



## Full wwPDB EM Validation Report ⓘ

Jul 15, 2025 – 03:22 PM EDT

PDB ID : 8VOO / pdb\_00008voo  
EMDB ID : EMD-43387  
Title : Escherichia coli transcription-translation loosely coupled complex (TTC-LC) containing mRNA with a 39 nt long spacer, ops signal, RfaH, NusA, and fMet-tRNAs in E-site and P-site  
Authors : Molodtsov, V.; Wang, C.; Ebright, R.H.  
Deposited on : 2024-01-15  
Resolution : 3.40 Å(reported)

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

We welcome your comments at [validation@mail.wwpdb.org](mailto: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>.

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

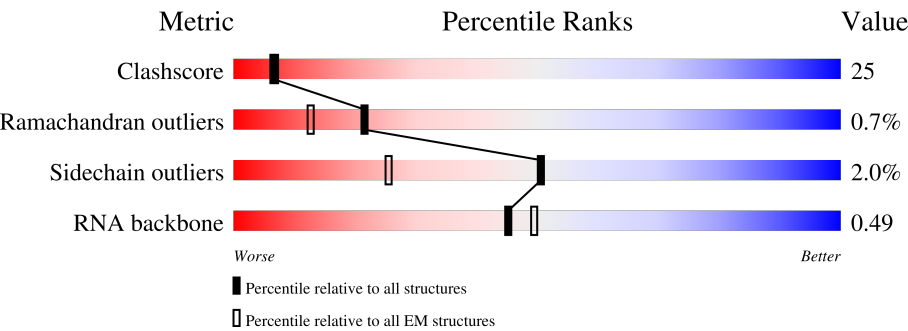
EMDB validation analysis : 0.0.1.dev118  
MolProbity : 4-5-2 with Phenix2.0rc1  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
MapQ : 1.9.13  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.44

# 1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:  
*ELECTRON MICROSCOPY*

The reported resolution of this entry is 3.40 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)
Clashscore	210492	15764
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  $\geq 3$ , 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	0	103	<div><div>9%</div><div>60%</div><div>40%</div></div>
2	1	110	<div><div>58%</div><div>42%</div></div>
3	2	100	<div><div>6%</div><div>51%</div><div>42%</div><div>6%</div></div>
4	3	104	<div><div>12%</div><div>66%</div><div>33%</div></div>
5	4	94	<div><div>10%</div><div>44%</div><div>56%</div></div>
6	5	38	<div><div>50%</div><div>55%</div><div>37%</div><div>8%</div></div>
7	6	38	<div><div>42%</div><div>63%</div><div>29%</div><div>8%</div></div>

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Mol	Chain	Length	Quality of chain
8	7	56	
9	9	165	
10	A	76	
10	B	76	
11	AA	1342	
12	AB	162	
13	AC	329	
13	AD	329	
14	AE	1407	
15	AF	91	
16	AG	495	
17	C	75	
18	D	1542	
19	E	87	
20	F	71	
21	G	241	
22	H	557	
23	I	233	
24	J	206	
25	K	167	
26	L	135	
27	M	179	
28	N	130	
29	O	130	
30	P	103	

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Mol	Chain	Length	Quality of chain
31	Q	129	
32	R	124	
33	S	101	
34	T	89	
35	U	82	
36	V	84	
37	W	92	
38	X	118	
39	Y	142	
40	Z	121	
41	a	2904	
42	b	85	
43	c	78	
44	d	120	
45	e	63	
46	f	59	
47	g	70	
48	h	273	
49	i	57	
50	j	209	
51	k	55	
52	l	201	
53	m	46	
54	n	179	
55	o	65	

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Mol	Chain	Length	Quality of chain
56	p	177	
57	q	38	
58	r	149	
59	s	142	
60	t	123	
61	u	144	
62	v	136	
63	w	127	
64	x	117	
65	y	115	
66	z	118	

The following table lists non-polymeric compounds, carbohydrate monomers and non-standard residues in protein, DNA, RNA chains that are outliers for geometric or electron-density-fit criteria:

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
68	ZN	AE	1502	-	-	X	-

## 2 Entry composition [i](#)

There are 68 unique types of molecules in this entry. The entry contains 280046 atoms, of which 98638 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 Ribosomal protein L21.

Mol	Chain	Residues	Atoms						AltConf	Trace
1	0	103	Total	C	H	N	O	S	0	0
			1655	516	839	153	145	2		

- Molecule 2 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms						AltConf	Trace
2	1	110	Total	C	H	N	O	S	0	0
			1779	532	922	166	156	3		

- Molecule 3 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms						AltConf	Trace
3	2	94	Total	C	H	N	O	S	0	0
			1557	470	811	140	134	2		

- Molecule 4 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms						AltConf	Trace
4	3	103	Total	C	H	N	O		0	0
			1632	498	844	148	142			

- Molecule 5 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms						AltConf	Trace
5	4	94	Total	C	H	N	O	S	0	0
			1533	479	780	137	134	3		

- Molecule 6 is a DNA chain called NT DNA ops.

Mol	Chain	Residues	Atoms						AltConf	Trace
6	5	35	Total	C	N	O	P		0	0
			726	342	141	208	35			

- Molecule 7 is a DNA chain called T DNA ops.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	6	35	Total	C	N	O	P	0	0
			703	336	117	215	35		

- Molecule 8 is a RNA chain called mRNA with 39 nt long spacer.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	7	25	Total	C	N	O	P	0	0
			527	235	80	187	25		

- Molecule 9 is a protein called 50S ribosomal protein L10.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	9	148	Total	C	N	O	S	0	0
			1117	705	196	209	7		

- Molecule 10 is a RNA chain called E-site and P-site tRNA (fMet).

Mol	Chain	Residues	Atoms						AltConf	Trace
10	A	76	Total 2446	C 723	H 826	N 295	O 527	P 75	0	0
10	B	76	Total 2433	C 723	H 813	N 295	O 527	P 75	0	0

- Molecule 11 is a protein called DNA-directed RNA polymerase subunit beta.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	AA	1316	Total	C	N	O	S	0	0
			10381	6514	1810	2014	43		

- Molecule 12 is a protein called Transcription antitermination protein RfaH.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	AB	161	Total	C	N	O	S	0	0
			1286	828	222	232	4		

- Molecule 13 is a protein called DNA-directed RNA polymerase subunit alpha.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	AC	221	Total	C	N	O	S	0	0
			1698	1060	299	333	6		

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Mol	Chain	Residues	Atoms					AltConf	Trace
13	AD	299	Total	C	N	O	S	0	0
			2078	1287	378	407	6		

- Molecule 14 is a protein called DNA-directed RNA polymerase subunit beta'.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	AE	1337	Total	C	N	O	S	0	0
			10404	6535	1856	1963	50		

- Molecule 15 is a protein called DNA-directed RNA polymerase subunit omega.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	AF	82	Total	C	N	O	S	0	0
			650	396	122	131	1		

- Molecule 16 is a protein called Transcription termination/antitermination protein NusA.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	AG	495	Total	C	N	O	S	0	0
			3852	2396	669	774	13		

- Molecule 17 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	C	66	Total	C	H	N	O	S	0
			1103	344	559	102	97	1	0

- Molecule 18 is a RNA chain called 16S rRNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	D	1524	Total	C	H	N	O	P	0
			49126	14585	16423	6003	10591	1524	0

- Molecule 19 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	E	86	Total	C	H	N	O	S	0
			1388	414	719	138	114	3	0

- Molecule 20 is a protein called 30S ribosomal protein S21.



Mol	Chain	Residues	Atoms						AltConf	Trace
20	F	70	Total	C	H	N	O	S	0	0
			1218	366	629	125	97	1		

- Molecule 21 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms						AltConf	Trace
21	G	225	Total	C	H	N	O	S	0	0
			3545	1113	1785	316	323	8		

- Molecule 22 is a protein called 30S ribosomal protein S1.

Mol	Chain	Residues	Atoms						AltConf	Trace
22	H	259	Total	C	H	N	O	S	0	0
			3184	1073	1454	305	349	3		

- Molecule 23 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms						AltConf	Trace
23	I	208	Total	C	H	N	O	S	0	0
			3346	1036	1710	307	290	3		

- Molecule 24 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms						AltConf	Trace
24	J	205	Total	C	H	N	O	S	0	0
			3350	1026	1707	315	298	4		

- Molecule 25 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms						AltConf	Trace
25	K	156	Total	C	H	N	O	S	0	0
			2348	717	1196	217	212	6		

- Molecule 26 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms						AltConf	Trace
26	L	104	Total	C	H	N	O	S	0	0
			1694	536	846	153	152	7		

- Molecule 27 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms						AltConf	Trace
27	M	151	Total	C	H	N	O	S	0	0
			2416	735	1235	227	215	4		

- Molecule 28 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms						AltConf	Trace
28	N	129	Total	C	H	N	O	S	0	0
			2010	616	1031	173	184	6		

- Molecule 29 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms						AltConf	Trace
29	O	127	Total	C	H	N	O	S	0	0
			2092	634	1070	206	179	3		

- Molecule 30 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms						AltConf	Trace
30	P	99	Total	C	H	N	O	S	0	0
			1621	495	831	151	143	1		

- Molecule 31 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms						AltConf	Trace
31	Q	117	Total	C	H	N	O	S	0	0
			1764	540	887	174	160	3		

- Molecule 32 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms						AltConf	Trace
32	R	121	Total	C	H	N	O	S	0	0
			1940	580	1001	194	161	4		

- Molecule 33 is a protein called 30S ribosomal protein S14.

Mol	Chain	Residues	Atoms						AltConf	Trace
33	S	100	Total	C	H	N	O	S	0	0
			1649	499	844	164	139	3		

- Molecule 34 is a protein called Small ribosomal subunit protein uS15.

Mol	Chain	Residues	Atoms						AltConf	Trace
34	T	88	Total	C	H	N	O	S	0	0
			1448	439	734	144	130	1		

- Molecule 35 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms						AltConf	Trace
35	U	82	Total	C	H	N	O	S	0	0
			1315	406	666	128	114	1		

- Molecule 36 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms						AltConf	Trace
36	V	80	Total	C	H	N	O	S	0	0
			1339	411	691	121	113	3		

- Molecule 37 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms						AltConf	Trace
37	W	83	Total	C	H	N	O	S	0	0
			1351	424	688	126	111	2		

- Molecule 38 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms						AltConf	Trace
38	X	116	Total	C	H	N	O	S	0	0
			1864	558	964	181	158	3		

- Molecule 39 is a protein called 50S ribosomal protein L11.

Mol	Chain	Residues	Atoms					AltConf	Trace
39	Y	141	Total	C	N	O	S	0	0
			1032	651	179	196	6		

- Molecule 40 is a protein called 50S ribosomal protein L7/L12.

Mol	Chain	Residues	Atoms					AltConf	Trace
40	Z	30	Total	C	N	O	S	0	0
			227	144	33	47	3		

- Molecule 41 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms						AltConf	Trace
41	a	2880	Total	C	H	N	O	P	0	0
			92918	27587	31077	11398	19976	2880		

There is a discrepancy between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
a	887	A	U	conflict	GB 937521852

- Molecule 42 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms						AltConf	Trace
42	b	76	Total	C	H	N	O	S	0	0
			1181	360	599	117	104	1		

- Molecule 43 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms						AltConf	Trace
43	c	77	Total	C	H	N	O	S	0	0
			1277	388	652	129	106	2		

- Molecule 44 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms						AltConf	Trace
44	d	120	Total	C	H	N	O	P	0	0
			3870	1144	1301	468	837	120		

- Molecule 45 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms						AltConf	Trace
45	e	62	Total	C	H	N	O	S	0	0
			1032	308	531	98	94	1		

- Molecule 46 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms						AltConf	Trace
46	f	58	Total	C	H	N	O	S	0	0
			936	281	488	87	78	2		

- Molecule 47 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms						AltConf	Trace
47	g	66	Total	C	H	N	O	S	0	0
			1042	323	520	99	94	6		

- Molecule 48 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms						AltConf	Trace
48	h	271	Total	C	H	N	O	S	0	0
			4236	1288	2154	423	364	7		

- Molecule 49 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms						AltConf	Trace
49	i	56	Total	C	H	N	O	S	0	0
			903	269	459	94	80	1		

- Molecule 50 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms						AltConf	Trace
50	j	209	Total	C	H	N	O	S	0	0
			3182	979	1617	288	294	4		

- Molecule 51 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms						AltConf	Trace
51	k	52	Total	C	H	N	O		0	0
			890	275	464	78	73			

- Molecule 52 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms						AltConf	Trace
52	l	201	Total	C	H	N	O	S	0	0
			3171	974	1619	283	290	5		

- Molecule 53 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms						AltConf	Trace
53	m	46	Total	C	H	N	O	S	0	0
			795	228	418	90	57	2		

- Molecule 54 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms						AltConf	Trace
54	n	177	Total	C	H	N	O	S	0	0
			2853	899	1443	249	256	6		

- Molecule 55 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms						AltConf	Trace
55	o	64	Total	C	H	N	O	S	0	0
			1076	323	572	105	74	2		

- Molecule 56 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms						AltConf	Trace
56	p	175	Total	C	H	N	O	S	0	0
			2671	826	1358	241	244	2		

- Molecule 57 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms						AltConf	Trace
57	q	38	Total	C	H	N	O	S	0	0
			645	185	343	65	48	4		

- Molecule 58 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms						AltConf	Trace
58	r	149	Total	C	H	N	O	S	0	0
			2259	699	1148	197	214	1		

- Molecule 59 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms						AltConf	Trace
59	s	142	Total	C	H	N	O	S	0	0
			2291	714	1162	212	199	4		

- Molecule 60 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms						AltConf	Trace
60	t	123	Total	C	H	N	O	S	0	0
			1969	593	1023	181	166	6		

- Molecule 61 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms						AltConf	Trace
61	u	144	Total	C	H	N	O	S	0	0
			2182	654	1129	207	190	2		

- Molecule 62 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms						AltConf	Trace
62	v	136	Total	C	H	N	O	S	0	0
			2231	686	1157	205	177	6		

- Molecule 63 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms						AltConf	Trace
63	w	119	Total	C	H	N	O	S	0	0
			1945	588	994	195	163	5		

- Molecule 64 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms					AltConf	Trace
64	x	116	Total	C	H	N	O	0	0
			1815	552	923	178	162		

- Molecule 65 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms						AltConf	Trace
65	y	114	Total	C	H	N	O	S	0	0
			1879	574	962	179	163	1		

- Molecule 66 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					AltConf	Trace
66	z	117	Total	C	H	N	O	0	0
			1967	604	1020	192	151		

- Molecule 67 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		AltConf
67	AE	1	Total	Mg	0
			1	1	

- Molecule 68 is ZINC ION (CCD ID: ZN) (formula: Zn).

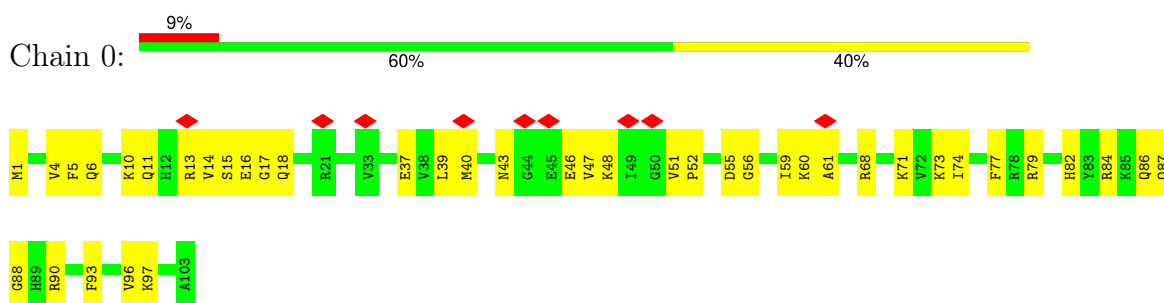
Mol	Chain	Residues	Atoms		AltConf
68	AE	2	Total	Zn	0
			2	2	



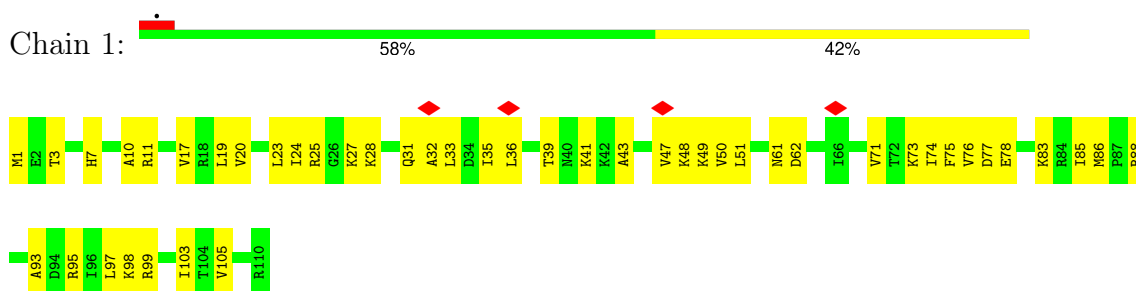
### 3 Residue-property plots [i](#)

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.

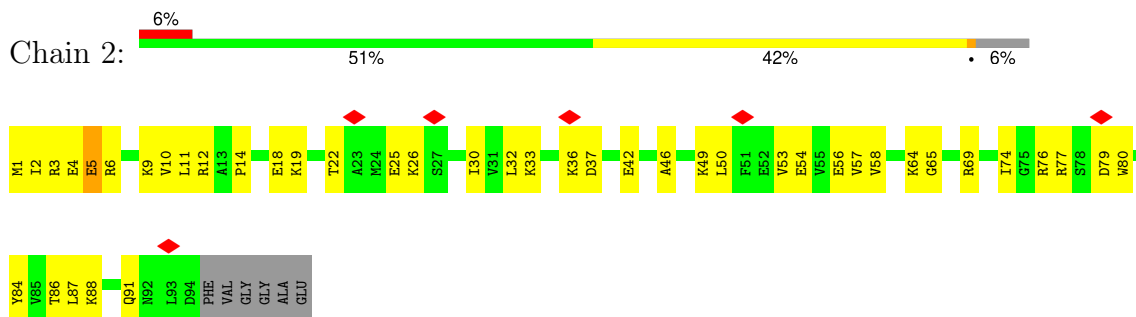
#### • Molecule 1: Ribosomal protein L21



#### • Molecule 2: 50S ribosomal protein L22



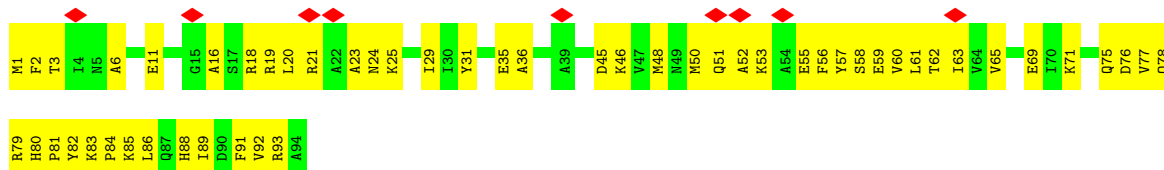
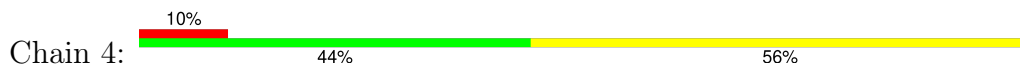
#### • Molecule 3: 50S ribosomal protein L23



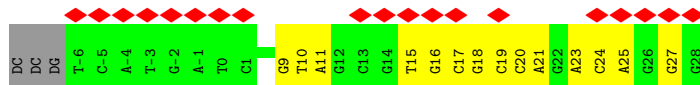
#### • Molecule 4: 50S ribosomal protein L24



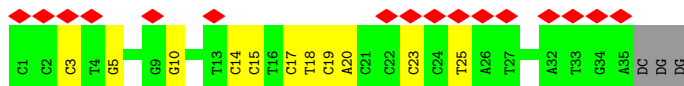
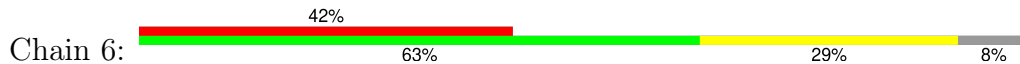
- Molecule 5: 50S ribosomal protein L25



- Molecule 6: NT DNA ops



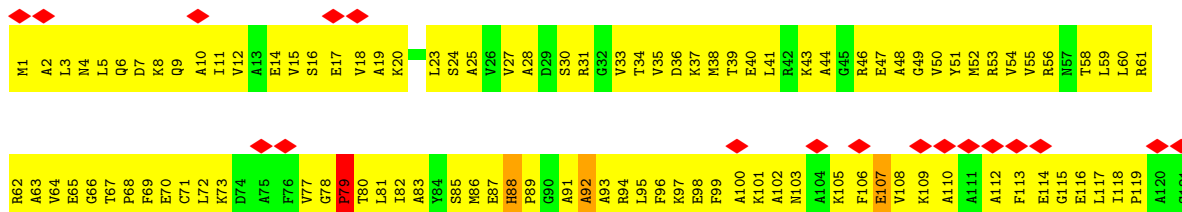
- Molecule 7: T DNA ops



- Molecule 8: mRNA with 39 nt long spacer

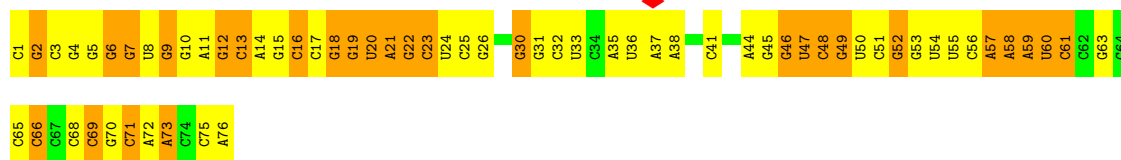
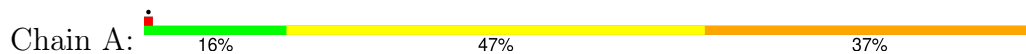


- Molecule 9: 50S ribosomal protein L10

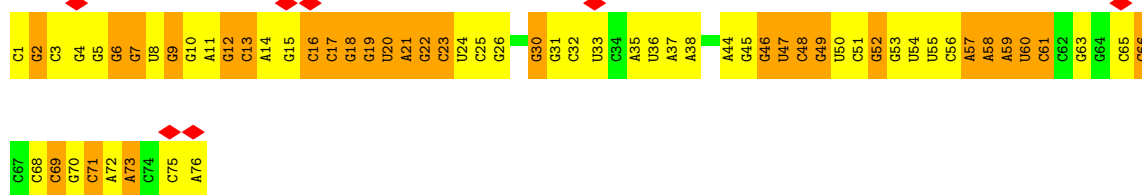
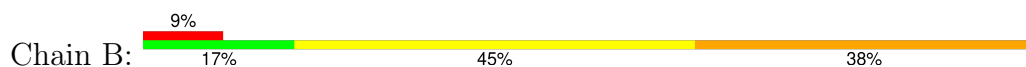




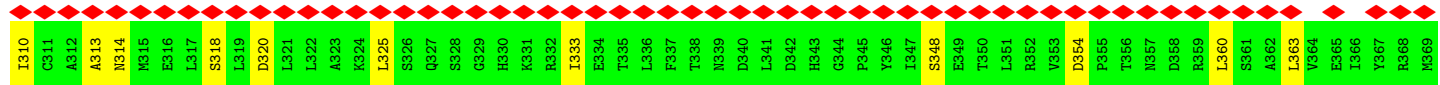
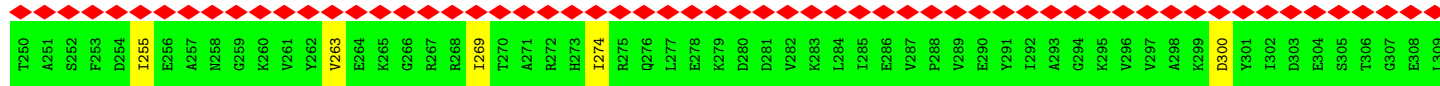
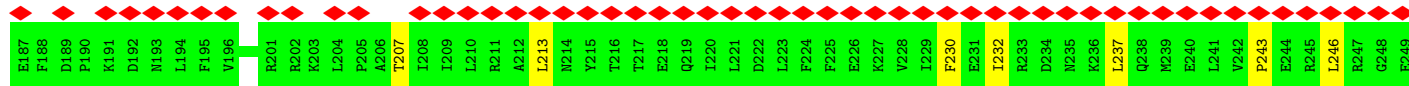
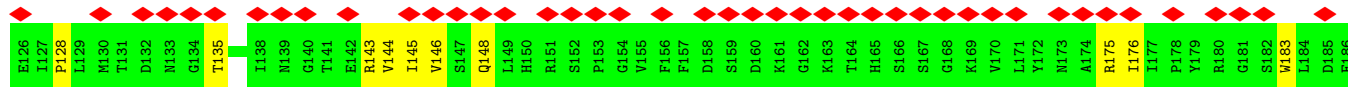
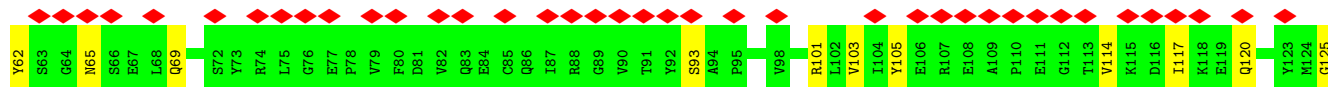
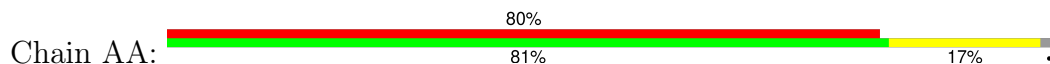
• Molecule 10: E-site and P-site tRNA (fMet)



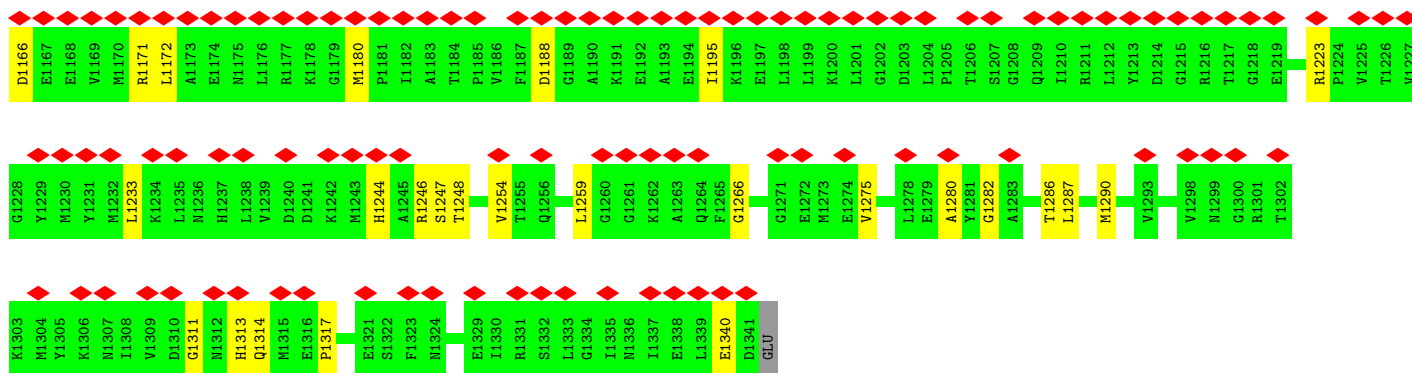
• Molecule 10: E-site and P-site tRNA (fMet)



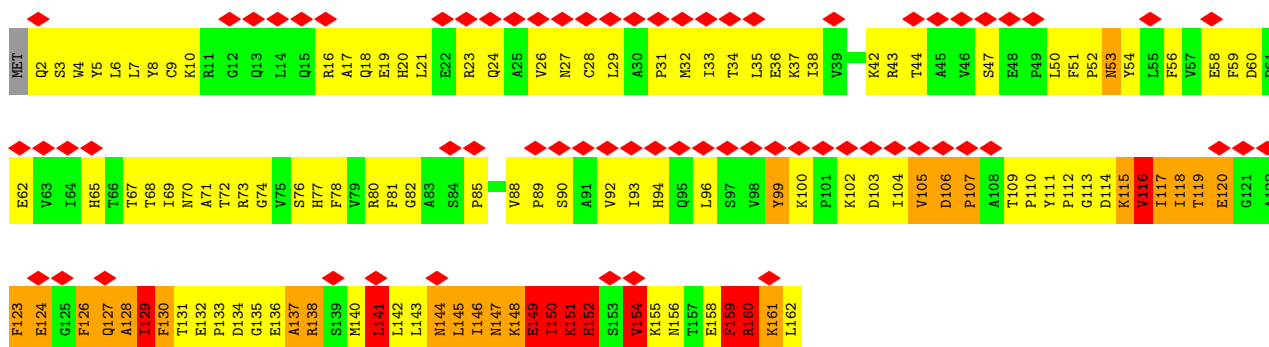
• Molecule 11: DNA-directed RNA polymerase subunit beta



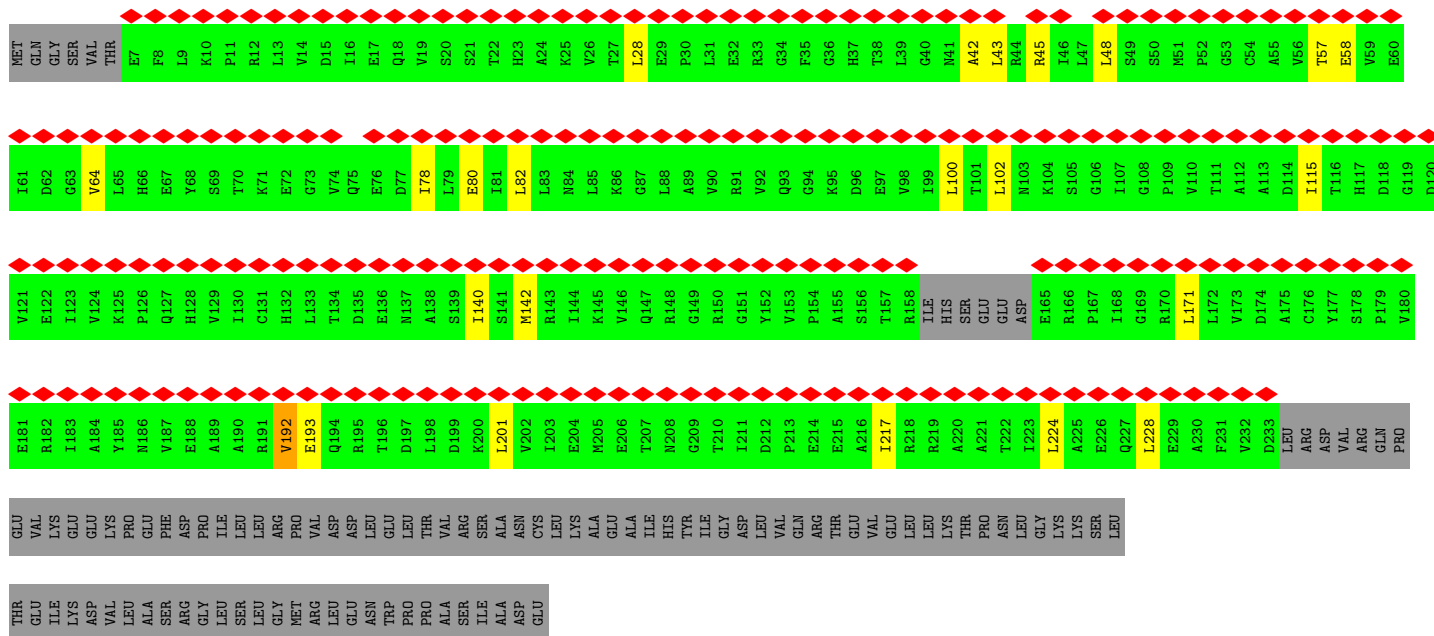
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L1042	A1043	P1044	G1045	A986	L1047	K1048	I1049	V1050	K1051	V1052	Y1053	L1054	A1055	V1056	K1057	R1058	I1059	I1060	Q1061	P1062	G1063	D1064	A1067	G1068	R1069	H1070	G1071	I1072	K1073	I1076	S1077	K1078	I1079	N1080	P1081	I1082	E1083	D1084	M1085	D1088	E1089	N1090	G1091	T1092	P1093	I1096	V1097	L1098	N1099	P1100	L1101	G1102	V1103	F1164	S1105																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				
G982	Q983	V984	E985	A986	E987	K988	L989	D990	K991	L992	P993	R994	D995	R996	H997	L998	E999	G1000	G1001	L1002	T1003	D1004	E1005	E1006	K1007	Q1008	N1009	L1010	L1011	E1012	Q1013	L1014	A1015	E1016	Q1017	Y1018	D1019	S961	E1020	L1021	K1022	H1023	Q965	E1024	F1025	E1026	K1027	K1028	L1029	E1030	A1031	K1032	R1033	R1034	K1035	I1036	T1037	Q1038	G1039	D1040	D1041																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																														
N922	Q923	V924	S925	G926	T927	V928	R929	D930	R931	Q932	V933	F934	T935	R336	D937	Q938	V939	E940	K941	D942	K943	R944	A945	L946	E947	I948	E949	E950	M951	Q952	L953	K954	Q955	A956	K957	K958	D959	L960	S961	E962	E963	L964	Q965	E1024	F1025	E1026	E968	A969	Q970	L971	F972	S973	R974	I975	R976	A977	V978	L979	V980	A981																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
L862	S863	R864	L865	D866	E867	S868	G869	I870	N871	Y872	I873	G874	A875	E876	R877	T878	Q879	E880	G881	I882	L883	V884	G885	K886	R887	T888	E889	L889	GLY	THR	GLN	LEU	THR	GLU	C838	V839	S840	R841	D842	PHE	GLY	GLU	LYS	ALA	ASP	VAL	K914	D915	S916	S917	L918	R919	V920	P921																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																					</



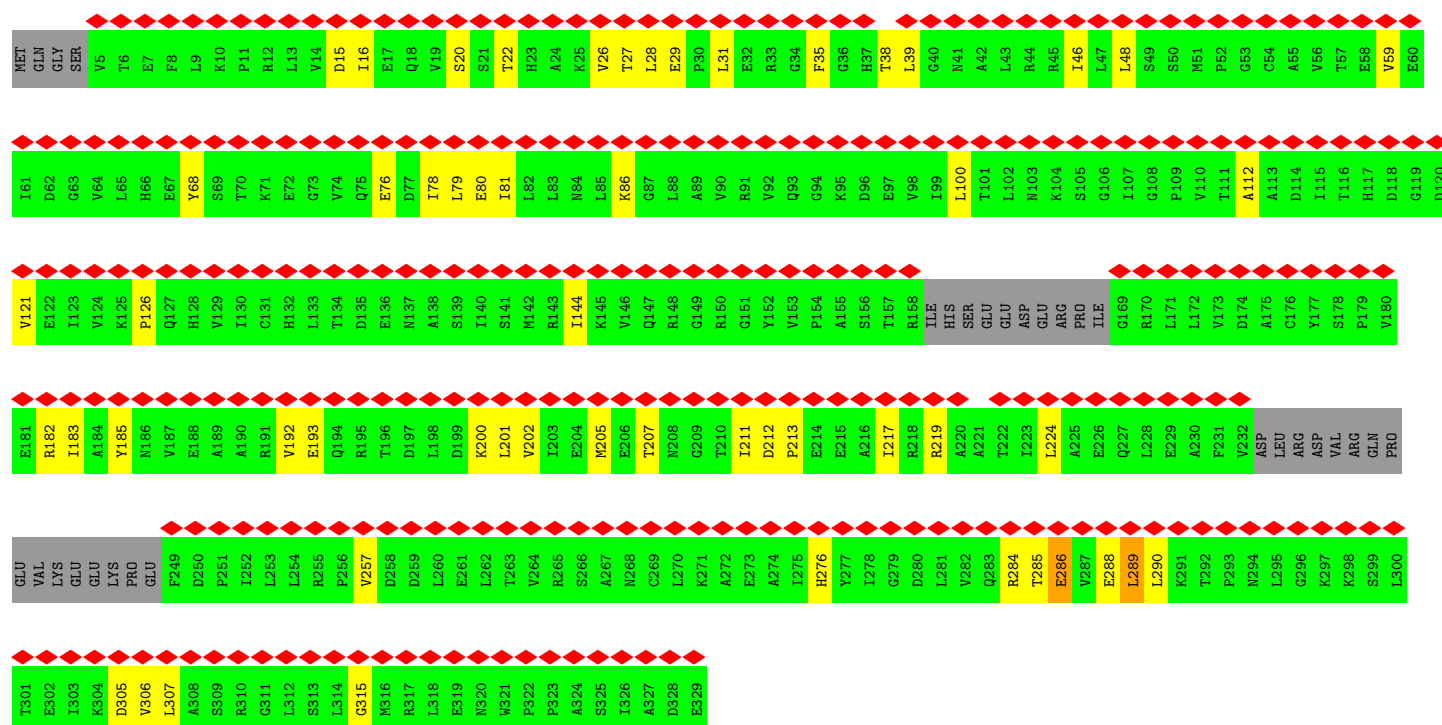
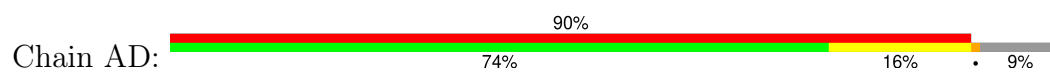
• Molecule 12: Transcription antitermination protein RfaH



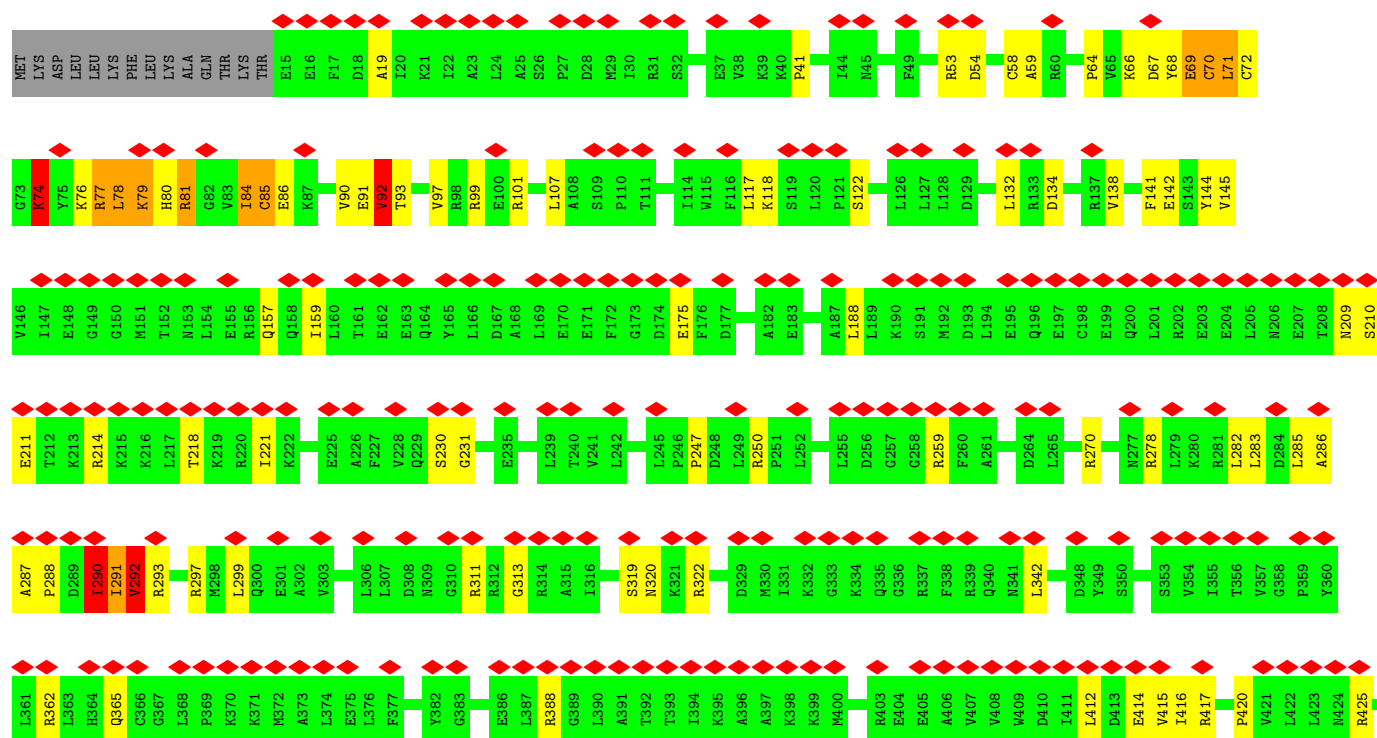
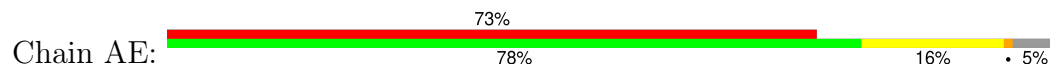
• Molecule 13: DNA-directed RNA polymerase subunit alpha



