	1623	U	Sidechain
85	AA	1624	U	Sidechain
85	AA	1626	U	Sidechain
85	AA	1629	C	Sidechain
85	AA	163	C	Sidechain
85	AA	1630	U	Sidechain
85	AA	1631	C	Sidechain
85	AA	1632	G	Sidechain
85	AA	1634	U	Sidechain
85	AA	1635	C	Sidechain
85	AA	1636	C	Sidechain
85	AA	1637	C	Sidechain
85	AA	164	G	Sidechain
85	AA	1641	A	Sidechain
85	AA	1642	A	Sidechain
85	AA	1643	U	Sidechain
85	AA	1644	G	Sidechain
85	AA	1645	G	Sidechain
85	AA	1649	U	Sidechain
85	AA	1650	G	Sidechain
85	AA	1653	U	Sidechain
85	AA	1655	G	Sidechain
85	AA	1656	C	Sidechain
85	AA	1657	C	Sidechain
85	AA	166	C	Sidechain
85	AA	1660	U	Sidechain
85	AA	1661	U	Sidechain
85	AA	1662	U	Sidechain
85	AA	1663	U	Sidechain
85	AA	1666	U	Sidechain
85	AA	1667	C	Sidechain
85	AA	1668	G	Sidechain
85	AA	1669	G	Sidechain
85	AA	1670	U	Sidechain
85	AA	1671	G	Sidechain
85	AA	1672	G	Sidechain
85	AA	1673	A	Sidechain
85	AA	1674	G	Sidechain
85	AA	1676	G	Sidechain
85	AA	1677	A	Sidechain
85	AA	1679	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1680	U	Sidechain
85	AA	1681	G	Sidechain
85	AA	1682	U	Sidechain
85	AA	1683	U	Sidechain
85	AA	1684	U	Sidechain
85	AA	1685	G	Sidechain
85	AA	1686	G	Sidechain
85	AA	1687	U	Sidechain
85	AA	1688	U	Sidechain
85	AA	1689	G	Sidechain
85	AA	169	G	Sidechain
85	AA	1690	A	Sidechain
85	AA	1691	U	Sidechain
85	AA	1693	C	Sidechain
85	AA	1694	C	Sidechain
85	AA	1696	U	Sidechain
85	AA	1698	A	Sidechain
85	AA	1699	A	Sidechain
85	AA	17	C	Sidechain
85	AA	1700	C	Sidechain
85	AA	1703	A	Sidechain
85	AA	1704	C	Sidechain
85	AA	1705	G	Sidechain
85	AA	1706	A	Sidechain
85	AA	1707	G	Sidechain
85	AA	1708	A	Sidechain
85	AA	1709	U	Sidechain
85	AA	171	U	Sidechain
85	AA	1710	C	Sidechain
85	AA	1711	C	Sidechain
85	AA	1712	A	Sidechain
85	AA	1713	A	Sidechain
85	AA	1714	G	Sidechain
85	AA	1715	C	Sidechain
85	AA	1716	U	Sidechain
85	AA	1717	G	Sidechain
85	AA	1718	C	Sidechain
85	AA	1719	C	Sidechain
85	AA	1720	C	Sidechain
85	AA	1721	A	Sidechain
85	AA	1726	G	Sidechain
85	AA	1727	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1728	G	Sidechain
85	AA	1729	C	Sidechain
85	AA	173	A	Sidechain
85	AA	1730	C	Sidechain
85	AA	1731	G	Sidechain
85	AA	1732	G	Sidechain
85	AA	1733	G	Sidechain
85	AA	1734	A	Sidechain
85	AA	1735	U	Sidechain
85	AA	1736	U	Sidechain
85	AA	174	U	Sidechain
85	AA	175	A	Sidechain
85	AA	1751	G	Sidechain
85	AA	1752	C	Sidechain
85	AA	1754	G	Sidechain
85	AA	1755	U	Sidechain
85	AA	1756	C	Sidechain
85	AA	1759	U	Sidechain
85	AA	176	C	Sidechain
85	AA	1762	G	Sidechain
85	AA	1766	G	Sidechain
85	AA	1768	G	Sidechain
85	AA	1771	U	Sidechain
85	AA	1779	C	Sidechain
85	AA	178	U	Sidechain
85	AA	1781	A	Sidechain
85	AA	1782	C	Sidechain
85	AA	1784	G	Sidechain
85	AA	1785	U	Sidechain
85	AA	1786	G	Sidechain
85	AA	1787	G	Sidechain
85	AA	1789	C	Sidechain
85	AA	179	G	Sidechain
85	AA	1790	G	Sidechain
85	AA	1791	U	Sidechain
85	AA	1793	A	Sidechain
85	AA	1794	U	Sidechain
85	AA	1795	C	Sidechain
85	AA	1797	U	Sidechain
85	AA	1799	C	Sidechain
85	AA	18	C	Sidechain
85	AA	180	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1800	U	Sidechain
85	AA	1802	U	Sidechain
85	AA	1805	A	Sidechain
85	AA	1806	C	Sidechain
85	AA	1808	G	Sidechain
85	AA	1809	G	Sidechain
85	AA	181	A	Sidechain
85	AA	1811	C	Sidechain
85	AA	1812	C	Sidechain
85	AA	1814	U	Sidechain
85	AA	1815	U	Sidechain
85	AA	1816	C	Sidechain
85	AA	1817	U	Sidechain
85	AA	1819	U	Sidechain
85	AA	182	C	Sidechain
85	AA	1820	G	Sidechain
85	AA	1822	G	Sidechain
85	AA	1824	G	Sidechain
85	AA	1825	A	Sidechain
85	AA	1826	U	Sidechain
85	AA	1829	C	Sidechain
85	AA	1830	U	Sidechain
85	AA	1831	U	Sidechain
85	AA	1833	C	Sidechain
85	AA	1835	U	Sidechain
85	AA	1836	U	Sidechain
85	AA	1839	G	Sidechain
85	AA	184	A	Sidechain
85	AA	1840	C	Sidechain
85	AA	1842	C	Sidechain
85	AA	1844	A	Sidechain
85	AA	1845	G	Sidechain
85	AA	1846	G	Sidechain
85	AA	1848	G	Sidechain
85	AA	1849	A	Sidechain
85	AA	185	A	Sidechain
85	AA	1850	G	Sidechain
85	AA	1851	A	Sidechain
85	AA	1852	U	Sidechain
85	AA	1853	U	Sidechain
85	AA	1854	U	Sidechain
85	AA	1858	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1860	A	Sidechain
85	AA	1861	A	Sidechain
85	AA	1862	C	Sidechain
85	AA	1863	A	Sidechain
85	AA	1864	G	Sidechain
85	AA	1865	C	Sidechain
85	AA	1866	A	Sidechain
85	AA	1869	U	Sidechain
85	AA	187	C	Sidechain
85	AA	1872	G	Sidechain
85	AA	1873	U	Sidechain
85	AA	1874	G	Sidechain
85	AA	1875	A	Sidechain
85	AA	1876	U	Sidechain
85	AA	1878	C	Sidechain
85	AA	1879	U	Sidechain
85	AA	188	G	Sidechain
85	AA	1881	C	Sidechain
85	AA	1882	U	Sidechain
85	AA	1883	C	Sidechain
85	AA	1885	A	Sidechain
85	AA	1887	G	Sidechain
85	AA	1888	U	Sidechain
85	AA	1889	U	Sidechain
85	AA	1890	C	Sidechain
85	AA	1893	G	Sidechain
85	AA	1894	G	Sidechain
85	AA	1896	G	Sidechain
85	AA	1898	C	Sidechain
85	AA	1899	A	Sidechain
85	AA	190	A	Sidechain
85	AA	1900	C	Sidechain
85	AA	1902	C	Sidechain
85	AA	1904	C	Sidechain
85	AA	1906	C	Sidechain
85	AA	1907	U	Sidechain
85	AA	1908	A	Sidechain
85	AA	191	C	Sidechain
85	AA	1910	A	Sidechain
85	AA	1913	G	Sidechain
85	AA	1918	U	Sidechain
85	AA	1919	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	192	G	Sidechain
85	AA	1920	A	Sidechain
85	AA	1921	G	Sidechain
85	AA	1922	A	Sidechain
85	AA	1923	A	Sidechain
85	AA	1924	C	Sidechain
85	AA	1925	A	Sidechain
85	AA	1926	A	Sidechain
85	AA	1927	G	Sidechain
85	AA	1929	G	Sidechain
85	AA	193	C	Sidechain
85	AA	1930	U	Sidechain
85	AA	1931	C	Sidechain
85	AA	1933	G	Sidechain
85	AA	1935	G	Sidechain
85	AA	1938	G	Sidechain
85	AA	194	U	Sidechain
85	AA	1940	A	Sidechain
85	AA	1943	U	Sidechain
85	AA	1947	A	Sidechain
85	AA	1948	A	Sidechain
85	AA	1949	U	Sidechain
85	AA	195	C	Sidechain
85	AA	1951	U	Sidechain
85	AA	1953	G	Sidechain
85	AA	1954	C	Sidechain
85	AA	1955	U	Sidechain
85	AA	1956	C	Sidechain
85	AA	1959	G	Sidechain
85	AA	196	U	Sidechain
85	AA	1960	C	Sidechain
85	AA	1961	U	Sidechain
85	AA	1962	U	Sidechain
85	AA	1963	G	Sidechain
85	AA	1964	A	Sidechain
85	AA	1968	A	Sidechain
85	AA	1969	A	Sidechain
85	AA	197	C	Sidechain
85	AA	1972	A	Sidechain
85	AA	1973	G	Sidechain
85	AA	1976	G	Sidechain
85	AA	1978	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	1979	A	Sidechain
85	AA	198	U	Sidechain
85	AA	1981	A	Sidechain
85	AA	1982	C	Sidechain
85	AA	1983	C	Sidechain
85	AA	1984	A	Sidechain
85	AA	1985	C	Sidechain
85	AA	1986	G	Sidechain
85	AA	1987	G	Sidechain
85	AA	1989	A	Sidechain
85	AA	199	U	Sidechain
85	AA	1991	C	Sidechain
85	AA	1992	A	Sidechain
85	AA	1993	C	Sidechain
85	AA	1995	U	Sidechain
85	AA	1996	A	Sidechain
85	AA	1997	G	Sidechain
85	AA	1999	C	Sidechain
85	AA	2	A	Sidechain
85	AA	20	G	Sidechain
85	AA	200	U	Sidechain
85	AA	2000	C	Sidechain
85	AA	2001	C	Sidechain
85	AA	2002	A	Sidechain
85	AA	2003	C	Sidechain
85	AA	2004	U	Sidechain
85	AA	2005	U	Sidechain
85	AA	2006	G	Sidechain
85	AA	2007	G	Sidechain
85	AA	2008	G	Sidechain
85	AA	201	U	Sidechain
85	AA	2010	C	Sidechain
85	AA	2011	C	Sidechain
85	AA	2013	A	Sidechain
85	AA	2014	G	Sidechain
85	AA	2015	U	Sidechain
85	AA	2017	U	Sidechain
85	AA	2018	U	Sidechain
85	AA	2019	G	Sidechain
85	AA	202	U	Sidechain
85	AA	2020	C	Sidechain
85	AA	2021	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	2022	A	Sidechain
85	AA	2023	U	Sidechain
85	AA	2024	U	Sidechain
85	AA	2026	U	Sidechain
85	AA	2029	G	Sidechain
85	AA	203	C	Sidechain
85	AA	2030	U	Sidechain
85	AA	2031	C	Sidechain
85	AA	2032	G	Sidechain
85	AA	2034	G	Sidechain
85	AA	2035	C	Sidechain
85	AA	2036	A	Sidechain
85	AA	2038	C	Sidechain
85	AA	2039	G	Sidechain
85	AA	204	U	Sidechain
85	AA	2041	G	Sidechain
85	AA	2042	G	Sidechain
85	AA	2044	A	Sidechain
85	AA	2045	U	Sidechain
85	AA	2046	G	Sidechain
85	AA	2047	U	Sidechain
85	AA	2048	C	Sidechain
85	AA	205	A	Sidechain
85	AA	2052	U	Sidechain
85	AA	2053	A	Sidechain
85	AA	2054	G	Sidechain
85	AA	2055	G	Sidechain
85	AA	2057	G	Sidechain
85	AA	2058	C	Sidechain
85	AA	2059	A	Sidechain
85	AA	206	U	Sidechain
85	AA	2060	G	Sidechain
85	AA	2061	C	Sidechain
85	AA	2062	U	Sidechain
85	AA	2063	C	Sidechain
85	AA	2067	A	Sidechain
85	AA	207	G	Sidechain
85	AA	2073	U	Sidechain
85	AA	2074	G	Sidechain
85	AA	2075	C	Sidechain
85	AA	2077	G	Sidechain
85	AA	2078	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	2079	U	Sidechain
85	AA	2080	U	Sidechain
85	AA	2082	C	Sidechain
85	AA	2084	U	Sidechain
85	AA	2085	C	Sidechain
85	AA	2086	C	Sidechain
85	AA	2087	C	Sidechain
85	AA	2088	U	Sidechain
85	AA	2089	G	Sidechain
85	AA	209	C	Sidechain
85	AA	2090	C	Sidechain
85	AA	2094	U	Sidechain
85	AA	2095	U	Sidechain
85	AA	2096	G	Sidechain
85	AA	2098	A	Sidechain
85	AA	2099	C	Sidechain
85	AA	21	U	Sidechain
85	AA	210	G	Sidechain
85	AA	2100	A	Sidechain
85	AA	2102	A	Sidechain
85	AA	2104	C	Sidechain
85	AA	2106	C	Sidechain
85	AA	2107	C	Sidechain
85	AA	2109	G	Sidechain
85	AA	211	C	Sidechain
85	AA	2110	U	Sidechain
85	AA	2112	G	Sidechain
85	AA	2114	U	Sidechain
85	AA	2116	U	Sidechain
85	AA	2117	U	Sidechain
85	AA	2118	U	Sidechain
85	AA	2119	C	Sidechain
85	AA	2120	C	Sidechain
85	AA	2121	G	Sidechain
85	AA	2122	A	Sidechain
85	AA	2123	U	Sidechain
85	AA	2124	G	Sidechain
85	AA	2125	A	Sidechain
85	AA	2126	U	Sidechain
85	AA	2128	G	Sidechain
85	AA	2129	U	Sidechain
85	AA	2130	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	2135	A	Sidechain
85	AA	2136	C	Sidechain
85	AA	2138	G	Sidechain
85	AA	2140	U	Sidechain
85	AA	2141	G	Sidechain
85	AA	2142	A	Sidechain
85	AA	2143	U	Sidechain
85	AA	2144	C	Sidechain
85	AA	2145	G	Sidechain
85	AA	2146	G	Sidechain
85	AA	2148	C	Sidechain
85	AA	2149	C	Sidechain
85	AA	215	U	Sidechain
85	AA	2150	G	Sidechain
85	AA	2152	C	Sidechain
85	AA	2153	G	Sidechain
85	AA	2155	U	Sidechain
85	AA	2156	C	Sidechain
85	AA	2158	U	Sidechain
85	AA	2159	C	Sidechain
85	AA	2161	C	Sidechain
85	AA	2162	G	Sidechain
85	AA	2163	G	Sidechain
85	AA	2164	G	Sidechain
85	AA	2166	G	Sidechain
85	AA	2167	A	Sidechain
85	AA	2169	C	Sidechain
85	AA	2170	G	Sidechain
85	AA	2171	A	Sidechain
85	AA	2172	A	Sidechain
85	AA	2174	G	Sidechain
85	AA	2175	U	Sidechain
85	AA	2176	U	Sidechain
85	AA	2178	A	Sidechain
85	AA	2179	C	Sidechain
85	AA	218	U	Sidechain
85	AA	2180	C	Sidechain
85	AA	2181	G	Sidechain
85	AA	2182	A	Sidechain
85	AA	2183	U	Sidechain
85	AA	2185	U	Sidechain
85	AA	2186	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	2189	U	Sidechain
85	AA	219	G	Sidechain
85	AA	2190	U	Sidechain
85	AA	2192	A	Sidechain
85	AA	2193	A	Sidechain
85	AA	2194	U	Sidechain
85	AA	2195	A	Sidechain
85	AA	2196	G	Sidechain
85	AA	2199	G	Sidechain
85	AA	22	A	Sidechain
85	AA	2201	A	Sidechain
85	AA	2202	G	Sidechain
85	AA	2206	A	Sidechain
85	AA	2207	A	Sidechain
85	AA	2208	G	Sidechain
85	AA	2209	U	Sidechain
85	AA	2211	G	Sidechain
85	AA	2213	A	Sidechain
85	AA	2214	A	Sidechain
85	AA	2216	A	Sidechain
85	AA	2217	A	Sidechain
85	AA	2218	G	Sidechain
85	AA	2219	G	Sidechain
85	AA	2222	G	Sidechain
85	AA	2223	C	Sidechain
85	AA	2224	U	Sidechain
85	AA	2226	U	Sidechain
85	AA	2227	A	Sidechain
85	AA	2228	G	Sidechain
85	AA	2229	G	Sidechain
85	AA	2230	U	Sidechain
85	AA	2231	G	Sidechain
85	AA	2232	A	Sidechain
85	AA	2233	A	Sidechain
85	AA	2234	C	Sidechain
85	AA	2236	U	Sidechain
85	AA	2237	G	Sidechain
85	AA	2238	C	Sidechain
85	AA	2240	G	Sidechain
85	AA	2241	C	Sidechain
85	AA	2245	A	Sidechain
85	AA	2246	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	2248	A	Sidechain
85	AA	2250	U	Sidechain
85	AA	228	C	Sidechain
85	AA	229	U	Sidechain
85	AA	23	G	Sidechain
85	AA	233	C	Sidechain
85	AA	234	G	Sidechain
85	AA	236	G	Sidechain
85	AA	237	G	Sidechain
85	AA	239	G	Sidechain
85	AA	24	U	Sidechain
85	AA	240	A	Sidechain
85	AA	242	G	Sidechain
85	AA	243	A	Sidechain
85	AA	244	G	Sidechain
85	AA	245	A	Sidechain
85	AA	246	C	Sidechain
85	AA	248	U	Sidechain
85	AA	249	C	Sidechain
85	AA	250	C	Sidechain
85	AA	251	A	Sidechain
85	AA	252	G	Sidechain
85	AA	253	C	Sidechain
85	AA	254	G	Sidechain
85	AA	255	A	Sidechain
85	AA	259	A	Sidechain
85	AA	26	A	Sidechain
85	AA	260	A	Sidechain
85	AA	261	U	Sidechain
85	AA	264	A	Sidechain
85	AA	265	A	Sidechain
85	AA	266	U	Sidechain
85	AA	267	U	Sidechain
85	AA	268	A	Sidechain
85	AA	269	G	Sidechain
85	AA	27	U	Sidechain
85	AA	270	A	Sidechain
85	AA	271	A	Sidechain
85	AA	275	A	Sidechain
85	AA	276	C	Sidechain
85	AA	277	G	Sidechain
85	AA	281	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	282	C	Sidechain
85	AA	283	A	Sidechain
85	AA	286	C	Sidechain
85	AA	287	G	Sidechain
85	AA	288	G	Sidechain
85	AA	289	G	Sidechain
85	AA	290	G	Sidechain
85	AA	293	A	Sidechain
85	AA	294	G	Sidechain
85	AA	296	A	Sidechain
85	AA	297	A	Sidechain
85	AA	298	C	Sidechain
85	AA	299	A	Sidechain
85	AA	30	G	Sidechain
85	AA	300	C	Sidechain
85	AA	301	U	Sidechain
85	AA	302	C	Sidechain
85	AA	303	A	Sidechain
85	AA	304	G	Sidechain
85	AA	306	C	Sidechain
85	AA	307	G	Sidechain
85	AA	308	U	Sidechain
85	AA	309	G	Sidechain
85	AA	311	U	Sidechain
85	AA	312	G	Sidechain
85	AA	313	A	Sidechain
85	AA	314	C	Sidechain
85	AA	316	C	Sidechain
85	AA	318	A	Sidechain
85	AA	319	U	Sidechain
85	AA	32	U	Sidechain
85	AA	321	C	Sidechain
85	AA	322	A	Sidechain
85	AA	323	U	Sidechain
85	AA	325	C	Sidechain
85	AA	326	C	Sidechain
85	AA	327	G	Sidechain
85	AA	328	U	Sidechain
85	AA	329	G	Sidechain
85	AA	33	U	Sidechain
85	AA	330	C	Sidechain
85	AA	331	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	332	A	Sidechain
85	AA	333	A	Sidechain
85	AA	334	A	Sidechain
85	AA	335	G	Sidechain
85	AA	336	C	Sidechain
85	AA	337	C	Sidechain
85	AA	339	A	Sidechain
85	AA	34	G	Sidechain
85	AA	342	C	Sidechain
85	AA	343	U	Sidechain
85	AA	345	U	Sidechain
85	AA	346	U	Sidechain
85	AA	35	U	Sidechain
85	AA	351	C	Sidechain
85	AA	353	G	Sidechain
85	AA	354	C	Sidechain
85	AA	355	G	Sidechain
85	AA	356	U	Sidechain
85	AA	358	U	Sidechain
85	AA	359	A	Sidechain
85	AA	36	U	Sidechain
85	AA	362	G	Sidechain
85	AA	363	A	Sidechain
85	AA	364	C	Sidechain
85	AA	366	A	Sidechain
85	AA	368	C	Sidechain
85	AA	369	A	Sidechain
85	AA	37	U	Sidechain
85	AA	371	C	Sidechain
85	AA	375	C	Sidechain
85	AA	376	C	Sidechain
85	AA	377	U	Sidechain
85	AA	378	A	Sidechain
85	AA	379	U	Sidechain
85	AA	380	C	Sidechain
85	AA	381	A	Sidechain
85	AA	383	C	Sidechain
85	AA	384	C	Sidechain
85	AA	387	U	Sidechain
85	AA	388	G	Sidechain
85	AA	389	A	Sidechain
85	AA	39	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	390	U	Sidechain
85	AA	391	G	Sidechain
85	AA	392	G	Sidechain
85	AA	393	C	Sidechain
85	AA	396	U	Sidechain
85	AA	397	G	Sidechain
85	AA	398	U	Sidechain
85	AA	399	A	Sidechain
85	AA	4	C	Sidechain
85	AA	400	G	Sidechain
85	AA	401	U	Sidechain
85	AA	402	G	Sidechain
85	AA	403	G	Sidechain
85	AA	404	A	Sidechain
85	AA	405	C	Sidechain
85	AA	407	G	Sidechain
85	AA	409	C	Sidechain
85	AA	41	G	Sidechain
85	AA	410	A	Sidechain
85	AA	411	U	Sidechain
85	AA	412	G	Sidechain
85	AA	413	G	Sidechain
85	AA	414	C	Sidechain
85	AA	416	U	Sidechain
85	AA	417	U	Sidechain
85	AA	418	G	Sidechain
85	AA	419	A	Sidechain
85	AA	42	G	Sidechain
85	AA	422	G	Sidechain
85	AA	424	A	Sidechain
85	AA	425	G	Sidechain
85	AA	426	C	Sidechain
85	AA	427	G	Sidechain
85	AA	428	G	Sidechain
85	AA	429	G	Sidechain
85	AA	430	G	Sidechain
85	AA	431	G	Sidechain
85	AA	432	A	Sidechain
85	AA	433	U	Sidechain
85	AA	436	G	Sidechain
85	AA	437	G	Sidechain
85	AA	438	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	439	U	Sidechain
85	AA	44	C	Sidechain
85	AA	440	U	Sidechain
85	AA	441	C	Sidechain
85	AA	443	A	Sidechain
85	AA	444	U	Sidechain
85	AA	445	U	Sidechain
85	AA	447	C	Sidechain
85	AA	448	G	Sidechain
85	AA	45	U	Sidechain
85	AA	450	A	Sidechain
85	AA	451	G	Sidechain
85	AA	453	G	Sidechain
85	AA	454	G	Sidechain
85	AA	455	G	Sidechain
85	AA	457	G	Sidechain
85	AA	46	U	Sidechain
85	AA	460	U	Sidechain
85	AA	461	G	Sidechain
85	AA	462	A	Sidechain
85	AA	463	G	Sidechain
85	AA	464	A	Sidechain
85	AA	465	A	Sidechain
85	AA	466	A	Sidechain
85	AA	467	U	Sidechain
85	AA	469	G	Sidechain
85	AA	470	C	Sidechain
85	AA	471	U	Sidechain
85	AA	472	A	Sidechain
85	AA	475	A	Sidechain
85	AA	476	C	Sidechain
85	AA	478	U	Sidechain
85	AA	48	G	Sidechain
85	AA	480	U	Sidechain
85	AA	481	A	Sidechain
85	AA	482	C	Sidechain
85	AA	483	G	Sidechain
85	AA	485	A	Sidechain
85	AA	486	G	Sidechain
85	AA	487	G	Sidechain
85	AA	488	G	Sidechain
85	AA	489	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	490	A	Sidechain
85	AA	491	G	Sidechain
85	AA	492	C	Sidechain
85	AA	494	G	Sidechain
85	AA	495	G	Sidechain
85	AA	496	C	Sidechain
85	AA	498	C	Sidechain
85	AA	499	G	Sidechain
85	AA	50	C	Sidechain
85	AA	500	C	Sidechain
85	AA	501	A	Sidechain
85	AA	502	A	Sidechain
85	AA	503	A	Sidechain
85	AA	504	U	Sidechain
85	AA	506	G	Sidechain
85	AA	507	C	Sidechain
85	AA	508	C	Sidechain
85	AA	509	C	Sidechain
85	AA	51	A	Sidechain
85	AA	510	A	Sidechain
85	AA	511	A	Sidechain
85	AA	512	U	Sidechain
85	AA	513	G	Sidechain
85	AA	514	U	Sidechain
85	AA	515	C	Sidechain
85	AA	516	G	Sidechain
85	AA	517	A	Sidechain
85	AA	52	U	Sidechain
85	AA	520	A	Sidechain
85	AA	521	A	Sidechain
85	AA	523	U	Sidechain
85	AA	526	G	Sidechain
85	AA	527	A	Sidechain
85	AA	528	U	Sidechain
85	AA	529	G	Sidechain
85	AA	53	G	Sidechain
85	AA	530	A	Sidechain
85	AA	531	G	Sidechain
85	AA	532	G	Sidechain
85	AA	533	C	Sidechain
85	AA	534	A	Sidechain
85	AA	535	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	536	C	Sidechain
85	AA	537	G	Sidechain
85	AA	538	A	Sidechain
85	AA	539	A	Sidechain
85	AA	54	C	Sidechain
85	AA	540	A	Sidechain
85	AA	541	A	Sidechain
85	AA	542	G	Sidechain
85	AA	544	A	Sidechain
85	AA	545	A	Sidechain
85	AA	546	U	Sidechain
85	AA	547	A	Sidechain
85	AA	548	G	Sidechain
85	AA	550	G	Sidechain
85	AA	552	C	Sidechain
85	AA	555	C	Sidechain
85	AA	556	C	Sidechain
85	AA	557	G	Sidechain
85	AA	559	G	Sidechain
85	AA	561	C	Sidechain
85	AA	563	U	Sidechain
85	AA	564	A	Sidechain
85	AA	565	G	Sidechain
85	AA	566	U	Sidechain
85	AA	567	G	Sidechain
85	AA	569	A	Sidechain
85	AA	57	G	Sidechain
85	AA	570	U	Sidechain
85	AA	571	G	Sidechain
85	AA	572	G	Sidechain
85	AA	573	U	Sidechain
85	AA	575	G	Sidechain
85	AA	576	U	Sidechain
85	AA	577	U	Sidechain
85	AA	578	U	Sidechain
85	AA	579	U	Sidechain
85	AA	582	A	Sidechain
85	AA	584	G	Sidechain
85	AA	585	G	Sidechain
85	AA	586	G	Sidechain
85	AA	587	G	Sidechain
85	AA	588	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	589	A	Sidechain
85	AA	59	C	Sidechain
85	AA	590	U	Sidechain
85	AA	591	A	Sidechain
85	AA	592	C	Sidechain
85	AA	593	U	Sidechain
85	AA	595	A	Sidechain
85	AA	596	A	Sidechain
85	AA	597	A	Sidechain
85	AA	6	G	Sidechain
85	AA	60	U	Sidechain
85	AA	600	C	Sidechain
85	AA	601	A	Sidechain
85	AA	602	U	Sidechain
85	AA	603	C	Sidechain
85	AA	604	C	Sidechain
85	AA	605	A	Sidechain
85	AA	609	U	Sidechain
85	AA	61	C	Sidechain
85	AA	610	C	Sidechain
85	AA	611	G	Sidechain
85	AA	613	G	Sidechain
85	AA	614	U	Sidechain
85	AA	615	A	Sidechain
85	AA	616	A	Sidechain
85	AA	617	C	Sidechain
85	AA	618	A	Sidechain
85	AA	62	A	Sidechain
85	AA	620	U	Sidechain
85	AA	621	U	Sidechain
85	AA	623	G	Sidechain
85	AA	624	A	Sidechain
85	AA	625	G	Sidechain
85	AA	626	G	Sidechain
85	AA	628	C	Sidechain
85	AA	629	A	Sidechain
85	AA	63	G	Sidechain
85	AA	630	A	Sidechain
85	AA	631	G	Sidechain
85	AA	634	U	Sidechain
85	AA	635	G	Sidechain
85	AA	636	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	637	U	Sidechain
85	AA	638	G	Sidechain
85	AA	64	A	Sidechain
85	AA	640	C	Sidechain
85	AA	641	A	Sidechain
85	AA	642	G	Sidechain
85	AA	646	C	Sidechain
85	AA	647	C	Sidechain
85	AA	648	G	Sidechain
85	AA	65	A	Sidechain
85	AA	650	G	Sidechain
85	AA	651	G	Sidechain
85	AA	652	U	Sidechain
85	AA	654	A	Sidechain
85	AA	655	U	Sidechain
85	AA	656	U	Sidechain
85	AA	657	C	Sidechain
85	AA	658	C	Sidechain
85	AA	659	A	Sidechain
85	AA	66	U	Sidechain
85	AA	661	C	Sidechain
85	AA	662	U	Sidechain
85	AA	664	C	Sidechain
85	AA	665	A	Sidechain
85	AA	666	A	Sidechain
85	AA	667	A	Sidechain
85	AA	668	A	Sidechain
85	AA	669	G	Sidechain
85	AA	67	C	Sidechain
85	AA	672	U	Sidechain
85	AA	673	A	Sidechain
85	AA	674	U	Sidechain
85	AA	675	A	Sidechain
85	AA	676	U	Sidechain
85	AA	677	U	Sidechain
85	AA	679	A	Sidechain
85	AA	68	A	Sidechain
85	AA	680	U	Sidechain
85	AA	681	G	Sidechain
85	AA	682	C	Sidechain
85	AA	683	U	Sidechain
85	AA	684	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	685	U	Sidechain
85	AA	686	U	Sidechain
85	AA	687	G	Sidechain
85	AA	688	C	Sidechain
85	AA	689	U	Sidechain
85	AA	69	C	Sidechain
85	AA	691	U	Sidechain
85	AA	692	U	Sidechain
85	AA	693	A	Sidechain
85	AA	694	A	Sidechain
85	AA	695	A	Sidechain
85	AA	696	G	Sidechain
85	AA	697	G	Sidechain
85	AA	698	G	Sidechain
85	AA	699	U	Sidechain
85	AA	7	G	Sidechain
85	AA	70	U	Sidechain
85	AA	700	U	Sidechain
85	AA	702	G	Sidechain
85	AA	703	U	Sidechain
85	AA	704	A	Sidechain
85	AA	706	U	Sidechain
85	AA	707	U	Sidechain
85	AA	708	G	Sidechain
85	AA	710	A	Sidechain
85	AA	711	C	Sidechain
85	AA	712	U	Sidechain
85	AA	713	G	Sidechain
85	AA	714	U	Sidechain
85	AA	72	C	Sidechain
85	AA	721	C	Sidechain
85	AA	722	G	Sidechain
85	AA	723	U	Sidechain
85	AA	725	G	Sidechain
85	AA	726	U	Sidechain
85	AA	727	U	Sidechain
85	AA	728	U	Sidechain
85	AA	730	G	Sidechain
85	AA	731	U	Sidechain
85	AA	732	G	Sidechain
85	AA	733	C	Sidechain
85	AA	735	G	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	736	U	Sidechain
85	AA	737	G	Sidechain
85	AA	738	C	Sidechain
85	AA	739	C	Sidechain
85	AA	740	A	Sidechain
85	AA	741	G	Sidechain
85	AA	742	U	Sidechain
85	AA	743	C	Sidechain
85	AA	744	C	Sidechain
85	AA	747	U	Sidechain
85	AA	748	C	Sidechain
85	AA	750	A	Sidechain
85	AA	751	C	Sidechain
85	AA	752	C	Sidechain
85	AA	754	C	Sidechain
85	AA	755	G	Sidechain
85	AA	756	G	Sidechain
85	AA	757	A	Sidechain
85	AA	758	C	Sidechain
85	AA	759	G	Sidechain
85	AA	76	G	Sidechain
85	AA	760	U	Sidechain
85	AA	761	G	Sidechain
85	AA	762	U	Sidechain
85	AA	764	U	Sidechain
85	AA	765	U	Sidechain
85	AA	766	G	Sidechain
85	AA	767	A	Sidechain
85	AA	768	C	Sidechain
85	AA	769	C	Sidechain
85	AA	77	C	Sidechain
85	AA	770	C	Sidechain
85	AA	771	A	Sidechain
85	AA	772	C	Sidechain
85	AA	773	G	Sidechain
85	AA	774	C	Sidechain
85	AA	775	C	Sidechain
85	AA	776	C	Sidechain
85	AA	777	U	Sidechain
85	AA	778	C	Sidechain
85	AA	779	G	Sidechain
85	AA	78	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	780	U	Sidechain
85	AA	781	G	Sidechain
85	AA	782	G	Sidechain
85	AA	785	C	Sidechain
85	AA	787	U	Sidechain
85	AA	788	G	Sidechain
85	AA	789	A	Sidechain
85	AA	79	G	Sidechain
85	AA	790	A	Sidechain
85	AA	792	A	Sidechain
85	AA	793	C	Sidechain
85	AA	794	A	Sidechain
85	AA	795	C	Sidechain
85	AA	796	U	Sidechain
85	AA	797	C	Sidechain
85	AA	799	G	Sidechain
85	AA	8	U	Sidechain
85	AA	800	A	Sidechain
85	AA	801	U	Sidechain
85	AA	802	A	Sidechain
85	AA	803	C	Sidechain
85	AA	804	A	Sidechain
85	AA	805	A	Sidechain
85	AA	806	G	Sidechain
85	AA	809	A	Sidechain
85	AA	812	C	Sidechain
85	AA	814	G	Sidechain
85	AA	815	G	Sidechain
85	AA	816	A	Sidechain
85	AA	817	G	Sidechain
85	AA	818	C	Sidechain
85	AA	819	G	Sidechain
85	AA	82	A	Sidechain
85	AA	820	G	Sidechain
85	AA	821	U	Sidechain
85	AA	822	U	Sidechain
85	AA	825	U	Sidechain
85	AA	827	C	Sidechain
85	AA	828	U	Sidechain
85	AA	83	U	Sidechain
85	AA	830	A	Sidechain
85	AA	832	U	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	833	U	Sidechain
85	AA	834	U	Sidechain
85	AA	836	A	Sidechain
85	AA	837	C	Sidechain
85	AA	838	G	Sidechain
85	AA	839	C	Sidechain
85	AA	84	C	Sidechain
85	AA	841	U	Sidechain
85	AA	842	G	Sidechain
85	AA	845	A	Sidechain
85	AA	846	U	Sidechain
85	AA	847	G	Sidechain
85	AA	848	C	Sidechain
85	AA	85	U	Sidechain
85	AA	850	U	Sidechain
85	AA	851	G	Sidechain
85	AA	852	C	Sidechain
85	AA	853	G	Sidechain
85	AA	854	A	Sidechain
85	AA	855	G	Sidechain
85	AA	856	G	Sidechain
85	AA	857	G	Sidechain
85	AA	858	G	Sidechain
85	AA	859	G	Sidechain
85	AA	86	G	Sidechain
85	AA	860	C	Sidechain
85	AA	861	G	Sidechain
85	AA	862	U	Sidechain
85	AA	863	C	Sidechain
85	AA	864	C	Sidechain
85	AA	866	U	Sidechain
85	AA	867	G	Sidechain
85	AA	868	A	Sidechain
85	AA	869	A	Sidechain
85	AA	87	C	Sidechain
85	AA	871	U	Sidechain
85	AA	872	U	Sidechain
85	AA	873	U	Sidechain
85	AA	875	C	Sidechain
85	AA	878	U	Sidechain
85	AA	879	G	Sidechain
85	AA	880	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	881	C	Sidechain
85	AA	882	C	Sidechain
85	AA	883	A	Sidechain
85	AA	884	A	Sidechain
85	AA	885	A	Sidechain
85	AA	887	A	Sidechain
85	AA	888	A	Sidechain
85	AA	889	G	Sidechain
85	AA	89	C	Sidechain
85	AA	890	U	Sidechain
85	AA	891	G	Sidechain
85	AA	892	C	Sidechain
85	AA	893	G	Sidechain
85	AA	894	A	Sidechain
85	AA	896	C	Sidechain
85	AA	897	A	Sidechain
85	AA	898	A	Sidechain
85	AA	899	A	Sidechain
85	AA	9	U	Sidechain
85	AA	90	A	Sidechain
85	AA	900	G	Sidechain
85	AA	901	C	Sidechain
85	AA	902	A	Sidechain
85	AA	903	G	Sidechain
85	AA	906	U	Sidechain
85	AA	908	C	Sidechain
85	AA	91	U	Sidechain
85	AA	910	G	Sidechain
85	AA	911	A	Sidechain
85	AA	912	C	Sidechain
85	AA	913	U	Sidechain
85	AA	914	U	Sidechain
85	AA	916	A	Sidechain
85	AA	917	A	Sidechain
85	AA	918	U	Sidechain
85	AA	92	G	Sidechain
85	AA	920	A	Sidechain
85	AA	921	C	Sidechain
85	AA	922	A	Sidechain
85	AA	924	A	Sidechain
85	AA	925	G	Sidechain
85	AA	926	C	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	929	G	Sidechain
85	AA	93	G	Sidechain
85	AA	930	G	Sidechain
85	AA	931	G	Sidechain
85	AA	932	A	Sidechain
85	AA	933	U	Sidechain
85	AA	934	A	Sidechain
85	AA	935	A	Sidechain
85	AA	936	C	Sidechain
85	AA	937	G	Sidechain
85	AA	939	A	Sidechain
85	AA	94	C	Sidechain
85	AA	940	G	Sidechain
85	AA	941	C	Sidechain
85	AA	943	U	Sidechain
85	AA	944	C	Sidechain
85	AA	945	A	Sidechain
85	AA	95	U	Sidechain
85	AA	957	A	Sidechain
85	AA	958	C	Sidechain
85	AA	959	C	Sidechain
85	AA	96	C	Sidechain
85	AA	960	G	Sidechain
85	AA	961	U	Sidechain
85	AA	963	U	Sidechain
85	AA	964	C	Sidechain
85	AA	966	G	Sidechain
85	AA	968	U	Sidechain
85	AA	969	U	Sidechain
85	AA	97	A	Sidechain
85	AA	970	U	Sidechain
85	AA	971	U	Sidechain
85	AA	972	G	Sidechain
85	AA	973	U	Sidechain
85	AA	974	U	Sidechain
85	AA	976	G	Sidechain
85	AA	977	U	Sidechain
85	AA	978	U	Sidechain
85	AA	979	U	Sidechain
85	AA	98	U	Sidechain
85	AA	980	U	Sidechain
85	AA	981	A	Sidechain

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Mol	Chain	Res	Type	Group
85	AA	982	G	Sidechain
85	AA	983	A	Sidechain
85	AA	985	G	Sidechain
85	AA	986	U	Sidechain
85	AA	987	C	Sidechain
85	AA	989	U	Sidechain
85	AA	99	U	Sidechain
85	AA	991	G	Sidechain
85	AA	993	G	Sidechain
85	AA	994	A	Sidechain
85	AA	995	G	Sidechain
85	AA	997	U	Sidechain
85	AA	998	U	Sidechain
86	AB	1	G	Sidechain
86	AB	11	C	Sidechain
86	AB	12	U	Sidechain
86	AB	14	A	Sidechain
86	AB	18	G	Sidechain
86	AB	2	C	Sidechain
86	AB	20	U	Sidechain
86	AB	21	A	Sidechain
86	AB	22	G	Sidechain
86	AB	23	A	Sidechain
86	AB	24	G	Sidechain
86	AB	25	C	Sidechain
86	AB	26	A	Sidechain
86	AB	27	G	Sidechain
86	AB	28	G	Sidechain
86	AB	32	U	Sidechain
86	AB	37	A	Sidechain
86	AB	39	U	Sidechain
86	AB	40	C	Sidechain
86	AB	43	C	Sidechain
86	AB	45	U	Sidechain
86	AB	46	G	Sidechain
86	AB	49	C	Sidechain
86	AB	5	G	Sidechain
86	AB	50	U	Sidechain
86	AB	51	U	Sidechain
86	AB	54	U	Sidechain
86	AB	56	C	Sidechain
86	AB	57	G	Sidechain

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Mol	Chain	Res	Type	Group
86	AB	58	A	Sidechain
86	AB	6	G	Sidechain
86	AB	61	C	Sidechain
86	AB	62	C	Sidechain
86	AB	63	G	Sidechain
86	AB	65	G	Sidechain
86	AB	66	U	Sidechain
86	AB	67	C	Sidechain
86	AB	68	C	Sidechain
86	AB	69	G	Sidechain
86	AB	7	A	Sidechain
86	AB	70	G	Sidechain
86	AB	72	C	Sidechain
86	AB	73	A	Sidechain
86	AB	8	U	Sidechain
86	AB	9	A	Sidechain
11	AC	118	PHE	Sidechain
11	AC	139	PHE	Sidechain
11	AC	144	PHE	Sidechain
11	AC	156	ARG	Mainchain,Sidechain
11	AC	193	VAL	Mainchain
11	AC	202	ARG	Sidechain
11	AC	204	ARG	Sidechain
11	AC	211	TYR	Sidechain
11	AC	216	ARG	Sidechain
11	AC	222	ARG	Peptide
11	AC	225	ILE	Mainchain
11	AC	228	SER	Mainchain
11	AC	229	VAL	Peptide
11	AC	238	PHE	Mainchain
11	AC	240	TYR	Sidechain
11	AC	241	ARG	Sidechain
11	AC	73	TYR	Sidechain
11	AC	74	ILE	Peptide
11	AC	77	ARG	Mainchain
11	AC	99	ARG	Sidechain
12	AD	14	TYR	Sidechain
12	AD	18	PHE	Sidechain
12	AD	3	THR	Mainchain
12	AD	4	TYR	Sidechain
12	AD	54	ARG	Sidechain
12	AD	59	ARG	Sidechain

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Mol	Chain	Res	Type	Group
12	AD	66	PHE	Sidechain
12	AD	69	ARG	Sidechain
12	AD	88	TYR	Sidechain
12	AD	94	VAL	Peptide
13	AE	103	ARG	Sidechain
13	AE	104	ARG	Sidechain
13	AE	106	TYR	Sidechain
13	AE	113	TYR	Sidechain
13	AE	116	TYR	Sidechain
13	AE	119	ARG	Sidechain
13	AE	120	HIS	Sidechain
13	AE	121	ARG	Sidechain
13	AE	131	PHE	Sidechain
13	AE	145	ARG	Sidechain
13	AE	152	ARG	Sidechain
13	AE	153	TYR	Sidechain
13	AE	171	PHE	Sidechain
13	AE	24	TYR	Sidechain
13	AE	44	ARG	Sidechain
13	AE	45	SER	Peptide
13	AE	50	TYR	Sidechain
13	AE	58	PHE	Sidechain
13	AE	69	TYR	Sidechain
13	AE	72	ARG	Sidechain
13	AE	76	PHE	Sidechain
13	AE	83	ARG	Sidechain
13	AE	88	ARG	Sidechain
13	AE	97	ARG	Sidechain
13	AE	98	ARG	Sidechain
14	AF	108	ARG	Sidechain
14	AF	116	ARG	Sidechain
14	AF	132	ARG	Sidechain
14	AF	41	ARG	Sidechain
14	AF	55	ARG	Sidechain
14	AF	60	ARG	Sidechain
15	AG	104	ARG	Sidechain
15	AG	113	PHE	Sidechain
15	AG	121	ARG	Sidechain
15	AG	124	ARG	Sidechain
15	AG	129	TYR	Sidechain
15	AG	131	ARG	Sidechain
15	AG	134	GLN	Peptide,Mainchain

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Mol	Chain	Res	Type	Group
15	AG	18	TYR	Sidechain
15	AG	21	THR	Peptide
15	AG	3	ARG	Sidechain
15	AG	31	ARG	Sidechain
15	AG	55	ARG	Sidechain
15	AG	69	ARG	Sidechain
15	AG	77	HIS	Sidechain
15	AG	89	TYR	Sidechain
15	AG	94	ARG	Sidechain
16	AH	10	TYR	Sidechain
16	AH	114	ARG	Sidechain
16	AH	13	SER	Peptide
16	AH	21	TYR	Sidechain
16	AH	27	TYR	Sidechain
16	AH	36	HIS	Sidechain
16	AH	60	ASP	Mainchain
16	AH	65	TYR	Sidechain
17	AI	104	TYR	Sidechain
17	AI	122	TYR	Sidechain
17	AI	126	PHE	Sidechain
17	AI	130	TYR	Sidechain
17	AI	144	HIS	Peptide
17	AI	19	PHE	Sidechain
17	AI	25	ARG	Sidechain
17	AI	58	ARG	Sidechain
17	AI	66	ARG	Sidechain
17	AI	68	ARG	Sidechain
18	AJ	12	ARG	Sidechain
18	AJ	128	PHE	Sidechain
18	AJ	129	PHE	Sidechain
18	AJ	19	ARG	Sidechain
18	AJ	20	ARG	Sidechain
18	AJ	46	TYR	Sidechain
18	AJ	54	ASP	Peptide
18	AJ	56	HIS	Sidechain
18	AJ	57	ARG	Sidechain
18	AJ	68	ARG	Sidechain
18	AJ	78	ARG	Sidechain
19	AK	115	TYR	Sidechain
19	AK	126	ASP	Peptide,Mainchain
19	AK	128	ARG	Sidechain
19	AK	129	ARG	Mainchain,Sidechain

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Mol	Chain	Res	Type	Group
19	AK	137	ARG	Sidechain
19	AK	141	ARG	Sidechain
19	AK	148	TYR	Sidechain
19	AK	149	ARG	Sidechain
19	AK	33	CYS	Peptide
19	AK	36	ARG	Sidechain
19	AK	47	PRO	Mainchain
19	AK	65	TYR	Sidechain
19	AK	68	ARG	Sidechain
19	AK	70	ARG	Sidechain
19	AK	74	ARG	Sidechain
19	AK	78	SER	Peptide
19	AK	9	LEU	Peptide
19	AK	98	TYR	Sidechain
20	AL	29	TYR	Sidechain
20	AL	33	ARG	Sidechain
20	AL	53	TYR	Sidechain
20	AL	78	ARG	Sidechain
20	AL	84	TYR	Sidechain
20	AL	90	HIS	Sidechain
21	AM	107	ARG	Sidechain
21	AM	119	ARG	Peptide
21	AM	12	HIS	Peptide
21	AM	121	HIS	Sidechain
21	AM	125	ARG	Sidechain
21	AM	128	TYR	Sidechain
21	AM	131	ARG	Sidechain
21	AM	133	ARG	Peptide
21	AM	142	ARG	Sidechain
21	AM	25	ARG	Sidechain
21	AM	29	PHE	Sidechain
21	AM	38	GLY	Peptide
21	AM	39	ILE	Peptide
21	AM	42	ALA	Peptide
21	AM	80	ILE	Peptide
21	AM	89	ARG	Sidechain
21	AM	9	SER	Peptide
21	AM	90	ASP	Peptide
22	AO	104	SER	Mainchain
22	AO	107	GLU	Mainchain
22	AO	111	ARG	Sidechain
22	AO	118	HIS	Sidechain

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Mol	Chain	Res	Type	Group
22	AO	138	HIS	Sidechain
22	AO	139	ARG	Sidechain
22	AO	158	ARG	Sidechain
22	AO	163	GLU	Peptide
22	AO	27	HIS	Sidechain
22	AO	29	TRP	Mainchain
22	AO	58	HIS	Sidechain
22	AO	60	ARG	Sidechain
22	AO	70	TYR	Sidechain
22	AO	79	ARG	Sidechain
22	AO	84	ARG	Sidechain
22	AO	89	TYR	Sidechain
22	AO	95	ARG	Sidechain
23	AP	108	PHE	Sidechain
23	AP	120	HIS	Sidechain
23	AP	136	ARG	Sidechain
23	AP	153	TYR	Sidechain
23	AP	179	LEU	Mainchain
23	AP	184	ARG	Sidechain
23	AP	207	TYR	Sidechain
23	AP	214	THR	Mainchain
23	AP	215	ARG	Sidechain
23	AP	247	ARG	Sidechain
23	AP	253	HIS	Sidechain
23	AP	59	THR	Mainchain
23	AP	68	SER	Mainchain
23	AP	82	ARG	Sidechain
24	AQ	101	PHE	Peptide
24	AQ	41	ARG	Mainchain,Sidechain
24	AQ	48	ARG	Sidechain
24	AQ	66	PRO	Peptide
24	AQ	67	CYS	Peptide
24	AQ	77	TYR	Sidechain
25	AR	17	TYR	Sidechain
25	AR	20	ARG	Sidechain
25	AR	26	ASN	Peptide,Mainchain
25	AR	34	HIS	Sidechain
25	AR	45	ASP	Mainchain
25	AR	50	ILE	Mainchain
25	AR	62	TYR	Sidechain
25	AR	69	SER	Mainchain
25	AR	7	TYR	Sidechain

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Mol	Chain	Res	Type	Group
25	AR	71	HIS	Sidechain
25	AR	84	ARG	Sidechain
26	AS	119	ARG	Sidechain
26	AS	131	TYR	Sidechain
26	AS	17	ARG	Sidechain
26	AS	18	ARG	Sidechain
26	AS	21	ARG	Sidechain
26	AS	42	GLY	Peptide
26	AS	71	ARG	Sidechain
27	AT	111	ARG	Sidechain
27	AT	114	VAL	Peptide
27	AT	118	ARG	Sidechain
27	AT	120	ARG	Sidechain
27	AT	123	LYS	Peptide
27	AT	125	ARG	Sidechain
27	AT	13	ARG	Sidechain
27	AT	132	MET	Peptide
27	AT	17	PHE	Sidechain
27	AT	25	ARG	Sidechain
27	AT	34	HIS	Sidechain
27	AT	35	PRO	Peptide
27	AT	36	GLY	Peptide,Mainchain
27	AT	37	TRP	Peptide
27	AT	38	CYS	Peptide
27	AT	40	THR	Peptide
27	AT	41	VAL	Peptide
27	AT	54	TYR	Sidechain
27	AT	58	ASP	Peptide,Mainchain
27	AT	65	PHE	Peptide,Mainchain,Sidechain
27	AT	8	ALA	Peptide
27	AT	83	TYR	Sidechain
27	AT	97	ARG	Sidechain
28	AU	105	ARG	Sidechain
28	AU	48	PHE	Sidechain
28	AU	60	VAL	Peptide
28	AU	63	TYR	Sidechain
28	AU	88	HIS	Sidechain
28	AU	91	ARG	Sidechain
28	AU	99	SER	Peptide
29	AV	10	ARG	Sidechain
29	AV	100	TYR	Sidechain
29	AV	17	ARG	Sidechain

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Mol	Chain	Res	Type	Group
29	AV	19	ARG	Sidechain
29	AV	21	ARG	Sidechain
29	AV	30	ARG	Sidechain
29	AV	40	ARG	Sidechain
29	AV	44	ARG	Sidechain
29	AV	5	ARG	Sidechain
29	AV	66	PHE	Peptide,Sidechain
29	AV	72	TYR	Sidechain
29	AV	98	ARG	Sidechain
30	AW	10	TYR	Peptide,Mainchain
30	AW	14	ARG	Peptide,Mainchain,Sidechain
30	AW	17	ARG	Sidechain
30	AW	22	ARG	Sidechain
30	AW	23	ARG	Peptide
30	AW	24	ARG	Sidechain
30	AW	43	ASN	Peptide
30	AW	5	ASP	Peptide
30	AW	58	ASN	Mainchain
30	AW	80	PHE	Sidechain
31	AX	155	TYR	Sidechain
31	AX	166	PHE	Sidechain
31	AX	17	TYR	Sidechain
31	AX	177	ARG	Sidechain
31	AX	191	GLN	Peptide
31	AX	39	ARG	Sidechain
31	AX	55	ARG	Sidechain
31	AX	63	ARG	Sidechain
31	AX	78	TYR	Peptide
31	AX	8	ARG	Sidechain
32	AY	31	ARG	Sidechain
32	AY	33	ARG	Sidechain
32	AY	40	TYR	Sidechain
32	AY	44	TYR	Peptide,Sidechain
32	AY	46	ALA	Mainchain
32	AY	47	LYS	Mainchain
32	AY	55	LEU	Peptide
32	AY	56	ARG	Sidechain
32	AY	59	LYS	Peptide,Mainchain
32	AY	60	GLN	Peptide
32	AY	64	LYS	Peptide
33	AZ	49	ARG	Sidechain
33	AZ	60	ARG	Sidechain

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Mol	Chain	Res	Type	Group
33	AZ	78	ARG	Sidechain
34	BA	1	C	Sidechain
34	BA	10	G	Sidechain
34	BA	100	A	Sidechain
34	BA	1001	G	Sidechain
34	BA	1004	U	Sidechain
34	BA	1005	C	Sidechain
34	BA	1006	G	Sidechain
34	BA	1007	G	Sidechain
34	BA	1009	G	Sidechain
34	BA	101	G	Sidechain
34	BA	1011	G	Sidechain
34	BA	1013	A	Sidechain
34	BA	1014	A	Sidechain
34	BA	1015	G	Sidechain
34	BA	1018	U	Sidechain
34	BA	1019	C	Sidechain
34	BA	102	G	Sidechain
34	BA	1020	A	Sidechain
34	BA	1021	U	Sidechain
34	BA	1022	C	Sidechain
34	BA	1023	G	Sidechain
34	BA	1024	A	Sidechain
34	BA	1027	C	Sidechain
34	BA	1028	A	Sidechain
34	BA	1029	C	Sidechain
34	BA	103	G	Sidechain
34	BA	1030	C	Sidechain
34	BA	1032	A	Sidechain
34	BA	1033	G	Sidechain
34	BA	1035	A	Sidechain
34	BA	1036	G	Sidechain
34	BA	1037	C	Sidechain
34	BA	1038	U	Sidechain
34	BA	1039	G	Sidechain
34	BA	1040	G	Sidechain
34	BA	1041	U	Sidechain
34	BA	1042	U	Sidechain
34	BA	1044	A	Sidechain
34	BA	1045	C	Sidechain
34	BA	1046	G	Sidechain
34	BA	1049	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	105	U	Sidechain
34	BA	1050	A	Sidechain
34	BA	1051	A	Sidechain
34	BA	1052	G	Sidechain
34	BA	1053	U	Sidechain
34	BA	1054	U	Sidechain
34	BA	1055	U	Sidechain
34	BA	1056	C	Sidechain
34	BA	1057	C	Sidechain
34	BA	1058	C	Sidechain
34	BA	1059	U	Sidechain
34	BA	106	U	Sidechain
34	BA	1062	G	Sidechain
34	BA	1063	G	Sidechain
34	BA	1064	A	Sidechain
34	BA	1065	U	Sidechain
34	BA	1066	A	Sidechain
34	BA	1068	C	Sidechain
34	BA	1069	U	Sidechain
34	BA	107	C	Sidechain
34	BA	1070	G	Sidechain
34	BA	1071	G	Sidechain
34	BA	1072	U	Sidechain
34	BA	1073	G	Sidechain
34	BA	1075	U	Sidechain
34	BA	1077	G	Sidechain
34	BA	1078	U	Sidechain
34	BA	1079	C	Sidechain
34	BA	108	A	Sidechain
34	BA	1080	U	Sidechain
34	BA	1081	U	Sidechain
34	BA	1082	U	Sidechain
34	BA	1083	A	Sidechain
34	BA	1084	A	Sidechain
34	BA	1085	G	Sidechain
34	BA	1086	A	Sidechain
34	BA	1087	A	Sidechain
34	BA	1088	G	Sidechain
34	BA	1089	U	Sidechain
34	BA	109	A	Sidechain
34	BA	1091	U	Sidechain
34	BA	1092	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1093	G	Sidechain
34	BA	1094	U	Sidechain
34	BA	1095	G	Sidechain
34	BA	1096	C	Sidechain
34	BA	1097	G	Sidechain
34	BA	1098	G	Sidechain
34	BA	1099	U	Sidechain
34	BA	11	U	Sidechain
34	BA	110	C	Sidechain
34	BA	1105	A	Sidechain
34	BA	1106	A	Sidechain
34	BA	1107	A	Sidechain
34	BA	1108	U	Sidechain
34	BA	1109	G	Sidechain
34	BA	111	U	Sidechain
34	BA	1110	A	Sidechain
34	BA	1111	U	Sidechain
34	BA	1112	U	Sidechain
34	BA	1113	A	Sidechain
34	BA	1114	G	Sidechain
34	BA	1116	G	Sidechain
34	BA	1117	G	Sidechain
34	BA	1119	A	Sidechain
34	BA	112	C	Sidechain
34	BA	1120	U	Sidechain
34	BA	1122	G	Sidechain
34	BA	1127	U	Sidechain
34	BA	1128	C	Sidechain
34	BA	1129	U	Sidechain
34	BA	113	G	Sidechain
34	BA	1130	U	Sidechain
34	BA	1131	G	Sidechain
34	BA	1133	A	Sidechain
34	BA	1134	A	Sidechain
34	BA	1136	A	Sidechain
34	BA	1137	U	Sidechain
34	BA	1138	C	Sidechain
34	BA	1139	G	Sidechain
34	BA	114	U	Sidechain
34	BA	1140	A	Sidechain
34	BA	1142	C	Sidechain
34	BA	1143	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1144	A	Sidechain
34	BA	1145	U	Sidechain
34	BA	1146	U	Sidechain
34	BA	1148	U	Sidechain
34	BA	1149	C	Sidechain
34	BA	115	U	Sidechain
34	BA	1151	A	Sidechain
34	BA	1152	A	Sidechain
34	BA	1153	C	Sidechain
34	BA	1154	U	Sidechain
34	BA	1155	U	Sidechain
34	BA	1156	U	Sidechain
34	BA	1157	A	Sidechain
34	BA	1159	A	Sidechain
34	BA	1161	G	Sidechain
34	BA	1162	U	Sidechain
34	BA	1165	A	Sidechain
34	BA	1167	A	Sidechain
34	BA	1168	C	Sidechain
34	BA	1170	A	Sidechain
34	BA	1171	C	Sidechain
34	BA	1173	C	Sidechain
34	BA	1174	A	Sidechain
34	BA	1175	G	Sidechain
34	BA	1176	C	Sidechain
34	BA	1177	C	Sidechain
34	BA	1178	U	Sidechain
34	BA	1179	U	Sidechain
34	BA	118	C	Sidechain
34	BA	1180	A	Sidechain
34	BA	1182	U	Sidechain
34	BA	1183	U	Sidechain
34	BA	1184	A	Sidechain
34	BA	1186	U	Sidechain
34	BA	1187	U	Sidechain
34	BA	1188	U	Sidechain
34	BA	1189	A	Sidechain
34	BA	1190	A	Sidechain
34	BA	1192	A	Sidechain
34	BA	1193	A	Sidechain
34	BA	1194	G	Sidechain
34	BA	1195	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1196	C	Sidechain
34	BA	1197	U	Sidechain
34	BA	1198	U	Sidechain
34	BA	1199	U	Sidechain
34	BA	12	G	Sidechain
34	BA	120	A	Sidechain
34	BA	1203	G	Sidechain
34	BA	1204	U	Sidechain
34	BA	1205	A	Sidechain
34	BA	1206	C	Sidechain
34	BA	1208	U	Sidechain
34	BA	1209	A	Sidechain
34	BA	121	A	Sidechain
34	BA	1210	A	Sidechain
34	BA	1211	G	Sidechain
34	BA	1212	A	Sidechain
34	BA	1214	U	Sidechain
34	BA	1215	U	Sidechain
34	BA	1216	G	Sidechain
34	BA	1217	A	Sidechain
34	BA	1219	G	Sidechain
34	BA	122	U	Sidechain
34	BA	1220	C	Sidechain
34	BA	1222	C	Sidechain
34	BA	1224	A	Sidechain
34	BA	1225	A	Sidechain
34	BA	1226	G	Sidechain
34	BA	1227	U	Sidechain
34	BA	1229	G	Sidechain
34	BA	123	C	Sidechain
34	BA	1230	G	Sidechain
34	BA	1231	C	Sidechain
34	BA	1232	C	Sidechain
34	BA	1233	U	Sidechain
34	BA	1236	U	Sidechain
34	BA	1239	G	Sidechain
34	BA	124	G	Sidechain
34	BA	1240	G	Sidechain
34	BA	1241	U	Sidechain
34	BA	1242	A	Sidechain
34	BA	1243	A	Sidechain
34	BA	1244	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1245	C	Sidechain
34	BA	1246	G	Sidechain
34	BA	1247	G	Sidechain
34	BA	1248	A	Sidechain
34	BA	125	G	Sidechain
34	BA	1250	C	Sidechain
34	BA	1252	G	Sidechain
34	BA	1253	G	Sidechain
34	BA	1255	G	Sidechain
34	BA	1256	A	Sidechain
34	BA	1257	U	Sidechain
34	BA	1258	G	Sidechain
34	BA	1259	C	Sidechain
34	BA	126	G	Sidechain
34	BA	1260	G	Sidechain
34	BA	1261	G	Sidechain
34	BA	1263	A	Sidechain
34	BA	1264	U	Sidechain
34	BA	1265	G	Sidechain
34	BA	1267	A	Sidechain
34	BA	1271	C	Sidechain
34	BA	1272	U	Sidechain
34	BA	1273	U	Sidechain
34	BA	1274	A	Sidechain
34	BA	1276	G	Sidechain
34	BA	1279	U	Sidechain
34	BA	128	C	Sidechain
34	BA	1280	A	Sidechain
34	BA	1281	U	Sidechain
34	BA	1282	G	Sidechain
34	BA	1283	U	Sidechain
34	BA	1284	G	Sidechain
34	BA	1285	G	Sidechain
34	BA	1287	G	Sidechain
34	BA	1288	U	Sidechain
34	BA	1289	C	Sidechain
34	BA	129	U	Sidechain
34	BA	1290	A	Sidechain
34	BA	1291	A	Sidechain
34	BA	1293	A	Sidechain
34	BA	1294	C	Sidechain
34	BA	1295	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1296	U	Sidechain
34	BA	1297	G	Sidechain
34	BA	1298	U	Sidechain
34	BA	1299	G	Sidechain
34	BA	13	U	Sidechain
34	BA	130	U	Sidechain
34	BA	1300	G	Sidechain
34	BA	1302	C	Sidechain
34	BA	1303	U	Sidechain
34	BA	1305	A	Sidechain
34	BA	1306	U	Sidechain
34	BA	1307	U	Sidechain
34	BA	1310	C	Sidechain
34	BA	1311	G	Sidechain
34	BA	1312	A	Sidechain
34	BA	1314	A	Sidechain
34	BA	1315	C	Sidechain
34	BA	1316	G	Sidechain
34	BA	1317	U	Sidechain
34	BA	1318	G	Sidechain
34	BA	1319	A	Sidechain
34	BA	132	U	Sidechain
34	BA	1320	A	Sidechain
34	BA	1321	A	Sidechain
34	BA	1323	G	Sidechain
34	BA	1324	G	Sidechain
34	BA	1325	G	Sidechain
34	BA	1326	U	Sidechain
34	BA	1329	U	Sidechain
34	BA	133	A	Sidechain
34	BA	1330	G	Sidechain
34	BA	1331	G	Sidechain
34	BA	1332	U	Sidechain
34	BA	1333	G	Sidechain
34	BA	1334	G	Sidechain
34	BA	1335	A	Sidechain
34	BA	1336	U	Sidechain
34	BA	1339	G	Sidechain
34	BA	134	U	Sidechain
34	BA	1340	G	Sidechain
34	BA	1342	C	Sidechain
34	BA	1343	A	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1345	U	Sidechain
34	BA	1346	U	Sidechain
34	BA	1347	G	Sidechain
34	BA	1348	G	Sidechain
34	BA	1349	A	Sidechain
34	BA	135	G	Sidechain
34	BA	1352	G	Sidechain
34	BA	1353	U	Sidechain
34	BA	1354	G	Sidechain
34	BA	1355	G	Sidechain
34	BA	1357	C	Sidechain
34	BA	1359	U	Sidechain
34	BA	1362	A	Sidechain
34	BA	1363	A	Sidechain
34	BA	1366	C	Sidechain
34	BA	1370	A	Sidechain
34	BA	1371	U	Sidechain
34	BA	1372	C	Sidechain
34	BA	1373	C	Sidechain
34	BA	1374	G	Sidechain
34	BA	1377	A	Sidechain
34	BA	1378	A	Sidechain
34	BA	1379	G	Sidechain
34	BA	138	C	Sidechain
34	BA	1380	G	Sidechain
34	BA	1382	G	Sidechain
34	BA	1383	U	Sidechain
34	BA	1384	G	Sidechain
34	BA	1385	U	Sidechain
34	BA	1386	G	Sidechain
34	BA	1387	U	Sidechain
34	BA	139	U	Sidechain
34	BA	1390	C	Sidechain
34	BA	1392	A	Sidechain
34	BA	1395	C	Sidechain
34	BA	1396	A	Sidechain
34	BA	1397	C	Sidechain
34	BA	1399	A	Sidechain
34	BA	14	G	Sidechain
34	BA	140	C	Sidechain
34	BA	1402	C	Sidechain
34	BA	1403	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1406	U	Sidechain
34	BA	1407	C	Sidechain
34	BA	1409	A	Sidechain
34	BA	141	G	Sidechain
34	BA	1410	C	Sidechain
34	BA	1411	C	Sidechain
34	BA	1412	G	Sidechain
34	BA	1413	G	Sidechain
34	BA	1414	C	Sidechain
34	BA	1415	C	Sidechain
34	BA	1416	C	Sidechain
34	BA	1417	C	Sidechain
34	BA	1418	G	Sidechain
34	BA	1419	A	Sidechain
34	BA	142	A	Sidechain
34	BA	1420	A	Sidechain
34	BA	1421	A	Sidechain
34	BA	1422	A	Sidechain
34	BA	1423	U	Sidechain
34	BA	1424	G	Sidechain
34	BA	1425	G	Sidechain
34	BA	1426	A	Sidechain
34	BA	1427	U	Sidechain
34	BA	143	A	Sidechain
34	BA	1430	C	Sidechain
34	BA	1431	G	Sidechain
34	BA	1432	C	Sidechain
34	BA	1433	U	Sidechain
34	BA	1434	U	Sidechain
34	BA	1435	A	Sidechain
34	BA	1437	G	Sidechain
34	BA	1439	C	Sidechain
34	BA	144	C	Sidechain
34	BA	1440	C	Sidechain
34	BA	1441	C	Sidechain
34	BA	1442	A	Sidechain
34	BA	1443	U	Sidechain
34	BA	1444	G	Sidechain
34	BA	1446	G	Sidechain
34	BA	1448	G	Sidechain
34	BA	145	U	Sidechain
34	BA	1450	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1451	A	Sidechain
34	BA	1453	U	Sidechain
34	BA	1454	G	Sidechain
34	BA	1455	C	Sidechain
34	BA	1456	C	Sidechain
34	BA	1458	A	Sidechain
34	BA	1459	U	Sidechain
34	BA	146	G	Sidechain
34	BA	1460	U	Sidechain
34	BA	1461	A	Sidechain
34	BA	1462	U	Sidechain
34	BA	1463	U	Sidechain
34	BA	1464	C	Sidechain
34	BA	1466	U	Sidechain
34	BA	1467	U	Sidechain
34	BA	1468	U	Sidechain
34	BA	1469	G	Sidechain
34	BA	147	U	Sidechain
34	BA	1470	G	Sidechain
34	BA	1472	G	Sidechain
34	BA	1474	G	Sidechain
34	BA	1476	G	Sidechain
34	BA	1477	C	Sidechain
34	BA	1478	G	Sidechain
34	BA	148	G	Sidechain
34	BA	1482	A	Sidechain
34	BA	1483	U	Sidechain
34	BA	1485	U	Sidechain
34	BA	1486	U	Sidechain
34	BA	1487	U	Sidechain
34	BA	1488	C	Sidechain
34	BA	149	G	Sidechain
34	BA	1490	U	Sidechain
34	BA	1491	U	Sidechain
34	BA	1492	G	Sidechain
34	BA	1493	U	Sidechain
34	BA	1494	G	Sidechain
34	BA	1496	G	Sidechain
34	BA	1497	A	Sidechain
34	BA	1498	A	Sidechain
34	BA	15	G	Sidechain
34	BA	150	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1500	G	Sidechain
34	BA	1501	U	Sidechain
34	BA	1502	G	Sidechain
34	BA	1503	U	Sidechain
34	BA	1504	A	Sidechain
34	BA	1505	G	Sidechain
34	BA	1506	C	Sidechain
34	BA	1507	C	Sidechain
34	BA	1508	C	Sidechain
34	BA	1509	U	Sidechain
34	BA	151	A	Sidechain
34	BA	1511	C	Sidechain
34	BA	1512	C	Sidechain
34	BA	1513	G	Sidechain
34	BA	1514	A	Sidechain
34	BA	1515	U	Sidechain
34	BA	1516	G	Sidechain
34	BA	1517	U	Sidechain
34	BA	1518	A	Sidechain
34	BA	1519	G	Sidechain
34	BA	152	C	Sidechain
34	BA	1522	G	Sidechain
34	BA	1523	U	Sidechain
34	BA	1524	G	Sidechain
34	BA	1525	G	Sidechain
34	BA	1526	C	Sidechain
34	BA	1527	G	Sidechain
34	BA	1530	G	Sidechain
34	BA	1534	U	Sidechain
34	BA	1536	A	Sidechain
34	BA	1537	G	Sidechain
34	BA	1538	G	Sidechain
34	BA	154	A	Sidechain
34	BA	1540	C	Sidechain
34	BA	1541	G	Sidechain
34	BA	1542	A	Sidechain
34	BA	1544	G	Sidechain
34	BA	1546	C	Sidechain
34	BA	1549	U	Sidechain
34	BA	155	U	Sidechain
34	BA	1550	G	Sidechain
34	BA	1551	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1552	C	Sidechain
34	BA	1553	G	Sidechain
34	BA	1554	C	Sidechain
34	BA	1555	G	Sidechain
34	BA	1556	A	Sidechain
34	BA	1557	G	Sidechain
34	BA	156	U	Sidechain
34	BA	1561	C	Sidechain
34	BA	1562	G	Sidechain
34	BA	1563	G	Sidechain
34	BA	1564	A	Sidechain
34	BA	1565	U	Sidechain
34	BA	1566	G	Sidechain
34	BA	1567	G	Sidechain
34	BA	1569	C	Sidechain
34	BA	157	U	Sidechain
34	BA	1570	C	Sidechain
34	BA	1571	C	Sidechain
34	BA	1572	G	Sidechain
34	BA	1573	C	Sidechain
34	BA	1577	U	Sidechain
34	BA	1579	G	Sidechain
34	BA	1580	U	Sidechain
34	BA	1582	C	Sidechain
34	BA	1585	A	Sidechain
34	BA	1586	U	Sidechain
34	BA	1587	C	Sidechain
34	BA	1588	U	Sidechain
34	BA	1589	U	Sidechain
34	BA	159	U	Sidechain
34	BA	1590	G	Sidechain
34	BA	1591	G	Sidechain
34	BA	1592	U	Sidechain
34	BA	1595	G	Sidechain
34	BA	1596	C	Sidechain
34	BA	1597	G	Sidechain
34	BA	1598	U	Sidechain
34	BA	1599	A	Sidechain
34	BA	16	C	Sidechain
34	BA	160	G	Sidechain
34	BA	1600	G	Sidechain
34	BA	1601	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1603	A	Sidechain
34	BA	1604	A	Sidechain
34	BA	1605	G	Sidechain
34	BA	1606	A	Sidechain
34	BA	1607	U	Sidechain
34	BA	1609	U	Sidechain
34	BA	161	U	Sidechain
34	BA	1611	A	Sidechain
34	BA	1612	C	Sidechain
34	BA	1613	G	Sidechain
34	BA	1614	G	Sidechain
34	BA	1615	A	Sidechain
34	BA	1616	A	Sidechain
34	BA	1617	U	Sidechain
34	BA	1618	A	Sidechain
34	BA	1619	U	Sidechain
34	BA	162	G	Sidechain
34	BA	1621	U	Sidechain
34	BA	1623	U	Sidechain
34	BA	1624	U	Sidechain
34	BA	1626	U	Sidechain
34	BA	1627	U	Sidechain
34	BA	1628	A	Sidechain
34	BA	163	G	Sidechain
34	BA	1630	A	Sidechain
34	BA	1631	U	Sidechain
34	BA	1632	G	Sidechain
34	BA	1633	C	Sidechain
34	BA	1634	A	Sidechain
34	BA	1636	C	Sidechain
34	BA	1637	G	Sidechain
34	BA	1638	U	Sidechain
34	BA	1639	U	Sidechain
34	BA	1640	G	Sidechain
34	BA	1641	G	Sidechain
34	BA	1642	A	Sidechain
34	BA	1643	U	Sidechain
34	BA	1646	U	Sidechain
34	BA	1647	G	Sidechain
34	BA	1648	G	Sidechain
34	BA	1649	A	Sidechain
34	BA	1650	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1652	G	Sidechain
34	BA	1653	G	Sidechain
34	BA	1654	G	Sidechain
34	BA	1655	G	Sidechain
34	BA	1659	G	Sidechain
34	BA	166	G	Sidechain
34	BA	1660	A	Sidechain
34	BA	1661	U	Sidechain
34	BA	1662	U	Sidechain
34	BA	1663	U	Sidechain
34	BA	1664	C	Sidechain
34	BA	1666	U	Sidechain
34	BA	1668	C	Sidechain
34	BA	1669	C	Sidechain
34	BA	167	U	Sidechain
34	BA	1671	A	Sidechain
34	BA	1672	C	Sidechain
34	BA	1673	G	Sidechain
34	BA	1674	G	Sidechain
34	BA	1675	C	Sidechain
34	BA	1676	A	Sidechain
34	BA	1677	C	Sidechain
34	BA	1678	U	Sidechain
34	BA	1679	C	Sidechain
34	BA	168	U	Sidechain
34	BA	1680	G	Sidechain
34	BA	1681	U	Sidechain
34	BA	1683	C	Sidechain
34	BA	1684	A	Sidechain
34	BA	1685	C	Sidechain
34	BA	1686	G	Sidechain
34	BA	1687	A	Sidechain
34	BA	1688	G	Sidechain
34	BA	1689	U	Sidechain
34	BA	1690	U	Sidechain
34	BA	1691	G	Sidechain
34	BA	1692	U	Sidechain
34	BA	1693	U	Sidechain
34	BA	1694	C	Sidechain
34	BA	1695	G	Sidechain
34	BA	1696	G	Sidechain
34	BA	1697	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1699	A	Sidechain
34	BA	17	A	Sidechain
34	BA	170	U	Sidechain
34	BA	1700	C	Sidechain
34	BA	1701	U	Sidechain
34	BA	1702	G	Sidechain
34	BA	1703	A	Sidechain
34	BA	1704	G	Sidechain
34	BA	1705	C	Sidechain
34	BA	1707	C	Sidechain
34	BA	1708	A	Sidechain
34	BA	1709	A	Sidechain
34	BA	171	U	Sidechain
34	BA	1710	C	Sidechain
34	BA	1711	G	Sidechain
34	BA	1712	U	Sidechain
34	BA	1713	U	Sidechain
34	BA	1716	A	Sidechain
34	BA	1718	C	Sidechain
34	BA	1719	G	Sidechain
34	BA	1720	U	Sidechain
34	BA	1721	U	Sidechain
34	BA	1722	U	Sidechain
34	BA	1723	U	Sidechain
34	BA	1724	G	Sidechain
34	BA	1726	U	Sidechain
34	BA	1727	A	Sidechain
34	BA	1728	G	Sidechain
34	BA	1729	G	Sidechain
34	BA	173	U	Sidechain
34	BA	1730	A	Sidechain
34	BA	1731	A	Sidechain
34	BA	1732	A	Sidechain
34	BA	1733	G	Sidechain
34	BA	1734	U	Sidechain
34	BA	1735	G	Sidechain
34	BA	1738	G	Sidechain
34	BA	1739	G	Sidechain
34	BA	174	A	Sidechain
34	BA	1740	U	Sidechain
34	BA	1741	G	Sidechain
34	BA	1742	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1743	U	Sidechain
34	BA	1744	C	Sidechain
34	BA	1745	G	Sidechain
34	BA	1746	G	Sidechain
34	BA	1747	C	Sidechain
34	BA	1748	G	Sidechain
34	BA	175	G	Sidechain
34	BA	1754	C	Sidechain
34	BA	1756	C	Sidechain
34	BA	1759	U	Sidechain
34	BA	1765	G	Sidechain
34	BA	177	G	Sidechain
34	BA	1771	U	Sidechain
34	BA	1772	G	Sidechain
34	BA	1775	U	Sidechain
34	BA	1779	U	Sidechain
34	BA	178	C	Sidechain
34	BA	1780	U	Sidechain
34	BA	1781	A	Sidechain
34	BA	1783	C	Sidechain
34	BA	1784	G	Sidechain
34	BA	1785	G	Sidechain
34	BA	1786	C	Sidechain
34	BA	1787	U	Sidechain
34	BA	1788	U	Sidechain
34	BA	179	U	Sidechain
34	BA	1790	U	Sidechain
34	BA	1792	U	Sidechain
34	BA	1793	G	Sidechain
34	BA	1795	A	Sidechain
34	BA	1796	A	Sidechain
34	BA	1797	A	Sidechain
34	BA	1798	G	Sidechain
34	BA	1799	G	Sidechain
34	BA	18	G	Sidechain
34	BA	180	G	Sidechain
34	BA	1800	G	Sidechain
34	BA	1801	G	Sidechain
34	BA	1802	C	Sidechain
34	BA	1803	A	Sidechain
34	BA	1806	A	Sidechain
34	BA	1807	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	1808	A	Sidechain
34	BA	1809	G	Sidechain
34	BA	181	G	Sidechain
34	BA	1814	U	Sidechain
34	BA	1815	G	Sidechain
34	BA	1816	G	Sidechain
34	BA	1817	G	Sidechain
34	BA	1818	A	Sidechain
34	BA	1819	U	Sidechain
34	BA	182	U	Sidechain
34	BA	1820	G	Sidechain
34	BA	1821	A	Sidechain
34	BA	1822	U	Sidechain
34	BA	1823	A	Sidechain
34	BA	1824	U	Sidechain
34	BA	1825	U	Sidechain
34	BA	1826	C	Sidechain
34	BA	1827	C	Sidechain
34	BA	1828	A	Sidechain
34	BA	1829	A	Sidechain
34	BA	183	G	Sidechain
34	BA	1831	A	Sidechain
34	BA	1833	G	Sidechain
34	BA	1834	A	Sidechain
34	BA	1836	A	Sidechain
34	BA	1837	U	Sidechain
34	BA	1838	U	Sidechain
34	BA	1839	G	Sidechain
34	BA	1841	A	Sidechain
34	BA	1842	U	Sidechain
34	BA	1844	U	Sidechain
34	BA	1845	G	Sidechain
34	BA	1846	G	Sidechain
34	BA	1847	G	Sidechain
34	BA	185	A	Sidechain
34	BA	186	G	Sidechain
34	BA	188	C	Sidechain
34	BA	189	G	Sidechain
34	BA	19	G	Sidechain
34	BA	190	U	Sidechain
34	BA	192	G	Sidechain
34	BA	193	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	194	G	Sidechain
34	BA	195	G	Sidechain
34	BA	197	A	Sidechain
34	BA	198	U	Sidechain
34	BA	199	U	Sidechain
34	BA	2	A	Sidechain
34	BA	20	A	Sidechain
34	BA	201	A	Sidechain
34	BA	202	A	Sidechain
34	BA	203	U	Sidechain
34	BA	204	U	Sidechain
34	BA	205	G	Sidechain
34	BA	206	C	Sidechain
34	BA	207	A	Sidechain
34	BA	208	A	Sidechain
34	BA	209	A	Sidechain
34	BA	21	C	Sidechain
34	BA	210	G	Sidechain
34	BA	211	C	Sidechain
34	BA	212	A	Sidechain
34	BA	213	A	Sidechain
34	BA	214	A	Sidechain
34	BA	215	C	Sidechain
34	BA	216	C	Sidechain
34	BA	217	C	Sidechain
34	BA	218	G	Sidechain
34	BA	22	C	Sidechain
34	BA	220	U	Sidechain
34	BA	221	G	Sidechain
34	BA	222	C	Sidechain
34	BA	224	G	Sidechain
34	BA	225	A	Sidechain
34	BA	226	A	Sidechain
34	BA	227	C	Sidechain
34	BA	228	A	Sidechain
34	BA	229	C	Sidechain
34	BA	23	A	Sidechain
34	BA	230	A	Sidechain
34	BA	231	U	Sidechain
34	BA	232	U	Sidechain
34	BA	233	U	Sidechain
34	BA	235	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	236	A	Sidechain
34	BA	237	A	Sidechain
34	BA	238	C	Sidechain
34	BA	24	C	Sidechain
34	BA	240	C	Sidechain
34	BA	241	U	Sidechain
34	BA	242	U	Sidechain
34	BA	243	C	Sidechain
34	BA	244	A	Sidechain
34	BA	245	U	Sidechain
34	BA	246	G	Sidechain
34	BA	247	U	Sidechain
34	BA	248	G	Sidechain
34	BA	249	A	Sidechain
34	BA	25	C	Sidechain
34	BA	250	G	Sidechain
34	BA	251	U	Sidechain
34	BA	252	A	Sidechain
34	BA	253	U	Sidechain
34	BA	254	U	Sidechain
34	BA	255	G	Sidechain
34	BA	256	A	Sidechain
34	BA	257	G	Sidechain
34	BA	259	C	Sidechain
34	BA	26	C	Sidechain
34	BA	260	A	Sidechain
34	BA	261	A	Sidechain
34	BA	262	A	Sidechain
34	BA	263	G	Sidechain
34	BA	264	A	Sidechain
34	BA	265	A	Sidechain
34	BA	266	G	Sidechain
34	BA	267	G	Sidechain
34	BA	268	U	Sidechain
34	BA	269	G	Sidechain
34	BA	270	U	Sidechain
34	BA	271	C	Sidechain
34	BA	273	G	Sidechain
34	BA	275	C	Sidechain
34	BA	277	A	Sidechain
34	BA	278	U	Sidechain
34	BA	279	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	280	A	Sidechain
34	BA	281	C	Sidechain
34	BA	282	A	Sidechain
34	BA	283	U	Sidechain
34	BA	284	U	Sidechain
34	BA	285	C	Sidechain
34	BA	286	C	Sidechain
34	BA	288	U	Sidechain
34	BA	289	A	Sidechain
34	BA	29	U	Sidechain
34	BA	290	G	Sidechain
34	BA	291	C	Sidechain
34	BA	293	A	Sidechain
34	BA	294	C	Sidechain
34	BA	295	G	Sidechain
34	BA	296	G	Sidechain
34	BA	297	A	Sidechain
34	BA	298	G	Sidechain
34	BA	299	C	Sidechain
34	BA	3	G	Sidechain
34	BA	30	A	Sidechain
34	BA	300	C	Sidechain
34	BA	301	U	Sidechain
34	BA	302	A	Sidechain
34	BA	303	C	Sidechain
34	BA	304	G	Sidechain
34	BA	309	U	Sidechain
34	BA	31	A	Sidechain
34	BA	312	U	Sidechain
34	BA	313	C	Sidechain
34	BA	314	A	Sidechain
34	BA	315	U	Sidechain
34	BA	316	G	Sidechain
34	BA	318	U	Sidechain
34	BA	319	C	Sidechain
34	BA	32	A	Sidechain
34	BA	320	G	Sidechain
34	BA	321	G	Sidechain
34	BA	322	U	Sidechain
34	BA	323	C	Sidechain
34	BA	324	C	Sidechain
34	BA	325	A	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	326	A	Sidechain
34	BA	327	G	Sidechain
34	BA	328	A	Sidechain
34	BA	329	G	Sidechain
34	BA	331	G	Sidechain
34	BA	332	U	Sidechain
34	BA	334	G	Sidechain
34	BA	335	C	Sidechain
34	BA	336	A	Sidechain
34	BA	338	U	Sidechain
34	BA	339	G	Sidechain
34	BA	34	U	Sidechain
34	BA	340	U	Sidechain
34	BA	341	U	Sidechain
34	BA	342	U	Sidechain
34	BA	343	G	Sidechain
34	BA	344	G	Sidechain
34	BA	345	G	Sidechain
34	BA	348	U	Sidechain
34	BA	349	G	Sidechain
34	BA	35	U	Sidechain
34	BA	352	G	Sidechain
34	BA	353	U	Sidechain
34	BA	355	U	Sidechain
34	BA	358	A	Sidechain
34	BA	359	G	Sidechain
34	BA	36	A	Sidechain
34	BA	360	C	Sidechain
34	BA	361	C	Sidechain
34	BA	362	G	Sidechain
34	BA	363	G	Sidechain
34	BA	364	C	Sidechain
34	BA	365	A	Sidechain
34	BA	366	G	Sidechain
34	BA	367	G	Sidechain
34	BA	368	U	Sidechain
34	BA	369	A	Sidechain
34	BA	370	U	Sidechain
34	BA	371	U	Sidechain
34	BA	372	U	Sidechain
34	BA	376	U	Sidechain
34	BA	377	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	378	C	Sidechain
34	BA	379	C	Sidechain
34	BA	38	G	Sidechain
34	BA	382	G	Sidechain
34	BA	383	G	Sidechain
34	BA	384	U	Sidechain
34	BA	385	U	Sidechain
34	BA	386	A	Sidechain
34	BA	387	A	Sidechain
34	BA	388	A	Sidechain
34	BA	39	C	Sidechain
34	BA	390	A	Sidechain
34	BA	391	U	Sidechain
34	BA	393	G	Sidechain
34	BA	395	G	Sidechain
34	BA	396	U	Sidechain
34	BA	397	A	Sidechain
34	BA	398	G	Sidechain
34	BA	399	G	Sidechain
34	BA	400	A	Sidechain
34	BA	401	A	Sidechain
34	BA	402	G	Sidechain
34	BA	403	A	Sidechain
34	BA	405	C	Sidechain
34	BA	406	G	Sidechain
34	BA	407	A	Sidechain
34	BA	408	U	Sidechain
34	BA	409	A	Sidechain
34	BA	41	U	Sidechain
34	BA	410	G	Sidechain
34	BA	411	C	Sidechain
34	BA	413	A	Sidechain
34	BA	415	C	Sidechain
34	BA	416	A	Sidechain
34	BA	417	A	Sidechain
34	BA	418	G	Sidechain
34	BA	419	U	Sidechain
34	BA	42	A	Sidechain
34	BA	420	A	Sidechain
34	BA	422	C	Sidechain
34	BA	423	G	Sidechain
34	BA	424	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	425	G	Sidechain
34	BA	426	A	Sidechain
34	BA	427	G	Sidechain
34	BA	429	G	Sidechain
34	BA	43	U	Sidechain
34	BA	431	A	Sidechain
34	BA	433	G	Sidechain
34	BA	434	U	Sidechain
34	BA	435	U	Sidechain
34	BA	436	U	Sidechain
34	BA	437	G	Sidechain
34	BA	438	A	Sidechain
34	BA	441	A	Sidechain
34	BA	442	G	Sidechain
34	BA	443	U	Sidechain
34	BA	445	C	Sidechain
34	BA	446	U	Sidechain
34	BA	448	U	Sidechain
34	BA	449	G	Sidechain
34	BA	453	A	Sidechain
34	BA	454	G	Sidechain
34	BA	455	A	Sidechain
34	BA	457	A	Sidechain
34	BA	458	G	Sidechain
34	BA	46	C	Sidechain
34	BA	460	G	Sidechain
34	BA	462	C	Sidechain
34	BA	463	A	Sidechain
34	BA	464	U	Sidechain
34	BA	466	G	Sidechain
34	BA	467	A	Sidechain
34	BA	469	C	Sidechain
34	BA	471	U	Sidechain
34	BA	472	G	Sidechain
34	BA	473	A	Sidechain
34	BA	474	A	Sidechain
34	BA	475	A	Sidechain
34	BA	476	U	Sidechain
34	BA	477	C	Sidechain
34	BA	478	G	Sidechain
34	BA	479	U	Sidechain
34	BA	48	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	480	G	Sidechain
34	BA	481	A	Sidechain
34	BA	484	A	Sidechain
34	BA	485	C	Sidechain
34	BA	486	G	Sidechain
34	BA	488	C	Sidechain
34	BA	49	A	Sidechain
34	BA	490	A	Sidechain
34	BA	491	U	Sidechain
34	BA	492	G	Sidechain
34	BA	495	A	Sidechain
34	BA	496	G	Sidechain
34	BA	498	A	Sidechain
34	BA	499	C	Sidechain
34	BA	500	C	Sidechain
34	BA	501	U	Sidechain
34	BA	502	U	Sidechain
34	BA	503	C	Sidechain
34	BA	504	A	Sidechain
34	BA	505	U	Sidechain
34	BA	506	U	Sidechain
34	BA	507	U	Sidechain
34	BA	51	C	Sidechain
34	BA	510	U	Sidechain
34	BA	511	U	Sidechain
34	BA	512	U	Sidechain
34	BA	513	U	Sidechain
34	BA	514	U	Sidechain
34	BA	515	U	Sidechain
34	BA	516	U	Sidechain
34	BA	517	A	Sidechain
34	BA	518	C	Sidechain
34	BA	519	G	Sidechain
34	BA	52	G	Sidechain
34	BA	520	G	Sidechain
34	BA	521	C	Sidechain
34	BA	522	C	Sidechain
34	BA	523	A	Sidechain
34	BA	525	A	Sidechain
34	BA	526	C	Sidechain
34	BA	527	C	Sidechain
34	BA	528	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	529	A	Sidechain
34	BA	53	G	Sidechain
34	BA	530	A	Sidechain
34	BA	531	C	Sidechain
34	BA	532	C	Sidechain
34	BA	533	U	Sidechain
34	BA	534	C	Sidechain
34	BA	535	G	Sidechain
34	BA	537	C	Sidechain
34	BA	538	G	Sidechain
34	BA	539	C	Sidechain
34	BA	540	G	Sidechain
34	BA	541	C	Sidechain
34	BA	542	A	Sidechain
34	BA	543	A	Sidechain
34	BA	544	U	Sidechain
34	BA	546	U	Sidechain
34	BA	548	G	Sidechain
34	BA	549	G	Sidechain
34	BA	55	G	Sidechain
34	BA	550	U	Sidechain
34	BA	555	C	Sidechain
34	BA	556	A	Sidechain
34	BA	557	U	Sidechain
34	BA	559	C	Sidechain
34	BA	56	G	Sidechain
34	BA	560	U	Sidechain
34	BA	563	A	Sidechain
34	BA	564	C	Sidechain
34	BA	565	U	Sidechain
34	BA	566	G	Sidechain
34	BA	567	U	Sidechain
34	BA	568	G	Sidechain
34	BA	569	C	Sidechain
34	BA	57	A	Sidechain
34	BA	570	G	Sidechain
34	BA	571	G	Sidechain
34	BA	572	G	Sidechain
34	BA	573	U	Sidechain
34	BA	574	U	Sidechain
34	BA	575	U	Sidechain
34	BA	578	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	579	U	Sidechain
34	BA	58	A	Sidechain
34	BA	581	U	Sidechain
34	BA	582	U	Sidechain
34	BA	583	G	Sidechain
34	BA	584	A	Sidechain
34	BA	585	G	Sidechain
34	BA	586	G	Sidechain
34	BA	588	C	Sidechain
34	BA	589	A	Sidechain
34	BA	59	A	Sidechain
34	BA	590	U	Sidechain
34	BA	591	G	Sidechain
34	BA	592	G	Sidechain
34	BA	593	G	Sidechain
34	BA	594	G	Sidechain
34	BA	595	U	Sidechain
34	BA	596	G	Sidechain
34	BA	597	C	Sidechain
34	BA	598	G	Sidechain
34	BA	599	U	Sidechain
34	BA	6	C	Sidechain
34	BA	60	A	Sidechain
34	BA	600	G	Sidechain
34	BA	601	A	Sidechain
34	BA	602	G	Sidechain
34	BA	605	G	Sidechain
34	BA	606	G	Sidechain
34	BA	607	C	Sidechain
34	BA	608	G	Sidechain
34	BA	61	G	Sidechain
34	BA	610	A	Sidechain
34	BA	611	A	Sidechain
34	BA	613	A	Sidechain
34	BA	615	A	Sidechain
34	BA	616	G	Sidechain
34	BA	621	G	Sidechain
34	BA	622	G	Sidechain
34	BA	623	U	Sidechain
34	BA	624	G	Sidechain
34	BA	625	U	Sidechain
34	BA	626	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	627	U	Sidechain
34	BA	628	U	Sidechain
34	BA	629	G	Sidechain
34	BA	63	A	Sidechain
34	BA	630	U	Sidechain
34	BA	631	G	Sidechain
34	BA	633	G	Sidechain
34	BA	641	U	Sidechain
34	BA	649	A	Sidechain
34	BA	651	U	Sidechain
34	BA	652	C	Sidechain
34	BA	653	U	Sidechain
34	BA	654	C	Sidechain
34	BA	656	U	Sidechain
34	BA	657	C	Sidechain
34	BA	659	U	Sidechain
34	BA	660	C	Sidechain
34	BA	661	C	Sidechain
34	BA	662	U	Sidechain
34	BA	663	U	Sidechain
34	BA	664	C	Sidechain
34	BA	665	C	Sidechain
34	BA	666	C	Sidechain
34	BA	667	U	Sidechain
34	BA	668	G	Sidechain
34	BA	669	U	Sidechain
34	BA	67	A	Sidechain
34	BA	670	U	Sidechain
34	BA	671	C	Sidechain
34	BA	672	G	Sidechain
34	BA	673	U	Sidechain
34	BA	674	G	Sidechain
34	BA	675	C	Sidechain
34	BA	676	G	Sidechain
34	BA	677	U	Sidechain
34	BA	678	C	Sidechain
34	BA	679	U	Sidechain
34	BA	68	A	Sidechain
34	BA	680	C	Sidechain
34	BA	681	G	Sidechain
34	BA	682	A	Sidechain
34	BA	683	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	684	G	Sidechain
34	BA	685	C	Sidechain
34	BA	686	U	Sidechain
34	BA	687	G	Sidechain
34	BA	688	G	Sidechain
34	BA	689	C	Sidechain
34	BA	69	C	Sidechain
34	BA	690	G	Sidechain
34	BA	691	A	Sidechain
34	BA	692	U	Sidechain
34	BA	693	G	Sidechain
34	BA	695	A	Sidechain
34	BA	696	A	Sidechain
34	BA	697	A	Sidechain
34	BA	699	G	Sidechain
34	BA	7	U	Sidechain
34	BA	70	C	Sidechain
34	BA	700	G	Sidechain
34	BA	701	G	Sidechain
34	BA	703	U	Sidechain
34	BA	704	G	Sidechain
34	BA	705	C	Sidechain
34	BA	706	C	Sidechain
34	BA	707	C	Sidechain
34	BA	708	C	Sidechain
34	BA	709	C	Sidechain
34	BA	71	G	Sidechain
34	BA	710	A	Sidechain
34	BA	711	C	Sidechain
34	BA	712	C	Sidechain
34	BA	713	C	Sidechain
34	BA	714	G	Sidechain
34	BA	715	U	Sidechain
34	BA	716	C	Sidechain
34	BA	717	U	Sidechain
34	BA	718	U	Sidechain
34	BA	719	G	Sidechain
34	BA	72	U	Sidechain
34	BA	721	A	Sidechain
34	BA	723	C	Sidechain
34	BA	725	C	Sidechain
34	BA	726	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	727	G	Sidechain
34	BA	728	A	Sidechain
34	BA	73	G	Sidechain
34	BA	730	C	Sidechain
34	BA	731	A	Sidechain
34	BA	732	A	Sidechain
34	BA	733	G	Sidechain
34	BA	736	G	Sidechain
34	BA	738	C	Sidechain
34	BA	739	A	Sidechain
34	BA	740	A	Sidechain
34	BA	741	A	Sidechain
34	BA	742	C	Sidechain
34	BA	743	A	Sidechain
34	BA	744	G	Sidechain
34	BA	745	A	Sidechain
34	BA	746	C	Sidechain
34	BA	747	G	Sidechain
34	BA	75	U	Sidechain
34	BA	751	A	Sidechain
34	BA	753	G	Sidechain
34	BA	755	G	Sidechain
34	BA	756	A	Sidechain
34	BA	757	G	Sidechain
34	BA	758	G	Sidechain
34	BA	759	A	Sidechain
34	BA	76	U	Sidechain
34	BA	761	U	Sidechain
34	BA	762	A	Sidechain
34	BA	763	U	Sidechain
34	BA	764	G	Sidechain
34	BA	765	U	Sidechain
34	BA	766	A	Sidechain
34	BA	767	U	Sidechain
34	BA	769	U	Sidechain
34	BA	77	C	Sidechain
34	BA	770	G	Sidechain
34	BA	771	A	Sidechain
34	BA	772	G	Sidechain
34	BA	773	A	Sidechain
34	BA	774	A	Sidechain
34	BA	775	C	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	776	U	Sidechain
34	BA	777	C	Sidechain
34	BA	779	U	Sidechain
34	BA	78	U	Sidechain
34	BA	780	U	Sidechain
34	BA	781	U	Sidechain
34	BA	782	C	Sidechain
34	BA	783	U	Sidechain
34	BA	785	G	Sidechain
34	BA	786	U	Sidechain
34	BA	787	A	Sidechain
34	BA	788	C	Sidechain
34	BA	789	U	Sidechain
34	BA	79	C	Sidechain
34	BA	790	G	Sidechain
34	BA	792	A	Sidechain
34	BA	794	G	Sidechain
34	BA	796	G	Sidechain
34	BA	798	G	Sidechain
34	BA	799	A	Sidechain
34	BA	8	G	Sidechain
34	BA	80	U	Sidechain
34	BA	800	G	Sidechain
34	BA	801	U	Sidechain
34	BA	802	G	Sidechain
34	BA	803	U	Sidechain
34	BA	804	G	Sidechain
34	BA	805	A	Sidechain
34	BA	806	U	Sidechain
34	BA	807	U	Sidechain
34	BA	808	U	Sidechain
34	BA	812	A	Sidechain
34	BA	814	C	Sidechain
34	BA	815	C	Sidechain
34	BA	816	G	Sidechain
34	BA	817	U	Sidechain
34	BA	819	G	Sidechain
34	BA	82	A	Sidechain
34	BA	820	C	Sidechain
34	BA	821	G	Sidechain
34	BA	822	U	Sidechain
34	BA	823	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	824	C	Sidechain
34	BA	825	G	Sidechain
34	BA	828	A	Sidechain
34	BA	829	U	Sidechain
34	BA	83	G	Sidechain
34	BA	830	U	Sidechain
34	BA	832	C	Sidechain
34	BA	833	U	Sidechain
34	BA	834	C	Sidechain
34	BA	835	U	Sidechain
34	BA	836	U	Sidechain
34	BA	837	U	Sidechain
34	BA	838	U	Sidechain
34	BA	839	U	Sidechain
34	BA	84	U	Sidechain
34	BA	840	U	Sidechain
34	BA	842	U	Sidechain
34	BA	844	U	Sidechain
34	BA	845	U	Sidechain
34	BA	847	U	Sidechain
34	BA	848	U	Sidechain
34	BA	849	G	Sidechain
34	BA	852	C	Sidechain
34	BA	853	A	Sidechain
34	BA	854	A	Sidechain
34	BA	855	C	Sidechain
34	BA	858	C	Sidechain
34	BA	859	G	Sidechain
34	BA	86	A	Sidechain
34	BA	860	G	Sidechain
34	BA	862	C	Sidechain
34	BA	863	G	Sidechain
34	BA	864	G	Sidechain
34	BA	87	G	Sidechain
34	BA	870	C	Sidechain
34	BA	871	G	Sidechain
34	BA	872	U	Sidechain
34	BA	873	G	Sidechain
34	BA	874	G	Sidechain
34	BA	875	G	Sidechain
34	BA	876	C	Sidechain
34	BA	877	U	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	878	G	Sidechain
34	BA	879	C	Sidechain
34	BA	88	C	Sidechain
34	BA	880	G	Sidechain
34	BA	882	G	Sidechain
34	BA	883	C	Sidechain
34	BA	884	G	Sidechain
34	BA	886	G	Sidechain
34	BA	887	U	Sidechain
34	BA	889	U	Sidechain
34	BA	89	G	Sidechain
34	BA	890	G	Sidechain
34	BA	893	U	Sidechain
34	BA	894	G	Sidechain
34	BA	895	U	Sidechain
34	BA	896	U	Sidechain
34	BA	897	U	Sidechain
34	BA	898	G	Sidechain
34	BA	899	G	Sidechain
34	BA	9	A	Sidechain
34	BA	90	G	Sidechain
34	BA	900	A	Sidechain
34	BA	901	C	Sidechain
34	BA	902	C	Sidechain
34	BA	903	C	Sidechain
34	BA	904	G	Sidechain
34	BA	905	A	Sidechain
34	BA	906	A	Sidechain
34	BA	909	G	Sidechain
34	BA	91	C	Sidechain
34	BA	910	U	Sidechain
34	BA	912	G	Sidechain
34	BA	913	U	Sidechain
34	BA	914	G	Sidechain
34	BA	915	A	Sidechain
34	BA	916	A	Sidechain
34	BA	917	C	Sidechain
34	BA	918	U	Sidechain
34	BA	919	A	Sidechain
34	BA	92	G	Sidechain
34	BA	920	U	Sidechain
34	BA	921	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	923	C	Sidechain
34	BA	924	U	Sidechain
34	BA	925	G	Sidechain
34	BA	926	A	Sidechain
34	BA	928	C	Sidechain
34	BA	929	A	Sidechain
34	BA	93	A	Sidechain
34	BA	930	A	Sidechain
34	BA	931	G	Sidechain
34	BA	932	G	Sidechain
34	BA	933	U	Sidechain
34	BA	934	G	Sidechain
34	BA	937	G	Sidechain
34	BA	939	C	Sidechain
34	BA	94	G	Sidechain
34	BA	940	C	Sidechain
34	BA	941	G	Sidechain
34	BA	942	G	Sidechain
34	BA	944	G	Sidechain
34	BA	945	A	Sidechain
34	BA	946	A	Sidechain
34	BA	947	A	Sidechain
34	BA	95	C	Sidechain
34	BA	950	C	Sidechain
34	BA	952	G	Sidechain
34	BA	953	G	Sidechain
34	BA	954	U	Sidechain
34	BA	956	G	Sidechain
34	BA	957	A	Sidechain
34	BA	959	G	Sidechain
34	BA	96	G	Sidechain
34	BA	960	C	Sidechain
34	BA	961	C	Sidechain
34	BA	962	U	Sidechain
34	BA	963	G	Sidechain
34	BA	964	U	Sidechain
34	BA	965	A	Sidechain
34	BA	968	G	Sidechain
34	BA	97	A	Sidechain
34	BA	970	U	Sidechain
34	BA	972	C	Sidechain
34	BA	974	G	Sidechain