• Molecule 13: DNA-directed RNA polymerase subunit alpha



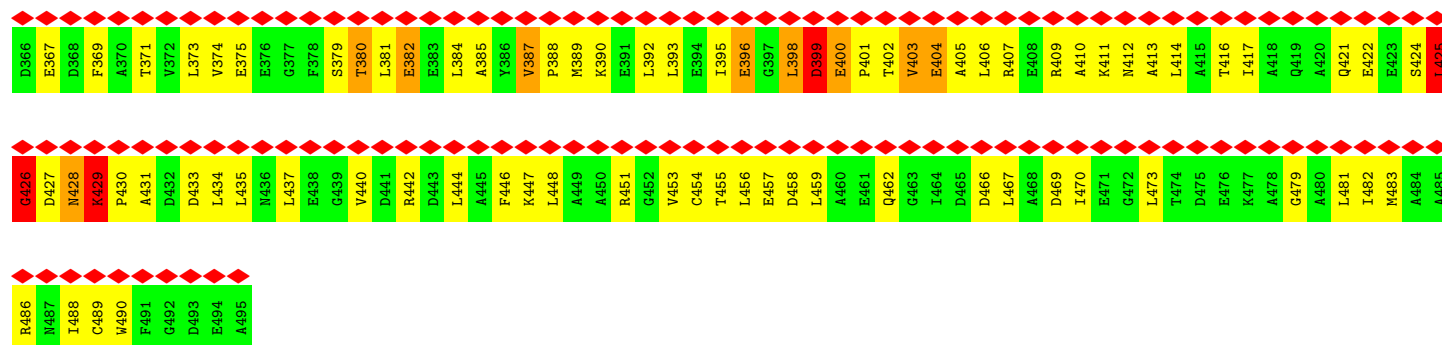
• Molecule 14: DNA-directed RNA polymerase subunit beta'



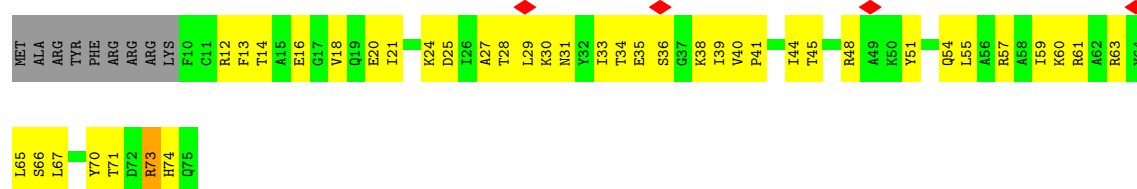
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F620	A621	D622	Q623	I624	M625	Y626	T627	G628	F629	A630	Y631	A632	A633	R634	S635	G636	A637	S638	V639	G640	I641	D642	V643	M644	V645	I646	P647	E648	K649	K650	H651	I652	I653	I654	S655	E656	A657	E658	A659	E660	V661	A662	E663	I664	Q665	E666	Q667	F668	Q669	S670	G671	L672	G673	T674	A675	G676	E677	R678	Y679	
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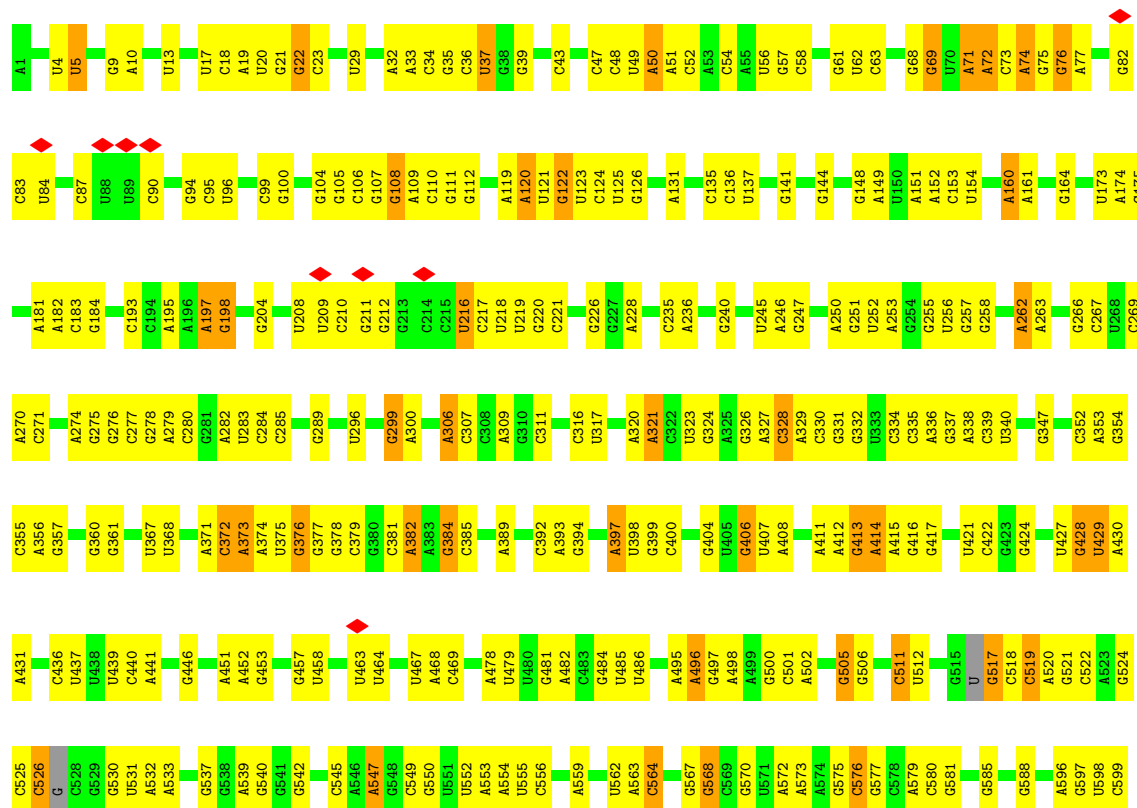




• Molecule 17: 30S ribosomal protein S18

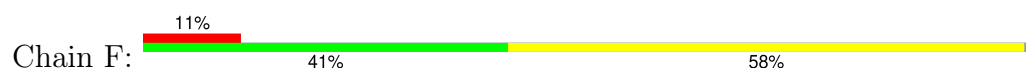


• Molecule 18: 16S rRNA

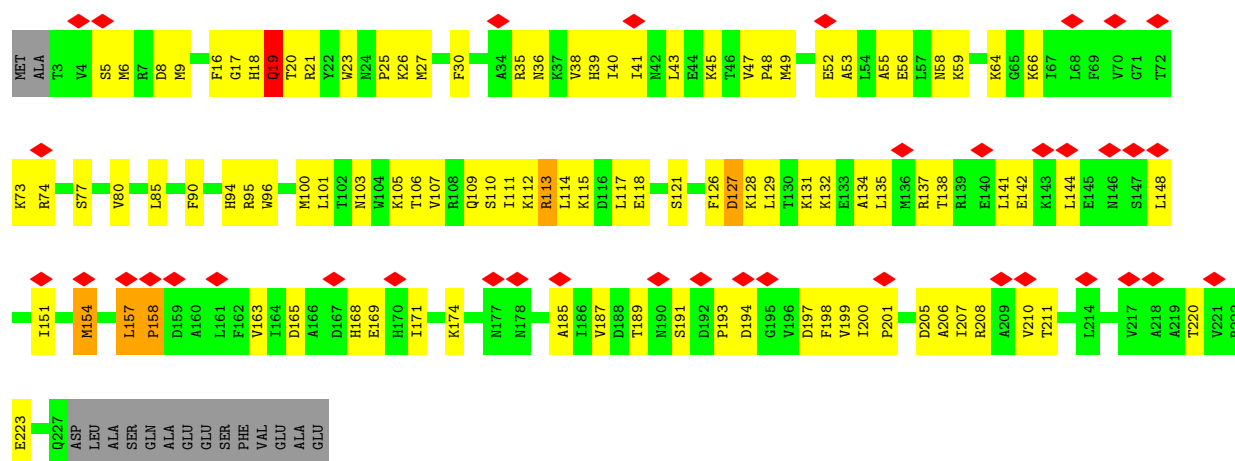




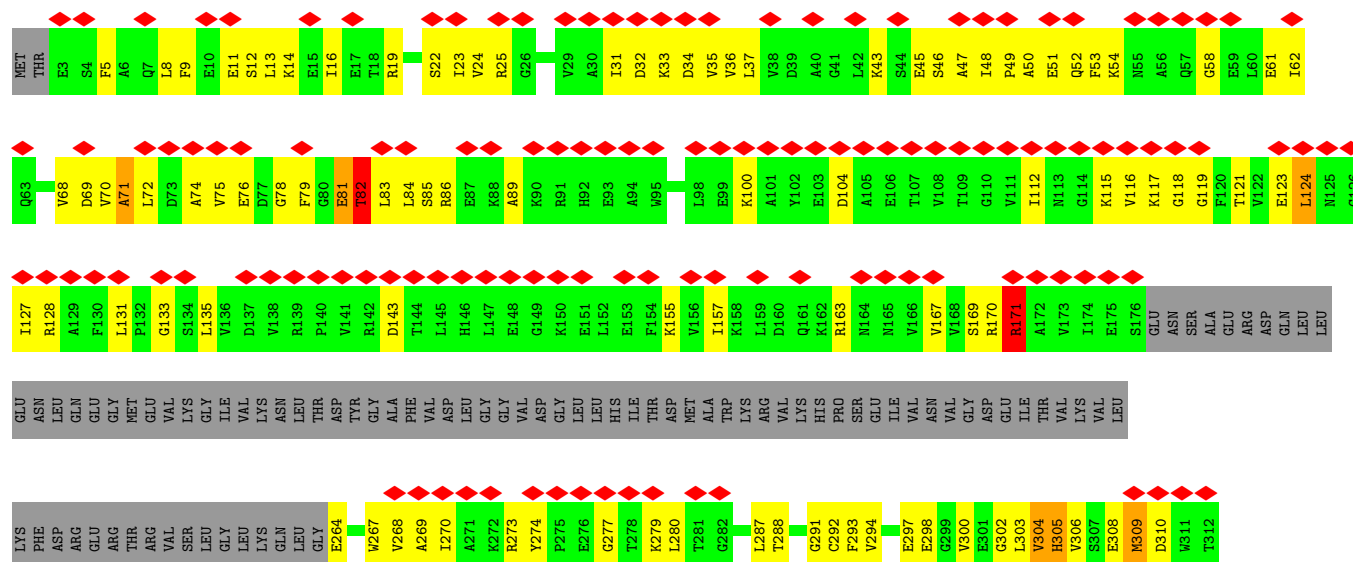
WORLDWIDE  
PDB  
PROTEIN DATA BANK

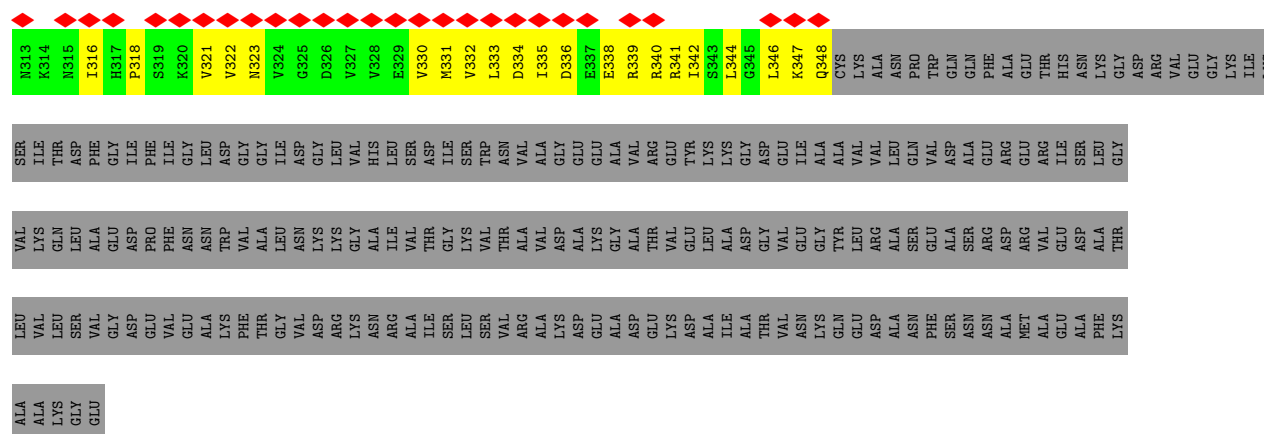


• Molecule 21: 30S ribosomal protein S2

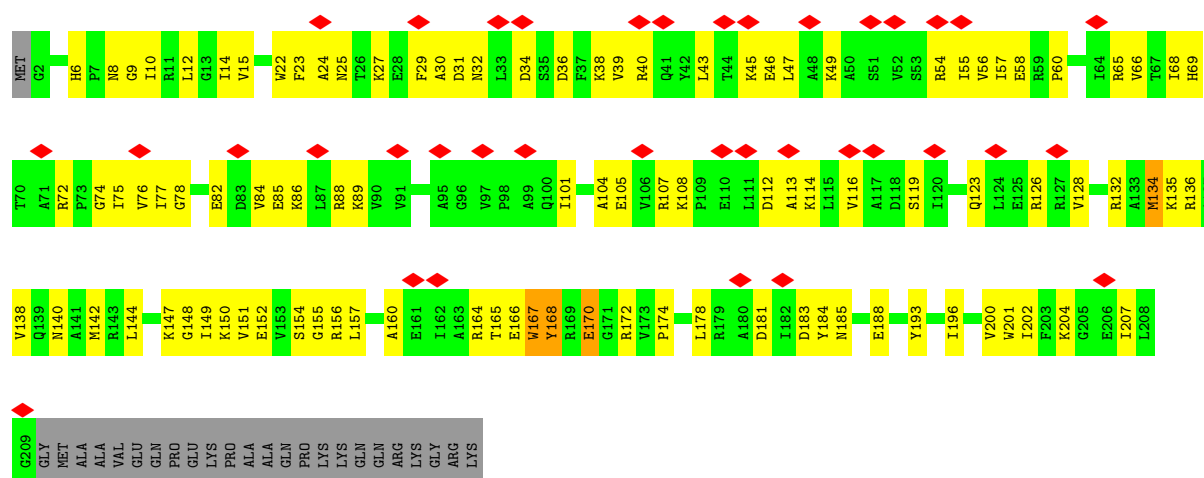
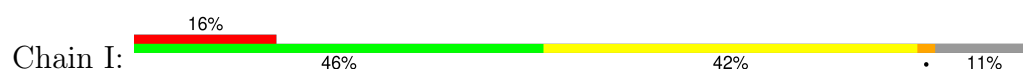


• Molecule 22: 30S ribosomal protein S1

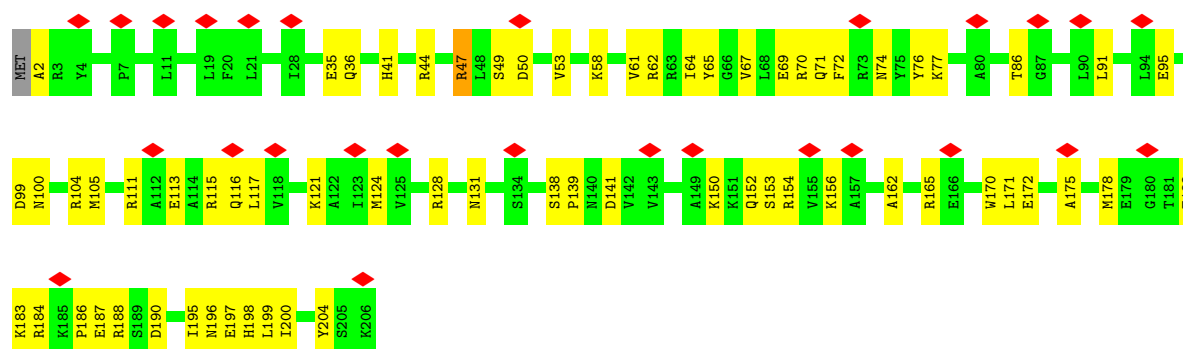




• Molecule 23: 30S ribosomal protein S3

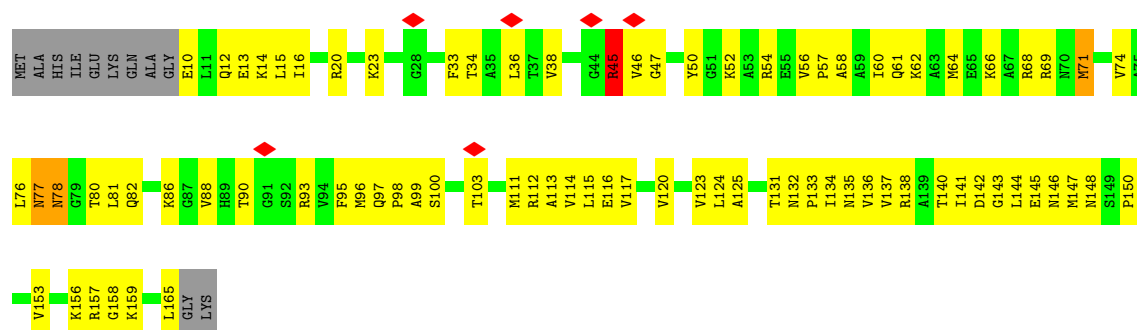


• Molecule 24: 30S ribosomal protein S4

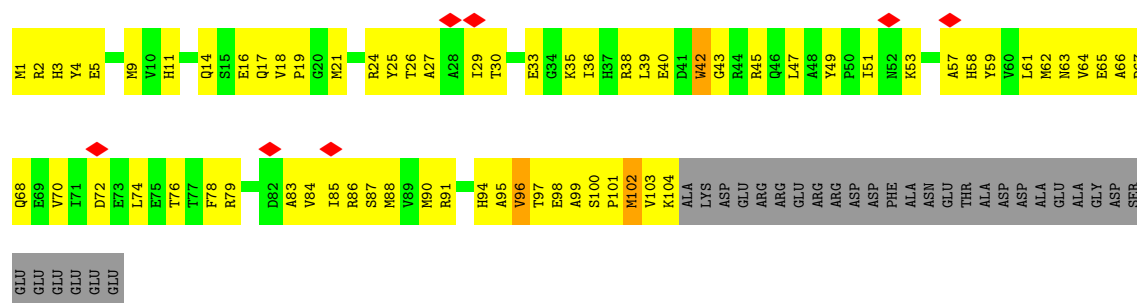


• Molecule 25: 30S ribosomal protein S5

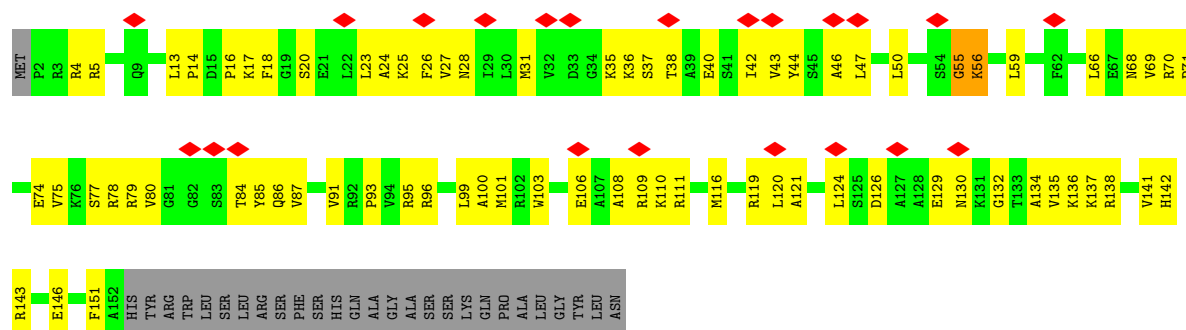
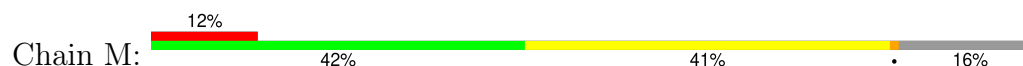




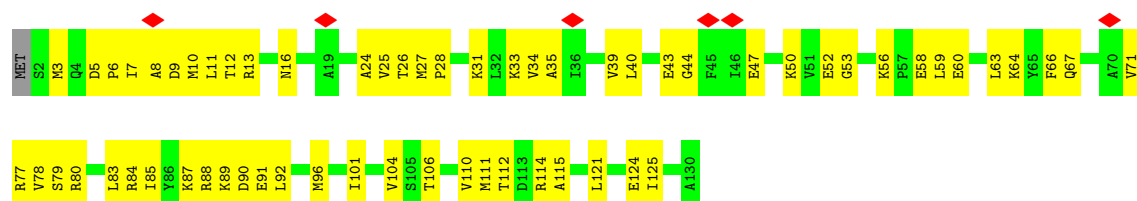
- Molecule 26: 30S ribosomal protein S6



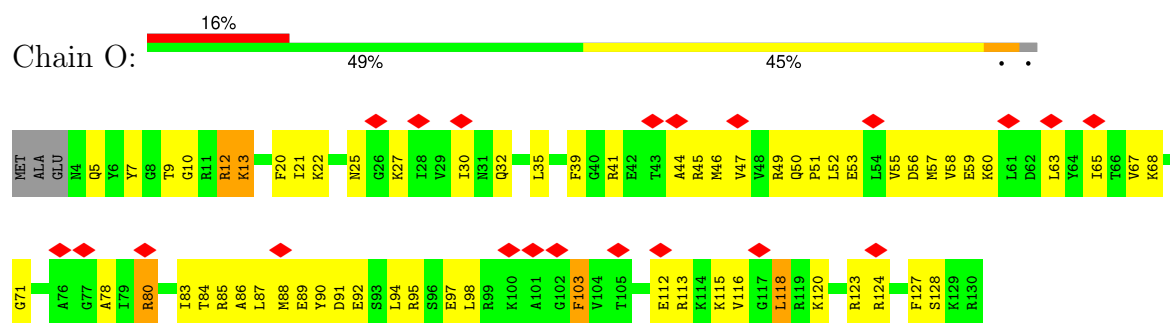
- Molecule 27: 30S ribosomal protein S7



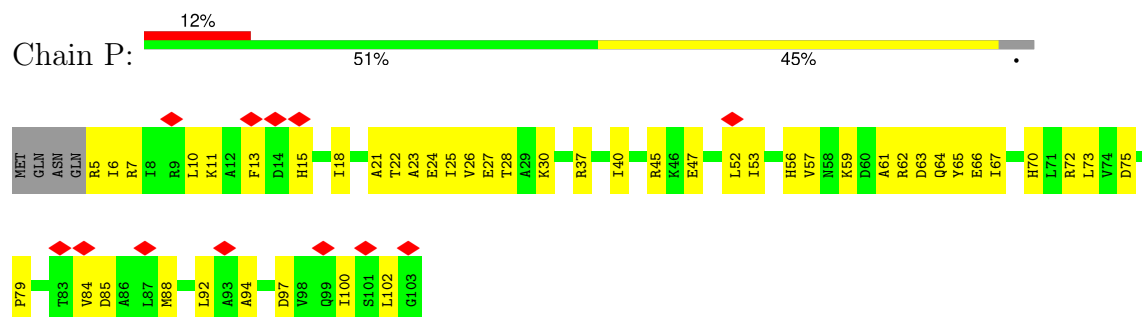
- Molecule 28: 30S ribosomal protein S8



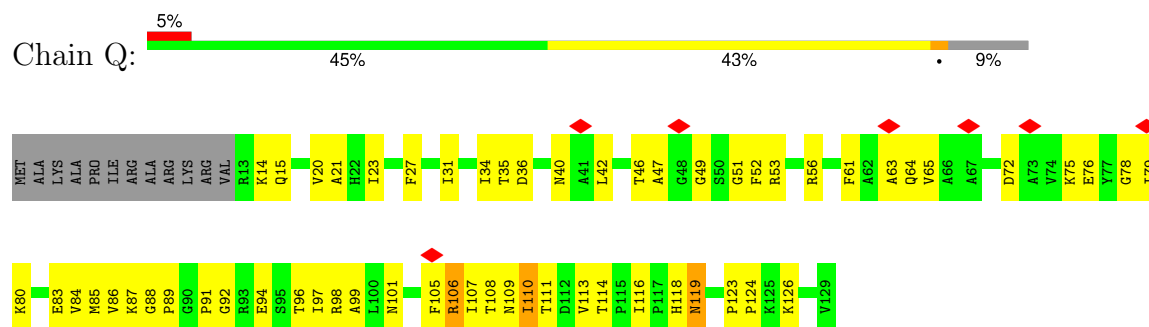
- Molecule 29: 30S ribosomal protein S9



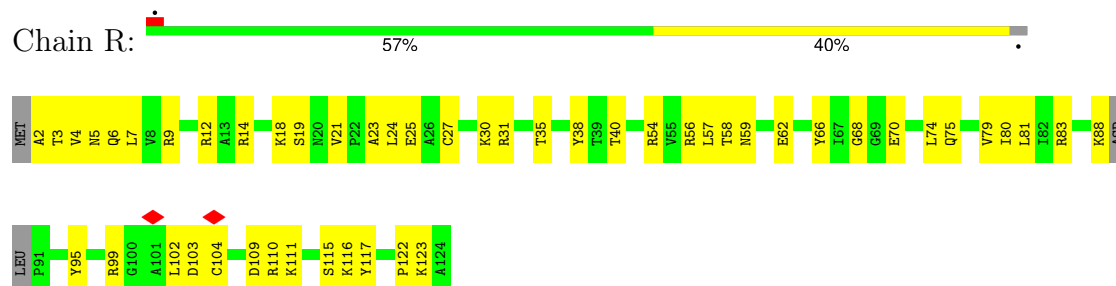
• Molecule 30: 30S ribosomal protein S10



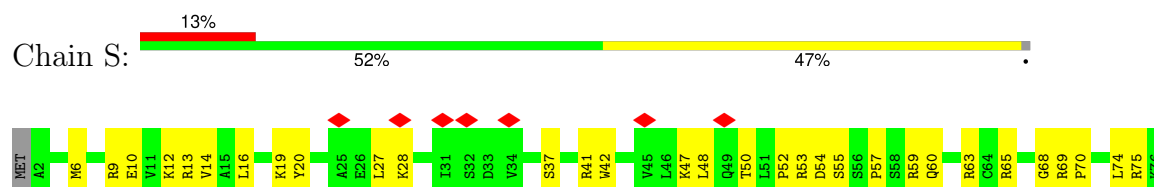
• Molecule 31: 30S ribosomal protein S11



• Molecule 32: 30S ribosomal protein S12

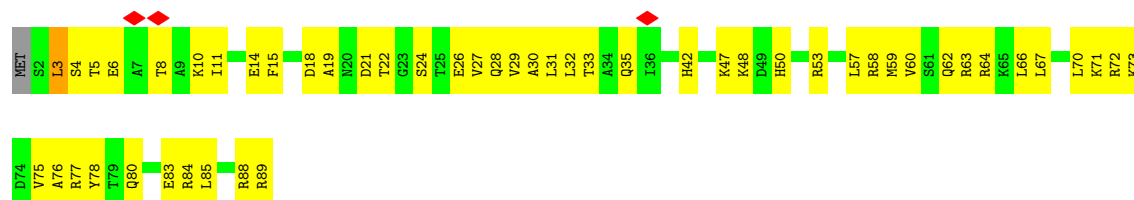


• Molecule 33: 30S ribosomal protein S14

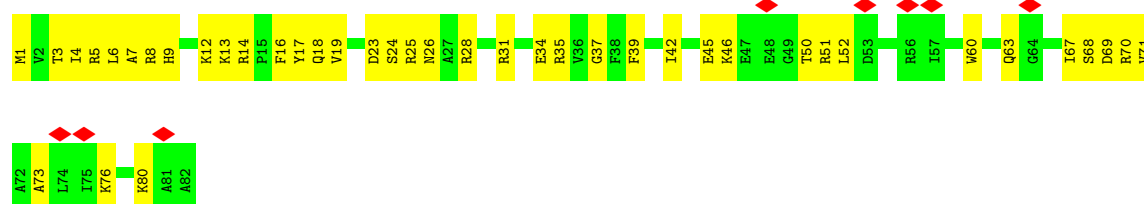




- Molecule 34: Small ribosomal subunit protein uS15



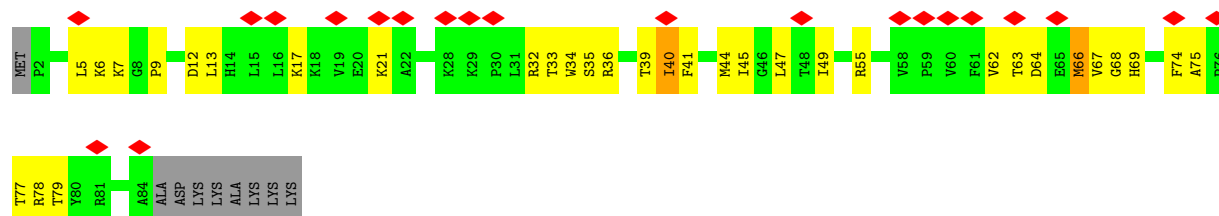
- Molecule 35: 30S ribosomal protein S16



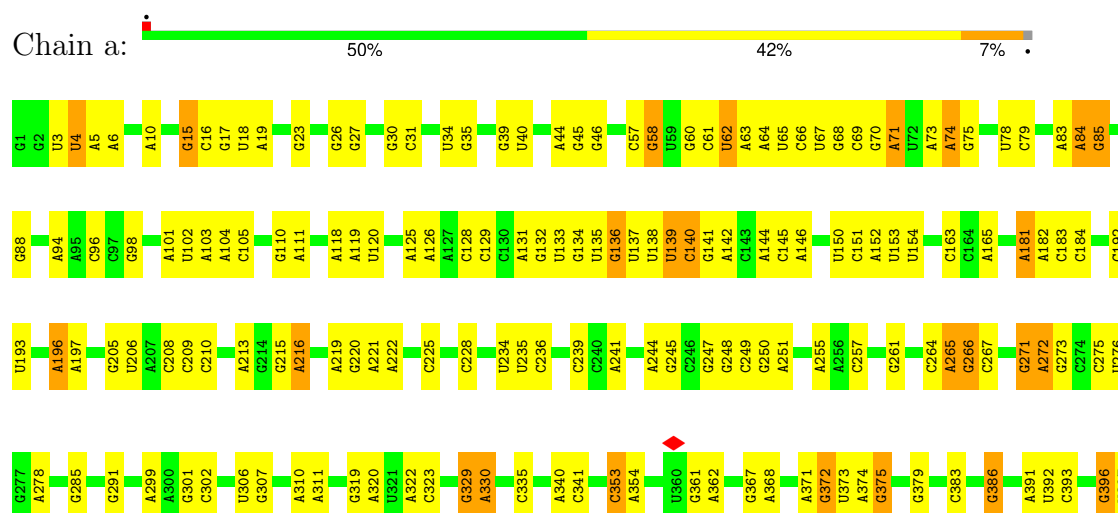
- Molecule 36: 30S ribosomal protein S17



- Molecule 37: 30S ribosomal protein S19



- Molecule 38: 30S ribosomal protein S13

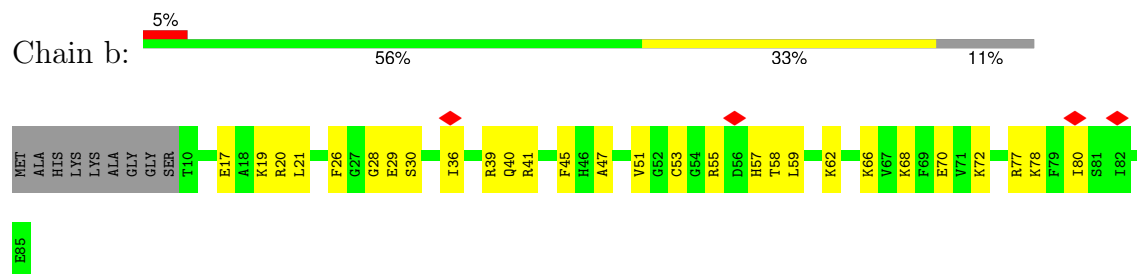




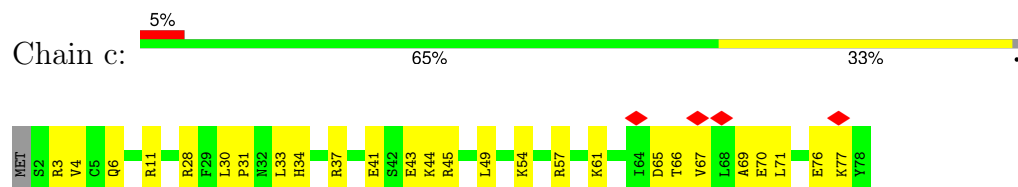
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G2603	U2604	U2606	U2607	U2608	U2609	C2610	C2611	C2612	U2613	U2614	U2615	U2616	U2617	U2618	U2619	U2620	U2627	U2628	U2629	U2630	U2636	U2637	U2638	U2639	U2640	U2646	U2647	U2648	U2649	U2650	U2651	U2663	U2664	U2665	U2666	U2669	U2670	U2671	U2672	U2673	U2674	U2677	U2678	U2682	U2683	U2684	U2685	U2686	U2687	U2688	U2689	U2690																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																										
A2518	U2519	C2520	C2521	U2525	U2526	U2529	U2530	U2531	U2532	U2535	U2536	U2537	U2538	U2539	U2542	U2547	U2548	U2549	U2550	U2555	U2556	U2557	U2558	U2566	U2567	U2568	U2569	U2570	U2571	U2572	U2573	U2574	U2567	U2568	U2569	U2570	U2571	U2572	U2573	U2574	U2579	U2580	U2581	U2582	U2583	U2584	U2585	U2586	U2587	U2588	U2589	U2590	U2591	U2592	U2593	U2594	U2595	U2596	U2597	U2598	U2599	U2600	U2601	U2602	U2603	U2604	U2605	U2606	U2607	U2608	U2609	U2610	U2611	U2612	U2613	U2614	U2615	U2616	U2617	U2618	U2619	U2620	U2621	U2622	U2623	U2624	U2625	U2626	U2627	U2628	U2629	U2630	U2631	U2632	U2633	U2634	U2635	U2636	U2637	U2638	U2639	U2640	U2641	U2642	U2643	U2644	U2645	U2646	U2647	U2648	U2649	U2650	U2651	U2652	U2653	U2654	U2655	U2656	U2657	U2658	U2659	U2660	U2661	U2662	U2663	U2664	U2665	U2666	U2667	U2668	U2669	U2670	U2671	U2672	U2673	U2674	U2675	U2676	U2677	U2678	U2679	U2680	U2681	U2682	U2683	U2684	U2685	U2686	U2687	U2688	U2689	U2690	U2691	U2692	U2693	U2694	U2695	U2696	U2697	U2698	U2699	U2700	U2701	U2702	U2703	U2704	U2705	U2706	U2707	U2708	U2709	U2710	U2711	U2712	U2713	U2714	U2715	U2716	U2717	U2718	U2719	U2720	U2721	U2722	U2723	U2724	U2725	U2726	U2727	U2728	U2729	U2730	U2731	U2732	U2733	U2734	U2735	U2736	U2737	U2738	U2739	U2740	U2741	U2742	U2743	U2744	U2745	U2746	U2747	U2748	U2749	U2750	U2751	U2752	U2753	U2754	U2755	U2756	U2757	U2758	U2759	U2760	U2761	U2762	U2763	U2764	U2765	U2766	U2767	U2768	U2769	U2770	U2771	U2772	U2773	U2774	U2775	U2776	U2777	U2778	U2779	U2780	U2781	U2782	U2783	U2784	U2785	U2786	U2787	U2788	U2789	U2790	U2791	U2792	U2793	U2794	U2795	U2796	U2797	U2798	U2799	U2800	U2801	U2802	U2803	U2804	U2805	U2806	U2807	U2808	U2809	U2810	U2811	U2812	U2813	U2814	U2815	U2816	U2817	U2818	U2819	U2820	U2821	U2822	U2823	U2824	U2825	U2826	U2827	U2828	U2829	U2830	U2831	U2832	U2833	U2834	U2835	U2836	U2837	U2838	U2839	U2840	U2841	U2842	U2843	U2844	U2845	U2846	U2847	U2848	U2849	U2850	U2851	U2852	U2853	U2854	U2855	U2856	U2857	U2858	U2859	U2860	U2861	U2862	U2863	U2864	U2865	U2866	U2867	U2868	U2869	U2870	U2871	U2872	U2873	U2874	U2875	U2876	U2877	U2878	U2879	U2880	U2881	U2882	U2883	U2884	U2885	U2886	U2887	U2888	U2889	U2890	U2891	U2892	U2893	U2894	U2895	U2896	U2897	U2898	U2899	U2900	U2901	U2902	U2903	U2904	U2905	U2906	U2907	U2908	U2909	U2910	U2911	U2912	U2913	U2914	U2915	U2916	U2917	U2918	U2919	U2920	U2921	U2922	U2923	U2924	U2925	U2926	U2927	U2928	U2929	U2930	U2931	U2932	U2933	U2934	U2935	U2936	U2937	U2938	U2939	U2940	U2941	U2942	U2943	U2944	U2945	U2946	U2947	U2948	U2949	U2950	U2951	U2952	U2953	U2954	U2955	U2956	U2957	U2958	U2959	U2960	U2961	U2962	U2963	U2964	U2965	U2966	U2967	U2968	U2969	U2970	U2971	U2972	U2973	U2974	U2975	U2976	U2977	U2978	U2979	U2980	U2981	U2982	U2983	U2984	U2985	U2986	U2987	U2988	U2989	U2990	U2991	U2992	U2993	U2994	U2995	U2996	U2997	U2998	U2999	U3000	U3001	U3002	U3003	U3004	U3005	U3006	U3007	U3008	U3009	U3010	U3011	U3012	U3013	U3014	U3015	U3016	U3017	U3018	U3019	U3020	U3021	U3022	U3023	U3024	U3025	U3026	U3027	U3028	U3029	U3030	U3031	U3032	U3033	U3034	U3035	U3036	U3037	U3038	U3039	U3040	U3041	U3042	U3043	U3044	U3045	U3046	U3047	U3048	U3049	U3050	U3051	U3052	U3053	U3054	U3055	U3056	U3057	U3058	U3059	U3060	U3061	U3062	U3063	U3064	U3065	U3066	U3067	U3068	U3069	U3070	U3071	U3072	U3073	U3074	U3075	U3076	U3077	U3078	U3079	U3080	U3081	U3082	U3083	U3084	U3085	U3086	U3087	U3088	U3089	U3090	U3091	U3092	U3093	U3094	U3095	U3096	U3097	U3098	U3099	U3100	U3101	U3102	U3103	U3104	U3105	U3106	U3107	U3108	U3109	U3110	U3111	U3112	U3113	U3114	U3115	U3116	U3117	U3118	U3119	U3120	U3121	U3122	U3123	U3124	U3125	U3126	U3127	U3128	U3129	U3130	U3131	U3132	U3133	U3134	U3135	U3136	U3137	U3138	U3139	U3140	U3141	U3142	U3143	U3144	U3145	U3146	U3147	U3148	U3149	U3150	U3151	U3152	U3153	U3154	U3155	U3156	U3157	U3158	U3159	U3160	U3161	U3162	U3163	U3164	U3165	U3166	U3167	U3168	U3169	U3170	U3171	U3172	U3173	U3174	U3175	U3176	U3177	U3178	U3179	U3180	U3181	U3182	U3183	U3184	U3185	U3186	U3187	U3188	U3189	U3190	U3191	U3192	U3193	U3194	U3195	U3196	U3197	U3198	U3199	U3200	U3201	U3202	U3203	U3204	U3205	U3206	U3207	U3208	U3209	U3210	U3211	U3212	U3213	U3214	U3215	U3216	U3217	U3218	U3219	U3220	U3221	U3222	U3223	U3224	U3225	U3226	U3227	U3228	U3229	U3230	U3231	U3232	U3233	U3234	U3235	U3236	U3237	U3238	U3239	U3240	U3241	U3242	U3243	U3244	U3245	U3246	U3247	U3248	U3249	U3250	U3251	U3252	U3253	U3254	U3255	U3256	U3257	U3258	U3259	U3260	U3261	U3262	U3263	U3264	U3265	U3266	U3267	U3268	U3269	U3270	U3271	U3272	U3273	U3274	U3275	U3276	U3277	U3278	U3279	U3280	U3281	U3282	U3283	U3284	U3285	U3286	U3287	U3288	U3289	U3290	U3291	U3292	U3293	U3294	U3295	U3296	U3297	U3298	U3299	U3300	U3301	U3302	U3303	U3304	U3305	U3306	U3307	U3308	U3309	U3310	U3311	U3312	U3313	U3314	U3315	U3316	U3317	U3318	U3319	U3320	U3321	U3322	U3323	U3324	U3325	U3326	U3327	U3328	U3329	U3330	U3331	U3332	U3333	U3334	U3335	U3336	U3337	U3338	U3339	U3340	U3341	U3342	U3343	U3344	U3345	U3346	U3347	U3348	U3349	U3350	U3351	U3352	U3353	U3354	U3355	U3356	U3357	U3358	U3359	U3360	U3361	U3362	U3363	U3364	U3365	U3366	U3367	U3368	U3369	U3370	U3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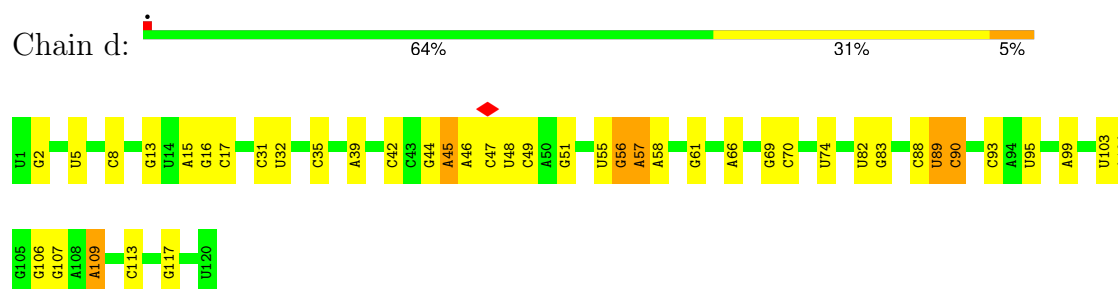
- Molecule 42: 50S ribosomal protein L27



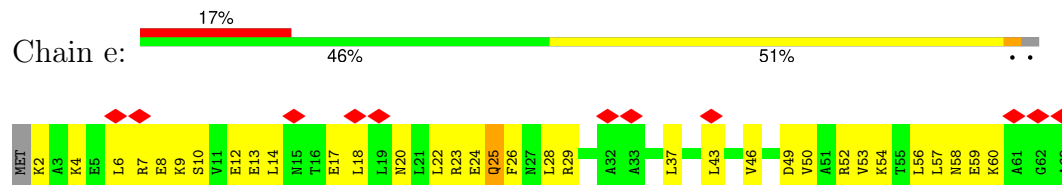
- Molecule 43: 50S ribosomal protein L28



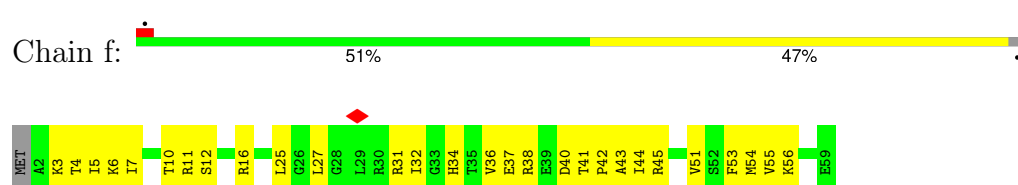
- Molecule 44: 5S rRNA



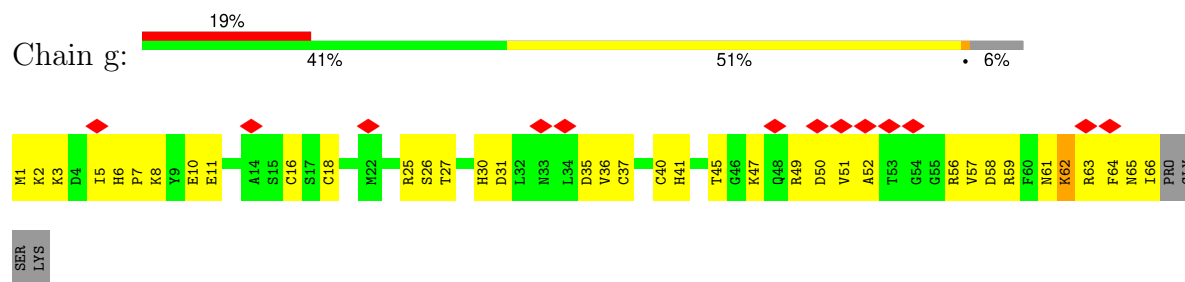
- Molecule 45: 50S ribosomal protein L29



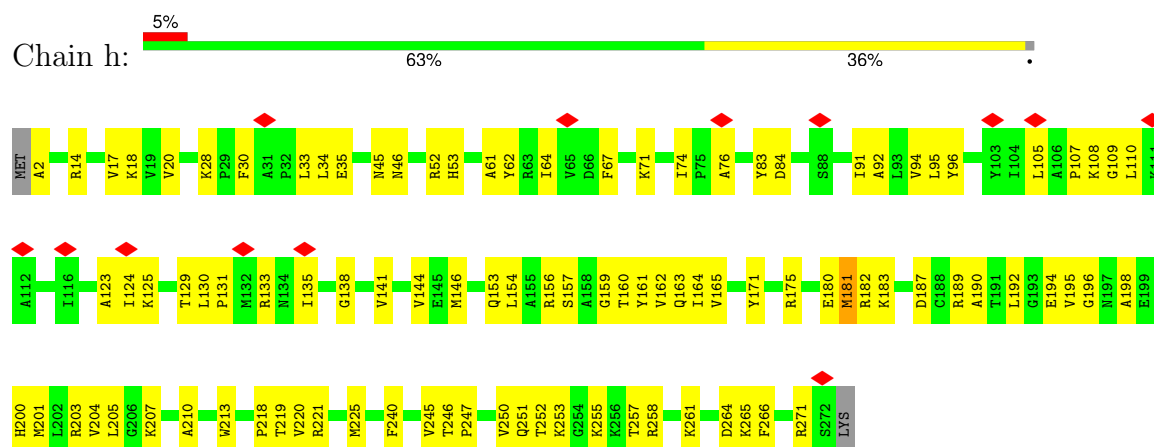
- Molecule 46: 50S ribosomal protein L30



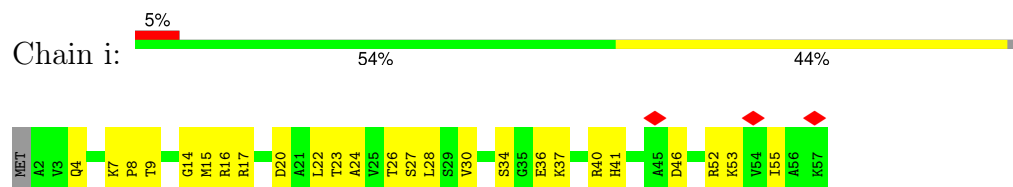
- Molecule 47: 50S ribosomal protein L31



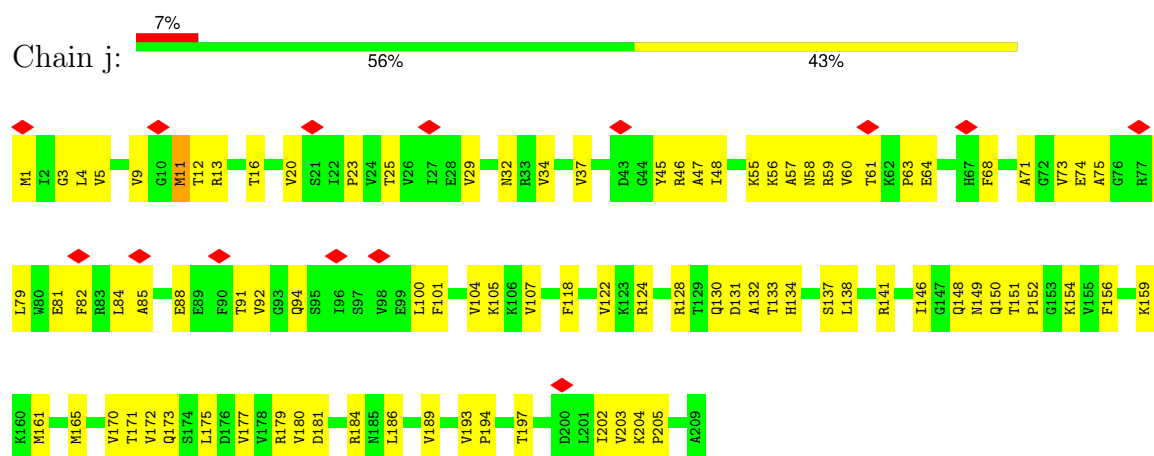
- Molecule 48: 50S ribosomal protein L2



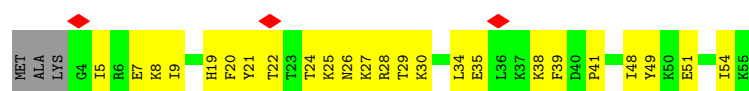
- Molecule 49: 50S ribosomal protein L32



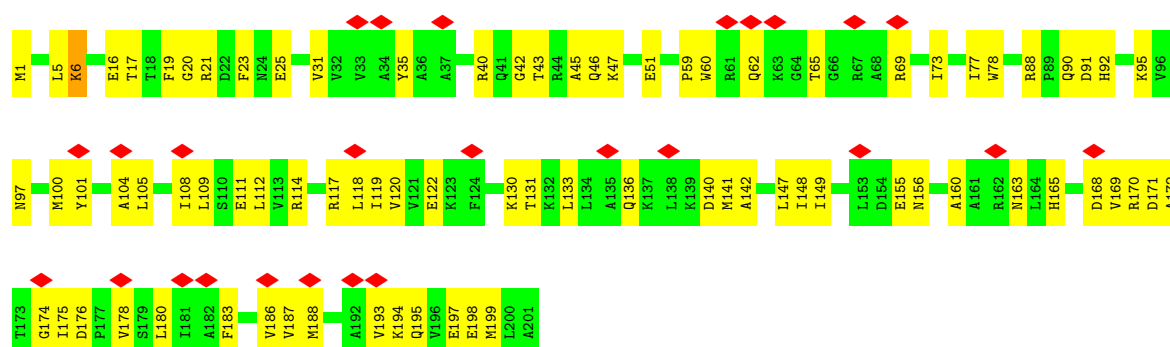
- Molecule 50: 50S ribosomal protein L3



- Molecule 51: 50S ribosomal protein L33



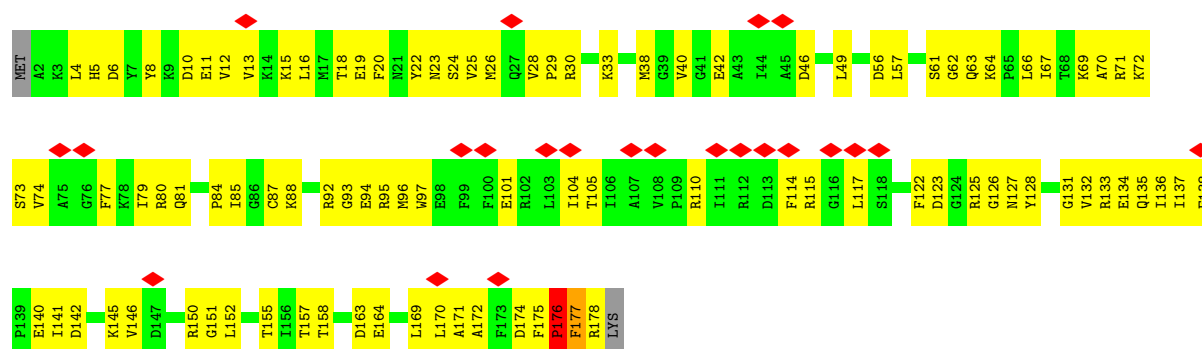
• Molecule 52: 50S ribosomal protein L4



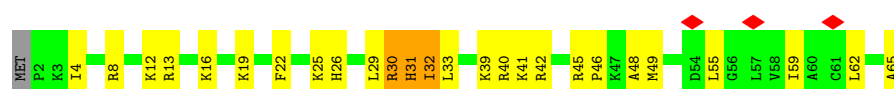
• Molecule 53: 50S ribosomal protein L34



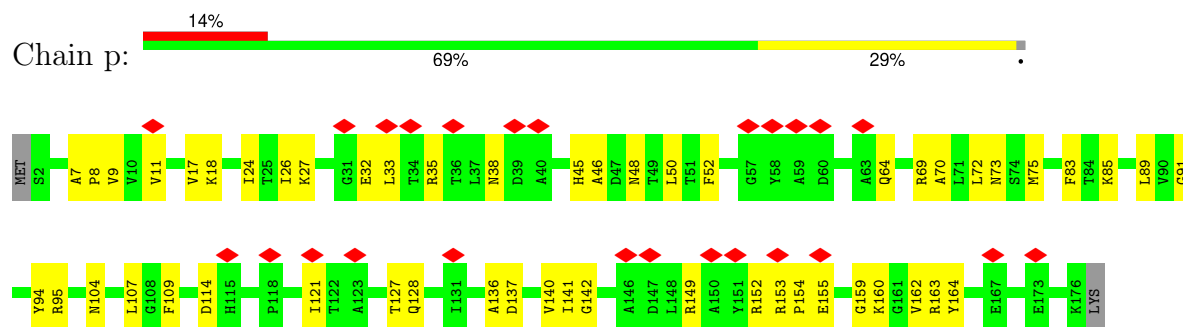
• Molecule 54: 50S ribosomal protein L5



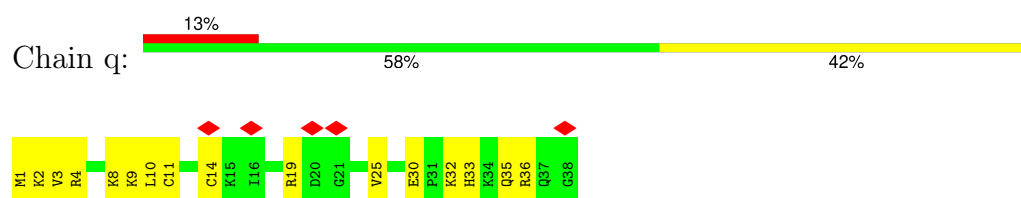
• Molecule 55: 50S ribosomal protein L35



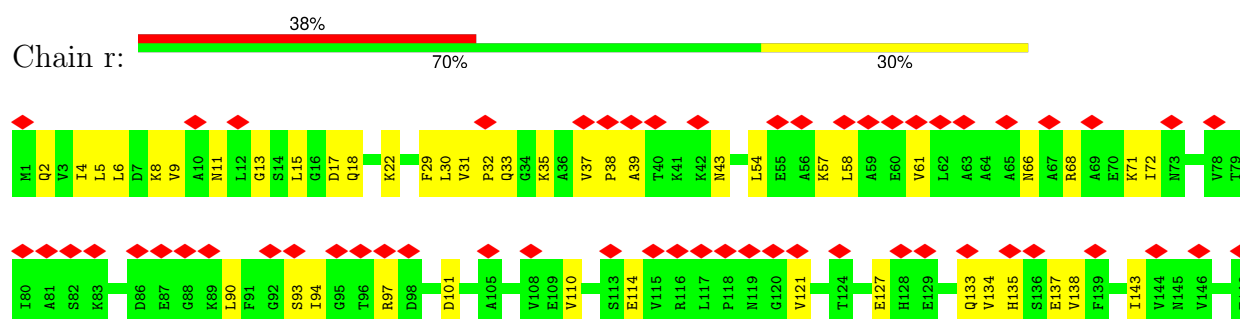
- Molecule 56: 50S ribosomal protein L6



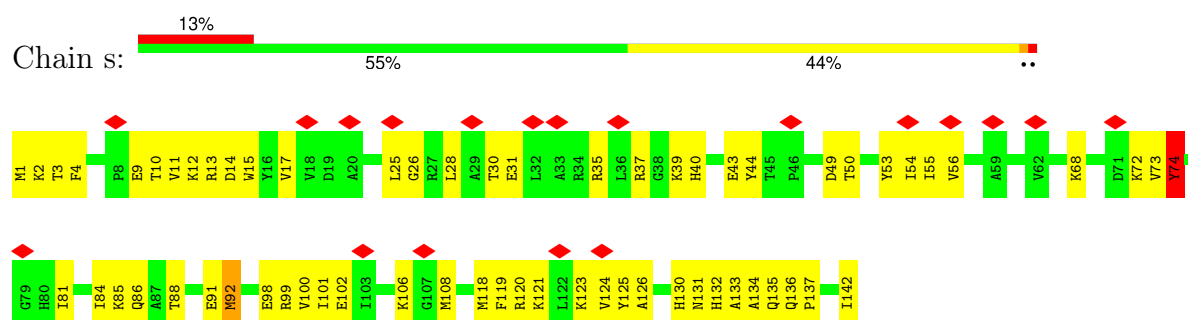
- Molecule 57: 50S ribosomal protein L36



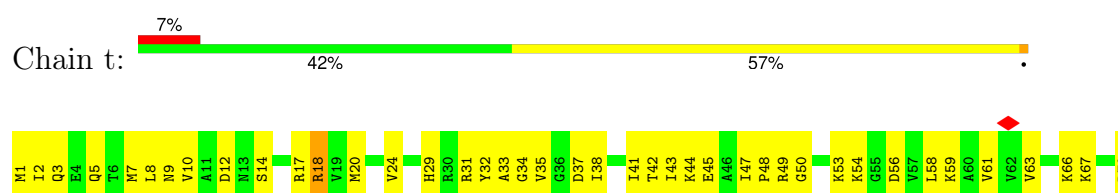
- Molecule 58: 50S ribosomal protein L9

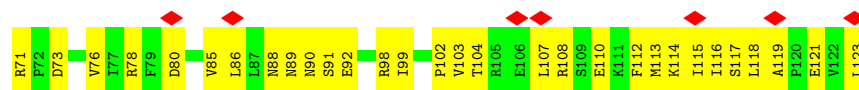


- Molecule 59: 50S ribosomal protein L13

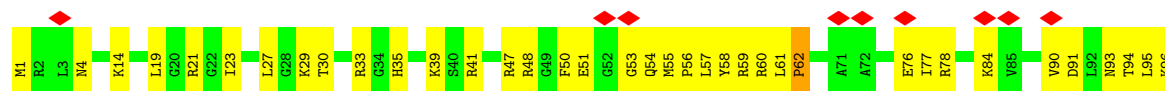


- Molecule 60: 50S ribosomal protein L14

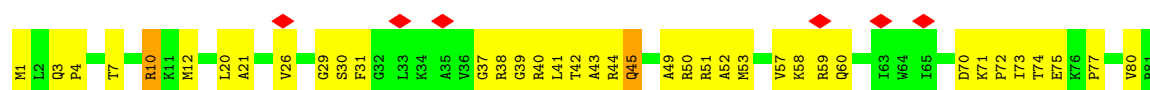




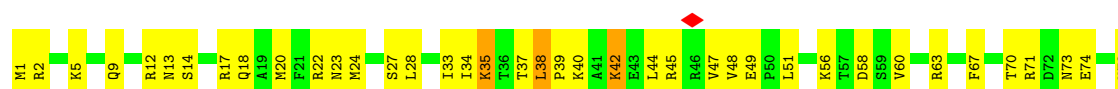
- Molecule 61: 50S ribosomal protein L15



- Molecule 62: 50S ribosomal protein L16



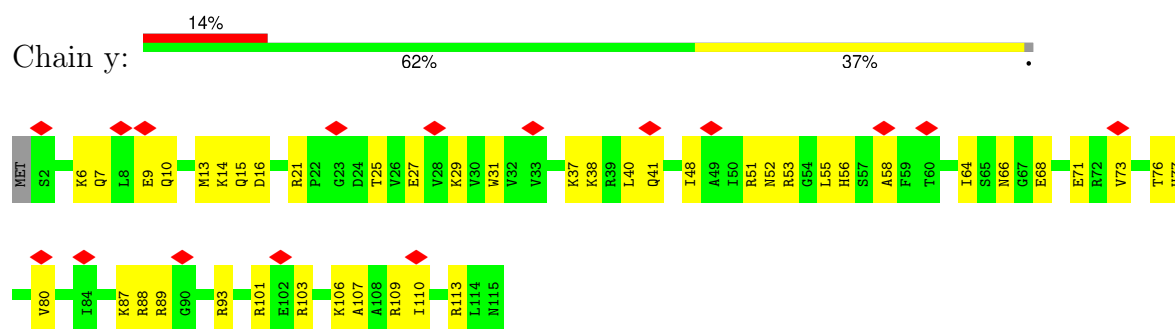
- Molecule 63: 50S ribosomal protein L17



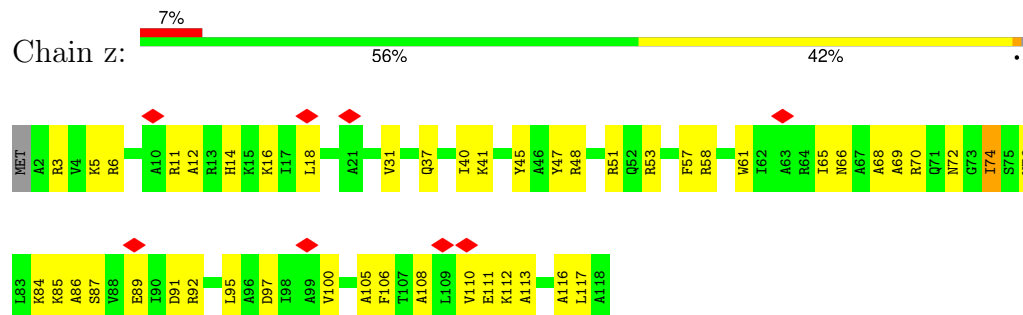
- Molecule 64: 50S ribosomal protein L18



- Molecule 65: 50S ribosomal protein L19



• Molecule 66: 50S ribosomal protein L20





## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	15810	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	50	Depositor
Minimum defocus (nm)	1250	Depositor
Maximum defocus (nm)	2500	Depositor
Magnification	Not provided	
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor
Maximum map value	0.019	Depositor
Minimum map value	-0.087	Depositor
Average map value	-0.006	Depositor
Map value standard deviation	0.004	Depositor
Recommended contour level	0.000001	Depositor
Map size (Å)	533.2, 533.2, 533.2	wwPDB
Map dimensions	620, 620, 620	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.86, 0.86, 0.86	Depositor

## 5 Model quality ⓘ

### 5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: MG, ZN

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	0	0.43	0/829	0.51	0/1107
2	1	0.60	0/864	0.63	0/1156
3	2	0.55	1/752 (0.1%)	0.67	0/1005
4	3	0.36	0/796	0.49	0/1062
5	4	0.57	0/766	0.63	0/1025
6	5	0.57	0/816	0.78	0/1259
7	6	0.57	0/783	0.78	0/1203
8	7	0.61	2/585 (0.3%)	0.91	3/906 (0.3%)
9	9	0.37	0/1131	0.73	2/1524 (0.1%)
10	A	0.27	0/1810	0.58	0/2821
10	B	0.27	0/1810	0.58	0/2821
11	AA	0.38	0/10547	0.62	0/14232
12	AB	0.53	0/1317	1.20	17/1786 (1.0%)
13	AC	0.38	0/1718	0.62	0/2328
13	AD	0.37	0/2096	0.69	3/2854 (0.1%)
14	AE	0.39	0/10561	0.64	5/14258 (0.0%)
15	AF	0.30	0/652	0.61	0/879
16	AG	0.97	6/3897 (0.2%)	1.40	49/5273 (0.9%)
17	C	0.67	0/553	0.81	1/743 (0.1%)
18	D	0.29	0/36610	0.41	9/57091 (0.0%)
19	E	0.61	0/675	0.72	0/895
20	F	0.53	0/597	0.61	0/792
21	G	0.68	5/1791 (0.3%)	0.79	8/2413 (0.3%)
22	H	0.38	0/1746	0.81	0/2382
23	I	0.57	1/1663 (0.1%)	0.68	4/2241 (0.2%)
24	J	0.46	0/1665	0.58	0/2227
25	K	0.72	2/1165 (0.2%)	0.86	6/1568 (0.4%)
26	L	0.68	1/867 (0.1%)	0.79	1/1171 (0.1%)
27	M	0.56	0/1195	0.69	2/1602 (0.1%)
28	N	0.50	0/989	0.58	0/1326
29	O	0.66	3/1034 (0.3%)	0.79	3/1375 (0.2%)
30	P	0.51	0/800	0.73	0/1082

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
31	Q	0.72	2/893 (0.2%)	0.76	2/1205 (0.2%)
32	R	0.49	0/952	0.59	0/1274
33	S	0.53	0/817	0.61	0/1088
34	T	0.57	1/722 (0.1%)	0.68	0/964
35	U	0.42	0/659	0.53	0/884
36	V	0.49	0/657	0.62	0/881
37	W	0.56	1/680 (0.1%)	0.62	1/915 (0.1%)
38	X	0.45	0/909	0.72	1/1215 (0.1%)
39	Y	0.38	2/1046 (0.2%)	0.66	2/1410 (0.1%)
40	Z	0.13	0/227	0.41	0/304
41	a	0.30	0/69247	0.43	10/107985 (0.0%)
42	b	0.42	0/589	0.55	0/779
43	c	0.51	0/635	0.61	0/848
44	d	0.25	0/2872	0.37	0/4478
45	e	0.69	2/502 (0.4%)	0.65	1/667 (0.1%)
46	f	0.52	0/452	0.57	0/605
47	g	0.45	1/531 (0.2%)	0.64	0/709
48	h	0.50	1/2121 (0.0%)	0.59	1/2852 (0.0%)
49	i	0.41	0/450	0.55	0/599
50	j	0.51	0/1586	0.57	1/2134 (0.0%)
51	k	0.46	0/433	0.63	0/576
52	l	0.51	1/1571 (0.1%)	0.62	0/2113
53	m	0.45	0/380	0.56	0/498
54	n	0.46	0/1434	0.66	1/1926 (0.1%)
55	o	0.51	0/513	0.71	1/676 (0.1%)
56	p	0.42	0/1333	0.62	2/1805 (0.1%)
57	q	0.45	0/303	0.64	0/397
58	r	0.28	0/1122	0.48	0/1515
59	s	0.70	2/1152 (0.2%)	0.74	2/1551 (0.1%)
60	t	0.53	0/955	0.75	3/1279 (0.2%)
61	u	0.43	0/1062	0.59	0/1413
62	v	0.60	1/1093 (0.1%)	0.77	3/1460 (0.2%)
63	w	0.83	2/964 (0.2%)	0.80	3/1289 (0.2%)
64	x	0.34	0/902	0.50	0/1209
65	y	0.41	0/929	0.54	0/1242
66	z	0.62	2/960 (0.2%)	0.62	0/1278
All	All	0.41	39/194733 (0.0%)	0.56	147/286430 (0.1%)

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
9	9	0	3
12	AB	0	1
13	AC	0	1
13	AD	0	2
14	AE	0	4
16	AG	0	6
21	G	0	1
22	H	0	5
23	I	0	1
25	K	0	2
29	O	0	1
38	X	0	1
39	Y	0	1
54	n	0	1
55	o	0	1
59	s	0	1
61	u	0	2
All	All	0	34

All (39) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
16	AG	429	LYS	C-N	13.93	1.51	1.33
63	w	35	LYS	CE-NZ	-12.76	1.11	1.49
29	O	80	ARG	CD-NE	-10.52	1.31	1.46
63	w	42	LYS	CD-CE	-9.42	1.24	1.52
25	K	45	ARG	CG-CD	7.72	1.75	1.52
66	z	74	ILE	CG1-CD1	7.61	1.81	1.51
8	7	55	G	C1'-N9	-7.52	1.36	1.48
21	G	19	GLN	CB-CG	-7.52	1.29	1.52
29	O	118	LEU	CG-CD2	-7.32	1.28	1.52
37	W	66	MET	CG-SD	-7.21	1.62	1.80
26	L	42	TRP	CB-CG	-7.15	1.28	1.50
59	s	74	TYR	CZ-OH	-7.15	1.23	1.38
45	e	46	VAL	CB-CG1	-7.11	1.29	1.52
8	7	2	U	C1'-N1	7.02	1.58	1.48
3	2	5	GLU	CG-CD	-7.00	1.34	1.52
16	AG	246	ASP	C-N	6.98	1.50	1.33
31	Q	106	ARG	CD-NE	-6.41	1.37	1.46
29	O	80	ARG	CZ-NH2	6.30	1.41	1.33
21	G	158	PRO	CA-C	-6.25	1.44	1.52
16	AG	355	ALA	N-CA	6.24	1.54	1.46
48	h	181	MET	CG-SD	-6.05	1.65	1.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
66	z	74	ILE	CB-CG2	-6.04	1.32	1.52
21	G	158	PRO	CG-CD	6.01	1.71	1.50
39	Y	5	GLN	CB-CG	-6.01	1.34	1.52
59	s	92	MET	CG-SD	-5.93	1.66	1.80
16	AG	355	ALA	CA-C	5.87	1.60	1.52
25	K	45	ARG	CD-NE	5.84	1.54	1.46
52	l	6	LYS	CE-NZ	-5.81	1.31	1.49
21	G	157	LEU	C-O	-5.71	1.15	1.24
34	T	3	LEU	CG-CD2	-5.62	1.33	1.52
39	Y	5	GLN	CA-C	-5.59	1.45	1.52
47	g	62	LYS	CE-NZ	-5.53	1.32	1.49
45	e	25	GLN	CB-CG	-5.38	1.36	1.52
23	I	134	MET	CB-CG	-5.30	1.36	1.52
16	AG	99	SER	N-CA	5.21	1.49	1.46
31	Q	119	ASN	CG-ND2	5.14	1.44	1.33
21	G	19	GLN	CD-NE2	-5.13	1.22	1.33
62	v	45	GLN	CB-CG	-5.11	1.37	1.52
16	AG	293	VAL	N-CA	5.07	1.52	1.46

All (147) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
16	AG	104	ARG	CA-C-N	17.97	154.32	121.97
16	AG	104	ARG	C-N-CA	17.97	154.32	121.97
16	AG	246	ASP	CA-C-N	14.49	137.95	119.84
16	AG	246	ASP	C-N-CA	14.49	137.95	119.84
16	AG	354	ALA	O-C-N	-13.69	107.30	122.09
16	AG	429	LYS	CA-C-N	11.45	132.17	119.92
16	AG	429	LYS	C-N-CA	11.45	132.17	119.92
25	K	71	MET	CG-SD-CE	-11.35	75.92	100.90
21	G	158	PRO	N-CD-CG	-10.95	86.78	103.20
12	AB	154	VAL	CA-C-N	10.81	140.09	121.29
12	AB	154	VAL	C-N-CA	10.81	140.09	121.29
16	AG	24	PHE	CA-CB-CG	-10.58	103.22	113.80
16	AG	105	ILE	CA-C-N	10.15	136.84	120.60
16	AG	105	ILE	C-N-CA	10.15	136.84	120.60
26	L	102	MET	CA-CB-CG	-9.62	94.86	114.10
59	s	92	MET	CG-SD-CE	-9.37	80.29	100.90
21	G	113	ARG	NE-CZ-NH2	9.26	127.53	119.20
16	AG	103	ASP	CA-CB-CG	9.02	121.62	112.60
8	7	1	A	O3'-P-O5'	-8.64	91.05	104.00
25	K	78	ASN	N-CA-CB	-8.51	96.12	110.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
12	AB	129	ILE	N-CA-C	8.43	120.24	107.77
16	AG	35	THR	CA-CB-OG1	-8.27	97.19	109.60
8	7	4	U	C2'-C3'-O3'	8.27	121.90	109.50
16	AG	292	ASP	O-C-N	-8.20	107.51	122.44
16	AG	109	THR	CB-CA-C	8.16	124.73	110.85
16	AG	108	GLN	N-CA-CB	8.07	123.55	110.40
60	t	18	ARG	NE-CZ-NH2	8.02	126.42	119.20
12	AB	128	ALA	CA-C-N	-7.83	111.86	122.98
12	AB	128	ALA	C-N-CA	-7.83	111.86	122.98
55	o	30	ARG	NE-CZ-NH1	-7.64	113.86	121.50
17	C	73	ARG	NE-CZ-NH2	7.54	125.99	119.20
16	AG	33	THR	CA-C-N	7.49	135.84	121.54
16	AG	33	THR	C-N-CA	7.49	135.84	121.54
16	AG	102	PHE	CA-C-N	7.47	135.82	121.54
16	AG	102	PHE	C-N-CA	7.47	135.82	121.54
12	AB	159	PHE	CA-C-N	7.45	135.76	121.54
12	AB	159	PHE	C-N-CA	7.45	135.76	121.54
63	w	112	TYR	CZ-CE2-CD2	-7.38	106.32	119.60
16	AG	108	GLN	N-CA-C	-7.36	103.77	112.89
29	O	80	ARG	CG-CD-NE	7.35	128.18	112.00
23	I	134	MET	CB-CG-SD	-7.21	91.08	112.70
16	AG	426	GLY	O-C-N	-7.16	113.40	122.70
21	G	158	PRO	CA-N-CD	-7.11	102.05	112.00
62	v	96	ILE	CA-C-N	7.05	132.32	121.03
62	v	96	ILE	C-N-CA	7.05	132.32	121.03
21	G	19	GLN	CA-CB-CG	6.99	128.07	114.10
60	t	18	ARG	NE-CZ-NH1	-6.96	114.55	121.50
41	a	2334	U	OP2-P-O3'	-6.95	87.16	108.00
29	O	80	ARG	CB-CG-CD	-6.94	95.33	111.30
12	AB	159	PHE	CB-CA-C	6.86	124.07	110.42
16	AG	102	PHE	N-CA-C	6.77	125.21	110.80
18	D	37	U	N3-C4-O4	-6.63	99.50	119.40
16	AG	104	ARG	O-C-N	6.62	131.40	122.59
63	w	112	TYR	OH-CZ-CE2	-6.57	100.20	119.90
18	D	718	A	P-O5'-C5'	-6.55	111.07	120.90
31	Q	110	ILE	N-CA-C	-6.51	98.99	108.11
16	AG	106	THR	OG1-CB-CG2	-6.49	96.31	109.30
12	AB	150	ILE	N-CA-C	6.47	122.79	109.34
13	AD	289	LEU	CA-C-N	-6.45	109.01	121.58
13	AD	289	LEU	C-N-CA	-6.45	109.01	121.58
16	AG	355	ALA	CB-CA-C	6.41	123.19	110.42
18	D	827	U	N3-C4-O4	-6.41	100.16	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	a	1019	U	N3-C4-O4	-6.38	100.25	119.40
18	D	1358	U	N3-C4-O4	-6.36	100.33	119.40
16	AG	103	ASP	CA-C-N	6.34	133.66	121.54
16	AG	103	ASP	C-N-CA	6.34	133.66	121.54
37	W	40	ILE	CA-CB-CG1	-6.31	99.67	110.40
59	s	92	MET	CB-CG-SD	-6.31	93.78	112.70
12	AB	99	TYR	CA-C-N	6.29	137.15	121.80
12	AB	99	TYR	C-N-CA	6.29	137.15	121.80
16	AG	109	THR	N-CA-C	-6.28	104.52	111.36
16	AG	403	VAL	N-CA-C	-6.22	104.22	111.00
41	a	1019	U	C5-C4-O4	6.16	144.38	125.90
16	AG	108	GLN	CB-CA-C	-6.08	97.28	109.99
16	AG	404	GLU	N-CA-C	-6.08	104.96	112.38
16	AG	71	PRO	N-CA-C	6.07	122.62	113.81
12	AB	106	ASP	CA-C-N	6.07	127.43	119.84
12	AB	106	ASP	C-N-CA	6.07	127.43	119.84
12	AB	141	LEU	N-CA-C	6.05	123.69	110.80
16	AG	33	THR	CA-CB-CG2	6.01	120.72	110.50
29	O	103	PHE	CD1-CE1-CZ	-6.01	109.18	120.00
41	a	1082	U	N3-C4-O4	-5.96	101.53	119.40
41	a	2613	U	N3-C4-O4	-5.91	101.67	119.40
18	D	37	U	C5-C4-O4	5.89	143.57	125.90
41	a	1141	U	C5-C4-O4	5.89	143.56	125.90
54	n	176	PRO	N-CA-C	5.87	121.24	114.20
21	G	157	LEU	C-N-CD	-5.87	100.94	125.00
18	D	1358	U	C5-C4-O4	5.87	143.50	125.90
23	I	168	TYR	N-CA-CB	5.82	120.20	110.59
38	X	81	MET	CG-SD-CE	-5.82	88.10	100.90
16	AG	66	ASP	N-CA-C	-5.80	103.93	111.71
27	M	55	GLY	CA-C-N	5.79	132.60	121.54
27	M	55	GLY	C-N-CA	5.79	132.60	121.54
18	D	884	U	N3-C4-O4	-5.77	102.08	119.40
12	AB	152	HIS	N-CA-C	5.76	123.08	110.80
14	AE	853	THR	CA-C-N	5.75	131.02	122.74
14	AE	853	THR	C-N-CA	5.75	131.02	122.74
18	D	827	U	C5-C4-O4	5.72	143.07	125.90
21	G	154	MET	CA-CB-CG	-5.72	102.66	114.10
16	AG	220	VAL	N-CA-C	-5.71	108.28	113.71
23	I	134	MET	CG-SD-CE	-5.67	88.44	100.90
25	K	45	ARG	CA-C-N	5.64	130.36	122.23
25	K	45	ARG	C-N-CA	5.64	130.36	122.23
21	G	154	MET	CG-SD-CE	-5.64	88.49	100.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	K	45	ARG	CB-CG-CD	-5.64	98.34	111.30
9	9	130	PRO	CA-N-CD	-5.63	104.11	112.00
12	AB	107	PRO	O-C-N	5.62	130.22	122.64
16	AG	101	THR	CA-CB-OG1	5.57	117.96	109.60
60	t	18	ARG	CG-CD-NE	-5.56	99.77	112.00
41	a	1141	U	N3-C4-O4	-5.51	102.86	119.40
12	AB	151	LYS	N-CA-C	5.50	116.55	107.20
50	j	11	MET	CB-CG-SD	-5.49	96.22	112.70
14	AE	1184	ASP	N-CA-C	5.47	121.91	109.81
62	v	10	ARG	NE-CZ-NH1	-5.46	116.04	121.50
16	AG	355	ALA	N-CA-CB	5.44	119.69	110.49
16	AG	74	GLU	CB-CA-C	-5.43	101.78	110.14
9	9	129	LEU	C-N-CD	-5.42	102.77	125.00
16	AG	354	ALA	CA-C-N	-5.42	111.19	121.54
16	AG	354	ALA	C-N-CA	-5.42	111.19	121.54
63	w	38	LEU	N-CA-C	-5.37	105.16	112.35
31	Q	106	ARG	NE-CZ-NH1	-5.30	116.20	121.50
25	K	45	ARG	O-C-N	-5.28	116.78	123.01
8	7	10	U	C2'-C3'-O3'	5.26	117.40	109.50
39	Y	5	GLN	CB-CA-C	-5.25	99.97	110.42
16	AG	105	ILE	N-CA-CB	-5.24	102.58	111.23
16	AG	114	ILE	N-CA-C	-5.22	105.41	110.42
56	p	46	ALA	CA-C-N	5.21	131.50	121.54
56	p	46	ALA	C-N-CA	5.21	131.50	121.54
41	a	2613	U	C5-C4-O4	5.20	141.49	125.90
16	AG	107	THR	CA-C-N	-5.17	112.09	121.14
16	AG	107	THR	C-N-CA	-5.17	112.09	121.14
45	e	25	GLN	CG-CD-NE2	-5.16	108.66	116.40
39	Y	5	GLN	CA-CB-CG	5.16	124.42	114.10
16	AG	105	ILE	O-C-N	5.13	128.98	122.57
23	I	170	GLU	CB-CG-CD	5.13	121.31	112.60
16	AG	356	ILE	N-CA-C	-5.10	107.77	111.90
18	D	884	U	C5-C4-O4	5.10	141.20	125.90
21	G	158	PRO	N-CA-CB	-5.09	98.03	103.38
41	a	2334	U	C2'-C3'-O3'	-5.09	101.86	109.50
41	a	2602	A	C4'-C3'-O3'	5.09	117.04	109.40
14	AE	709	ARG	CA-C-N	5.08	135.45	126.45
14	AE	709	ARG	C-N-CA	5.08	135.45	126.45
13	AD	289	LEU	CA-C-O	-5.07	113.26	120.51
16	AG	107	THR	CA-C-O	-5.06	113.16	119.18
16	AG	65	VAL	CA-C-N	5.05	131.61	121.81
16	AG	65	VAL	C-N-CA	5.05	131.61	121.81

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	h	181	MET	CG-SD-CE	-5.02	89.86	100.90

There are no chirality outliers.

All (34) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
9	9	107	GLU	Peptide
9	9	79	PRO	Peptide
9	9	92	ALA	Peptide
12	AB	149	GLU	Mainchain
13	AC	192	VAL	Peptide
13	AD	20	SER	Peptide
13	AD	284	ARG	Mainchain
14	AE	1326	GLN	Peptide
14	AE	1344	LEU	Peptide
14	AE	313	GLY	Peptide
14	AE	416	ILE	Peptide
16	AG	102	PHE	Peptide
16	AG	104	ARG	Peptide,Mainchain
16	AG	11	ALA	Peptide
16	AG	292	ASP	Mainchain
16	AG	426	GLY	Mainchain
21	G	19	GLN	Sidechain
22	H	124	LEU	Peptide
22	H	171	ARG	Peptide
22	H	274	TYR	Peptide
22	H	81	GLU	Peptide
22	H	82	THR	Peptide
23	I	167	TRP	Mainchain
25	K	45	ARG	Mainchain
25	K	77	ASN	Peptide
29	O	12	ARG	Peptide
38	X	65	VAL	Peptide
39	Y	5	GLN	Peptide
54	n	176	PRO	Peptide
55	o	31	HIS	Peptide
59	s	74	TYR	Sidechain
61	u	35	HIS	Peptide
61	u	62	PRO	Peptide

## 5.2 Too-close contacts ⓘ

In the following table, the Non-H and H(model) columns list the number of non-hydrogen atoms and hydrogen atoms in the chain respectively. The H(added) column lists the number of hydrogen atoms added and optimized by MolProbity. The Clashes column lists the number of clashes within the asymmetric unit, whereas Symm-Clashes lists symmetry-related clashes.

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	0	816	839	839	53	0
2	1	857	922	922	73	0
3	2	746	811	811	60	0
4	3	788	844	844	44	0
5	4	753	780	780	87	0
6	5	726	0	392	112	0
7	6	703	0	396	35	0
8	7	527	0	262	82	0
9	9	1117	0	1155	254	0
10	A	1620	826	827	109	0
10	B	1620	813	827	111	0
11	AA	10381	0	10388	382	0
12	AB	1286	0	1302	375	0
13	AC	1698	0	1718	13	0
13	AD	2078	0	1891	37	0
14	AE	10404	0	10625	273	0
15	AF	650	0	658	10	0
16	AG	3852	0	3828	899	0
17	C	544	559	560	110	0
18	D	32703	16423	16451	744	0
19	E	669	719	719	79	0
20	F	589	629	629	60	0
21	G	1760	1785	1787	152	0
22	H	1730	1454	1455	164	0
23	I	1636	1710	1710	129	0
24	J	1643	1707	1707	88	0
25	K	1152	1196	1196	119	0
26	L	848	846	846	105	0
27	M	1181	1235	1238	123	0
28	N	979	1031	1031	91	0
29	O	1022	1070	1070	92	0
30	P	790	831	831	94	0
31	Q	877	887	887	96	0
32	R	939	1001	1001	94	0
33	S	805	844	844	74	0
34	T	714	734	734	93	0
35	U	649	666	666	62	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
36	V	648	691	691	71	0
37	W	663	688	688	46	0
38	X	900	964	965	111	0
39	Y	1032	0	1088	211	0
40	Z	227	0	237	30	0
41	a	61841	31077	31120	1314	0
42	b	582	599	599	48	0
43	c	625	652	652	32	0
44	d	2569	1301	1301	43	0
45	e	501	531	531	43	0
46	f	448	488	488	37	0
47	g	522	520	520	65	0
48	h	2082	2154	2154	136	0
49	i	444	459	458	47	0
50	j	1565	1617	1616	121	0
51	k	426	464	464	42	0
52	l	1552	1619	1619	126	0
53	m	377	418	418	29	0
54	n	1410	1443	1444	152	0
55	o	504	572	572	53	0
56	p	1313	1358	1358	74	0
57	q	302	343	343	27	0
58	r	1111	1148	1148	51	0
59	s	1129	1162	1162	123	0
60	t	946	1023	1023	107	0
61	u	1053	1129	1129	78	0
62	v	1074	1157	1157	124	0
63	w	951	994	994	98	0
64	x	892	923	923	48	0
65	y	917	962	962	54	0
66	z	947	1020	1019	82	0
67	AE	1	0	0	0	0
68	AE	2	0	0	3	0
All	All	181408	98638	132670	7674	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 25.