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Mol	Chain	Res	Type	Group
34	BA	975	A	Sidechain
34	BA	976	C	Sidechain
34	BA	977	G	Sidechain
34	BA	98	A	Sidechain
34	BA	980	C	Sidechain
34	BA	982	A	Sidechain
34	BA	984	U	Sidechain
34	BA	985	C	Sidechain
34	BA	986	G	Sidechain
34	BA	987	C	Sidechain
34	BA	99	G	Sidechain
34	BA	991	U	Sidechain
34	BA	992	A	Sidechain
34	BA	993	C	Sidechain
34	BA	994	G	Sidechain
34	BA	996	U	Sidechain
34	BA	997	U	Sidechain
34	BA	998	U	Sidechain
34	BA	999	G	Sidechain
35	BB	1	U	Sidechain
35	BB	100	A	Sidechain
35	BB	1001	G	Sidechain
35	BB	1003	G	Sidechain
35	BB	1004	A	Sidechain
35	BB	1005	A	Sidechain
35	BB	1007	U	Sidechain
35	BB	101	U	Sidechain
35	BB	1012	G	Sidechain
35	BB	1014	U	Sidechain
35	BB	1015	U	Sidechain
35	BB	1016	C	Sidechain
35	BB	1017	U	Sidechain
35	BB	1018	U	Sidechain
35	BB	1019	C	Sidechain
35	BB	102	G	Sidechain
35	BB	1020	U	Sidechain
35	BB	1021	C	Sidechain
35	BB	1022	C	Sidechain
35	BB	1023	G	Sidechain
35	BB	1024	G	Sidechain
35	BB	1025	A	Sidechain
35	BB	1026	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1027	U	Sidechain
35	BB	1028	C	Sidechain
35	BB	1029	U	Sidechain
35	BB	1030	U	Sidechain
35	BB	1031	G	Sidechain
35	BB	1032	U	Sidechain
35	BB	1033	U	Sidechain
35	BB	1034	U	Sidechain
35	BB	1035	C	Sidechain
35	BB	1036	G	Sidechain
35	BB	1037	A	Sidechain
35	BB	1038	G	Sidechain
35	BB	104	G	Sidechain
35	BB	1040	C	Sidechain
35	BB	1041	A	Sidechain
35	BB	1042	U	Sidechain
35	BB	1044	U	Sidechain
35	BB	1045	G	Sidechain
35	BB	1048	A	Sidechain
35	BB	1049	G	Sidechain
35	BB	105	U	Sidechain
35	BB	1051	U	Sidechain
35	BB	1053	G	Sidechain
35	BB	1054	G	Sidechain
35	BB	1055	G	Sidechain
35	BB	1056	A	Sidechain
35	BB	1057	G	Sidechain
35	BB	1058	U	Sidechain
35	BB	1059	U	Sidechain
35	BB	1060	U	Sidechain
35	BB	1062	G	Sidechain
35	BB	1063	C	Sidechain
35	BB	1064	U	Sidechain
35	BB	1066	G	Sidechain
35	BB	1068	G	Sidechain
35	BB	1070	G	Sidechain
35	BB	1071	G	Sidechain
35	BB	1072	C	Sidechain
35	BB	1073	A	Sidechain
35	BB	1074	U	Sidechain
35	BB	1075	A	Sidechain
35	BB	1078	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1079	G	Sidechain
35	BB	108	G	Sidechain
35	BB	1081	U	Sidechain
35	BB	1082	A	Sidechain
35	BB	1083	C	Sidechain
35	BB	1084	A	Sidechain
35	BB	1085	C	Sidechain
35	BB	1086	G	Sidechain
35	BB	1087	A	Sidechain
35	BB	1088	C	Sidechain
35	BB	109	U	Sidechain
35	BB	1090	A	Sidechain
35	BB	1092	G	Sidechain
35	BB	1093	C	Sidechain
35	BB	1094	A	Sidechain
35	BB	1095	G	Sidechain
35	BB	1097	U	Sidechain
35	BB	1098	G	Sidechain
35	BB	1099	U	Sidechain
35	BB	110	U	Sidechain
35	BB	1100	C	Sidechain
35	BB	1102	U	Sidechain
35	BB	1103	A	Sidechain
35	BB	1104	A	Sidechain
35	BB	1105	G	Sidechain
35	BB	1106	G	Sidechain
35	BB	1107	C	Sidechain
35	BB	1108	G	Sidechain
35	BB	1109	A	Sidechain
35	BB	111	C	Sidechain
35	BB	1110	G	Sidechain
35	BB	1112	U	Sidechain
35	BB	1114	A	Sidechain
35	BB	1115	G	Sidechain
35	BB	1116	U	Sidechain
35	BB	112	G	Sidechain
35	BB	1120	A	Sidechain
35	BB	1122	C	Sidechain
35	BB	1123	A	Sidechain
35	BB	1124	G	Sidechain
35	BB	1125	A	Sidechain
35	BB	1126	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1128	U	Sidechain
35	BB	1129	C	Sidechain
35	BB	1130	U	Sidechain
35	BB	1131	C	Sidechain
35	BB	1132	A	Sidechain
35	BB	1134	G	Sidechain
35	BB	1136	G	Sidechain
35	BB	1137	G	Sidechain
35	BB	1138	A	Sidechain
35	BB	114	A	Sidechain
35	BB	1140	C	Sidechain
35	BB	1142	C	Sidechain
35	BB	1143	A	Sidechain
35	BB	1144	A	Sidechain
35	BB	1145	G	Sidechain
35	BB	1148	U	Sidechain
35	BB	1149	A	Sidechain
35	BB	115	A	Sidechain
35	BB	1151	A	Sidechain
35	BB	1152	U	Sidechain
35	BB	1153	G	Sidechain
35	BB	1154	C	Sidechain
35	BB	1155	U	Sidechain
35	BB	1156	U	Sidechain
35	BB	1159	U	Sidechain
35	BB	116	G	Sidechain
35	BB	1160	U	Sidechain
35	BB	1161	G	Sidechain
35	BB	1163	U	Sidechain
35	BB	1164	U	Sidechain
35	BB	1165	A	Sidechain
35	BB	1166	A	Sidechain
35	BB	1167	C	Sidechain
35	BB	1168	G	Sidechain
35	BB	1169	A	Sidechain
35	BB	1170	U	Sidechain
35	BB	1172	U	Sidechain
35	BB	1174	C	Sidechain
35	BB	1175	A	Sidechain
35	BB	1177	U	Sidechain
35	BB	1178	A	Sidechain
35	BB	1180	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1181	A	Sidechain
35	BB	1183	U	Sidechain
35	BB	1184	C	Sidechain
35	BB	1185	G	Sidechain
35	BB	1186	A	Sidechain
35	BB	1187	G	Sidechain
35	BB	119	G	Sidechain
35	BB	1190	U	Sidechain
35	BB	1192	C	Sidechain
35	BB	1193	G	Sidechain
35	BB	1194	A	Sidechain
35	BB	1195	A	Sidechain
35	BB	1196	A	Sidechain
35	BB	1197	G	Sidechain
35	BB	12	G	Sidechain
35	BB	1200	A	Sidechain
35	BB	1201	G	Sidechain
35	BB	1202	G	Sidechain
35	BB	1203	C	Sidechain
35	BB	1205	A	Sidechain
35	BB	1207	C	Sidechain
35	BB	1208	G	Sidechain
35	BB	1209	A	Sidechain
35	BB	121	A	Sidechain
35	BB	1210	U	Sidechain
35	BB	1211	C	Sidechain
35	BB	1212	C	Sidechain
35	BB	1213	U	Sidechain
35	BB	1214	U	Sidechain
35	BB	1215	U	Sidechain
35	BB	1216	G	Sidechain
35	BB	1217	C	Sidechain
35	BB	1218	G	Sidechain
35	BB	122	U	Sidechain
35	BB	1220	A	Sidechain
35	BB	1221	G	Sidechain
35	BB	1222	A	Sidechain
35	BB	1223	A	Sidechain
35	BB	1225	A	Sidechain
35	BB	1226	G	Sidechain
35	BB	1227	G	Sidechain
35	BB	1229	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	123	U	Sidechain
35	BB	1230	A	Sidechain
35	BB	1232	A	Sidechain
35	BB	1233	U	Sidechain
35	BB	1234	G	Sidechain
35	BB	1236	A	Sidechain
35	BB	1240	A	Sidechain
35	BB	1241	U	Sidechain
35	BB	1243	A	Sidechain
35	BB	1244	U	Sidechain
35	BB	1246	C	Sidechain
35	BB	1248	A	Sidechain
35	BB	1249	G	Sidechain
35	BB	125	G	Sidechain
35	BB	1251	G	Sidechain
35	BB	1252	G	Sidechain
35	BB	1253	U	Sidechain
35	BB	1254	G	Sidechain
35	BB	1256	C	Sidechain
35	BB	1257	A	Sidechain
35	BB	1258	G	Sidechain
35	BB	1259	A	Sidechain
35	BB	1261	U	Sidechain
35	BB	1262	A	Sidechain
35	BB	1265	U	Sidechain
35	BB	1266	A	Sidechain
35	BB	1267	C	Sidechain
35	BB	127	U	Sidechain
35	BB	1270	C	Sidechain
35	BB	1271	A	Sidechain
35	BB	1272	G	Sidechain
35	BB	1273	G	Sidechain
35	BB	1274	G	Sidechain
35	BB	1276	U	Sidechain
35	BB	1279	C	Sidechain
35	BB	128	C	Sidechain
35	BB	1280	U	Sidechain
35	BB	1281	G	Sidechain
35	BB	1282	G	Sidechain
35	BB	1283	C	Sidechain
35	BB	1284	U	Sidechain
35	BB	1285	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1286	G	Sidechain
35	BB	1287	U	Sidechain
35	BB	1289	G	Sidechain
35	BB	129	U	Sidechain
35	BB	1290	C	Sidechain
35	BB	1291	G	Sidechain
35	BB	1292	G	Sidechain
35	BB	1293	C	Sidechain
35	BB	1295	A	Sidechain
35	BB	1297	G	Sidechain
35	BB	1299	G	Sidechain
35	BB	13	A	Sidechain
35	BB	1300	U	Sidechain
35	BB	1301	U	Sidechain
35	BB	1302	C	Sidechain
35	BB	1304	U	Sidechain
35	BB	1305	A	Sidechain
35	BB	1307	C	Sidechain
35	BB	1310	C	Sidechain
35	BB	1312	U	Sidechain
35	BB	1315	C	Sidechain
35	BB	1316	U	Sidechain
35	BB	1317	U	Sidechain
35	BB	1318	U	Sidechain
35	BB	1319	U	Sidechain
35	BB	132	G	Sidechain
35	BB	1320	U	Sidechain
35	BB	1321	G	Sidechain
35	BB	1323	U	Sidechain
35	BB	1324	C	Sidechain
35	BB	1326	U	Sidechain
35	BB	1327	U	Sidechain
35	BB	133	G	Sidechain
35	BB	1330	A	Sidechain
35	BB	1331	U	Sidechain
35	BB	1332	G	Sidechain
35	BB	1333	U	Sidechain
35	BB	1334	C	Sidechain
35	BB	1335	G	Sidechain
35	BB	1337	C	Sidechain
35	BB	1338	U	Sidechain
35	BB	1339	C	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1340	U	Sidechain
35	BB	1341	U	Sidechain
35	BB	1342	C	Sidechain
35	BB	1343	C	Sidechain
35	BB	1345	A	Sidechain
35	BB	1346	A	Sidechain
35	BB	1348	C	Sidechain
35	BB	135	C	Sidechain
35	BB	1350	A	Sidechain
35	BB	1353	G	Sidechain
35	BB	1354	C	Sidechain
35	BB	1356	G	Sidechain
35	BB	1357	C	Sidechain
35	BB	1360	A	Sidechain
35	BB	1361	A	Sidechain
35	BB	1362	G	Sidechain
35	BB	1364	C	Sidechain
35	BB	1367	U	Sidechain
35	BB	1368	A	Sidechain
35	BB	1369	A	Sidechain
35	BB	137	A	Sidechain
35	BB	1370	G	Sidechain
35	BB	1372	G	Sidechain
35	BB	1374	U	Sidechain
35	BB	1375	G	Sidechain
35	BB	1376	G	Sidechain
35	BB	1377	A	Sidechain
35	BB	1378	U	Sidechain
35	BB	1379	U	Sidechain
35	BB	138	A	Sidechain
35	BB	1381	U	Sidechain
35	BB	1382	U	Sidechain
35	BB	1383	C	Sidechain
35	BB	1384	A	Sidechain
35	BB	1385	C	Sidechain
35	BB	1386	C	Sidechain
35	BB	1387	C	Sidechain
35	BB	1388	A	Sidechain
35	BB	139	G	Sidechain
35	BB	1390	U	Sidechain
35	BB	1391	G	Sidechain
35	BB	1392	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1393	C	Sidechain
35	BB	1394	A	Sidechain
35	BB	1395	G	Sidechain
35	BB	1396	G	Sidechain
35	BB	1397	G	Sidechain
35	BB	1399	A	Sidechain
35	BB	14	C	Sidechain
35	BB	1400	C	Sidechain
35	BB	1401	G	Sidechain
35	BB	1402	U	Sidechain
35	BB	1403	G	Sidechain
35	BB	1404	A	Sidechain
35	BB	1406	C	Sidechain
35	BB	1407	U	Sidechain
35	BB	1408	G	Sidechain
35	BB	1409	G	Sidechain
35	BB	1410	G	Sidechain
35	BB	1411	U	Sidechain
35	BB	1413	U	Sidechain
35	BB	1415	G	Sidechain
35	BB	1416	A	Sidechain
35	BB	1417	C	Sidechain
35	BB	1418	C	Sidechain
35	BB	1419	G	Sidechain
35	BB	142	G	Sidechain
35	BB	1420	U	Sidechain
35	BB	1422	G	Sidechain
35	BB	1423	U	Sidechain
35	BB	1424	G	Sidechain
35	BB	1425	A	Sidechain
35	BB	1426	G	Sidechain
35	BB	1427	A	Sidechain
35	BB	1429	A	Sidechain
35	BB	143	G	Sidechain
35	BB	1430	G	Sidechain
35	BB	1432	U	Sidechain
35	BB	1433	U	Sidechain
35	BB	1434	G	Sidechain
35	BB	1435	G	Sidechain
35	BB	1436	U	Sidechain
35	BB	1437	U	Sidechain
35	BB	1438	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1439	U	Sidechain
35	BB	144	G	Sidechain
35	BB	1441	C	Sidechain
35	BB	1442	C	Sidechain
35	BB	1443	C	Sidechain
35	BB	1445	A	Sidechain
35	BB	1446	C	Sidechain
35	BB	1447	U	Sidechain
35	BB	1448	U	Sidechain
35	BB	1452	U	Sidechain
35	BB	1453	G	Sidechain
35	BB	1454	G	Sidechain
35	BB	1455	A	Sidechain
35	BB	1456	G	Sidechain
35	BB	1458	U	Sidechain
35	BB	1459	U	Sidechain
35	BB	1461	C	Sidechain
35	BB	1462	G	Sidechain
35	BB	1464	G	Sidechain
35	BB	1465	U	Sidechain
35	BB	1466	A	Sidechain
35	BB	1467	A	Sidechain
35	BB	1468	A	Sidechain
35	BB	1469	A	Sidechain
35	BB	1470	G	Sidechain
35	BB	1472	U	Sidechain
35	BB	1473	U	Sidechain
35	BB	1474	A	Sidechain
35	BB	1475	U	Sidechain
35	BB	1477	C	Sidechain
35	BB	1478	G	Sidechain
35	BB	148	C	Sidechain
35	BB	1480	G	Sidechain
35	BB	1481	C	Sidechain
35	BB	1482	A	Sidechain
35	BB	1484	A	Sidechain
35	BB	1485	G	Sidechain
35	BB	1487	G	Sidechain
35	BB	1488	G	Sidechain
35	BB	1489	A	Sidechain
35	BB	149	A	Sidechain
35	BB	1490	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	1492	C	Sidechain
35	BB	1494	G	Sidechain
35	BB	1495	U	Sidechain
35	BB	1496	C	Sidechain
35	BB	1497	C	Sidechain
35	BB	1498	G	Sidechain
35	BB	15	C	Sidechain
35	BB	1500	U	Sidechain
35	BB	1501	U	Sidechain
35	BB	1502	U	Sidechain
35	BB	1503	U	Sidechain
35	BB	1505	U	Sidechain
35	BB	1506	C	Sidechain
35	BB	1507	U	Sidechain
35	BB	1508	G	Sidechain
35	BB	1509	G	Sidechain
35	BB	1510	G	Sidechain
35	BB	1511	U	Sidechain
35	BB	1512	C	Sidechain
35	BB	1513	U	Sidechain
35	BB	1517	G	Sidechain
35	BB	1518	U	Sidechain
35	BB	152	G	Sidechain
35	BB	1520	C	Sidechain
35	BB	1521	G	Sidechain
35	BB	1523	U	Sidechain
35	BB	1527	A	Sidechain
35	BB	1528	U	Sidechain
35	BB	1529	G	Sidechain
35	BB	1530	U	Sidechain
35	BB	1532	C	Sidechain
35	BB	1533	U	Sidechain
35	BB	1534	U	Sidechain
35	BB	1538	G	Sidechain
35	BB	1539	C	Sidechain
35	BB	154	A	Sidechain
35	BB	1541	G	Sidechain
35	BB	1542	C	Sidechain
35	BB	1543	C	Sidechain
35	BB	1547	U	Sidechain
35	BB	1548	C	Sidechain
35	BB	156	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	157	G	Sidechain
35	BB	16	G	Sidechain
35	BB	163	G	Sidechain
35	BB	164	U	Sidechain
35	BB	168	U	Sidechain
35	BB	17	U	Sidechain
35	BB	2	C	Sidechain
35	BB	20	U	Sidechain
35	BB	21	C	Sidechain
35	BB	23	U	Sidechain
35	BB	24	C	Sidechain
35	BB	25	A	Sidechain
35	BB	256	G	Sidechain
35	BB	265	C	Sidechain
35	BB	266	C	Sidechain
35	BB	267	C	Sidechain
35	BB	27	C	Sidechain
35	BB	273	G	Sidechain
35	BB	274	U	Sidechain
35	BB	278	U	Sidechain
35	BB	279	A	Sidechain
35	BB	281	U	Sidechain
35	BB	285	C	Sidechain
35	BB	289	U	Sidechain
35	BB	29	C	Sidechain
35	BB	292	U	Sidechain
35	BB	298	G	Sidechain
35	BB	3	C	Sidechain
35	BB	30	A	Sidechain
35	BB	305	U	Sidechain
35	BB	307	A	Sidechain
35	BB	31	U	Sidechain
35	BB	311	C	Sidechain
35	BB	314	A	Sidechain
35	BB	315	C	Sidechain
35	BB	317	C	Sidechain
35	BB	32	C	Sidechain
35	BB	323	C	Sidechain
35	BB	325	G	Sidechain
35	BB	326	G	Sidechain
35	BB	328	G	Sidechain
35	BB	33	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	332	U	Sidechain
35	BB	339	C	Sidechain
35	BB	34	G	Sidechain
35	BB	341	U	Sidechain
35	BB	345	U	Sidechain
35	BB	348	G	Sidechain
35	BB	353	G	Sidechain
35	BB	357	C	Sidechain
35	BB	358	U	Sidechain
35	BB	359	A	Sidechain
35	BB	36	U	Sidechain
35	BB	361	A	Sidechain
35	BB	362	A	Sidechain
35	BB	364	U	Sidechain
35	BB	365	U	Sidechain
35	BB	366	G	Sidechain
35	BB	367	C	Sidechain
35	BB	368	C	Sidechain
35	BB	369	A	Sidechain
35	BB	37	C	Sidechain
35	BB	370	A	Sidechain
35	BB	371	C	Sidechain
35	BB	372	U	Sidechain
35	BB	373	C	Sidechain
35	BB	374	A	Sidechain
35	BB	375	G	Sidechain
35	BB	376	A	Sidechain
35	BB	377	A	Sidechain
35	BB	38	C	Sidechain
35	BB	380	G	Sidechain
35	BB	382	U	Sidechain
35	BB	383	U	Sidechain
35	BB	384	A	Sidechain
35	BB	386	G	Sidechain
35	BB	387	G	Sidechain
35	BB	388	C	Sidechain
35	BB	389	G	Sidechain
35	BB	39	C	Sidechain
35	BB	390	G	Sidechain
35	BB	391	G	Sidechain
35	BB	392	G	Sidechain
35	BB	393	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	395	U	Sidechain
35	BB	396	C	Sidechain
35	BB	398	A	Sidechain
35	BB	399	A	Sidechain
35	BB	40	C	Sidechain
35	BB	400	C	Sidechain
35	BB	401	U	Sidechain
35	BB	402	G	Sidechain
35	BB	403	U	Sidechain
35	BB	404	A	Sidechain
35	BB	405	U	Sidechain
35	BB	406	A	Sidechain
35	BB	407	A	Sidechain
35	BB	409	U	Sidechain
35	BB	41	A	Sidechain
35	BB	413	A	Sidechain
35	BB	414	C	Sidechain
35	BB	415	A	Sidechain
35	BB	416	U	Sidechain
35	BB	417	A	Sidechain
35	BB	418	G	Sidechain
35	BB	419	G	Sidechain
35	BB	42	A	Sidechain
35	BB	420	U	Sidechain
35	BB	421	U	Sidechain
35	BB	423	G	Sidechain
35	BB	424	U	Sidechain
35	BB	425	G	Sidechain
35	BB	427	U	Sidechain
35	BB	428	G	Sidechain
35	BB	429	C	Sidechain
35	BB	43	G	Sidechain
35	BB	431	U	Sidechain
35	BB	432	C	Sidechain
35	BB	433	C	Sidechain
35	BB	434	A	Sidechain
35	BB	435	A	Sidechain
35	BB	436	G	Sidechain
35	BB	437	U	Sidechain
35	BB	439	G	Sidechain
35	BB	440	U	Sidechain
35	BB	441	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	442	U	Sidechain
35	BB	443	A	Sidechain
35	BB	444	U	Sidechain
35	BB	445	G	Sidechain
35	BB	448	G	Sidechain
35	BB	449	C	Sidechain
35	BB	45	A	Sidechain
35	BB	450	A	Sidechain
35	BB	451	A	Sidechain
35	BB	452	A	Sidechain
35	BB	454	U	Sidechain
35	BB	455	G	Sidechain
35	BB	456	A	Sidechain
35	BB	457	U	Sidechain
35	BB	458	U	Sidechain
35	BB	459	U	Sidechain
35	BB	46	U	Sidechain
35	BB	460	C	Sidechain
35	BB	461	U	Sidechain
35	BB	462	G	Sidechain
35	BB	464	C	Sidechain
35	BB	466	A	Sidechain
35	BB	469	G	Sidechain
35	BB	47	C	Sidechain
35	BB	471	U	Sidechain
35	BB	472	C	Sidechain
35	BB	473	U	Sidechain
35	BB	474	G	Sidechain
35	BB	475	A	Sidechain
35	BB	476	A	Sidechain
35	BB	477	U	Sidechain
35	BB	478	G	Sidechain
35	BB	48	G	Sidechain
35	BB	481	A	Sidechain
35	BB	484	G	Sidechain
35	BB	486	G	Sidechain
35	BB	487	A	Sidechain
35	BB	488	G	Sidechain
35	BB	489	A	Sidechain
35	BB	490	G	Sidechain
35	BB	491	A	Sidechain
35	BB	492	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	493	U	Sidechain
35	BB	497	C	Sidechain
35	BB	498	G	Sidechain
35	BB	499	A	Sidechain
35	BB	5	A	Sidechain
35	BB	50	A	Sidechain
35	BB	502	C	Sidechain
35	BB	503	G	Sidechain
35	BB	504	C	Sidechain
35	BB	507	G	Sidechain
35	BB	508	U	Sidechain
35	BB	51	U	Sidechain
35	BB	510	A	Sidechain
35	BB	512	C	Sidechain
35	BB	514	G	Sidechain
35	BB	515	C	Sidechain
35	BB	516	G	Sidechain
35	BB	517	G	Sidechain
35	BB	519	A	Sidechain
35	BB	52	G	Sidechain
35	BB	522	A	Sidechain
35	BB	524	C	Sidechain
35	BB	525	U	Sidechain
35	BB	526	A	Sidechain
35	BB	528	G	Sidechain
35	BB	53	C	Sidechain
35	BB	530	C	Sidechain
35	BB	533	U	Sidechain
35	BB	534	C	Sidechain
35	BB	536	U	Sidechain
35	BB	537	A	Sidechain
35	BB	538	A	Sidechain
35	BB	539	G	Sidechain
35	BB	54	U	Sidechain
35	BB	540	G	Sidechain
35	BB	544	C	Sidechain
35	BB	545	C	Sidechain
35	BB	546	A	Sidechain
35	BB	547	A	Sidechain
35	BB	549	U	Sidechain
35	BB	550	G	Sidechain
35	BB	551	C	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	553	U	Sidechain
35	BB	555	G	Sidechain
35	BB	556	U	Sidechain
35	BB	557	C	Sidechain
35	BB	558	U	Sidechain
35	BB	559	U	Sidechain
35	BB	56	U	Sidechain
35	BB	560	C	Sidechain
35	BB	561	C	Sidechain
35	BB	563	A	Sidechain
35	BB	564	U	Sidechain
35	BB	565	U	Sidechain
35	BB	57	G	Sidechain
35	BB	571	C	Sidechain
35	BB	572	G	Sidechain
35	BB	574	G	Sidechain
35	BB	575	C	Sidechain
35	BB	576	A	Sidechain
35	BB	577	U	Sidechain
35	BB	578	G	Sidechain
35	BB	579	A	Sidechain
35	BB	58	G	Sidechain
35	BB	580	A	Sidechain
35	BB	581	U	Sidechain
35	BB	582	G	Sidechain
35	BB	583	G	Sidechain
35	BB	585	U	Sidechain
35	BB	587	A	Sidechain
35	BB	588	A	Sidechain
35	BB	589	U	Sidechain
35	BB	59	U	Sidechain
35	BB	590	G	Sidechain
35	BB	591	A	Sidechain
35	BB	592	G	Sidechain
35	BB	593	A	Sidechain
35	BB	594	U	Sidechain
35	BB	595	U	Sidechain
35	BB	597	C	Sidechain
35	BB	598	C	Sidechain
35	BB	599	U	Sidechain
35	BB	6	A	Sidechain
35	BB	60	A	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	600	C	Sidechain
35	BB	601	U	Sidechain
35	BB	603	U	Sidechain
35	BB	604	C	Sidechain
35	BB	606	C	Sidechain
35	BB	607	G	Sidechain
35	BB	608	A	Sidechain
35	BB	61	A	Sidechain
35	BB	610	U	Sidechain
35	BB	611	U	Sidechain
35	BB	612	A	Sidechain
35	BB	614	U	Sidechain
35	BB	615	A	Sidechain
35	BB	616	U	Sidechain
35	BB	618	U	Sidechain
35	BB	62	C	Sidechain
35	BB	621	C	Sidechain
35	BB	622	G	Sidechain
35	BB	623	A	Sidechain
35	BB	624	A	Sidechain
35	BB	626	C	Sidechain
35	BB	628	A	Sidechain
35	BB	629	C	Sidechain
35	BB	631	G	Sidechain
35	BB	632	U	Sidechain
35	BB	633	C	Sidechain
35	BB	634	A	Sidechain
35	BB	638	G	Sidechain
35	BB	639	A	Sidechain
35	BB	64	U	Sidechain
35	BB	641	C	Sidechain
35	BB	643	G	Sidechain
35	BB	644	A	Sidechain
35	BB	645	C	Sidechain
35	BB	646	U	Sidechain
35	BB	647	U	Sidechain
35	BB	649	A	Sidechain
35	BB	650	A	Sidechain
35	BB	651	G	Sidechain
35	BB	652	G	Sidechain
35	BB	653	G	Sidechain
35	BB	654	C	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	655	U	Sidechain
35	BB	658	G	Sidechain
35	BB	659	C	Sidechain
35	BB	66	G	Sidechain
35	BB	660	G	Sidechain
35	BB	661	G	Sidechain
35	BB	663	G	Sidechain
35	BB	664	A	Sidechain
35	BB	665	A	Sidechain
35	BB	666	A	Sidechain
35	BB	667	G	Sidechain
35	BB	668	A	Sidechain
35	BB	67	A	Sidechain
35	BB	670	G	Sidechain
35	BB	671	A	Sidechain
35	BB	672	C	Sidechain
35	BB	674	C	Sidechain
35	BB	675	U	Sidechain
35	BB	676	G	Sidechain
35	BB	677	U	Sidechain
35	BB	678	U	Sidechain
35	BB	680	A	Sidechain
35	BB	681	G	Sidechain
35	BB	682	U	Sidechain
35	BB	683	U	Sidechain
35	BB	684	U	Sidechain
35	BB	685	G	Sidechain
35	BB	686	A	Sidechain
35	BB	687	C	Sidechain
35	BB	688	U	Sidechain
35	BB	69	A	Sidechain
35	BB	692	G	Sidechain
35	BB	693	U	Sidechain
35	BB	694	C	Sidechain
35	BB	695	U	Sidechain
35	BB	696	G	Sidechain
35	BB	697	G	Sidechain
35	BB	698	C	Sidechain
35	BB	699	U	Sidechain
35	BB	70	A	Sidechain
35	BB	700	C	Sidechain
35	BB	701	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	702	G	Sidechain
35	BB	703	U	Sidechain
35	BB	706	G	Sidechain
35	BB	709	G	Sidechain
35	BB	71	A	Sidechain
35	BB	711	C	Sidechain
35	BB	713	U	Sidechain
35	BB	714	U	Sidechain
35	BB	715	G	Sidechain
35	BB	716	G	Sidechain
35	BB	717	A	Sidechain
35	BB	719	G	Sidechain
35	BB	72	G	Sidechain
35	BB	723	A	Sidechain
35	BB	724	G	Sidechain
35	BB	726	A	Sidechain
35	BB	728	A	Sidechain
35	BB	729	G	Sidechain
35	BB	734	A	Sidechain
35	BB	736	G	Sidechain
35	BB	737	C	Sidechain
35	BB	738	G	Sidechain
35	BB	74	U	Sidechain
35	BB	741	A	Sidechain
35	BB	744	U	Sidechain
35	BB	746	A	Sidechain
35	BB	748	A	Sidechain
35	BB	749	U	Sidechain
35	BB	750	G	Sidechain
35	BB	751	A	Sidechain
35	BB	753	A	Sidechain
35	BB	754	U	Sidechain
35	BB	756	C	Sidechain
35	BB	757	C	Sidechain
35	BB	760	C	Sidechain
35	BB	762	C	Sidechain
35	BB	763	U	Sidechain
35	BB	764	C	Sidechain
35	BB	766	G	Sidechain
35	BB	767	A	Sidechain
35	BB	768	A	Sidechain
35	BB	769	C	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	770	G	Sidechain
35	BB	772	U	Sidechain
35	BB	773	G	Sidechain
35	BB	774	C	Sidechain
35	BB	775	U	Sidechain
35	BB	776	U	Sidechain
35	BB	777	C	Sidechain
35	BB	778	A	Sidechain
35	BB	779	C	Sidechain
35	BB	780	U	Sidechain
35	BB	781	U	Sidechain
35	BB	782	A	Sidechain
35	BB	783	U	Sidechain
35	BB	785	G	Sidechain
35	BB	786	A	Sidechain
35	BB	788	U	Sidechain
35	BB	789	G	Sidechain
35	BB	790	A	Sidechain
35	BB	791	A	Sidechain
35	BB	792	G	Sidechain
35	BB	793	A	Sidechain
35	BB	794	G	Sidechain
35	BB	795	A	Sidechain
35	BB	797	C	Sidechain
35	BB	798	A	Sidechain
35	BB	799	A	Sidechain
35	BB	8	U	Sidechain
35	BB	80	C	Sidechain
35	BB	800	U	Sidechain
35	BB	801	G	Sidechain
35	BB	802	G	Sidechain
35	BB	803	U	Sidechain
35	BB	804	U	Sidechain
35	BB	805	G	Sidechain
35	BB	807	U	Sidechain
35	BB	808	U	Sidechain
35	BB	809	U	Sidechain
35	BB	81	A	Sidechain
35	BB	810	G	Sidechain
35	BB	812	G	Sidechain
35	BB	815	G	Sidechain
35	BB	816	U	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	818	U	Sidechain
35	BB	822	G	Sidechain
35	BB	825	U	Sidechain
35	BB	826	G	Sidechain
35	BB	827	U	Sidechain
35	BB	828	G	Sidechain
35	BB	83	G	Sidechain
35	BB	830	G	Sidechain
35	BB	831	C	Sidechain
35	BB	832	C	Sidechain
35	BB	833	G	Sidechain
35	BB	834	U	Sidechain
35	BB	835	C	Sidechain
35	BB	836	U	Sidechain
35	BB	837	A	Sidechain
35	BB	838	G	Sidechain
35	BB	84	G	Sidechain
35	BB	840	C	Sidechain
35	BB	842	G	Sidechain
35	BB	843	G	Sidechain
35	BB	844	G	Sidechain
35	BB	845	C	Sidechain
35	BB	846	A	Sidechain
35	BB	847	U	Sidechain
35	BB	85	A	Sidechain
35	BB	850	U	Sidechain
35	BB	851	U	Sidechain
35	BB	852	G	Sidechain
35	BB	853	U	Sidechain
35	BB	854	G	Sidechain
35	BB	855	G	Sidechain
35	BB	856	U	Sidechain
35	BB	859	U	Sidechain
35	BB	860	U	Sidechain
35	BB	862	U	Sidechain
35	BB	864	U	Sidechain
35	BB	866	A	Sidechain
35	BB	868	C	Sidechain
35	BB	869	G	Sidechain
35	BB	870	C	Sidechain
35	BB	877	A	Sidechain
35	BB	878	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	879	G	Sidechain
35	BB	88	U	Sidechain
35	BB	881	G	Sidechain
35	BB	882	U	Sidechain
35	BB	884	U	Sidechain
35	BB	885	U	Sidechain
35	BB	886	G	Sidechain
35	BB	891	U	Sidechain
35	BB	893	U	Sidechain
35	BB	894	A	Sidechain
35	BB	895	U	Sidechain
35	BB	896	C	Sidechain
35	BB	897	C	Sidechain
35	BB	898	U	Sidechain
35	BB	899	C	Sidechain
35	BB	90	G	Sidechain
35	BB	901	U	Sidechain
35	BB	903	U	Sidechain
35	BB	91	G	Sidechain
35	BB	910	C	Sidechain
35	BB	915	U	Sidechain
35	BB	93	A	Sidechain
35	BB	931	U	Sidechain
35	BB	934	U	Sidechain
35	BB	94	A	Sidechain
35	BB	943	U	Sidechain
35	BB	95	A	Sidechain
35	BB	953	G	Sidechain
35	BB	956	G	Sidechain
35	BB	958	C	Sidechain
35	BB	959	C	Sidechain
35	BB	96	A	Sidechain
35	BB	964	G	Sidechain
35	BB	965	G	Sidechain
35	BB	966	C	Sidechain
35	BB	967	G	Sidechain
35	BB	968	C	Sidechain
35	BB	97	U	Sidechain
35	BB	971	A	Sidechain
35	BB	972	C	Sidechain
35	BB	973	G	Sidechain
35	BB	975	G	Sidechain

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Mol	Chain	Res	Type	Group
35	BB	976	U	Sidechain
35	BB	977	G	Sidechain
35	BB	978	C	Sidechain
35	BB	979	G	Sidechain
35	BB	98	A	Sidechain
35	BB	981	A	Sidechain
35	BB	984	U	Sidechain
35	BB	986	C	Sidechain
35	BB	987	U	Sidechain
35	BB	988	G	Sidechain
35	BB	989	C	Sidechain
35	BB	99	G	Sidechain
35	BB	991	C	Sidechain
35	BB	992	C	Sidechain
35	BB	993	A	Sidechain
35	BB	994	A	Sidechain
35	BB	995	C	Sidechain
35	BB	996	G	Sidechain
35	BB	997	G	Sidechain
35	BB	998	G	Sidechain
36	BC	1	A	Sidechain
36	BC	100	U	Sidechain
36	BC	101	U	Sidechain
36	BC	102	G	Sidechain
36	BC	103	A	Sidechain
36	BC	104	A	Sidechain
36	BC	106	G	Sidechain
36	BC	107	C	Sidechain
36	BC	109	A	Sidechain
36	BC	11	G	Sidechain
36	BC	111	C	Sidechain
36	BC	119	G	Sidechain
36	BC	12	A	Sidechain
36	BC	120	G	Sidechain
36	BC	121	G	Sidechain
36	BC	123	G	Sidechain
36	BC	125	A	Sidechain
36	BC	126	G	Sidechain
36	BC	128	U	Sidechain
36	BC	129	C	Sidechain
36	BC	13	U	Sidechain
36	BC	130	U	Sidechain

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Mol	Chain	Res	Type	Group
36	BC	131	C	Sidechain
36	BC	134	G	Sidechain
36	BC	136	G	Sidechain
36	BC	138	C	Sidechain
36	BC	139	A	Sidechain
36	BC	14	G	Sidechain
36	BC	140	U	Sidechain
36	BC	141	C	Sidechain
36	BC	142	C	Sidechain
36	BC	143	C	Sidechain
36	BC	144	C	Sidechain
36	BC	145	G	Sidechain
36	BC	146	U	Sidechain
36	BC	147	G	Sidechain
36	BC	148	C	Sidechain
36	BC	149	A	Sidechain
36	BC	15	G	Sidechain
36	BC	150	U	Sidechain
36	BC	151	G	Sidechain
36	BC	152	C	Sidechain
36	BC	154	A	Sidechain
36	BC	155	C	Sidechain
36	BC	156	A	Sidechain
36	BC	157	U	Sidechain
36	BC	158	U	Sidechain
36	BC	159	U	Sidechain
36	BC	16	A	Sidechain
36	BC	160	C	Sidechain
36	BC	163	A	Sidechain
36	BC	164	G	Sidechain
36	BC	165	U	Sidechain
36	BC	166	G	Sidechain
36	BC	167	U	Sidechain
36	BC	168	C	Sidechain
36	BC	169	G	Sidechain
36	BC	17	U	Sidechain
36	BC	18	G	Sidechain
36	BC	19	A	Sidechain
36	BC	2	A	Sidechain
36	BC	20	C	Sidechain
36	BC	21	U	Sidechain
36	BC	22	U	Sidechain

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Mol	Chain	Res	Type	Group
36	BC	23	G	Sidechain
36	BC	25	C	Sidechain
36	BC	26	U	Sidechain
36	BC	27	U	Sidechain
36	BC	28	C	Sidechain
36	BC	3	C	Sidechain
36	BC	30	U	Sidechain
36	BC	31	A	Sidechain
36	BC	33	U	Sidechain
36	BC	34	U	Sidechain
36	BC	36	G	Sidechain
36	BC	37	U	Sidechain
36	BC	38	U	Sidechain
36	BC	4	G	Sidechain
36	BC	40	A	Sidechain
36	BC	41	A	Sidechain
36	BC	42	G	Sidechain
36	BC	43	A	Sidechain
36	BC	44	A	Sidechain
36	BC	45	C	Sidechain
36	BC	47	C	Sidechain
36	BC	48	A	Sidechain
36	BC	49	G	Sidechain
36	BC	5	U	Sidechain
36	BC	50	C	Sidechain
36	BC	51	A	Sidechain
36	BC	53	A	Sidechain
36	BC	54	G	Sidechain
36	BC	56	G	Sidechain
36	BC	58	G	Sidechain
36	BC	59	A	Sidechain
36	BC	6	G	Sidechain
36	BC	60	U	Sidechain
36	BC	61	A	Sidechain
36	BC	62	A	Sidechain
36	BC	63	G	Sidechain
36	BC	64	U	Sidechain
36	BC	68	A	Sidechain
36	BC	69	U	Sidechain
36	BC	7	U	Sidechain
36	BC	70	C	Sidechain
36	BC	71	A	Sidechain

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Mol	Chain	Res	Type	Group
36	BC	74	U	Sidechain
36	BC	75	G	Sidechain
36	BC	76	C	Sidechain
36	BC	78	G	Sidechain
36	BC	79	A	Sidechain
36	BC	8	C	Sidechain
36	BC	80	A	Sidechain
36	BC	81	U	Sidechain
36	BC	82	C	Sidechain
36	BC	83	A	Sidechain
36	BC	84	U	Sidechain
36	BC	86	U	Sidechain
36	BC	88	A	Sidechain
36	BC	89	U	Sidechain
36	BC	9	G	Sidechain
36	BC	90	U	Sidechain
36	BC	91	G	Sidechain
36	BC	92	C	Sidechain
36	BC	94	C	Sidechain
36	BC	95	A	Sidechain
36	BC	97	U	Sidechain
36	BC	99	U	Sidechain
37	BD	1	G	Sidechain
37	BD	100	A	Sidechain
37	BD	101	A	Sidechain
37	BD	102	C	Sidechain
37	BD	103	C	Sidechain
37	BD	104	C	Sidechain
37	BD	105	G	Sidechain
37	BD	106	G	Sidechain
37	BD	108	G	Sidechain
37	BD	109	U	Sidechain
37	BD	11	A	Sidechain
37	BD	110	G	Sidechain
37	BD	112	U	Sidechain
37	BD	113	G	Sidechain
37	BD	117	U	Sidechain
37	BD	118	C	Sidechain
37	BD	119	U	Sidechain
37	BD	12	U	Sidechain
37	BD	14	C	Sidechain
37	BD	15	U	Sidechain

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Mol	Chain	Res	Type	Group
37	BD	16	U	Sidechain
37	BD	17	G	Sidechain
37	BD	18	G	Sidechain
37	BD	2	G	Sidechain
37	BD	22	A	Sidechain
37	BD	24	U	Sidechain
37	BD	25	G	Sidechain
37	BD	27	A	Sidechain
37	BD	28	C	Sidechain
37	BD	29	C	Sidechain
37	BD	3	G	Sidechain
37	BD	30	A	Sidechain
37	BD	31	U	Sidechain
37	BD	32	A	Sidechain
37	BD	34	C	Sidechain
37	BD	35	C	Sidechain
37	BD	36	C	Sidechain
37	BD	37	G	Sidechain
37	BD	38	U	Sidechain
37	BD	4	U	Sidechain
37	BD	41	G	Sidechain
37	BD	46	G	Sidechain
37	BD	47	U	Sidechain
37	BD	48	G	Sidechain
37	BD	49	A	Sidechain
37	BD	5	A	Sidechain
37	BD	52	U	Sidechain
37	BD	53	U	Sidechain
37	BD	54	A	Sidechain
37	BD	56	G	Sidechain
37	BD	57	C	Sidechain
37	BD	58	G	Sidechain
37	BD	61	C	Sidechain
37	BD	63	C	Sidechain
37	BD	64	A	Sidechain
37	BD	65	G	Sidechain
37	BD	66	G	Sidechain
37	BD	68	C	Sidechain
37	BD	69	U	Sidechain
37	BD	7	G	Sidechain
37	BD	70	C	Sidechain
37	BD	71	G	Sidechain

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Mol	Chain	Res	Type	Group
37	BD	72	U	Sidechain
37	BD	73	U	Sidechain
37	BD	74	A	Sidechain
37	BD	75	G	Sidechain
37	BD	76	U	Sidechain
37	BD	77	A	Sidechain
37	BD	78	C	Sidechain
37	BD	8	A	Sidechain
37	BD	80	G	Sidechain
37	BD	81	C	Sidechain
37	BD	82	G	Sidechain
37	BD	83	A	Sidechain
37	BD	84	U	Sidechain
37	BD	85	C	Sidechain
37	BD	86	A	Sidechain
37	BD	87	G	Sidechain
37	BD	88	U	Sidechain
37	BD	89	G	Sidechain
37	BD	9	C	Sidechain
37	BD	90	A	Sidechain
37	BD	92	G	Sidechain
37	BD	93	G	Sidechain
37	BD	94	C	Sidechain
37	BD	95	G	Sidechain
37	BD	96	C	Sidechain
37	BD	97	U	Sidechain
37	BD	99	G	Sidechain
38	BE	1	U	Sidechain
38	BE	10	G	Sidechain
38	BE	100	U	Sidechain
38	BE	101	C	Sidechain
38	BE	102	U	Sidechain
38	BE	104	G	Sidechain
38	BE	105	A	Sidechain
38	BE	106	C	Sidechain
38	BE	107	U	Sidechain
38	BE	108	U	Sidechain
38	BE	109	C	Sidechain
38	BE	11	A	Sidechain
38	BE	110	U	Sidechain
38	BE	111	C	Sidechain
38	BE	112	G	Sidechain