All (7674) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5:11:DA:C5'	12:AB:67:THR:HG22	1.20	1.62
25:K:45:ARG:CD	25:K:45:ARG:CG	1.75	1.58
66:z:74:ILE:CG1	66:z:74:ILE:CD1	1.81	1.58
16:AG:425:LEU:CD2	16:AG:429:LYS:HE2	1.21	1.57
6:5:10:DT:C7	11:AA:62:TYR:CD2	1.84	1.57
16:AG:287:ALA:CA	16:AG:331:LEU:HD12	1.18	1.56
6:5:9:DG:H5''	11:AA:473:ARG:CB	1.29	1.55
6:5:9:DG:C5'	11:AA:473:ARG:HB3	1.34	1.55
11:AA:846:GLY:C	16:AG:104:ARG:HH12	1.02	1.54
16:AG:287:ALA:HA	16:AG:331:LEU:CD1	1.38	1.53
7:6:23:DC:C5'	14:AE:259:ARG:HH12	1.21	1.53
11:AA:855:PRO:HB2	16:AG:35:THR:CB	1.38	1.53
16:AG:393:LEU:CB	16:AG:398:LEU:CD2	1.83	1.53
16:AG:168:LEU:CD2	16:AG:231:GLY:HA3	1.34	1.52
12:AB:81:PHE:CD2	14:AE:293:ARG:NH1	1.77	1.51
11:AA:855:PRO:HB2	16:AG:35:THR:CG2	1.37	1.50
6:5:10:DT:C6	11:AA:62:TYR:CE2	2.00	1.50
6:5:10:DT:C5	11:AA:62:TYR:CD2	1.95	1.50
16:AG:287:ALA:CB	16:AG:331:LEU:HD12	1.41	1.50
11:AA:887:VAL:CB	16:AG:105:ILE:HD12	1.42	1.50
16:AG:363:LEU:CD2	16:AG:410:ALA:H	1.24	1.49
11:AA:861:ALA:H	16:AG:38:LYS:NZ	1.00	1.49
14:AE:79:LYS:CA	16:AG:144:LYS:HZ2	1.28	1.47
16:AG:425:LEU:CA	16:AG:429:LYS:HE3	1.40	1.47
14:AE:79:LYS:HA	16:AG:144:LYS:CE	1.42	1.47
14:AE:72:CYS:SG	68:AE:1502:ZN:ZN	1.03	1.46
16:AG:187:ARG:HB3	16:AG:188:PRO:CD	1.45	1.46
16:AG:425:LEU:HD23	16:AG:429:LYS:CE	1.45	1.45
6:5:10:DT:C5	11:AA:62:TYR:CE2	2.04	1.44
16:AG:168:LEU:HD22	16:AG:231:GLY:CA	1.49	1.43
12:AB:115:LYS:NZ	12:AB:129:ILE:CG2	1.82	1.42
16:AG:363:LEU:CD2	16:AG:410:ALA:N	1.82	1.42
12:AB:81:PHE:CE1	14:AE:293:ARG:CB	1.92	1.42
6:5:11:DA:P	12:AB:68:THR:HA	1.58	1.42
16:AG:168:LEU:CD2	16:AG:231:GLY:CA	1.98	1.41
11:AA:380:ALA:N	12:AB:65:HIS:CE1	1.84	1.40
14:AE:79:LYS:CA	16:AG:144:LYS:NZ	1.81	1.39
11:AA:887:VAL:HB	16:AG:105:ILE:CD1	1.52	1.39
16:AG:393:LEU:CD2	16:AG:398:LEU:CD2	1.99	1.38
12:AB:115:LYS:NZ	12:AB:129:ILE:HG21	1.05	1.37
11:AA:846:GLY:C	16:AG:104:ARG:NH1	1.82	1.37
6:5:9:DG:OP2	11:AA:473:ARG:CB	1.73	1.36

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:437:LEU:HD21	16:AG:456:LEU:CD1	1.54	1.35
11:AA:858:GLY:HA3	16:AG:38:LYS:CB	1.40	1.34
16:AG:393:LEU:CA	16:AG:398:LEU:CD2	2.05	1.34
16:AG:434:LEU:HD22	16:AG:459:LEU:CD1	1.58	1.33
16:AG:385:ALA:CB	16:AG:411:LYS:HE3	1.56	1.33
16:AG:387:VAL:HG12	16:AG:388:PRO:CD	1.57	1.32
16:AG:363:LEU:HD23	16:AG:410:ALA:N	1.38	1.31
11:AA:856:ASN:N	16:AG:35:THR:HB	1.42	1.31
16:AG:393:LEU:CA	16:AG:398:LEU:HD22	1.59	1.31
12:AB:115:LYS:CE	12:AB:129:ILE:HG21	1.61	1.31
16:AG:434:LEU:CD2	16:AG:456:LEU:HA	1.58	1.31
12:AB:51:PHE:HE2	14:AE:288:PRO:CG	1.43	1.30
14:AE:79:LYS:HA	16:AG:144:LYS:NZ	1.03	1.30
12:AB:82:GLY:CA	14:AE:142:GLU:HB2	1.62	1.30
16:AG:448:LEU:HD21	16:AG:473:LEU:CD1	1.61	1.30
14:AE:79:LYS:CB	16:AG:144:LYS:HZ2	1.44	1.30
16:AG:393:LEU:CD2	16:AG:398:LEU:HD21	1.57	1.30
14:AE:287:ALA:CB	14:AE:292:VAL:HG13	1.61	1.30
11:AA:377:THR:OG1	12:AB:62:GLU:CB	1.80	1.29
6:5:11:DA:OP2	12:AB:68:THR:CA	1.81	1.29
11:AA:858:GLY:CA	16:AG:38:LYS:HB2	1.62	1.28
16:AG:434:LEU:HA	16:AG:456:LEU:CD2	1.64	1.27
11:AA:387:ASN:CG	11:AA:394:ARG:NH1	1.92	1.27
11:AA:377:THR:OG1	12:AB:62:GLU:HB3	1.20	1.27
11:AA:861:ALA:N	16:AG:38:LYS:HZ1	1.31	1.27
16:AG:362:TYR:O	16:AG:413:ALA:HB2	1.35	1.27
6:5:11:DA:C3'	12:AB:67:THR:HG21	1.63	1.26
16:AG:353:HIS:O	16:AG:356:ILE:HG23	1.13	1.26
6:5:11:DA:C5'	12:AB:67:THR:CG2	2.12	1.26
11:AA:846:GLY:HA3	16:AG:104:ARG:NH2	1.48	1.26
12:AB:81:PHE:HA	14:AE:141:PHE:O	1.27	1.26
11:AA:856:ASN:ND2	16:AG:35:THR:HG22	1.47	1.25
11:AA:887:VAL:CB	16:AG:105:ILE:CD1	2.11	1.25
16:AG:393:LEU:CG	16:AG:398:LEU:HD21	1.63	1.25
12:AB:115:LYS:CD	12:AB:129:ILE:HG13	1.67	1.25
6:5:10:DT:C1'	12:AB:71:ALA:H	1.46	1.25
16:AG:425:LEU:HA	16:AG:429:LYS:CE	1.64	1.25
8:7:11:U:H4'	25:K:20:ARG:NH2	1.51	1.25
14:AE:288:PRO:HD2	14:AE:291:ILE:CG2	1.66	1.25
11:AA:481:LEU:CD2	12:AB:16:ARG:CD	2.01	1.25
11:AA:856:ASN:OD1	16:AG:35:THR:HA	1.08	1.25

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:123:ARG:HH12	16:AG:191:ARG:CA	1.36	1.24
16:AG:387:VAL:CG1	16:AG:388:PRO:HD2	1.67	1.24
16:AG:433:ASP:O	16:AG:456:LEU:HD11	1.35	1.23
12:AB:51:PHE:CE2	14:AE:288:PRO:CG	2.22	1.23
12:AB:81:PHE:CE1	14:AE:293:ARG:HB3	1.32	1.23
11:AA:855:PRO:CB	16:AG:35:THR:CB	2.16	1.23
16:AG:363:LEU:HD23	16:AG:409:ARG:C	1.63	1.23
16:AG:287:ALA:CB	16:AG:331:LEU:CD1	2.16	1.23
11:AA:861:ALA:N	16:AG:38:LYS:NZ	1.84	1.23
16:AG:355:ALA:CB	16:AG:382:GLU:CD	2.10	1.23
16:AG:425:LEU:CG	16:AG:429:LYS:HE2	1.69	1.22
6:5:10:DT:H71	11:AA:62:TYR:CG	1.75	1.22
16:AG:353:HIS:C	16:AG:356:ILE:HG23	1.62	1.22
16:AG:393:LEU:HB3	16:AG:398:LEU:CD2	1.51	1.21
12:AB:119:THR:OG1	12:AB:126:PHE:HE2	1.23	1.20
16:AG:123:ARG:NH1	16:AG:191:ARG:C	1.98	1.20
6:5:11:DA:OP2	12:AB:68:THR:N	1.73	1.20
7:6:25:DT:OP1	11:AA:496:LYS:CA	1.80	1.19
16:AG:442:ARG:O	16:AG:446:PHE:CD2	1.95	1.19
12:AB:115:LYS:HD3	12:AB:129:ILE:CD1	1.73	1.19
11:AA:853:ASP:O	16:AG:103:ASP:OD1	1.58	1.19
16:AG:381:LEU:HA	16:AG:384:LEU:HD12	1.21	1.19
16:AG:355:ALA:HA	16:AG:382:GLU:OE2	1.37	1.18
11:AA:887:VAL:CA	16:AG:105:ILE:HD12	1.73	1.18
16:AG:393:LEU:HD22	16:AG:398:LEU:CD1	1.73	1.18
16:AG:437:LEU:CD2	16:AG:456:LEU:HD13	1.74	1.18
11:AA:855:PRO:CB	16:AG:35:THR:OG1	1.90	1.18
11:AA:856:ASN:N	16:AG:35:THR:CB	2.02	1.18
16:AG:353:HIS:O	16:AG:357:ASP:N	1.77	1.18
6:5:9:DG:C5'	11:AA:473:ARG:CG	2.21	1.17
11:AA:887:VAL:C	16:AG:105:ILE:CD1	2.16	1.17
58:r:135:HIS:ND1	58:r:137:GLU:OE1	1.77	1.17
7:6:23:DC:C5'	14:AE:259:ARG:NH1	2.05	1.17
11:AA:855:PRO:C	16:AG:35:THR:HB	1.68	1.17
16:AG:355:ALA:HB2	16:AG:382:GLU:OE1	1.40	1.17
16:AG:355:ALA:HA	16:AG:382:GLU:CD	1.71	1.16
56:p:159:GLY:O	56:p:163:ARG:NH1	1.77	1.16
38:X:81:MET:HE1	38:X:92:ARG:HB3	1.20	1.16
6:5:10:DT:H73	11:AA:62:TYR:CD2	1.77	1.15
6:5:9:DG:OP2	11:AA:473:ARG:HB2	0.97	1.15
6:5:10:DT:H1'	12:AB:71:ALA:N	1.21	1.15

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:285:ILE:HG23	16:AG:293:VAL:HG12	1.25	1.15
11:AA:887:VAL:C	16:AG:105:ILE:HD12	1.72	1.14
16:AG:393:LEU:CB	16:AG:398:LEU:HD21	1.58	1.14
12:AB:35:LEU:HA	12:AB:106:ASP:OD1	1.47	1.14
12:AB:140:MET:HE2	30:P:102:LEU:HG	1.18	1.14
16:AG:393:LEU:HD22	16:AG:398:LEU:CD2	1.71	1.14
12:AB:115:LYS:HD2	12:AB:129:ILE:HG13	1.13	1.13
14:AE:287:ALA:HB2	14:AE:292:VAL:CG1	1.78	1.13
6:5:9:DG:C5'	11:AA:473:ARG:CB	2.02	1.13
6:5:9:DG:H5'	11:AA:473:ARG:CG	1.76	1.12
6:5:11:DA:OP2	12:AB:68:THR:HA	1.36	1.12
16:AG:123:ARG:NH1	16:AG:191:ARG:O	1.80	1.13
16:AG:393:LEU:HD22	16:AG:398:LEU:HD11	1.28	1.13
16:AG:430:PRO:HG2	16:AG:435:LEU:HD21	1.29	1.13
36:V:7:THR:OG1	36:V:60:GLU:OE2	1.66	1.12
12:AB:115:LYS:HA	12:AB:129:ILE:HD12	1.28	1.12
10:B:75:C:OP2	41:a:2602:A:O2'	1.67	1.12
12:AB:115:LYS:CE	12:AB:129:ILE:CG2	2.25	1.12
16:AG:393:LEU:CD2	16:AG:398:LEU:CD1	2.27	1.12
6:5:10:DT:C7	11:AA:62:TYR:CG	2.31	1.12
10:A:76:A:O2'	41:a:2394:C:N3	1.81	1.12
16:AG:279:ASN:OD1	16:AG:280:PRO:HD3	1.50	1.12
6:5:10:DT:O3'	12:AB:67:THR:O	1.67	1.12
12:AB:102:LYS:CE	14:AE:285:LEU:HG	1.78	1.12
41:a:2335:A:OP2	64:x:13:ARG:NE	1.81	1.12
16:AG:355:ALA:HB1	16:AG:382:GLU:CG	1.77	1.11
11:AA:846:GLY:CA	16:AG:104:ARG:HH22	1.63	1.11
37:W:40:ILE:HD13	37:W:66:MET:SD	1.89	1.11
58:r:135:HIS:HD1	58:r:137:GLU:CD	1.59	1.11
11:AA:380:ALA:N	12:AB:65:HIS:HE1	1.28	1.11
12:AB:54:TYR:CE2	14:AE:291:ILE:HG23	1.84	1.11
16:AG:187:ARG:HB3	16:AG:188:PRO:HD2	1.22	1.11
16:AG:353:HIS:O	16:AG:356:ILE:CG2	1.98	1.11
16:AG:422:GLU:HA	16:AG:426:GLY:CA	1.79	1.11
16:AG:425:LEU:CD2	16:AG:429:LYS:CE	2.13	1.11
9:9:50:VAL:HG22	39:Y:119:ALA:HB3	1.32	1.10
16:AG:424:SER:O	16:AG:429:LYS:HG2	1.47	1.10
16:AG:434:LEU:CD2	16:AG:459:LEU:HD12	1.81	1.10
12:AB:28:CYS:HA	12:AB:56:PHE:O	1.52	1.10
16:AG:393:LEU:HB3	16:AG:398:LEU:HD23	1.26	1.10
11:AA:887:VAL:HB	16:AG:105:ILE:CG1	1.81	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:285:ILE:CG1	16:AG:293:VAL:HG11	1.79	1.10
46:f:4:THR:OG1	46:f:37:GLU:OE2	1.68	1.10
16:AG:393:LEU:HA	16:AG:398:LEU:CD2	1.69	1.09
7:6:25:DT:C3'	11:AA:496:LYS:HG3	1.74	1.09
12:AB:82:GLY:HA2	14:AE:142:GLU:HB2	1.13	1.09
12:AB:115:LYS:HD3	12:AB:129:ILE:HD12	1.31	1.09
16:AG:425:LEU:HA	16:AG:429:LYS:CG	1.80	1.09
39:Y:5:GLN:HG2	39:Y:61:TYR:CE1	1.87	1.09
16:AG:181:GLY:HA2	16:AG:204:MET:SD	1.92	1.09
16:AG:287:ALA:CA	16:AG:331:LEU:CD1	2.02	1.09
16:AG:363:LEU:HD21	16:AG:410:ALA:H	0.94	1.09
16:AG:434:LEU:HD21	16:AG:456:LEU:HA	1.26	1.09
11:AA:855:PRO:CB	16:AG:35:THR:CG2	2.30	1.09
11:AA:855:PRO:CB	16:AG:35:THR:HG21	1.82	1.09
14:AE:288:PRO:HD2	14:AE:291:ILE:HG21	1.10	1.08
16:AG:123:ARG:HH12	16:AG:191:ARG:C	1.57	1.08
11:AA:469:VAL:O	11:AA:473:ARG:HG2	1.53	1.08
11:AA:887:VAL:CG2	16:AG:105:ILE:HG13	1.84	1.08
12:AB:51:PHE:CZ	14:AE:288:PRO:HD3	1.88	1.08
16:AG:168:LEU:HD21	16:AG:231:GLY:CA	1.82	1.08
6:5:9:DG:C5'	11:AA:473:ARG:HG3	1.82	1.08
8:7:11:U:C4'	25:K:20:ARG:NH2	2.16	1.07
11:AA:914:LYS:N	16:AG:105:ILE:HA	1.68	1.07
12:AB:81:PHE:CE2	14:AE:293:ARG:NH1	1.88	1.07
16:AG:425:LEU:HA	16:AG:429:LYS:HG3	1.33	1.07
16:AG:451:ARG:HH12	16:AG:467:LEU:C	1.62	1.07
16:AG:302:LYS:H	16:AG:302:LYS:HE3	1.16	1.07
16:AG:451:ARG:NH1	16:AG:467:LEU:HA	1.68	1.07
12:AB:115:LYS:CD	12:AB:129:ILE:CG1	2.33	1.07
16:AG:434:LEU:CD2	16:AG:456:LEU:CA	2.33	1.07
7:6:23:DC:H5''	14:AE:259:ARG:HH12	1.14	1.07
20:F:37:PHE:HB2	31:Q:126:LYS:HE2	1.35	1.07
26:L:9:MET:HE1	26:L:85:ILE:HB	1.17	1.07
39:Y:14:ALA:HB1	39:Y:50:LYS:HD2	1.37	1.07
6:5:11:DA:H5'	12:AB:67:THR:HG22	1.13	1.07
11:AA:855:PRO:HB2	16:AG:35:THR:HG21	1.15	1.07
9:9:46:ARG:HG3	9:9:94:ARG:HH21	1.12	1.06
11:AA:856:ASN:CG	16:AG:35:THR:HA	1.80	1.06
27:M:4:ARG:HG3	27:M:5:ARG:HG3	1.33	1.06
11:AA:387:ASN:CG	11:AA:394:ARG:HH11	1.55	1.06
16:AG:393:LEU:CB	16:AG:398:LEU:HD23	1.80	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:96:LYS:HE3	39:Y:136:GLY:HA3	1.29	1.06
6:5:9:DG:H5'	11:AA:473:ARG:HG3	1.08	1.06
12:AB:82:GLY:HA2	14:AE:142:GLU:CB	1.85	1.06
53:m:22:MET:O	53:m:28:ARG:NH1	1.87	1.06
61:u:122:VAL:HG12	61:u:125:LEU:HD21	1.35	1.06
7:6:23:DC:O5'	14:AE:259:ARG:NH1	1.88	1.06
12:AB:102:LYS:HE2	14:AE:285:LEU:HG	1.07	1.06
16:AG:434:LEU:HA	16:AG:456:LEU:HD21	1.25	1.06
16:AG:425:LEU:CG	16:AG:429:LYS:CE	2.34	1.06
9:9:136:ILE:HD12	9:9:139:LEU:HD12	1.38	1.05
11:AA:847:PRO:HD2	16:AG:104:ARG:NH2	1.69	1.05
16:AG:365:ILE:HD13	16:AG:406:LEU:CD2	1.85	1.05
6:5:10:DT:H71	11:AA:62:TYR:CD2	1.77	1.05
16:AG:385:ALA:CB	16:AG:411:LYS:CE	2.33	1.05
12:AB:115:LYS:HD3	12:AB:129:ILE:CG1	1.87	1.05
16:AG:187:ARG:HB3	16:AG:188:PRO:HD3	1.32	1.05
16:AG:285:ILE:HG23	16:AG:293:VAL:CG1	1.87	1.05
18:D:1309:G:OP2	38:X:98:ARG:NE	1.88	1.05
9:9:103:ASN:ND2	9:9:107:GLU:OE2	1.90	1.04
11:AA:887:VAL:HG21	16:AG:105:ILE:HG13	1.39	1.04
16:AG:285:ILE:HG13	16:AG:293:VAL:CG1	1.86	1.04
6:5:11:DA:H3'	12:AB:67:THR:HG21	1.05	1.04
11:AA:856:ASN:OD1	16:AG:35:THR:CA	2.04	1.04
50:j:184:ARG:NH2	65:y:7:GLN:OE1	1.90	1.04
11:AA:481:LEU:HD23	12:AB:16:ARG:CD	1.87	1.04
16:AG:434:LEU:HD23	16:AG:456:LEU:HD23	1.10	1.04
16:AG:434:LEU:HD23	16:AG:456:LEU:HA	1.34	1.04
16:AG:448:LEU:CD2	16:AG:473:LEU:CD1	2.36	1.04
18:D:564:C:OP2	32:R:12:ARG:NH2	1.91	1.04
16:AG:365:ILE:HD13	16:AG:406:LEU:HD22	1.37	1.03
16:AG:385:ALA:HB1	16:AG:411:LYS:HE3	1.35	1.03
20:F:25:LYS:NZ	22:H:336:ASP:OD1	1.90	1.03
26:L:9:MET:HE1	26:L:85:ILE:CB	1.86	1.03
52:l:1:MET:HE2	52:l:16:GLU:HA	1.40	1.03
9:9:31:ARG:NH1	41:a:1054:A:O4'	1.91	1.03
12:AB:51:PHE:HE2	14:AE:288:PRO:HG2	1.20	1.03
16:AG:363:LEU:HG	16:AG:410:ALA:HB2	1.37	1.03
16:AG:434:LEU:HD21	16:AG:456:LEU:CA	1.89	1.03
9:9:48:ALA:HB3	9:9:51:TYR:HE1	1.20	1.03
47:g:58:ASP:OD1	47:g:62:LYS:NZ	1.90	1.03
56:p:149:ARG:NH2	56:p:152:ARG:O	1.91	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:387:VAL:HG12	16:AG:388:PRO:HD2	1.06	1.03
8:7:4:U:O2'	18:D:1403:C:O2	1.75	1.02
12:AB:81:PHE:CA	14:AE:141:PHE:O	2.06	1.02
23:I:152:GLU:OE2	23:I:154:SER:OG	1.77	1.02
6:5:11:DA:P	12:AB:68:THR:CA	2.44	1.02
7:6:18:DT:H3	8:7:52:A:N6	1.55	1.02
16:AG:393:LEU:HD22	16:AG:398:LEU:HD21	1.25	1.02
16:AG:422:GLU:HA	16:AG:426:GLY:N	1.74	1.02
16:AG:422:GLU:HA	16:AG:426:GLY:HA3	1.33	1.02
16:AG:425:LEU:C	16:AG:429:LYS:HE3	1.84	1.02
26:L:42:TRP:CH2	26:L:102:MET:HG3	1.94	1.02
50:j:179:ARG:NH1	50:j:180:VAL:O	1.92	1.02
12:AB:102:LYS:HE2	14:AE:285:LEU:CG	1.90	1.02
23:I:123:GLN:OE1	23:I:126:ARG:NH2	1.91	1.02
7:6:25:DT:H3'	11:AA:496:LYS:HG3	1.41	1.02
65:y:51:ARG:NH1	65:y:56:HIS:O	1.91	1.02
14:AE:287:ALA:HB2	14:AE:292:VAL:HG13	1.08	1.02
16:AG:287:ALA:HA	16:AG:331:LEU:HD11	1.41	1.02
16:AG:365:ILE:CG2	16:AG:409:ARG:NH1	2.23	1.02
39:Y:93:ASN:HA	39:Y:135:MET:CE	1.89	1.02
8:7:3:G:H5''	18:D:1500:A:N6	1.75	1.01
12:AB:82:GLY:CA	14:AE:142:GLU:CB	2.38	1.01
16:AG:355:ALA:HB1	16:AG:382:GLU:HG3	1.41	1.01
16:AG:361:LYS:CE	16:AG:417:ILE:HG12	1.90	1.01
21:G:49:MET:HE1	21:G:201:PRO:HD3	1.42	1.01
8:7:11:U:O2'	25:K:33:PHE:CE1	2.10	1.01
11:AA:889:PRO:HG3	16:AG:104:ARG:HD3	1.39	1.01
39:Y:56:VAL:HA	39:Y:71:LYS:HZ1	1.22	1.01
59:s:72:LYS:HD3	59:s:74:TYR:OH	1.59	1.01
9:9:3:LEU:HD12	9:9:4:ASN:H	1.25	1.01
9:9:43:LYS:HG3	9:9:46:ARG:HH12	1.21	1.01
18:D:377:G:OP1	35:U:5:ARG:NH1	1.94	1.01
16:AG:361:LYS:HE2	16:AG:417:ILE:HG12	1.39	1.01
16:AG:363:LEU:HD23	16:AG:409:ARG:CA	1.89	1.01
16:AG:393:LEU:CD2	16:AG:398:LEU:HD11	1.90	1.01
12:AB:148:LYS:HZ2	12:AB:150:ILE:CD1	1.73	1.00
16:AG:363:LEU:CD2	16:AG:409:ARG:HB2	1.91	1.00
16:AG:393:LEU:HD23	16:AG:398:LEU:HD22	1.43	1.00
16:AG:425:LEU:CA	16:AG:429:LYS:CE	2.30	1.00
52:l:1:MET:HE1	52:l:20:GLY:HA3	1.43	1.00
9:9:30:SER:O	9:9:31:ARG:NE	1.94	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:187:ARG:CB	16:AG:188:PRO:CD	2.37	1.00
9:9:88:HIS:HB3	9:9:89:PRO:HD3	1.44	1.00
16:AG:393:LEU:CG	16:AG:398:LEU:CD2	2.29	1.00
6:5:11:DA:H3'	12:AB:67:THR:CG2	1.91	0.99
11:AA:887:VAL:HG11	16:AG:105:ILE:HG21	1.41	0.99
14:AE:70:CYS:SG	14:AE:85:CYS:SG	2.60	0.99
16:AG:434:LEU:HD23	16:AG:456:LEU:CD2	1.91	0.99
16:AG:442:ARG:O	16:AG:446:PHE:HD2	1.34	0.99
26:L:42:TRP:HH2	26:L:102:MET:HG3	1.25	0.99
6:5:10:DT:C1'	12:AB:71:ALA:N	2.10	0.99
16:AG:362:TYR:O	16:AG:413:ALA:CB	2.10	0.99
14:AE:70:CYS:SG	14:AE:72:CYS:SG	2.61	0.99
16:AG:131:ARG:O	16:AG:134:GLU:HG3	1.61	0.99
16:AG:168:LEU:HD21	16:AG:231:GLY:N	1.76	0.99
16:AG:393:LEU:HD22	16:AG:398:LEU:CG	1.93	0.99
16:AG:434:LEU:CD2	16:AG:456:LEU:HD23	1.92	0.99
39:Y:60:VAL:HG22	39:Y:66:PHE:HB3	1.43	0.99
16:AG:168:LEU:HD22	16:AG:231:GLY:HA2	1.40	0.99
16:AG:336:LEU:HD12	16:AG:336:LEU:H	1.26	0.99
16:AG:437:LEU:CD2	16:AG:456:LEU:CD1	2.34	0.99
6:5:11:DA:H5'	12:AB:67:THR:CG2	1.81	0.98
11:AA:481:LEU:HD21	12:AB:16:ARG:CD	1.60	0.98
14:AE:288:PRO:CD	14:AE:291:ILE:HG21	1.92	0.98
14:AE:79:LYS:HA	16:AG:144:LYS:HE3	1.40	0.98
58:r:38:PRO:O	58:r:43:ASN:ND2	1.96	0.98
6:5:10:DT:O5'	11:AA:62:TYR:OH	1.80	0.98
14:AE:79:LYS:HE2	16:AG:153:ASP:OD1	1.63	0.98
21:G:154:MET:HE2	21:G:158:PRO:HD3	1.44	0.98
6:5:11:DA:H5''	12:AB:67:THR:CG2	1.86	0.98
11:AA:387:ASN:ND2	11:AA:394:ARG:HH12	1.61	0.98
14:AE:287:ALA:CB	14:AE:292:VAL:CG1	2.38	0.98
7:6:25:DT:OP1	11:AA:496:LYS:HA	1.12	0.98
16:AG:228:ARG:HA	16:AG:327:LEU:HD11	1.46	0.97
16:AG:355:ALA:HB2	16:AG:382:GLU:CD	1.83	0.97
12:AB:119:THR:OG1	12:AB:126:PHE:CE2	2.13	0.97
33:S:86:GLU:OE1	33:S:90:ARG:NH1	1.96	0.97
39:Y:93:ASN:HA	39:Y:135:MET:HE1	1.00	0.97
14:AE:79:LYS:CA	16:AG:144:LYS:CE	2.32	0.97
16:AG:279:ASN:CG	16:AG:280:PRO:CD	2.38	0.97
6:5:10:DT:C7	11:AA:62:TYR:HD2	1.69	0.97
18:D:1344:C:OP1	29:O:124:ARG:NH1	1.98	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:S:92:GLU:OE1	33:S:92:GLU:N	1.98	0.97
39:Y:93:ASN:CA	39:Y:135:MET:HE1	1.94	0.97
59:s:12:LYS:NZ	59:s:14:ASP:OD1	1.97	0.97
16:AG:214:PRO:O	16:AG:218:GLU:OE2	1.83	0.97
16:AG:380:THR:O	16:AG:384:LEU:CD1	2.13	0.97
16:AG:424:SER:O	16:AG:429:LYS:CG	2.13	0.97
40:Z:14:MET:HB3	40:Z:17:MET:HG2	1.46	0.97
16:AG:168:LEU:HD21	16:AG:230:PRO:C	1.89	0.96
41:a:2093:G:OP1	58:r:22:LYS:NZ	1.98	0.96
6:5:11:DA:H5''	12:AB:67:THR:HG22	0.97	0.96
12:AB:51:PHE:CE2	14:AE:288:PRO:CD	2.48	0.96
16:AG:287:ALA:HB2	16:AG:331:LEU:CD1	1.92	0.96
6:5:11:DA:C3'	12:AB:67:THR:CG2	2.42	0.96
11:AA:846:GLY:CA	16:AG:104:ARG:NH2	2.25	0.96
18:D:562:U:O2'	32:R:14:ARG:NH1	1.99	0.96
9:9:95:LEU:HD13	9:9:98:GLU:OE2	1.65	0.96
16:AG:393:LEU:CA	16:AG:398:LEU:HD23	1.88	0.96
23:I:164:ARG:NH2	23:I:166:GLU:OE2	1.99	0.96
11:AA:846:GLY:CA	16:AG:104:ARG:HH12	1.78	0.96
16:AG:433:ASP:O	16:AG:456:LEU:CD1	2.12	0.95
39:Y:5:GLN:HG2	39:Y:61:TYR:HE1	1.31	0.95
12:AB:115:LYS:HZ2	12:AB:129:ILE:CG2	1.58	0.95
39:Y:75:ALA:HB2	39:Y:112:LYS:HE3	1.48	0.95
64:x:111:ARG:NH2	64:x:117:PHE:OXT	1.99	0.95
14:AE:85:CYS:HG	68:AE:1502:ZN:ZN	0.75	0.95
16:AG:365:ILE:CD1	16:AG:406:LEU:HD22	1.97	0.95
8:7:2:U:P	18:D:1505:G:H8	1.88	0.95
12:AB:6:LEU:HD13	14:AE:291:ILE:HD11	1.46	0.95
16:AG:448:LEU:HD21	16:AG:473:LEU:HD13	1.48	0.95
12:AB:51:PHE:CE2	14:AE:288:PRO:HG3	1.98	0.95
12:AB:140:MET:O	12:AB:150:ILE:HB	1.67	0.95
36:V:7:THR:OG1	36:V:61:ILE:O	1.85	0.95
9:9:31:ARG:HD2	41:a:1053:C:O2'	1.66	0.95
11:AA:861:ALA:H	16:AG:38:LYS:HZ3	1.14	0.94
16:AG:440:VAL:CG2	16:AG:481:LEU:HD22	1.97	0.94
12:AB:103:ASP:HB3	12:AB:105:VAL:HG23	1.47	0.94
12:AB:145:LEU:HD11	30:P:84:VAL:HG12	1.47	0.94
14:AE:79:LYS:HA	16:AG:144:LYS:HZ1	1.31	0.94
9:9:88:HIS:HB3	9:9:89:PRO:CD	1.97	0.94
11:AA:887:VAL:HB	16:AG:105:ILE:HD12	0.97	0.94
9:9:23:LEU:HD13	9:9:92:ALA:HB1	1.49	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:846:GLY:HA3	16:AG:104:ARG:HH22	1.12	0.94
16:AG:393:LEU:HA	16:AG:398:LEU:HD22	0.96	0.94
41:a:30:G:OP2	66:z:5:LYS:NZ	2.00	0.94
16:AG:258:ARG:HH11	16:AG:258:ARG:HG2	1.31	0.94
1:0:18:GLN:N	1:0:18:GLN:OE1	2.01	0.94
9:9:61:ARG:NH1	41:a:1047:G:N7	2.15	0.94
11:AA:481:LEU:HD21	12:AB:16:ARG:HD3	1.47	0.94
11:AA:846:GLY:HA3	16:AG:104:ARG:CZ	1.97	0.94
11:AA:887:VAL:CG2	16:AG:105:ILE:CD1	2.45	0.94
12:AB:82:GLY:N	14:AE:142:GLU:CB	2.30	0.94
16:AG:393:LEU:HD23	16:AG:398:LEU:CD2	1.93	0.94
26:L:9:MET:CE	26:L:85:ILE:HB	1.97	0.94
56:p:38:ASN:ND2	56:p:64:GLN:OE1	2.01	0.94
11:AA:858:GLY:HA3	16:AG:38:LYS:HB2	0.94	0.94
39:Y:33:ASN:HB3	39:Y:36:GLU:HG2	1.48	0.94
9:9:58:THR:HG21	9:9:81:LEU:HD23	1.48	0.94
16:AG:451:ARG:NH1	16:AG:467:LEU:CA	2.30	0.94
9:9:50:VAL:HG22	39:Y:119:ALA:CB	1.98	0.93
14:AE:79:LYS:CG	16:AG:144:LYS:HZ2	1.81	0.93
9:9:3:LEU:HD12	9:9:4:ASN:N	1.82	0.93
16:AG:393:LEU:CD2	16:AG:398:LEU:HD22	1.89	0.93
6:5:11:DA:OP2	12:AB:67:THR:C	2.11	0.93
16:AG:171:GLU:OE2	16:AG:267:GLY:HA3	1.68	0.93
16:AG:279:ASN:CG	16:AG:280:PRO:HD3	1.93	0.93
6:5:9:DG:OP2	11:AA:473:ARG:CG	2.16	0.93
6:5:10:DT:C6	11:AA:62:TYR:HE2	1.85	0.93
11:AA:858:GLY:CA	16:AG:38:LYS:CB	2.27	0.93
16:AG:387:VAL:HG12	16:AG:388:PRO:HD3	1.47	0.93
41:a:1042:G:HO2'	41:a:1043:C:H6	1.09	0.93
12:AB:110:PRO:HA	12:AB:114:ASP:OD2	1.69	0.93
18:D:526:C:OP2	32:R:88:LYS:NZ	2.01	0.93
7:6:18:DT:H3	8:7:52:A:H61	1.05	0.93
7:6:25:DT:H2'	11:AA:496:LYS:HE3	1.51	0.93
11:AA:387:ASN:ND2	11:AA:394:ARG:NH1	2.15	0.93
16:AG:448:LEU:HD21	16:AG:473:LEU:HD11	1.48	0.93
60:t:90:ASN:OD1	60:t:91:SER:N	2.02	0.93
9:9:23:LEU:HD13	9:9:92:ALA:CB	1.98	0.92
14:AE:59:ALA:CB	14:AE:71:LEU:HD21	1.99	0.92
39:Y:14:ALA:O	39:Y:50:LYS:HE3	1.68	0.92
60:t:20:MET:HB3	60:t:44:LYS:HZ3	1.32	0.92
16:AG:363:LEU:HG	16:AG:410:ALA:CB	1.97	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:30:G:HO2'	18:D:1339:A:H2	1.01	0.92
16:AG:123:ARG:HH12	16:AG:191:ARG:HA	1.32	0.92
16:AG:434:LEU:HD22	16:AG:459:LEU:HD12	0.93	0.92
16:AG:168:LEU:HD22	16:AG:231:GLY:HA3	1.00	0.92
10:B:76:A:N3	41:a:2493:U:H4'	1.84	0.92
14:AE:79:LYS:CA	16:AG:144:LYS:HE3	1.99	0.92
16:AG:363:LEU:CG	16:AG:410:ALA:HB2	1.99	0.92
31:Q:51:GLY:O	31:Q:56:ARG:NH2	2.03	0.92
6:5:9:DG:P	11:AA:473:ARG:HB2	2.10	0.92
16:AG:123:ARG:CZ	16:AG:191:ARG:O	2.17	0.92
16:AG:174:ARG:HB3	16:AG:175:PRO:HD2	1.51	0.92
27:M:50:LEU:HD21	27:M:121:ALA:HA	1.52	0.92
16:AG:422:GLU:HA	16:AG:426:GLY:H	1.33	0.92
12:AB:115:LYS:HZ3	12:AB:129:ILE:CG2	1.68	0.91
16:AG:434:LEU:HD21	16:AG:455:THR:O	1.69	0.91
41:a:2293:G:OP1	64:x:94:ARG:NH1	2.04	0.91
47:g:27:THR:OG1	54:n:140:GLU:OE1	1.88	0.91
11:AA:481:LEU:CD2	12:AB:16:ARG:HD3	1.98	0.91
16:AG:363:LEU:HD22	16:AG:409:ARG:HB2	1.53	0.91
12:AB:148:LYS:NZ	12:AB:150:ILE:CD1	2.32	0.91
16:AG:183:LEU:HG	16:AG:197:VAL:HG13	1.51	0.91
16:AG:425:LEU:HA	16:AG:429:LYS:CD	2.00	0.91
41:a:1818:U:OP2	48:h:156:ARG:NH1	2.03	0.91
25:K:45:ARG:CD	25:K:45:ARG:CB	2.49	0.91
9:9:92:ALA:O	9:9:129:LEU:HD13	1.71	0.91
16:AG:200:SER:O	16:AG:230:PRO:HG2	1.71	0.91
41:a:2682:A:O4'	50:j:11:MET:SD	2.29	0.91
8:7:11:U:H4'	25:K:20:ARG:HH22	1.18	0.91
8:7:47:U:H5'	11:AA:1259:LEU:HD21	1.52	0.91
12:AB:115:LYS:HA	12:AB:129:ILE:CD1	1.99	0.91
10:B:38:A:H4'	18:D:790:A:O4'	1.71	0.91
16:AG:123:ARG:NH1	16:AG:191:ARG:CA	2.10	0.91
16:AG:451:ARG:NH1	16:AG:466:ASP:O	2.02	0.91
16:AG:240:THR:OG1	16:AG:247:PRO:HG3	1.68	0.90
18:D:1526:G:N7	20:F:40:LYS:NZ	2.18	0.90
16:AG:453:VAL:HG13	16:AG:458:ASP:CB	2.00	0.90
16:AG:453:VAL:HG23	16:AG:462:GLN:NE2	1.87	0.90
6:5:10:DT:C5	11:AA:62:TYR:HD2	1.81	0.90
17:C:38:LYS:NZ	18:D:718:A:N1	2.19	0.90
27:M:66:LEU:HD13	27:M:101:MET:CE	2.02	0.90
12:AB:82:GLY:N	14:AE:142:GLU:HB3	1.86	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:240:THR:OG1	16:AG:247:PRO:CG	2.20	0.90
8:7:2:U:P	18:D:1505:G:C8	2.64	0.90
11:AA:889:PRO:HG3	16:AG:104:ARG:HH11	1.36	0.90
12:AB:115:LYS:CD	12:AB:129:ILE:HG21	2.01	0.90
36:V:27:ARG:NH1	36:V:42:THR:OG1	2.03	0.90
16:AG:363:LEU:HD23	16:AG:409:ARG:CB	2.00	0.90
65:y:89:ARG:NE	65:y:113:ARG:O	2.05	0.90
16:AG:434:LEU:HA	16:AG:456:LEU:HD23	1.53	0.90
61:u:93:ASN:OD1	61:u:94:THR:N	2.05	0.90
6:5:10:DT:H2''	12:AB:70:ASN:N	1.87	0.90
16:AG:425:LEU:CA	16:AG:429:LYS:HG3	2.02	0.90
39:Y:129:GLU:HB3	39:Y:133:ARG:NH1	1.87	0.90
52:l:170:ARG:NH1	52:l:174:GLY:O	2.05	0.90
27:M:13:LEU:HD12	27:M:14:PRO:HD2	1.54	0.90
64:x:36:TYR:OH	64:x:38:GLN:NE2	2.05	0.90
16:AG:385:ALA:HB3	16:AG:411:LYS:CE	2.00	0.89
39:Y:45:THR:CG2	39:Y:50:LYS:HD3	2.02	0.89
39:Y:79:LEU:HD21	39:Y:105:LEU:HD21	1.52	0.89
6:5:11:DA:P	12:AB:67:THR:O	2.30	0.89
12:AB:3:SER:O	12:AB:58:GLU:HA	1.71	0.89
12:AB:81:PHE:HD2	14:AE:293:ARG:NH1	1.35	0.89
41:a:404:A:O2'	41:a:405:U:OP2	1.90	0.89
11:AA:380:ALA:H	12:AB:65:HIS:CE1	1.61	0.89
11:AA:855:PRO:CG	16:AG:105:ILE:HG22	2.02	0.89
16:AG:285:ILE:HG13	16:AG:293:VAL:HG11	1.45	0.89
20:F:31:GLU:OE2	20:F:35:ARG:NE	2.06	0.89
45:e:23:ARG:NH1	45:e:24:GLU:OE2	2.06	0.89
14:AE:59:ALA:CB	14:AE:71:LEU:CD2	2.51	0.89
18:D:545:C:H5'	24:J:69:GLU:HG3	1.55	0.89
33:S:90:ARG:HG3	33:S:92:GLU:OE1	1.73	0.89
13:AD:286:GLU:H	13:AD:315:GLY:HA2	1.36	0.89
16:AG:355:ALA:HB1	16:AG:382:GLU:CD	1.94	0.89
10:A:76:A:N6	41:a:2422:C:O4'	2.06	0.88
11:AA:846:GLY:CA	16:AG:104:ARG:NH1	2.36	0.88
16:AG:302:LYS:HE3	16:AG:302:LYS:N	1.87	0.88
41:a:2756:U:OP2	57:q:19:ARG:NE	2.06	0.88
17:C:27:ALA:O	17:C:30:LYS:HG2	1.74	0.88
27:M:4:ARG:HG2	27:M:5:ARG:CZ	2.03	0.88
12:AB:81:PHE:CE1	14:AE:293:ARG:HB2	1.91	0.88
16:AG:187:ARG:CB	16:AG:188:PRO:HD2	2.03	0.88
16:AG:365:ILE:HG21	16:AG:409:ARG:NH1	1.87	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:422:GLU:CA	16:AG:426:GLY:HA3	2.03	0.88
17:C:60:LYS:NZ	18:D:735:C:O4'	2.06	0.88
39:Y:100:ILE:HD12	39:Y:105:LEU:HD11	1.54	0.88
3:2:54:GLU:OE1	3:2:91:GLN:NE2	2.06	0.88
41:a:2780:G:OP2	59:s:120:ARG:NH1	2.07	0.88
41:a:585:G:N7	66:z:6:ARG:NH1	2.21	0.88
59:s:98:GLU:OE2	59:s:126:ALA:N	2.07	0.88
16:AG:434:LEU:HD23	16:AG:456:LEU:CA	2.00	0.88
38:X:81:MET:HE2	41:a:888:C:C5	2.09	0.88
31:Q:15:GLN:NE2	31:Q:76:GLU:O	2.06	0.88
41:a:2393:U:OP1	55:o:30:ARG:NE	2.05	0.88
62:v:111:GLU:O	62:v:114:ARG:HG2	1.71	0.88
9:9:48:ALA:HB3	9:9:51:TYR:CE1	2.09	0.88
11:AA:887:VAL:CG2	16:AG:105:ILE:CG1	2.51	0.88
12:AB:129:ILE:C	12:AB:142:LEU:HD21	1.98	0.88
12:AB:148:LYS:HZ2	12:AB:150:ILE:HD11	1.34	0.88
16:AG:363:LEU:HG	16:AG:410:ALA:CA	2.04	0.88
41:a:78:U:OP1	45:e:4:LYS:NZ	2.06	0.88
18:D:1439:G:OP1	19:E:33:LYS:NZ	2.05	0.88
39:Y:129:GLU:HB3	39:Y:133:ARG:HH12	1.38	0.88
12:AB:117:ILE:HG23	12:AB:160:ARG:O	1.73	0.87
18:D:280:C:O2	36:V:40:ARG:CZ	2.21	0.87
47:g:16:CYS:HB2	47:g:37:CYS:HB3	1.54	0.87
9:9:8:LYS:O	9:9:12:VAL:HG23	1.73	0.87
12:AB:140:MET:HG3	30:P:102:LEU:HD23	1.56	0.87
16:AG:453:VAL:HG13	16:AG:458:ASP:HB2	1.55	0.87
24:J:67:VAL:HB	24:J:72:PHE:CZ	2.09	0.87
59:s:37:ARG:HA	59:s:118:MET:HE3	1.54	0.87
61:u:90:VAL:HG13	61:u:95:LEU:HD21	1.56	0.87
16:AG:434:LEU:HD21	16:AG:455:THR:C	1.97	0.87
16:AG:486:ARG:O	16:AG:490:TRP:HD1	1.57	0.87
52:l:170:ARG:NH1	52:l:176:ASP:OD1	2.08	0.87
12:AB:51:PHE:HE2	14:AE:288:PRO:CD	1.85	0.87
16:AG:248:VAL:HG13	16:AG:273:ILE:HB	1.56	0.87
16:AG:254:MET:HA	16:AG:254:MET:HE3	1.55	0.87
11:AA:847:PRO:O	16:AG:104:ARG:NH1	2.07	0.87
11:AA:887:VAL:CG1	16:AG:105:ILE:CG2	2.53	0.87
39:Y:99:LYS:HD2	39:Y:140:GLU:OE1	1.75	0.87
59:s:13:ARG:NH1	59:s:49:ASP:O	2.07	0.87
16:AG:362:TYR:C	16:AG:413:ALA:CB	2.48	0.87
16:AG:362:TYR:C	16:AG:413:ALA:HB2	1.99	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:95:LEU:O	62:v:100:LYS:NZ	2.08	0.87
63:w:24:MET:HA	63:w:24:MET:HE3	1.54	0.87
12:AB:143:LEU:HD22	30:P:88:MET:HE1	1.54	0.86
16:AG:171:GLU:CD	16:AG:267:GLY:HA3	1.99	0.86
12:AB:115:LYS:HD2	12:AB:129:ILE:CG1	1.97	0.86
7:6:25:DT:C2'	11:AA:496:LYS:HG3	2.05	0.86
11:AA:856:ASN:HD21	16:AG:35:THR:HG22	1.37	0.86
16:AG:285:ILE:CG2	16:AG:293:VAL:HG12	2.06	0.86
39:Y:100:ILE:HD12	39:Y:105:LEU:CD1	2.05	0.86
16:AG:61:ARG:HD3	16:AG:63:LEU:HD21	1.57	0.86
16:AG:302:LYS:O	16:AG:302:LYS:NZ	2.08	0.86
39:Y:15:GLY:HA2	39:Y:50:LYS:HG3	1.57	0.86
28:N:5:ASP:OD2	28:N:77:ARG:NH1	2.07	0.86
16:AG:434:LEU:HD12	16:AG:454:CYS:O	1.75	0.86
23:I:58:GLU:OE1	23:I:65:ARG:NH1	2.08	0.86
36:V:77:ARG:NH2	36:V:78:VAL:O	2.08	0.86
1:0:87:GLN:NE2	1:0:88:GLY:O	2.09	0.85
6:5:10:DT:O4'	11:AA:62:TYR:OH	1.94	0.85
16:AG:430:PRO:HG2	16:AG:435:LEU:CD2	2.05	0.85
12:AB:6:LEU:CD1	14:AE:291:ILE:HD11	2.05	0.85
14:AE:79:LYS:CE	16:AG:153:ASP:OD1	2.23	0.85
29:O:84:THR:HG21	29:O:103:PHE:HB3	1.58	0.85
16:AG:355:ALA:CA	16:AG:382:GLU:CD	2.49	0.85
16:AG:451:ARG:HH11	16:AG:467:LEU:HA	1.42	0.85
6:5:9:DG:H5''	11:AA:473:ARG:CG	1.95	0.85
9:9:58:THR:CG2	9:9:81:LEU:HD23	2.06	0.85
11:AA:853:ASP:O	16:AG:103:ASP:CG	2.19	0.85
16:AG:168:LEU:HD23	16:AG:231:GLY:HA3	1.56	0.85
36:V:25:ILE:O	36:V:27:ARG:NH2	2.10	0.85
47:g:26:SER:OG	54:n:140:GLU:OE2	1.94	0.85
1:0:68:ARG:NH1	41:a:1223:G:OP1	2.10	0.85
10:A:33:U:OP1	27:M:77:SER:OG	1.95	0.85
54:n:23:ASN:OD1	54:n:24:SER:N	2.09	0.85
12:AB:140:MET:CE	30:P:102:LEU:HG	2.02	0.85
12:AB:148:LYS:NZ	12:AB:150:ILE:HD11	1.90	0.85
42:b:36:ILE:HG21	42:b:39:ARG:HD3	1.59	0.85
62:v:77:PRO:HG2	62:v:80:VAL:HG21	1.57	0.85
11:AA:855:PRO:CA	16:AG:35:THR:OG1	2.25	0.85
12:AB:81:PHE:HE1	14:AE:293:ARG:CB	1.55	0.85
11:AA:887:VAL:HG23	16:AG:105:ILE:HD11	1.56	0.84
12:AB:145:LEU:HD11	30:P:84:VAL:CG1	2.06	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:213:VAL:HG11	16:AG:251:CYS:HA	1.59	0.84
19:E:9:LYS:O	19:E:13:GLN:NE2	2.10	0.84
11:AA:854:ILE:O	16:AG:102:PHE:CE2	2.30	0.84
16:AG:141:VAL:HG22	16:AG:178:ARG:HG2	1.57	0.84
16:AG:385:ALA:HB3	16:AG:411:LYS:HE3	1.57	0.84
6:5:10:DT:C6	11:AA:62:TYR:CZ	2.65	0.84
12:AB:51:PHE:CE2	14:AE:288:PRO:HD3	2.08	0.84
16:AG:448:LEU:CD2	16:AG:473:LEU:HD11	2.05	0.84
44:d:55:U:H1'	54:n:26:MET:HE2	1.57	0.84
61:u:122:VAL:CG1	61:u:125:LEU:HD21	2.07	0.84
62:v:7:THR:OG1	62:v:10:ARG:NH1	2.11	0.84
5:4:58:SER:OG	5:4:59:GLU:OE1	1.95	0.84
8:7:2:U:OP2	18:D:1505:G:C8	2.30	0.84
16:AG:243:LYS:HB3	16:AG:245:ILE:HD11	1.59	0.84
6:5:9:DG:P	11:AA:473:ARG:CB	2.66	0.84
12:AB:110:PRO:CA	12:AB:114:ASP:OD2	2.25	0.84
6:5:10:DT:C2	12:AB:72:THR:OG1	2.30	0.84
16:AG:387:VAL:CB	16:AG:388:PRO:HD2	2.06	0.84
12:AB:51:PHE:HZ	14:AE:288:PRO:HD3	1.39	0.84
59:s:102:GLU:OE1	59:s:102:GLU:N	2.10	0.84
39:Y:96:LYS:CE	39:Y:136:GLY:HA3	2.08	0.84
11:AA:860:ALA:H	16:AG:38:LYS:CE	1.90	0.83
16:AG:210:ARG:HG3	16:AG:216:ILE:HB	1.59	0.83
16:AG:287:ALA:HA	16:AG:331:LEU:HD12	0.87	0.83
16:AG:287:ALA:HB2	16:AG:331:LEU:HD13	1.58	0.83
10:A:6:G:O2'	10:A:7:G:O5'	1.96	0.83
11:AA:377:THR:CB	12:AB:62:GLU:HB3	2.07	0.83
18:D:742:G:OP1	34:T:58:ARG:NH2	2.11	0.83
39:Y:48:ILE:HG13	39:Y:49:GLU:H	1.41	0.83
63:w:38:LEU:HD13	63:w:111:ALA:HB2	1.59	0.83
16:AG:354:ALA:HA	16:AG:357:ASP:HB3	1.60	0.83
38:X:81:MET:HE1	38:X:92:ARG:CB	2.08	0.83
41:a:1257:C:H4'	52:l:78:TRP:CE2	2.14	0.83
41:a:2393:U:H5''	55:o:30:ARG:CZ	2.09	0.83
48:h:175:ARG:HG3	48:h:181:MET:HE1	1.58	0.83
9:9:43:LYS:O	9:9:46:ARG:HG2	1.78	0.83
66:z:78:LYS:HD3	66:z:117:LEU:CD2	2.09	0.83
16:AG:203:GLU:HA	16:AG:206:ILE:HG12	1.60	0.83
16:AG:433:ASP:C	16:AG:456:LEU:HG	2.03	0.83
39:Y:116:MET:HG2	39:Y:124:MET:HG2	1.59	0.83
11:AA:887:VAL:CB	16:AG:105:ILE:CG1	2.49	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:z:74:ILE:HD11	66:z:79:PHE:HA	1.60	0.83
12:AB:54:TYR:HE2	14:AE:291:ILE:HG23	1.44	0.83
58:r:137:GLU:HG2	58:r:138:VAL:HG23	1.61	0.83
21:G:105:LYS:O	21:G:109:GLN:NE2	2.12	0.83
26:L:102:MET:N	26:L:102:MET:SD	2.46	0.83
6:5:11:DA:C2'	12:AB:67:THR:HG21	2.08	0.82
12:AB:38:ILE:HG12	12:AB:43:ARG:HG2	1.61	0.82
14:AE:79:LYS:HG3	16:AG:144:LYS:NZ	1.94	0.82
39:Y:75:ALA:CB	39:Y:112:LYS:HE3	2.07	0.82
11:AA:387:ASN:CB	11:AA:394:ARG:HH11	1.93	0.82
11:AA:481:LEU:HD23	12:AB:16:ARG:HD2	1.60	0.82
16:AG:279:ASN:CG	16:AG:280:PRO:HD2	2.04	0.82
16:AG:285:ILE:CG2	16:AG:293:VAL:CG1	2.57	0.82
16:AG:365:ILE:HG21	16:AG:409:ARG:HH12	1.42	0.82
56:p:7:ALA:O	56:p:69:ARG:NE	2.13	0.82
10:B:6:G:O2'	10:B:7:G:O5'	1.96	0.82
23:I:22:TRP:CZ3	23:I:24:ALA:HB2	2.14	0.82
12:AB:119:THR:HG1	12:AB:126:PHE:HE2	0.83	0.82
16:AG:12:VAL:HB	16:AG:16:LYS:HG3	1.61	0.82
16:AG:127:VAL:CG2	16:AG:192:GLY:HA2	2.09	0.82
16:AG:353:HIS:CA	16:AG:356:ILE:HG23	2.10	0.82
16:AG:362:TYR:HA	16:AG:413:ALA:CB	2.09	0.82
16:AG:425:LEU:HG	16:AG:429:LYS:CE	2.07	0.82
9:9:46:ARG:HG3	9:9:94:ARG:NH2	1.94	0.82
9:9:43:LYS:HG3	9:9:46:ARG:NH1	1.94	0.82
11:AA:856:ASN:ND2	16:AG:35:THR:CG2	2.38	0.82
12:AB:138:ARG:HD2	12:AB:155:LYS:N	1.93	0.82
14:AE:287:ALA:HB1	14:AE:292:VAL:HG13	1.60	0.82
11:AA:855:PRO:HG3	16:AG:105:ILE:HG22	1.62	0.82
16:AG:133:HIS:ND1	16:AG:136:GLU:OE1	2.13	0.82
16:AG:226:ALA:HA	16:AG:236:ILE:HG13	1.61	0.82
16:AG:285:ILE:CB	16:AG:293:VAL:HG11	2.09	0.82
11:AA:847:PRO:CD	16:AG:104:ARG:NH2	2.42	0.82
39:Y:14:ALA:HB3	39:Y:51:GLY:H	1.45	0.82
54:n:73:SER:OG	54:n:79:ILE:O	1.96	0.82
60:t:12:ASP:OD2	60:t:85:VAL:HG13	1.80	0.82
9:9:1:MET:HE1	9:9:7:ASP:HB3	1.61	0.82
38:X:81:MET:CE	38:X:92:ARG:HB3	2.06	0.82
50:j:152:PRO:HG3	50:j:156:PHE:CZ	2.14	0.82
9:9:129:LEU:H	9:9:130:PRO:HD2	1.45	0.82
12:AB:115:LYS:CD	12:AB:129:ILE:CG2	2.55	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:393:LEU:HB3	16:AG:398:LEU:HD21	1.25	0.81
41:a:2298:A:OP1	54:n:72:LYS:NZ	2.11	0.81
41:a:2682:A:O2'	50:j:13:ARG:NE	2.12	0.81
48:h:130:LEU:HA	48:h:189:ARG:HH11	1.45	0.81
16:AG:361:LYS:CD	16:AG:417:ILE:HG12	2.10	0.81
48:h:53:HIS:HE2	48:h:219:THR:HA	1.45	0.81
8:7:3:G:C5'	18:D:1500:A:H62	1.92	0.81
9:9:52:MET:HG2	9:9:95:LEU:HD11	1.62	0.81
11:AA:889:PRO:CG	16:AG:104:ARG:HH11	1.92	0.81
14:AE:79:LYS:C	16:AG:144:LYS:HE3	2.04	0.81
16:AG:362:TYR:HA	16:AG:413:ALA:HB1	1.62	0.81
26:L:70:VAL:O	26:L:74:LEU:HD23	1.81	0.81
39:Y:74:PRO:HG2	39:Y:77:VAL:HB	1.60	0.81
41:a:1613:G:O2'	53:m:3:ARG:NH2	2.12	0.81
41:a:2356:U:H5''	42:b:20:ARG:HD2	1.61	0.81
12:AB:120:GLU:HB2	12:AB:162:LEU:HG	1.63	0.81
16:AG:434:LEU:CG	16:AG:455:THR:C	2.54	0.81
6:5:10:DT:C4	11:AA:62:TYR:CD2	2.68	0.81
18:D:107:G:O6	19:E:10:ARG:NE	2.13	0.81
20:F:37:PHE:HB2	31:Q:126:LYS:CE	2.10	0.81
39:Y:45:THR:HG22	39:Y:50:LYS:HD3	1.59	0.81
41:a:2356:U:O3'	42:b:20:ARG:NH1	2.14	0.81
52:l:112:LEU:HD12	52:l:117:ARG:HB2	1.62	0.81
56:p:104:ASN:ND2	56:p:114:ASP:OD1	2.12	0.81
8:7:2:U:OP1	18:D:1505:G:C8	2.33	0.81
11:AA:387:ASN:CG	11:AA:394:ARG:HH12	1.81	0.81
61:u:122:VAL:HG13	61:u:125:LEU:HD11	1.61	0.81
11:AA:855:PRO:C	16:AG:35:THR:CB	2.47	0.81
16:AG:451:ARG:NE	16:AG:470:ILE:HG13	1.95	0.81
56:p:121:ILE:HD12	56:p:141:ILE:HG22	1.60	0.81
60:t:7:MET:HG3	60:t:18:ARG:CZ	2.10	0.81
14:AE:81:ARG:HH21	14:AE:81:ARG:HG3	1.44	0.81
11:AA:887:VAL:HG23	16:AG:105:ILE:CD1	2.08	0.81
39:Y:9:LYS:HE2	39:Y:55:PRO:HB3	1.63	0.81
43:c:30:LEU:HD12	43:c:31:PRO:HD2	1.63	0.81
52:l:1:MET:HE1	52:l:20:GLY:CA	2.11	0.81
14:AE:79:LYS:CG	16:AG:144:LYS:NZ	2.44	0.80