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Mol	Chain	Res	Type	Group
38	BE	113	C	Sidechain
38	BE	114	G	Sidechain
38	BE	115	U	Sidechain
38	BE	116	U	Sidechain
38	BE	117	A	Sidechain
38	BE	119	U	Sidechain
38	BE	120	C	Sidechain
38	BE	121	G	Sidechain
38	BE	122	G	Sidechain
38	BE	123	A	Sidechain
38	BE	124	G	Sidechain
38	BE	125	C	Sidechain
38	BE	126	G	Sidechain
38	BE	127	G	Sidechain
38	BE	128	G	Sidechain
38	BE	129	G	Sidechain
38	BE	13	A	Sidechain
38	BE	130	G	Sidechain
38	BE	132	U	Sidechain
38	BE	133	C	Sidechain
38	BE	134	A	Sidechain
38	BE	135	A	Sidechain
38	BE	136	G	Sidechain
38	BE	14	C	Sidechain
38	BE	140	G	Sidechain
38	BE	141	A	Sidechain
38	BE	142	A	Sidechain
38	BE	143	A	Sidechain
38	BE	145	A	Sidechain
38	BE	146	U	Sidechain
38	BE	147	G	Sidechain
38	BE	149	A	Sidechain
38	BE	15	A	Sidechain
38	BE	150	G	Sidechain
38	BE	152	U	Sidechain
38	BE	153	C	Sidechain
38	BE	154	A	Sidechain
38	BE	155	C	Sidechain
38	BE	158	U	Sidechain
38	BE	159	A	Sidechain
38	BE	160	C	Sidechain
38	BE	161	G	Sidechain

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Mol	Chain	Res	Type	Group
38	BE	163	A	Sidechain
38	BE	164	C	Sidechain
38	BE	165	U	Sidechain
38	BE	167	U	Sidechain
38	BE	168	C	Sidechain
38	BE	169	C	Sidechain
38	BE	17	U	Sidechain
38	BE	170	U	Sidechain
38	BE	171	U	Sidechain
38	BE	172	U	Sidechain
38	BE	173	G	Sidechain
38	BE	174	U	Sidechain
38	BE	175	U	Sidechain
38	BE	176	G	Sidechain
38	BE	177	U	Sidechain
38	BE	178	G	Sidechain
38	BE	179	A	Sidechain
38	BE	18	U	Sidechain
38	BE	180	G	Sidechain
38	BE	181	U	Sidechain
38	BE	182	U	Sidechain
38	BE	183	C	Sidechain
38	BE	184	G	Sidechain
38	BE	185	G	Sidechain
38	BE	186	C	Sidechain
38	BE	187	G	Sidechain
38	BE	188	C	Sidechain
38	BE	19	G	Sidechain
38	BE	190	U	Sidechain
38	BE	191	U	Sidechain
38	BE	192	A	Sidechain
38	BE	193	A	Sidechain
38	BE	194	A	Sidechain
38	BE	195	G	Sidechain
38	BE	196	C	Sidechain
38	BE	198	A	Sidechain
38	BE	20	C	Sidechain
38	BE	200	A	Sidechain
38	BE	201	A	Sidechain
38	BE	202	C	Sidechain
38	BE	203	C	Sidechain
38	BE	204	U	Sidechain

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Mol	Chain	Res	Type	Group
38	BE	205	G	Sidechain
38	BE	206	G	Sidechain
38	BE	207	G	Sidechain
38	BE	208	G	Sidechain
38	BE	21	C	Sidechain
38	BE	22	A	Sidechain
38	BE	23	G	Sidechain
38	BE	24	G	Sidechain
38	BE	25	U	Sidechain
38	BE	26	G	Sidechain
38	BE	27	A	Sidechain
38	BE	29	C	Sidechain
38	BE	30	C	Sidechain
38	BE	31	A	Sidechain
38	BE	32	U	Sidechain
38	BE	33	C	Sidechain
38	BE	35	A	Sidechain
38	BE	36	U	Sidechain
38	BE	37	C	Sidechain
38	BE	4	A	Sidechain
38	BE	40	C	Sidechain
38	BE	41	C	Sidechain
38	BE	43	A	Sidechain
38	BE	44	C	Sidechain
38	BE	46	G	Sidechain
38	BE	47	U	Sidechain
38	BE	48	G	Sidechain
38	BE	49	A	Sidechain
38	BE	5	A	Sidechain
38	BE	51	C	Sidechain
38	BE	52	U	Sidechain
38	BE	53	U	Sidechain
38	BE	54	U	Sidechain
38	BE	55	C	Sidechain
38	BE	57	U	Sidechain
38	BE	58	U	Sidechain
38	BE	59	U	Sidechain
38	BE	6	A	Sidechain
38	BE	60	C	Sidechain
38	BE	61	A	Sidechain
38	BE	63	C	Sidechain
38	BE	64	A	Sidechain

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Mol	Chain	Res	Type	Group
38	BE	65	U	Sidechain
38	BE	67	A	Sidechain
38	BE	68	U	Sidechain
38	BE	69	C	Sidechain
38	BE	7	U	Sidechain
38	BE	70	C	Sidechain
38	BE	71	A	Sidechain
38	BE	72	C	Sidechain
38	BE	74	U	Sidechain
38	BE	75	C	Sidechain
38	BE	76	U	Sidechain
38	BE	79	G	Sidechain
38	BE	80	G	Sidechain
38	BE	82	C	Sidechain
38	BE	83	U	Sidechain
38	BE	86	C	Sidechain
38	BE	87	U	Sidechain
38	BE	88	G	Sidechain
38	BE	89	G	Sidechain
38	BE	9	C	Sidechain
38	BE	90	G	Sidechain
38	BE	91	G	Sidechain
38	BE	92	C	Sidechain
38	BE	93	U	Sidechain
38	BE	94	U	Sidechain
38	BE	96	G	Sidechain
38	BE	97	G	Sidechain
38	BE	98	C	Sidechain
39	BF	10	A	Sidechain
39	BF	12	U	Sidechain
39	BF	13	U	Sidechain
39	BF	14	C	Sidechain
39	BF	15	U	Sidechain
39	BF	16	C	Sidechain
39	BF	17	U	Sidechain
39	BF	19	A	Sidechain
39	BF	2	G	Sidechain
39	BF	20	U	Sidechain
39	BF	21	C	Sidechain
39	BF	22	U	Sidechain
39	BF	23	G	Sidechain
39	BF	24	G	Sidechain

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Mol	Chain	Res	Type	Group
39	BF	25	G	Sidechain
39	BF	26	U	Sidechain
39	BF	27	G	Sidechain
39	BF	28	C	Sidechain
39	BF	29	U	Sidechain
39	BF	3	A	Sidechain
39	BF	30	C	Sidechain
39	BF	32	G	Sidechain
39	BF	33	C	Sidechain
39	BF	34	C	Sidechain
39	BF	38	C	Sidechain
39	BF	39	C	Sidechain
39	BF	4	A	Sidechain
39	BF	40	U	Sidechain
39	BF	41	U	Sidechain
39	BF	43	U	Sidechain
39	BF	44	C	Sidechain
39	BF	46	G	Sidechain
39	BF	48	G	Sidechain
39	BF	5	U	Sidechain
39	BF	50	C	Sidechain
39	BF	51	C	Sidechain
39	BF	52	A	Sidechain
39	BF	53	G	Sidechain
39	BF	54	U	Sidechain
39	BF	55	A	Sidechain
39	BF	56	C	Sidechain
39	BF	57	C	Sidechain
39	BF	58	U	Sidechain
39	BF	59	U	Sidechain
39	BF	6	C	Sidechain
39	BF	60	C	Sidechain
39	BF	61	A	Sidechain
39	BF	62	U	Sidechain
39	BF	63	U	Sidechain
39	BF	64	U	Sidechain
39	BF	65	U	Sidechain
39	BF	66	C	Sidechain
39	BF	67	A	Sidechain
39	BF	68	C	Sidechain
39	BF	69	A	Sidechain
39	BF	7	G	Sidechain

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Mol	Chain	Res	Type	Group
39	BF	70	A	Sidechain
39	BF	71	G	Sidechain
39	BF	72	A	Sidechain
39	BF	8	C	Sidechain
39	BF	9	C	Sidechain
40	BG	1	G	Sidechain
40	BG	10	U	Sidechain
40	BG	100	G	Sidechain
40	BG	101	G	Sidechain
40	BG	102	G	Sidechain
40	BG	103	C	Sidechain
40	BG	104	A	Sidechain
40	BG	106	G	Sidechain
40	BG	107	U	Sidechain
40	BG	11	G	Sidechain
40	BG	110	U	Sidechain
40	BG	111	C	Sidechain
40	BG	112	C	Sidechain
40	BG	113	G	Sidechain
40	BG	114	A	Sidechain
40	BG	115	C	Sidechain
40	BG	118	U	Sidechain
40	BG	119	A	Sidechain
40	BG	12	A	Sidechain
40	BG	120	U	Sidechain
40	BG	121	C	Sidechain
40	BG	122	G	Sidechain
40	BG	123	C	Sidechain
40	BG	124	A	Sidechain
40	BG	125	C	Sidechain
40	BG	126	G	Sidechain
40	BG	127	G	Sidechain
40	BG	128	U	Sidechain
40	BG	129	G	Sidechain
40	BG	13	A	Sidechain
40	BG	130	G	Sidechain
40	BG	131	U	Sidechain
40	BG	132	U	Sidechain
40	BG	133	C	Sidechain
40	BG	135	C	Sidechain
40	BG	136	G	Sidechain
40	BG	137	G	Sidechain

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Mol	Chain	Res	Type	Group
40	BG	138	C	Sidechain
40	BG	139	U	Sidechain
40	BG	140	G	Sidechain
40	BG	143	C	Sidechain
40	BG	144	G	Sidechain
40	BG	145	C	Sidechain
40	BG	146	C	Sidechain
40	BG	147	U	Sidechain
40	BG	148	C	Sidechain
40	BG	149	U	Sidechain
40	BG	15	G	Sidechain
40	BG	150	A	Sidechain
40	BG	151	A	Sidechain
40	BG	152	G	Sidechain
40	BG	153	C	Sidechain
40	BG	154	C	Sidechain
40	BG	155	A	Sidechain
40	BG	156	G	Sidechain
40	BG	157	A	Sidechain
40	BG	158	A	Sidechain
40	BG	159	A	Sidechain
40	BG	160	C	Sidechain
40	BG	163	G	Sidechain
40	BG	164	U	Sidechain
40	BG	166	C	Sidechain
40	BG	167	C	Sidechain
40	BG	168	A	Sidechain
40	BG	169	A	Sidechain
40	BG	17	A	Sidechain
40	BG	170	G	Sidechain
40	BG	171	A	Sidechain
40	BG	172	C	Sidechain
40	BG	173	C	Sidechain
40	BG	174	G	Sidechain
40	BG	176	G	Sidechain
40	BG	177	U	Sidechain
40	BG	178	G	Sidechain
40	BG	181	C	Sidechain
40	BG	182	G	Sidechain
40	BG	2	U	Sidechain
40	BG	20	U	Sidechain
40	BG	21	C	Sidechain

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Mol	Chain	Res	Type	Group
40	BG	22	G	Sidechain
40	BG	24	A	Sidechain
40	BG	25	G	Sidechain
40	BG	26	G	Sidechain
40	BG	27	C	Sidechain
40	BG	29	U	Sidechain
40	BG	31	G	Sidechain
40	BG	32	U	Sidechain
40	BG	33	G	Sidechain
40	BG	34	A	Sidechain
40	BG	35	G	Sidechain
40	BG	36	G	Sidechain
40	BG	37	G	Sidechain
40	BG	39	A	Sidechain
40	BG	4	A	Sidechain
40	BG	40	G	Sidechain
40	BG	41	U	Sidechain
40	BG	42	A	Sidechain
40	BG	43	U	Sidechain
40	BG	45	G	Sidechain
40	BG	47	G	Sidechain
40	BG	48	U	Sidechain
40	BG	49	A	Sidechain
40	BG	5	G	Sidechain
40	BG	50	G	Sidechain
40	BG	53	C	Sidechain
40	BG	54	G	Sidechain
40	BG	56	G	Sidechain
40	BG	58	G	Sidechain
40	BG	60	A	Sidechain
40	BG	61	A	Sidechain
40	BG	62	C	Sidechain
40	BG	63	U	Sidechain
40	BG	64	C	Sidechain
40	BG	65	C	Sidechain
40	BG	66	C	Sidechain
40	BG	68	U	Sidechain
40	BG	69	G	Sidechain
40	BG	70	C	Sidechain
40	BG	71	C	Sidechain
40	BG	72	G	Sidechain
40	BG	73	U	Sidechain

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Mol	Chain	Res	Type	Group
40	BG	74	G	Sidechain
40	BG	75	C	Sidechain
40	BG	76	C	Sidechain
40	BG	77	U	Sidechain
40	BG	78	C	Sidechain
40	BG	79	U	Sidechain
40	BG	8	U	Sidechain
40	BG	80	G	Sidechain
40	BG	81	G	Sidechain
40	BG	83	U	Sidechain
40	BG	84	U	Sidechain
40	BG	85	C	Sidechain
40	BG	86	U	Sidechain
40	BG	89	A	Sidechain
40	BG	9	G	Sidechain
40	BG	90	G	Sidechain
40	BG	93	U	Sidechain
40	BG	94	G	Sidechain
40	BG	95	U	Sidechain
40	BG	96	C	Sidechain
40	BG	97	G	Sidechain
40	BG	98	A	Sidechain
40	BG	99	A	Sidechain
41	BH	10	U	Sidechain
41	BH	100	A	Sidechain
41	BH	101	A	Sidechain
41	BH	102	C	Sidechain
41	BH	104	U	Sidechain
41	BH	105	U	Sidechain
41	BH	106	G	Sidechain
41	BH	107	A	Sidechain
41	BH	108	U	Sidechain
41	BH	109	G	Sidechain
41	BH	11	C	Sidechain
41	BH	110	C	Sidechain
41	BH	113	G	Sidechain
41	BH	114	G	Sidechain
41	BH	115	A	Sidechain
41	BH	116	A	Sidechain
41	BH	117	U	Sidechain
41	BH	118	U	Sidechain
41	BH	119	U	Sidechain

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Mol	Chain	Res	Type	Group
41	BH	12	U	Sidechain
41	BH	120	C	Sidechain
41	BH	121	A	Sidechain
41	BH	123	G	Sidechain
41	BH	124	C	Sidechain
41	BH	126	C	Sidechain
41	BH	127	A	Sidechain
41	BH	128	G	Sidechain
41	BH	130	G	Sidechain
41	BH	132	C	Sidechain
41	BH	133	U	Sidechain
41	BH	134	U	Sidechain
41	BH	135	U	Sidechain
41	BH	15	A	Sidechain
41	BH	16	A	Sidechain
41	BH	19	G	Sidechain
41	BH	2	U	Sidechain
41	BH	21	G	Sidechain
41	BH	22	A	Sidechain
41	BH	23	G	Sidechain
41	BH	24	U	Sidechain
41	BH	25	A	Sidechain
41	BH	26	C	Sidechain
41	BH	27	A	Sidechain
41	BH	28	U	Sidechain
41	BH	29	G	Sidechain
41	BH	3	U	Sidechain
41	BH	30	C	Sidechain
41	BH	31	A	Sidechain
41	BH	32	U	Sidechain
41	BH	33	G	Sidechain
41	BH	34	G	Sidechain
41	BH	36	C	Sidechain
41	BH	37	U	Sidechain
41	BH	38	G	Sidechain
41	BH	39	G	Sidechain
41	BH	4	U	Sidechain
41	BH	41	A	Sidechain
41	BH	42	U	Sidechain
41	BH	43	G	Sidechain
41	BH	44	A	Sidechain
41	BH	48	G	Sidechain

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Mol	Chain	Res	Type	Group
41	BH	49	C	Sidechain
41	BH	5	G	Sidechain
41	BH	50	A	Sidechain
41	BH	51	C	Sidechain
41	BH	52	G	Sidechain
41	BH	54	U	Sidechain
41	BH	6	U	Sidechain
41	BH	62	C	Sidechain
41	BH	63	G	Sidechain
41	BH	65	G	Sidechain
41	BH	7	C	Sidechain
41	BH	70	U	Sidechain
41	BH	71	C	Sidechain
41	BH	74	G	Sidechain
41	BH	75	G	Sidechain
41	BH	76	G	Sidechain
41	BH	79	A	Sidechain
41	BH	8	C	Sidechain
41	BH	81	U	Sidechain
41	BH	82	U	Sidechain
41	BH	84	A	Sidechain
41	BH	93	G	Sidechain
41	BH	94	G	Sidechain
41	BH	96	G	Sidechain
41	BH	98	U	Sidechain
41	BH	99	G	Sidechain
42	BI	101	ARG	Sidechain
42	BI	119	ARG	Sidechain
42	BI	147	ARG	Sidechain
42	BI	149	ARG	Sidechain
42	BI	15	VAL	Peptide,Mainchain
42	BI	16	ARG	Sidechain
42	BI	20	TYR	Sidechain
42	BI	25	TYR	Sidechain
42	BI	33	TYR	Sidechain
42	BI	39	ARG	Mainchain
42	BI	49	HIS	Sidechain
42	BI	51	ARG	Sidechain
42	BI	66	ARG	Sidechain
42	BI	71	MET	Peptide
42	BI	73	ARG	Sidechain
42	BI	77	TRP	Mainchain

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Mol	Chain	Res	Type	Group
42	BI	79	LYS	Peptide
42	BI	94	LEU	Mainchain
43	BJ	104	TYR	Sidechain
43	BJ	20	ARG	Sidechain
43	BJ	25	PHE	Peptide
43	BJ	45	ARG	Sidechain
43	BJ	57	ARG	Sidechain
43	BJ	59	ARG	Peptide
43	BJ	9	LEU	Peptide
44	BK	116	ARG	Sidechain
44	BK	119	TYR	Sidechain
44	BK	139	ARG	Sidechain
44	BK	168	SER	Peptide
44	BK	17	TYR	Sidechain
44	BK	174	THR	Peptide
44	BK	181	TYR	Sidechain
44	BK	185	ARG	Sidechain
44	BK	193	ARG	Sidechain
44	BK	196	HIS	Sidechain
44	BK	206	ILE	Peptide
44	BK	209	TYR	Sidechain
44	BK	21	ARG	Sidechain
44	BK	4	ARG	Sidechain
44	BK	40	ARG	Sidechain
44	BK	51	HIS	Sidechain
44	BK	56	GLU	Peptide
44	BK	69	ARG	Sidechain
44	BK	7	ARG	Sidechain
44	BK	75	TYR	Sidechain
44	BK	79	ARG	Sidechain
44	BK	85	PHE	Sidechain
44	BK	86	HIS	Sidechain
44	BK	88	ARG	Sidechain
44	BK	95	HIS	Sidechain
44	BK	99	ILE	Peptide
45	BL	113	ILE	Peptide
45	BL	116	HIS	Sidechain
45	BL	123	TYR	Sidechain
45	BL	140	ARG	Sidechain
45	BL	148	ARG	Sidechain
45	BL	15	PRO	Peptide
45	BL	151	ARG	Sidechain

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Mol	Chain	Res	Type	Group
45	BL	154	ARG	Sidechain
45	BL	159	HIS	Sidechain
45	BL	17	ARG	Sidechain
45	BL	174	HIS	Mainchain
45	BL	58	ARG	Sidechain
45	BL	62	ARG	Sidechain
45	BL	64	PHE	Sidechain
45	BL	67	ARG	Sidechain
47	BN	10	HIS	Sidechain
47	BN	101	ARG	Sidechain
47	BN	104	ARG	Sidechain
47	BN	105	ARG	Mainchain,Sidechain
47	BN	12	HIS	Sidechain
47	BN	124	TYR	Sidechain
47	BN	135	ARG	Sidechain
47	BN	14	ARG	Sidechain
47	BN	143	ALA	Peptide
47	BN	154	ASP	Peptide
47	BN	155	ARG	Mainchain
47	BN	157	ARG	Sidechain
47	BN	18	ASN	Mainchain
47	BN	187	TYR	Sidechain
47	BN	203	ARG	Sidechain
47	BN	204	ASN	Peptide
47	BN	214	GLU	Mainchain
47	BN	42	ARG	Sidechain
47	BN	44	ARG	Sidechain
47	BN	51	ILE	Peptide,Mainchain
47	BN	54	ARG	Sidechain
47	BN	60	ARG	Sidechain
47	BN	63	VAL	Peptide
47	BN	67	THR	Peptide,Mainchain
47	BN	69	ARG	Sidechain
47	BN	74	ARG	Sidechain
47	BN	80	PHE	Sidechain
47	BN	96	ARG	Sidechain
48	BO	108	ARG	Sidechain
48	BO	109	TYR	Sidechain
48	BO	118	ARG	Sidechain
48	BO	119	GLN	Mainchain
48	BO	123	TYR	Sidechain
48	BO	127	PRO	Peptide

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Mol	Chain	Res	Type	Group
48	BO	144	ARG	Sidechain
48	BO	153	ARG	Sidechain
48	BO	154	PHE	Sidechain
48	BO	179	ARG	Sidechain
48	BO	186	HIS	Sidechain
48	BO	201	ARG	Sidechain
48	BO	219	PHE	Sidechain
48	BO	23	ASP	Peptide
48	BO	24	ILE	Mainchain
48	BO	55	ARG	Sidechain
48	BO	67	ARG	Sidechain
48	BO	72	TYR	Sidechain
48	BO	89	PHE	Mainchain,Sidechain
48	BO	91	HIS	Sidechain
49	BP	11	ARG	Sidechain
49	BP	110	ARG	Sidechain
49	BP	120	ARG	Sidechain
49	BP	122	TYR	Sidechain
49	BP	127	ILE	Peptide
49	BP	135	ASN	Peptide
49	BP	14	ARG	Sidechain
49	BP	140	HIS	Sidechain
49	BP	150	ASN	Mainchain
49	BP	151	ALA	Mainchain
49	BP	155	ASP	Peptide
49	BP	157	THR	Peptide,Mainchain
49	BP	158	PRO	Peptide
49	BP	17	ARG	Sidechain
49	BP	171	ARG	Sidechain
49	BP	20	ARG	Sidechain
49	BP	3	LYS	Peptide
49	BP	35	ARG	Sidechain
49	BP	6	TYR	Sidechain
49	BP	8	ARG	Sidechain
49	BP	87	TYR	Sidechain
49	BP	96	HIS	Sidechain
49	BP	97	ARG	Sidechain
50	BQ	101	PRO	Peptide,Mainchain
50	BQ	109	ARG	Peptide,Sidechain
50	BQ	116	ARG	Sidechain
50	BQ	129	ASN	Mainchain
50	BQ	147	TYR	Sidechain

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Mol	Chain	Res	Type	Group
50	BQ	160	ARG	Sidechain
50	BQ	179	ARG	Sidechain
50	BQ	186	ARG	Sidechain
50	BQ	188	HIS	Sidechain
50	BQ	189	ARG	Sidechain
50	BQ	192	ARG	Sidechain
50	BQ	194	LYS	Peptide
50	BQ	204	SER	Mainchain
50	BQ	205	ARG	Sidechain
50	BQ	206	ARG	Sidechain
50	BQ	221	ARG	Sidechain
50	BQ	48	ARG	Sidechain
50	BQ	61	ARG	Sidechain
50	BQ	66	ARG	Sidechain
50	BQ	70	TYR	Sidechain
50	BQ	76	TYR	Sidechain
50	BQ	80	ARG	Sidechain
50	BQ	82	ARG	Sidechain
50	BQ	85	ARG	Sidechain
50	BQ	88	ARG	Sidechain
50	BQ	98	TYR	Sidechain
51	BR	102	ALA	Mainchain
51	BR	127	ARG	Sidechain
51	BR	131	ARG	Sidechain
51	BR	133	HIS	Sidechain
51	BR	141	ARG	Sidechain
51	BR	145	HIS	Sidechain
51	BR	149	PHE	Sidechain
51	BR	154	GLN	Peptide,Mainchain
51	BR	26	TYR	Sidechain
51	BR	39	MET	Peptide
51	BR	4	TYR	Sidechain
51	BR	48	TYR	Sidechain
51	BR	49	ARG	Sidechain
51	BR	56	ARG	Sidechain
51	BR	58	ILE	Mainchain
51	BR	6	ARG	Sidechain
51	BR	60	PHE	Sidechain
51	BR	63	TYR	Sidechain
51	BR	65	GLY	Peptide,Mainchain
52	BS	116	HIS	Sidechain
52	BS	119	ARG	Sidechain

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Mol	Chain	Res	Type	Group
52	BS	121	HIS	Sidechain
52	BS	134	HIS	Sidechain
52	BS	137	ARG	Sidechain
52	BS	138	ARG	Sidechain
52	BS	14	ARG	Sidechain
52	BS	156	ARG	Sidechain
52	BS	170	LYS	Peptide
52	BS	2	VAL	Peptide
52	BS	24	PRO	Peptide
52	BS	27	TYR	Sidechain
52	BS	29	PHE	Peptide
52	BS	3	ARG	Sidechain
52	BS	4	PRO	Peptide
52	BS	42	ARG	Sidechain
52	BS	57	HIS	Sidechain
52	BS	76	TYR	Sidechain
52	BS	89	TYR	Sidechain
52	BS	9	TYR	Sidechain
52	BS	96	PHE	Sidechain
53	BT	100	ARG	Sidechain
53	BT	103	ARG	Sidechain
53	BT	104	ARG	Sidechain
53	BT	117	ARG	Sidechain
53	BT	120	TYR	Sidechain
53	BT	121	ARG	Sidechain
53	BT	131	VAL	Mainchain
53	BT	153	ARG	Sidechain
53	BT	16	ARG	Sidechain
53	BT	160	ALA	Peptide
53	BT	171	ARG	Sidechain
53	BT	179	LYS	Peptide,Mainchain
53	BT	185	ARG	Sidechain
53	BT	189	ARG	Sidechain
53	BT	191	ASP	Peptide
53	BT	21	ARG	Sidechain
53	BT	38	ARG	Sidechain
53	BT	62	ARG	Sidechain
53	BT	74	ARG	Sidechain
53	BT	75	HIS	Sidechain
53	BT	80	ARG	Sidechain
53	BT	81	ARG	Sidechain
53	BT	97	ARG	Sidechain

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Mol	Chain	Res	Type	Group
54	BU	12	ARG	Sidechain
54	BU	120	LYS	Peptide
54	BU	123	THR	Peptide
54	BU	133	ARG	Peptide,Sidechain
54	BU	141	LYS	Peptide
54	BU	148	ARG	Sidechain
54	BU	150	THR	Peptide
54	BU	157	ILE	Peptide
54	BU	19	PHE	Sidechain
54	BU	20	ARG	Sidechain
54	BU	34	PHE	Sidechain
54	BU	39	TYR	Sidechain
54	BU	49	ARG	Sidechain
54	BU	56	TYR	Peptide,Sidechain
54	BU	57	TYR	Sidechain
54	BU	7	TYR	Sidechain
54	BU	81	ARG	Sidechain
54	BU	83	ARG	Sidechain
54	BU	88	ARG	Sidechain
55	BV	108	ARG	Sidechain
55	BV	121	TYR	Sidechain
55	BV	43	ILE	Peptide
55	BV	50	TYR	Sidechain
55	BV	51	PHE	Sidechain
55	BV	92	TYR	Sidechain
56	BW	134	HIS	Sidechain
56	BW	135	ALA	Peptide
56	BW	37	TYR	Sidechain
56	BW	44	TYR	Sidechain
56	BW	6	ALA	Peptide
56	BW	72	ARG	Sidechain
56	BW	73	ARG	Peptide,Mainchain
56	BW	84	ARG	Sidechain
56	BW	89	ARG	Sidechain
57	BX	125	TYR	Sidechain
57	BX	147	ARG	Sidechain
57	BX	46	ARG	Sidechain
57	BX	49	TYR	Sidechain
57	BX	60	TYR	Sidechain
57	BX	61	ARG	Mainchain,Sidechain
58	BY	17	HIS	Sidechain
58	BY	19	ARG	Sidechain

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Mol	Chain	Res	Type	Group
58	BY	2	ARG	Sidechain
58	BY	21	TYR	Sidechain
58	BY	29	THR	Peptide
58	BY	46	ARG	Sidechain
58	BY	53	ILE	Mainchain
58	BY	57	ARG	Sidechain
58	BY	59	TYR	Sidechain
58	BY	61	ARG	Sidechain
58	BY	62	ILE	Mainchain
58	BY	63	HIS	Sidechain
58	BY	66	THR	Peptide
58	BY	70	ARG	Sidechain
58	BY	84	ARG	Sidechain
59	BZ	111	ARG	Sidechain
59	BZ	121	ARG	Sidechain
59	BZ	13	ARG	Sidechain
59	BZ	24	ARG	Sidechain
59	BZ	25	ARG	Sidechain
59	BZ	54	ARG	Sidechain
59	BZ	57	ARG	Sidechain
59	BZ	7	ARG	Mainchain,Sidechain
59	BZ	71	TYR	Sidechain
59	BZ	72	ARG	Sidechain
59	BZ	78	HIS	Sidechain
60	Ba	100	ASP	Peptide
60	Ba	113	ARG	Sidechain
60	Ba	127	PHE	Sidechain
60	Ba	130	ARG	Sidechain
60	Ba	16	ARG	Sidechain
60	Ba	17	TYR	Sidechain
60	Ba	3	PHE	Sidechain
60	Ba	37	TYR	Sidechain
60	Ba	59	ARG	Peptide
60	Ba	62	THR	Mainchain
60	Ba	64	ARG	Sidechain
60	Ba	70	PHE	Sidechain
60	Ba	72	ARG	Sidechain
60	Ba	76	HIS	Sidechain
60	Ba	78	HIS	Sidechain
60	Ba	79	PHE	Sidechain
60	Ba	84	TYR	Sidechain
60	Ba	90	ARG	Sidechain

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Mol	Chain	Res	Type	Group
60	Ba	91	GLU	Peptide
60	Ba	95	ARG	Sidechain
61	Bb	106	TYR	Sidechain
61	Bb	110	LEU	Mainchain
61	Bb	117	ALA	Peptide
61	Bb	12	ARG	Sidechain
61	Bb	122	LYS	Mainchain
61	Bb	14	SER	Peptide
61	Bb	19	TYR	Sidechain
61	Bb	26	ARG	Sidechain
61	Bb	32	ARG	Sidechain
61	Bb	42	ARG	Sidechain
61	Bb	48	TYR	Sidechain
61	Bb	53	PHE	Sidechain
61	Bb	61	TYR	Sidechain
61	Bb	69	TRP	Mainchain
61	Bb	9	ARG	Sidechain
62	Bc	107	TYR	Sidechain
62	Bc	109	ARG	Sidechain
62	Bc	127	ARG	Sidechain
62	Bc	13	ARG	Sidechain
62	Bc	130	ARG	Sidechain
62	Bc	135	HIS	Peptide
62	Bc	137	HIS	Sidechain
62	Bc	142	ARG	Sidechain
62	Bc	18	PHE	Sidechain
62	Bc	22	ARG	Sidechain
62	Bc	26	ARG	Mainchain
62	Bc	27	LEU	Mainchain
62	Bc	32	PHE	Sidechain
62	Bc	34	ASN	Peptide
62	Bc	41	ARG	Sidechain
62	Bc	45	PHE	Sidechain
62	Bc	51	ALA	Mainchain
62	Bc	59	ARG	Peptide,Sidechain
62	Bc	83	GLU	Peptide
62	Bc	93	ARG	Peptide,Mainchain,Sidechain
62	Bc	96	ALA	Peptide,Mainchain
62	Bc	97	ASP	Peptide
62	Bc	99	ARG	Peptide
63	Bd	10	HIS	Sidechain
63	Bd	18	ARG	Sidechain

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Mol	Chain	Res	Type	Group
63	Bd	19	ASN	Peptide
63	Bd	28	TYR	Sidechain
63	Bd	31	ASN	Peptide,Mainchain
63	Bd	34	ARG	Sidechain
63	Bd	49	LYS	Peptide
63	Bd	59	ARG	Sidechain
63	Bd	61	GLU	Sidechain
64	Be	128	ARG	Sidechain
64	Be	133	TYR	Sidechain
64	Be	145	ARG	Sidechain
64	Be	147	ARG	Sidechain
64	Be	163	ARG	Sidechain
64	Be	186	PHE	Sidechain
64	Be	189	PHE	Mainchain
64	Be	200	ARG	Sidechain
64	Be	204	ARG	Mainchain,Sidechain
64	Be	207	VAL	Mainchain
64	Be	211	HIS	Sidechain
64	Be	221	HIS	Sidechain
64	Be	245	ARG	Sidechain
64	Be	250	ARG	Sidechain
64	Be	253	ARG	Sidechain
64	Be	71	ARG	Sidechain
64	Be	72	VAL	Mainchain
64	Be	83	PHE	Sidechain
64	Be	89	LEU	Peptide,Mainchain
65	Bf	100	TYR	Sidechain
65	Bf	106	HIS	Sidechain
65	Bf	111	VAL	Mainchain
65	Bf	150	HIS	Sidechain
65	Bf	167	ARG	Sidechain
65	Bf	168	ARG	Sidechain
65	Bf	173	TRP	Peptide
65	Bf	184	ARG	Sidechain
65	Bf	189	ARG	Sidechain
65	Bf	212	ARG	Sidechain
65	Bf	216	HIS	Sidechain
65	Bf	223	ARG	Sidechain
65	Bf	226	ARG	Sidechain
65	Bf	260	ARG	Sidechain
65	Bf	265	PHE	Sidechain
65	Bf	295	PRO	Mainchain

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Mol	Chain	Res	Type	Group
65	Bf	296	ARG	Sidechain
65	Bf	299	HIS	Sidechain
65	Bf	300	ARG	Sidechain
65	Bf	312	HIS	Sidechain
65	Bf	315	ARG	Sidechain
65	Bf	318	TYR	Sidechain
65	Bf	339	TYR	Sidechain
65	Bf	343	ARG	Sidechain
65	Bf	348	GLU	Mainchain
65	Bf	356	TYR	Sidechain
65	Bf	373	GLY	Peptide
65	Bf	376	ARG	Sidechain
65	Bf	392	ARG	Sidechain
65	Bf	397	ARG	Sidechain
65	Bf	404	THR	Peptide
65	Bf	407	LYS	Peptide
65	Bf	412	ILE	Mainchain
65	Bf	416	PHE	Sidechain
65	Bf	425	HIS	Sidechain
65	Bf	427	ARG	Sidechain
65	Bf	428	PHE	Sidechain
65	Bf	438	TYR	Sidechain
65	Bf	445	ARG	Sidechain
65	Bf	447	ARG	Mainchain,Sidechain
65	Bf	451	ARG	Sidechain
65	Bf	453	ARG	Sidechain
65	Bf	54	HIS	Sidechain
65	Bf	57	PHE	Sidechain
65	Bf	62	HIS	Sidechain
65	Bf	77	ARG	Sidechain
65	Bf	79	ARG	Sidechain
66	Bg	105	ASP	Peptide,Mainchain
66	Bg	34	TYR	Sidechain
66	Bg	46	ARG	Sidechain
66	Bg	59	CYS	Mainchain
66	Bg	79	HIS	Sidechain
66	Bg	81	TYR	Sidechain
66	Bg	99	CYS	Mainchain
67	Bh	121	LYS	Peptide
67	Bh	134	PHE	Sidechain
67	Bh	146	ARG	Sidechain
67	Bh	148	ARG	Sidechain

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Mol	Chain	Res	Type	Group
67	Bh	164	LYS	Peptide,Mainchain
67	Bh	167	TYR	Sidechain
67	Bh	175	VAL	Mainchain
67	Bh	178	PHE	Sidechain
67	Bh	183	THR	Mainchain
67	Bh	29	ALA	Mainchain
67	Bh	3	ARG	Sidechain
67	Bh	40	ARG	Sidechain
67	Bh	60	ALA	Peptide,Mainchain
67	Bh	64	ARG	Sidechain
67	Bh	71	ALA	Mainchain
67	Bh	73	LEU	Mainchain
67	Bh	74	PHE	Sidechain
67	Bh	79	ARG	Sidechain
67	Bh	82	ASP	Peptide
67	Bh	91	HIS	Sidechain
67	Bh	99	ARG	Sidechain
68	Bi	106	ARG	Sidechain
68	Bi	111	ARG	Sidechain
68	Bi	114	HIS	Sidechain
68	Bi	119	ARG	Sidechain
68	Bi	3	LYS	Peptide
68	Bi	45	ARG	Sidechain
68	Bi	46	ARG	Sidechain
68	Bi	47	TYR	Sidechain
68	Bi	5	PHE	Sidechain
68	Bi	62	ARG	Sidechain
68	Bi	68	THR	Mainchain
68	Bi	84	TYR	Sidechain
68	Bi	99	HIS	Sidechain
69	Bj	10	ARG	Sidechain
69	Bj	14	TYR	Sidechain
69	Bj	17	ARG	Sidechain
69	Bj	2	SER	Mainchain
69	Bj	20	ARG	Sidechain
69	Bj	22	ARG	Sidechain
69	Bj	3	CYS	Peptide
69	Bj	30	ARG	Sidechain
69	Bj	36	ARG	Sidechain
69	Bj	39	ARG	Sidechain
69	Bj	45	THR	Peptide
69	Bj	46	PRO	Mainchain

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Mol	Chain	Res	Type	Group
69	Bj	5	ARG	Sidechain
69	Bj	53	ARG	Sidechain
69	Bj	6	VAL	Mainchain
69	Bj	62	HIS	Sidechain
69	Bj	66	ARG	Sidechain
69	Bj	77	ARG	Sidechain
69	Bj	8	TYR	Sidechain
69	Bj	85	HIS	Mainchain,Sidechain
69	Bj	9	ARG	Sidechain
69	Bj	94	ARG	Sidechain
70	Bk	106	ARG	Sidechain
70	Bk	110	ARG	Sidechain
70	Bk	116	HIS	Sidechain
70	Bk	123	TYR	Sidechain
70	Bk	47	ARG	Sidechain
70	Bk	67	ARG	Sidechain
70	Bk	81	SER	Peptide,Mainchain
70	Bk	87	LEU	Mainchain
70	Bk	88	ARG	Sidechain
70	Bk	94	ARG	Sidechain
70	Bk	95	ARG	Sidechain
71	Bl	101	ARG	Sidechain
71	Bl	102	ARG	Sidechain
71	Bl	103	SER	Peptide
71	Bl	106	ARG	Sidechain
71	Bl	114	ARG	Sidechain
71	Bl	116	HIS	Sidechain
71	Bl	130	SER	Peptide
71	Bl	139	ARG	Sidechain
71	Bl	148	ARG	Sidechain
71	Bl	34	LYS	Peptide
71	Bl	35	SER	Peptide
71	Bl	39	TYR	Sidechain
71	Bl	47	TYR	Sidechain
71	Bl	75	TYR	Sidechain
71	Bl	79	ARG	Sidechain
71	Bl	82	TYR	Sidechain
71	Bl	84	TYR	Sidechain
71	Bl	89	ILE	Peptide
71	Bl	91	ARG	Sidechain
71	Bl	93	VAL	Peptide
71	Bl	94	ARG	Sidechain

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Mol	Chain	Res	Type	Group
71	Bl	98	ALA	Peptide,Mainchain
72	Bm	12	ARG	Sidechain
72	Bm	13	THR	Peptide
72	Bm	27	ARG	Sidechain
72	Bm	28	ARG	Sidechain
72	Bm	31	ARG	Sidechain
72	Bm	37	ARG	Sidechain
72	Bm	38	TYR	Sidechain
72	Bm	42	HIS	Mainchain
72	Bm	63	ARG	Sidechain
72	Bm	70	ARG	Sidechain
72	Bm	76	ARG	Mainchain,Sidechain
72	Bm	77	ALA	Peptide
72	Bm	84	ARG	Sidechain
72	Bm	9	PRO	Peptide
72	Bm	94	ARG	Sidechain
73	Bn	11	ARG	Sidechain
73	Bn	21	ARG	Sidechain
73	Bn	31	TRP	Mainchain
73	Bn	45	ARG	Sidechain
73	Bn	58	ARG	Sidechain
73	Bn	66	TYR	Sidechain
73	Bn	72	ARG	Sidechain
73	Bn	73	ARG	Sidechain
73	Bn	75	ARG	Sidechain
73	Bn	76	ASN	Peptide,Mainchain
73	Bn	78	PHE	Peptide,Mainchain
73	Bn	79	LYS	Mainchain
74	Bo	14	TYR	Sidechain
74	Bo	17	ARG	Sidechain
74	Bo	18	TYR	Sidechain
74	Bo	25	ARG	Sidechain
74	Bo	4	ARG	Sidechain
74	Bo	49	ARG	Sidechain
74	Bo	57	CYS	Mainchain
74	Bo	59	GLY	Peptide,Mainchain
74	Bo	69	TYR	Sidechain
74	Bo	85	ARG	Sidechain
74	Bo	87	ARG	Sidechain
75	Bp	11	PHE	Sidechain
75	Bp	21	ARG	Sidechain
75	Bp	30	LYS	Peptide

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Mol	Chain	Res	Type	Group
75	Bp	37	ARG	Sidechain
75	Bp	72	ARG	Sidechain
75	Bp	79	THR	Peptide
76	Bq	13	TYR	Sidechain
76	Bq	21	ARG	Sidechain
76	Bq	25	TYR	Sidechain
76	Bq	3	ARG	Sidechain
76	Bq	30	ARG	Sidechain
76	Bq	42	ARG	Sidechain
76	Bq	43	HIS	Sidechain
76	Bq	45	ARG	Peptide,Sidechain
77	Br	10	TYR	Sidechain
77	Br	103	PRO	Peptide,Mainchain
77	Br	107	PHE	Peptide
77	Br	109	ARG	Sidechain
77	Br	132	LEU	Peptide,Mainchain
77	Br	139	ARG	Sidechain
77	Br	160	TYR	Sidechain
77	Br	169	PHE	Sidechain
77	Br	174	GLY	Peptide,Mainchain
77	Br	198	ARG	Sidechain
77	Br	203	ARG	Sidechain
77	Br	204	ARG	Sidechain
77	Br	217	ARG	Peptide,Sidechain
77	Br	219	PHE	Sidechain
77	Br	220	ARG	Sidechain
77	Br	225	LEU	Mainchain
77	Br	246	ARG	Sidechain
77	Br	250	TRP	Peptide
77	Br	262	PHE	Sidechain
77	Br	264	THR	Mainchain
77	Br	265	PHE	Sidechain
77	Br	275	PHE	Sidechain
77	Br	289	ARG	Sidechain
77	Br	297	ARG	Sidechain
77	Br	298	ARG	Sidechain
77	Br	299	VAL	Peptide
77	Br	309	ARG	Sidechain
77	Br	313	TYR	Sidechain
77	Br	32	ARG	Sidechain
77	Br	320	ILE	Peptide
77	Br	323	ARG	Sidechain

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Mol	Chain	Res	Type	Group
77	Br	326	ARG	Sidechain
77	Br	328	ARG	Sidechain
77	Br	33	HIS	Sidechain
77	Br	349	ARG	Mainchain,Sidechain
77	Br	364	ARG	Sidechain
77	Br	4	ARG	Sidechain
77	Br	48	ARG	Sidechain
77	Br	55	ARG	Sidechain
77	Br	74	ARG	Sidechain
77	Br	79	HIS	Sidechain
78	Bs	2	MET	Mainchain
78	Bs	21	ARG	Sidechain
78	Bs	22	ARG	Sidechain
78	Bs	24	TYR	Sidechain
78	Bs	41	HIS	Sidechain
79	Bt	15	GLU	Peptide
79	Bt	16	ARG	Sidechain
79	Bt	2	VAL	Mainchain
79	Bt	20	HIS	Sidechain
79	Bt	23	PHE	Sidechain
79	Bt	28	TYR	Sidechain
79	Bt	4	TYR	Sidechain
79	Bt	48	SER	Mainchain
79	Bt	58	PHE	Sidechain
79	Bt	61	LYS	Mainchain
79	Bt	90	HIS	Mainchain,Sidechain
79	Bt	91	PHE	Sidechain
80	Bu	108	ARG	Sidechain
80	Bu	12	TYR	Sidechain
80	Bu	13	TYR	Sidechain
80	Bu	132	VAL	Peptide
80	Bu	133	ARG	Sidechain
80	Bu	145	ARG	Sidechain
80	Bu	146	PHE	Sidechain
80	Bu	147	PRO	Peptide
80	Bu	148	PHE	Sidechain
80	Bu	164	ARG	Sidechain
80	Bu	182	ARG	Sidechain
80	Bu	189	TYR	Sidechain
80	Bu	202	ARG	Sidechain
80	Bu	204	ARG	Sidechain
80	Bu	206	PHE	Sidechain

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Mol	Chain	Res	Type	Group
80	Bu	213	TYR	Sidechain
80	Bu	218	ARG	Sidechain
80	Bu	226	ASP	Peptide
80	Bu	23	ARG	Sidechain
80	Bu	232	PHE	Sidechain
80	Bu	236	ILE	Mainchain
80	Bu	238	ALA	Mainchain
80	Bu	24	ARG	Sidechain
80	Bu	249	TYR	Sidechain
80	Bu	257	ARG	Sidechain
80	Bu	264	ARG	Sidechain
80	Bu	278	TYR	Sidechain
80	Bu	30	TYR	Sidechain
80	Bu	33	ARG	Sidechain
80	Bu	44	PHE	Sidechain
80	Bu	54	ARG	Sidechain
80	Bu	79	TYR	Sidechain
80	Bu	81	HIS	Sidechain
80	Bu	99	TYR	Sidechain
81	Bv	147	ARG	Sidechain
81	Bv	15	LYS	Peptide
81	Bv	170	GLU	Peptide
81	Bv	174	TYR	Sidechain
81	Bv	176	HIS	Sidechain
81	Bv	179	PHE	Sidechain
81	Bv	18	ARG	Sidechain
81	Bv	188	HIS	Sidechain
81	Bv	189	ARG	Sidechain
81	Bv	23	TYR	Sidechain
81	Bv	27	ARG	Sidechain
81	Bv	3	ALA	Peptide
81	Bv	42	ARG	Sidechain
81	Bv	44	ARG	Sidechain
81	Bv	46	ARG	Sidechain
81	Bv	64	GLY	Peptide
81	Bv	68	TYR	Peptide,Mainchain
81	Bv	74	ARG	Sidechain
81	Bv	79	ARG	Sidechain
81	Bv	8	ALA	Peptide
81	Bv	87	ARG	Sidechain
82	Bw	103	ARG	Sidechain
82	Bw	138	MET	Mainchain

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Mol	Chain	Res	Type	Group
82	Bw	151	TYR	Sidechain
82	Bw	160	ARG	Sidechain
82	Bw	164	TYR	Sidechain
82	Bw	166	ARG	Sidechain
82	Bw	168	HIS	Mainchain,Sidechain
82	Bw	18	ALA	Mainchain
82	Bw	188	TYR	Mainchain
82	Bw	204	TYR	Sidechain
82	Bw	224	PRO	Peptide,Mainchain
82	Bw	240	GLY	Peptide
82	Bw	245	ARG	Sidechain
82	Bw	251	ARG	Sidechain
82	Bw	255	ARG	Sidechain
82	Bw	27	ALA	Peptide
82	Bw	30	ARG	Sidechain
82	Bw	36	GLN	Mainchain
82	Bw	40	ASN	Mainchain
82	Bw	41	PHE	Sidechain
82	Bw	42	LYS	Peptide,Mainchain
82	Bw	44	ALA	Mainchain
82	Bw	60	TYR	Sidechain
82	Bw	70	TYR	Sidechain
82	Bw	81	ARG	Sidechain
82	Bw	82	ARG	Sidechain
82	Bw	85	ARG	Sidechain
82	Bw	90	TYR	Sidechain
82	Bw	91	TYR	Sidechain
83	Bx	118	ARG	Sidechain
83	Bx	121	ARG	Sidechain
83	Bx	162	ARG	Sidechain
83	Bx	171	ASP	Peptide
83	Bx	198	ARG	Sidechain
83	Bx	215	ASP	Mainchain
83	Bx	234	ARG	Sidechain
83	Bx	265	ARG	Sidechain
83	Bx	269	ASN	Mainchain
83	Bx	273	ALA	Peptide
83	Bx	38	SER	Peptide
83	Bx	43	ARG	Sidechain
83	Bx	45	LYS	Peptide,Mainchain
83	Bx	47	PHE	Sidechain
83	Bx	57	ARG	Sidechain