9:9:77:VAL:HG13	9:9:114:GLU:OE1	1.81	0.80
11:AA:887:VAL:CG1	16:AG:105:ILE:HG21	2.11	0.80
16:AG:381:LEU:HA	16:AG:384:LEU:CD1	2.09	0.80
7:6:23:DC:H5'	14:AE:259:ARG:HH12	1.43	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:440:VAL:CG2	16:AG:481:LEU:CD2	2.60	0.80
18:D:652:U:O3'	28:N:56:LYS:NZ	2.13	0.80
41:a:2682:A:C8	50:j:11:MET:CE	2.65	0.80
16:AG:252:VAL:HA	16:AG:259:VAL:HG23	1.61	0.80
16:AG:287:ALA:HB1	16:AG:331:LEU:HD12	1.62	0.80
16:AG:425:LEU:HG	16:AG:429:LYS:CD	2.10	0.80
39:Y:58:ILE:HG22	39:Y:60:VAL:HG23	1.63	0.80
9:9:118:ILE:HB	9:9:119:PRO:HD3	1.64	0.80
12:AB:117:ILE:HD13	12:AB:161:LYS:HZ1	1.47	0.80
42:b:28:GLY:O	42:b:66:LYS:NZ	2.13	0.80
9:9:25:ALA:HB3	9:9:99:PHE:CD2	2.16	0.80
18:D:311:C:OP1	35:U:31:ARG:NH2	2.13	0.80
28:N:9:ASP:OD2	28:N:13:ARG:NH1	2.15	0.80
9:9:3:LEU:HD11	9:9:5:LEU:HD23	1.61	0.80
11:AA:860:ALA:N	16:AG:38:LYS:CE	2.45	0.80
16:AG:425:LEU:CA	16:AG:429:LYS:CG	2.60	0.80
39:Y:81:LYS:HZ2	39:Y:86:LYS:HE2	1.47	0.80
41:a:1655:A:H1'	50:j:118:PHE:HE2	1.46	0.80
6:5:10:DT:C2	12:AB:71:ALA:HB3	1.84	0.80
16:AG:133:HIS:HD1	16:AG:136:GLU:CD	1.90	0.80
4:3:46:GLN:OE1	4:3:56:GLY:HA2	1.82	0.80
9:9:81:LEU:HD21	41:a:1107:G:H4'	1.64	0.80
11:AA:855:PRO:HB2	16:AG:35:THR:HB	1.61	0.80
16:AG:380:THR:O	16:AG:384:LEU:HD11	1.80	0.80
16:AG:381:LEU:CA	16:AG:384:LEU:HD12	2.08	0.80
16:AG:451:ARG:NH2	16:AG:470:ILE:HG13	1.97	0.80
41:a:1818:U:O5'	48:h:156:ARG:NH2	2.14	0.80
9:9:5:LEU:O	9:9:9:GLN:NE2	2.15	0.80
11:AA:887:VAL:HG11	16:AG:105:ILE:CG2	2.12	0.80
12:AB:54:TYR:CE2	14:AE:291:ILE:CG2	2.65	0.80
1:0:86:GLN:OE1	41:a:1225:G:H4'	1.81	0.79
11:AA:855:PRO:CG	16:AG:35:THR:HG21	2.12	0.79
39:Y:40:ALA:HB1	39:Y:68:PHE:CE1	2.16	0.79
60:t:98:ARG:NH2	60:t:118:LEU:O	2.15	0.79
6:5:11:DA:C4'	12:AB:67:THR:HG22	2.11	0.79
16:AG:379:SER:OG	16:AG:384:LEU:HD21	1.81	0.79
66:z:74:ILE:HD13	66:z:79:PHE:HD1	1.46	0.79
41:a:2636:C:H4'	50:j:81:GLU:OE1	1.82	0.79
66:z:69:ALA:HB1	66:z:74:ILE:HG23	1.63	0.79
16:AG:451:ARG:NH1	16:AG:467:LEU:C	2.40	0.79
49:i:46:ASP:OD2	49:i:53:LYS:NZ	2.15	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:t:20:MET:HB3	60:t:44:LYS:NZ	1.98	0.79
6:5:10:DT:H73	11:AA:62:TYR:HD2	1.28	0.79
12:AB:51:PHE:CE2	14:AE:288:PRO:HG2	2.06	0.79
11:AA:855:PRO:CA	16:AG:35:THR:CB	2.61	0.79
52:l:111:GLU:OE2	52:l:114:ARG:NH1	2.14	0.79
16:AG:353:HIS:C	16:AG:356:ILE:CG2	2.50	0.79
16:AG:430:PRO:CG	16:AG:435:LEU:HD21	2.11	0.79
22:H:306:VAL:HA	22:H:309:MET:SD	2.23	0.79
41:a:1276:A:O2'	63:w:20:MET:CE	2.30	0.79
41:a:2393:U:P	55:o:30:ARG:HE	2.04	0.79
6:5:10:DT:C2'	12:AB:71:ALA:H	1.95	0.79
12:AB:110:PRO:HB3	12:AB:114:ASP:OD2	1.83	0.79
16:AG:447:LYS:O	16:AG:470:ILE:CD1	2.31	0.79
17:C:38:LYS:HE2	17:C:38:LYS:HA	1.64	0.79
24:J:62:ARG:NH1	24:J:69:GLU:OE2	2.16	0.79
41:a:1453:A:N1	63:w:74:GLU:OE1	2.16	0.79
8:7:2:U:OP1	18:D:1505:G:H8	1.65	0.79
16:AG:363:LEU:CG	16:AG:410:ALA:N	2.45	0.79
16:AG:425:LEU:HD23	16:AG:429:LYS:HE2	0.81	0.79
54:n:117:LEU:HD13	54:n:176:PRO:HG2	1.64	0.79
9:9:3:LEU:CD1	9:9:4:ASN:H	1.96	0.78
16:AG:437:LEU:HD21	16:AG:456:LEU:HD13	0.81	0.78
39:Y:102:ARG:HD2	39:Y:141:ASP:CA	2.13	0.78
48:h:258:ARG:NH1	48:h:264:ASP:OD1	2.14	0.78
9:9:112:ALA:O	9:9:123:ILE:HD11	1.83	0.78
11:AA:377:THR:OG1	12:AB:62:GLU:CA	2.30	0.78
14:AE:79:LYS:HB3	16:AG:159:GLU:OE2	1.82	0.78
16:AG:235:LYS:HD3	16:AG:327:LEU:HG	1.65	0.78
16:AG:365:ILE:CG2	16:AG:409:ARG:HH12	1.91	0.78
16:AG:434:LEU:CD2	16:AG:456:LEU:N	2.46	0.78
33:S:12:LYS:O	33:S:16:LEU:HD23	1.83	0.78
12:AB:51:PHE:CZ	14:AE:288:PRO:CD	2.66	0.78
12:AB:136:GLU:CD	12:AB:160:ARG:NH1	2.42	0.78
35:U:3:THR:HG21	35:U:5:ARG:CZ	2.13	0.78
37:W:40:ILE:HG23	37:W:44:MET:SD	2.24	0.78
41:a:2393:U:H5''	55:o:30:ARG:NH1	1.97	0.78
47:g:18:CYS:HB3	47:g:40:CYS:CB	2.13	0.78
11:AA:856:ASN:CG	16:AG:35:THR:CA	2.54	0.78
16:AG:363:LEU:HD23	16:AG:409:ARG:HB2	1.62	0.78
16:AG:387:VAL:CG1	16:AG:388:PRO:CD	2.41	0.78
27:M:20:SER:HB3	27:M:23:LEU:HD13	1.63	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1:98:LYS:HD2	41:a:2012:G:OP1	1.83	0.78
59:s:37:ARG:HA	59:s:118:MET:CE	2.14	0.78
10:A:18:G:N2	10:A:55:U:O2	2.17	0.78
11:AA:847:PRO:HD2	16:AG:104:ARG:HH22	1.43	0.78
12:AB:120:GLU:HB2	12:AB:162:LEU:CG	2.14	0.78
12:AB:129:ILE:O	12:AB:142:LEU:HD21	1.84	0.78
56:p:107:LEU:HB3	56:p:152:ARG:HE	1.47	0.78
7:6:25:DT:H3'	11:AA:496:LYS:CG	2.13	0.78
9:9:34:THR:O	9:9:38:MET:HG2	1.83	0.78
41:a:1406:U:O2'	41:a:1407:G:O5'	2.02	0.78
60:t:8:LEU:C	60:t:18:ARG:HH12	1.92	0.78
9:9:19:ALA:HA	9:9:70:GLU:HG2	1.66	0.78
16:AG:393:LEU:CD2	16:AG:398:LEU:CG	2.58	0.78
18:D:673:A:H2'	18:D:674:G:C8	2.18	0.78
54:n:142:ASP:HB3	54:n:145:LYS:HE2	1.65	0.78
12:AB:9:CYS:H	12:AB:53:ASN:HB3	1.48	0.78
16:AG:451:ARG:HH12	16:AG:467:LEU:CA	1.92	0.78
18:D:275:G:H4'	36:V:16:LYS:HD2	1.65	0.78
28:N:27:MET:N	28:N:58:GLU:OE2	2.16	0.78
16:AG:189:GLU:HB3	16:AG:194:GLN:HG3	1.64	0.78
27:M:40:GLU:OE2	29:O:41:ARG:NH2	2.17	0.78
6:5:10:DT:C2	12:AB:71:ALA:CB	2.61	0.77
9:9:27:VAL:HG13	9:9:83:ALA:HB3	1.66	0.77
11:AA:846:GLY:C	16:AG:104:ARG:CZ	2.58	0.77
11:AA:887:VAL:CG1	16:AG:105:ILE:HG23	2.13	0.77
9:9:134:GLU:OE2	9:9:136:ILE:HG22	1.84	0.77
16:AG:425:LEU:HD23	16:AG:429:LYS:NZ	1.99	0.77
19:E:28:MET:HE1	19:E:58:VAL:HA	1.63	0.77
40:Z:14:MET:HG3	40:Z:15:SER:H	1.48	0.77
59:s:15:TRP:HZ3	59:s:135:GLN:HA	1.50	0.77
11:AA:887:VAL:C	16:AG:105:ILE:HD11	2.08	0.77
16:AG:354:ALA:HA	16:AG:357:ASP:CB	2.14	0.77
16:AG:402:THR:O	16:AG:405:ALA:HB3	1.84	0.77
26:L:42:TRP:HZ3	26:L:45:ARG:HH22	1.32	0.77
39:Y:9:LYS:HE2	39:Y:55:PRO:CG	2.14	0.77
9:9:60:LEU:HD23	9:9:64:VAL:HG21	1.65	0.77
12:AB:115:LYS:CE	12:AB:129:ILE:HG23	2.14	0.77
16:AG:365:ILE:CD1	16:AG:406:LEU:CD2	2.57	0.77
18:D:375:U:OP2	35:U:70:ARG:NH2	2.17	0.77
18:D:375:U:P	35:U:70:ARG:HE	2.07	0.77
57:q:2:LYS:NZ	57:q:32:LYS:O	2.17	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:94:ARG:HB2	9:9:97:LYS:HE2	1.67	0.77
16:AG:233:ARG:HB2	16:AG:327:LEU:HD23	1.66	0.77
16:AG:434:LEU:HD11	16:AG:455:THR:O	1.85	0.77
18:D:1227:A:OP1	38:X:93:ARG:NH2	2.18	0.77
16:AG:284:VAL:HG12	16:AG:296:ILE:HD13	1.66	0.77
11:AA:855:PRO:HB3	16:AG:35:THR:OG1	1.81	0.77
16:AG:412:ASN:O	16:AG:416:THR:HG23	1.83	0.77
64:x:104:GLN:NE2	64:x:108:ASP:OD2	2.16	0.77
16:AG:258:ARG:HG2	16:AG:258:ARG:NH1	1.99	0.77
11:AA:887:VAL:HB	16:AG:105:ILE:CG2	2.14	0.77
10:B:18:G:N2	10:B:55:U:O2	2.17	0.76
17:C:38:LYS:HD3	18:D:719:C:H1'	1.67	0.76
18:D:739:C:O2'	34:T:42:HIS:ND1	2.18	0.76
31:Q:84:VAL:HG23	31:Q:107:ILE:HD11	1.67	0.76
31:Q:87:LYS:CG	31:Q:113:VAL:HG23	2.15	0.76
35:U:3:THR:HG21	35:U:5:ARG:NH1	1.99	0.76
22:H:52:GLN:OE1	22:H:86:ARG:CB	2.34	0.76
41:a:686:U:O4	53:m:12:ARG:HD3	1.86	0.76
41:a:2511:U:C4'	50:j:130:GLN:OE1	2.33	0.76
16:AG:228:ARG:HD2	16:AG:236:ILE:HB	1.67	0.76
16:AG:451:ARG:CZ	16:AG:470:ILE:HG13	2.16	0.76
40:Z:2:ILE:HG22	40:Z:4:LYS:H	1.50	0.76
41:a:196:A:OP2	61:u:47:ARG:NH1	2.18	0.76
41:a:1801:A:P	48:h:146:MET:HE1	2.25	0.76
41:a:2297:A:N1	41:a:2321:U:C4	2.53	0.76
2:1:11:ARG:CZ	2:1:98:LYS:HE3	2.15	0.76
14:AE:79:LYS:CB	16:AG:144:LYS:NZ	2.33	0.76
29:O:56:ASP:O	29:O:60:LYS:NZ	2.16	0.76
41:a:2335:A:OP2	64:x:13:ARG:CZ	2.34	0.76
42:b:21:LEU:HD21	42:b:41:ARG:HE	1.51	0.76
2:1:24:ILE:HG21	2:1:36:LEU:HD11	1.68	0.76
21:G:30:PHE:CD1	21:G:201:PRO:HG3	2.21	0.76
39:Y:14:ALA:CB	39:Y:50:LYS:HD2	2.16	0.76
65:y:71:GLU:OE1	65:y:101:ARG:NE	2.19	0.76
11:AA:856:ASN:CG	16:AG:35:THR:HG22	2.11	0.76
11:AA:855:PRO:HD2	16:AG:105:ILE:HG21	1.67	0.76
16:AG:283:PHE:CZ	16:AG:331:LEU:O	2.39	0.76
16:AG:354:ALA:O	16:AG:356:ILE:N	2.19	0.76
18:D:376:G:H5''	35:U:5:ARG:HD2	1.67	0.76
27:M:18:PHE:HE2	27:M:47:LEU:HD21	1.48	0.76
39:Y:93:ASN:C	39:Y:94:LYS:HD2	2.10	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2314:A:OP1	54:n:88:LYS:NZ	2.18	0.76
16:AG:131:ARG:HA	16:AG:186:VAL:HG11	1.68	0.76
16:AG:385:ALA:HB1	16:AG:411:LYS:CE	2.04	0.76
37:W:44:MET:SD	37:W:62:VAL:HG11	2.25	0.76
41:a:1828:G:O6	48:h:221:ARG:NH1	2.19	0.76
5:4:50:MET:O	5:4:56:PHE:CE2	2.38	0.76
18:D:550:G:O2'	32:R:116:LYS:NZ	2.18	0.76
9:9:37:LYS:O	9:9:41:LEU:HG	1.86	0.75
25:K:142:ASP:O	25:K:145:GLU:HG3	1.86	0.75
6:5:10:DT:C4	11:AA:62:TYR:CE2	2.74	0.75
6:5:11:DA:C4'	12:AB:67:THR:CG2	2.64	0.75
9:9:77:VAL:HG12	9:9:82:ILE:HG21	1.68	0.75
14:AE:59:ALA:HB1	14:AE:71:LEU:CD2	2.16	0.75
21:G:20:THR:HA	21:G:39:HIS:CE1	2.21	0.75
27:M:18:PHE:CE2	27:M:47:LEU:HD21	2.21	0.75
38:X:57:ARG:NH2	47:g:35:ASP:OD1	2.19	0.75
42:b:68:LYS:NZ	42:b:70:GLU:OE1	2.19	0.75
6:5:10:DT:H2''	12:AB:69:ILE:C	2.11	0.75
11:AA:887:VAL:HB	16:AG:105:ILE:HG23	1.69	0.75
16:AG:354:ALA:C	16:AG:356:ILE:H	1.93	0.75
18:D:404:G:OP1	24:J:115:ARG:NH1	2.19	0.75
38:X:28:THR:HG23	38:X:31:LYS:HE3	1.69	0.75
41:a:1405:U:H2'	41:a:1406:U:C6	2.20	0.75
8:7:11:U:HO2'	25:K:33:PHE:HE1	1.32	0.75
16:AG:62:TRP:HB3	16:AG:75:ILE:HD11	1.69	0.75
16:AG:201:LYS:HD2	16:AG:201:LYS:N	2.01	0.75
16:AG:240:THR:CG2	16:AG:247:PRO:HG3	2.16	0.75
41:a:1276:A:O2'	63:w:20:MET:HE2	1.87	0.75
16:AG:307:ILE:HB	16:AG:338:VAL:HA	1.66	0.75
16:AG:425:LEU:CB	16:AG:429:LYS:HE3	2.16	0.75
39:Y:56:VAL:CA	39:Y:71:LYS:HZ1	1.99	0.75
8:7:3:G:H5''	18:D:1500:A:H62	1.49	0.75
9:9:36:ASP:O	9:9:39:THR:HG22	1.87	0.75
11:AA:847:PRO:N	16:AG:104:ARG:NH1	2.33	0.75
12:AB:89:PRO:HD2	14:AE:290:ILE:HG21	1.68	0.75
16:AG:434:LEU:CD2	16:AG:455:THR:C	2.59	0.75
21:G:49:MET:CE	21:G:201:PRO:HD3	2.15	0.75
62:v:1:MET:HA	62:v:1:MET:HE3	1.67	0.75
59:s:15:TRP:CZ3	59:s:135:GLN:HA	2.21	0.75
12:AB:18:GLN:HG3	12:AB:21:LEU:HD12	1.69	0.75
16:AG:248:VAL:HG22	16:AG:273:ILE:HG22	1.68	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:73:ARG:NH1	31:Q:113:VAL:HG12	2.02	0.75
22:H:36:VAL:O	22:H:37:LEU:HD12	1.87	0.75
24:J:58:LYS:NZ	24:J:69:GLU:OE2	2.15	0.74
39:Y:126:ARG:NH2	41:a:1083:U:OP1	2.19	0.74
16:AG:198:THR:HB	16:AG:201:LYS:HD3	1.70	0.74
16:AG:365:ILE:HG23	16:AG:409:ARG:HH11	1.52	0.74
16:AG:385:ALA:HB3	16:AG:411:LYS:HE2	1.68	0.74
18:D:1217:C:OP2	33:S:9:ARG:NH2	2.19	0.74
31:Q:87:LYS:HG2	31:Q:113:VAL:HG23	1.67	0.74
41:a:537:G:OP2	59:s:2:LYS:NZ	2.20	0.74
41:a:742:A:H2'	41:a:743:A:C8	2.21	0.74
54:n:4:LEU:HD11	54:n:101:GLU:CA	2.16	0.74
63:w:12:ARG:O	63:w:17:ARG:NH2	2.19	0.74
5:4:57:TYR:HB3	62:v:136:MET:CE	2.16	0.74
8:7:3:G:C5'	18:D:1500:A:N6	2.46	0.74
9:9:59:LEU:HD22	9:9:62:ARG:NE	2.02	0.74
16:AG:354:ALA:O	16:AG:355:ALA:C	2.26	0.74
16:AG:354:ALA:C	16:AG:356:ILE:N	2.43	0.74
50:j:46:ARG:NH2	50:j:88:GLU:O	2.19	0.74
11:AA:860:ALA:H	16:AG:38:LYS:CD	1.98	0.74
12:AB:137:ALA:O	12:AB:155:LYS:C	2.30	0.74
16:AG:371:THR:O	16:AG:375:GLU:HG3	1.86	0.74
41:a:1278:C:H4'	63:w:24:MET:HE1	1.69	0.74
16:AG:451:ARG:HE	16:AG:470:ILE:HG13	1.51	0.74
39:Y:9:LYS:HE2	39:Y:55:PRO:CB	2.17	0.74
11:AA:387:ASN:OD1	11:AA:394:ARG:NH1	2.21	0.74
22:H:303:LEU:O	22:H:305:HIS:N	2.20	0.74
54:n:56:ASP:OD2	54:n:150:ARG:NH1	2.20	0.74
9:9:5:LEU:C	9:9:9:GLN:HE22	1.94	0.74
12:AB:133:PRO:HA	12:AB:159:PHE:HZ	1.50	0.74
16:AG:279:ASN:CB	16:AG:280:PRO:CD	2.66	0.74
16:AG:313:ASN:N	16:AG:313:ASN:OD1	2.19	0.74
17:C:18:VAL:O	17:C:51:TYR:OH	2.06	0.74
2:1:86:MET:HE1	2:1:88:ARG:HD3	1.70	0.74
16:AG:365:ILE:HG23	16:AG:409:ARG:NH1	2.02	0.74
34:T:64:ARG:CZ	34:T:89:ARG:HH22	2.00	0.74
39:Y:100:ILE:HG13	39:Y:139:VAL:CG2	2.18	0.74
39:Y:102:ARG:CD	39:Y:141:ASP:HA	2.18	0.74
52:l:90:GLN:NE2	52:l:91:ASP:O	2.20	0.74
5:4:45:ASP:OD1	5:4:46:LYS:N	2.21	0.74
9:9:23:LEU:HD11	9:9:96:PHE:CD2	2.23	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:g:18:CYS:CB	47:g:40:CYS:HB3	2.17	0.74
60:t:17:ARG:HB2	60:t:45:GLU:OE1	1.88	0.74
16:AG:363:LEU:CD1	16:AG:410:ALA:HB2	2.17	0.73
48:h:130:LEU:HA	48:h:189:ARG:NH1	2.03	0.73
58:r:6:LEU:HD11	58:r:37:VAL:HG23	1.69	0.73
6:5:10:DT:C5	11:AA:62:TYR:HE2	1.97	0.73
16:AG:442:ARG:HB3	16:AG:446:PHE:HE2	1.53	0.73
18:D:375:U:OP1	35:U:70:ARG:HG2	1.88	0.73
20:F:63:GLU:OE1	20:F:67:ARG:NH1	2.20	0.73
49:i:55:ILE:HD13	63:w:112:TYR:CZ	2.23	0.73
56:p:69:ARG:NH1	56:p:73:ASN:OD1	2.21	0.73
2:l:25:ARG:HH22	41:a:520:G:P	2.10	0.73
9:9:59:LEU:HD13	9:9:62:ARG:HD2	1.70	0.73
9:9:129:LEU:HD12	9:9:130:PRO:HD3	1.70	0.73
16:AG:297:VAL:HG23	16:AG:306:ASP:HB2	1.70	0.73
16:AG:425:LEU:O	16:AG:429:LYS:HE3	1.87	0.73
21:G:154:MET:CE	21:G:158:PRO:HD3	2.16	0.73
34:T:76:ALA:O	34:T:80:GLN:NE2	2.22	0.73
9:9:43:LYS:NZ	9:9:98:GLU:HB2	2.04	0.73
16:AG:365:ILE:CG2	16:AG:409:ARG:HH11	2.01	0.73
16:AG:486:ARG:O	16:AG:490:TRP:CD1	2.41	0.73
10:B:76:A:C2	41:a:2493:U:H4'	2.24	0.73
24:J:100:ASN:OD1	24:J:111:ARG:NH1	2.21	0.73
39:Y:100:ILE:HD11	39:Y:139:VAL:HB	1.69	0.73
16:AG:393:LEU:C	16:AG:398:LEU:HD23	2.13	0.73
18:D:636:U:H4'	36:V:6:ARG:CZ	2.19	0.73
24:J:105:MET:CE	24:J:171:LEU:HD13	2.18	0.73
6:5:10:DT:N3	12:AB:72:THR:OG1	2.20	0.73
16:AG:429:LYS:H	16:AG:429:LYS:HD2	1.53	0.73
16:AG:219:GLU:HA	16:AG:219:GLU:OE2	1.87	0.73
27:M:55:GLY:O	27:M:56:LYS:O	2.07	0.73
5:4:75:GLN:HG3	5:4:92:VAL:HG23	1.70	0.73
10:A:53:G:C2	10:A:54:U:C5	2.77	0.73
18:D:1158:C:O2'	21:G:132:LYS:NZ	2.21	0.73
62:v:97:GLN:N	62:v:100:LYS:HE3	2.03	0.73
63:w:33:ILE:HD11	63:w:112:TYR:CD1	2.23	0.73
12:AB:89:PRO:CD	14:AE:290:ILE:HG21	2.18	0.73
12:AB:148:LYS:NZ	12:AB:150:ILE:HD13	2.03	0.73
16:AG:365:ILE:HD13	16:AG:406:LEU:HD21	1.70	0.73
16:AG:413:ALA:O	16:AG:416:THR:OG1	2.05	0.73
38:X:81:MET:HE2	41:a:888:C:C6	2.24	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:Z:6:GLN:OE1	40:Z:10:ALA:HB2	1.88	0.73
2:1:25:ARG:NH2	41:a:520:G:H5'	2.03	0.73
12:AB:36:GLU:H	12:AB:106:ASP:CG	1.96	0.73
47:g:18:CYS:CB	47:g:40:CYS:CB	2.67	0.73
55:o:25:LYS:HB3	61:u:62:PRO:HG2	1.71	0.73
62:v:73:ILE:HD11	62:v:93:VAL:HG22	1.70	0.73
10:B:53:G:C2	10:B:54:U:C5	2.77	0.72
24:J:184:ARG:NH1	24:J:187:GLU:OE1	2.22	0.72
41:a:465:G:OP1	53:m:12:ARG:NH2	2.22	0.72
62:v:50:ARG:HA	62:v:53:MET:SD	2.29	0.72
63:w:60:VAL:HG22	63:w:63:ARG:NH1	2.03	0.72
11:AA:847:PRO:CD	16:AG:104:ARG:HH22	2.00	0.72
17:C:57:ARG:HE	17:C:61:ARG:HH12	1.37	0.72
12:AB:24:GLN:HB2	12:AB:26:VAL:HG23	1.71	0.72
16:AG:223:ILE:O	16:AG:223:ILE:HD13	1.89	0.72
16:AG:434:LEU:CD2	16:AG:459:LEU:CD1	2.54	0.72
18:D:13:U:C4	18:D:915:A:N6	2.57	0.72
18:D:653:U:C6	28:N:56:LYS:HE3	2.25	0.72
9:9:78:GLY:N	9:9:79:PRO:HD2	2.04	0.72
11:AA:377:THR:CG2	12:AB:62:GLU:HB3	2.18	0.72
11:AA:380:ALA:H	12:AB:65:HIS:HE1	1.00	0.72
16:AG:262:VAL:HA	16:AG:265:GLU:HB2	1.72	0.72
17:C:60:LYS:HZ1	18:D:735:C:C4'	2.02	0.72
8:7:13:U:H3	24:J:47:ARG:HH11	1.37	0.72
12:AB:21:LEU:HB3	12:AB:26:VAL:HG11	1.70	0.72
16:AG:393:LEU:CD2	16:AG:398:LEU:HD13	2.19	0.72
16:AG:434:LEU:HG	16:AG:455:THR:C	2.14	0.72
26:L:36:ILE:CD1	26:L:64:VAL:HG22	2.19	0.72
39:Y:102:ARG:CZ	39:Y:139:VAL:HG11	2.19	0.72
11:AA:861:ALA:N	16:AG:38:LYS:HZ3	1.72	0.72
12:AB:149:GLU:HA	12:AB:149:GLU:OE2	1.87	0.72
23:I:142:MET:HE3	23:I:170:GLU:OE1	1.89	0.72
44:d:42:C:O5'	54:n:64:LYS:NZ	2.20	0.72
48:h:123:ALA:HB1	48:h:125:LYS:NZ	2.04	0.72
14:AE:68:TYR:HA	14:AE:92:VAL:HG23	1.72	0.72
23:I:22:TRP:HZ3	23:I:24:ALA:HB2	1.54	0.72
23:I:39:VAL:O	23:I:43:LEU:HD23	1.90	0.72
30:P:18:ILE:O	30:P:22:THR:HG23	1.90	0.72
32:R:56:ARG:HG2	32:R:62:GLU:OE2	1.89	0.72
20:F:32:VAL:HG12	20:F:36:GLU:OE1	1.90	0.72
41:a:1825:U:OP2	48:h:52:ARG:NH1	2.23	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2328:A:H2'	41:a:2329:U:C6	2.25	0.72
56:p:17:VAL:O	56:p:18:LYS:HE2	1.89	0.72
9:9:27:VAL:CG1	9:9:83:ALA:HB3	2.20	0.72
9:9:47:GLU:HB3	9:9:51:TYR:CZ	2.24	0.72
11:AA:889:PRO:HG3	16:AG:104:ARG:NH1	2.04	0.72
16:AG:434:LEU:HD21	16:AG:456:LEU:N	2.03	0.72
16:AG:434:LEU:CA	16:AG:456:LEU:HD21	2.12	0.72
2:1:83:LYS:HB2	2:1:95:ARG:HD3	1.72	0.72
9:9:56:ARG:O	9:9:56:ARG:HG2	1.88	0.72
12:AB:140:MET:HE2	30:P:102:LEU:CG	2.10	0.72
14:AE:64:PRO:HG3	14:AE:90:VAL:CG1	2.19	0.72
16:AG:220:VAL:O	16:AG:240:THR:HG22	1.90	0.72
16:AG:254:MET:HA	16:AG:254:MET:CE	2.17	0.72
24:J:76:TYR:CE2	24:J:204:TYR:HB2	2.24	0.72
43:c:71:LEU:HB3	43:c:76:GLU:OE2	1.88	0.72
60:t:8:LEU:C	60:t:18:ARG:NH1	2.48	0.72
14:AE:85:CYS:SG	68:AE:1502:ZN:ZN	1.79	0.71
16:AG:352:ALA:O	16:AG:356:ILE:HG22	1.90	0.71
47:g:5:ILE:HD12	54:n:64:LYS:HE3	1.72	0.71
1:0:10:LYS:HE2	41:a:994:C:O2'	1.90	0.71
2:1:23:LEU:HD11	49:i:24:ALA:HB2	1.70	0.71
12:AB:38:ILE:CD1	12:AB:160:ARG:NH1	2.54	0.71
16:AG:355:ALA:CB	16:AG:382:GLU:OE1	2.18	0.71
19:E:21:ASN:C	19:E:25:ARG:HE	1.97	0.71
21:G:19:GLN:NE2	22:H:75:VAL:O	2.23	0.71
25:K:69:ARG:CZ	25:K:69:ARG:HA	2.20	0.71
29:O:47:VAL:HA	29:O:50:GLN:HG2	1.70	0.71
41:a:2884:U:OP2	49:i:40:ARG:NH2	2.22	0.71
16:AG:403:VAL:HG22	16:AG:406:LEU:HD12	1.71	0.71
18:D:1208:C:H2'	18:D:1209:C:H6	1.56	0.71
56:p:85:LYS:HE3	56:p:164:TYR:CD1	2.24	0.71
3:2:56:GLU:OE2	3:2:88:LYS:HA	1.91	0.71
12:AB:137:ALA:O	12:AB:155:LYS:HA	1.90	0.71
16:AG:167:MET:HE1	16:AG:199:ARG:HG3	1.72	0.71
66:z:92:ARG:HG3	66:z:95:LEU:HD12	1.71	0.71
9:9:142:THR:OG1	40:Z:7:ILE:HG21	1.90	0.71
16:AG:127:VAL:HG22	16:AG:192:GLY:C	2.15	0.71
18:D:736:C:C5'	26:L:90:MET:HE2	2.21	0.71
20:F:13:ASP:OD1	20:F:14:VAL:N	2.22	0.71
25:K:111:MET:O	25:K:115:LEU:HD23	1.91	0.71
40:Z:14:MET:HB3	40:Z:17:MET:CG	2.21	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:v:39:GLY:N	62:v:96:ILE:O	2.22	0.71
14:AE:77:ARG:NE	14:AE:77:ARG:HA	2.03	0.71
18:D:973:G:O4'	30:P:57:VAL:HG22	1.90	0.71
19:E:75:HIS:CE1	19:E:79:LEU:HD11	2.25	0.71
32:R:5:ASN:OD1	36:V:36:LYS:NZ	2.24	0.71
34:T:63:ARG:O	34:T:67:LEU:HD23	1.90	0.71
41:a:2780:G:P	59:s:120:ARG:HH11	2.13	0.71
11:AA:887:VAL:O	16:AG:105:ILE:HD11	1.90	0.71
14:AE:59:ALA:HB1	14:AE:71:LEU:HD22	1.71	0.71
16:AG:362:TYR:CA	16:AG:413:ALA:CB	2.69	0.71
41:a:2356:U:H5''	42:b:20:ARG:CD	2.21	0.71
41:a:2469:A:N6	41:a:2481:G:O2'	2.23	0.71
52:l:6:LYS:NZ	52:l:120:VAL:O	2.24	0.71
11:AA:854:ILE:C	16:AG:102:PHE:CD2	2.69	0.71
12:AB:110:PRO:CB	12:AB:114:ASP:OD2	2.38	0.71
16:AG:393:LEU:HD23	16:AG:398:LEU:CD1	2.20	0.71
16:AG:433:ASP:C	16:AG:456:LEU:CG	2.64	0.71
18:D:13:U:O4	18:D:21:G:C2	2.43	0.71
23:I:74:GLY:O	23:I:78:GLY:N	2.24	0.71
23:I:166:GLU:OE1	23:I:166:GLU:N	2.23	0.71
33:S:90:ARG:HG3	33:S:92:GLU:CD	2.15	0.71
41:a:1824:G:OP1	48:h:52:ARG:NH2	2.23	0.71
54:n:134:GLU:HG2	54:n:137:ILE:HG23	1.72	0.71
27:M:16:PRO:HB3	29:O:46:MET:HG3	1.73	0.71
39:Y:57:VAL:O	39:Y:69:VAL:HG22	1.91	0.71
10:B:13:C:OP2	41:a:1907:G:N2	2.23	0.71
18:D:1328:C:O2'	38:X:29:ARG:NH2	2.23	0.71
22:H:332:VAL:HG22	22:H:342:ILE:HD11	1.73	0.71
34:T:64:ARG:NH1	34:T:64:ARG:HA	2.05	0.71
59:s:15:TRP:CZ3	59:s:135:GLN:HG3	2.25	0.71
2:l:25:ARG:HH22	41:a:520:G:C5'	2.04	0.70
9:9:56:ARG:NH2	41:a:1084:A:O4'	2.23	0.70
9:9:94:ARG:HB3	9:9:97:LYS:HZ3	1.55	0.70
10:A:6:G:HO2'	10:A:7:G:C5'	2.03	0.70
11:AA:377:THR:OG1	12:AB:62:GLU:CG	2.38	0.70
16:AG:434:LEU:CA	16:AG:456:LEU:CD2	2.58	0.70
26:L:42:TRP:CH2	26:L:102:MET:CG	2.73	0.70
39:Y:81:LYS:HD2	39:Y:81:LYS:N	2.05	0.70
41:a:987:C:O3'	46:f:11:ARG:NH1	2.23	0.70
52:l:6:LYS:NZ	52:l:120:VAL:N	2.39	0.70
6:5:9:DG:OP2	11:AA:470:ARG:HA	1.90	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:122:GLN:HG2	9:9:123:ILE:N	2.06	0.70
12:AB:138:ARG:CD	12:AB:155:LYS:N	2.53	0.70
23:I:147:LYS:NZ	23:I:204:LYS:O	2.22	0.70
5:4:77:VAL:CG2	5:4:79:ARG:HE	2.03	0.70
11:AA:846:GLY:CA	16:AG:104:ARG:CZ	2.63	0.70
12:AB:138:ARG:HG3	12:AB:154:VAL:C	2.16	0.70
14:AE:79:LYS:CA	16:AG:144:LYS:HZ1	1.95	0.70
16:AG:389:MET:O	16:AG:404:GLU:OE2	2.09	0.70
16:AG:422:GLU:CB	16:AG:426:GLY:HA3	2.21	0.70
29:O:112:GLU:OE2	29:O:115:LYS:NZ	2.23	0.70
41:a:2682:A:C8	50:j:11:MET:HE2	2.25	0.70
4:3:46:GLN:NE2	4:3:47:LYS:O	2.24	0.70
9:9:6:GLN:HA	9:9:9:GLN:NE2	2.06	0.70
9:9:43:LYS:HZ2	9:9:101:LYS:HE2	1.55	0.70
11:AA:856:ASN:CG	16:AG:35:THR:CB	2.64	0.70
12:AB:6:LEU:HD13	14:AE:291:ILE:CD1	2.21	0.70
12:AB:18:GLN:HA	12:AB:21:LEU:HB2	1.74	0.70
16:AG:353:HIS:HA	16:AG:356:ILE:CG2	2.22	0.70
29:O:7:TYR:N	29:O:89:GLU:OE2	2.24	0.70
39:Y:100:ILE:HG13	39:Y:139:VAL:HG23	1.74	0.70
41:a:871:U:H2'	41:a:872:U:C6	2.26	0.70
11:AA:860:ALA:H	16:AG:38:LYS:HE2	1.56	0.70
16:AG:282:GLN:NE2	16:AG:282:GLN:HA	2.04	0.70
18:D:375:U:OP1	35:U:70:ARG:NE	2.24	0.70
38:X:71:ARG:CZ	38:X:72:GLU:HG3	2.22	0.70
41:a:1012:U:O4	59:s:30:THR:HG21	1.92	0.70
41:a:2350:C:OP1	55:o:45:ARG:NH2	2.25	0.70
54:n:56:ASP:HB3	54:n:150:ARG:NH2	2.07	0.70
58:r:33:GLN:OE1	58:r:35:LYS:NZ	2.24	0.70
11:AA:481:LEU:CD2	12:AB:16:ARG:HD2	2.12	0.70
18:D:274:A:H4'	36:V:16:LYS:CE	2.22	0.70
18:D:311:C:P	35:U:31:ARG:HH22	2.13	0.70
25:K:93:ARG:HB3	25:K:93:ARG:CZ	2.21	0.70
38:X:66:GLU:OE1	38:X:66:GLU:N	2.23	0.70
39:Y:116:MET:CG	39:Y:124:MET:HG2	2.21	0.70
56:p:107:LEU:O	56:p:152:ARG:NH1	2.25	0.70
62:v:105:MET:SD	62:v:117:PHE:CZ	2.85	0.70
2:1:7:HIS:CD2	2:1:10:ALA:HB2	2.27	0.70
12:AB:118:ILE:HG13	12:AB:142:LEU:CD2	2.22	0.70
12:AB:133:PRO:HA	12:AB:159:PHE:CZ	2.26	0.70
18:D:564:C:P	32:R:12:ARG:NH2	2.64	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:R:7:LEU:HD21	32:R:12:ARG:NH2	2.07	0.70
11:AA:860:ALA:N	16:AG:38:LYS:HZ3	1.89	0.70
17:C:60:LYS:NZ	18:D:734:G:O3'	2.20	0.70
17:C:60:LYS:HZ2	18:D:734:G:H2'	1.57	0.70
41:a:1653:G:H3'	63:w:2:ARG:HD2	1.74	0.70
41:a:1816:C:N4	48:h:35:GLU:OE2	2.20	0.70
51:k:21:TYR:OH	51:k:39:PHE:O	2.10	0.70
62:v:53:MET:HE3	62:v:120:ALA:HB2	1.74	0.70
59:s:15:TRP:NE1	59:s:53:TYR:HD2	1.90	0.70
59:s:73:VAL:C	59:s:74:TYR:CD2	2.70	0.70
39:Y:19:PRO:HG2	39:Y:23:VAL:HG23	1.73	0.69
41:a:2742:G:OP1	57:q:36:ARG:NH1	2.25	0.69
47:g:16:CYS:CB	47:g:37:CYS:HB3	2.21	0.69
9:9:44:ALA:HB1	9:9:52:MET:HB3	1.74	0.69
9:9:52:MET:HE1	9:9:85:SER:C	2.17	0.69
19:E:45:ALA:O	19:E:48:GLN:HG3	1.92	0.69
30:P:47:GLU:N	30:P:47:GLU:OE1	2.23	0.69
38:X:28:THR:HG23	38:X:31:LYS:CE	2.22	0.69
63:w:42:LYS:N	63:w:42:LYS:HD3	2.05	0.69
11:AA:855:PRO:CD	16:AG:105:ILE:HG21	2.22	0.69
14:AE:1033:GLY:HA3	14:AE:1081:VAL:O	1.93	0.69
27:M:66:LEU:HG	27:M:70:ARG:HE	1.58	0.69
54:n:69:LYS:HD3	54:n:84:PRO:HA	1.74	0.69
57:q:25:VAL:HB	57:q:35:GLN:OE1	1.92	0.69
7:6:25:DT:H2'	11:AA:496:LYS:HG3	1.73	0.69
12:AB:104:ILE:O	12:AB:111:TYR:OH	2.10	0.69
16:AG:235:LYS:HZ2	16:AG:331:LEU:HG	1.58	0.69
19:E:25:ARG:CD	19:E:66:LEU:HD11	2.22	0.69
23:I:24:ALA:HB3	23:I:29:PHE:CE2	2.27	0.69
23:I:142:MET:HE1	23:I:148:GLY:HA2	1.72	0.69
34:T:21:ASP:O	34:T:22:THR:OG1	2.07	0.69
39:Y:100:ILE:CG1	39:Y:139:VAL:HB	2.22	0.69
39:Y:102:ARG:HD2	39:Y:141:ASP:HA	1.75	0.69
16:AG:285:ILE:CD1	16:AG:293:VAL:HG11	2.22	0.69
16:AG:433:ASP:O	16:AG:437:LEU:HG	1.93	0.69
10:B:58:A:O2'	10:B:59:A:H3'	1.93	0.69
19:E:12:ILE:O	19:E:15:GLU:HG3	1.92	0.69
25:K:111:MET:CE	25:K:125:ALA:HB1	2.23	0.69
39:Y:91:LYS:NZ	39:Y:97:VAL:HG21	2.07	0.69
16:AG:60:ARG:HG3	16:AG:98:GLU:HG3	1.74	0.69
34:T:28:GLN:O	34:T:32:LEU:HG	1.93	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:g:5:ILE:HB	54:n:64:LYS:HD3	1.75	0.69
8:7:56:G:OP1	11:AA:1073:LYS:NZ	2.24	0.69
10:A:37:A:O2'	10:A:38:A:H5'	1.93	0.69
12:AB:102:LYS:HE2	14:AE:285:LEU:HA	1.74	0.69
16:AG:279:ASN:CB	16:AG:280:PRO:HD2	2.22	0.69
16:AG:321:ASN:OD1	16:AG:321:ASN:N	2.25	0.69
16:AG:453:VAL:CG2	16:AG:462:GLN:NE2	2.56	0.69
18:D:636:U:H4'	36:V:6:ARG:NH2	2.06	0.69
31:Q:52:PHE:HD2	31:Q:56:ARG:HG2	1.56	0.69
32:R:27:CYS:HB2	32:R:30:LYS:HE2	1.75	0.69
8:7:8:U:OP2	18:D:1397:C:N3	2.25	0.69
11:AA:854:ILE:O	16:AG:102:PHE:CD2	2.45	0.69
16:AG:266:LEU:H	16:AG:266:LEU:CD2	2.06	0.69
16:AG:462:GLN:OE1	16:AG:467:LEU:HD21	1.93	0.69
17:C:38:LYS:NZ	18:D:718:A:C2	2.57	0.69
19:E:9:LYS:C	19:E:13:GLN:HE22	2.00	0.69
21:G:27:MET:SD	21:G:30:PHE:HB2	2.33	0.69
21:G:100:MET:HE1	21:G:101:LEU:HD21	1.75	0.69
23:I:134:MET:HE3	23:I:151:VAL:HG23	1.75	0.69
33:S:20:TYR:HE2	33:S:52:PRO:HG2	1.58	0.69
40:Z:2:ILE:HG22	40:Z:5:ASP:H	1.58	0.69
41:a:2314:A:H1'	54:n:155:THR:HG21	1.74	0.69
41:a:2511:U:H4'	50:j:130:GLN:OE1	1.91	0.69
52:l:17:THR:HA	52:l:21:ARG:HH21	1.58	0.69
54:n:127:ASN:OD1	54:n:158:THR:N	2.18	0.69
63:w:45:ARG:O	63:w:49:GLU:OE1	2.11	0.69
5:4:53:LYS:HD2	5:4:56:PHE:CE2	2.28	0.69
9:9:92:ALA:HB1	9:9:129:LEU:HD22	1.74	0.69
16:AG:285:ILE:HG13	16:AG:293:VAL:HG13	1.71	0.69
16:AG:355:ALA:C	16:AG:380:THR:HG21	2.18	0.69
10:B:37:A:O2'	10:B:38:A:H5'	1.93	0.69
17:C:60:LYS:HZ1	18:D:735:C:C5'	2.05	0.69
18:D:1308:U:OP1	38:X:100:GLN:NE2	2.26	0.69
21:G:26:LYS:HD2	21:G:193:PRO:HG2	1.74	0.69
22:H:22:SER:N	22:H:69:ASP:OD2	2.26	0.69
52:l:101:TYR:CE2	52:l:105:LEU:HD11	2.28	0.69
62:v:1:MET:HE1	62:v:43:ALA:C	2.18	0.69
11:AA:65:ASN:HB3	11:AA:105:TYR:HB2	1.73	0.69
11:AA:887:VAL:CB	16:AG:105:ILE:CG2	2.71	0.69
18:D:1376:U:OP2	27:M:25:LYS:NZ	2.24	0.69
26:L:9:MET:HE1	26:L:85:ILE:CG2	2.22	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:860:ALA:H	16:AG:38:LYS:HD3	1.54	0.68
16:AG:107:THR:HA	16:AG:110:ALA:HB3	1.74	0.68
10:B:6:G:HO2'	10:B:7:G:C5'	2.06	0.68
21:G:126:PHE:O	21:G:127:ASP:O	2.10	0.68
26:L:35:LYS:NZ	26:L:36:ILE:O	2.26	0.68
36:V:16:LYS:HD3	36:V:16:LYS:C	2.17	0.68
39:Y:19:PRO:O	39:Y:24:GLY:HA2	1.92	0.68
41:a:1190:G:OP1	61:u:30:THR:OG1	2.10	0.68
48:h:133:ARG:NE	48:h:187:ASP:OD2	2.25	0.68
61:u:55:MET:HE1	61:u:60:ARG:HG2	1.75	0.68
16:AG:171:GLU:OE1	16:AG:267:GLY:HA3	1.92	0.68
16:AG:425:LEU:CB	16:AG:429:LYS:CE	2.71	0.68
22:H:308:GLU:O	22:H:310:ASP:N	2.26	0.68
39:Y:23:VAL:HG23	39:Y:24:GLY:H	1.58	0.68
61:u:123:ARG:NH2	61:u:143:GLU:OE2	2.25	0.68
10:A:58:A:O2'	10:A:59:A:H3'	1.93	0.68
12:AB:136:GLU:HA	12:AB:159:PHE:HE1	1.57	0.68
16:AG:102:PHE:HB2	16:AG:106:THR:HG21	1.75	0.68
20:F:8:GLU:OE2	20:F:9:ASN:ND2	2.26	0.68
27:M:4:ARG:HG2	27:M:5:ARG:NH1	2.08	0.68
27:M:109:ARG:HD2	27:M:116:MET:HE1	1.76	0.68
46:f:16:ARG:NE	46:f:54:MET:SD	2.66	0.68
62:v:1:MET:HE1	62:v:44:ARG:N	2.09	0.68
62:v:111:GLU:O	62:v:115:GLU:OE1	2.10	0.68
1:0:52:PRO:HB3	66:z:89:GLU:OE2	1.94	0.68
12:AB:140:MET:O	12:AB:150:ILE:CB	2.38	0.68
14:AE:59:ALA:HB2	14:AE:71:LEU:HD21	1.74	0.68
16:AG:434:LEU:HD11	16:AG:455:THR:C	2.17	0.68
48:h:180:GLU:C	48:h:181:MET:HE2	2.17	0.68
60:t:103:VAL:HG22	60:t:104:THR:H	1.58	0.68
63:w:38:LEU:HD13	63:w:111:ALA:CB	2.24	0.68
19:E:79:LEU:HA	19:E:82:GLN:OE1	1.94	0.68
20:F:12:PHE:CE1	31:Q:107:ILE:HG22	2.29	0.68
25:K:115:LEU:HD12	25:K:120:VAL:HG11	1.75	0.68
48:h:205:LEU:HD12	48:h:210:ALA:HB1	1.75	0.68
60:t:5:GLN:HA	60:t:20:MET:HE2	1.74	0.68
16:AG:171:GLU:OE2	16:AG:267:GLY:CA	2.41	0.68
16:AG:207:GLU:HA	16:AG:210:ARG:HB2	1.76	0.68
16:AG:298:VAL:HA	16:AG:303:HIS:HD1	1.58	0.68
16:AG:323:GLN:HA	16:AG:323:GLN:HE21	1.59	0.68
16:AG:440:VAL:HG23	16:AG:481:LEU:HD22	1.76	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:24:U:O2'	41:a:1922:G:O3'	2.12	0.68
10:B:38:A:O2'	18:D:790:A:OP1	2.09	0.68
22:H:304:VAL:O	22:H:304:VAL:HG12	1.94	0.68
33:S:19:LYS:HG3	33:S:20:TYR:CD1	2.29	0.68
19:E:78:ASN:O	19:E:82:GLN:OE1	2.10	0.68
34:T:76:ALA:C	34:T:80:GLN:HE22	2.00	0.68
38:X:90:ARG:CD	38:X:97:VAL:HA	2.23	0.68
2:1:19:LEU:HD21	49:i:22:LEU:HB2	1.75	0.68
5:4:63:ILE:HD11	5:4:91:PHE:CD2	2.28	0.68
5:4:89:ILE:HG21	5:4:91:PHE:CE1	2.29	0.68
26:L:42:TRP:HZ2	26:L:102:MET:SD	2.17	0.68
39:Y:116:MET:HG2	39:Y:124:MET:CG	2.24	0.68
41:a:1056:G:O2'	41:a:1103:A:N6	2.27	0.68
9:9:43:LYS:HZ3	9:9:98:GLU:HB2	1.58	0.68
12:AB:137:ALA:O	12:AB:155:LYS:CA	2.41	0.68
13:AD:48:LEU:HB2	13:AD:183:ILE:HD11	1.75	0.68
31:Q:80:LYS:HB3	31:Q:106:ARG:NH1	2.09	0.68
33:S:42:TRP:CZ3	47:g:66:ILE:HG23	2.29	0.68
34:T:27:VAL:O	34:T:31:LEU:HD23	1.93	0.68
58:r:37:VAL:HG12	58:r:43:ASN:ND2	2.09	0.68
62:v:1:MET:HE1	62:v:44:ARG:HA	1.76	0.68
12:AB:89:PRO:CG	14:AE:290:ILE:HG21	2.24	0.68
18:D:1308:U:P	38:X:100:GLN:HE22	2.16	0.68
28:N:87:LYS:HD2	28:N:91:GLU:HB3	1.74	0.68
37:W:36:ARG:NH2	37:W:75:ALA:O	2.26	0.68
39:Y:118:GLY:O	39:Y:124:MET:HE3	1.92	0.68
41:a:1655:A:HI1'	50:j:118:PHE:CE2	2.29	0.68
5:4:50:MET:HA	5:4:56:PHE:CZ	2.29	0.67
12:AB:34:THR:HA	12:AB:47:SER:HA	1.76	0.67
10:B:75:C:P	41:a:2602:A:HO2'	2.12	0.67
19:E:25:ARG:HD2	19:E:66:LEU:HD11	1.75	0.67
24:J:76:TYR:HE2	24:J:204:TYR:HB2	1.57	0.67
27:M:20:SER:CB	27:M:23:LEU:HD13	2.24	0.67
63:w:24:MET:HE2	63:w:34:ILE:CD1	2.24	0.67
9:9:24:SER:CB	9:9:116:GLU:HG3	2.24	0.67
9:9:43:LYS:NZ	9:9:101:LYS:HE2	2.09	0.67
9:9:43:LYS:HA	9:9:46:ARG:CZ	2.24	0.67
9:9:44:ALA:HA	9:9:47:GLU:OE1	1.95	0.67
11:AA:887:VAL:CB	16:AG:105:ILE:HG23	2.23	0.67
26:L:38:ARG:NH1	26:L:99:ALA:HB2	2.09	0.67
27:M:23:LEU:HD21	27:M:59:LEU:HD21	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Q:72:ASP:O	31:Q:75:LYS:NZ	2.21	0.67
41:a:752:A:OP1	53:m:1:MET:HE1	1.95	0.67
16:AG:239:LYS:HG3	16:AG:276:TRP:HB3	1.75	0.67
22:H:23:ILE:N	22:H:69:ASP:OD2	2.27	0.67
39:Y:37:PHE:CZ	39:Y:56:VAL:HG11	2.29	0.67
52:l:6:LYS:HE3	52:l:119:ILE:HG23	1.75	0.67
60:t:112:PHE:HD2	60:t:115:ILE:HD12	1.59	0.67
34:T:80:GLN:O	34:T:83:GLU:HG3	1.95	0.67
41:a:927:A:H2'	41:a:928:A:C8	2.30	0.67
41:a:2682:A:O4'	50:j:11:MET:HE1	1.95	0.67
66:z:68:ALA:HB1	66:z:106:PHE:CE2	2.29	0.67
17:C:13:PHE:HB3	17:C:51:TYR:CE1	2.30	0.67
27:M:46:ALA:CB	27:M:120:LEU:HD22	2.25	0.67
38:X:90:ARG:HD2	38:X:97:VAL:HA	1.77	0.67
56:p:85:LYS:CE	56:p:164:TYR:HD1	2.07	0.67
9:9:73:LYS:HG2	9:9:117:LEU:HD12	1.76	0.67
12:AB:27:ASN:CG	12:AB:58:GLU:HB2	2.20	0.67
12:AB:120:GLU:HB2	12:AB:162:LEU:HB2	1.77	0.67
16:AG:284:VAL:CG1	16:AG:296:ILE:HD13	2.25	0.67
16:AG:353:HIS:C	16:AG:357:ASP:H	2.01	0.67
18:D:274:A:H4'	36:V:16:LYS:HE3	1.76	0.67
26:L:18:VAL:HA	26:L:21:MET:HE2	1.75	0.67
27:M:111:ARG:O	27:M:119:ARG:NH2	2.27	0.67
39:Y:48:ILE:HG13	39:Y:49:GLU:N	2.10	0.67
46:f:40:ASP:C	46:f:45:ARG:HH22	2.03	0.67
46:f:41:THR:O	46:f:44:ILE:HG22	1.94	0.67
62:v:45:GLN:NE2	62:v:125:PRO:HD3	2.10	0.67
8:7:8:U:H5'	18:D:1397:C:N3	2.10	0.67
16:AG:453:VAL:HG13	16:AG:458:ASP:HB3	1.77	0.67
22:H:163:ARG:N	22:H:298:GLU:OE2	2.28	0.67
4:3:72:ILE:HD12	4:3:83:VAL:HG21	1.76	0.67
16:AG:228:ARG:HA	16:AG:327:LEU:CD1	2.22	0.67
23:I:134:MET:HE1	23:I:167:TRP:HA	1.76	0.67
57:q:1:MET:HE1	57:q:35:GLN:CA	2.25	0.67
16:AG:168:LEU:HB2	16:AG:171:GLU:HB2	1.76	0.67
16:AG:302:LYS:H	16:AG:302:LYS:CE	2.00	0.67
16:AG:363:LEU:CD2	16:AG:409:ARG:CB	2.62	0.67
16:AG:447:LYS:O	16:AG:470:ILE:HD13	1.95	0.67
10:B:75:C:OP2	41:a:2602:A:C2'	2.43	0.67
18:D:564:C:OP1	32:R:12:ARG:NH1	2.27	0.67
18:D:945:G:C2	18:D:946:A:C8	2.83	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:F:31:GLU:OE1	20:F:34:ARG:NH2	2.28	0.67
28:N:67:GLN:OE1	28:N:67:GLN:N	2.28	0.67
39:Y:85:ILE:H	39:Y:85:ILE:HD12	1.59	0.67
41:a:1802:A:H2'	41:a:1803:A:C8	2.30	0.67
41:a:2682:A:C8	50:j:11:MET:HE1	2.30	0.67
16:AG:380:THR:O	16:AG:384:LEU:HD12	1.95	0.67
18:D:1344:C:P	29:O:124:ARG:HH11	2.16	0.67
25:K:156:LYS:HE3	28:N:71:VAL:HG13	1.76	0.67
41:a:1665:A:O5'	60:t:66:LYS:NZ	2.28	0.67
52:l:118:LEU:HD11	52:l:188:MET:SD	2.35	0.67
5:4:84:PRO:O	5:4:85:LYS:HE2	1.96	0.66
10:A:71:C:H3'	10:A:72:A:C8	2.31	0.66
16:AG:235:LYS:NZ	16:AG:331:LEU:HG	2.11	0.66
16:AG:429:LYS:CD	16:AG:429:LYS:H	2.05	0.66
17:C:70:TYR:CE2	18:D:674:G:H4'	2.30	0.66
18:D:673:A:H5''	26:L:86:ARG:NH1	2.09	0.66
29:O:86:ALA:O	29:O:89:GLU:HG2	1.95	0.66
33:S:42:TRP:HH2	47:g:66:ILE:HA	1.60	0.66
35:U:45:GLU:HG3	35:U:46:LYS:HD3	1.76	0.66
39:Y:32:VAL:HG22	39:Y:60:VAL:HG21	1.77	0.66
41:a:2420:C:H5''	51:k:8:LYS:NZ	2.09	0.66
59:s:13:ARG:HH22	59:s:50:THR:HA	1.60	0.66
60:t:42:THR:CG2	60:t:44:LYS:HZ1	2.07	0.66
7:6:20:DA:N6	8:7:51:G:N1	2.42	0.66
10:A:32:C:H4'	27:M:86:GLN:NE2	2.10	0.66
10:B:71:C:H3'	10:B:72:A:C8	2.30	0.66
24:J:121:LYS:HG2	24:J:131:ASN:HD21	1.59	0.66
41:a:1454:C:O4'	63:w:63:ARG:NE	2.28	0.66
5:4:50:MET:HB3	5:4:56:PHE:CE1	2.31	0.66
9:9:47:GLU:HG3	9:9:94:ARG:CZ	2.24	0.66
19:E:54:MET:HE1	19:E:75:HIS:CE1	2.31	0.66
23:I:151:VAL:HG12	23:I:200:VAL:HG23	1.77	0.66
43:c:71:LEU:HB3	43:c:76:GLU:CD	2.20	0.66
48:h:53:HIS:NE2	48:h:219:THR:HA	2.11	0.66
50:j:3:GLY:O	50:j:4:LEU:HD12	1.95	0.66
59:s:15:TRP:NE1	59:s:53:TYR:CD2	2.63	0.66
5:4:75:GLN:HG3	5:4:92:VAL:CG2	2.25	0.66
9:9:23:LEU:HD13	9:9:92:ALA:HB2	1.77	0.66
12:AB:115:LYS:HE2	12:AB:129:ILE:HG23	1.76	0.66
39:Y:79:LEU:HD21	39:Y:105:LEU:CD2	2.24	0.66
48:h:76:ALA:HB1	48:h:94:VAL:HG12	1.76	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:z:79:PHE:HE1	66:z:110:VAL:HA	1.60	0.66
21:G:101:LEU:HD22	21:G:157:LEU:CD1	2.26	0.66
41:a:111:A:O3'	45:e:58:ASN:ND2	2.28	0.66
41:a:666:A:H1'	55:o:4:ILE:HD12	1.76	0.66
2:1:25:ARG:HH22	41:a:520:G:H5'	1.58	0.66
6:5:10:DT:H73	11:AA:62:TYR:CB	2.25	0.66
16:AG:227:ALA:HB3	16:AG:331:LEU:HA	1.78	0.66
16:AG:308:ALA:HB2	16:AG:344:LEU:HD11	1.78	0.66
18:D:43:C:OP1	35:U:12:LYS:NZ	2.29	0.66
21:G:49:MET:HE1	21:G:201:PRO:CD	2.23	0.66
21:G:90:PHE:CD2	21:G:154:MET:HE3	2.31	0.66
22:H:308:GLU:HG2	22:H:309:MET:H	1.61	0.66
38:X:81:MET:CE	41:a:888:C:C5	2.78	0.66
40:Z:20:VAL:O	40:Z:20:VAL:HG12	1.94	0.66
41:a:2420:C:H5''	51:k:8:LYS:CE	2.25	0.66
60:t:38:ILE:HD11	60:t:112:PHE:HZ	1.60	0.66
60:t:99:ILE:HD13	60:t:118:LEU:HB2	1.77	0.66
5:4:56:PHE:O	5:4:61:LEU:HD11	1.95	0.66
6:5:10:DT:C7	11:AA:62:TYR:CB	2.74	0.66
9:9:24:SER:OG	9:9:86:MET:HE2	1.96	0.66
9:9:139:LEU:HD21	40:Z:11:VAL:HG22	1.76	0.66
16:AG:213:VAL:HG22	16:AG:216:ILE:HG13	1.78	0.66
21:G:134:ALA:O	21:G:137:ARG:HG2	1.96	0.66
48:h:129:THR:HG22	48:h:189:ARG:HD2	1.77	0.66
59:s:43:GLU:OE1	59:s:43:GLU:N	2.29	0.66
12:AB:113:GLY:H	12:AB:133:PRO:HD2	1.58	0.66
16:AG:448:LEU:CD2	16:AG:473:LEU:HD12	2.24	0.66
19:E:21:ASN:C	19:E:25:ARG:NE	2.54	0.66
39:Y:24:GLY:O	39:Y:27:LEU:HG	1.96	0.66
39:Y:45:THR:HG21	39:Y:50:LYS:HD3	1.76	0.66
54:n:94:GLU:HA	54:n:97:TRP:HE3	1.58	0.66
60:t:1:MET:HE1	60:t:32:TYR:CD1	2.31	0.66
62:v:45:GLN:NE2	62:v:125:PRO:CD	2.59	0.66
9:9:28:ALA:H	9:9:110:ALA:HA	1.61	0.66
18:D:246:A:C2	18:D:282:A:C5	2.84	0.66
18:D:375:U:P	35:U:70:ARG:NE	2.69	0.66
18:D:1151:A:HO2'	18:D:1152:A:P	2.18	0.66
21:G:53:ALA:HA	21:G:56:GLU:OE1	1.96	0.66
41:a:2175:C:H2'	41:a:2176:A:C8	2.31	0.66
52:l:40:ARG:HD2	52:l:92:HIS:ND1	2.11	0.66
63:w:49:GLU:OE1	63:w:49:GLU:N	2.26	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:55:ASP:OD1	1:0:56:GLY:N	2.27	0.66
5:4:77:VAL:HG23	5:4:89:ILE:HG12	1.76	0.66
11:AA:847:PRO:O	16:AG:104:ARG:CZ	2.44	0.66
12:AB:146:ILE:HG21	30:P:88:MET:O	1.96	0.66
16:AG:232:SER:HB2	16:AG:323:GLN:OE1	1.96	0.66
23:I:45:LYS:NZ	23:I:46:GLU:OE1	2.29	0.66
29:O:5:GLN:NE2	29:O:20:PHE:HB3	2.10	0.66
41:a:2355:G:C2'	41:a:2356:U:H5'	2.26	0.66
9:9:43:LYS:HD3	9:9:98:GLU:HG3	1.78	0.65
16:AG:133:HIS:ND1	16:AG:136:GLU:CD	2.55	0.65
16:AG:355:ALA:CB	16:AG:382:GLU:CG	2.54	0.65
17:C:51:TYR:CD1	17:C:54:GLN:NE2	2.63	0.65
28:N:10:MET:HG3	28:N:27:MET:HE1	1.78	0.65
41:a:998:C:OP2	66:z:92:ARG:NH2	2.29	0.65
52:l:149:ILE:HD11	52:l:188:MET:HG2	1.78	0.65
10:A:76:A:N6	41:a:2422:C:O5'	2.29	0.65
14:AE:58:CYS:SG	14:AE:59:ALA:N	2.69	0.65
16:AG:123:ARG:HG2	16:AG:191:ARG:O	1.96	0.65
18:D:876:C:H1'	28:N:12:THR:HG21	1.76	0.65
21:G:23:TRP:CD1	21:G:39:HIS:HE1	2.14	0.65
21:G:90:PHE:CE2	21:G:154:MET:HE3	2.31	0.65
24:J:196:ASN:OD1	24:J:198:HIS:CE1	2.48	0.65
39:Y:20:SER:HB2	39:Y:21:PRO:HD3	1.79	0.65
6:5:9:DG:O5'	11:AA:473:ARG:HB3	1.95	0.65
9:9:139:LEU:CD2	40:Z:11:VAL:HG22	2.26	0.65
10:A:32:C:H4'	27:M:86:GLN:HE21	1.62	0.65
16:AG:393:LEU:O	16:AG:398:LEU:HD23	1.95	0.65
22:H:303:LEU:O	22:H:303:LEU:HG	1.96	0.65
26:L:25:TYR:O	26:L:29:ILE:HD12	1.97	0.65
33:S:53:ARG:HB3	33:S:59:ARG:NH1	2.11	0.65
33:S:87:ALA:CA	33:S:90:ARG:NH1	2.60	0.65
37:W:62:VAL:HG13	37:W:66:MET:HE3	1.77	0.65
66:z:85:LYS:HD2	66:z:116:ALA:HB1	1.79	0.65
9:9:14:GLU:O	9:9:18:VAL:HG23	1.96	0.65
12:AB:9:CYS:HB3	12:AB:74:GLY:HA2	1.79	0.65
12:AB:35:LEU:CA	12:AB:106:ASP:OD1	2.36	0.65
52:l:147:LEU:HB2	52:l:183:PHE:HD2	1.62	0.65
52:l:194:LYS:O	52:l:198:GLU:OE1	2.15	0.65
58:r:137:GLU:HG2	58:r:138:VAL:N	2.11	0.65
6:5:10:DT:H1'	12:AB:71:ALA:H	0.83	0.65
9:9:25:ALA:HB2	9:9:96:PHE:CE1	2.31	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:46:ARG:CG	9:9:94:ARG:HH21	2.02	0.65
12:AB:115:LYS:HZ3	12:AB:129:ILE:HG22	1.56	0.65
12:AB:155:LYS:HG3	12:AB:156:ASN:H	1.62	0.65
18:D:1228:C:OP2	38:X:110:LYS:NZ	2.28	0.65
34:T:64:ARG:NH1	34:T:89:ARG:HH22	1.94	0.65
41:a:1824:G:OP1	48:h:52:ARG:CZ	2.44	0.65
41:a:2682:A:H2'	41:a:2683:C:O4'	1.96	0.65
47:g:58:ASP:C	47:g:62:LYS:NZ	2.55	0.65
60:t:42:THR:O	60:t:44:LYS:NZ	2.19	0.65
9:9:129:LEU:N	9:9:130:PRO:HD2	2.12	0.65
12:AB:118:ILE:CD1	12:AB:142:LEU:HD22	2.27	0.65
23:I:132:ARG:O	23:I:135:LYS:HG2	1.97	0.65
32:R:30:LYS:HB3	32:R:57:LEU:HD11	1.78	0.65
39:Y:14:ALA:C	39:Y:50:LYS:HE3	2.22	0.65
39:Y:81:LYS:NZ	39:Y:86:LYS:HE2	2.12	0.65
40:Z:22:LEU:O	40:Z:26:MET:HE1	1.97	0.65
41:a:617:G:H4'	52:l:199:MET:HE1	1.78	0.65
3:2:3:ARG:NH1	3:2:5:GLU:OE2	2.30	0.65
4:3:26:LYS:HB2	4:3:35:ILE:HG23	1.78	0.65
4:3:34:VAL:HG13	4:3:67:VAL:HG22	1.78	0.65
16:AG:174:ARG:HB3	16:AG:175:PRO:CD	2.25	0.65
17:C:73:ARG:HH12	31:Q:113:VAL:HA	1.62	0.65
18:D:588:G:H4'	28:N:3:MET:HE1	1.77	0.65
18:D:1290:G:H4'	27:M:35:LYS:HZ3	1.62	0.65
29:O:83:ILE:O	29:O:87:LEU:HD23	1.96	0.65
36:V:60:GLU:OE1	36:V:76:VAL:CG2	2.45	0.65
41:a:2308:G:C5	54:n:77:PHE:CZ	2.85	0.65
52:l:62:GLN:O	52:l:69:ARG:NE	2.30	0.65
60:t:42:THR:HG23	60:t:44:LYS:NZ	2.12	0.65
61:u:84:LYS:HZ2	61:u:98:ALA:C	2.02	0.65
16:AG:226:ALA:O	16:AG:228:ARG:HD3	1.97	0.65
27:M:103:TRP:CG	27:M:137:LYS:HZ1	2.14	0.65
38:X:28:THR:HA	38:X:31:LYS:HE3	1.78	0.65
41:a:1278:C:C4'	63:w:24:MET:HE1	2.27	0.65
41:a:1839:G:H1'	41:a:1927:A:C8	2.32	0.65
54:n:117:LEU:HD13	54:n:176:PRO:CG	2.26	0.65
62:v:31:PHE:N	62:v:104:GLU:OE2	2.30	0.65
9:9:25:ALA:HB3	9:9:99:PHE:HD2	1.61	0.65
10:A:19:G:C2	41:a:2112:G:O4'	2.50	0.65
16:AG:425:LEU:HG	16:AG:429:LYS:HE2	1.65	0.65
17:C:73:ARG:HH12	31:Q:113:VAL:CB	2.09	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:E:16:LYS:O	19:E:19:LYS:HG2	1.97	0.65
20:F:58:LYS:O	20:F:62:ARG:HG2	1.96	0.65
39:Y:5:GLN:HG2	39:Y:61:TYR:CD1	2.32	0.65
39:Y:102:ARG:HD2	39:Y:141:ASP:CB	2.27	0.65
62:v:111:GLU:HG2	62:v:112:LEU:HD12	1.78	0.65
34:T:53:ARG:O	34:T:57:LEU:HD23	1.97	0.65
8:7:11:U:H4'	25:K:20:ARG:HH21	1.53	0.64
9:9:47:GLU:HB3	9:9:51:TYR:OH	1.97	0.64
12:AB:136:GLU:OE1	12:AB:160:ARG:NH1	2.31	0.64
16:AG:64:VAL:HG22	16:AG:75:ILE:HD12	1.78	0.64
16:AG:393:LEU:HD23	16:AG:398:LEU:HD13	1.78	0.64
19:E:28:MET:SD	19:E:61:GLN:HG3	2.36	0.64
32:R:25:GLU:CD	32:R:59:ASN:HD22	2.05	0.64
33:S:10:GLU:O	33:S:14:VAL:HG23	1.97	0.64
39:Y:102:ARG:HG3	39:Y:140:GLU:O	1.96	0.64
54:n:93:GLY:O	54:n:96:MET:N	2.29	0.64
6:5:10:DT:H73	11:AA:62:TYR:CG	2.18	0.64
16:AG:425:LEU:O	16:AG:429:LYS:CE	2.45	0.64
18:D:280:C:H1'	36:V:40:ARG:NH2	2.11	0.64
18:D:675:A:H1'	31:Q:118:HIS:CD2	2.33	0.64
18:D:738:C:H5''	26:L:68:GLN:NE2	2.12	0.64
22:H:277:GLY:HA2	22:H:331:MET:CE	2.27	0.64
23:I:156:ARG:HH11	23:I:193:TYR:HB3	1.63	0.64
25:K:148:ASN:OD1	28:N:96:MET:CE	2.45	0.64
41:a:2419:U:H4'	51:k:22:THR:HG21	1.79	0.64
41:a:2572:A:N7	50:j:150:GLN:NE2	2.45	0.64
52:l:112:LEU:HG	52:l:118:LEU:HB2	1.79	0.64
56:p:149:ARG:CZ	56:p:162:VAL:O	2.45	0.64
60:t:1:MET:HE1	60:t:32:TYR:CG	2.31	0.64
8:7:11:U:C4'	25:K:20:ARG:HH22	1.93	0.64
16:AG:183:LEU:HA	16:AG:197:VAL:HA	1.78	0.64
16:AG:349:GLN:C	16:AG:351:GLU:H	2.06	0.64
22:H:119:GLY:HA2	22:H:133:GLY:HA2	1.80	0.64
41:a:2334:U:H4'	41:a:2335:A:OP2	1.95	0.64
55:o:30:ARG:HH12	61:u:62:PRO:CB	2.10	0.64
60:t:42:THR:HG23	60:t:44:LYS:HZ1	1.62	0.64
65:y:64:ILE:HG13	65:y:64:ILE:O	1.97	0.64
14:AE:79:LYS:NZ	16:AG:153:ASP:OD1	2.30	0.64
37:W:40:ILE:CD1	37:W:66:MET:SD	2.77	0.64
41:a:2723:C:H5''	63:w:1:MET:HE3	1.78	0.64
54:n:135:GLN:HG3	54:n:150:ARG:NH2	2.12	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:t:9:ASN:OD1	60:t:10:VAL:N	2.27	0.64
14:AE:84:ILE:HG22	14:AE:91:GLU:HG3	1.79	0.64
16:AG:353:HIS:CA	16:AG:356:ILE:CG2	2.76	0.64
16:AG:437:LEU:CD2	16:AG:456:LEU:HD11	2.27	0.64
17:C:60:LYS:HZ3	18:D:734:G:C3'	2.10	0.64
19:E:81:ALA:C	19:E:85:LYS:HZ3	2.04	0.64
20:F:10:GLU:OE2	20:F:18:ARG:NE	2.30	0.64
20:F:58:LYS:O	20:F:62:ARG:CG	2.45	0.64
24:J:99:ASP:OD1	24:J:100:ASN:N	2.31	0.64
27:M:26:PHE:HD1	27:M:101:MET:SD	2.19	0.64
33:S:87:ALA:HA	33:S:90:ARG:CZ	2.27	0.64
39:Y:126:ARG:NH2	41:a:1083:U:P	2.70	0.64
48:h:52:ARG:HE	48:h:53:HIS:HE1	1.44	0.64
56:p:52:PHE:CD1	56:p:69:ARG:HD3	2.32	0.64
8:7:47:U:C5'	11:AA:1259:LEU:HD21	2.27	0.64
9:9:73:LYS:NZ	9:9:115:GLY:O	2.31	0.64
12:AB:38:ILE:HD12	12:AB:160:ARG:HH11	1.63	0.64
12:AB:115:LYS:HZ2	12:AB:129:ILE:HG21	0.77	0.64
12:AB:138:ARG:HD3	12:AB:155:LYS:CA	2.27	0.64
16:AG:205:LEU:HA	16:AG:208:LEU:HB2	1.80	0.64
16:AG:361:LYS:CD	16:AG:417:ILE:CG1	2.75	0.64
16:AG:434:LEU:HD12	16:AG:454:CYS:C	2.22	0.64
18:D:1158:C:O2'	21:G:132:LYS:CE	2.45	0.64
36:V:31:HIS:CD2	36:V:34:TYR:H	2.15	0.64
39:Y:55:PRO:HB2	39:Y:71:LYS:HE2	1.80	0.64
39:Y:86:LYS:HD3	39:Y:86:LYS:C	2.23	0.64
52:l:45:ALA:HB1	52:l:88:ARG:NH2	2.12	0.64
61:u:122:VAL:CG1	61:u:125:LEU:HD11	2.27	0.64
62:v:105:MET:CE	62:v:108:VAL:HG21	2.28	0.64
2:1:25:ARG:NH2	41:a:520:G:C5'	2.60	0.64
10:A:22:G:H2'	10:A:23:C:C6	2.32	0.64
12:AB:38:ILE:HD13	12:AB:160:ARG:NH1	2.12	0.64
16:AG:354:ALA:CA	16:AG:357:ASP:H	2.10	0.64
16:AG:433:ASP:O	16:AG:456:LEU:CG	2.45	0.64
18:D:718:A:H5'	31:Q:119:ASN:ND2	2.13	0.64
24:J:138:SER:O	24:J:141:ASP:CG	2.40	0.64
25:K:61:GLN:HA	25:K:64:MET:CE	2.27	0.64
27:M:23:LEU:HD21	27:M:59:LEU:CD2	2.27	0.64
27:M:23:LEU:HD12	27:M:23:LEU:N	2.13	0.64
39:Y:99:LYS:NZ	39:Y:140:GLU:OE2	2.26	0.64
44:d:55:U:H1'	54:n:26:MET:CE	2.26	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5:11:DA:P	12:AB:67:THR:C	2.79	0.64
16:AG:131:ARG:HG2	16:AG:186:VAL:HG12	1.79	0.64
18:D:1308:U:P	38:X:100:GLN:NE2	2.70	0.64
18:D:1323:G:H2'	18:D:1324:A:C8	2.33	0.64
21:G:27:MET:HG2	21:G:189:THR:HA	1.78	0.64
41:a:2356:U:O3'	42:b:20:ARG:CZ	2.45	0.64
52:l:6:LYS:HE3	52:l:119:ILE:CG2	2.28	0.64
52:l:69:ARG:NH1	52:l:69:ARG:HA	2.13	0.64
60:t:8:LEU:N	60:t:18:ARG:NH1	2.46	0.64
6:5:16:DG:H1	11:AA:451:ARG:NH2	1.94	0.64
8:7:9:U:N3	18:D:1196:A:C8	2.66	0.64
9:9:117:LEU:O	9:9:117:LEU:HD23	1.98	0.64
10:B:22:G:H2'	10:B:23:C:C6	2.32	0.64
26:L:2:ARG:CZ	26:L:68:GLN:NE2	2.60	0.64
27:M:4:ARG:CG	27:M:5:ARG:CZ	2.75	0.64
34:T:3:LEU:HD22	34:T:8:THR:CG2	2.28	0.64
39:Y:102:ARG:NE	39:Y:102:ARG:HA	2.11	0.64
8:7:9:U:N3	18:D:1196:A:N7	2.45	0.64
17:C:34:THR:N	17:C:38:LYS:O	2.30	0.64
21:G:19:GLN:HG3	22:H:76:GLU:HB3	1.79	0.64
24:J:150:LYS:HE2	24:J:178:MET:HE2	1.79	0.64
43:c:66:THR:HG22	43:c:70:GLU:OE2	1.97	0.64
59:s:53:TYR:CE1	59:s:121:LYS:HD2	2.32	0.64
66:z:41:LYS:HG3	66:z:45:TYR:CE2	2.32	0.64
9:9:47:GLU:HG2	9:9:95:LEU:HD21	1.78	0.63
21:G:26:LYS:NZ	21:G:193:PRO:HG2	2.12	0.63
22:H:19:ARG:O	22:H:72:LEU:HB2	1.98	0.63
41:a:1067:A:O2'	41:a:1068:G:O4'	2.16	0.63
41:a:1596:A:H2'	41:a:1597:A:C8	2.32	0.63
41:a:2013:A:N1	41:a:2613:U:O4	2.31	0.63
56:p:107:LEU:O	56:p:152:ARG:CZ	2.45	0.63
64:x:114:GLY:O	64:x:116:GLN:NE2	2.31	0.63
1:0:51:VAL:HG23	66:z:86:ALA:O	1.98	0.63
5:4:80:HIS:CE1	5:4:82:TYR:H	2.17	0.63
7:6:23:DC:H5''	14:AE:259:ARG:NH1	1.92	0.63
16:AG:63:LEU:HD22	16:AG:92:TYR:HD1	1.63	0.63
16:AG:385:ALA:HB2	16:AG:411:LYS:HE3	1.70	0.63
17:C:33:ILE:O	22:H:338:GLU:CD	2.42	0.63
18:D:406:G:H21	24:J:116:GLN:HE22	1.46	0.63
39:Y:102:ARG:NE	39:Y:139:VAL:HG21	2.13	0.63
41:a:2335:A:P	64:x:13:ARG:CZ	2.87	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:50:MET:O	5:4:52:ALA:N	2.31	0.63
9:9:24:SER:OG	9:9:116:GLU:HG3	1.97	0.63
9:9:50:VAL:HG23	39:Y:120:ASP:OD2	1.98	0.63
16:AG:363:LEU:CD2	16:AG:409:ARG:CA	2.71	0.63
18:D:714:G:H2'	18:D:715:A:C8	2.32	0.63
39:Y:99:LYS:NZ	39:Y:138:VAL:HG13	2.14	0.63
41:a:2393:U:P	55:o:30:ARG:NE	2.70	0.63
2:1:71:VAL:HB	2:1:74:ILE:HD11	1.79	0.63
4:3:68:SER:OG	41:a:335:C:O2	2.15	0.63
11:AA:120:GLN:NE2	11:AA:490:GLN:OE1	2.31	0.63
11:AA:314:ASN:HD21	11:AA:348:SER:HA	1.62	0.63
12:AB:115:LYS:HE2	12:AB:129:ILE:CG2	2.27	0.63
16:AG:424:SER:C	16:AG:429:LYS:CG	2.72	0.63
32:R:80:ILE:HD12	32:R:104:CYS:HB2	1.79	0.63
39:Y:5:GLN:HE22	39:Y:60:VAL:N	1.96	0.63
39:Y:60:VAL:HG22	39:Y:66:PHE:CB	2.23	0.63
41:a:196:A:C2	61:u:50:PHE:HZ	2.16	0.63
41:a:2682:A:O5'	50:j:11:MET:HE1	1.98	0.63
11:AA:855:PRO:CG	16:AG:105:ILE:CG2	2.76	0.63
10:B:31:G:O3'	18:D:1340:A:O2'	2.17	0.63
18:D:1226:C:H2'	38:X:102:THR:OG1	1.98	0.63
18:D:1228:C:OP1	38:X:107:ARG:NH1	2.32	0.63
41:a:1800:C:OP2	48:h:182:ARG:NH1	2.27	0.63
61:u:84:LYS:NZ	61:u:98:ALA:O	2.20	0.63
1:0:84:ARG:NH1	41:a:813:U:OP1	2.31	0.63
12:AB:31:PRO:HB2	12:AB:50:LEU:HB3	1.80	0.63
10:B:54:U:H3	10:B:58:A:N6	1.96	0.63
10:B:76:A:H1'	41:a:2494:G:OP1	1.98	0.63
22:H:70:VAL:HG12	22:H:83:LEU:O	1.98	0.63
24:J:62:ARG:HG2	24:J:72:PHE:CZ	2.33	0.63
25:K:111:MET:HE1	25:K:125:ALA:HB1	1.80	0.63
33:S:77:PHE:HE1	33:S:93:ILE:HG21	1.63	0.63
38:X:16:VAL:HG23	38:X:17:ILE:HD12	1.80	0.63
41:a:995:C:O2	59:s:3:THR:OG1	2.17	0.63
41:a:2682:A:O4'	50:j:11:MET:CE	2.47	0.63
41:a:2725:A:C4	41:a:2727:A:C8	2.87	0.63
52:l:23:PHE:HB2	52:l:111:GLU:OE2	1.99	0.63
16:AG:425:LEU:HG	16:AG:429:LYS:HD3	1.78	0.63
17:C:13:PHE:CE2	17:C:21:ILE:HD11	2.34	0.63
38:X:16:VAL:HG12	38:X:34:LEU:HD12	1.80	0.63
39:Y:31:GLY:C	39:Y:60:VAL:HG11	2.23	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:99:LYS:HD2	39:Y:140:GLU:CD	2.22	0.63
50:j:55:LYS:HZ1	50:j:59:ARG:C	2.06	0.63
8:7:9:U:O4	18:D:1196:A:N9	2.32	0.63
9:9:56:ARG:H	9:9:56:ARG:HD3	1.63	0.63
9:9:100:ALA:HB2	9:9:125:ARG:NE	2.13	0.63
12:AB:136:GLU:O	12:AB:156:ASN:HA	1.97	0.63
16:AG:162:ILE:HD11	16:AG:199:ARG:HD2	1.79	0.63
16:AG:433:ASP:C	16:AG:456:LEU:CD1	2.72	0.63
22:H:61:GLU:HG2	22:H:62:ILE:HG23	1.81	0.63
25:K:61:GLN:HA	25:K:64:MET:HE3	1.81	0.63
26:L:102:MET:HG2	26:L:103:VAL:N	2.05	0.63
34:T:88:ARG:NH2	41:a:714:U:OP2	2.31	0.63
35:U:18:GLN:OE1	35:U:37:GLY:O	2.17	0.63
39:Y:14:ALA:HB3	39:Y:51:GLY:N	2.12	0.63
39:Y:37:PHE:CE1	39:Y:56:VAL:HG11	2.33	0.63
46:f:10:THR:HB	46:f:56:LYS:HZ2	1.61	0.63
63:w:33:ILE:CG1	63:w:112:TYR:HD1	2.11	0.63
7:6:18:DT:N3	8:7:52:A:N6	2.36	0.63
11:AA:1282:GLY:HA3	15:AF:17:PHE:HE1	1.63	0.63
16:AG:287:ALA:HB1	16:AG:331:LEU:HB3	1.81	0.63
16:AG:301:ASP:OD1	16:AG:301:ASP:N	2.32	0.63
16:AG:422:GLU:HG2	16:AG:426:GLY:HA3	1.80	0.63
21:G:49:MET:HE3	21:G:199:VAL:O	1.99	0.63
39:Y:79:LEU:CD1	39:Y:132:ALA:HA	2.29	0.63
39:Y:102:ARG:HG3	39:Y:141:ASP:HA	1.80	0.63
48:h:251:GLN:HE22	48:h:253:LYS:C	2.07	0.63
49:i:17:ARG:HA	49:i:20:ASP:OD1	1.99	0.63
64:x:90:VAL:HG21	64:x:115:LEU:HD21	1.81	0.63
6:5:10:DT:C4	11:AA:62:TYR:HD2	2.12	0.62
11:AA:888:THR:N	16:AG:105:ILE:CD1	2.62	0.62
16:AG:434:LEU:CD1	16:AG:454:CYS:C	2.72	0.62
25:K:137:VAL:O	25:K:140:THR:OG1	2.15	0.62
41:a:754:U:H2'	41:a:755:U:C6	2.34	0.62
65:y:31:TRP:CZ3	65:y:38:LYS:HD3	2.33	0.62
5:4:77:VAL:HG22	5:4:79:ARG:HE	1.64	0.62
6:5:11:DA:H5'	12:AB:67:THR:CB	2.27	0.62
9:9:77:VAL:HG12	9:9:77:VAL:O	1.99	0.62
12:AB:38:ILE:HD12	12:AB:160:ARG:NH1	2.13	0.62
12:AB:80:ARG:HB3	12:AB:82:GLY:O	1.99	0.62
16:AG:323:GLN:HE21	16:AG:323:GLN:CA	2.12	0.62
25:K:140:THR:O	25:K:144:LEU:HD23	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:P:24:GLU:HA	30:P:27:GLU:OE1	1.98	0.62
37:W:66:MET:SD	37:W:66:MET:O	2.57	0.62
39:Y:21:PRO:HB2	39:Y:22:PRO:HD3	1.80	0.62
39:Y:102:ARG:HA	39:Y:102:ARG:HE	1.64	0.62
41:a:537:G:OP1	59:s:2:LYS:HE2	1.98	0.62
45:e:13:GLU:O	45:e:17:GLU:OE1	2.16	0.62
54:n:38:MET:SD	54:n:150:ARG:HD2	2.39	0.62
3:2:36:LYS:HE3	3:2:79:ASP:OD2	1.98	0.62
6:5:10:DT:C5'	11:AA:62:TYR:OH	2.48	0.62
9:9:23:LEU:HD11	9:9:96:PHE:CE2	2.34	0.62
13:AD:211:ILE:HG12	13:AD:219:ARG:HH12	1.65	0.62
16:AG:266:LEU:H	16:AG:266:LEU:HD23	1.64	0.62
16:AG:389:MET:HE3	16:AG:407:ARG:HD2	1.81	0.62
18:D:337:G:H2'	18:D:338:A:C8	2.34	0.62
21:G:114:LEU:HD13	21:G:144:LEU:CB	2.28	0.62
39:Y:80:LYS:HG3	39:Y:86:LYS:O	2.00	0.62
16:AG:354:ALA:HA	16:AG:357:ASP:CA	2.28	0.62
18:D:1152:A:OP1	30:P:70:HIS:ND1	2.29	0.62
41:a:1138:G:N3	59:s:108:MET:HE2	2.14	0.62
5:4:77:VAL:HG11	5:4:79:ARG:HH21	1.64	0.62
16:AG:453:VAL:HG21	16:AG:462:GLN:CD	2.25	0.62
18:D:1145:A:O2'	18:D:1146:A:O5'	2.18	0.62
21:G:19:GLN:OE1	21:G:21:ARG:HG2	2.00	0.62
21:G:126:PHE:O	21:G:126:PHE:CD1	2.53	0.62
22:H:135:LEU:O	22:H:169:SER:HA	2.00	0.62
27:M:18:PHE:HD2	27:M:59:LEU:HD21	1.65	0.62
35:U:5:ARG:HG2	35:U:68:SER:HB2	1.79	0.62
51:k:21:TYR:CE2	51:k:38:LYS:HG2	2.34	0.62
62:v:88:ASN:OD1	62:v:89:VAL:N	2.32	0.62
1:0:82:HIS:CG	1:0:82:HIS:O	2.52	0.62
9:9:47:GLU:HA	9:9:94:ARG:HH22	1.65	0.62
12:AB:143:LEU:CD2	30:P:88:MET:HE1	2.27	0.62
16:AG:223:ILE:HA	16:AG:238:VAL:HG12	1.81	0.62
16:AG:288:MET:SD	16:AG:293:VAL:N	2.73	0.62
18:D:932:C:H5'	27:M:4:ARG:HH11	1.63	0.62
21:G:73:LYS:O	21:G:74:ARG:HG2	2.00	0.62
21:G:118:GLU:O	21:G:121:SER:OG	2.15	0.62
22:H:135:LEU:CB	22:H:167:VAL:CB	2.78	0.62
27:M:66:LEU:HD13	27:M:101:MET:HE1	1.81	0.62
32:R:75:GLN:OE1	32:R:75:GLN:N	2.32	0.62
41:a:2311:A:N3	54:n:85:ILE:HD11	2.13	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:37:ARG:NH2	59:s:44:TYR:OH	2.32	0.62
62:v:38:ARG:HA	62:v:96:ILE:O	2.00	0.62
6:5:9:DG:P	11:AA:473:ARG:CG	2.88	0.62
16:AG:428:ASN:O	16:AG:428:ASN:ND2	2.32	0.62
41:a:985:C:C2	41:a:986:C:C5	2.87	0.62
63:w:35:LYS:HZ2	63:w:112:TYR:HE2	1.47	0.62
63:w:37:THR:OG1	63:w:40:LYS:HD2	2.00	0.62
8:7:7:U:H6	8:7:7:U:H5''	1.65	0.62
9:9:7:ASP:O	9:9:11:ILE:HD12	1.98	0.62
11:AA:387:ASN:HB3	11:AA:394:ARG:HH11	1.63	0.62
11:AA:859:GLU:C	16:AG:38:LYS:HZ3	2.07	0.62
11:AA:860:ALA:N	16:AG:38:LYS:NZ	2.48	0.62
16:AG:27:LEU:HD22	16:AG:114:ILE:HG23	1.82	0.62
28:N:43:GLU:CD	28:N:101:ILE:HG21	2.25	0.62
41:a:1406:U:C2	41:a:1407:G:C8	2.88	0.62
41:a:1824:G:C3'	48:h:52:ARG:HH12	2.13	0.62
6:5:10:DT:N1	11:AA:62:TYR:CE2	2.65	0.62
9:9:113:PHE:O	9:9:123:ILE:HG13	1.99	0.62
12:AB:138:ARG:HD3	12:AB:155:LYS:HB2	1.81	0.62
16:AG:440:VAL:HG23	16:AG:481:LEU:CD2	2.29	0.62
21:G:110:SER:HA	21:G:113:ARG:HE	1.65	0.62
39:Y:9:LYS:HG2	39:Y:55:PRO:HB3	1.80	0.62
39:Y:100:ILE:CD1	39:Y:139:VAL:HB	2.29	0.62
41:a:614:A:O2'	41:a:615:U:OP2	2.12	0.62
62:v:1:MET:HE1	62:v:44:ARG:CA	2.30	0.62
8:7:11:U:O4'	25:K:20:ARG:NH2	2.32	0.62
11:AA:889:PRO:CG	16:AG:104:ARG:NH1	2.62	0.62
12:AB:102:LYS:HE2	14:AE:285:LEU:CB	2.29	0.62
14:AE:288:PRO:HD2	14:AE:291:ILE:HG22	1.74	0.62
16:AG:425:LEU:O	16:AG:429:LYS:NZ	2.33	0.62
18:D:1124:G:O2'	18:D:1145:A:N6	2.33	0.62
22:H:16:ILE:HG22	22:H:25:ARG:NH2	2.14	0.62
25:K:113:ALA:HA	25:K:116:GLU:OE1	2.00	0.62
41:a:2315:G:H4'	54:n:127:ASN:ND2	2.14	0.62
58:r:5:LEU:HD12	58:r:17:ASP:O	2.00	0.62
10:B:36:U:H2'	10:B:37:A:C8	2.35	0.61
18:D:1157:A:C2	18:D:1181:G:C4	2.88	0.61
33:S:19:LYS:HG3	33:S:20:TYR:CE1	2.35	0.61
47:g:1:MET:HE2	47:g:6:HIS:CD2	2.35	0.61
54:n:33:LYS:HA	54:n:96:MET:SD	2.39	0.61
59:s:11:VAL:HG11	59:s:13:ARG:NH1	2.15	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2:9:LYS:O	3:2:12:ARG:NH1	2.30	0.61
3:2:56:GLU:OE1	3:2:56:GLU:N	2.33	0.61
12:AB:140:MET:O	12:AB:150:ILE:CG2	2.48	0.61
14:AE:741:ALA:O	14:AE:762:ASN:ND2	2.32	0.61
16:AG:285:ILE:HD13	16:AG:285:ILE:N	2.15	0.61
18:D:562:U:C2'	32:R:14:ARG:HH11	2.13	0.61
23:I:14:ILE:HG22	23:I:15:VAL:HG13	1.82	0.61
27:M:111:ARG:O	27:M:119:ARG:NH1	2.34	0.61
28:N:88:ARG:HB2	28:N:91:GLU:OE1	2.00	0.61
29:O:123:ARG:HG2	29:O:123:ARG:O	1.98	0.61
34:T:67:LEU:HD12	34:T:78:TYR:CE1	2.35	0.61
60:t:20:MET:CB	60:t:44:LYS:HZ3	2.09	0.61
63:w:47:VAL:O	63:w:51:LEU:HD23	2.00	0.61
3:2:56:GLU:OE2	3:2:87:LEU:C	2.42	0.61
10:A:54:U:H3	10:A:58:A:N6	1.96	0.61
18:D:13:U:O2	18:D:914:A:C8	2.52	0.61
25:K:81:LEU:HG	25:K:96:MET:HE1	1.82	0.61
27:M:17:LYS:HG3	27:M:18:PHE:CD1	2.35	0.61
38:X:28:THR:CG2	38:X:31:LYS:HE3	2.30	0.61
39:Y:5:GLN:CG	39:Y:61:TYR:CE1	2.76	0.61
41:a:2307:G:O4'	41:a:2308:G:C5	2.52	0.61
46:f:41:THR:O	46:f:44:ILE:N	2.30	0.61
38:X:22:ILE:HB	38:X:25:VAL:HG12	1.80	0.61
41:a:548:G:H3'	41:a:549:G:O4'	2.01	0.61
48:h:141:VAL:HG23	48:h:162:VAL:CG1	2.30	0.61
10:A:5:G:H2'	10:A:6:G:H5'	1.82	0.61
10:A:65:C:H2'	10:A:66:C:H5'	1.83	0.61
11:AA:472:GLU:HG3	11:AA:473:ARG:HD3	1.82	0.61
11:AA:887:VAL:CB	16:AG:105:ILE:HG13	2.21	0.61
16:AG:253:GLY:HA3	16:AG:258:ARG:HD2	1.82	0.61
16:AG:421:GLN:O	16:AG:425:LEU:N	2.34	0.61
10:B:5:G:H2'	10:B:6:G:H5'	1.82	0.61
17:C:45:THR:HA	22:H:340:ARG:NH2	2.15	0.61
25:K:45:ARG:HB3	25:K:71:MET:HE2	1.81	0.61
33:S:87:ALA:HA	33:S:90:ARG:NH1	2.15	0.61
41:a:2002:G:OP1	63:w:17:ARG:NH2	2.26	0.61
51:k:34:LEU:HB3	51:k:51:GLU:HG3	1.82	0.61
9:9:25:ALA:HB2	9:9:96:PHE:CD1	2.36	0.61
11:AA:855:PRO:CD	16:AG:105:ILE:CG2	2.79	0.61
11:AA:860:ALA:N	16:AG:38:LYS:HD3	2.02	0.61
12:AB:31:PRO:O	12:AB:50:LEU:N	2.31	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:120:GLU:HA	12:AB:124:GLU:HA	1.80	0.61
16:AG:127:VAL:CG2	16:AG:192:GLY:CA	2.78	0.61
16:AG:168:LEU:HD21	16:AG:230:PRO:O	2.00	0.61
16:AG:197:VAL:HG11	16:AG:199:ARG:HH21	1.66	0.61
16:AG:380:THR:C	16:AG:384:LEU:HD11	2.24	0.61
16:AG:451:ARG:CZ	16:AG:469:ASP:HB2	2.31	0.61
21:G:30:PHE:CD1	21:G:201:PRO:CG	2.83	0.61
23:I:58:GLU:OE2	30:P:94:ALA:CB	2.48	0.61
25:K:69:ARG:HA	25:K:69:ARG:NE	2.15	0.61
31:Q:84:VAL:HG23	31:Q:107:ILE:CD1	2.30	0.61
38:X:81:MET:HE1	38:X:92:ARG:HD3	1.83	0.61
39:Y:19:PRO:O	39:Y:24:GLY:CA	2.48	0.61
41:a:1839:G:C8	41:a:1927:A:H1'	2.35	0.61
41:a:2363:G:OP2	55:o:40:ARG:HD2	2.01	0.61
47:g:5:ILE:HD12	54:n:64:LYS:CE	2.31	0.61
8:7:11:U:O2'	25:K:33:PHE:HE1	1.71	0.61
9:9:47:GLU:HG3	9:9:94:ARG:NH1	2.16	0.61
9:9:52:MET:C	9:9:52:MET:HE3	2.25	0.61
12:AB:146:ILE:HG22	30:P:88:MET:HA	1.82	0.61