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Mol	Chain	Res	Type	Group
83	Bx	61	ARG	Sidechain
83	Bx	64	ARG	Sidechain
83	Bx	91	PHE	Sidechain
84	By	120	ARG	Sidechain
84	By	121	ARG	Sidechain
84	By	127	ASP	Peptide,Mainchain
84	By	131	TYR	Sidechain
84	By	132	ARG	Sidechain
84	By	138	VAL	Peptide,Mainchain
84	By	140	ASP	Mainchain
84	By	154	ARG	Sidechain
84	By	171	ARG	Sidechain
84	By	186	GLU	Mainchain
84	By	2	LYS	Mainchain
84	By	38	ARG	Sidechain
84	By	41	GLN	Peptide
84	By	45	ARG	Sidechain
84	By	47	ASN	Mainchain
84	By	51	ARG	Sidechain
84	By	6	HIS	Peptide,Sidechain
84	By	7	ASP	Peptide,Mainchain
84	By	74	HIS	Sidechain
84	By	76	ARG	Sidechain
84	By	8	GLN	Peptide
84	By	94	TYR	Sidechain
84	By	96	HIS	Sidechain

5.2 Too-close contacts [i](#)

Due to software issues we are unable to calculate clashes - this section is therefore empty.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

The Analysed column shows the number of residues for which the backbone conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A0	217/256 (85%)	180 (83%)	22 (10%)	15 (7%)	1	11
2	A1	246/273 (90%)	189 (77%)	38 (15%)	19 (8%)	1	9
3	A2	185/190 (97%)	159 (86%)	17 (9%)	9 (5%)	2	16
4	A3	248/250 (99%)	204 (82%)	24 (10%)	20 (8%)	1	9
5	A4	190/202 (94%)	148 (78%)	24 (13%)	18 (10%)	0	8
6	A5	191/220 (87%)	158 (83%)	18 (9%)	15 (8%)	1	9
7	A6	185/190 (97%)	135 (73%)	30 (16%)	20 (11%)	0	6
8	A7	313/318 (98%)	256 (82%)	39 (12%)	18 (6%)	1	13
9	A8	40/57 (70%)	34 (85%)	3 (8%)	3 (8%)	1	10
10	A9	64/153 (42%)	50 (78%)	9 (14%)	5 (8%)	1	9
11	AC	202/277 (73%)	162 (80%)	25 (12%)	15 (7%)	1	10
12	AD	102/172 (59%)	79 (78%)	9 (9%)	14 (14%)	0	4
13	AE	158/174 (91%)	126 (80%)	12 (8%)	20 (13%)	0	4
14	AF	119/144 (83%)	105 (88%)	11 (9%)	3 (2%)	4	26
15	AG	139/151 (92%)	120 (86%)	12 (9%)	7 (5%)	1	15
16	AH	124/144 (86%)	99 (80%)	14 (11%)	11 (9%)	0	8
17	AI	132/152 (87%)	100 (76%)	21 (16%)	11 (8%)	0	9
18	AJ	127/130 (98%)	114 (90%)	7 (6%)	6 (5%)	2	16
19	AK	146/149 (98%)	118 (81%)	18 (12%)	10 (7%)	1	11
20	AL	125/142 (88%)	94 (75%)	19 (15%)	12 (10%)	0	7
21	AM	151/153 (99%)	114 (76%)	16 (11%)	21 (14%)	0	4
22	AO	147/167 (88%)	128 (87%)	10 (7%)	9 (6%)	1	13
23	AP	222/266 (84%)	171 (77%)	29 (13%)	22 (10%)	0	7
24	AQ	103/117 (88%)	86 (84%)	10 (10%)	7 (7%)	1	11
25	AR	79/194 (41%)	59 (75%)	12 (15%)	8 (10%)	0	7
26	AS	140/143 (98%)	111 (79%)	18 (13%)	11 (8%)	1	9
27	AT	129/137 (94%)	100 (78%)	15 (12%)	14 (11%)	0	6
28	AU	84/113 (74%)	66 (79%)	12 (14%)	6 (7%)	1	10
29	AV	99/111 (89%)	64 (65%)	20 (20%)	15 (15%)	0	3
30	AW	81/86 (94%)	61 (75%)	11 (14%)	9 (11%)	0	6
31	AX	204/214 (95%)	176 (86%)	17 (8%)	11 (5%)	1	14
32	AY	63/66 (96%)	46 (73%)	8 (13%)	9 (14%)	0	3

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
33	AZ	66/103 (64%)	56 (85%)	7 (11%)	3 (4%)	2	17
42	BI	190/193 (98%)	126 (66%)	37 (20%)	27 (14%)	0	3
43	BJ	212/214 (99%)	186 (88%)	20 (9%)	6 (3%)	4	24
44	BK	210/213 (99%)	161 (77%)	26 (12%)	23 (11%)	0	6
45	BL	168/194 (87%)	134 (80%)	21 (12%)	13 (8%)	1	9
46	BM	137/164 (84%)	103 (75%)	25 (18%)	9 (7%)	1	12
47	BN	214/218 (98%)	158 (74%)	20 (9%)	36 (17%)	0	3
48	BO	199/222 (90%)	164 (82%)	21 (11%)	14 (7%)	1	11
49	BP	182/189 (96%)	149 (82%)	19 (10%)	14 (8%)	1	9
50	BQ	201/221 (91%)	160 (80%)	21 (10%)	20 (10%)	0	7
51	BR	153/166 (92%)	121 (79%)	21 (14%)	11 (7%)	1	10
52	BS	177/179 (99%)	133 (75%)	15 (8%)	29 (16%)	0	3
53	BT	198/260 (76%)	168 (85%)	20 (10%)	10 (5%)	1	15
54	BU	156/159 (98%)	118 (76%)	19 (12%)	19 (12%)	0	4
55	BV	102/130 (78%)	84 (82%)	13 (13%)	5 (5%)	2	16
56	BW	136/139 (98%)	103 (76%)	18 (13%)	15 (11%)	0	6
57	BX	119/164 (73%)	97 (82%)	9 (8%)	13 (11%)	0	6
58	BY	98/125 (78%)	81 (83%)	9 (9%)	8 (8%)	1	9
59	BZ	123/143 (86%)	104 (85%)	12 (10%)	7 (6%)	1	14
60	Ba	130/133 (98%)	93 (72%)	21 (16%)	16 (12%)	0	4
61	Bb	142/145 (98%)	108 (76%)	19 (13%)	15 (11%)	0	6
62	Bc	139/146 (95%)	103 (74%)	18 (13%)	18 (13%)	0	4
63	Bd	68/71 (96%)	50 (74%)	8 (12%)	10 (15%)	0	3
64	Be	184/260 (71%)	135 (73%)	32 (17%)	17 (9%)	0	8
65	Bf	412/429 (96%)	296 (72%)	66 (16%)	50 (12%)	0	4
66	Bg	94/105 (90%)	88 (94%)	5 (5%)	1 (1%)	12	46
67	Bh	186/188 (99%)	155 (83%)	13 (7%)	18 (10%)	0	7
68	Bi	127/132 (96%)	98 (77%)	18 (14%)	11 (9%)	0	9
69	Bj	160/170 (94%)	123 (77%)	22 (14%)	15 (9%)	0	8
70	Bk	82/127 (65%)	56 (68%)	14 (17%)	12 (15%)	0	3
71	Bl	114/149 (76%)	84 (74%)	10 (9%)	20 (18%)	0	2

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
72	Bm	105/109 (96%)	79 (75%)	8 (8%)	18 (17%)	0	2
73	Bn	81/84 (96%)	56 (69%)	15 (18%)	10 (12%)	0	4
74	Bo	90/93 (97%)	71 (79%)	6 (7%)	13 (14%)	0	3
75	Bp	79/82 (96%)	59 (75%)	10 (13%)	10 (13%)	0	4
76	Bq	48/51 (94%)	34 (71%)	6 (12%)	8 (17%)	0	3
77	Br	366/374 (98%)	274 (75%)	56 (15%)	36 (10%)	0	7
78	Bs	50/128 (39%)	44 (88%)	4 (8%)	2 (4%)	2	18
79	Bt	103/106 (97%)	75 (73%)	16 (16%)	12 (12%)	0	4
80	Bu	297/308 (96%)	216 (73%)	46 (16%)	35 (12%)	0	4
81	Bv	154/192 (80%)	114 (74%)	25 (16%)	15 (10%)	0	7
82	Bw	255/257 (99%)	200 (78%)	33 (13%)	22 (9%)	0	9
83	Bx	238/276 (86%)	188 (79%)	26 (11%)	24 (10%)	0	7
84	By	187/189 (99%)	155 (83%)	15 (8%)	17 (9%)	0	8
All	All	11687/13211 (88%)	9172 (78%)	1415 (12%)	1100 (9%)	1	8

All (1100) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A0	37	ASN
1	A0	119	TRP
1	A0	145	THR
1	A0	212	LEU
1	A0	225	PHE
2	A1	65	GLY
2	A1	66	LEU
2	A1	171	LYS
2	A1	172	ILE
2	A1	174	ASN
3	A2	39	ALA
3	A2	43	GLN
3	A2	149	PHE
4	A3	44	GLU
4	A3	71	PRO
4	A3	72	SER
4	A3	108	ASP
4	A3	161	ILE
4	A3	171	LYS

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Mol	Chain	Res	Type
4	A3	177	PRO
4	A3	178	LYS
5	A4	3	ALA
5	A4	36	HIS
5	A4	80	GLN
5	A4	114	LEU
5	A4	191	MET
6	A5	23	LYS
6	A5	25	MET
6	A5	36	THR
6	A5	41	ARG
6	A5	173	LYS
7	A6	14	PRO
7	A6	35	LEU
7	A6	64	ASN
7	A6	131	ARG
7	A6	139	ILE
7	A6	148	ARG
7	A6	149	VAL
7	A6	155	ILE
7	A6	157	PHE
7	A6	168	PRO
7	A6	176	ARG
8	A7	6	GLU
8	A7	30	LYS
8	A7	47	PRO
8	A7	51	SER
8	A7	89	HIS
8	A7	131	ASN
8	A7	215	ASP
9	A8	34	LYS
10	A9	100	ALA
10	A9	125	ASN
11	AC	132	THR
11	AC	232	GLU
11	AC	239	PHE
11	AC	241	ARG
12	AD	6	PRO
13	AE	18	VAL
13	AE	21	GLU
13	AE	29	ALA
13	AE	58	PHE

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Mol	Chain	Res	Type
13	AE	60	THR
13	AE	166	LYS
13	AE	172	SER
15	AG	5	HIS
15	AG	23	PRO
15	AG	25	TRP
15	AG	62	GLN
16	AH	13	SER
16	AH	60	ASP
16	AH	61	GLU
17	AI	56	ALA
17	AI	148	PHE
18	AJ	29	PRO
18	AJ	30	SER
19	AK	10	LYS
19	AK	31	PRO
19	AK	131	GLU
20	AL	23	LYS
20	AL	41	ALA
20	AL	83	ASP
21	AM	27	VAL
21	AM	34	VAL
21	AM	35	LYS
21	AM	39	ILE
21	AM	43	TYR
21	AM	90	ASP
21	AM	91	PRO
21	AM	119	ARG
21	AM	143	ARG
21	AM	146	THR
22	AO	52	GLU
23	AP	87	ARG
23	AP	102	ALA
23	AP	118	ASN
23	AP	162	HIS
23	AP	184	ARG
24	AQ	64	LYS
24	AQ	67	CYS
24	AQ	68	GLY
25	AR	23	HIS
25	AR	30	THR
26	AS	61	GLN

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Mol	Chain	Res	Type
26	AS	65	ALA
26	AS	87	ASN
26	AS	107	ARG
26	AS	136	GLN
27	AT	35	PRO
27	AT	42	PRO
27	AT	73	GLY
27	AT	75	LYS
27	AT	127	LYS
28	AU	47	MET
28	AU	103	LYS
29	AV	17	ARG
29	AV	26	PHE
29	AV	47	LEU
29	AV	91	ARG
29	AV	93	VAL
30	AW	6	SER
30	AW	7	ASP
30	AW	21	LYS
30	AW	60	CYS
30	AW	61	ALA
31	AX	64	ARG
31	AX	80	GLU
31	AX	192	SER
31	AX	205	THR
32	AY	4	VAL
32	AY	62	PRO
32	AY	64	LYS
33	AZ	84	CYS
42	BI	4	ASP
42	BI	12	LYS
42	BI	19	THR
42	BI	20	TYR
42	BI	21	SER
42	BI	41	ASN
42	BI	85	PRO
42	BI	97	VAL
42	BI	104	ALA
42	BI	150	LYS
42	BI	178	LYS
42	BI	179	GLU
42	BI	192	HIS

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Mol	Chain	Res	Type
43	BJ	58	PRO
44	BK	31	ILE
44	BK	81	ASN
44	BK	93	PRO
44	BK	102	MET
44	BK	119	TYR
44	BK	132	GLY
44	BK	172	GLY
44	BK	176	ILE
45	BL	16	MET
45	BL	18	GLU
45	BL	32	GLU
45	BL	59	LEU
45	BL	71	LYS
45	BL	95	GLU
46	BM	11	ILE
47	BN	6	ASN
47	BN	9	PRO
47	BN	11	VAL
47	BN	14	ARG
47	BN	19	PRO
47	BN	20	CYS
47	BN	23	GLN
47	BN	66	PRO
47	BN	67	THR
47	BN	81	SER
47	BN	90	VAL
47	BN	155	ARG
47	BN	174	PRO
47	BN	176	LYS
47	BN	201	ARG
48	BO	88	PRO
48	BO	136	ARG
48	BO	140	PRO
48	BO	151	GLU
48	BO	203	GLU
49	BP	6	TYR
49	BP	32	ASP
49	BP	47	TRP
50	BQ	58	ARG
50	BQ	104	ALA
50	BQ	111	LEU

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Mol	Chain	Res	Type
50	BQ	175	HIS
50	BQ	199	SER
51	BR	40	LYS
51	BR	54	LYS
52	BS	2	VAL
52	BS	3	ARG
52	BS	5	HIS
52	BS	6	LEU
52	BS	15	GLU
52	BS	71	LEU
52	BS	118	ALA
52	BS	155	GLN
52	BS	157	ARG
52	BS	158	VAL
52	BS	161	PRO
52	BS	166	VAL
52	BS	167	ILE
52	BS	169	VAL
52	BS	173	SER
53	BT	5	LYS
53	BT	75	HIS
53	BT	130	ASN
53	BT	134	ASN
53	BT	179	LYS
53	BT	191	ASP
54	BU	19	PHE
54	BU	46	ALA
54	BU	47	ALA
54	BU	56	TYR
54	BU	99	SER
54	BU	126	PRO
54	BU	149	ARG
54	BU	150	THR
56	BW	7	ASN
56	BW	11	CYS
56	BW	91	ASP
56	BW	114	SER
56	BW	120	VAL
57	BX	60	TYR
57	BX	69	SER
57	BX	72	VAL
57	BX	75	ILE

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Mol	Chain	Res	Type
58	BY	68	THR
58	BY	99	ALA
59	BZ	47	VAL
59	BZ	48	ARG
59	BZ	51	ASP
60	Ba	14	SER
60	Ba	53	VAL
60	Ba	56	MET
60	Ba	67	VAL
60	Ba	96	ILE
60	Ba	101	ALA
60	Ba	121	ALA
60	Ba	126	TRP
61	Bb	35	ALA
61	Bb	50	PRO
61	Bb	71	PRO
61	Bb	107	SER
61	Bb	111	GLY
62	Bc	16	SER
62	Bc	17	LYS
62	Bc	41	ARG
62	Bc	66	SER
62	Bc	84	PRO
62	Bc	95	VAL
62	Bc	134	MET
63	Bd	5	LYS
63	Bd	21	ILE
63	Bd	26	PRO
63	Bd	33	LYS
63	Bd	50	ASN
64	Be	70	ARG
64	Be	84	THR
64	Be	96	LEU
64	Be	217	GLN
65	Bf	61	ARG
65	Bf	76	ILE
65	Bf	81	ARG
65	Bf	86	ASP
65	Bf	133	PRO
65	Bf	134	PRO
65	Bf	225	ASN
65	Bf	232	ALA

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Mol	Chain	Res	Type
65	Bf	243	THR
65	Bf	323	ALA
65	Bf	348	GLU
65	Bf	349	PRO
65	Bf	360	ALA
65	Bf	387	ALA
65	Bf	405	SER
65	Bf	427	ARG
66	Bg	108	ASP
67	Bh	7	LYS
67	Bh	97	LYS
67	Bh	100	THR
67	Bh	122	THR
67	Bh	140	VAL
67	Bh	159	GLU
67	Bh	174	PRO
67	Bh	175	VAL
68	Bi	12	LYS
68	Bi	50	GLN
68	Bi	87	LEU
69	Bj	4	PRO
69	Bj	5	ARG
69	Bj	14	TYR
69	Bj	16	THR
69	Bj	43	PRO
69	Bj	47	TRP
69	Bj	85	HIS
70	Bk	75	SER
70	Bk	80	ARG
70	Bk	89	ALA
70	Bk	92	THR
70	Bk	97	LEU
70	Bk	104	LYS
70	Bk	122	ILE
71	Bl	36	PRO
71	Bl	89	ILE
71	Bl	91	ARG
71	Bl	93	VAL
71	Bl	94	ARG
71	Bl	96	SER
71	Bl	99	PRO
71	Bl	102	ARG

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Mol	Chain	Res	Type
71	Bl	130	SER
71	Bl	132	PRO
71	Bl	146	PRO
72	Bm	16	ILE
72	Bm	17	ALA
72	Bm	27	ARG
72	Bm	29	ALA
72	Bm	33	SER
72	Bm	39	ALA
72	Bm	43	LYS
72	Bm	87	SER
73	Bn	15	THR
73	Bn	36	ALA
73	Bn	40	PRO
73	Bn	77	HIS
74	Bo	19	GLY
74	Bo	68	ALA
75	Bp	20	ALA
75	Bp	63	SER
75	Bp	71	SER
75	Bp	76	ASN
76	Bq	6	PRO
76	Bq	35	ILE
76	Bq	38	ASN
76	Bq	39	GLU
76	Bq	40	LYS
77	Br	13	SER
77	Br	59	MET
77	Br	93	ASN
77	Br	94	MET
77	Br	143	ILE
77	Br	206	PRO
77	Br	216	THR
77	Br	228	ALA
77	Br	230	VAL
77	Br	278	PRO
77	Br	293	SER
77	Br	300	LEU
77	Br	304	LYS
77	Br	308	LYS
77	Br	344	GLN
77	Br	366	ASN

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Mol	Chain	Res	Type
78	Bs	3	GLU
79	Bt	5	PRO
79	Bt	14	ASP
79	Bt	17	CYS
79	Bt	33	ALA
79	Bt	48	SER
79	Bt	62	ALA
79	Bt	92	GLU
80	Bu	57	ASN
80	Bu	58	ARG
80	Bu	121	GLY
80	Bu	142	ASP
80	Bu	146	PHE
80	Bu	227	GLU
80	Bu	229	THR
80	Bu	239	LYS
80	Bu	264	ARG
80	Bu	269	ARG
80	Bu	299	VAL
81	Bv	10	GLU
81	Bv	18	ARG
81	Bv	59	PRO
81	Bv	170	GLU
81	Bv	177	SER
81	Bv	179	PHE
81	Bv	185	ASP
82	Bw	4	THR
82	Bw	15	VAL
82	Bw	19	LYS
82	Bw	22	PRO
82	Bw	29	LYS
82	Bw	225	PRO
82	Bw	235	HIS
82	Bw	242	TYR
83	Bx	44	PRO
83	Bx	47	PHE
83	Bx	49	ILE
83	Bx	52	ASP
83	Bx	56	ALA
83	Bx	90	GLN
83	Bx	195	ASP
83	Bx	216	VAL

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Mol	Chain	Res	Type
83	Bx	271	ALA
83	Bx	272	ALA
84	By	7	ASP
84	By	8	GLN
84	By	12	PRO
84	By	19	VAL
84	By	43	ASP
84	By	57	ARG
84	By	59	PHE
1	A0	53	GLY
1	A0	108	THR
2	A1	32	PRO
2	A1	33	HIS
2	A1	134	PRO
2	A1	151	THR
2	A1	245	ASN
3	A2	38	SER
3	A2	54	ILE
3	A2	150	ARG
4	A3	29	GLY
4	A3	45	ALA
4	A3	123	PRO
4	A3	136	LEU
4	A3	151	GLY
4	A3	155	ASP
5	A4	47	HIS
5	A4	61	LYS
5	A4	105	THR
6	A5	9	HIS
6	A5	13	ILE
6	A5	16	GLY
6	A5	22	ARG
6	A5	168	HIS
7	A6	17	PRO
7	A6	97	SER
7	A6	135	VAL
7	A6	145	PHE
7	A6	186	GLY
8	A7	50	HIS
8	A7	55	SER
8	A7	154	TRP
9	A8	22	CYS

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Mol	Chain	Res	Type
10	A9	129	GLY
11	AC	43	VAL
11	AC	79	SER
11	AC	176	VAL
11	AC	203	GLY
12	AD	88	TYR
12	AD	92	ASP
12	AD	95	PRO
12	AD	103	LYS
13	AE	24	TYR
13	AE	55	GLY
13	AE	75	PRO
13	AE	173	LYS
16	AH	41	SER
17	AI	55	HIS
17	AI	74	VAL
17	AI	81	LYS
18	AJ	82	THR
18	AJ	98	GLN
19	AK	30	ALA
19	AK	49	THR
20	AL	99	ILE
20	AL	122	PRO
21	AM	85	LEU
21	AM	120	ALA
21	AM	133	ARG
21	AM	145	LYS
21	AM	149	VAL
22	AO	57	SER
22	AO	64	PRO
22	AO	159	LYS
22	AO	164	THR
23	AP	68	SER
23	AP	82	ARG
23	AP	117	GLY
23	AP	188	ILE
23	AP	215	ARG
24	AQ	65	THR
25	AR	12	VAL
25	AR	81	GLY
26	AS	37	LYS
26	AS	105	PHE

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Mol	Chain	Res	Type
26	AS	126	SER
27	AT	10	VAL
27	AT	115	LYS
28	AU	33	TRP
28	AU	75	LEU
28	AU	101	SER
29	AV	6	ARG
29	AV	28	CYS
29	AV	49	GLN
29	AV	64	SER
29	AV	88	VAL
30	AW	52	THR
31	AX	93	ARG
31	AX	194	ILE
33	AZ	82	GLY
42	BI	10	LYS
42	BI	99	MET
42	BI	168	HIS
44	BK	10	ARG
44	BK	14	ASN
44	BK	57	LEU
44	BK	108	ALA
44	BK	115	MET
44	BK	160	PRO
45	BL	154	ARG
45	BL	159	HIS
46	BM	40	LYS
47	BN	78	ARG
47	BN	130	LEU
47	BN	139	GLN
47	BN	156	SER
47	BN	172	GLU
47	BN	202	ALA
48	BO	85	LYS
48	BO	91	HIS
48	BO	142	ALA
48	BO	210	GLN
49	BP	33	ALA
49	BP	92	VAL
49	BP	159	ALA
49	BP	184	VAL
50	BQ	51	HIS

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Mol	Chain	Res	Type
50	BQ	72	THR
50	BQ	99	GLY
50	BQ	108	GLY
50	BQ	130	LEU
50	BQ	154	PRO
50	BQ	174	LYS
50	BQ	214	ARG
51	BR	8	PRO
51	BR	41	LEU
51	BR	65	GLY
52	BS	26	VAL
52	BS	138	ARG
52	BS	178	VAL
53	BT	55	VAL
53	BT	74	ARG
53	BT	77	GLY
54	BU	18	LYS
54	BU	131	SER
54	BU	141	LYS
54	BU	145	VAL
54	BU	157	ILE
55	BV	44	LEU
55	BV	61	LYS
56	BW	4	ASP
56	BW	8	VAL
56	BW	18	ALA
56	BW	72	ARG
56	BW	118	GLY
56	BW	136	PRO
57	BX	47	LYS
59	BZ	7	ARG
59	BZ	50	ASP
60	Ba	3	PHE
60	Ba	18	ALA
60	Ba	34	GLU
60	Ba	54	ARG
60	Ba	60	SER
61	Bb	15	THR
61	Bb	32	ARG
61	Bb	51	GLY
61	Bb	65	LYS
61	Bb	113	GLY

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Mol	Chain	Res	Type
62	Bc	39	THR
62	Bc	59	ARG
62	Bc	97	ASP
63	Bd	23	PRO
63	Bd	24	PRO
63	Bd	29	MET
64	Be	117	GLU
64	Be	118	ALA
64	Be	162	CYS
64	Be	173	GLY
64	Be	180	LEU
64	Be	216	HIS
65	Bf	242	GLY
65	Bf	268	SER
65	Bf	295	PRO
65	Bf	304	LYS
65	Bf	315	ARG
65	Bf	420	SER
65	Bf	444	ASP
67	Bh	3	ARG
67	Bh	44	LYS
67	Bh	56	VAL
67	Bh	99	ARG
67	Bh	124	ASP
67	Bh	160	GLY
67	Bh	162	LYS
69	Bj	48	VAL
69	Bj	56	GLY
69	Bj	115	GLN
70	Bk	103	GLU
70	Bk	126	LYS
71	Bl	103	SER
72	Bm	8	ALA
72	Bm	22	GLY
72	Bm	25	THR
72	Bm	34	SER
72	Bm	40	LEU
72	Bm	78	LEU
73	Bn	41	ARG
73	Bn	81	ASP
74	Bo	5	THR
74	Bo	8	MET

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Mol	Chain	Res	Type
74	Bo	40	SER
74	Bo	72	SER
74	Bo	92	SER
75	Bp	18	LYS
75	Bp	60	ILE
75	Bp	72	ARG
76	Bq	22	PRO
76	Bq	47	THR
77	Br	14	GLU
77	Br	63	ALA
77	Br	217	ARG
77	Br	232	SER
77	Br	269	SER
77	Br	314	ARG
78	Bs	34	CYS
79	Bt	75	SER
79	Bt	103	ASP
79	Bt	104	PRO
80	Bu	8	LYS
80	Bu	115	LEU
80	Bu	132	VAL
80	Bu	139	GLN
80	Bu	154	VAL
80	Bu	160	THR
80	Bu	194	ASP
80	Bu	298	ALA
81	Bv	41	GLY
82	Bw	6	LEU
82	Bw	94	ALA
82	Bw	97	LYS
82	Bw	174	GLN
82	Bw	234	ARG
82	Bw	240	GLY
82	Bw	241	ASP
83	Bx	42	ALA
83	Bx	95	LEU
83	Bx	136	THR
83	Bx	139	THR
83	Bx	246	TRP
83	Bx	263	ARG
83	Bx	267	ALA
84	By	5	SER

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Mol	Chain	Res	Type
84	By	21	ASP
84	By	45	ARG
84	By	105	ASN
84	By	168	LYS
1	A0	34	ALA
1	A0	36	LYS
1	A0	78	ASN
1	A0	206	ILE
1	A0	226	ASP
2	A1	19	LYS
3	A2	36	PRO
3	A2	173	SER
5	A4	100	ALA
5	A4	120	SER
6	A5	33	PRO
6	A5	38	LEU
6	A5	156	SER
7	A6	117	LEU
8	A7	11	GLY
8	A7	28	ALA
8	A7	65	GLY
8	A7	239	PRO
8	A7	276	PRO
9	A8	35	TYR
10	A9	92	ARG
10	A9	124	PRO
11	AC	58	HIS
11	AC	102	ALA
11	AC	140	ILE
11	AC	164	ARG
12	AD	29	PRO
12	AD	31	GLY
12	AD	35	GLY
12	AD	45	PRO
12	AD	67	ALA
13	AE	38	SER
13	AE	69	TYR
13	AE	164	ALA
14	AF	76	TYR
16	AH	29	SER
16	AH	40	MET
16	AH	59	ARG

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Mol	Chain	Res	Type
16	AH	92	ALA
16	AH	131	ASP
17	AI	78	GLU
18	AJ	59	GLY
19	AK	5	LYS
20	AL	6	THR
20	AL	65	PRO
20	AL	92	ASP
21	AM	28	PRO
22	AO	163	GLU
23	AP	70	PRO
23	AP	185	GLY
23	AP	240	TRP
23	AP	248	ASP
25	AR	15	ASP
26	AS	135	ARG
27	AT	65	PHE
27	AT	103	MET
27	AT	107	LYS
29	AV	5	ARG
29	AV	100	TYR
30	AW	53	SER
30	AW	83	LYS
31	AX	61	LYS
32	AY	10	ARG
42	BI	5	LEU
42	BI	8	VAL
42	BI	61	PRO
42	BI	149	ARG
42	BI	152	GLY
43	BJ	11	GLU
43	BJ	27	GLU
43	BJ	45	ARG
43	BJ	146	LEU
44	BK	100	ASN
44	BK	169	LYS
45	BL	17	ARG
45	BL	115	GLU
45	BL	147	HIS
46	BM	60	CYS
47	BN	104	ARG
47	BN	175	ARG

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Mol	Chain	Res	Type
47	BN	180	GLU
48	BO	132	ARG
48	BO	166	TRP
48	BO	199	ASN
48	BO	205	LEU
48	BO	209	PRO
49	BP	17	ARG
50	BQ	74	GLN
50	BQ	129	ASN
50	BQ	203	PRO
51	BR	5	SER
52	BS	68	ASP
52	BS	84	SER
53	BT	133	ARG
54	BU	22	HIS
54	BU	32	THR
54	BU	124	PRO
56	BW	138	ILE
57	BX	57	PRO
57	BX	63	PRO
57	BX	66	VAL
57	BX	102	ASN
58	BY	10	HIS
58	BY	64	ARG
58	BY	71	VAL
61	Bb	116	GLN
62	Bc	65	LYS
62	Bc	67	GLY
62	Bc	87	LYS
62	Bc	103	ALA
63	Bd	25	LEU
63	Bd	35	GLY
64	Be	77	VAL
64	Be	212	GLY
65	Bf	65	LEU
65	Bf	82	SER
65	Bf	314	SER
65	Bf	353	THR
65	Bf	443	LYS
67	Bh	27	ALA
68	Bi	14	ARG
68	Bi	20	ARG

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Mol	Chain	Res	Type
68	Bi	25	LEU
68	Bi	49	GLY
68	Bi	59	GLY
68	Bi	126	LYS
69	Bj	7	GLN
70	Bk	77	LYS
71	Bl	97	LYS
72	Bm	7	GLU
72	Bm	23	TYR
72	Bm	77	ALA
73	Bn	32	GLU
74	Bo	51	ALA
77	Br	91	PHE
77	Br	110	TRP
77	Br	146	VAL
77	Br	220	ARG
77	Br	233	LEU
77	Br	309	ARG
79	Bt	6	LYS
79	Bt	58	PHE
80	Bu	92	LEU
80	Bu	120	VAL
80	Bu	131	ALA
80	Bu	144	SER
80	Bu	163	ALA
80	Bu	222	SER
80	Bu	262	LYS
80	Bu	283	LEU
81	Bv	28	LYS
82	Bw	111	PRO
82	Bw	233	ARG
83	Bx	114	THR
84	By	13	GLU
84	By	175	ASP
1	A0	79	GLU
1	A0	208	PRO
2	A1	51	TYR
2	A1	139	HIS
2	A1	147	PRO
4	A3	20	ASP
4	A3	90	GLY
4	A3	153	THR

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Mol	Chain	Res	Type
5	A4	113	LYS
7	A6	16	ARG
7	A6	133	ILE
8	A7	194	GLY
12	AD	2	THR
12	AD	44	LEU
12	AD	102	HIS
13	AE	19	GLN
13	AE	70	ILE
13	AE	80	VAL
15	AG	26	LEU
15	AG	107	LYS
17	AI	132	PRO
17	AI	133	VAL
18	AJ	3	MET
19	AK	32	GLN
19	AK	133	LYS
19	AK	148	TYR
20	AL	24	LEU
21	AM	129	GLY
21	AM	134	GLY
21	AM	142	ARG
22	AO	98	SER
23	AP	183	PRO
23	AP	234	PHE
24	AQ	49	GLY
24	AQ	66	PRO
25	AR	47	ASN
25	AR	85	ILE
26	AS	110	HIS
27	AT	37	TRP
27	AT	110	ALA
27	AT	131	LYS
28	AU	80	ALA
30	AW	3	PHE
31	AX	58	LEU
32	AY	22	SER
32	AY	46	ALA
42	BI	134	ALA
44	BK	13	LYS
44	BK	206	ILE
45	BL	70	GLU

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Mol	Chain	Res	Type
46	BM	90	ARG
46	BM	93	LYS
46	BM	141	CYS
47	BN	8	ILE
47	BN	13	GLN
47	BN	16	HIS
47	BN	28	LYS
47	BN	188	LYS
49	BP	136	PRO
50	BQ	31	LYS
50	BQ	143	THR
50	BQ	176	ARG
52	BS	146	PRO
52	BS	162	ARG
52	BS	176	ALA
54	BU	81	ARG
55	BV	45	SER
55	BV	60	ARG
55	BV	115	ASP
56	BW	44	TYR
56	BW	93	THR
58	BY	17	HIS
58	BY	87	VAL
59	BZ	56	LYS
60	Ba	57	SER
60	Ba	91	GLU
62	Bc	40	LYS
62	Bc	42	GLN
62	Bc	74	LYS
64	Be	83	PHE
64	Be	120	VAL
64	Be	229	ALA
65	Bf	60	PRO
65	Bf	66	GLY
65	Bf	148	VAL
65	Bf	174	LYS
65	Bf	322	ARG
65	Bf	347	MET
65	Bf	350	ASN
65	Bf	351	GLN
65	Bf	354	THR
65	Bf	390	ARG

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Mol	Chain	Res	Type
67	Bh	161	GLY
68	Bi	10	ILE
68	Bi	100	THR
70	Bk	107	LYS
71	Bl	51	LEU
71	Bl	87	TYR
71	Bl	88	LYS
71	Bl	119	SER
73	Bn	20	ARG
74	Bo	46	ALA
75	Bp	30	LYS
77	Br	147	ALA
77	Br	174	GLY
77	Br	176	ILE
80	Bu	5	LYS
80	Bu	260	PRO
81	Bv	7	LYS
82	Bw	43	LYS
83	Bx	249	LEU
84	By	22	ARG
2	A1	74	ARG
2	A1	92	THR
2	A1	238	LYS
5	A4	37	LYS
5	A4	48	ILE
5	A4	104	ILE
5	A4	108	PRO
6	A5	35	ASN
8	A7	45	PRO
11	AC	109	ASP
11	AC	141	PRO
11	AC	195	ILE
14	AF	121	CYS
16	AH	121	ARG
17	AI	25	ARG
19	AK	142	THR
20	AL	66	VAL
20	AL	67	ARG
20	AL	85	VAL
22	AO	54	VAL
22	AO	63	ALA
23	AP	190	ALA

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Mol	Chain	Res	Type
23	AP	257	LEU
24	AQ	42	ASN
25	AR	17	TYR
27	AT	9	GLU
29	AV	90	ALA
31	AX	77	ASN
42	BI	15	VAL
42	BI	105	LEU
42	BI	139	THR
43	BJ	40	PRO
44	BK	15	LYS
44	BK	110	ARG
44	BK	131	ILE
47	BN	80	PHE
47	BN	103	ASP
47	BN	167	VAL
49	BP	152	LYS
49	BP	157	THR
49	BP	158	PRO
50	BQ	218	LEU
51	BR	10	VAL
51	BR	61	LYS
51	BR	78	GLN
52	BS	136	VAL
52	BS	137	ARG
52	BS	165	ARG
52	BS	168	PHE
54	BU	54	HIS
54	BU	142	LYS
56	BW	116	ILE
58	BY	37	ARG
59	BZ	57	ARG
61	Bb	70	LYS
61	Bb	91	LYS
61	Bb	115	LEU
64	Be	87	SER
64	Be	109	GLU
65	Bf	59	HIS
65	Bf	224	ASN
65	Bf	266	GLN
65	Bf	377	ASN
65	Bf	423	ILE

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Mol	Chain	Res	Type
65	Bf	430	THR
69	Bj	6	VAL
69	Bj	79	TYR
70	Bk	102	ASN
71	Bl	76	VAL
71	Bl	131	VAL
73	Bn	63	ARG
74	Bo	60	CYS
74	Bo	70	THR
75	Bp	48	ALA
75	Bp	80	PRO
77	Br	270	THR
80	Bu	38	LEU
80	Bu	245	MET
80	Bu	270	PRO
81	Bv	58	GLY
81	Bv	186	ALA
82	Bw	230	ARG
83	Bx	48	GLY
83	Bx	128	GLU
83	Bx	148	GLY
84	By	116	GLU
1	A0	223	PRO
2	A1	34	LYS
3	A2	172	ASN
4	A3	91	TYR
4	A3	135	ARG
6	A5	59	ARG
12	AD	30	LEU
16	AH	12	SER
17	AI	60	ALA
23	AP	230	LYS
32	AY	27	LYS
32	AY	61	PRO
42	BI	100	THR
47	BN	164	GLY
47	BN	165	GLY
49	BP	4	ALA
51	BR	68	GLY
57	BX	45	LYS
57	BX	54	PHE
57	BX	67	LYS

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Mol	Chain	Res	Type
61	Bb	118	PRO
62	Bc	37	ASN
65	Bf	298	THR
65	Bf	344	ALA
65	Bf	369	PHE
65	Bf	425	HIS
77	Br	98	GLY
77	Br	271	VAL
77	Br	310	PRO
81	Bv	187	PRO
82	Bw	227	GLY
83	Bx	273	ALA
84	By	112	ASN
2	A1	131	GLY
5	A4	18	PRO
5	A4	44	PRO
8	A7	138	VAL
13	AE	30	VAL
21	AM	13	ILE
23	AP	237	PRO
31	AX	59	GLY
31	AX	201	PRO
32	AY	21	VAL
42	BI	137	ALA
44	BK	204	GLY
46	BM	14	VAL
47	BN	29	VAL
69	Bj	37	GLY
69	Bj	162	ALA
71	Bl	66	VAL
73	Bn	62	GLY
74	Bo	52	VAL
76	Bq	24	PRO
77	Br	75	ILE
13	AE	34	ILE
26	AS	40	PRO
29	AV	79	ILE
44	BK	29	PRO
45	BL	15	PRO
46	BM	88	PRO
47	BN	53	PRO
52	BS	4	PRO

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Mol	Chain	Res	Type
52	BS	152	PRO
57	BX	62	ARG
60	Ba	61	ILE
65	Bf	229	VAL
65	Bf	305	VAL
72	Bm	41	PRO
74	Bo	67	GLY
80	Bu	207	GLY
81	Bv	16	VAL
82	Bw	45	ILE
83	Bx	94	VAL
7	A6	174	VAL
23	AP	45	PRO
33	AZ	83	PRO
51	BR	38	GLY
80	Bu	70	VAL
80	Bu	162	GLY
5	A4	99	VAL
13	AE	46	GLY
14	AF	105	GLY
15	AG	17	PRO
17	AI	61	PRO
21	AM	132	VAL
29	AV	92	PRO
49	BP	154	VAL
67	Bh	176	PRO
82	Bw	104	ILE
4	A3	179	ILE
23	AP	71	ILE
46	BM	104	ILE
81	Bv	54	LEU

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A0	189/218 (87%)	159 (84%)	30 (16%)	2	10
2	A1	209/231 (90%)	172 (82%)	37 (18%)	1	8
3	A2	158/160 (99%)	132 (84%)	26 (16%)	2	10
4	A3	207/207 (100%)	174 (84%)	33 (16%)	2	10
5	A4	176/187 (94%)	144 (82%)	32 (18%)	1	8
6	A5	158/180 (88%)	140 (89%)	18 (11%)	4	17
7	A6	162/166 (98%)	137 (85%)	25 (15%)	2	11
8	A7	264/267 (99%)	232 (88%)	32 (12%)	4	16
9	A8	36/49 (74%)	32 (89%)	4 (11%)	5	18
10	A9	57/126 (45%)	52 (91%)	5 (9%)	8	25
11	AC	179/243 (74%)	143 (80%)	36 (20%)	1	6
12	AD	92/131 (70%)	81 (88%)	11 (12%)	4	16
13	AE	143/156 (92%)	126 (88%)	17 (12%)	4	16
14	AF	102/120 (85%)	99 (97%)	3 (3%)	37	56
15	AG	124/131 (95%)	110 (89%)	14 (11%)	4	17
16	AH	95/112 (85%)	83 (87%)	12 (13%)	3	15
17	AI	110/128 (86%)	102 (93%)	8 (7%)	11	31
18	AJ	108/109 (99%)	90 (83%)	18 (17%)	2	9
19	AK	123/124 (99%)	105 (85%)	18 (15%)	2	12
20	AL	111/122 (91%)	101 (91%)	10 (9%)	8	24
21	AM	133/133 (100%)	121 (91%)	12 (9%)	8	24
22	AO	123/137 (90%)	107 (87%)	16 (13%)	3	14
23	AP	185/204 (91%)	154 (83%)	31 (17%)	1	9
24	AQ	94/104 (90%)	84 (89%)	10 (11%)	5	19
25	AR	66/150 (44%)	55 (83%)	11 (17%)	2	9
26	AS	117/118 (99%)	105 (90%)	12 (10%)	6	20
27	AT	110/116 (95%)	82 (74%)	28 (26%)	0	3
28	AU	73/94 (78%)	67 (92%)	6 (8%)	9	28
29	AV	87/97 (90%)	74 (85%)	13 (15%)	2	12
30	AW	71/75 (95%)	58 (82%)	13 (18%)	1	8

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
31	AX	173/180 (96%)	155 (90%)	18 (10%)	5	19
32	AY	52/53 (98%)	41 (79%)	11 (21%)	1	5
33	AZ	57/84 (68%)	52 (91%)	5 (9%)	8	25
42	BI	164/165 (99%)	140 (85%)	24 (15%)	2	12
43	BJ	201/201 (100%)	189 (94%)	12 (6%)	16	37
44	BK	184/185 (100%)	154 (84%)	30 (16%)	2	10
45	BL	146/167 (87%)	120 (82%)	26 (18%)	1	8
46	BM	114/137 (83%)	106 (93%)	8 (7%)	12	32
47	BN	185/188 (98%)	142 (77%)	43 (23%)	0	4
48	BO	175/195 (90%)	150 (86%)	25 (14%)	2	12
49	BP	156/160 (98%)	139 (89%)	17 (11%)	5	18
50	BQ	176/193 (91%)	147 (84%)	29 (16%)	2	10
51	BR	132/144 (92%)	106 (80%)	26 (20%)	1	7
52	BS	160/160 (100%)	130 (81%)	30 (19%)	1	7
53	BT	170/198 (86%)	144 (85%)	26 (15%)	2	11
54	BU	133/134 (99%)	115 (86%)	18 (14%)	3	13
55	BV	95/116 (82%)	85 (90%)	10 (10%)	5	19
56	BW	107/108 (99%)	86 (80%)	21 (20%)	1	7
57	BX	108/136 (79%)	97 (90%)	11 (10%)	6	20
58	BY	85/102 (83%)	78 (92%)	7 (8%)	9	28
59	BZ	110/125 (88%)	98 (89%)	12 (11%)	5	18
60	Ba	116/117 (99%)	100 (86%)	16 (14%)	3	13
61	Bb	115/116 (99%)	95 (83%)	20 (17%)	1	9
62	Bc	124/130 (95%)	95 (77%)	29 (23%)	0	4
63	Bd	58/59 (98%)	54 (93%)	4 (7%)	13	33
64	Be	145/204 (71%)	122 (84%)	23 (16%)	2	10
65	Bf	349/360 (97%)	305 (87%)	44 (13%)	3	15
66	Bg	84/92 (91%)	72 (86%)	12 (14%)	2	12
67	Bh	162/162 (100%)	136 (84%)	26 (16%)	2	10
68	Bi	113/117 (97%)	96 (85%)	17 (15%)	2	11
69	Bj	130/137 (95%)	105 (81%)	25 (19%)	1	7

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
70	Bk	75/114 (66%)	60 (80%)	15 (20%)	1	6
71	Bl	97/126 (77%)	81 (84%)	16 (16%)	2	10
72	Bm	87/90 (97%)	72 (83%)	15 (17%)	1	9
73	Bn	70/71 (99%)	55 (79%)	15 (21%)	1	5
74	Bo	74/76 (97%)	59 (80%)	15 (20%)	1	6
75	Bp	76/77 (99%)	68 (90%)	8 (10%)	5	19
76	Bq	46/47 (98%)	35 (76%)	11 (24%)	0	4
77	Br	304/310 (98%)	251 (83%)	53 (17%)	1	9
78	Bs	46/111 (41%)	38 (83%)	8 (17%)	1	9
79	Bt	94/95 (99%)	83 (88%)	11 (12%)	4	16
80	Bu	238/247 (96%)	199 (84%)	39 (16%)	2	10
81	Bv	132/160 (82%)	112 (85%)	20 (15%)	2	11
82	Bw	213/213 (100%)	176 (83%)	37 (17%)	1	9
83	Bx	203/229 (89%)	173 (85%)	30 (15%)	2	12
84	By	172/172 (100%)	138 (80%)	34 (20%)	1	7
All	All	10068/11158 (90%)	8575 (85%)	1493 (15%)	5	12

All (1493) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A0	33	VAL
1	A0	65	VAL
1	A0	79	GLU
1	A0	83	TYR
1	A0	96	ARG
1	A0	98	LEU
1	A0	106	ASP
1	A0	110	ASP
1	A0	113	TYR
1	A0	114	TYR
1	A0	116	LEU
1	A0	119	TRP
1	A0	121	THR
1	A0	127	VAL
1	A0	133	ASP
1	A0	142	ILE
1	A0	145	THR

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Mol	Chain	Res	Type
1	A0	150	ASN
1	A0	153	SER
1	A0	159	LYS
1	A0	165	TRP
1	A0	168	MET
1	A0	201	LYS
1	A0	204	ASN
1	A0	207	ILE
1	A0	211	ASP
1	A0	215	ARG
1	A0	221	ARG
1	A0	224	LYS
1	A0	231	ILE
2	A1	17	LEU
2	A1	21	THR
2	A1	23	VAL
2	A1	27	ARG
2	A1	35	LEU
2	A1	39	LEU
2	A1	45	ILE
2	A1	51	TYR
2	A1	56	LEU
2	A1	63	ARG
2	A1	68	CYS
2	A1	83	PHE
2	A1	89	ILE
2	A1	96	PHE
2	A1	101	ASP
2	A1	103	LYS
2	A1	111	VAL
2	A1	112	SER
2	A1	121	MET
2	A1	127	TYR
2	A1	128	THR
2	A1	139	HIS
2	A1	153	ARG
2	A1	157	LEU
2	A1	162	LYS
2	A1	168	ASP
2	A1	211	SER
2	A1	215	PHE
2	A1	221	ASN

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Mol	Chain	Res	Type
2	A1	227	LYS
2	A1	229	MET
2	A1	239	GLN
2	A1	242	LEU
2	A1	245	ASN
2	A1	246	VAL
2	A1	252	GLU
2	A1	253	LYS
3	A2	9	PHE
3	A2	20	THR
3	A2	27	TYR
3	A2	28	ILE
3	A2	30	ARG
3	A2	35	VAL
3	A2	37	HIS
3	A2	47	PHE
3	A2	52	ILE
3	A2	56	GLU
3	A2	60	ASN
3	A2	69	ASN
3	A2	72	LYS
3	A2	84	GLU
3	A2	94	PRO
3	A2	113	ARG
3	A2	120	VAL
3	A2	126	ASP
3	A2	128	SER
3	A2	130	MET
3	A2	139	LEU
3	A2	153	LYS
3	A2	174	TYR
3	A2	184	ARG
3	A2	187	LYS
3	A2	190	ARG
4	A3	7	TYR
4	A3	15	GLN
4	A3	19	THR
4	A3	21	GLU
4	A3	26	VAL
4	A3	32	ARG
4	A3	33	LEU
4	A3	54	ARG

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Mol	Chain	Res	Type
4	A3	57	SER
4	A3	59	LYS
4	A3	66	GLN
4	A3	80	ARG
4	A3	85	PHE
4	A3	88	PHE
4	A3	91	TYR
4	A3	95	ARG
4	A3	115	THR
4	A3	116	VAL
4	A3	122	GLN
4	A3	127	VAL
4	A3	134	ARG
4	A3	140	ARG
4	A3	152	ARG
4	A3	161	ILE
4	A3	173	ARG
4	A3	175	LYS
4	A3	177	PRO
4	A3	180	GLN
4	A3	194	LYS
4	A3	210	ARG
4	A3	215	LEU
4	A3	237	ASN
4	A3	239	GLN
5	A4	2	SER
5	A4	6	HIS
5	A4	10	LEU
5	A4	14	LYS
5	A4	27	ARG
5	A4	29	LEU
5	A4	30	PHE
5	A4	32	LEU
5	A4	49	ASN
5	A4	50	THR
5	A4	62	THR
5	A4	71	ARG
5	A4	79	ILE
5	A4	90	ARG
5	A4	97	VAL
5	A4	102	ARG
5	A4	105	THR

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Mol	Chain	Res	Type
5	A4	110	ASP
5	A4	112	TYR
5	A4	114	LEU
5	A4	116	GLN
5	A4	121	ARG
5	A4	130	ILE
5	A4	135	ILE
5	A4	138	CYS
5	A4	143	ARG
5	A4	162	ARG
5	A4	165	LYS
5	A4	178	TYR
5	A4	183	HIS
5	A4	191	MET
5	A4	192	TRP
6	A5	19	LYS
6	A5	31	ARG
6	A5	33	PRO
6	A5	35	ASN
6	A5	38	LEU
6	A5	42	ARG
6	A5	49	ARG
6	A5	56	ARG
6	A5	61	ASP
6	A5	69	THR
6	A5	70	GLU
6	A5	75	ARG
6	A5	77	ARG
6	A5	88	ASN
6	A5	127	LYS
6	A5	172	GLU
6	A5	202	LEU
6	A5	207	GLU
7	A6	2	ARG
7	A6	4	TYR
7	A6	14	PRO
7	A6	19	GLU
7	A6	22	ARG
7	A6	27	MET
7	A6	30	CYS
7	A6	36	ARG
7	A6	48	LEU

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Mol	Chain	Res	Type
7	A6	52	ARG
7	A6	60	THR
7	A6	86	ASP
7	A6	99	THR
7	A6	101	PRO
7	A6	104	LEU
7	A6	105	GLU
7	A6	122	HIS
7	A6	125	ARG
7	A6	139	ILE
7	A6	144	SER
7	A6	147	VAL
7	A6	148	ARG
7	A6	149	VAL
7	A6	173	ARG
7	A6	175	LYS
8	A7	10	THR
8	A7	13	ARG
8	A7	19	LEU
8	A7	34	THR
8	A7	56	TYR
8	A7	69	PHE
8	A7	70	VAL
8	A7	75	LEU
8	A7	80	ASN
8	A7	88	ASP
8	A7	91	LEU
8	A7	100	GLN
8	A7	104	LYS
8	A7	122	ARG
8	A7	130	ASP
8	A7	135	VAL
8	A7	158	VAL
8	A7	159	ARG
8	A7	160	PHE
8	A7	174	TRP
8	A7	198	TYR
8	A7	207	ASP
8	A7	211	CYS
8	A7	239	PRO
8	A7	252	MET
8	A7	253	CYS

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Mol	Chain	Res	Type
8	A7	276	PRO
8	A7	277	GLU
8	A7	278	HIS
8	A7	279	GLN
8	A7	300	LEU
8	A7	307	ASN
9	A8	23	VAL
9	A8	27	ASN
9	A8	28	GLN
9	A8	33	ARG
10	A9	91	HIS
10	A9	97	LYS
10	A9	120	ARG
10	A9	132	VAL
10	A9	152	MET
11	AC	46	MET
11	AC	49	SER
11	AC	57	MET
11	AC	63	THR
11	AC	65	ASN
11	AC	67	SER
11	AC	74	ILE
11	AC	77	ARG
11	AC	84	ILE
11	AC	91	TRP
11	AC	95	ILE
11	AC	96	LEU
11	AC	108	GLN
11	AC	116	ARG
11	AC	124	PHE
11	AC	126	PHE
11	AC	129	LEU
11	AC	144	PHE
11	AC	145	THR
11	AC	147	GLN
11	AC	160	VAL
11	AC	171	ARG
11	AC	177	ASN
11	AC	178	ILE
11	AC	194	ASP
11	AC	205	HIS
11	AC	207	ILE

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Mol	Chain	Res	Type
11	AC	211	TYR
11	AC	217	GLU
11	AC	221	LEU
11	AC	227	ARG
11	AC	237	LEU
11	AC	238	PHE
11	AC	239	PHE
11	AC	241	ARG
11	AC	242	ASP
12	AD	11	ASP
12	AD	17	PHE
12	AD	33	TRP
12	AD	36	THR
12	AD	41	THR
12	AD	47	LEU
12	AD	65	GLN
12	AD	80	ILE
12	AD	89	LEU
12	AD	95	PRO
12	AD	102	HIS
13	AE	18	VAL
13	AE	27	GLN
13	AE	39	LYS
13	AE	58	PHE
13	AE	65	ILE
13	AE	70	ILE
13	AE	80	VAL
13	AE	83	ARG
13	AE	85	ARG
13	AE	96	MET
13	AE	117	GLN
13	AE	135	PRO
13	AE	143	GLN
13	AE	153	TYR
13	AE	154	ASN
13	AE	172	SER
13	AE	173	LYS
14	AF	39	LYS
14	AF	72	GLU
14	AF	123	CYS
15	AG	4	MET
15	AG	23	PRO

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Mol	Chain	Res	Type
15	AG	46	SER
15	AG	52	MET
15	AG	53	GLN
15	AG	55	ARG
15	AG	64	LYS
15	AG	65	ASN
15	AG	71	ILE
15	AG	75	LEU
15	AG	97	GLN
15	AG	115	LEU
15	AG	131	ARG
15	AG	141	TYR
16	AH	23	VAL
16	AH	32	ASP
16	AH	43	ARG
16	AH	56	LYS
16	AH	59	ARG
16	AH	69	MET
16	AH	73	ASP
16	AH	75	VAL
16	AH	81	CYS
16	AH	123	GLU
16	AH	126	THR
16	AH	130	THR
17	AI	55	HIS
17	AI	97	VAL
17	AI	104	TYR
17	AI	107	ARG
17	AI	119	ILE
17	AI	130	TYR
17	AI	131	LYS
17	AI	137	ARG
18	AJ	20	ARG
18	AJ	23	ARG
18	AJ	39	GLN
18	AJ	41	MET
18	AJ	49	GLU
18	AJ	51	GLU
18	AJ	52	ILE
18	AJ	54	ASP
18	AJ	57	ARG
18	AJ	60	LYS

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Mol	Chain	Res	Type
18	AJ	81	CYS
18	AJ	85	ASP
18	AJ	98	GLN
18	AJ	104	LEU
18	AJ	106	THR
18	AJ	108	LEU
18	AJ	111	MET
18	AJ	112	ASP
19	AK	5	LYS
19	AK	11	GLN
19	AK	12	VAL
19	AK	28	THR
19	AK	40	VAL
19	AK	49	THR
19	AK	61	VAL
19	AK	72	ASP
19	AK	74	ARG
19	AK	78	SER
19	AK	80	GLN
19	AK	81	VAL
19	AK	89	GLN
19	AK	115	TYR
19	AK	121	PHE
19	AK	124	ILE
19	AK	129	ARG
19	AK	141	ARG
20	AL	5	ARG
20	AL	17	VAL
20	AL	21	PHE
20	AL	29	TYR
20	AL	55	THR
20	AL	71	LEU
20	AL	73	LEU
20	AL	75	GLU
20	AL	92	ASP
20	AL	100	ARG
21	AM	28	PRO
21	AM	40	ARG
21	AM	60	LEU
21	AM	87	ARG
21	AM	107	ARG
21	AM	121	HIS

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Mol	Chain	Res	Type
21	AM	122	ARG
21	AM	126	HIS
21	AM	132	VAL
21	AM	133	ARG
21	AM	139	THR
21	AM	140	SER
22	AO	45	MET
22	AO	46	MET
22	AO	52	GLU
22	AO	62	ARG
22	AO	73	ARG
22	AO	74	CYS
22	AO	87	VAL
22	AO	95	ARG
22	AO	96	PHE
22	AO	106	PRO
22	AO	108	ILE
22	AO	140	VAL
22	AO	143	LEU
22	AO	160	MET
22	AO	163	GLU
22	AO	167	LYS
23	AP	43	TRP
23	AP	44	VAL
23	AP	51	ARG
23	AP	62	GLU
23	AP	69	MET
23	AP	74	HIS
23	AP	75	GLN
23	AP	77	VAL
23	AP	78	ASP
23	AP	90	MET
23	AP	97	GLN
23	AP	105	ARG
23	AP	112	ASN
23	AP	133	LEU
23	AP	139	MET
23	AP	160	GLU
23	AP	168	VAL
23	AP	178	ARG
23	AP	183	PRO
23	AP	193	VAL

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Mol	Chain	Res	Type
23	AP	206	VAL
23	AP	210	SER
23	AP	217	ARG
23	AP	232	TYR
23	AP	236	THR
23	AP	238	ASP
23	AP	239	LEU
23	AP	245	VAL
23	AP	247	ARG
23	AP	256	PHE
23	AP	259	MET
24	AQ	16	ARG
24	AQ	21	MET
24	AQ	24	ARG
24	AQ	37	LEU
24	AQ	38	LEU
24	AQ	47	LEU
24	AQ	48	ARG
24	AQ	52	ARG
24	AQ	76	THR
24	AQ	114	PRO
25	AR	15	ASP
25	AR	29	ILE
25	AR	32	PHE
25	AR	39	ILE
25	AR	46	PRO
25	AR	49	VAL
25	AR	51	ASP
25	AR	59	ILE
25	AR	73	ILE
25	AR	78	ILE
25	AR	85	ILE
26	AS	4	THR
26	AS	7	GLN
26	AS	20	ASN
26	AS	30	SER
26	AS	39	ASN
26	AS	55	ILE
26	AS	66	ILE
26	AS	88	ASP
26	AS	101	LEU
26	AS	120	PHE

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Mol	Chain	Res	Type
26	AS	123	VAL
26	AS	134	TYR
27	AT	7	LYS
27	AT	9	GLU
27	AT	11	THR
27	AT	13	ARG
27	AT	15	SER
27	AT	25	ARG
27	AT	26	LYS
27	AT	31	GLU
27	AT	34	HIS
27	AT	35	PRO
27	AT	37	TRP
27	AT	38	CYS
27	AT	40	THR
27	AT	42	PRO
27	AT	48	ASN
27	AT	50	LEU
27	AT	59	GLU
27	AT	65	PHE
27	AT	67	PHE
27	AT	76	THR
27	AT	89	MET
27	AT	103	MET
27	AT	111	ARG
27	AT	118	ARG
27	AT	129	LYS
27	AT	132	MET
27	AT	133	GLN
27	AT	134	THR
28	AU	34	SER
28	AU	72	SER
28	AU	95	ILE
28	AU	105	ARG
28	AU	106	VAL
28	AU	108	THR
29	AV	5	ARG
29	AV	12	LYS
29	AV	17	ARG
29	AV	19	ARG
29	AV	28	CYS
29	AV	44	ARG

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Mol	Chain	Res	Type
29	AV	57	GLU
29	AV	61	ILE
29	AV	66	PHE
29	AV	68	MET
29	AV	74	LYS
29	AV	78	CYS
29	AV	94	GLU
30	AW	3	PHE
30	AW	4	PHE
30	AW	11	PRO
30	AW	14	ARG
30	AW	20	HIS
30	AW	23	ARG
30	AW	25	LEU
30	AW	30	ASN
30	AW	35	ASP
30	AW	48	TYR
30	AW	57	CYS
30	AW	66	ARG
30	AW	83	LYS
31	AX	10	MET
31	AX	22	GLU
31	AX	36	VAL
31	AX	37	GLU
31	AX	39	ARG
31	AX	43	THR
31	AX	51	SER
31	AX	54	THR
31	AX	66	ARG
31	AX	72	LEU
31	AX	75	ARG
31	AX	108	LEU
31	AX	136	THR
31	AX	144	GLN
31	AX	147	LYS
31	AX	152	ARG
31	AX	156	MET
31	AX	201	PRO
32	AY	4	VAL
32	AY	18	THR
32	AY	25	GLU
32	AY	26	LYS

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Mol	Chain	Res	Type
32	AY	31	ARG
32	AY	36	LYS
32	AY	38	LEU
32	AY	41	THR
32	AY	49	LEU
32	AY	59	LYS
32	AY	60	GLN
33	AZ	55	ASN
33	AZ	57	THR
33	AZ	64	MET
33	AZ	86	GLU
33	AZ	97	ARG
42	BI	13	ARG
42	BI	32	LEU
42	BI	48	ILE
42	BI	57	ASN
42	BI	69	VAL
42	BI	70	CYS
42	BI	72	ARG
42	BI	75	THR
42	BI	77	TRP
42	BI	80	LYS
42	BI	95	ASP
42	BI	96	ASP
42	BI	101	ARG
42	BI	108	CYS
42	BI	112	PHE
42	BI	115	SER
42	BI	119	ARG
42	BI	143	THR
42	BI	144	MET
42	BI	147	ARG
42	BI	151	SER
42	BI	171	PRO
42	BI	179	GLU
42	BI	193	LYS
43	BJ	30	ASP
43	BJ	38	TYR
43	BJ	41	GLN
43	BJ	60	MET
43	BJ	63	CYS
43	BJ	78	ASP

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Mol	Chain	Res	Type
43	BJ	93	ASN
43	BJ	94	LYS
43	BJ	95	LYS
43	BJ	159	VAL
43	BJ	177	ARG
43	BJ	188	VAL
44	BK	8	CYS
44	BK	14	ASN
44	BK	17	TYR
44	BK	24	ARG
44	BK	30	ARG
44	BK	49	CYS
44	BK	53	VAL
44	BK	54	SER
44	BK	59	GLN
44	BK	69	ARG
44	BK	86	HIS
44	BK	87	MET
44	BK	90	ARG
44	BK	99	ILE
44	BK	101	LYS
44	BK	102	MET
44	BK	103	LEU
44	BK	115	MET
44	BK	121	LYS
44	BK	128	ARG
44	BK	142	ASP
44	BK	143	THR
44	BK	151	SER
44	BK	153	ARG
44	BK	169	LYS
44	BK	177	LEU
44	BK	185	ARG
44	BK	191	GLN
44	BK	196	HIS
44	BK	200	ILE
45	BL	17	ARG
45	BL	21	VAL
45	BL	24	LEU
45	BL	25	CYS
45	BL	51	THR
45	BL	53	VAL

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Mol	Chain	Res	Type
45	BL	58	ARG
45	BL	67	ARG
45	BL	74	VAL
45	BL	77	THR
45	BL	88	GLU
45	BL	113	ILE
45	BL	117	ILE
45	BL	132	MET
45	BL	133	ASP
45	BL	140	ARG
45	BL	141	ARG
45	BL	148	ARG
45	BL	151	ARG
45	BL	153	SER
45	BL	157	PHE
45	BL	159	HIS
45	BL	162	ARG
45	BL	167	MET
45	BL	175	ASP
45	BL	180	GLN
46	BM	9	GLN
46	BM	12	THR
46	BM	45	ASP
46	BM	67	ARG
46	BM	80	ARG
46	BM	87	GLU
46	BM	99	LYS
46	BM	100	HIS
47	BN	6	ASN
47	BN	10	HIS
47	BN	11	VAL
47	BN	14	ARG
47	BN	16	HIS
47	BN	23	GLN
47	BN	28	LYS
47	BN	32	ASN
47	BN	52	PHE
47	BN	62	GLN
47	BN	64	ASN
47	BN	65	CYS
47	BN	66	PRO
47	BN	71	ASN

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Mol	Chain	Res	Type
47	BN	72	MET
47	BN	74	ARG
47	BN	78	ARG
47	BN	80	PHE
47	BN	84	GLU
47	BN	91	LYS
47	BN	93	ARG
47	BN	96	ARG
47	BN	103	ASP
47	BN	104	ARG
47	BN	105	ARG
47	BN	111	GLU
47	BN	112	GLU
47	BN	114	MET
47	BN	119	GLN
47	BN	126	SER
47	BN	135	ARG
47	BN	149	LYS
47	BN	158	TYR
47	BN	167	VAL
47	BN	171	ARG
47	BN	177	VAL
47	BN	189	PHE
47	BN	192	LYS
47	BN	193	ASN
47	BN	194	HIS
47	BN	204	ASN
47	BN	210	LYS
47	BN	214	GLU
48	BO	23	ASP
48	BO	25	ILE
48	BO	34	LEU
48	BO	56	CYS
48	BO	67	ARG
48	BO	75	PHE
48	BO	99	ILE
48	BO	103	ARG
48	BO	112	ARG
48	BO	132	ARG
48	BO	137	VAL
48	BO	140	PRO
48	BO	144	ARG

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Mol	Chain	Res	Type
48	BO	155	THR
48	BO	162	LYS
48	BO	176	GLU
48	BO	179	ARG
48	BO	189	LYS
48	BO	193	VAL
48	BO	197	TRP
48	BO	201	ARG
48	BO	202	LYS
48	BO	206	LYS
48	BO	207	LYS
48	BO	208	MET
49	BP	8	ARG
49	BP	13	VAL
49	BP	17	ARG
49	BP	19	PRO
49	BP	20	ARG
49	BP	38	VAL
49	BP	59	LEU
49	BP	62	CYS
49	BP	66	SER
49	BP	69	CYS
49	BP	74	LEU
49	BP	112	GLN
49	BP	120	ARG
49	BP	128	PHE
49	BP	133	LYS
49	BP	140	HIS
49	BP	183	LYS
50	BQ	20	SER
50	BQ	24	LEU
50	BQ	35	VAL
50	BQ	40	GLN
50	BQ	48	ARG
50	BQ	63	GLU
50	BQ	64	LYS
50	BQ	66	ARG
50	BQ	67	MET
50	BQ	68	VAL
50	BQ	72	THR
50	BQ	88	ARG
50	BQ	92	VAL

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Mol	Chain	Res	Type
50	BQ	94	LYS
50	BQ	102	ASN
50	BQ	106	VAL
50	BQ	112	ASN
50	BQ	137	TRP
50	BQ	145	LEU
50	BQ	153	ASP
50	BQ	172	VAL
50	BQ	178	GLN
50	BQ	188	HIS
50	BQ	189	ARG
50	BQ	192	ARG
50	BQ	211	ARG
50	BQ	219	ARG
50	BQ	220	LYS
50	BQ	221	ARG
51	BR	8	PRO
51	BR	9	GLN
51	BR	12	SER
51	BR	14	THR
51	BR	28	ASN
51	BR	41	LEU
51	BR	49	ARG
51	BR	55	THR
51	BR	56	ARG
51	BR	59	PRO
51	BR	61	LYS
51	BR	62	ARG
51	BR	66	LYS
51	BR	69	ASN
51	BR	75	GLU
51	BR	78	GLN
51	BR	79	THR
51	BR	103	ILE
51	BR	112	MET
51	BR	125	VAL
51	BR	127	ARG
51	BR	139	TYR
51	BR	141	ARG
51	BR	144	CYS
51	BR	147	GLN
51	BR	149	PHE

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Mol	Chain	Res	Type
52	BS	2	VAL
52	BS	3	ARG
52	BS	6	LEU
52	BS	11	VAL
52	BS	14	ARG
52	BS	16	THR
52	BS	42	ARG
52	BS	46	MET
52	BS	48	ARG
52	BS	53	VAL
52	BS	57	HIS
52	BS	64	LYS
52	BS	66	VAL
52	BS	68	ASP
52	BS	70	LYS
52	BS	71	LEU
52	BS	82	TYR
52	BS	86	ARG
52	BS	89	TYR
52	BS	90	THR
52	BS	92	MET
52	BS	110	ASN
52	BS	117	ARG
52	BS	121	HIS
52	BS	137	ARG
52	BS	145	HIS
52	BS	151	PHE
52	BS	158	VAL
52	BS	162	ARG
52	BS	175	ARG
53	BT	4	LEU
53	BT	6	LEU
53	BT	7	GLN
53	BT	16	ARG
53	BT	21	ARG
53	BT	26	PRO
53	BT	37	SER
53	BT	39	LYS
53	BT	42	ARG
53	BT	43	LYS
53	BT	63	TRP
53	BT	74	ARG

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Mol	Chain	Res	Type
53	BT	78	THR
53	BT	81	ARG
53	BT	89	MET
53	BT	90	PRO
53	BT	95	TRP
53	BT	99	LEU
53	BT	110	ARG
53	BT	113	LYS
53	BT	116	ASP
53	BT	137	ASN
53	BT	144	LYS
53	BT	154	GLN
53	BT	175	LYS
53	BT	185	ARG
54	BU	7	TYR
54	BU	12	ARG
54	BU	13	HIS
54	BU	19	PHE
54	BU	30	ILE
54	BU	61	THR
54	BU	65	TRP
54	BU	69	PRO
54	BU	75	ILE
54	BU	86	ARG
54	BU	87	LYS
54	BU	88	ARG
54	BU	97	ARG
54	BU	113	GLU
54	BU	115	PHE
54	BU	127	PRO
54	BU	149	ARG
54	BU	153	TYR
55	BV	23	LYS
55	BV	27	LYS
55	BV	31	SER
55	BV	48	GLN
55	BV	54	ASN
55	BV	55	VAL
55	BV	86	ARG
55	BV	87	LYS
55	BV	93	LEU
55	BV	110	LEU

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Mol	Chain	Res	Type
56	BW	4	ASP
56	BW	13	PHE
56	BW	14	ARG
56	BW	16	SER
56	BW	37	TYR
56	BW	47	ARG
56	BW	50	ARG
56	BW	56	LEU
56	BW	59	MET
56	BW	73	ARG
56	BW	75	VAL
56	BW	77	ASN
56	BW	81	ILE
56	BW	84	ARG
56	BW	89	ARG
56	BW	97	PHE
56	BW	105	VAL
56	BW	106	ASN
56	BW	129	PRO
56	BW	131	ILE
56	BW	136	PRO
57	BX	62	ARG
57	BX	63	PRO
57	BX	71	ASN
57	BX	82	PHE
57	BX	95	MET
57	BX	109	ASP
57	BX	134	THR
57	BX	135	LEU
57	BX	143	LYS
57	BX	156	ASP
57	BX	160	LYS
58	BY	17	HIS
58	BY	22	VAL
58	BY	45	MET
58	BY	47	LYS
58	BY	60	ARG
58	BY	64	ARG
58	BY	67	THR
59	BZ	13	ARG
59	BZ	22	HIS
59	BZ	33	SER

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Mol	Chain	Res	Type
59	BZ	39	LYS
59	BZ	53	VAL
59	BZ	57	ARG
59	BZ	63	ARG
59	BZ	84	CYS
59	BZ	92	VAL
59	BZ	97	HIS
59	BZ	99	SER
59	BZ	105	LYS
60	Ba	3	PHE
60	Ba	8	LYS
60	Ba	30	THR
60	Ba	31	ARG
60	Ba	52	VAL
60	Ba	53	VAL
60	Ba	62	THR
60	Ba	63	ARG
60	Ba	80	LEU
60	Ba	82	THR
60	Ba	85	ASN
60	Ba	88	LEU
60	Ba	89	SER
60	Ba	96	ILE
60	Ba	100	ASP
60	Ba	132	ARG
61	Bb	4	ARG
61	Bb	7	LYS
61	Bb	11	GLN
61	Bb	16	PHE
61	Bb	22	VAL
61	Bb	25	HIS
61	Bb	40	HIS
61	Bb	48	TYR
61	Bb	49	HIS
61	Bb	58	MET
61	Bb	76	ASP
61	Bb	77	ASN
61	Bb	100	ASP
61	Bb	102	LEU
61	Bb	112	ASN
61	Bb	114	HIS
61	Bb	116	GLN

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Mol	Chain	Res	Type
61	Bb	124	ARG
61	Bb	128	LYS
61	Bb	135	ARG
62	Bc	10	LEU
62	Bc	12	VAL
62	Bc	13	ARG
62	Bc	21	LYS
62	Bc	28	SER
62	Bc	30	ASP
62	Bc	33	ASN
62	Bc	40	LYS
62	Bc	45	PHE
62	Bc	52	VAL
62	Bc	55	THR
62	Bc	58	ASP
62	Bc	69	THR
62	Bc	76	MET
62	Bc	77	TYR
62	Bc	79	LYS
62	Bc	83	GLU
62	Bc	84	PRO
62	Bc	89	SER
62	Bc	93	ARG
62	Bc	99	ARG
62	Bc	101	ASP
62	Bc	104	LYS
62	Bc	109	ARG
62	Bc	122	LYS
62	Bc	124	CYS
62	Bc	132	SER
62	Bc	135	HIS
62	Bc	138	ARG
63	Bd	10	HIS
63	Bd	24	PRO
63	Bd	26	PRO
63	Bd	28	TYR
64	Be	69	TYR
64	Be	71	ARG
64	Be	80	GLU
64	Be	83	PHE
64	Be	84	THR
64	Be	96	LEU

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Mol	Chain	Res	Type
64	Be	98	ILE
64	Be	100	ASN
64	Be	123	ARG
64	Be	132	ASP
64	Be	135	ILE
64	Be	140	ASN
64	Be	145	ARG
64	Be	146	THR
64	Be	148	LEU
64	Be	186	PHE
64	Be	192	LYS
64	Be	205	ASN
64	Be	208	GLU
64	Be	216	HIS
64	Be	234	LYS
64	Be	241	ARG
64	Be	242	ARG
65	Bf	56	LYS
65	Bf	58	GLU
65	Bf	73	SER
65	Bf	90	GLN
65	Bf	91	LYS
65	Bf	106	HIS
65	Bf	107	ILE
65	Bf	112	ASP
65	Bf	123	VAL
65	Bf	125	GLU
65	Bf	144	ARG
65	Bf	146	THR
65	Bf	147	PRO
65	Bf	152	THR
65	Bf	153	ILE
65	Bf	159	HIS
65	Bf	162	SER
65	Bf	164	GLU
65	Bf	165	PHE
65	Bf	168	ARG
65	Bf	169	PHE
65	Bf	184	ARG
65	Bf	223	ARG
65	Bf	238	GLN
65	Bf	240	ASN

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Mol	Chain	Res	Type
65	Bf	250	LEU
65	Bf	257	LYS
65	Bf	258	GLU
65	Bf	267	GLN
65	Bf	277	THR
65	Bf	288	ARG
65	Bf	296	ARG
65	Bf	298	THR
65	Bf	302	LEU
65	Bf	315	ARG
65	Bf	351	GLN
65	Bf	354	THR
65	Bf	363	ILE
65	Bf	364	THR
65	Bf	383	LYS
65	Bf	386	VAL
65	Bf	392	ARG
65	Bf	403	GLN
65	Bf	432	LYS
66	Bg	22	ASN
66	Bg	29	MET
66	Bg	36	LEU
66	Bg	45	LEU
66	Bg	52	LEU
66	Bg	58	ASN
66	Bg	62	ILE
66	Bg	77	PRO
66	Bg	81	TYR
66	Bg	96	PHE
66	Bg	97	ARG
66	Bg	102	SER
67	Bh	1	MET
67	Bh	15	LYS
67	Bh	25	LYS
67	Bh	35	ASP
67	Bh	41	VAL
67	Bh	53	TYR
67	Bh	59	THR
67	Bh	64	ARG
67	Bh	79	ARG
67	Bh	81	PRO
67	Bh	86	MET

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Mol	Chain	Res	Type
67	Bh	90	ILE
67	Bh	95	LEU
67	Bh	109	ILE
67	Bh	122	THR
67	Bh	126	ARG
67	Bh	131	LEU
67	Bh	133	THR
67	Bh	148	ARG
67	Bh	159	GLU
67	Bh	163	ARG
67	Bh	171	SER
67	Bh	174	PRO
67	Bh	176	PRO
67	Bh	180	ASN
67	Bh	184	LYS
68	Bi	4	PRO
68	Bi	5	PHE
68	Bi	9	ASN
68	Bi	13	LYS
68	Bi	15	THR
68	Bi	24	GLU
68	Bi	34	ARG
68	Bi	61	ASP
68	Bi	63	ARG
68	Bi	69	PRO
68	Bi	77	VAL
68	Bi	85	MET
68	Bi	88	MET
68	Bi	100	THR
68	Bi	101	VAL
68	Bi	106	ARG
68	Bi	122	ASN
69	Bj	2	SER
69	Bj	3	CYS
69	Bj	10	ARG
69	Bj	17	ARG
69	Bj	20	ARG
69	Bj	27	PRO
69	Bj	30	ARG
69	Bj	36	ARG
69	Bj	40	SER
69	Bj	43	PRO

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Mol	Chain	Res	Type
69	Bj	44	HIS
69	Bj	53	ARG
69	Bj	54	LEU
69	Bj	58	LYS
69	Bj	63	THR
69	Bj	66	ARG
69	Bj	69	PRO
69	Bj	71	HIS
69	Bj	74	THR
69	Bj	77	ARG
69	Bj	85	HIS
69	Bj	87	GLN
69	Bj	90	ASP
69	Bj	114	VAL
69	Bj	125	ARG
70	Bk	47	ARG
70	Bk	49	ARG
70	Bk	59	MET
70	Bk	60	THR
70	Bk	62	LEU
70	Bk	68	GLU
70	Bk	73	PHE
70	Bk	74	TYR
70	Bk	77	LYS
70	Bk	83	LYS
70	Bk	92	THR
70	Bk	93	HIS
70	Bk	103	GLU
70	Bk	111	GLN
70	Bk	116	HIS
71	Bl	37	ARG
71	Bl	69	THR
71	Bl	89	ILE
71	Bl	96	SER
71	Bl	99	PRO
71	Bl	103	SER
71	Bl	112	ILE
71	Bl	114	ARG
71	Bl	115	PRO
71	Bl	118	THR
71	Bl	119	SER
71	Bl	121	THR

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Mol	Chain	Res	Type
71	Bl	124	VAL
71	Bl	132	PRO
71	Bl	134	SER
71	Bl	147	SER
72	Bm	12	ARG
72	Bm	20	ASN
72	Bm	26	THR
72	Bm	36	ASP
72	Bm	37	ARG
72	Bm	51	ILE
72	Bm	61	MET
72	Bm	78	LEU
72	Bm	85	LEU
72	Bm	87	SER
72	Bm	99	GLU
72	Bm	100	GLU
72	Bm	105	GLN
72	Bm	106	THR
72	Bm	107	LYS
73	Bn	2	THR
73	Bn	12	HIS
73	Bn	17	ILE
73	Bn	19	CYS
73	Bn	33	ARG
73	Bn	43	GLN
73	Bn	45	ARG
73	Bn	46	ARG
73	Bn	47	TYR
73	Bn	49	TRP
73	Bn	58	ARG
73	Bn	66	TYR
73	Bn	77	HIS
73	Bn	80	THR
73	Bn	81	ASP
74	Bo	8	MET
74	Bo	11	MET
74	Bo	30	GLU
74	Bo	38	PHE
74	Bo	41	PHE
74	Bo	42	CYS
74	Bo	47	PHE
74	Bo	49	ARG

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Mol	Chain	Res	Type
74	Bo	56	ARG
74	Bo	72	SER
74	Bo	73	THR
74	Bo	74	PRO
74	Bo	87	ARG
74	Bo	90	LYS
74	Bo	91	GLN
75	Bp	8	LEU
75	Bp	9	LYS
75	Bp	10	GLU
75	Bp	38	CYS
75	Bp	47	MET
75	Bp	51	LYS
75	Bp	58	ARG
75	Bp	73	SER
76	Bq	3	ARG
76	Bq	5	LYS
76	Bq	6	PRO
76	Bq	11	LYS
76	Bq	16	LYS
76	Bq	18	LYS
76	Bq	22	PRO
76	Bq	25	TYR
76	Bq	33	ASN
76	Bq	42	ARG
76	Bq	46	ARG
77	Br	8	SER
77	Br	16	LYS
77	Br	22	SER
77	Br	23	LEU
77	Br	26	VAL
77	Br	31	ILE
77	Br	32	ARG
77	Br	34	ASP
77	Br	35	VAL
77	Br	38	PHE
77	Br	40	HIS
77	Br	51	TYR
77	Br	55	ARG
77	Br	60	LYS
77	Br	76	PRO
77	Br	93	ASN

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Mol	Chain	Res	Type
77	Br	94	MET
77	Br	96	ARG
77	Br	100	MET
77	Br	103	PRO
77	Br	107	PHE
77	Br	110	TRP
77	Br	116	LEU
77	Br	124	VAL
77	Br	135	LEU
77	Br	137	MET
77	Br	144	GLU
77	Br	151	LEU
77	Br	155	ASP
77	Br	158	ARG
77	Br	161	GLU
77	Br	165	GLU
77	Br	169	PHE
77	Br	173	VAL
77	Br	178	ASP
77	Br	179	VAL
77	Br	188	ILE
77	Br	194	LYS
77	Br	195	MET
77	Br	203	ARG
77	Br	214	LYS
77	Br	222	ILE
77	Br	227	LEU
77	Br	246	ARG
77	Br	271	VAL
77	Br	273	SER
77	Br	289	ARG
77	Br	320	ILE
77	Br	323	ARG
77	Br	334	LYS
77	Br	339	MET
77	Br	343	MET
77	Br	365	LYS
78	Bs	2	MET
78	Bs	11	LYS
78	Bs	12	LYS
78	Bs	24	TYR
78	Bs	29	VAL

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Mol	Chain	Res	Type
78	Bs	46	ARG
78	Bs	47	MET
78	Bs	51	LEU
79	Bt	4	TYR
79	Bt	10	MET
79	Bt	15	GLU
79	Bt	20	HIS
79	Bt	32	LYS
79	Bt	34	ARG
79	Bt	56	PRO
79	Bt	63	LYS
79	Bt	73	GLN
79	Bt	83	ASN
79	Bt	84	VAL
80	Bu	6	VAL
80	Bu	7	VAL
80	Bu	10	LYS
80	Bu	13	TYR
80	Bu	15	ARG
80	Bu	30	TYR
80	Bu	40	ASP
80	Bu	46	THR
80	Bu	56	THR
80	Bu	64	ILE
80	Bu	65	VAL
80	Bu	66	HIS
80	Bu	68	LYS
80	Bu	79	TYR
80	Bu	84	PRO
80	Bu	90	HIS
80	Bu	99	TYR
80	Bu	117	GLU
80	Bu	142	ASP
80	Bu	143	GLU
80	Bu	145	ARG
80	Bu	164	ARG
80	Bu	169	LEU
80	Bu	170	LYS
80	Bu	187	PRO
80	Bu	192	GLU
80	Bu	220	GLU
80	Bu	231	GLN

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Mol	Chain	Res	Type
80	Bu	243	ASP
80	Bu	245	MET
80	Bu	248	MET
80	Bu	257	ARG
80	Bu	264	ARG
80	Bu	270	PRO
80	Bu	272	GLU
80	Bu	276	LYS
80	Bu	278	TYR
80	Bu	282	LYS
80	Bu	283	LEU
81	Bv	6	LYS
81	Bv	10	GLU
81	Bv	12	LYS
81	Bv	14	LYS
81	Bv	15	LYS
81	Bv	16	VAL
81	Bv	24	SER
81	Bv	27	ARG
81	Bv	30	CYS
81	Bv	31	LYS
81	Bv	53	GLN
81	Bv	66	MET
81	Bv	86	THR
81	Bv	101	GLU
81	Bv	167	LEU
81	Bv	169	LYS
81	Bv	174	TYR
81	Bv	185	ASP
81	Bv	188	HIS
81	Bv	190	LEU
82	Bw	1	MET
82	Bw	3	MET
82	Bw	6	LEU
82	Bw	7	ILE
82	Bw	11	LYS
82	Bw	28	ILE
82	Bw	34	LYS
82	Bw	41	PHE
82	Bw	60	TYR
82	Bw	73	THR
82	Bw	82	ARG

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Mol	Chain	Res	Type
82	Bw	103	ARG
82	Bw	105	ARG
82	Bw	107	ILE
82	Bw	114	GLN
82	Bw	132	VAL
82	Bw	133	ARG
82	Bw	135	ASN
82	Bw	141	MET
82	Bw	143	ARG
82	Bw	156	LEU
82	Bw	179	THR
82	Bw	183	MET
82	Bw	201	ASN
82	Bw	202	GLN
82	Bw	204	TYR
82	Bw	209	HIS
82	Bw	211	ARG
82	Bw	221	LYS
82	Bw	224	PRO
82	Bw	225	PRO
82	Bw	229	MET
82	Bw	230	ARG
82	Bw	235	HIS
82	Bw	238	GLU
82	Bw	241	ASP
82	Bw	251	ARG
83	Bx	37	PRO
83	Bx	46	ASP
83	Bx	54	PRO
83	Bx	84	VAL
83	Bx	85	PRO
83	Bx	97	ARG
83	Bx	107	VAL
83	Bx	116	LYS
83	Bx	119	ARG
83	Bx	120	ASP
83	Bx	150	GLN
83	Bx	160	THR
83	Bx	175	LEU
83	Bx	181	THR
83	Bx	184	ARG
83	Bx	193	VAL

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Mol	Chain	Res	Type
83	Bx	195	ASP
83	Bx	206	LYS
83	Bx	207	THR
83	Bx	209	THR
83	Bx	215	ASP
83	Bx	224	LEU
83	Bx	240	ASP
83	Bx	243	ARG
83	Bx	253	LEU
83	Bx	256	ARG
83	Bx	259	LEU
83	Bx	260	ARG
83	Bx	261	LYS
83	Bx	270	ASP
84	By	1	MET
84	By	2	LYS
84	By	7	ASP
84	By	8	GLN
84	By	13	GLU
84	By	16	THR
84	By	39	HIS
84	By	44	PHE
84	By	45	ARG
84	By	52	THR
84	By	54	THR
84	By	58	TRP
84	By	64	ASN
84	By	82	VAL
84	By	92	PHE
84	By	98	PRO
84	By	105	ASN
84	By	106	GLN
84	By	113	PHE
84	By	119	VAL
84	By	120	ARG
84	By	121	ARG
84	By	131	TYR
84	By	132	ARG
84	By	133	THR
84	By	134	ASP
84	By	145	GLU
84	By	150	GLU

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Mol	Chain	Res	Type
84	By	154	ARG
84	By	159	MET
84	By	170	ILE
84	By	172	LYS
84	By	175	ASP
84	By	181	THR

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (164) such sidechains are listed below:

Mol	Chain	Res	Type
1	A0	150	ASN
2	A1	33	HIS
2	A1	59	GLN
3	A2	43	GLN
3	A2	81	HIS
4	A3	27	ASN
4	A3	66	GLN
4	A3	231	HIS
4	A3	232	HIS
5	A4	6	HIS
5	A4	20	GLN
5	A4	36	HIS
5	A4	59	HIS
5	A4	183	HIS
6	A5	74	GLN
6	A5	88	ASN
6	A5	169	HIS
7	A6	3	ASN
8	A7	23	GLN
8	A7	89	HIS
8	A7	108	HIS
8	A7	121	ASN
8	A7	197	ASN
8	A7	278	HIS
8	A7	307	ASN
9	A8	21	HIS
9	A8	28	GLN
10	A9	147	HIS
11	AC	83	ASN
13	AE	19	GLN
13	AE	41	HIS
13	AE	79	ASN

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Mol	Chain	Res	Type
13	AE	120	HIS
13	AE	122	ASN
14	AF	86	GLN
15	AG	77	HIS
16	AH	87	HIS
17	AI	86	HIS
18	AJ	9	ASN
18	AJ	44	HIS
18	AJ	56	HIS
19	AK	32	GLN
19	AK	34	ASN
20	AL	30	GLN
20	AL	74	GLN
20	AL	90	HIS
21	AM	98	HIS
22	AO	138	HIS
23	AP	97	GLN
23	AP	112	ASN
24	AQ	12	GLN
24	AQ	115	HIS
25	AR	75	HIS
27	AT	16	GLN
27	AT	48	ASN
31	AX	163	HIS
32	AY	5	HIS
32	AY	29	GLN
32	AY	60	GLN
33	AZ	55	ASN
33	AZ	73	ASN
42	BI	17	HIS
42	BI	18	HIS
42	BI	57	ASN
42	BI	142	ASN
42	BI	172	HIS
42	BI	188	HIS
43	BJ	54	HIS
43	BJ	168	HIS
44	BK	86	HIS
45	BL	75	HIS
45	BL	102	ASN
45	BL	147	HIS
47	BN	6	ASN

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Mol	Chain	Res	Type
47	BN	10	HIS
48	BO	74	GLN
48	BO	91	HIS
48	BO	119	GLN
49	BP	51	GLN
49	BP	96	HIS
49	BP	140	HIS
50	BQ	49	HIS
50	BQ	102	ASN
50	BQ	156	HIS
50	BQ	170	ASN
50	BQ	196	HIS
50	BQ	213	ASN
51	BR	3	HIS
51	BR	9	GLN
51	BR	46	GLN
51	BR	69	ASN
51	BR	121	GLN
52	BS	57	HIS
52	BS	121	HIS
52	BS	145	HIS
53	BT	58	HIS
53	BT	75	HIS
53	BT	143	HIS
54	BU	3	HIS
54	BU	22	HIS
54	BU	58	HIS
57	BX	101	ASN
57	BX	133	ASN
58	BY	10	HIS
58	BY	17	HIS
59	BZ	97	HIS
61	Bb	25	HIS
61	Bb	60	HIS
62	Bc	135	HIS
62	Bc	137	HIS
63	Bd	6	ASN
63	Bd	16	ASN
63	Bd	43	ASN
64	Be	115	ASN
64	Be	205	ASN
64	Be	216	HIS

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Mol	Chain	Res	Type
64	Be	233	GLN
65	Bf	90	GLN
65	Bf	159	HIS
65	Bf	160	HIS
65	Bf	267	GLN
65	Bf	326	HIS
65	Bf	335	ASN
66	Bg	39	GLN
66	Bg	79	HIS
66	Bg	95	HIS
67	Bh	54	HIS
67	Bh	132	ASN
68	Bi	99	HIS
69	Bj	44	HIS
69	Bj	62	HIS
69	Bj	101	GLN
70	Bk	69	ASN
71	Bl	54	GLN
71	Bl	67	ASN
71	Bl	85	HIS
72	Bm	42	HIS
74	Bo	34	HIS
74	Bo	91	GLN
75	Bp	74	HIS
76	Bq	43	HIS
77	Br	54	ASN
77	Br	111	HIS
77	Br	145	ASN
77	Br	187	GLN
77	Br	237	HIS
77	Br	333	GLN
77	Br	353	HIS
79	Bt	20	HIS
79	Bt	83	ASN
80	Bu	39	GLN
80	Bu	57	ASN
80	Bu	90	HIS
80	Bu	139	GLN
80	Bu	253	HIS
81	Bv	176	HIS
81	Bv	188	HIS
82	Bw	87	HIS

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Mol	Chain	Res	Type
82	Bw	189	HIS
82	Bw	190	ASN
82	Bw	209	HIS
83	Bx	72	GLN
84	By	8	GLN
84	By	39	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
34	BA	1846/1847 (99%)	616 (33%)	225 (12%)
35	BB	1464/1465 (99%)	460 (31%)	143 (9%)
36	BC	169/169 (100%)	46 (27%)	20 (11%)
37	BD	118/119 (99%)	31 (26%)	9 (7%)
38	BE	209/210 (99%)	78 (37%)	27 (12%)
39	BF	73/73 (100%)	48 (65%)	23 (31%)
40	BG	181/182 (99%)	40 (22%)	8 (4%)
41	BH	134/135 (99%)	49 (36%)	16 (11%)
85	AA	2226/2251 (98%)	767 (34%)	312 (14%)
86	AB	72/73 (98%)	30 (41%)	7 (9%)
All	All	6492/6524 (99%)	2165 (33%)	790 (12%)

All (2165) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
34	BA	11	U
34	BA	13	U
34	BA	22	C
34	BA	23	A
34	BA	37	A
34	BA	39	C
34	BA	46	C
34	BA	56	G
34	BA	57	A
34	BA	62	A
34	BA	63	A
34	BA	70	C
34	BA	71	G
34	BA	72	U
34	BA	73	G
34	BA	80	U

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Mol	Chain	Res	Type
34	BA	85	C
34	BA	90	G
34	BA	91	C
34	BA	97	A
34	BA	107	C
34	BA	108	A
34	BA	109	A
34	BA	112	C
34	BA	115	U
34	BA	116	G
34	BA	117	C
34	BA	119	G
34	BA	120	A
34	BA	121	A
34	BA	130	U
34	BA	131	A
34	BA	132	U
34	BA	134	U
34	BA	141	G
34	BA	144	C
34	BA	145	U
34	BA	147	U
34	BA	149	G
34	BA	151	A
34	BA	152	C
34	BA	153	C
34	BA	154	A
34	BA	156	U
34	BA	157	U
34	BA	159	U
34	BA	160	G
34	BA	161	U
34	BA	163	G
34	BA	165	C
34	BA	166	G
34	BA	175	G
34	BA	178	C
34	BA	179	U
34	BA	180	G
34	BA	185	A
34	BA	188	C
34	BA	189	G

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Mol	Chain	Res	Type
34	BA	194	G
34	BA	195	G
34	BA	197	A
34	BA	201	A
34	BA	202	A
34	BA	206	C
34	BA	207	A
34	BA	208	A
34	BA	209	A
34	BA	214	A
34	BA	215	C
34	BA	216	C
34	BA	217	C
34	BA	218	G
34	BA	219	U
34	BA	224	G
34	BA	225	A
34	BA	226	A
34	BA	227	C
34	BA	248	G
34	BA	249	A
34	BA	250	G
34	BA	256	A
34	BA	257	G
34	BA	261	A
34	BA	262	A
34	BA	265	A
34	BA	269	G
34	BA	270	U
34	BA	271	C
34	BA	280	A
34	BA	282	A
34	BA	285	C
34	BA	287	U
34	BA	288	U
34	BA	290	G
34	BA	295	G
34	BA	296	G
34	BA	297	A
34	BA	298	G
34	BA	299	C
34	BA	302	A

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Mol	Chain	Res	Type
34	BA	303	C
34	BA	304	G
34	BA	314	A
34	BA	315	U
34	BA	316	G
34	BA	317	U
34	BA	318	U
34	BA	324	C
34	BA	325	A
34	BA	326	A
34	BA	328	A
34	BA	331	G
34	BA	332	U
34	BA	344	G
34	BA	345	G
34	BA	347	A
34	BA	348	U
34	BA	357	A
34	BA	358	A
34	BA	359	G
34	BA	360	C
34	BA	361	C
34	BA	368	U
34	BA	370	U
34	BA	386	A
34	BA	392	A
34	BA	393	G
34	BA	394	A
34	BA	395	G
34	BA	396	U
34	BA	397	A
34	BA	403	A
34	BA	404	C
34	BA	405	C
34	BA	411	C
34	BA	412	G
34	BA	413	A
34	BA	415	C
34	BA	416	A
34	BA	417	A
34	BA	435	U
34	BA	437	G

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Mol	Chain	Res	Type
34	BA	438	A
34	BA	439	A
34	BA	442	G
34	BA	461	A
34	BA	463	A
34	BA	464	U
34	BA	465	A
34	BA	467	A
34	BA	469	C
34	BA	483	A
34	BA	484	A
34	BA	487	A
34	BA	488	C
34	BA	489	A
34	BA	490	A
34	BA	492	G
34	BA	501	U
34	BA	502	U
34	BA	509	U
34	BA	510	U
34	BA	511	U
34	BA	512	U
34	BA	519	G
34	BA	520	G
34	BA	521	C
34	BA	523	A
34	BA	524	G
34	BA	526	C
34	BA	527	C
34	BA	529	A
34	BA	531	C
34	BA	532	C
34	BA	533	U
34	BA	534	C
34	BA	536	C
34	BA	539	C
34	BA	547	C
34	BA	548	G
34	BA	551	U
34	BA	553	A
34	BA	555	C
34	BA	557	U

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Mol	Chain	Res	Type
34	BA	559	C
34	BA	560	U
34	BA	562	C
34	BA	569	C
34	BA	571	G
34	BA	573	U
34	BA	575	U
34	BA	576	C
34	BA	577	U
34	BA	578	C
34	BA	579	U
34	BA	581	U
34	BA	584	A
34	BA	586	G
34	BA	587	U
34	BA	588	C
34	BA	589	A
34	BA	591	G
34	BA	592	G
34	BA	593	G
34	BA	594	G
34	BA	595	U
34	BA	596	G
34	BA	597	C
34	BA	599	U
34	BA	600	G
34	BA	603	U
34	BA	604	G
34	BA	605	G
34	BA	607	C
34	BA	608	G
34	BA	612	U
34	BA	617	G
34	BA	623	U
34	BA	624	G
34	BA	626	G
34	BA	628	U
34	BA	629	G
34	BA	630	U
34	BA	634	U
34	BA	635	G
34	BA	638	U

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Mol	Chain	Res	Type
34	BA	639	U
34	BA	647	U
34	BA	651	U
34	BA	656	U
34	BA	657	C
34	BA	659	U
34	BA	660	C
34	BA	666	C
34	BA	668	G
34	BA	673	U
34	BA	674	G
34	BA	675	C
34	BA	678	C
34	BA	680	C
34	BA	681	G
34	BA	682	A
34	BA	683	C
34	BA	684	G
34	BA	685	C
34	BA	686	U
34	BA	688	G
34	BA	690	G
34	BA	691	A
34	BA	692	U
34	BA	693	G
34	BA	694	G
34	BA	695	A
34	BA	700	G
34	BA	711	C
34	BA	712	C
34	BA	713	C
34	BA	720	A
34	BA	721	A
34	BA	722	A
34	BA	724	A
34	BA	731	A
34	BA	735	A
34	BA	738	C
34	BA	739	A
34	BA	751	A
34	BA	752	A
34	BA	753	G

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Mol	Chain	Res	Type
34	BA	756	A
34	BA	757	G
34	BA	758	G
34	BA	764	G
34	BA	765	U
34	BA	766	A
34	BA	769	U
34	BA	770	G
34	BA	771	A
34	BA	772	G
34	BA	773	A
34	BA	774	A
34	BA	776	U
34	BA	781	U
34	BA	786	U
34	BA	787	A
34	BA	790	G
34	BA	794	G
34	BA	797	A
34	BA	798	G
34	BA	799	A
34	BA	800	G
34	BA	805	A
34	BA	806	U
34	BA	807	U
34	BA	808	U
34	BA	814	C
34	BA	815	C
34	BA	816	G
34	BA	821	G
34	BA	823	G
34	BA	828	A
34	BA	834	C
34	BA	836	U
34	BA	846	U
34	BA	847	U
34	BA	848	U
34	BA	849	G
34	BA	850	C
34	BA	863	G
34	BA	864	G
34	BA	865	C

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Mol	Chain	Res	Type
34	BA	874	G
34	BA	876	C
34	BA	879	C
34	BA	883	C
34	BA	884	G
34	BA	885	A
34	BA	895	U
34	BA	896	U
34	BA	898	G
34	BA	899	G
34	BA	900	A
34	BA	905	A
34	BA	906	A
34	BA	916	A
34	BA	929	A
34	BA	936	A
34	BA	945	A
34	BA	946	A
34	BA	948	C
34	BA	958	G
34	BA	960	C
34	BA	970	U
34	BA	973	U
34	BA	978	U
34	BA	982	A
34	BA	983	A
34	BA	993	C
34	BA	995	A
34	BA	996	U
34	BA	1006	G
34	BA	1007	G
34	BA	1012	A
34	BA	1013	A
34	BA	1015	G
34	BA	1019	C
34	BA	1020	A
34	BA	1022	C
34	BA	1023	G
34	BA	1024	A
34	BA	1035	A
34	BA	1036	G
34	BA	1037	C

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Mol	Chain	Res	Type
34	BA	1043	C
34	BA	1046	G
34	BA	1058	C
34	BA	1059	U
34	BA	1080	U
34	BA	1081	U
34	BA	1082	U
34	BA	1083	A
34	BA	1084	A
34	BA	1085	G
34	BA	1086	A
34	BA	1093	G
34	BA	1097	G
34	BA	1098	G
34	BA	1105	A
34	BA	1106	A
34	BA	1107	A
34	BA	1110	A
34	BA	1114	G
34	BA	1118	C
34	BA	1119	A
34	BA	1120	U
34	BA	1121	U
34	BA	1122	G
34	BA	1124	U
34	BA	1125	G
34	BA	1128	C
34	BA	1130	U
34	BA	1132	U
34	BA	1133	A
34	BA	1136	A
34	BA	1140	A
34	BA	1141	C
34	BA	1151	A
34	BA	1153	C
34	BA	1155	U
34	BA	1156	U
34	BA	1161	G
34	BA	1168	C
34	BA	1169	A
34	BA	1175	G
34	BA	1176	C

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Mol	Chain	Res	Type
34	BA	1180	A
34	BA	1181	G
34	BA	1183	U
34	BA	1184	A
34	BA	1186	U
34	BA	1187	U
34	BA	1188	U
34	BA	1189	A
34	BA	1196	C
34	BA	1198	U
34	BA	1199	U
34	BA	1200	U
34	BA	1201	G
34	BA	1205	A
34	BA	1206	C
34	BA	1207	A
34	BA	1208	U
34	BA	1209	A
34	BA	1210	A
34	BA	1211	G
34	BA	1216	G
34	BA	1217	A
34	BA	1229	G
34	BA	1230	G
34	BA	1244	G
34	BA	1257	U
34	BA	1263	A
34	BA	1264	U
34	BA	1266	A
34	BA	1272	U
34	BA	1273	U
34	BA	1293	A
34	BA	1294	C
34	BA	1295	U
34	BA	1296	U
34	BA	1297	G
34	BA	1298	U
34	BA	1306	U
34	BA	1307	U
34	BA	1308	C
34	BA	1309	U
34	BA	1310	C

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Mol	Chain	Res	Type
34	BA	1313	U
34	BA	1318	G
34	BA	1320	A
34	BA	1327	G
34	BA	1331	G
34	BA	1334	G
34	BA	1336	U
34	BA	1338	G
34	BA	1339	G
34	BA	1340	G
34	BA	1341	A
34	BA	1342	C
34	BA	1343	A
34	BA	1345	U
34	BA	1350	C
34	BA	1351	G
34	BA	1352	G
34	BA	1353	U
34	BA	1354	G
34	BA	1355	G
34	BA	1356	C
34	BA	1357	C
34	BA	1359	U
34	BA	1360	G
34	BA	1361	G
34	BA	1362	A
34	BA	1363	A
34	BA	1364	G
34	BA	1365	U
34	BA	1366	C
34	BA	1367	G
34	BA	1369	C
34	BA	1376	U
34	BA	1380	G
34	BA	1381	A
34	BA	1382	G
34	BA	1383	U
34	BA	1384	G
34	BA	1385	U
34	BA	1389	A
34	BA	1391	A
34	BA	1395	C

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Mol	Chain	Res	Type
34	BA	1396	A
34	BA	1397	C
34	BA	1398	C
34	BA	1399	A
34	BA	1403	G
34	BA	1405	A
34	BA	1410	C
34	BA	1423	U
34	BA	1425	G
34	BA	1426	A
34	BA	1427	U
34	BA	1430	C
34	BA	1431	G
34	BA	1434	U
34	BA	1444	G
34	BA	1445	U
34	BA	1446	G
34	BA	1447	C
34	BA	1448	G
34	BA	1449	U
34	BA	1450	G
34	BA	1452	U
34	BA	1453	U
34	BA	1454	G
34	BA	1455	C
34	BA	1457	C
34	BA	1458	A
34	BA	1459	U
34	BA	1460	U
34	BA	1471	U
34	BA	1472	G
34	BA	1474	G
34	BA	1476	G
34	BA	1477	C
34	BA	1478	G
34	BA	1481	U
34	BA	1482	A
34	BA	1486	U
34	BA	1490	U
34	BA	1491	U
34	BA	1494	G
34	BA	1495	A

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Mol	Chain	Res	Type
34	BA	1496	G
34	BA	1497	A
34	BA	1498	A
34	BA	1504	A
34	BA	1506	C
34	BA	1510	C
34	BA	1515	U
34	BA	1524	G
34	BA	1525	G
34	BA	1536	A
34	BA	1541	G
34	BA	1548	A
34	BA	1549	U
34	BA	1550	G
34	BA	1563	G
34	BA	1564	A
34	BA	1565	U
34	BA	1568	A
34	BA	1569	C
34	BA	1578	A
34	BA	1584	G
34	BA	1587	C
34	BA	1593	U
34	BA	1596	C
34	BA	1598	U
34	BA	1599	A
34	BA	1600	G
34	BA	1605	G
34	BA	1615	A
34	BA	1616	A
34	BA	1617	U
34	BA	1623	U
34	BA	1624	U
34	BA	1625	C
34	BA	1629	A
34	BA	1632	G
34	BA	1634	A
34	BA	1645	C
34	BA	1648	G
34	BA	1649	A
34	BA	1652	G
34	BA	1655	G

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Mol	Chain	Res	Type
34	BA	1660	A
34	BA	1661	U
34	BA	1674	G
34	BA	1675	C
34	BA	1688	G
34	BA	1689	U
34	BA	1693	U
34	BA	1695	G
34	BA	1696	G
34	BA	1697	U
34	BA	1698	C
34	BA	1706	A
34	BA	1713	U
34	BA	1714	A
34	BA	1715	C
34	BA	1716	A
34	BA	1717	C
34	BA	1718	C
34	BA	1725	U
34	BA	1726	U
34	BA	1727	A
34	BA	1729	G
34	BA	1730	A
34	BA	1732	A
34	BA	1735	G
34	BA	1737	A
34	BA	1776	G
34	BA	1777	U
34	BA	1788	U
34	BA	1789	A
34	BA	1793	G
34	BA	1794	A
34	BA	1795	A
34	BA	1796	A
34	BA	1797	A
34	BA	1798	G
34	BA	1799	G
34	BA	1804	A
34	BA	1805	C
34	BA	1806	A
34	BA	1810	A
34	BA	1811	A

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Mol	Chain	Res	Type
34	BA	1817	G
34	BA	1818	A
34	BA	1819	U
34	BA	1820	G
34	BA	1825	U
34	BA	1826	C
34	BA	1828	A
34	BA	1830	A
34	BA	1831	A
34	BA	1832	A
34	BA	1842	U
34	BA	1847	G
35	BB	2	C
35	BB	3	C
35	BB	4	C
35	BB	13	A
35	BB	17	U
35	BB	18	A
35	BB	19	C
35	BB	22	A
35	BB	23	U
35	BB	24	C
35	BB	25	A
35	BB	28	G
35	BB	29	C
35	BB	30	A
35	BB	31	U
35	BB	32	C
35	BB	33	A
35	BB	38	C
35	BB	47	C
35	BB	48	G
35	BB	49	A
35	BB	59	U
35	BB	60	A
35	BB	61	A
35	BB	62	C
35	BB	63	A
35	BB	64	U
35	BB	69	A
35	BB	70	A
35	BB	76	C

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Mol	Chain	Res	Type
35	BB	77	A
35	BB	81	A
35	BB	90	G
35	BB	91	G
35	BB	97	U
35	BB	98	A
35	BB	111	C
35	BB	136	A
35	BB	137	A
35	BB	139	G
35	BB	144	G
35	BB	146	U
35	BB	147	C
35	BB	148	C
35	BB	149	A
35	BB	152	G
35	BB	157	G
35	BB	158	C
35	BB	159	C
35	BB	161	G
35	BB	162	U
35	BB	168	U
35	BB	169	U
35	BB	253	G
35	BB	254	A
35	BB	255	A
35	BB	262	C
35	BB	264	U
35	BB	265	C
35	BB	267	C
35	BB	268	G
35	BB	282	A
35	BB	283	A
35	BB	287	C
35	BB	289	U
35	BB	290	U
35	BB	291	C
35	BB	296	G
35	BB	312	U
35	BB	315	C
35	BB	319	C
35	BB	322	G

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Mol	Chain	Res	Type
35	BB	323	C
35	BB	324	A
35	BB	325	G
35	BB	329	U
35	BB	330	U
35	BB	331	U
35	BB	333	C
35	BB	337	U
35	BB	340	U
35	BB	344	U
35	BB	346	U
35	BB	354	C
35	BB	357	C
35	BB	358	U
35	BB	359	A
35	BB	360	C
35	BB	361	A
35	BB	362	A
35	BB	363	A
35	BB	364	U
35	BB	371	C
35	BB	372	U
35	BB	381	C
35	BB	382	U
35	BB	383	U
35	BB	384	A
35	BB	385	C
35	BB	388	C
35	BB	389	G
35	BB	390	G
35	BB	399	A
35	BB	407	A
35	BB	408	U
35	BB	416	U
35	BB	426	A
35	BB	431	U
35	BB	437	U
35	BB	438	G
35	BB	444	U
35	BB	447	C
35	BB	448	G
35	BB	455	G

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Mol	Chain	Res	Type
35	BB	456	A
35	BB	461	U
35	BB	469	G
35	BB	473	U
35	BB	474	G
35	BB	475	A
35	BB	476	A
35	BB	477	U
35	BB	478	G
35	BB	481	A
35	BB	484	G
35	BB	488	G
35	BB	489	A
35	BB	490	G
35	BB	492	U
35	BB	495	A
35	BB	511	A
35	BB	516	G
35	BB	517	G
35	BB	519	A
35	BB	520	G
35	BB	522	A
35	BB	523	A
35	BB	524	C
35	BB	526	A
35	BB	530	C
35	BB	539	G
35	BB	540	G
35	BB	547	A
35	BB	548	A
35	BB	549	U
35	BB	555	G
35	BB	561	C
35	BB	562	A
35	BB	563	A
35	BB	565	U
35	BB	566	A
35	BB	571	C
35	BB	573	C
35	BB	574	G
35	BB	575	C
35	BB	577	U

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Mol	Chain	Res	Type
35	BB	580	A
35	BB	581	U
35	BB	582	G
35	BB	586	U
35	BB	597	C
35	BB	602	G
35	BB	603	U
35	BB	627	G
35	BB	640	A
35	BB	641	C
35	BB	642	G
35	BB	650	A
35	BB	651	G
35	BB	652	G
35	BB	655	U
35	BB	660	G
35	BB	664	A
35	BB	665	A
35	BB	668	A
35	BB	669	A
35	BB	670	G
35	BB	671	A
35	BB	672	C
35	BB	673	C
35	BB	678	U
35	BB	685	G
35	BB	702	G
35	BB	704	G
35	BB	710	A
35	BB	712	U
35	BB	713	U
35	BB	714	U
35	BB	715	G
35	BB	716	G
35	BB	717	A
35	BB	718	G
35	BB	719	G
35	BB	720	U
35	BB	721	G
35	BB	722	U
35	BB	725	U
35	BB	726	A

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Mol	Chain	Res	Type
35	BB	727	U
35	BB	728	A
35	BB	730	G
35	BB	735	A
35	BB	736	G
35	BB	746	A
35	BB	747	A
35	BB	748	A
35	BB	751	A
35	BB	752	A
35	BB	753	A
35	BB	754	U
35	BB	755	A
35	BB	758	A
35	BB	759	C
35	BB	760	C
35	BB	766	G
35	BB	767	A
35	BB	768	A
35	BB	769	C
35	BB	770	G
35	BB	772	U
35	BB	774	C
35	BB	778	A
35	BB	780	U
35	BB	781	U
35	BB	782	A
35	BB	789	G
35	BB	790	A
35	BB	791	A
35	BB	792	G
35	BB	793	A
35	BB	796	C
35	BB	797	C
35	BB	800	U
35	BB	803	U
35	BB	804	U
35	BB	805	G
35	BB	806	U
35	BB	818	U
35	BB	819	U
35	BB	820	C

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Mol	Chain	Res	Type
35	BB	823	G
35	BB	834	U
35	BB	838	G
35	BB	839	G
35	BB	840	C
35	BB	841	U
35	BB	846	A
35	BB	847	U
35	BB	849	A
35	BB	850	U
35	BB	851	U
35	BB	852	G
35	BB	853	U
35	BB	854	G
35	BB	860	U
35	BB	861	C
35	BB	862	U
35	BB	863	U
35	BB	869	G
35	BB	870	C
35	BB	877	A
35	BB	878	G
35	BB	879	G
35	BB	882	U
35	BB	884	U
35	BB	885	U
35	BB	890	U
35	BB	892	U
35	BB	893	U
35	BB	894	A
35	BB	895	U
35	BB	897	C
35	BB	900	C
35	BB	901	U
35	BB	906	G
35	BB	913	C
35	BB	914	U
35	BB	926	C
35	BB	928	C
35	BB	929	C
35	BB	932	U
35	BB	933	U

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Mol	Chain	Res	Type
35	BB	934	U
35	BB	943	U
35	BB	955	U
35	BB	965	G
35	BB	966	C
35	BB	971	A
35	BB	972	C
35	BB	973	G
35	BB	974	C
35	BB	975	G
35	BB	977	G
35	BB	978	C
35	BB	979	G
35	BB	980	G
35	BB	982	A
35	BB	983	C
35	BB	984	U
35	BB	985	A
35	BB	986	C
35	BB	987	U
35	BB	988	G
35	BB	989	C
35	BB	990	G
35	BB	991	C
35	BB	992	C
35	BB	995	C
35	BB	996	G
35	BB	1023	G
35	BB	1025	A
35	BB	1026	G
35	BB	1027	U
35	BB	1030	U
35	BB	1031	G
35	BB	1032	U
35	BB	1033	U
35	BB	1040	C
35	BB	1043	C
35	BB	1044	U
35	BB	1045	G
35	BB	1053	G
35	BB	1054	G
35	BB	1061	G

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Mol	Chain	Res	Type
35	BB	1066	G
35	BB	1073	A
35	BB	1084	A
35	BB	1099	U
35	BB	1103	A
35	BB	1119	G
35	BB	1121	A
35	BB	1124	G
35	BB	1125	A
35	BB	1126	A
35	BB	1128	U
35	BB	1135	U
35	BB	1136	G
35	BB	1137	G
35	BB	1138	A
35	BB	1140	C
35	BB	1141	A
35	BB	1143	A
35	BB	1146	C
35	BB	1150	A
35	BB	1152	U
35	BB	1155	U
35	BB	1156	U
35	BB	1161	G
35	BB	1166	A
35	BB	1167	C
35	BB	1176	G
35	BB	1177	U
35	BB	1200	A
35	BB	1201	G
35	BB	1202	G
35	BB	1209	A
35	BB	1218	G
35	BB	1220	A
35	BB	1221	G
35	BB	1222	A
35	BB	1224	C
35	BB	1225	A
35	BB	1226	G
35	BB	1227	G
35	BB	1228	A
35	BB	1229	A

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Mol	Chain	Res	Type
35	BB	1230	A
35	BB	1231	U
35	BB	1235	A
35	BB	1236	A
35	BB	1237	C
35	BB	1254	G
35	BB	1257	A
35	BB	1258	G
35	BB	1259	A
35	BB	1260	A
35	BB	1268	C
35	BB	1274	G
35	BB	1275	A
35	BB	1276	U
35	BB	1277	A
35	BB	1287	U
35	BB	1300	U
35	BB	1301	U
35	BB	1302	C
35	BB	1303	A
35	BB	1307	C
35	BB	1318	U
35	BB	1325	C
35	BB	1329	G
35	BB	1331	U
35	BB	1333	U
35	BB	1336	G
35	BB	1337	C
35	BB	1340	U
35	BB	1345	A
35	BB	1346	A
35	BB	1347	C
35	BB	1354	C
35	BB	1355	C
35	BB	1356	G
35	BB	1357	C
35	BB	1358	A
35	BB	1372	G
35	BB	1381	U
35	BB	1385	C
35	BB	1393	C
35	BB	1394	A

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Mol	Chain	Res	Type
35	BB	1395	G
35	BB	1400	C
35	BB	1403	G
35	BB	1405	G
35	BB	1409	G
35	BB	1412	U
35	BB	1413	U
35	BB	1415	G
35	BB	1429	A
35	BB	1436	U
35	BB	1437	U
35	BB	1440	A
35	BB	1441	C
35	BB	1448	U
35	BB	1450	G
35	BB	1453	G
35	BB	1454	G
35	BB	1455	A
35	BB	1456	G
35	BB	1459	U
35	BB	1460	G
35	BB	1461	C
35	BB	1462	G
35	BB	1464	G
35	BB	1465	U
35	BB	1468	A
35	BB	1469	A
35	BB	1471	A
35	BB	1472	U
35	BB	1473	U
35	BB	1474	A
35	BB	1475	U
35	BB	1481	C
35	BB	1482	A
35	BB	1490	G
35	BB	1495	U
35	BB	1499	U
35	BB	1501	U
35	BB	1502	U
35	BB	1503	U
35	BB	1504	U
35	BB	1505	U

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Mol	Chain	Res	Type
35	BB	1506	C
35	BB	1507	U
35	BB	1512	C
35	BB	1517	G
35	BB	1518	U
35	BB	1521	G
35	BB	1533	U
35	BB	1535	G
35	BB	1536	G
35	BB	1541	G
36	BC	2	A
36	BC	3	C
36	BC	22	U
36	BC	23	G
36	BC	32	U
36	BC	33	U
36	BC	34	U
36	BC	35	C
36	BC	47	C
36	BC	48	A
36	BC	51	A
36	BC	59	A
36	BC	61	A
36	BC	62	A
36	BC	63	G
36	BC	74	U
36	BC	75	G
36	BC	80	A
36	BC	81	U
36	BC	82	C
36	BC	83	A
36	BC	84	U
36	BC	85	U
36	BC	86	U
36	BC	89	U
36	BC	91	G
36	BC	92	C
36	BC	94	C
36	BC	103	A
36	BC	104	A
36	BC	105	C
36	BC	110	A

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Mol	Chain	Res	Type
36	BC	111	C
36	BC	117	A
36	BC	124	A
36	BC	130	U
36	BC	131	C
36	BC	132	U
36	BC	147	G
36	BC	149	A
36	BC	157	U
36	BC	158	U
36	BC	159	U
36	BC	160	C
36	BC	165	U
36	BC	169	G
37	BD	7	G
37	BD	11	A
37	BD	12	U
37	BD	13	A
37	BD	14	C
37	BD	21	G
37	BD	22	A
37	BD	26	C
37	BD	39	C
37	BD	40	C
37	BD	41	G
37	BD	42	A
37	BD	48	G
37	BD	50	A
37	BD	52	U
37	BD	53	U
37	BD	63	C
37	BD	64	A
37	BD	72	U
37	BD	74	A
37	BD	83	A
37	BD	84	U
37	BD	88	U
37	BD	89	G
37	BD	90	A
37	BD	99	G
37	BD	100	A
37	BD	101	A

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Mol	Chain	Res	Type
37	BD	108	G
37	BD	110	G
37	BD	113	G
38	BE	2	G
38	BE	4	A
38	BE	11	A
38	BE	12	A
38	BE	13	A
38	BE	19	G
38	BE	20	C
38	BE	22	A
38	BE	23	G
38	BE	24	G
38	BE	25	U
38	BE	26	G
38	BE	28	C
38	BE	53	U
38	BE	55	C
38	BE	56	U
38	BE	57	U
38	BE	71	A
38	BE	72	C
38	BE	78	C
38	BE	80	G
38	BE	82	C
38	BE	83	U
38	BE	84	U
38	BE	86	C
38	BE	89	G
38	BE	92	C
38	BE	102	U
38	BE	103	C
38	BE	104	G
38	BE	108	U
38	BE	109	C
38	BE	112	G
38	BE	113	C
38	BE	114	G
38	BE	118	C
38	BE	119	U
38	BE	125	C
38	BE	129	G

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Mol	Chain	Res	Type
38	BE	130	G
38	BE	131	C
38	BE	134	A
38	BE	135	A
38	BE	136	G
38	BE	137	A
38	BE	141	A
38	BE	142	A
38	BE	143	A
38	BE	144	A
38	BE	152	U
38	BE	153	C
38	BE	155	C
38	BE	156	C
38	BE	157	C
38	BE	159	A
38	BE	160	C
38	BE	161	G
38	BE	162	U
38	BE	165	U
38	BE	172	U
38	BE	173	G
38	BE	174	U
38	BE	175	U
38	BE	176	G
38	BE	179	A
38	BE	180	G
38	BE	181	U
38	BE	182	U
38	BE	186	C
38	BE	187	G
38	BE	189	A
38	BE	191	U
38	BE	203	C
38	BE	204	U
38	BE	206	G
38	BE	207	G
38	BE	209	U
38	BE	210	G
39	BF	2	G
39	BF	3	A
39	BF	6	C

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Mol	Chain	Res	Type
39	BF	7	G
39	BF	8	C
39	BF	10	A
39	BF	11	C
39	BF	12	U
39	BF	13	U
39	BF	14	C
39	BF	15	U
39	BF	16	C
39	BF	17	U
39	BF	18	U
39	BF	19	A
39	BF	20	U
39	BF	21	C
39	BF	22	U
39	BF	23	G
39	BF	24	G
39	BF	25	G
39	BF	26	U
39	BF	29	U
39	BF	31	U
39	BF	33	C
39	BF	34	C
39	BF	40	U
39	BF	41	U
39	BF	42	G
39	BF	43	U
39	BF	44	C
39	BF	45	G
39	BF	52	A
39	BF	53	G
39	BF	54	U
39	BF	57	C
39	BF	59	U
39	BF	60	C
39	BF	61	A
39	BF	62	U
39	BF	63	U
39	BF	64	U
39	BF	66	C
39	BF	67	A
39	BF	68	C

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Mol	Chain	Res	Type
39	BF	69	A
39	BF	71	G
39	BF	73	U
40	BG	2	U
40	BG	3	G
40	BG	11	G
40	BG	12	A
40	BG	13	A
40	BG	14	G
40	BG	15	G
40	BG	21	C
40	BG	22	G
40	BG	23	C
40	BG	24	A
40	BG	25	G
40	BG	34	A
40	BG	40	G
40	BG	84	U
40	BG	85	C
40	BG	86	U
40	BG	87	G
40	BG	99	A
40	BG	106	G
40	BG	107	U
40	BG	108	G
40	BG	114	A
40	BG	120	U
40	BG	121	C
40	BG	141	A
40	BG	147	U
40	BG	150	A
40	BG	156	G
40	BG	158	A
40	BG	159	A
40	BG	166	C
40	BG	167	C
40	BG	169	A
40	BG	170	G
40	BG	171	A
40	BG	172	C
40	BG	173	C
40	BG	180	C

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Mol	Chain	Res	Type
40	BG	181	C
41	BH	2	U
41	BH	3	U
41	BH	11	C
41	BH	12	U
41	BH	15	A
41	BH	21	G
41	BH	24	U
41	BH	25	A
41	BH	26	C
41	BH	27	A
41	BH	28	U
41	BH	30	C
41	BH	34	G
41	BH	38	G
41	BH	39	G
41	BH	40	C
41	BH	41	A
41	BH	42	U
41	BH	43	G
41	BH	44	A
41	BH	49	C
41	BH	50	A
41	BH	51	C
41	BH	60	A
41	BH	61	C
41	BH	62	C
41	BH	72	G
41	BH	73	A
41	BH	74	G
41	BH	77	G
41	BH	79	A
41	BH	80	C
41	BH	81	U
41	BH	82	U
41	BH	84	A
41	BH	86	G
41	BH	88	C
41	BH	89	C
41	BH	90	C
41	BH	93	G
41	BH	101	A

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Mol	Chain	Res	Type
41	BH	102	C
41	BH	103	C
41	BH	115	A
41	BH	117	U
41	BH	122	U
41	BH	123	G
41	BH	129	G
41	BH	135	U
85	AA	2	A
85	AA	3	U
85	AA	4	C
85	AA	9	U
85	AA	25	C
85	AA	26	A
85	AA	28	A
85	AA	34	G
85	AA	39	A
85	AA	42	G
85	AA	43	A
85	AA	44	C
85	AA	45	U
85	AA	46	U
85	AA	47	A
85	AA	48	G
85	AA	51	A
85	AA	56	U
85	AA	57	G
85	AA	59	C
85	AA	60	U
85	AA	61	C
85	AA	66	U
85	AA	67	C
85	AA	68	A
85	AA	73	A
85	AA	74	U
85	AA	77	C
85	AA	82	A
85	AA	85	U
85	AA	86	G
85	AA	90	A
85	AA	91	U
85	AA	92	G

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Mol	Chain	Res	Type
85	AA	96	C
85	AA	97	A
85	AA	99	U
85	AA	101	C
85	AA	102	A
85	AA	103	U
85	AA	104	C
85	AA	107	A
85	AA	111	A
85	AA	112	A
85	AA	116	G
85	AA	121	C
85	AA	123	A
85	AA	124	A
85	AA	125	A
85	AA	126	U
85	AA	127	U
85	AA	128	U
85	AA	129	U
85	AA	136	U
85	AA	137	C
85	AA	142	U
85	AA	143	U
85	AA	144	A
85	AA	145	C
85	AA	162	A
85	AA	171	U
85	AA	174	U
85	AA	175	A
85	AA	179	G
85	AA	180	A
85	AA	181	A
85	AA	182	C
85	AA	183	C
85	AA	184	A
85	AA	185	A
85	AA	193	C
85	AA	197	C
85	AA	198	U
85	AA	200	U
85	AA	202	U
85	AA	205	A

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Mol	Chain	Res	Type
85	AA	207	G
85	AA	208	U
85	AA	209	C
85	AA	210	G
85	AA	211	C
85	AA	212	G
85	AA	222	U
85	AA	224	C
85	AA	225	G
85	AA	230	U
85	AA	236	G
85	AA	237	G
85	AA	239	G
85	AA	240	A
85	AA	241	U
85	AA	242	G
85	AA	243	A
85	AA	244	G
85	AA	246	C
85	AA	251	A
85	AA	253	C
85	AA	254	G
85	AA	263	A
85	AA	264	A
85	AA	265	A
85	AA	266	U
85	AA	267	U
85	AA	269	G
85	AA	280	U
85	AA	282	C
85	AA	284	C
85	AA	290	G
85	AA	294	G
85	AA	295	U
85	AA	296	A
85	AA	298	C
85	AA	299	A
85	AA	300	C
85	AA	301	U
85	AA	302	C
85	AA	303	A
85	AA	304	G

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Mol	Chain	Res	Type
85	AA	307	G
85	AA	308	U
85	AA	311	U
85	AA	312	G
85	AA	313	A
85	AA	315	U
85	AA	316	C
85	AA	318	A
85	AA	326	C
85	AA	327	G
85	AA	328	U
85	AA	329	G
85	AA	330	C
85	AA	331	G
85	AA	332	A
85	AA	333	A
85	AA	334	A
85	AA	338	G
85	AA	344	U
85	AA	345	U
85	AA	356	U
85	AA	357	C
85	AA	358	U
85	AA	363	A
85	AA	375	C
85	AA	378	A
85	AA	379	U
85	AA	380	C
85	AA	382	G
85	AA	385	A
85	AA	386	G
85	AA	387	U
85	AA	388	G
85	AA	389	A
85	AA	390	U
85	AA	394	C
85	AA	395	G
85	AA	404	A
85	AA	412	G
85	AA	417	U
85	AA	418	G
85	AA	419	A

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Mol	Chain	Res	Type
85	AA	425	G
85	AA	435	A
85	AA	442	G
85	AA	443	A
85	AA	444	U
85	AA	445	U
85	AA	446	C
85	AA	453	G
85	AA	455	G
85	AA	464	A
85	AA	465	A
85	AA	466	A
85	AA	467	U
85	AA	468	A
85	AA	469	G
85	AA	482	C
85	AA	483	G
85	AA	487	G
85	AA	489	C
85	AA	495	G
85	AA	504	U
85	AA	505	U
85	AA	510	A
85	AA	518	A
85	AA	519	A
85	AA	520	A
85	AA	521	A
85	AA	522	A
85	AA	524	A
85	AA	526	G
85	AA	529	G
85	AA	530	A
85	AA	533	C
85	AA	538	A
85	AA	557	G
85	AA	571	G
85	AA	576	U
85	AA	578	U
85	AA	580	C
85	AA	582	A
85	AA	589	A
85	AA	590	U

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Mol	Chain	Res	Type
85	AA	591	A
85	AA	603	C
85	AA	604	C
85	AA	605	A
85	AA	606	A
85	AA	607	U
85	AA	608	A
85	AA	612	A
85	AA	615	A
85	AA	616	A
85	AA	618	A
85	AA	619	A
85	AA	620	U
85	AA	625	G
85	AA	628	C
85	AA	629	A
85	AA	633	C
85	AA	634	U
85	AA	639	C
85	AA	640	C
85	AA	648	G
85	AA	652	U
85	AA	653	A
85	AA	654	A
85	AA	655	U
85	AA	657	C
85	AA	660	G
85	AA	668	A
85	AA	669	G
85	AA	681	G
85	AA	682	C
85	AA	683	U
85	AA	684	G
85	AA	685	U
85	AA	687	G
85	AA	688	C
85	AA	693	A
85	AA	694	A
85	AA	696	G
85	AA	697	G
85	AA	698	G
85	AA	709	A

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Mol	Chain	Res	Type
85	AA	711	C
85	AA	712	U
85	AA	713	G
85	AA	714	U
85	AA	715	G
85	AA	724	A
85	AA	725	G
85	AA	726	U
85	AA	727	U
85	AA	728	U
85	AA	736	U
85	AA	737	G
85	AA	738	C
85	AA	739	C
85	AA	740	A
85	AA	741	G
85	AA	742	U
85	AA	744	C
85	AA	749	C
85	AA	750	A
85	AA	752	C
85	AA	753	U
85	AA	754	C
85	AA	756	G
85	AA	760	U
85	AA	761	G
85	AA	762	U
85	AA	763	U
85	AA	764	U
85	AA	765	U
85	AA	766	G
85	AA	767	A
85	AA	770	C
85	AA	771	A
85	AA	772	C
85	AA	776	C
85	AA	777	U
85	AA	779	G
85	AA	786	G
85	AA	787	U
85	AA	789	A
85	AA	790	A

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Mol	Chain	Res	Type
85	AA	791	C
85	AA	792	A
85	AA	793	C
85	AA	795	C
85	AA	797	C
85	AA	798	A
85	AA	799	G
85	AA	801	U
85	AA	803	C
85	AA	804	A
85	AA	805	A
85	AA	806	G
85	AA	809	A
85	AA	810	C
85	AA	815	G
85	AA	816	A
85	AA	818	C
85	AA	825	U
85	AA	826	C
85	AA	828	U
85	AA	829	C
85	AA	830	A
85	AA	831	C
85	AA	832	U
85	AA	833	U
85	AA	834	U
85	AA	836	A
85	AA	837	C
85	AA	838	G
85	AA	845	A
85	AA	846	U
85	AA	847	G
85	AA	853	G
85	AA	854	A
85	AA	855	G
85	AA	856	G
85	AA	861	G
85	AA	862	U
85	AA	868	A
85	AA	869	A
85	AA	873	U
85	AA	874	A

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Mol	Chain	Res	Type
85	AA	875	C
85	AA	883	A
85	AA	884	A
85	AA	885	A
85	AA	886	A
85	AA	887	A
85	AA	888	A
85	AA	889	G
85	AA	890	U
85	AA	891	G
85	AA	892	C
85	AA	893	G
85	AA	895	C
85	AA	899	A
85	AA	901	C
85	AA	902	A
85	AA	903	G
85	AA	904	U
85	AA	906	U
85	AA	907	G
85	AA	908	C
85	AA	909	C
85	AA	910	G
85	AA	914	U
85	AA	919	U
85	AA	923	A
85	AA	924	A
85	AA	927	A
85	AA	931	G
85	AA	934	A
85	AA	935	A
85	AA	937	G
85	AA	938	A
85	AA	940	G
85	AA	941	C
85	AA	942	A
85	AA	943	U
85	AA	945	A
85	AA	957	A
85	AA	961	U
85	AA	964	C
85	AA	965	G

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Mol	Chain	Res	Type
85	AA	974	U
85	AA	988	C
85	AA	989	U
85	AA	990	U
85	AA	998	U
85	AA	999	A
85	AA	1000	U
85	AA	1005	C
85	AA	1010	U
85	AA	1012	C
85	AA	1013	C
85	AA	1014	U
85	AA	1017	G
85	AA	1020	C
85	AA	1022	G
85	AA	1023	U
85	AA	1026	U
85	AA	1030	U
85	AA	1034	U
85	AA	1037	U
85	AA	1040	U
85	AA	1043	U
85	AA	1051	A
85	AA	1052	C
85	AA	1054	U
85	AA	1055	U
85	AA	1058	G
85	AA	1060	U
85	AA	1061	C
85	AA	1062	U
85	AA	1063	U
85	AA	1068	A
85	AA	1073	U
85	AA	1075	U
85	AA	1076	U
85	AA	1077	U
85	AA	1078	A
85	AA	1079	C
85	AA	1080	A
85	AA	1086	U
85	AA	1091	C
85	AA	1098	C

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Mol	Chain	Res	Type
85	AA	1103	A
85	AA	1104	G
85	AA	1106	A
85	AA	1107	A
85	AA	1108	U
85	AA	1109	G
85	AA	1111	A
85	AA	1112	G
85	AA	1113	G
85	AA	1114	A
85	AA	1121	U
85	AA	1125	G
85	AA	1130	G
85	AA	1136	A
85	AA	1138	U
85	AA	1140	G
85	AA	1143	C
85	AA	1145	U
85	AA	1148	G
85	AA	1153	G
85	AA	1154	A
85	AA	1157	U
85	AA	1158	U
85	AA	1159	C
85	AA	1161	U
85	AA	1162	A
85	AA	1163	G
85	AA	1177	G
85	AA	1179	A
85	AA	1183	C
85	AA	1184	A
85	AA	1191	G
85	AA	1193	A
85	AA	1208	C
85	AA	1209	U
85	AA	1212	C
85	AA	1213	U
85	AA	1214	C
85	AA	1215	A
85	AA	1219	A
85	AA	1236	G
85	AA	1237	A

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Mol	Chain	Res	Type
85	AA	1241	A
85	AA	1242	A
85	AA	1246	G
85	AA	1249	U
85	AA	1250	A
85	AA	1253	G
85	AA	1272	G
85	AA	1275	A
85	AA	1276	A
85	AA	1278	C
85	AA	1279	A
85	AA	1281	G
85	AA	1287	C
85	AA	1288	A
85	AA	1290	G
85	AA	1291	A
85	AA	1292	A
85	AA	1293	U
85	AA	1294	U
85	AA	1299	A
85	AA	1300	A
85	AA	1302	A
85	AA	1303	U
85	AA	1311	U
85	AA	1315	C
85	AA	1319	U
85	AA	1321	G
85	AA	1330	U
85	AA	1331	G
85	AA	1352	U
85	AA	1353	U
85	AA	1354	A
85	AA	1357	U
85	AA	1358	A
85	AA	1359	U
85	AA	1360	C
85	AA	1368	G
85	AA	1372	C
85	AA	1373	U
85	AA	1379	A
85	AA	1381	C
85	AA	1383	C

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Mol	Chain	Res	Type
85	AA	1391	U
85	AA	1393	C
85	AA	1397	U
85	AA	1399	U
85	AA	1400	U
85	AA	1406	U
85	AA	1407	C
85	AA	1416	U
85	AA	1417	U
85	AA	1420	U
85	AA	1422	A
85	AA	1423	C
85	AA	1425	G
85	AA	1430	A
85	AA	1433	C
85	AA	1434	U
85	AA	1436	A
85	AA	1437	G
85	AA	1441	G
85	AA	1446	U
85	AA	1449	C
85	AA	1450	U
85	AA	1457	C
85	AA	1458	G
85	AA	1459	C
85	AA	1466	U
85	AA	1467	U
85	AA	1468	G
85	AA	1469	G
85	AA	1470	A
85	AA	1471	G
85	AA	1474	U
85	AA	1475	A
85	AA	1476	C
85	AA	1477	A
85	AA	1478	G
85	AA	1486	G
85	AA	1490	A
85	AA	1491	G
85	AA	1492	U
85	AA	1493	A
85	AA	1494	C

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Mol	Chain	Res	Type
85	AA	1495	G
85	AA	1515	A
85	AA	1527	G
85	AA	1528	A
85	AA	1532	G
85	AA	1535	C
85	AA	1536	C
85	AA	1537	A
85	AA	1547	G
85	AA	1560	A
85	AA	1561	A
85	AA	1563	U
85	AA	1567	C
85	AA	1570	A
85	AA	1571	A
85	AA	1572	C
85	AA	1573	A
85	AA	1574	C
85	AA	1576	G
85	AA	1577	G
85	AA	1578	G
85	AA	1579	A
85	AA	1580	A
85	AA	1584	U
85	AA	1585	A
85	AA	1594	G
85	AA	1595	G
85	AA	1596	A
85	AA	1597	C
85	AA	1605	G
85	AA	1606	G
85	AA	1614	G
85	AA	1616	U
85	AA	1618	G
85	AA	1620	G
85	AA	1621	U
85	AA	1622	G
85	AA	1623	U
85	AA	1633	A
85	AA	1634	U
85	AA	1635	C
85	AA	1638	C

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Mol	Chain	Res	Type
85	AA	1643	U
85	AA	1646	U
85	AA	1647	G
85	AA	1650	G
85	AA	1651	C
85	AA	1661	U
85	AA	1662	U
85	AA	1663	U
85	AA	1664	G
85	AA	1665	G
85	AA	1683	U
85	AA	1684	U
85	AA	1689	G
85	AA	1691	U
85	AA	1692	U
85	AA	1698	A
85	AA	1699	A
85	AA	1702	G
85	AA	1706	A
85	AA	1717	G
85	AA	1718	C
85	AA	1719	C
85	AA	1722	G
85	AA	1724	A
85	AA	1725	G
85	AA	1726	G
85	AA	1729	C
85	AA	1731	G
85	AA	1736	U
85	AA	1752	C
85	AA	1753	A
85	AA	1754	G
85	AA	1756	C
85	AA	1758	C
85	AA	1759	U
85	AA	1760	C
85	AA	1770	U
85	AA	1772	U
85	AA	1773	U
85	AA	1774	U
85	AA	1785	U
85	AA	1788	U

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Mol	Chain	Res	Type
85	AA	1789	C
85	AA	1791	U
85	AA	1792	C
85	AA	1793	A
85	AA	1798	U
85	AA	1800	U
85	AA	1803	U
85	AA	1804	U
85	AA	1808	G
85	AA	1809	G
85	AA	1810	C
85	AA	1812	C
85	AA	1813	C
85	AA	1814	U
85	AA	1815	U
85	AA	1817	U
85	AA	1820	G
85	AA	1826	U
85	AA	1827	U
85	AA	1833	C
85	AA	1834	U
85	AA	1836	U
85	AA	1837	U
85	AA	1839	G
85	AA	1841	G
85	AA	1849	A
85	AA	1851	A
85	AA	1852	U
85	AA	1853	U
85	AA	1854	U
85	AA	1860	A
85	AA	1861	A
85	AA	1866	A
85	AA	1869	U
85	AA	1870	C
85	AA	1872	G
85	AA	1874	G
85	AA	1875	A
85	AA	1884	A
85	AA	1885	A
85	AA	1887	G
85	AA	1897	A

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Mol	Chain	Res	Type
85	AA	1898	C
85	AA	1899	A
85	AA	1911	A
85	AA	1912	U
85	AA	1913	G
85	AA	1920	A
85	AA	1921	G
85	AA	1922	A
85	AA	1923	A
85	AA	1924	C
85	AA	1925	A
85	AA	1926	A
85	AA	1929	G
85	AA	1931	C
85	AA	1932	C
85	AA	1933	G
85	AA	1934	A
85	AA	1943	U
85	AA	1945	A
85	AA	1947	A
85	AA	1957	C
85	AA	1958	C
85	AA	1959	G
85	AA	1960	C
85	AA	1964	A
85	AA	1973	G
85	AA	1974	C
85	AA	1979	A
85	AA	1980	A
85	AA	1981	A
85	AA	1982	C
85	AA	1983	C
85	AA	1984	A
85	AA	1986	G
85	AA	1987	G
85	AA	1988	A
85	AA	1989	A
85	AA	1990	U
85	AA	1993	C
85	AA	2000	C
85	AA	2001	C
85	AA	2003	C

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Mol	Chain	Res	Type
85	AA	2004	U
85	AA	2005	U
85	AA	2006	G
85	AA	2008	G
85	AA	2009	A
85	AA	2010	C
85	AA	2020	C
85	AA	2023	U
85	AA	2024	U
85	AA	2025	A
85	AA	2026	U
85	AA	2034	G
85	AA	2035	C
85	AA	2040	A
85	AA	2041	G
85	AA	2050	C
85	AA	2051	G
85	AA	2057	G
85	AA	2062	U
85	AA	2068	A
85	AA	2081	A
85	AA	2086	C
85	AA	2098	A
85	AA	2099	C
85	AA	2101	C
85	AA	2102	A
85	AA	2103	C
85	AA	2104	C
85	AA	2105	G
85	AA	2125	A
85	AA	2133	A
85	AA	2144	C
85	AA	2146	G
85	AA	2147	A
85	AA	2159	C
85	AA	2160	U
85	AA	2161	C
85	AA	2172	A
85	AA	2197	A
85	AA	2208	G
85	AA	2211	G
85	AA	2212	U

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Mol	Chain	Res	Type
85	AA	2213	A
85	AA	2215	C
85	AA	2216	A
85	AA	2218	G
85	AA	2219	G
85	AA	2220	U
85	AA	2222	G
85	AA	2230	U
85	AA	2231	G
85	AA	2234	C
85	AA	2243	G
85	AA	2244	G
85	AA	2245	A
85	AA	2246	U
85	AA	2247	C
85	AA	2248	A
85	AA	2249	U
85	AA	2250	U
85	AA	2251	U
86	AB	2	C
86	AB	3	C
86	AB	4	C
86	AB	8	U
86	AB	9	A
86	AB	15	G
86	AB	17	C
86	AB	18	G
86	AB	19	G
86	AB	20	U
86	AB	23	A
86	AB	27	G
86	AB	35	A
86	AB	37	A
86	AB	43	C
86	AB	46	G
86	AB	47	U
86	AB	48	C
86	AB	50	U
86	AB	55	U
86	AB	57	G
86	AB	58	A
86	AB	59	U

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Mol	Chain	Res	Type
86	AB	60	U
86	AB	65	G
86	AB	66	U
86	AB	70	G
86	AB	71	G
86	AB	72	C
86	AB	73	A

All (790) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
34	BA	13	U
34	BA	21	C
34	BA	62	A
34	BA	70	C
34	BA	84	U
34	BA	91	C
34	BA	114	U
34	BA	116	G
34	BA	120	A
34	BA	131	A
34	BA	140	C
34	BA	148	G
34	BA	156	U
34	BA	160	G
34	BA	165	C
34	BA	174	A
34	BA	177	G
34	BA	179	U
34	BA	187	G
34	BA	188	C
34	BA	215	C
34	BA	217	C
34	BA	218	G
34	BA	224	G
34	BA	225	A
34	BA	226	A
34	BA	247	U
34	BA	261	A
34	BA	270	U
34	BA	279	U
34	BA	281	C

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Mol	Chain	Res	Type
34	BA	283	U
34	BA	289	A
34	BA	294	C
34	BA	298	G
34	BA	301	U
34	BA	302	A
34	BA	303	C
34	BA	316	G
34	BA	317	U
34	BA	323	C
34	BA	344	G
34	BA	347	A
34	BA	360	C
34	BA	367	G
34	BA	368	U
34	BA	392	A
34	BA	394	A
34	BA	404	C
34	BA	411	C
34	BA	415	C
34	BA	463	A
34	BA	464	U
34	BA	469	C
34	BA	482	C
34	BA	487	A
34	BA	488	C
34	BA	508	C
34	BA	510	U
34	BA	511	U
34	BA	518	C
34	BA	519	G
34	BA	521	C
34	BA	525	A
34	BA	526	C
34	BA	530	A
34	BA	531	C
34	BA	538	G
34	BA	547	C
34	BA	561	U
34	BA	570	G
34	BA	574	U
34	BA	575	U

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Mol	Chain	Res	Type
34	BA	577	U
34	BA	578	C
34	BA	579	U
34	BA	581	U
34	BA	588	C
34	BA	592	G
34	BA	595	U
34	BA	596	G
34	BA	599	U
34	BA	607	C
34	BA	611	A
34	BA	625	U
34	BA	628	U
34	BA	629	G
34	BA	647	U
34	BA	650	C
34	BA	667	U
34	BA	672	G
34	BA	674	G
34	BA	680	C
34	BA	683	C
34	BA	687	G
34	BA	690	G
34	BA	712	C
34	BA	720	A
34	BA	722	A
34	BA	738	C
34	BA	739	A
34	BA	751	A
34	BA	752	A
34	BA	764	G
34	BA	769	U
34	BA	770	G
34	BA	771	A
34	BA	772	G
34	BA	785	G
34	BA	786	U
34	BA	787	A
34	BA	797	A
34	BA	799	A
34	BA	805	A
34	BA	807	U

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Mol	Chain	Res	Type
34	BA	813	C
34	BA	827	A
34	BA	833	U
34	BA	845	U
34	BA	847	U
34	BA	848	U
34	BA	849	G
34	BA	863	G
34	BA	875	G
34	BA	885	A
34	BA	895	U
34	BA	897	U
34	BA	915	A
34	BA	953	G
34	BA	972	C
34	BA	982	A
34	BA	995	A
34	BA	1007	G
34	BA	1023	G
34	BA	1035	A
34	BA	1036	G
34	BA	1042	U
34	BA	1079	C
34	BA	1080	U
34	BA	1083	A
34	BA	1085	G
34	BA	1097	G
34	BA	1120	U
34	BA	1121	U
34	BA	1131	G
34	BA	1168	C
34	BA	1179	U
34	BA	1180	A
34	BA	1187	U
34	BA	1188	U
34	BA	1197	U
34	BA	1198	U
34	BA	1199	U
34	BA	1200	U
34	BA	1216	G
34	BA	1262	A
34	BA	1267	A

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Mol	Chain	Res	Type
34	BA	1293	A
34	BA	1295	U
34	BA	1296	U
34	BA	1297	G
34	BA	1305	A
34	BA	1306	U
34	BA	1307	U
34	BA	1309	U
34	BA	1313	U
34	BA	1319	A
34	BA	1326	U
34	BA	1331	G
34	BA	1340	G
34	BA	1341	A
34	BA	1355	G
34	BA	1359	U
34	BA	1363	A
34	BA	1381	A
34	BA	1421	A
34	BA	1422	A
34	BA	1425	G
34	BA	1443	U
34	BA	1447	C
34	BA	1452	U
34	BA	1453	U
34	BA	1454	G
34	BA	1457	C
34	BA	1459	U
34	BA	1470	G
34	BA	1471	U
34	BA	1475	G
34	BA	1477	C
34	BA	1490	U
34	BA	1494	G
34	BA	1495	A
34	BA	1496	G
34	BA	1497	A
34	BA	1505	G
34	BA	1509	U
34	BA	1515	U
34	BA	1563	G
34	BA	1615	A

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Mol	Chain	Res	Type
34	BA	1616	A
34	BA	1622	U
34	BA	1625	C
34	BA	1628	A
34	BA	1635	A
34	BA	1648	G
34	BA	1651	C
34	BA	1660	A
34	BA	1674	G
34	BA	1678	U
34	BA	1697	U
34	BA	1713	U
34	BA	1716	A
34	BA	1725	U
34	BA	1775	U
34	BA	1793	G
34	BA	1794	A
34	BA	1795	A
34	BA	1796	A
34	BA	1797	A
34	BA	1805	C
34	BA	1806	A
34	BA	1809	G
34	BA	1816	G
34	BA	1825	U
34	BA	1831	A
35	BB	3	C
35	BB	28	G
35	BB	29	C
35	BB	30	A
35	BB	31	U
35	BB	32	C
35	BB	48	G
35	BB	49	A
35	BB	60	A
35	BB	61	A
35	BB	62	C
35	BB	69	A
35	BB	97	U
35	BB	98	A
35	BB	116	G
35	BB	145	G

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Mol	Chain	Res	Type
35	BB	147	C
35	BB	148	C
35	BB	261	C
35	BB	267	C
35	BB	295	U
35	BB	353	G
35	BB	361	A
35	BB	371	C
35	BB	380	G
35	BB	381	C
35	BB	383	U
35	BB	388	C
35	BB	443	A
35	BB	447	C
35	BB	475	A
35	BB	477	U
35	BB	488	G
35	BB	494	C
35	BB	516	G
35	BB	523	A
35	BB	524	C
35	BB	548	A
35	BB	561	C
35	BB	602	G
35	BB	639	A
35	BB	641	C
35	BB	650	A
35	BB	659	C
35	BB	667	G
35	BB	671	A
35	BB	713	U
35	BB	715	G
35	BB	735	A
35	BB	747	A
35	BB	754	U
35	BB	780	U
35	BB	792	G
35	BB	795	A
35	BB	796	C
35	BB	797	C
35	BB	799	A
35	BB	803	U

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Mol	Chain	Res	Type
35	BB	833	G
35	BB	838	G
35	BB	845	C
35	BB	848	A
35	BB	850	U
35	BB	851	U
35	BB	853	U
35	BB	860	U
35	BB	861	C
35	BB	863	U
35	BB	877	A
35	BB	878	G
35	BB	883	G
35	BB	884	U
35	BB	885	U
35	BB	892	U
35	BB	894	A
35	BB	896	C
35	BB	912	C
35	BB	942	G
35	BB	970	C
35	BB	971	A
35	BB	972	C
35	BB	974	C
35	BB	976	U
35	BB	981	A
35	BB	985	A
35	BB	986	C
35	BB	990	G
35	BB	991	C
35	BB	992	C
35	BB	1024	G
35	BB	1026	G
35	BB	1029	U
35	BB	1030	U
35	BB	1031	G
35	BB	1032	U
35	BB	1039	A
35	BB	1044	U
35	BB	1063	C
35	BB	1098	G
35	BB	1124	G

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Mol	Chain	Res	Type
35	BB	1126	A
35	BB	1136	G
35	BB	1151	A
35	BB	1165	A
35	BB	1166	A
35	BB	1167	C
35	BB	1178	A
35	BB	1200	A
35	BB	1219	A
35	BB	1220	A
35	BB	1221	G
35	BB	1225	A
35	BB	1227	G
35	BB	1230	A
35	BB	1235	A
35	BB	1258	G
35	BB	1259	A
35	BB	1275	A
35	BB	1276	U
35	BB	1300	U
35	BB	1325	C
35	BB	1354	C
35	BB	1436	U
35	BB	1437	U
35	BB	1454	G
35	BB	1459	U
35	BB	1460	G
35	BB	1461	C
35	BB	1464	G
35	BB	1468	A
35	BB	1472	U
35	BB	1474	A
35	BB	1482	A
35	BB	1489	A
35	BB	1498	G
35	BB	1502	U
35	BB	1503	U
35	BB	1504	U
35	BB	1505	U
35	BB	1506	C
35	BB	1517	G
35	BB	1520	C

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Mol	Chain	Res	Type
35	BB	1535	G
36	BC	1	A
36	BC	2	A
36	BC	22	U
36	BC	31	A
36	BC	32	U
36	BC	33	U
36	BC	34	U
36	BC	60	U
36	BC	61	A
36	BC	74	U
36	BC	80	A
36	BC	81	U
36	BC	84	U
36	BC	98	C
36	BC	110	A
36	BC	123	G
36	BC	156	A
36	BC	157	U
36	BC	158	U
36	BC	159	U
37	BD	12	U
37	BD	13	A
37	BD	24	U
37	BD	47	U
37	BD	51	G
37	BD	71	G
37	BD	83	A
37	BD	88	U
37	BD	98	G
38	BE	18	U
38	BE	19	G
38	BE	21	C
38	BE	22	A
38	BE	26	G
38	BE	32	U
38	BE	52	U
38	BE	70	C
38	BE	80	G
38	BE	82	C
38	BE	88	G
38	BE	102	U

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Mol	Chain	Res	Type
38	BE	111	C
38	BE	117	A
38	BE	118	C
38	BE	129	G
38	BE	133	C
38	BE	134	A
38	BE	135	A
38	BE	136	G
38	BE	141	A
38	BE	142	A
38	BE	155	C
38	BE	175	U
38	BE	181	U
38	BE	186	C
38	BE	208	G
39	BF	1	C
39	BF	2	G
39	BF	5	U
39	BF	6	C
39	BF	7	G
39	BF	8	C
39	BF	9	C
39	BF	11	C
39	BF	18	U
39	BF	20	U
39	BF	23	G
39	BF	25	G
39	BF	32	G
39	BF	40	U
39	BF	41	U
39	BF	43	U
39	BF	51	C
39	BF	53	G
39	BF	60	C
39	BF	62	U
39	BF	63	U
39	BF	66	C
39	BF	71	G
40	BG	12	A
40	BG	14	G
40	BG	21	C
40	BG	84	U

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Mol	Chain	Res	Type
40	BG	106	G
40	BG	107	U
40	BG	149	U
40	BG	168	A
41	BH	24	U
41	BH	26	C
41	BH	27	A
41	BH	30	C
41	BH	33	G
41	BH	38	G
41	BH	41	A
41	BH	42	U
41	BH	62	C
41	BH	71	C
41	BH	72	G
41	BH	81	U
41	BH	85	C
41	BH	116	A
41	BH	117	U
41	BH	118	U
85	AA	1	G
85	AA	3	U
85	AA	8	U
85	AA	25	C
85	AA	28	A
85	AA	33	U
85	AA	39	A
85	AA	42	G
85	AA	44	C
85	AA	45	U
85	AA	46	U
85	AA	47	A
85	AA	56	U
85	AA	67	C
85	AA	82	A
85	AA	90	A
85	AA	96	C
85	AA	100	A
85	AA	101	C
85	AA	111	A
85	AA	123	A
85	AA	125	A

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Mol	Chain	Res	Type
85	AA	127	U
85	AA	143	U
85	AA	144	A
85	AA	145	C
85	AA	157	G
85	AA	174	U
85	AA	183	C
85	AA	192	G
85	AA	197	C
85	AA	206	U
85	AA	240	A
85	AA	243	A
85	AA	245	A
85	AA	262	G
85	AA	265	A
85	AA	266	U
85	AA	284	C
85	AA	293	A
85	AA	294	G
85	AA	295	U
85	AA	300	C
85	AA	303	A
85	AA	304	G
85	AA	312	G
85	AA	314	C
85	AA	315	U
85	AA	325	C
85	AA	326	C
85	AA	327	G
85	AA	329	G
85	AA	332	A
85	AA	333	A
85	AA	344	U
85	AA	357	C
85	AA	378	A
85	AA	385	A
85	AA	387	U
85	AA	388	G
85	AA	394	C
85	AA	403	G
85	AA	411	U
85	AA	416	U

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Mol	Chain	Res	Type
85	AA	418	G
85	AA	425	G
85	AA	435	A
85	AA	445	U
85	AA	452	A
85	AA	454	G
85	AA	463	G
85	AA	465	A
85	AA	467	U
85	AA	482	C
85	AA	486	G
85	AA	488	G
85	AA	503	A
85	AA	504	U
85	AA	505	U
85	AA	509	C
85	AA	519	A
85	AA	520	A
85	AA	521	A
85	AA	523	U
85	AA	529	G
85	AA	575	G
85	AA	589	A
85	AA	590	U
85	AA	591	A
85	AA	603	C
85	AA	604	C
85	AA	606	A
85	AA	617	C
85	AA	619	A
85	AA	628	C
85	AA	632	U
85	AA	639	C
85	AA	652	U
85	AA	653	A
85	AA	656	U
85	AA	668	A
85	AA	682	C
85	AA	687	G
85	AA	696	G
85	AA	708	G
85	AA	711	C

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Mol	Chain	Res	Type
85	AA	714	U
85	AA	725	G
85	AA	727	U
85	AA	735	G
85	AA	743	C
85	AA	752	C
85	AA	753	U
85	AA	759	G
85	AA	761	G
85	AA	764	U
85	AA	765	U
85	AA	775	C
85	AA	785	C
85	AA	789	A
85	AA	791	C
85	AA	808	A
85	AA	809	A
85	AA	824	C
85	AA	829	C
85	AA	830	A
85	AA	837	C
85	AA	845	A
85	AA	846	U
85	AA	852	C
85	AA	854	A
85	AA	860	C
85	AA	861	G
85	AA	867	G
85	AA	873	U
85	AA	882	C
85	AA	884	A
85	AA	886	A
85	AA	888	A
85	AA	891	G
85	AA	895	C
85	AA	900	G
85	AA	901	C
85	AA	906	U
85	AA	907	G
85	AA	909	C
85	AA	923	A
85	AA	926	C

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Mol	Chain	Res	Type
85	AA	930	G
85	AA	936	C
85	AA	940	G
85	AA	941	C
85	AA	942	A
85	AA	960	G
85	AA	964	C
85	AA	990	U
85	AA	998	U
85	AA	1009	G
85	AA	1012	C
85	AA	1016	G
85	AA	1022	G
85	AA	1032	U
85	AA	1035	C
85	AA	1051	A
85	AA	1053	A
85	AA	1057	G
85	AA	1061	C
85	AA	1062	U
85	AA	1067	G
85	AA	1074	U
85	AA	1076	U
85	AA	1090	A
85	AA	1103	A
85	AA	1105	G
85	AA	1106	A
85	AA	1110	A
85	AA	1111	A
85	AA	1112	G
85	AA	1147	A
85	AA	1153	G
85	AA	1158	U
85	AA	1180	C
85	AA	1182	A
85	AA	1183	C
85	AA	1208	C
85	AA	1212	C
85	AA	1213	U
85	AA	1237	A
85	AA	1252	A
85	AA	1264	U

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Mol	Chain	Res	Type
85	AA	1275	A
85	AA	1277	C
85	AA	1279	A
85	AA	1280	U
85	AA	1291	A
85	AA	1301	C
85	AA	1302	A
85	AA	1310	G
85	AA	1314	C
85	AA	1329	U
85	AA	1351	U
85	AA	1353	U
85	AA	1357	U
85	AA	1358	A
85	AA	1371	C
85	AA	1378	U
85	AA	1396	C
85	AA	1399	U
85	AA	1416	U
85	AA	1421	U
85	AA	1424	G
85	AA	1449	C
85	AA	1468	G
85	AA	1469	G
85	AA	1470	A
85	AA	1473	U
85	AA	1474	U
85	AA	1475	A
85	AA	1485	G
85	AA	1490	A
85	AA	1491	G
85	AA	1493	A
85	AA	1494	C
85	AA	1514	A
85	AA	1535	C
85	AA	1560	A
85	AA	1562	U
85	AA	1570	A
85	AA	1576	G
85	AA	1577	G
85	AA	1578	G
85	AA	1579	A

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Mol	Chain	Res	Type
85	AA	1584	U
85	AA	1593	C
85	AA	1604	A
85	AA	1605	G
85	AA	1622	G
85	AA	1632	G
85	AA	1633	A
85	AA	1646	U
85	AA	1650	G
85	AA	1661	U
85	AA	1662	U
85	AA	1664	G
85	AA	1682	U
85	AA	1683	U
85	AA	1690	A
85	AA	1691	U
85	AA	1698	A
85	AA	1716	U
85	AA	1718	C
85	AA	1725	G
85	AA	1728	G
85	AA	1730	C
85	AA	1735	U
85	AA	1751	G
85	AA	1752	C
85	AA	1753	A
85	AA	1784	G
85	AA	1787	G
85	AA	1791	U
85	AA	1803	U
85	AA	1807	A
85	AA	1810	C
85	AA	1825	A
85	AA	1835	U
85	AA	1852	U
85	AA	1853	U
85	AA	1860	A
85	AA	1874	G
85	AA	1884	A
85	AA	1886	U
85	AA	1896	G
85	AA	1898	C

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Mol	Chain	Res	Type
85	AA	1911	A
85	AA	1912	U
85	AA	1920	A
85	AA	1923	A
85	AA	1928	A
85	AA	1930	U
85	AA	1946	C
85	AA	1964	A
85	AA	1973	G
85	AA	1978	G
85	AA	1980	A
85	AA	1982	C
85	AA	1989	A
85	AA	2000	C
85	AA	2003	C
85	AA	2004	U
85	AA	2005	U
85	AA	2024	U
85	AA	2040	A
85	AA	2051	G
85	AA	2062	U
85	AA	2068	A
85	AA	2082	C
85	AA	2085	C
85	AA	2098	A
85	AA	2100	A
85	AA	2103	C
85	AA	2104	C
85	AA	2125	A
85	AA	2132	A
85	AA	2215	C
85	AA	2216	A
85	AA	2218	G
85	AA	2219	G
85	AA	2242	U
85	AA	2244	G
85	AA	2245	A
85	AA	2249	U
86	AB	3	C
86	AB	8	U
86	AB	20	U
86	AB	54	U

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Mol	Chain	Res	Type
86	AB	56	C
86	AB	65	G
86	AB	70	G

5.4 Non-standard residues in protein, DNA, RNA chains [i](#)

There are no non-standard protein/DNA/RNA residues in this entry.

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
34	BA	3
40	BG	2

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	BA	546:U	O3'	547:C	P	2.11
1	BA	557:U	O3'	558:C	P	1.87
1	BA	547:C	O3'	548:G	P	1.78
1	BG	24:A	O3'	25:G	P	1.39
1	BG	9:G	O3'	10:U	P	1.37

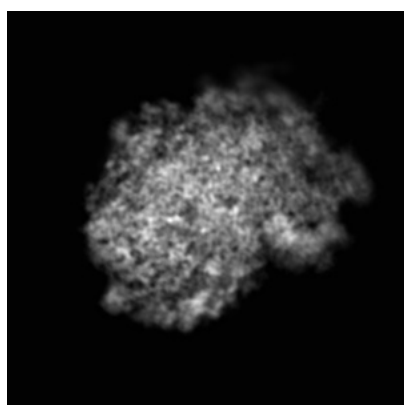
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-2239. These allow visual inspection of the internal detail of the map and identification of artifacts.

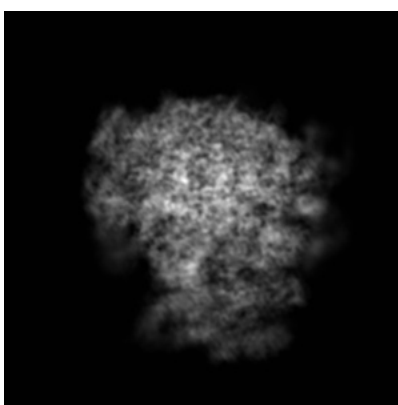
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

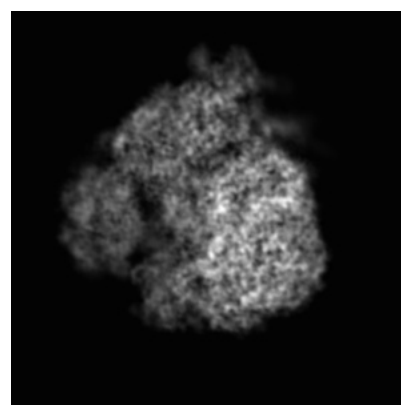
6.1.1 Primary map



X



Y

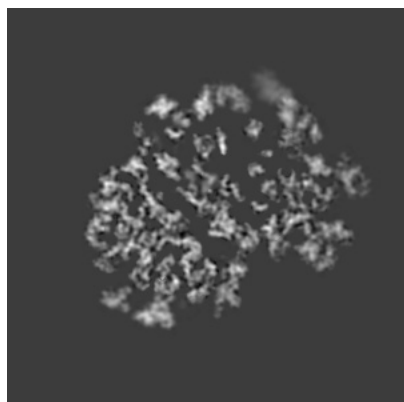


Z

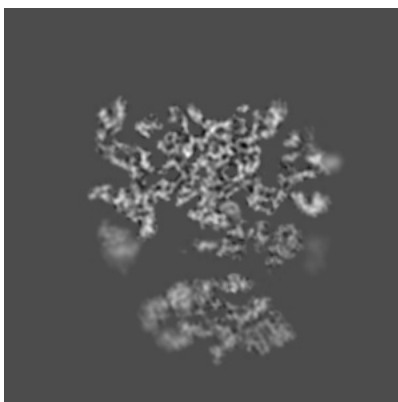
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

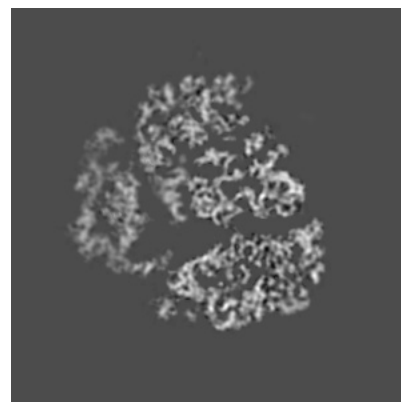
6.2.1 Primary map



X Index: 179



Y Index: 179

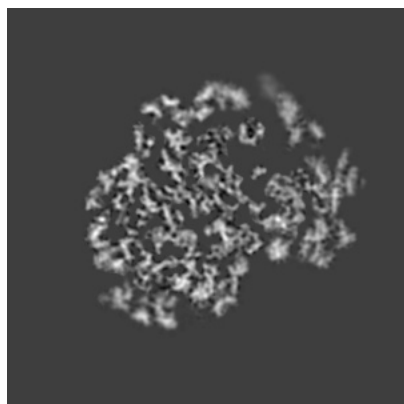


Z Index: 179

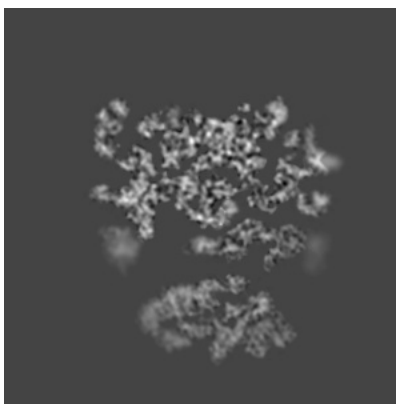
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

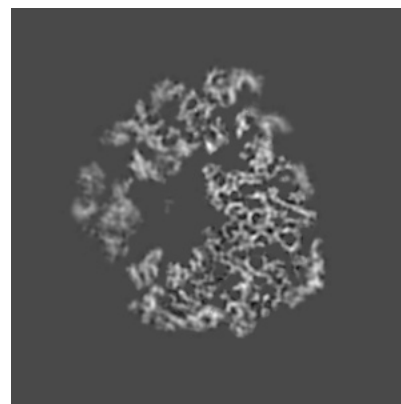
6.3.1 Primary map



X Index: 184



Y Index: 182

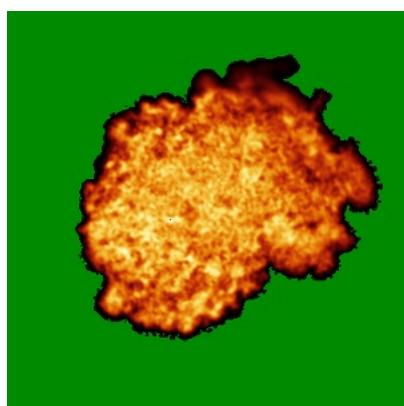


Z Index: 161

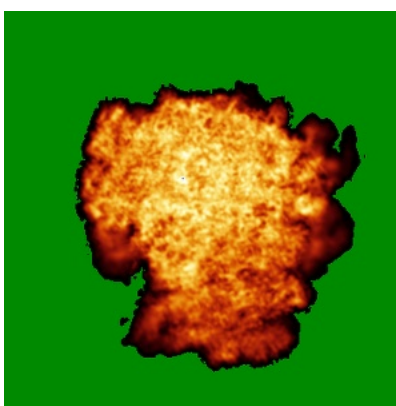
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) [i](#)

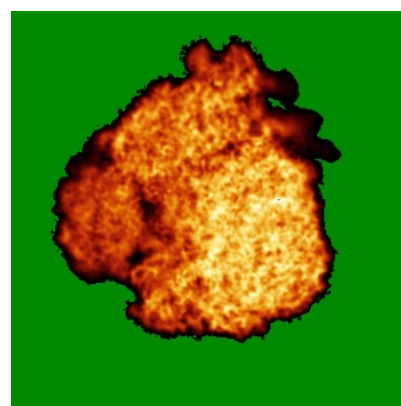
6.4.1 Primary map



X



Y

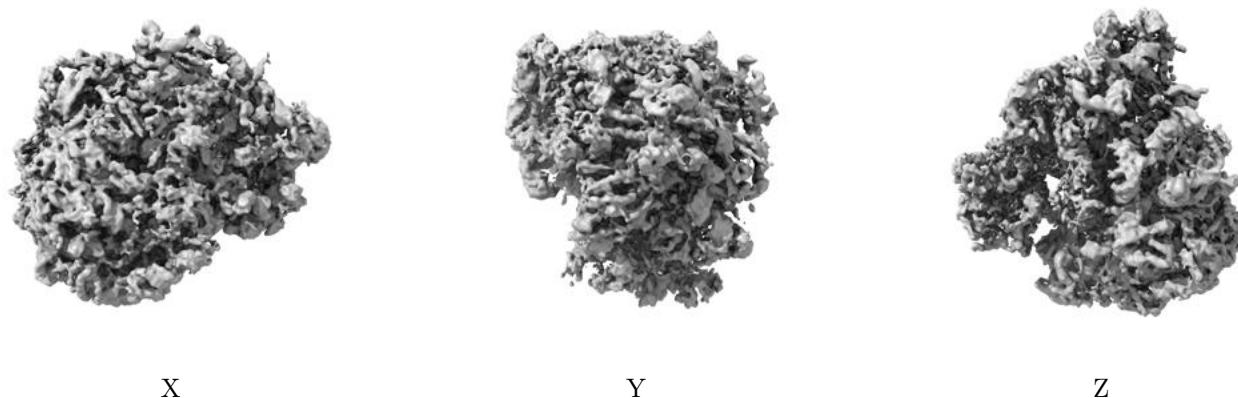


Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 108000.0. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

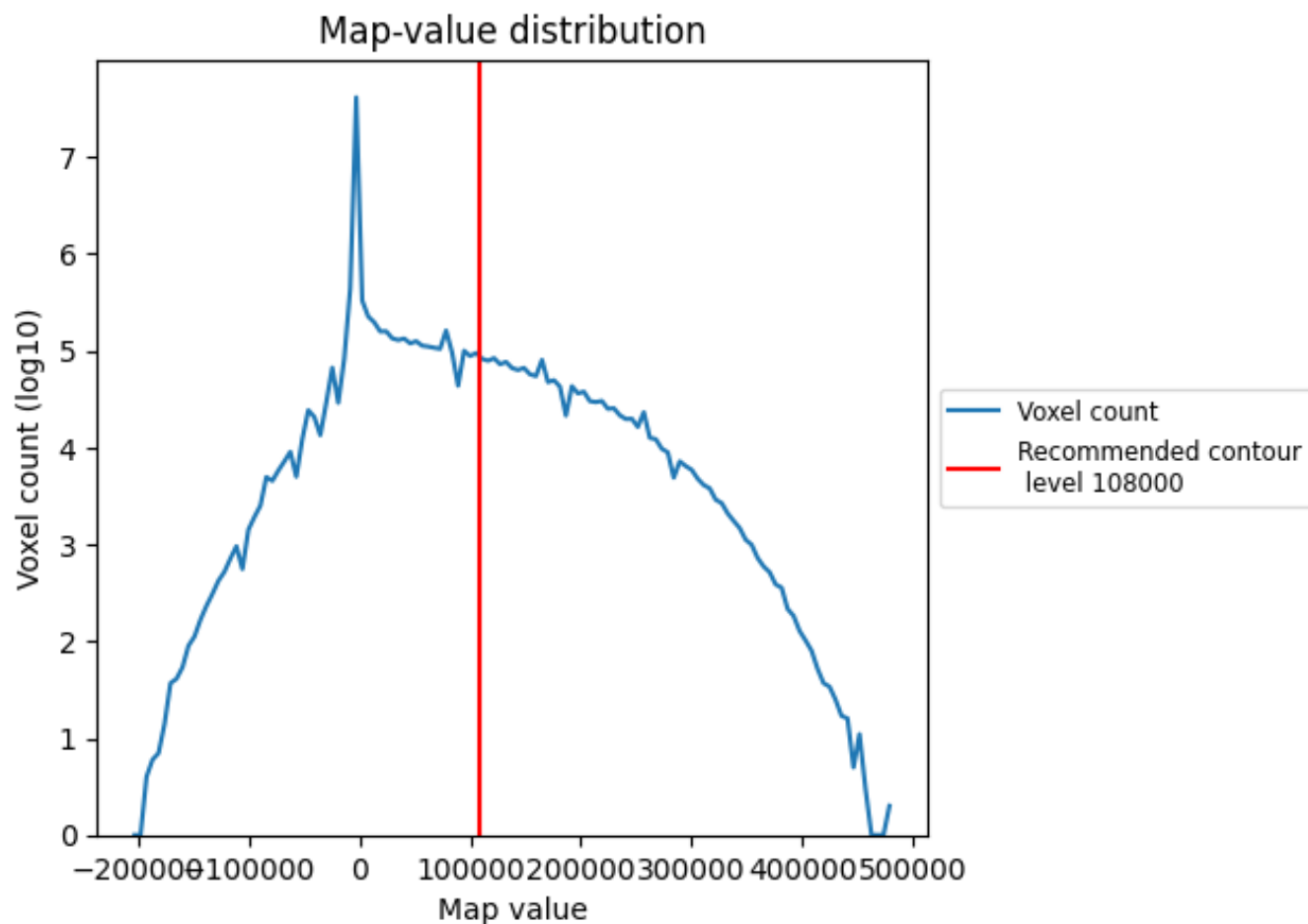
6.6 Mask visualisation [i](#)

This section was not generated. No masks/segmentation were deposited.

7 Map analysis [i](#)

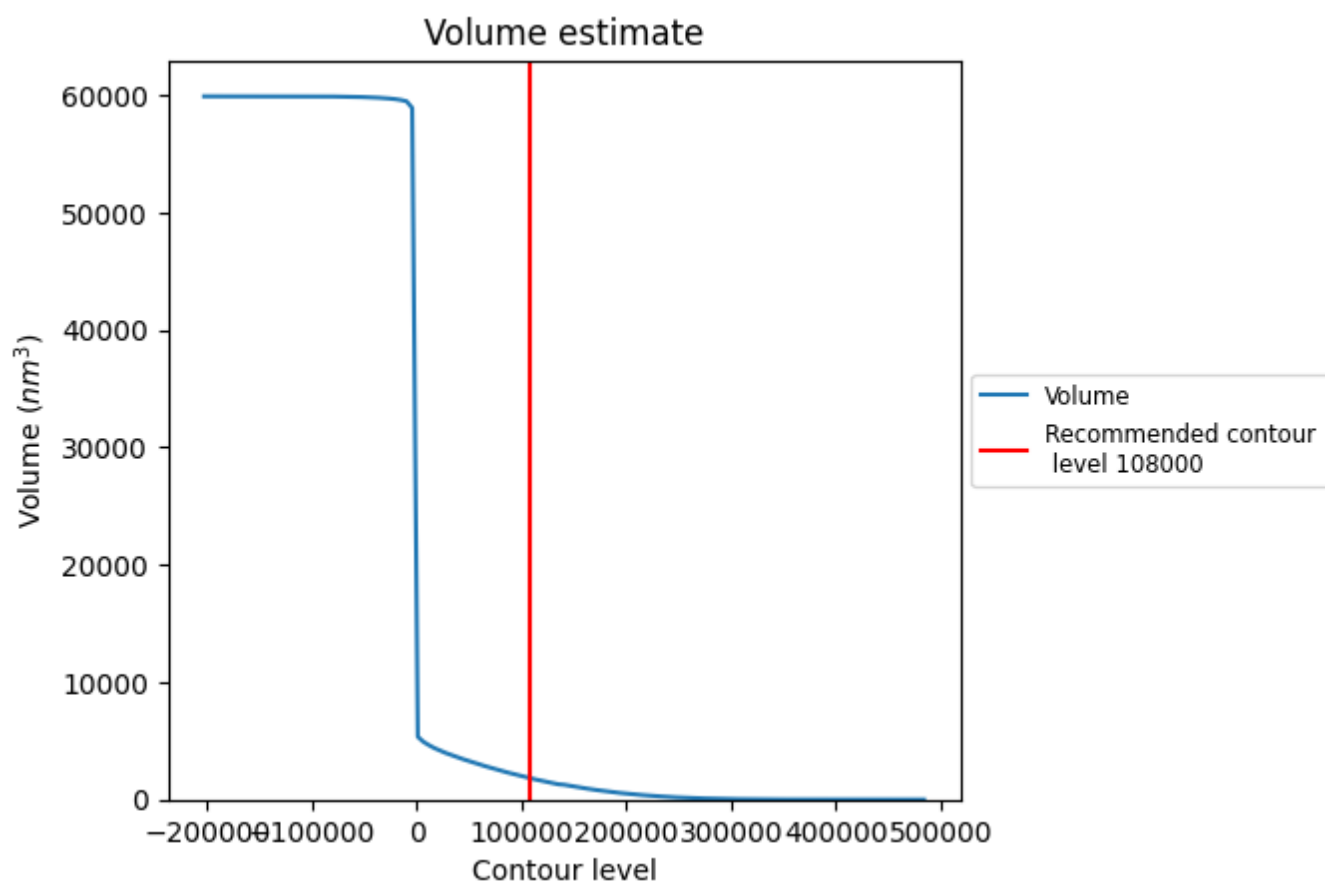
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

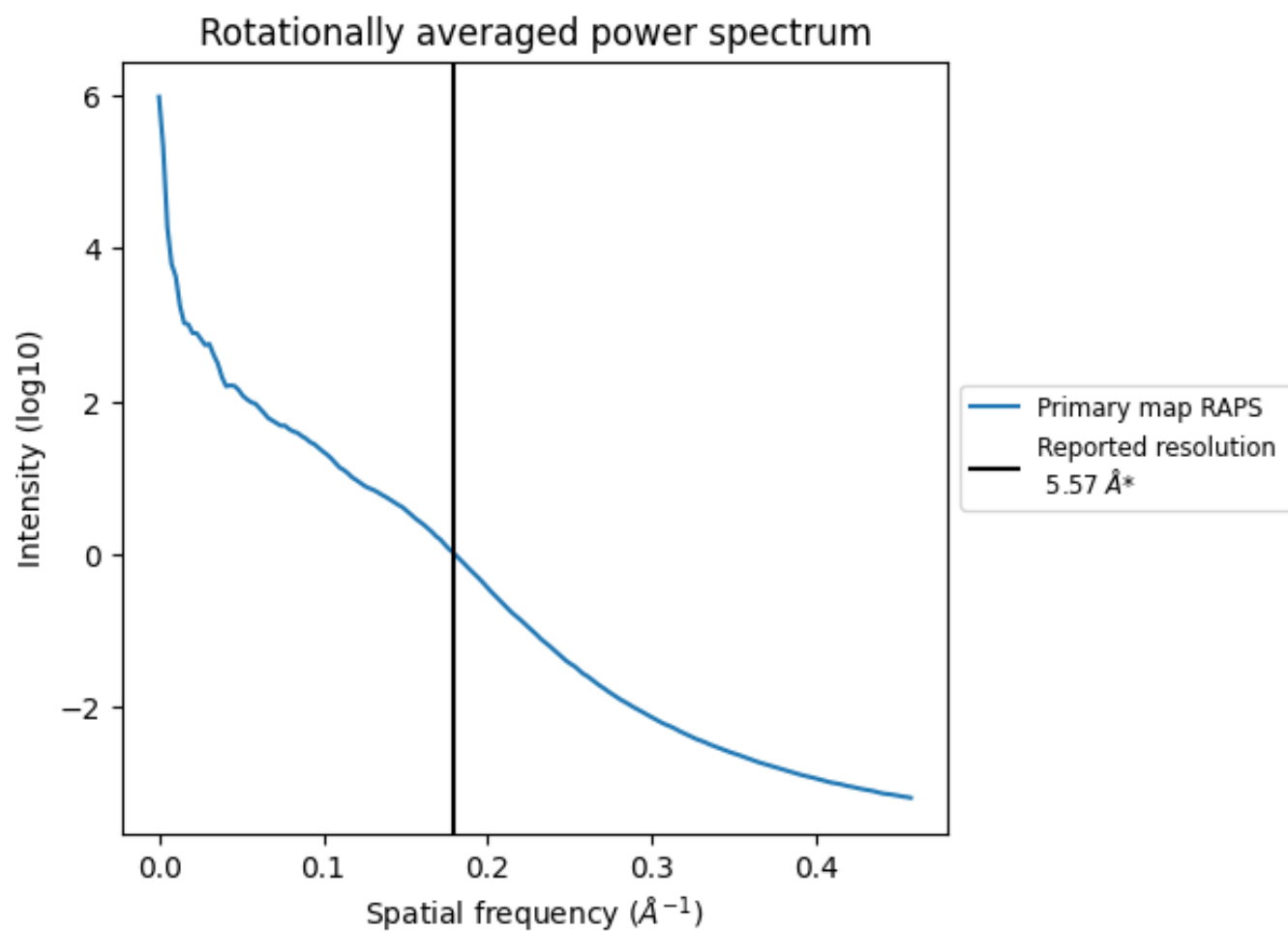
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1820 nm³; this corresponds to an approximate mass of 1644 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.

7.3 Rotationally averaged power spectrum ⓘ

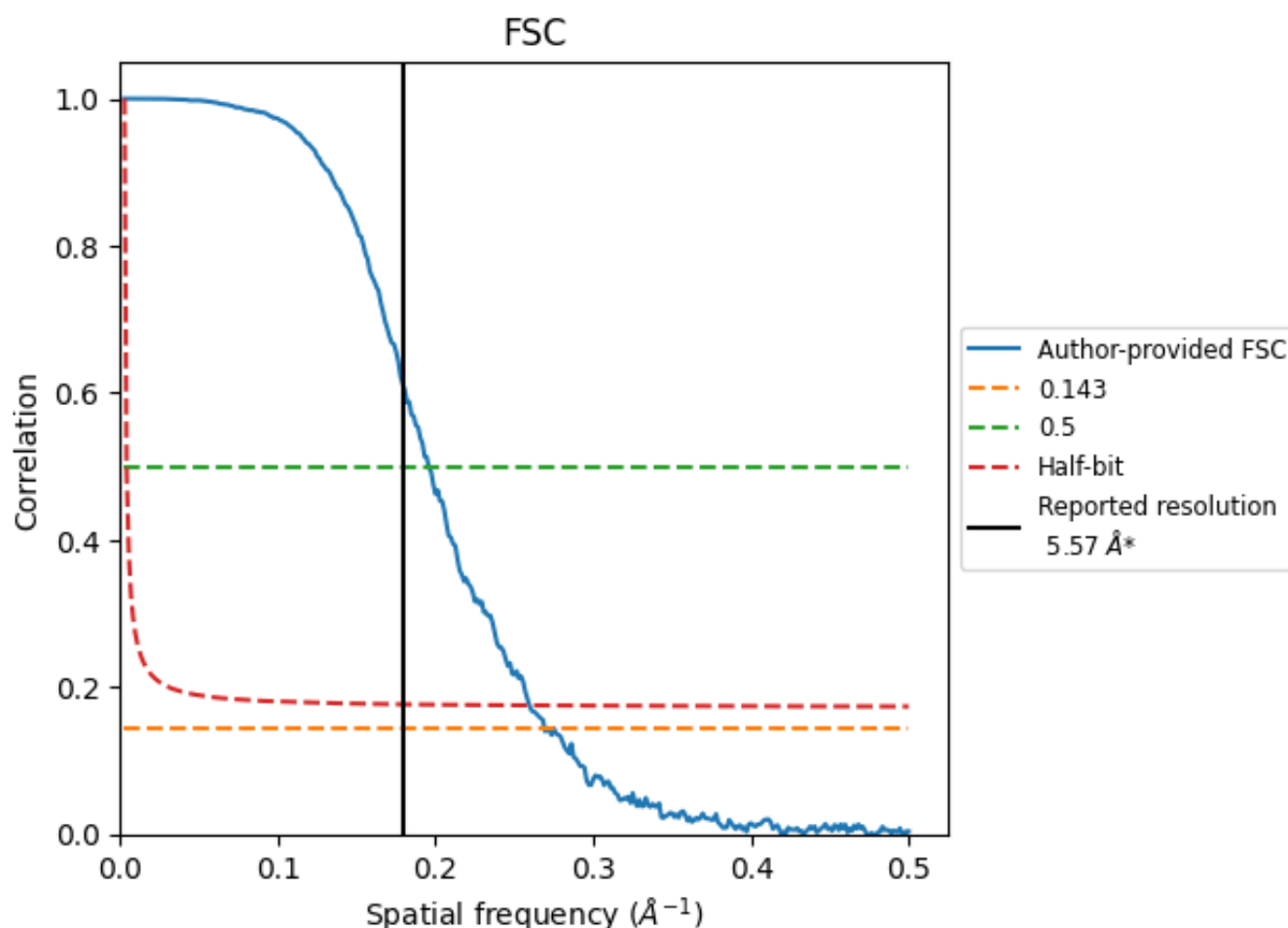


*Reported resolution corresponds to spatial frequency of 0.180 \AA^{-1}

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.180 Å⁻¹

8.2 Resolution estimates [i](#)

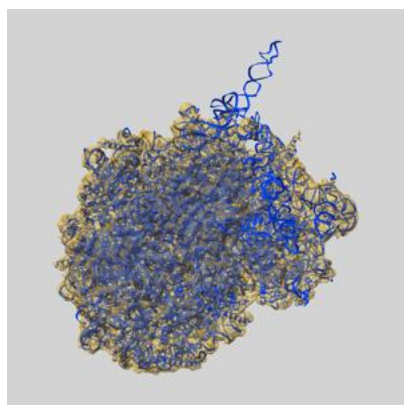
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	-	-	-
Author-provided FSC curve	3.72	5.11	3.85
Unmasked-calculated*	-	-	-

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps.

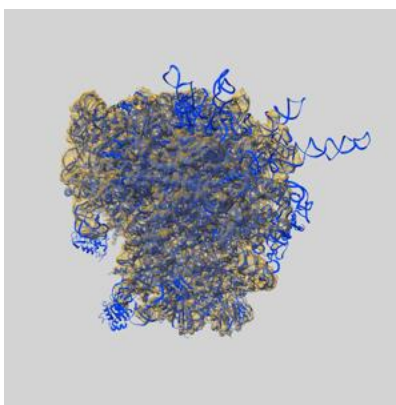
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-2239 and PDB model 4V8M. Per-residue inclusion information can be found in section 3 on page 21.

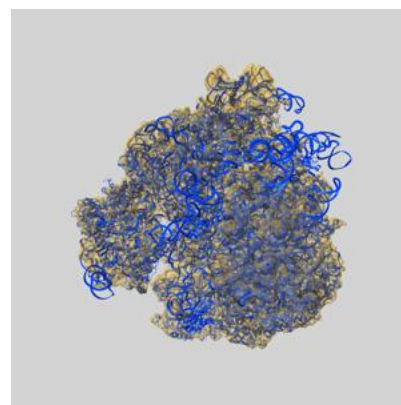
9.1 Map-model overlay [i](#)



X



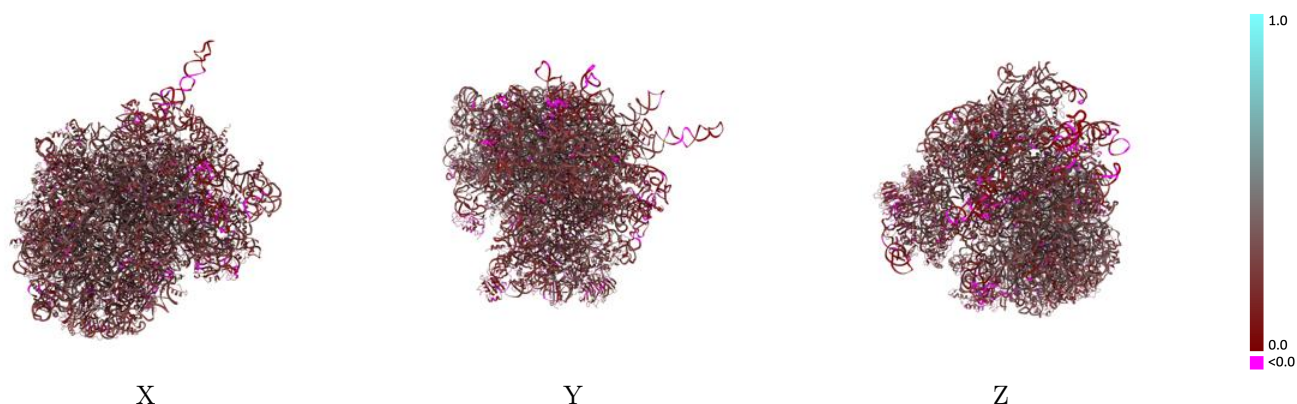
Y



Z

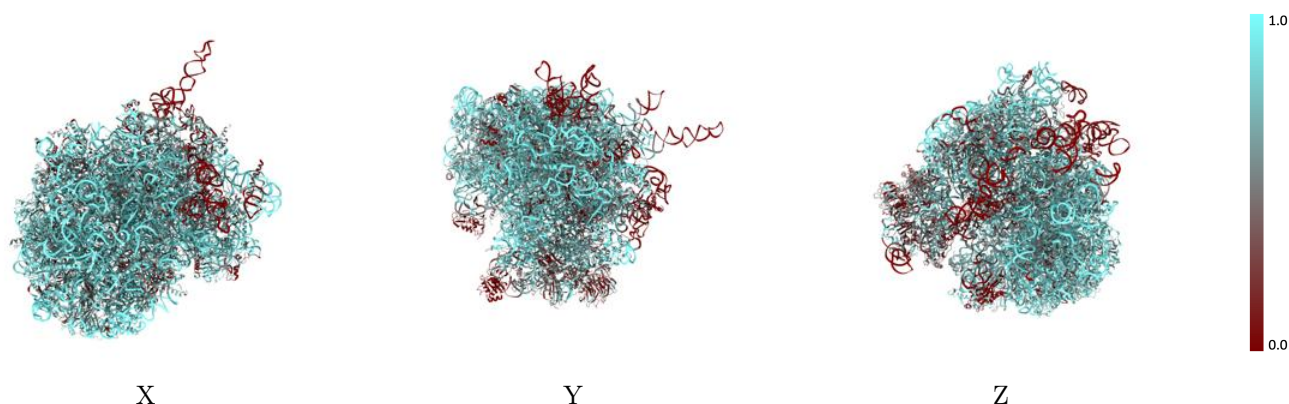
The images above show the 3D surface view of the map at the recommended contour level 108000.0 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

9.2 Q-score mapped to coordinate model [i](#)



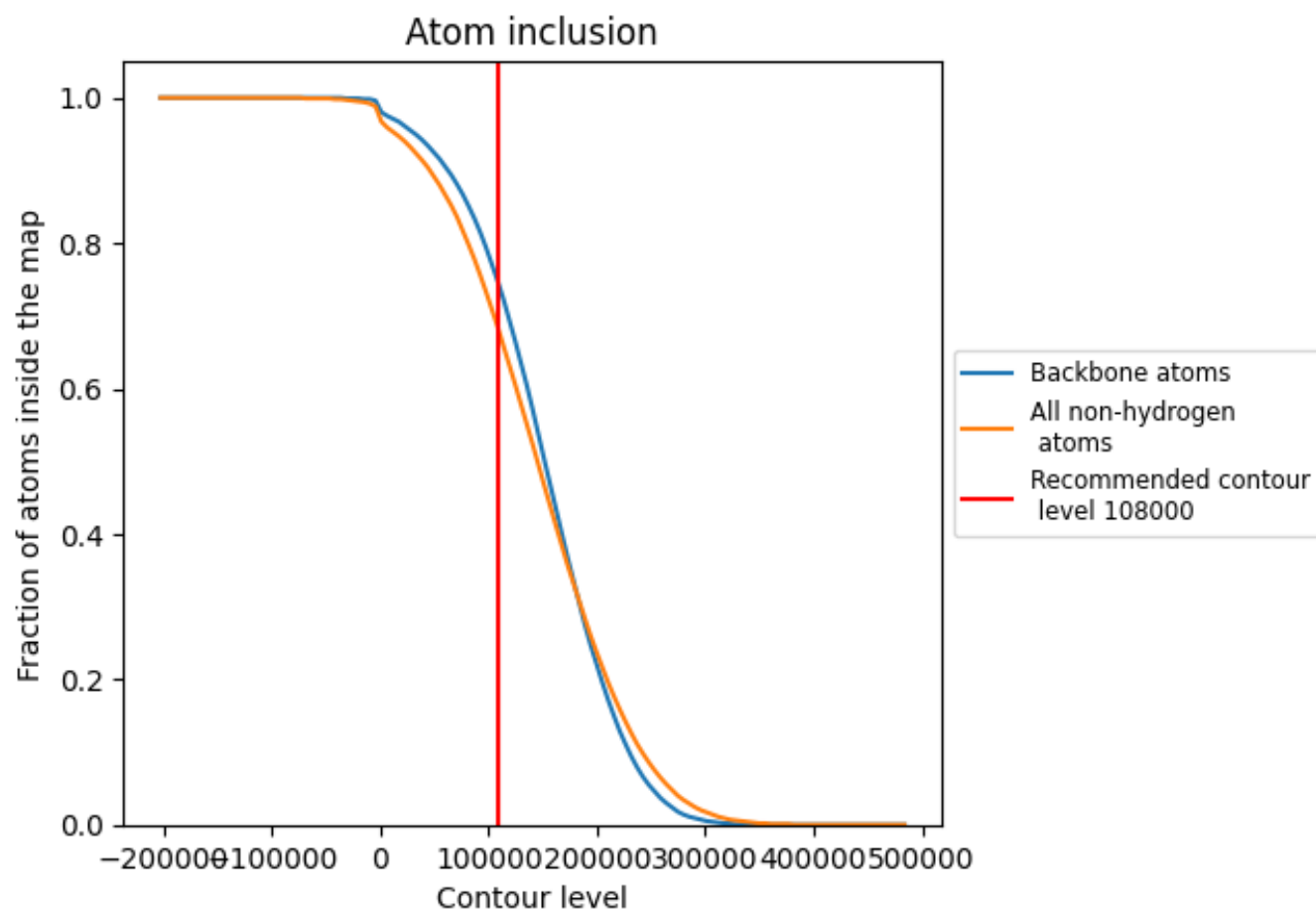
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions at which atoms can be resolved. Low Q-score values may therefore be expected for many entries.

9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (108000).







































































9.4 Atom inclusion [i](#)



At the recommended contour level, 75% of all backbone atoms, 69% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ





















































































The table lists the average atom inclusion at the recommended contour level (108000) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.6860	 0.2390
A0	 0.5170	 0.2360
A1	 0.5980	 0.2390
A2	 0.3430	 0.1910
A3	 0.6080	 0.2070
A4	 0.6130	 0.2320
A5	 0.4740	 0.2180
A6	 0.6720	 0.2270
A7	 0.3000	 0.1870
A8	 0.1980	 0.1170
A9	 0.1860	 0.1310
AA	 0.7340	 0.2390
AB	 0.4690	 0.1630
AC	 0.6380	 0.2440
AD	 0.3260	 0.1670
AE	 0.3730	 0.2080
AF	 0.0100	 0.1110
AG	 0.5170	 0.2330
AH	 0.3730	 0.1990
AI	 0.2750	 0.1480
AJ	 0.4700	 0.2300
AK	 0.3240	 0.2000
AL	 0.5290	 0.2220
AM	 0.2880	 0.1420
AO	 0.4990	 0.1980
AP	 0.6320	 0.2360
AQ	 0.3770	 0.1720
AR	 0.6530	 0.2550
AS	 0.4020	 0.1980
AT	 0.6470	 0.2000
AU	 0.3200	 0.1280
AV	 0.2260	 0.2090
AW	 0.6460	 0.2790
AX	 0.3660	 0.1860
AY	 0.5460	 0.1940























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Chain	Atom inclusion	Q-score
AZ	 0.4490	 0.2050
BA	 0.8610	 0.2760
BB	 0.7680	 0.2580
BC	 0.9190	 0.3090
BD	 0.9510	 0.2740
BE	 0.8330	 0.2680
BF	 0.8680	 0.2620
BG	 0.9360	 0.2990
BH	 0.7890	 0.2190
BI	 0.5580	 0.2150
BJ	 0.0020	 0.0520
BK	 0.6170	 0.2440
BL	 0.7250	 0.2310
BM	 0.0060	 0.0470
BN	 0.6690	 0.2310
BO	 0.5960	 0.2140
BP	 0.7160	 0.2300
BQ	 0.5560	 0.2370
BR	 0.5830	 0.2270
BS	 0.5430	 0.2280
BT	 0.6370	 0.2260
BU	 0.5080	 0.2240
BV	 0.0750	 0.0940
BW	 0.6440	 0.2610
BX	 0.6060	 0.2500
BY	 0.3600	 0.1320
BZ	 0.6920	 0.2300
Ba	 0.5800	 0.2450
Bb	 0.5890	 0.2360
Bc	 0.7570	 0.2500
Bd	 0.5270	 0.1590
Be	 0.4990	 0.2320
Bf	 0.5470	 0.2240
Bg	 0.7180	 0.2600
Bh	 0.5850	 0.2040
Bi	 0.4600	 0.2010
Bj	 0.4200	 0.1440
Bk	 0.6270	 0.2390
Bl	 0.5120	 0.2040
Bm	 0.6440	 0.2360
Bn	 0.6380	 0.2130
Bo	 0.5410	 0.2050

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Chain	Atom inclusion	Q-score
Bp	 0.7110	 0.2480
Bq	 0.5520	 0.2320
Br	 0.6450	 0.2410
Bs	 0.6410	 0.2440
Bt	 0.6610	 0.2570
Bu	 0.6820	 0.2240
Bv	 0.6580	 0.2470
Bw	 0.5990	 0.2310
Bx	 0.6090	 0.2340
By	 0.7460	 0.2370