13:AD:112:ALA:HB3	13:AD:126:PRO:HA	1.83	0.61
16:AG:233:ARG:CB	16:AG:327:LEU:HD23	2.28	0.61
16:AG:285:ILE:CG2	16:AG:293:VAL:HG11	2.28	0.61
18:D:563:A:C2	18:D:567:G:C5	2.89	0.61
27:M:50:LEU:HD21	27:M:121:ALA:CA	2.28	0.61
27:M:103:TRP:O	27:M:106:GLU:HG3	2.01	0.61
28:N:106:THR:HB	28:N:121:LEU:HD13	1.82	0.61
29:O:94:LEU:HB3	29:O:97:GLU:OE2	2.01	0.61
48:h:164:ILE:HG23	48:h:164:ILE:O	2.01	0.61
16:AG:303:HIS:HB3	16:AG:334:TRP:HZ3	1.66	0.61
16:AG:320:ARG:CG	16:AG:320:ARG:HH11	2.14	0.61
10:B:30:G:O2'	18:D:1339:A:C2	2.52	0.61
18:D:1309:G:P	38:X:98:ARG:HE	2.20	0.61
36:V:60:GLU:OE1	36:V:76:VAL:HG23	2.01	0.61
36:V:61:ILE:HG13	36:V:73:TRP:CZ3	2.36	0.61
9:9:23:LEU:HD21	9:9:96:PHE:CG	2.36	0.61
25:K:112:ARG:O	25:K:116:GLU:OE1	2.19	0.61
28:N:24:ALA:HB1	28:N:60:GLU:OE2	2.00	0.61
37:W:67:VAL:HG21	47:g:57:VAL:HG22	1.82	0.61
41:a:413:C:H2'	41:a:414:C:H6	1.66	0.61
54:n:122:PHE:O	54:n:123:ASP:OD1	2.19	0.61
63:w:42:LYS:HE2	63:w:97:ILE:HG21	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AB:115:LYS:NZ	12:AB:129:ILE:HG22	2.04	0.61
16:AG:279:ASN:HB3	16:AG:280:PRO:HD2	1.83	0.61
18:D:1125:U:C2	18:D:1127:G:C8	2.89	0.61
33:S:13:ARG:HB3	33:S:60:GLN:HG3	1.81	0.61
39:Y:36:GLU:HG3	39:Y:66:PHE:CE1	2.36	0.61
39:Y:72:THR:OG1	39:Y:73:PRO:HD2	2.01	0.61
40:Z:8:ILE:HG13	40:Z:9:GLU:N	2.15	0.61
47:g:65:ASN:OD1	47:g:66:ILE:N	2.30	0.61
58:r:114:GLU:OE2	58:r:134:VAL:N	2.33	0.61
62:v:111:GLU:HA	62:v:114:ARG:HE	1.66	0.61
8:7:11:U:O2'	25:K:33:PHE:CZ	2.52	0.60
9:9:61:ARG:NE	41:a:1046:A:O5'	2.34	0.60
12:AB:82:GLY:HA2	14:AE:142:GLU:OE1	2.01	0.60
14:AE:506:VAL:HG23	14:AE:628:GLY:HA3	1.83	0.60
16:AG:433:ASP:HB3	16:AG:456:LEU:HD12	1.83	0.60
16:AG:434:LEU:HG	16:AG:456:LEU:N	2.16	0.60
10:B:65:C:H2'	10:B:66:C:H5'	1.83	0.60
24:J:74:ASN:HA	24:J:77:LYS:HZ3	1.66	0.60
26:L:9:MET:HE2	26:L:51:ILE:HD11	1.83	0.60
29:O:55:VAL:O	29:O:57:MET:N	2.34	0.60
30:P:26:VAL:HG12	30:P:30:LYS:NZ	2.16	0.60
39:Y:79:LEU:CD2	39:Y:105:LEU:HD21	2.27	0.60
41:a:753:A:C2	41:a:754:U:C6	2.89	0.60
3:2:1:MET:N	41:a:142:A:O2'	2.34	0.60
3:2:56:GLU:OE2	3:2:87:LEU:O	2.18	0.60
10:A:36:U:H2'	10:A:37:A:C8	2.36	0.60
14:AE:79:LYS:HG2	16:AG:144:LYS:HD2	1.84	0.60
16:AG:240:THR:OG1	16:AG:247:PRO:HG2	2.01	0.60
16:AG:402:THR:O	16:AG:406:LEU:N	2.33	0.60
17:C:29:LEU:HD23	17:C:59:ILE:HD13	1.83	0.60
22:H:155:LYS:O	22:H:169:SER:N	2.34	0.60
36:V:10:GLY:HA3	36:V:25:ILE:HD13	1.83	0.60
58:r:135:HIS:ND1	58:r:137:GLU:CD	2.44	0.60
1:0:73:LYS:NZ	41:a:1225:G:OP1	2.34	0.60
3:2:56:GLU:OE1	3:2:86:THR:O	2.19	0.60
11:AA:55:SER:OG	11:AA:465:ARG:NH1	2.33	0.60
11:AA:528:ARG:NH2	11:AA:576:SER:O	2.29	0.60
14:AE:785:ASP:O	14:AE:789:LYS:HB2	2.01	0.60
14:AE:814:CYS:SG	14:AE:883:ARG:NH2	2.74	0.60
16:AG:424:SER:C	16:AG:429:LYS:HG3	2.26	0.60
17:C:60:LYS:NZ	18:D:734:G:C2'	2.64	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:687:A:C2	18:D:704:A:C5	2.88	0.60
21:G:38:VAL:CG2	22:H:75:VAL:HG21	2.31	0.60
21:G:114:LEU:HD13	21:G:144:LEU:HB2	1.84	0.60
29:O:56:ASP:O	29:O:57:MET:HE2	2.02	0.60
30:P:23:ALA:O	30:P:27:GLU:OE1	2.18	0.60
41:a:1406:U:O2'	41:a:1407:G:O4'	2.19	0.60
55:o:13:ARG:HD3	61:u:58:TYR:O	2.00	0.60
55:o:32:ILE:HG13	55:o:32:ILE:O	2.00	0.60
59:s:98:GLU:O	59:s:102:GLU:OE1	2.18	0.60
2:1:98:LYS:NZ	41:a:2011:U:O3'	2.34	0.60
19:E:28:MET:SD	19:E:57:ILE:HG22	2.42	0.60
23:I:142:MET:HE2	23:I:149:ILE:HG22	1.83	0.60
39:Y:56:VAL:HA	39:Y:71:LYS:NZ	2.08	0.60
41:a:271:G:C4	41:a:272:A:C8	2.89	0.60
41:a:639:U:H2'	41:a:640:C:C6	2.36	0.60
54:n:122:PHE:CZ	54:n:170:LEU:HD12	2.36	0.60
62:v:75:GLU:HG3	62:v:90:GLU:OE1	2.01	0.60
12:AB:115:LYS:CA	12:AB:129:ILE:CD1	2.79	0.60
14:AE:1161:GLY:HA3	14:AE:1179:PRO:HA	1.83	0.60
16:AG:422:GLU:CA	16:AG:426:GLY:CA	2.65	0.60
16:AG:422:GLU:CA	16:AG:426:GLY:H	2.08	0.60
10:B:69:C:H2'	10:B:70:G:O4'	2.02	0.60
18:D:718:A:O2'	20:F:34:ARG:NH2	2.34	0.60
24:J:91:LEU:HD21	24:J:195:ILE:HD12	1.84	0.60
41:a:323:C:H2'	52:l:163:ASN:OD1	2.02	0.60
41:a:1070:A:N7	41:a:1096:A:O2'	2.35	0.60
41:a:2189:U:P	41:a:2189:U:O4'	2.60	0.60
41:a:2191:A:H2'	41:a:2192:U:C6	2.36	0.60
41:a:2780:G:P	59:s:120:ARG:NH1	2.74	0.60
48:h:67:PHE:CE1	48:h:156:ARG:HD2	2.37	0.60
54:n:38:MET:HG3	54:n:57:LEU:HD21	1.83	0.60
59:s:13:ARG:NH2	59:s:49:ASP:O	2.34	0.60
59:s:102:GLU:HB3	59:s:119:PHE:HZ	1.67	0.60
60:t:8:LEU:N	60:t:18:ARG:HH12	1.98	0.60
61:u:51:GLU:OE1	61:u:51:GLU:N	2.34	0.60
1:0:68:ARG:HH12	41:a:1223:G:P	2.25	0.60
12:AB:129:ILE:O	12:AB:131:THR:N	2.35	0.60
13:AD:286:GLU:H	13:AD:315:GLY:CA	2.11	0.60
14:AE:92:VAL:HG22	14:AE:92:VAL:O	2.01	0.60
41:a:856:G:H2'	41:a:857:G:C8	2.37	0.60
41:a:1277:G:N3	63:w:23:ASN:ND2	2.47	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:57:TYR:OH	5:4:79:ARG:NH2	2.24	0.60
11:AA:889:PRO:CG	16:AG:104:ARG:HD3	2.23	0.60
12:AB:92:VAL:HA	12:AB:96:LEU:HB2	1.83	0.60
14:AE:86:GLU:OE1	14:AE:86:GLU:N	2.34	0.60
17:C:73:ARG:CZ	31:Q:113:VAL:HG12	2.32	0.60
28:N:43:GLU:HG2	28:N:101:ILE:HD13	1.84	0.60
29:O:57:MET:HA	29:O:60:LYS:HE2	1.83	0.60
39:Y:126:ARG:HH22	41:a:1083:U:P	2.25	0.60
41:a:468:G:N7	53:m:39:ARG:NH2	2.42	0.60
41:a:1172:C:H2'	41:a:1173:U:O4'	2.02	0.60
41:a:2298:A:C4	41:a:2321:U:C5	2.90	0.60
47:g:58:ASP:C	47:g:62:LYS:HZ3	2.07	0.60
59:s:13:ARG:HH22	59:s:50:THR:CA	2.14	0.60
63:w:60:VAL:HA	63:w:63:ARG:NH1	2.17	0.60
7:6:25:DT:H2'	11:AA:496:LYS:CE	2.28	0.60
9:9:60:LEU:HA	9:9:64:VAL:HB	1.84	0.60
11:AA:914:LYS:N	16:AG:105:ILE:CA	2.58	0.60
14:AE:1275:LEU:HB3	14:AE:1278:GLU:HB2	1.83	0.60
16:AG:10:GLU:HA	16:AG:12:VAL:HG22	1.84	0.60
16:AG:233:ARG:HG2	16:AG:270:ARG:HB2	1.84	0.60
17:C:14:THR:HG21	17:C:48:ARG:NE	2.17	0.60
25:K:159:LYS:NZ	28:N:44:GLY:HA3	2.17	0.60
50:j:25:THR:HG21	50:j:193:VAL:HG22	1.84	0.60
51:k:34:LEU:HB3	51:k:51:GLU:CG	2.31	0.60
55:o:30:ARG:HH12	61:u:62:PRO:CA	2.14	0.60
2:1:86:MET:CE	2:1:88:ARG:HD3	2.32	0.60
6:5:11:DA:H2'	12:AB:67:THR:HG21	1.82	0.60
9:9:94:ARG:CB	9:9:97:LYS:HE2	2.32	0.60
9:9:123:ILE:O	9:9:123:ILE:HG23	2.02	0.60
12:AB:81:PHE:HE1	14:AE:293:ARG:HB3	0.85	0.60
18:D:652:U:H4'	28:N:56:LYS:NZ	2.16	0.60
22:H:131:LEU:CB	22:H:167:VAL:HA	2.32	0.60
44:d:5:U:OP1	44:d:61:G:O2'	2.16	0.60
48:h:84:ASP:HB2	48:h:91:ILE:HD12	1.84	0.60
54:n:25:VAL:O	54:n:28:VAL:HG12	2.02	0.60
59:s:73:VAL:HG13	59:s:86:GLN:HG2	1.84	0.60
5:4:76:ASP:OD1	5:4:77:VAL:N	2.33	0.60
11:AA:856:ASN:CA	16:AG:35:THR:HB	2.31	0.60
12:AB:89:PRO:CG	14:AE:290:ILE:CG2	2.79	0.60
16:AG:139:THR:HG22	16:AG:180:ARG:HB3	1.83	0.60
18:D:1160:G:C2	18:D:1161:C:C6	2.89	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1432:G:H2'	41:a:1433:A:C8	2.37	0.60
9:9:33:VAL:HG22	9:9:36:ASP:CG	2.27	0.59
12:AB:118:ILE:HG13	12:AB:142:LEU:HD23	1.84	0.59
18:D:692:U:N3	18:D:695:A:OP2	2.34	0.59
18:D:1006:G:H2'	18:D:1007:U:C6	2.36	0.59
21:G:35:ARG:HB3	22:H:14:LYS:HZ3	1.67	0.59
33:S:63:ARG:NH2	33:S:68:GLY:O	2.32	0.59
38:X:93:ARG:HG2	38:X:95:LEU:HD23	1.84	0.59
39:Y:5:GLN:HE22	39:Y:60:VAL:C	2.10	0.59
41:a:1028:A:H2'	41:a:1029:A:C8	2.37	0.59
41:a:1257:C:H4'	52:l:78:TRP:CD2	2.37	0.59
41:a:1817:G:H3'	48:h:156:ARG:HH22	1.67	0.59
50:j:1:MET:HE2	50:j:100:LEU:HD11	1.83	0.59
61:u:76:GLU:HG3	61:u:111:ILE:HD13	1.84	0.59
9:9:7:ASP:OD1	9:9:8:LYS:N	2.35	0.59
9:9:23:LEU:HA	9:9:118:ILE:CG1	2.31	0.59
9:9:51:TYR:HB2	9:9:88:HIS:CB	2.32	0.59
9:9:98:GLU:HA	9:9:101:LYS:HG2	1.83	0.59
11:AA:1072:ASN:ND2	11:AA:1111:GLN:OE1	2.34	0.59
16:AG:353:HIS:O	16:AG:356:ILE:C	2.43	0.59
16:AG:434:LEU:CD1	16:AG:455:THR:C	2.74	0.59
18:D:736:C:O2'	26:L:88:MET:HE1	2.02	0.59
18:D:978:A:C5	18:D:1319:A:C2	2.90	0.59
22:H:280:LEU:N	22:H:330:VAL:O	2.31	0.59
39:Y:99:LYS:HD3	39:Y:138:VAL:O	2.02	0.59
51:k:19:HIS:ND1	51:k:41:PRO:CD	2.65	0.59
4:3:7:ARG:HH22	41:a:98:G:N2	2.00	0.59
5:4:63:ILE:HD11	5:4:91:PHE:CE2	2.37	0.59
9:9:14:GLU:OE1	9:9:62:ARG:O	2.19	0.59
9:9:103:ASN:HA	9:9:107:GLU:OE2	2.02	0.59
12:AB:102:LYS:CD	14:AE:285:LEU:HA	2.31	0.59
12:AB:154:VAL:CG1	12:AB:162:LEU:HD21	2.32	0.59
16:AG:127:VAL:HG21	16:AG:192:GLY:HA2	1.84	0.59
16:AG:354:ALA:HA	16:AG:357:ASP:N	2.18	0.59
16:AG:453:VAL:HG11	16:AG:459:LEU:HG	1.85	0.59
18:D:1290:G:H4'	27:M:35:LYS:NZ	2.17	0.59
24:J:62:ARG:HG2	24:J:72:PHE:CE1	2.38	0.59
27:M:50:LEU:HD11	27:M:124:LEU:HB2	1.84	0.59
35:U:5:ARG:HH22	35:U:24:SER:HA	1.66	0.59
41:a:993:G:OP2	66:z:51:ARG:NH2	2.36	0.59
41:a:1322:A:C5	41:a:1323:C:C5	2.90	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:h:30:PHE:O	48:h:34:LEU:HD23	2.02	0.59
12:AB:35:LEU:HB3	12:AB:37:LYS:HD3	1.82	0.59
12:AB:117:ILE:HD13	12:AB:161:LYS:NZ	2.17	0.59
12:AB:143:LEU:HD22	30:P:88:MET:CE	2.28	0.59
16:AG:187:ARG:CB	16:AG:188:PRO:HD3	2.20	0.59
18:D:1169:A:H2'	18:D:1170:A:C8	2.37	0.59
20:F:49:LYS:O	20:F:53:VAL:HG23	2.02	0.59
21:G:23:TRP:HZ3	21:G:25:PRO:HA	1.67	0.59
21:G:26:LYS:CD	21:G:193:PRO:HG2	2.32	0.59
23:I:54:ARG:NH2	23:I:56:VAL:HG21	2.18	0.59
30:P:25:ILE:O	30:P:28:THR:OG1	2.19	0.59
32:R:7:LEU:CD2	32:R:12:ARG:NH2	2.65	0.59
34:T:64:ARG:CZ	34:T:64:ARG:HB3	2.32	0.59
35:U:76:LYS:C	35:U:80:LYS:HZ3	2.10	0.59
40:Z:14:MET:HG3	40:Z:15:SER:N	2.17	0.59
41:a:947:A:H2'	41:a:948:C:C6	2.37	0.59
41:a:2460:U:C2	41:a:2461:A:C8	2.90	0.59
49:i:22:LEU:HG	49:i:23:THR:N	2.17	0.59
63:w:33:ILE:CG1	63:w:112:TYR:CD1	2.85	0.59
2:1:1:MET:SD	2:1:3:THR:OG1	2.59	0.59
8:7:10:U:H5''	8:7:10:U:O2	2.03	0.59
12:AB:154:VAL:CG1	12:AB:162:LEU:CD2	2.80	0.59
16:AG:240:THR:CB	16:AG:247:PRO:HG3	2.32	0.59
18:D:718:A:C4	31:Q:118:HIS:ND1	2.71	0.59
18:D:878:A:OP2	28:N:80:ARG:NH1	2.35	0.59
25:K:16:ILE:HD11	25:K:38:VAL:HG12	1.83	0.59
27:M:18:PHE:HB3	27:M:59:LEU:HD11	1.84	0.59
27:M:137:LYS:O	27:M:141:VAL:HG23	2.01	0.59
31:Q:46:THR:OG1	31:Q:49:GLY:N	2.34	0.59
33:S:42:TRP:CH2	47:g:66:ILE:HA	2.37	0.59
39:Y:9:LYS:HE2	39:Y:55:PRO:HG3	1.82	0.59
41:a:654:A:H61	55:o:19:LYS:HD2	1.68	0.59
48:h:62:TYR:CE2	48:h:64:ILE:HD13	2.38	0.59
59:s:55:ILE:HG13	59:s:123:LYS:NZ	2.17	0.59
10:A:33:U:O3'	27:M:84:THR:OG1	2.16	0.59
12:AB:142:LEU:HB2	12:AB:152:HIS:H	1.65	0.59
16:AG:31:LEU:HD23	16:AG:110:ALA:HB1	1.85	0.59
16:AG:63:LEU:HG	16:AG:73:LYS:HB3	1.83	0.59
16:AG:137:ILE:O	16:AG:137:ILE:HG22	2.03	0.59
18:D:61:G:N7	19:E:6:SER:OG	2.33	0.59
22:H:333:LEU:O	22:H:334:ASP:OD1	2.19	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:25:VAL:HG23	38:X:29:ARG:CB	2.32	0.59
38:X:55:THR:O	38:X:59:GLU:OE1	2.18	0.59
45:e:2:LYS:CE	45:e:52:ARG:HE	2.16	0.59
61:u:61:LEU:HB3	61:u:62:PRO:HD2	1.85	0.59
61:u:91:ASP:O	61:u:95:LEU:HD23	2.03	0.59
8:7:9:U:O4	18:D:1196:A:C8	2.55	0.59
9:9:17:GLU:HG2	9:9:88:HIS:NE2	2.18	0.59
10:A:69:C:H2'	10:A:70:G:O4'	2.02	0.59
12:AB:143:LEU:O	12:AB:145:LEU:N	2.36	0.59
16:AG:202:PRO:O	16:AG:205:LEU:HD12	2.02	0.59
18:D:13:U:N3	18:D:915:A:N6	2.51	0.59
18:D:921:U:H4'	25:K:23:LYS:NZ	2.18	0.59
21:G:35:ARG:HD2	22:H:14:LYS:HE3	1.84	0.59
21:G:128:LYS:HD2	21:G:128:LYS:O	2.03	0.59
25:K:10:GLU:OE1	25:K:10:GLU:N	2.35	0.59
28:N:8:ALA:O	28:N:12:THR:HG23	2.03	0.59
33:S:20:TYR:HD2	33:S:55:SER:OG	1.85	0.59
37:W:45:ILE:HA	37:W:62:VAL:CG1	2.31	0.59
41:a:2682:A:H8	50:j:11:MET:HE1	1.65	0.59
48:h:181:MET:HE2	48:h:181:MET:N	2.18	0.59
61:u:76:GLU:HG3	61:u:111:ILE:CD1	2.31	0.59
9:9:73:LYS:CG	9:9:117:LEU:HD12	2.32	0.59
9:9:97:LYS:NZ	9:9:130:PRO:HA	2.17	0.59
10:A:70:G:H2'	10:A:71:C:H5''	1.85	0.59
12:AB:89:PRO:HG2	14:AE:290:ILE:CG2	2.33	0.59
12:AB:115:LYS:HD2	12:AB:129:ILE:HG21	1.82	0.59
16:AG:318:ILE:HG12	16:AG:338:VAL:HG11	1.85	0.59
18:D:829:G:O3'	21:G:23:TRP:CZ2	2.56	0.59
18:D:1055:A:C2'	23:I:156:ARG:HH21	2.15	0.59
18:D:1107:C:P	23:I:172:ARG:HG3	2.43	0.59
41:a:15:G:O2'	49:i:15:MET:CE	2.51	0.59
46:f:40:ASP:C	46:f:45:ARG:NH2	2.60	0.59
66:z:74:ILE:HD11	66:z:79:PHE:CA	2.30	0.59
1:0:51:VAL:O	1:0:51:VAL:HG13	2.02	0.59
5:4:31:TYR:OH	44:d:103:U:O2'	2.20	0.59
6:5:11:DA:H2'	12:AB:67:THR:CG2	2.32	0.59
7:6:25:DT:OP1	11:AA:496:LYS:C	2.43	0.59
11:AA:861:ALA:CA	16:AG:38:LYS:HZ1	2.14	0.59
12:AB:77:HIS:NE2	14:AE:297:ARG:NH2	2.51	0.59
14:AE:81:ARG:HG3	14:AE:81:ARG:NH2	2.17	0.59
16:AG:434:LEU:N	16:AG:456:LEU:HG	2.17	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:12:ARG:CG	22:H:264:GLU:CB	2.81	0.59
18:D:972:C:O2'	30:P:57:VAL:HG13	2.03	0.59
39:Y:23:VAL:HG23	39:Y:24:GLY:N	2.17	0.59
41:a:1798:U:OP2	48:h:271:ARG:NH2	2.36	0.59
46:f:10:THR:HG22	46:f:54:MET:C	2.27	0.59
51:k:7:GLU:CD	51:k:7:GLU:H	2.11	0.59
9:9:31:ARG:CD	41:a:1053:C:O2'	2.47	0.59
9:9:78:GLY:N	9:9:79:PRO:CD	2.66	0.59
11:AA:856:ASN:CG	16:AG:35:THR:CG2	2.75	0.59
11:AA:889:PRO:HG3	16:AG:104:ARG:CD	2.23	0.59
14:AE:64:PRO:HG3	14:AE:90:VAL:HG12	1.83	0.59
16:AG:363:LEU:HD21	16:AG:410:ALA:N	1.79	0.59
17:C:51:TYR:HD1	17:C:54:GLN:NE2	2.00	0.59
18:D:496:A:N3	18:D:496:A:H2'	2.17	0.59
18:D:736:C:H5''	26:L:90:MET:HE2	1.85	0.59
18:D:890:G:O2'	18:D:906:A:N6	2.35	0.59
40:Z:8:ILE:HG13	40:Z:9:GLU:H	1.68	0.59
41:a:607:U:C5	41:a:620:G:C4	2.91	0.59
41:a:1277:G:H5'	63:w:20:MET:HE2	1.84	0.59
41:a:2225:A:H4'	41:a:2226:C:O5'	2.03	0.59
41:a:2259:U:N3	41:a:2260:C:C5	2.71	0.59
53:m:1:MET:HE3	53:m:3:ARG:HH11	1.67	0.59
60:t:88:ASN:ND2	60:t:90:ASN:OD1	2.36	0.59
66:z:74:ILE:CD1	66:z:74:ILE:CB	2.76	0.59
11:AA:707:ALA:O	11:AA:711:ASP:HB2	2.03	0.58
11:AA:1142:ARG:NH1	11:AA:1161:LEU:O	2.36	0.58
12:AB:90:SER:HB3	12:AB:93:ILE:HB	1.84	0.58
12:AB:145:LEU:HD22	12:AB:148:LYS:HD3	1.85	0.58
12:AB:148:LYS:HZ1	12:AB:150:ILE:HD13	1.67	0.58
16:AG:212:GLU:HB3	16:AG:258:ARG:HB3	1.84	0.58
26:L:42:TRP:CZ2	26:L:102:MET:HG3	2.36	0.58
28:N:104:VAL:HG23	28:N:125:ILE:HD13	1.83	0.58
52:l:170:ARG:CZ	52:l:175:ILE:HA	2.33	0.58
56:p:85:LYS:NZ	56:p:164:TYR:HD1	2.01	0.58
62:v:97:GLN:H	62:v:100:LYS:HZ1	1.51	0.58
11:AA:861:ALA:H	16:AG:38:LYS:HZ1	0.59	0.58
12:AB:120:GLU:HB2	12:AB:162:LEU:CB	2.32	0.58
16:AG:130:PHE:C	16:AG:186:VAL:HG11	2.28	0.58
17:C:73:ARG:NH1	31:Q:113:VAL:HA	2.18	0.58
18:D:695:A:H8	18:D:695:A:H5''	1.68	0.58
25:K:156:LYS:HA	28:N:66:PHE:CD2	2.39	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:N:7:ILE:HB	28:N:77:ARG:NH1	2.18	0.58
34:T:11:ILE:HG21	34:T:31:LEU:HD13	1.85	0.58
38:X:22:ILE:HG23	38:X:66:GLU:OE2	2.03	0.58
41:a:567:U:OP2	61:u:29:LYS:NZ	2.35	0.58
41:a:1668:A:O2'	41:a:1674:G:N7	2.32	0.58
44:d:42:C:C5'	54:n:64:LYS:HZ1	2.16	0.58
52:l:119:ILE:O	52:l:187:VAL:HA	2.03	0.58
53:m:24:THR:HG23	53:m:27:GLY:H	1.68	0.58
59:s:74:TYR:CD2	59:s:74:TYR:N	2.69	0.58
59:s:132:HIS:O	59:s:135:GLN:HB2	2.03	0.58
60:t:8:LEU:CA	60:t:18:ARG:HH12	2.15	0.58
3:2:56:GLU:OE1	3:2:87:LEU:C	2.46	0.58
12:AB:158:GLU:HB3	12:AB:162:LEU:HD22	1.85	0.58
18:D:522:C:OP2	32:R:66:TYR:OH	2.17	0.58
19:E:61:GLN:NE2	19:E:66:LEU:HD22	2.18	0.58
20:F:16:LEU:C	20:F:20:LYS:HZ3	2.10	0.58
21:G:66:LYS:HB3	21:G:154:MET:CE	2.34	0.58
21:G:106:THR:HA	21:G:109:GLN:NE2	2.18	0.58
22:H:81:GLU:O	22:H:82:THR:HG23	2.03	0.58
28:N:47:GLU:OE2	28:N:64:LYS:HA	2.02	0.58
41:a:70:G:H4'	41:a:71:A:OP1	2.02	0.58
58:r:6:LEU:HD11	58:r:37:VAL:CG2	2.33	0.58
58:r:30:LEU:HD23	58:r:35:LYS:HE2	1.85	0.58
1:0:15:SER:HB2	1:0:18:GLN:NE2	2.18	0.58
14:AE:1221:LEU:HD22	14:AE:1306:LEU:HB2	1.84	0.58
18:D:439:U:O2	18:D:440:C:C5	2.56	0.58
18:D:1151:A:O2'	18:D:1152:A:O5'	2.13	0.58
21:G:96:TRP:CH2	21:G:100:MET:HE3	2.38	0.58
24:J:74:ASN:HA	24:J:77:LYS:NZ	2.18	0.58
25:K:12:GLN:HB3	25:K:14:LYS:NZ	2.18	0.58
30:P:22:THR:O	30:P:26:VAL:HG23	2.03	0.58
41:a:575:A:C2	41:a:576:U:C5	2.90	0.58
41:a:685:A:O2'	41:a:773:U:O4	2.17	0.58
41:a:2189:U:O4'	41:a:2189:U:OP1	2.21	0.58
41:a:2815:C:C2	41:a:2816:G:C8	2.91	0.58
54:n:4:LEU:HD11	54:n:101:GLU:N	2.18	0.58
56:p:149:ARG:NH2	56:p:162:VAL:N	2.52	0.58
1:0:61:ALA:HB1	1:0:96:VAL:HG22	1.84	0.58
2:1:11:ARG:NH1	2:1:98:LYS:HE3	2.19	0.58
6:5:9:DG:O5'	11:AA:473:ARG:CB	2.49	0.58
12:AB:19:GLU:O	12:AB:23:ARG:HB2	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:437:LEU:HD22	16:AG:488:ILE:CD1	2.33	0.58
10:B:70:G:H2'	10:B:71:C:H5''	1.85	0.58
17:C:73:ARG:HH12	31:Q:113:VAL:CA	2.16	0.58
23:I:24:ALA:C	23:I:29:PHE:HE2	2.11	0.58
25:K:93:ARG:HB3	25:K:93:ARG:NH2	2.18	0.58
25:K:146:ASN:OD1	25:K:147:MET:N	2.37	0.58
27:M:4:ARG:HG3	27:M:5:ARG:CG	2.22	0.58
34:T:60:VAL:O	34:T:64:ARG:HG2	2.03	0.58
39:Y:109:ALA:HB2	39:Y:128:ILE:HG13	1.86	0.58
41:a:580:U:H2'	41:a:581:C:H6	1.69	0.58
41:a:832:U:H2'	41:a:833:A:H8	1.69	0.58
60:t:5:GLN:HA	60:t:20:MET:CE	2.32	0.58
62:v:1:MET:HE3	62:v:1:MET:CA	2.34	0.58
65:y:7:GLN:CD	65:y:7:GLN:C	2.71	0.58
5:4:60:VAL:HG12	5:4:71:LYS:HE2	1.86	0.58
6:5:10:DT:O3'	12:AB:68:THR:CA	2.51	0.58
6:5:11:DA:C2'	12:AB:67:THR:CG2	2.77	0.58
11:AA:838:CYS:HB2	11:AA:918:LEU:HD22	1.85	0.58
14:AE:1143:ASP:OD1	14:AE:1148:ARG:NH1	2.37	0.58
18:D:915:A:N6	18:D:916:U:C4	2.72	0.58
18:D:1308:U:P	38:X:98:ARG:HG3	2.43	0.58
19:E:32:ILE:HG12	19:E:75:HIS:CE1	2.37	0.58
22:H:291:GLY:HA2	22:H:305:HIS:HA	1.85	0.58
35:U:19:VAL:HG12	35:U:37:GLY:C	2.28	0.58
36:V:4:LYS:HD2	36:V:5:ILE:HG23	1.85	0.58
36:V:11:ARG:HE	36:V:58:VAL:CG2	2.16	0.58
58:r:37:VAL:HG12	58:r:43:ASN:HD21	1.68	0.58
59:s:102:GLU:CD	59:s:124:VAL:HG11	2.28	0.58
62:v:49:ALA:O	62:v:52:ALA:N	2.37	0.58
6:5:10:DT:O3'	12:AB:68:THR:HA	2.03	0.58
16:AG:453:VAL:CG2	16:AG:462:GLN:CD	2.76	0.58
10:B:15:G:N1	10:B:20:U:O2	2.37	0.58
18:D:974:A:OP1	33:S:69:ARG:NH1	2.37	0.58
31:Q:20:VAL:CG1	31:Q:85:MET:HE1	2.33	0.58
41:a:1021:A:N1	41:a:1141:U:O4	2.37	0.58
41:a:2349:G:C6	41:a:2369:A:C6	2.92	0.58
46:f:40:ASP:OD1	46:f:45:ARG:NH2	2.37	0.58
56:p:85:LYS:CE	56:p:164:TYR:CD1	2.86	0.58
58:r:68:ARG:O	58:r:72:ILE:HD12	2.03	0.58
59:s:31:GLU:OE2	59:s:35:ARG:NH1	2.37	0.58
9:9:24:SER:HB2	9:9:86:MET:HA	1.86	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:69:GLN:HE21	11:AA:101:ARG:HD2	1.67	0.58
12:AB:6:LEU:H	12:AB:88:VAL:HG21	1.69	0.58
13:AC:82:LEU:HD11	13:AC:171:LEU:HD23	1.86	0.58
16:AG:413:ALA:HA	16:AG:416:THR:OG1	2.03	0.58
10:B:60:U:H4'	10:B:61:C:OP2	2.04	0.58
27:M:26:PHE:CD1	27:M:101:MET:SD	2.97	0.58
28:N:52:GLU:OE1	28:N:53:GLY:N	2.37	0.58
39:Y:70:THR:O	39:Y:71:LYS:HG3	2.03	0.58
39:Y:102:ARG:CG	39:Y:141:ASP:HA	2.33	0.58
41:a:2395:C:H2'	41:a:2396:G:O4'	2.03	0.58
42:b:51:VAL:C	42:b:62:LYS:HZ3	2.11	0.58
44:d:106:G:H2'	44:d:107:G:O4'	2.04	0.58
51:k:35:GLU:OE2	51:k:48:ILE:HG23	2.03	0.58
52:l:62:GLN:O	52:l:69:ARG:HD3	2.03	0.58
7:6:20:DA:C6	8:7:51:G:C2	2.91	0.58
11:AA:13:LYS:HB2	11:AA:1180:MET:HE2	1.86	0.58
16:AG:25:GLU:HG3	16:AG:49:ILE:HD11	1.86	0.58
16:AG:61:ARG:HD2	16:AG:73:LYS:HD3	1.86	0.58
16:AG:354:ALA:CA	16:AG:357:ASP:HB3	2.32	0.58
16:AG:361:LYS:CE	16:AG:417:ILE:CG1	2.75	0.58
17:C:13:PHE:HB3	17:C:51:TYR:CD1	2.39	0.58
21:G:35:ARG:HD2	22:H:14:LYS:CE	2.34	0.58
22:H:273:ARG:CZ	22:H:297:GLU:HG3	2.33	0.58
25:K:148:ASN:OD1	28:N:96:MET:HE1	2.04	0.58
34:T:32:LEU:HD11	34:T:62:GLN:OE1	2.04	0.58
36:V:42:THR:HG22	36:V:44:LEU:CD1	2.34	0.58
41:a:563:A:OP1	66:z:41:LYS:NZ	2.35	0.58
41:a:1047:G:HO2'	41:a:1110:G:H1	1.52	0.58
41:a:2189:U:O2'	41:a:2190:G:O4'	2.22	0.58
51:k:34:LEU:H	51:k:51:GLU:CD	2.12	0.58
60:t:121:GLU:OE2	65:y:41:GLN:NE2	2.36	0.58
62:v:111:GLU:OE1	62:v:111:GLU:N	2.32	0.58
5:4:2:PHE:HE2	5:4:55:GLU:HG2	1.68	0.58
6:5:25:DA:C2	7:6:5:DG:C2	2.92	0.58
11:AA:855:PRO:CA	16:AG:35:THR:HB	2.28	0.58
14:AE:79:LYS:HG3	16:AG:144:LYS:HZ3	1.67	0.58
16:AG:63:LEU:HD23	16:AG:63:LEU:H	1.69	0.58
19:E:61:GLN:HE21	19:E:66:LEU:HD22	1.69	0.58
21:G:19:GLN:CG	22:H:76:GLU:HB3	2.34	0.58
28:N:66:PHE:CG	28:N:67:GLN:OE1	2.57	0.58
36:V:11:ARG:HE	36:V:58:VAL:HG22	1.69	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:25:VAL:HG23	38:X:29:ARG:HB2	1.85	0.58
39:Y:116:MET:HG2	39:Y:124:MET:CB	2.34	0.58
41:a:2243:U:H2'	41:a:2244:U:C6	2.38	0.58
45:e:2:LYS:HD2	45:e:6:LEU:HD11	1.85	0.58
52:l:136:GLN:NE2	52:l:140:ASP:OD2	2.37	0.58
54:n:46:ASP:OD1	54:n:49:LEU:HD13	2.03	0.58
61:u:55:MET:CE	61:u:60:ARG:HG2	2.33	0.58
66:z:82:GLY:CA	66:z:117:LEU:CD1	2.81	0.58
1:0:16:GLU:C	1:0:18:GLN:OE1	2.46	0.57
9:9:23:LEU:N	9:9:118:ILE:HD11	2.19	0.57
12:AB:99:TYR:O	12:AB:100:LYS:HG2	2.03	0.57
16:AG:434:LEU:HD11	16:AG:455:THR:N	2.19	0.57
29:O:49:ARG:HB3	29:O:53:GLU:OE2	2.02	0.57
29:O:94:LEU:O	29:O:98:LEU:HD13	2.04	0.57
34:T:28:GLN:HB2	34:T:66:LEU:HD21	1.86	0.57
41:a:811:U:H2'	61:u:21:ARG:HA	1.85	0.57
41:a:1141:U:H5	59:s:68:LYS:NZ	2.02	0.57
59:s:53:TYR:CD1	59:s:121:LYS:HD2	2.39	0.57
8:7:7:U:H5''	8:7:7:U:C6	2.39	0.57
11:AA:860:ALA:N	16:AG:38:LYS:CD	2.62	0.57
12:AB:102:LYS:CE	14:AE:285:LEU:HA	2.34	0.57
16:AG:233:ARG:HH12	16:AG:324:ASN:N	2.02	0.57
16:AG:392:LEU:CD2	16:AG:395:ILE:HD11	2.34	0.57
16:AG:433:ASP:C	16:AG:456:LEU:HD11	2.23	0.57
18:D:588:G:H4'	28:N:3:MET:CE	2.33	0.57
23:I:82:GLU:O	23:I:86:LYS:HD3	2.04	0.57
26:L:95:ALA:O	26:L:96:VAL:C	2.47	0.57
41:a:593:U:H2'	41:a:594:U:C6	2.39	0.57
2:1:73:LYS:C	2:1:74:ILE:HD12	2.29	0.57
9:9:50:VAL:CG2	39:Y:119:ALA:HB3	2.19	0.57
10:A:15:G:N1	10:A:20:U:O2	2.37	0.57
16:AG:387:VAL:HB	16:AG:388:PRO:HD2	1.85	0.57
18:D:911:U:H2'	18:D:912:C:C6	2.40	0.57
23:I:25:ASN:N	23:I:29:PHE:HE2	2.01	0.57
25:K:136:VAL:O	25:K:140:THR:HG23	2.04	0.57
27:M:66:LEU:HD11	27:M:100:ALA:HB3	1.85	0.57
27:M:68:ASN:ND2	27:M:130:ASN:OD1	2.38	0.57
41:a:839:U:C2	41:a:840:C:C5	2.93	0.57
41:a:1682:G:H2'	41:a:1683:U:C6	2.39	0.57
41:a:2682:A:H8	50:j:11:MET:CE	2.15	0.57
63:w:9:GLN:O	63:w:17:ARG:HD3	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:z:53:ARG:CG	66:z:57:PHE:HE2	2.18	0.57
4:3:6:ARG:NH1	41:a:84:A:O2'	2.36	0.57
11:AA:1254:VAL:O	14:AE:99:ARG:NH2	2.37	0.57
12:AB:147:ASN:C	12:AB:147:ASN:HD22	2.12	0.57
16:AG:233:ARG:O	16:AG:327:LEU:HD21	2.04	0.57
16:AG:379:SER:HG	16:AG:384:LEU:HD21	1.67	0.57
18:D:99:C:HO2'	18:D:100:G:H8	1.53	0.57
19:E:9:LYS:C	19:E:13:GLN:NE2	2.62	0.57
22:H:51:GLU:OE1	22:H:51:GLU:N	2.38	0.57
23:I:49:LYS:HZ3	23:I:75:ILE:HD11	1.69	0.57
27:M:24:ALA:O	27:M:27:VAL:HG22	2.04	0.57
27:M:28:ASN:HA	27:M:31:MET:HE3	1.85	0.57
30:P:21:ALA:O	30:P:24:GLU:HG3	2.04	0.57
30:P:63:ASP:OD2	30:P:64:GLN:N	2.38	0.57
31:Q:51:GLY:O	31:Q:56:ARG:CZ	2.51	0.57
34:T:32:LEU:CD1	34:T:62:GLN:OE1	2.53	0.57
39:Y:32:VAL:HG22	39:Y:60:VAL:CG2	2.34	0.57
39:Y:116:MET:HB3	39:Y:124:MET:HE2	1.86	0.57
41:a:1278:C:O2'	63:w:27:SER:OG	2.16	0.57
41:a:1869:G:N2	41:a:1871:A:O2'	2.37	0.57
41:a:2335:A:OP2	64:x:13:ARG:CD	2.51	0.57
41:a:2547:A:H2'	41:a:2548:U:C6	2.38	0.57
47:g:18:CYS:HB3	47:g:40:CYS:HB2	1.85	0.57
59:s:39:LYS:HA	59:s:44:TYR:CD1	2.39	0.57
6:5:9:DG:P	11:AA:470:ARG:HA	2.43	0.57
16:AG:210:ARG:HE	16:AG:216:ILE:HG22	1.69	0.57
16:AG:352:ALA:O	16:AG:356:ILE:CG2	2.51	0.57
17:C:55:LEU:O	17:C:59:ILE:HG12	2.04	0.57
21:G:107:VAL:O	21:G:111:ILE:HG12	2.04	0.57
34:T:10:LYS:O	34:T:14:GLU:OE1	2.23	0.57
41:a:5:A:H2'	41:a:6:A:H8	1.70	0.57
41:a:1824:G:H5''	48:h:52:ARG:HH22	1.69	0.57
41:a:2683:C:H4'	50:j:13:ARG:CZ	2.34	0.57
59:s:120:ARG:CZ	59:s:120:ARG:HB2	2.34	0.57
60:t:20:MET:HB3	60:t:44:LYS:CE	2.35	0.57
66:z:74:ILE:HD13	66:z:79:PHE:CD1	2.34	0.57
5:4:46:LYS:O	5:4:50:MET:HG2	2.04	0.57
8:7:9:U:O4	18:D:1196:A:O4'	2.21	0.57
9:9:94:ARG:CB	9:9:97:LYS:HZ3	2.17	0.57
12:AB:135:GLY:O	12:AB:137:ALA:N	2.37	0.57
16:AG:130:PHE:HB3	16:AG:186:VAL:HG22	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:287:ALA:CB	16:AG:331:LEU:HB3	2.33	0.57
16:AG:302:LYS:NZ	16:AG:302:LYS:HB2	2.20	0.57
17:C:16:GLU:OE1	17:C:18:VAL:N	2.36	0.57
41:a:5:A:H2'	41:a:6:A:C8	2.39	0.57
41:a:782:A:C2	48:h:225:MET:HE2	2.39	0.57
41:a:1065:U:O2'	41:a:1066:U:O5'	2.16	0.57
47:g:47:LYS:HG2	54:n:115:ARG:NH2	2.20	0.57
48:h:52:ARG:HE	48:h:53:HIS:CE1	2.23	0.57
64:x:21:LEU:HD22	64:x:23:ALA:HB2	1.87	0.57
65:y:51:ARG:HD3	65:y:58:ALA:HB3	1.86	0.57
9:9:38:MET:SD	39:Y:117:THR:HG22	2.45	0.57
12:AB:154:VAL:HG12	12:AB:158:GLU:HB3	1.86	0.57
18:D:562:U:C3'	32:R:14:ARG:HH11	2.17	0.57
18:D:662:U:O2'	18:D:836:G:OP1	2.21	0.57
22:H:116:VAL:CA	22:H:279:LYS:HD2	2.35	0.57
33:S:90:ARG:C	33:S:92:GLU:OE1	2.47	0.57
41:a:64:A:H2'	41:a:65:U:C6	2.40	0.57
42:b:55:ARG:NE	42:b:55:ARG:HA	2.20	0.57
43:c:66:THR:O	43:c:70:GLU:OE1	2.23	0.57
45:e:18:LEU:HD11	45:e:54:LYS:HE3	1.86	0.57
58:r:97:ARG:NH1	58:r:101:ASP:OD2	2.37	0.57
61:u:90:VAL:CG1	61:u:95:LEU:HD21	2.33	0.57
5:4:57:TYR:CD2	62:v:136:MET:HE2	2.40	0.57
11:AA:847:PRO:N	16:AG:104:ARG:CZ	2.67	0.57
11:AA:1122:LYS:NZ	11:AA:1126:ASP:OD1	2.38	0.57
16:AG:162:ILE:HD13	16:AG:167:MET:HE2	1.86	0.57
16:AG:285:ILE:CG1	16:AG:293:VAL:CG1	2.54	0.57
21:G:19:GLN:HG3	22:H:76:GLU:CB	2.35	0.57
22:H:273:ARG:HD3	22:H:280:LEU:HD11	1.85	0.57
23:I:6:HIS:HB3	33:S:89:MET:SD	2.45	0.57
31:Q:52:PHE:CD2	31:Q:56:ARG:HG2	2.39	0.57
32:R:110:ARG:NE	32:R:117:TYR:HD2	2.02	0.57
35:U:12:LYS:CD	35:U:13:LYS:HG2	2.33	0.57
37:W:40:ILE:HG21	37:W:66:MET:SD	2.45	0.57
38:X:93:ARG:CG	38:X:95:LEU:HD23	2.35	0.57
41:a:3:U:H2'	41:a:4:U:H6	1.69	0.57
41:a:2071:A:H2'	41:a:2072:C:C6	2.40	0.57
62:v:77:PRO:HG2	62:v:80:VAL:CG2	2.31	0.57
66:z:74:ILE:HG12	66:z:79:PHE:HB2	1.86	0.57
3:2:56:GLU:CD	3:2:87:LEU:C	2.73	0.57
11:AA:839:VAL:HG12	11:AA:1049:ILE:HG12	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:1223:ARG:NH2	14:AE:721:SER:OG	2.38	0.57
12:AB:119:THR:HB	12:AB:123:PHE:CD2	2.40	0.57
16:AG:362:TYR:CA	16:AG:413:ALA:HB1	2.30	0.57
16:AG:367:GLU:O	16:AG:371:THR:HG23	2.04	0.57
18:D:512:U:OP1	24:J:44:ARG:NH1	2.38	0.57
25:K:148:ASN:ND2	25:K:153:VAL:HG12	2.20	0.57
27:M:66:LEU:HD23	27:M:70:ARG:HH21	1.69	0.57
31:Q:87:LYS:HG3	31:Q:113:VAL:HG23	1.86	0.57
38:X:16:VAL:CG2	38:X:17:ILE:HD12	2.35	0.57
41:a:78:U:H5'	45:e:4:LYS:NZ	2.20	0.57
41:a:1197:G:H2'	41:a:1198:U:H6	1.68	0.57
41:a:1360:G:N7	41:a:1361:G:C8	2.73	0.57
41:a:1818:U:OP2	48:h:156:ARG:CZ	2.53	0.57
9:9:54:VAL:HG13	9:9:54:VAL:O	2.03	0.57
9:9:124:ASP:OD2	9:9:126:LEU:HD21	2.05	0.57
10:A:5:G:H2'	10:A:6:G:C5'	2.35	0.57
10:A:60:U:H4'	10:A:61:C:OP2	2.04	0.57
10:A:75:C:O2'	10:A:76:A:C8	2.58	0.57
11:AA:846:GLY:O	16:AG:104:ARG:NH1	2.13	0.57
16:AG:33:THR:HG23	16:AG:43:ILE:HD11	1.85	0.57
16:AG:440:VAL:HG22	16:AG:481:LEU:CD2	2.34	0.57
18:D:739:C:HO2'	34:T:42:HIS:CE1	2.19	0.57
18:D:1356:G:H2'	18:D:1357:A:C8	2.40	0.57
21:G:52:GLU:O	21:G:56:GLU:OE1	2.23	0.57
27:M:134:ALA:O	27:M:137:LYS:HG2	2.05	0.57
35:U:39:PHE:HD1	35:U:50:THR:HG22	1.70	0.57
41:a:16:C:C4'	49:i:15:MET:HE3	2.35	0.57
41:a:628:G:C6	41:a:636:G:C2	2.92	0.57
41:a:2816:G:O4'	49:i:41:HIS:CE1	2.57	0.57
5:4:48:MET:HE3	5:4:84:PRO:O	2.04	0.56
9:9:27:VAL:HG13	9:9:27:VAL:O	2.05	0.56
12:AB:26:VAL:HA	12:AB:58:GLU:OE1	2.05	0.56
12:AB:102:LYS:HE2	14:AE:285:LEU:CA	2.34	0.56
14:AE:144:TYR:OH	14:AE:293:ARG:NH2	2.37	0.56
16:AG:9:VAL:HA	16:AG:24:PHE:CZ	2.40	0.56
16:AG:266:LEU:HD23	16:AG:266:LEU:N	2.19	0.56
16:AG:285:ILE:HG23	16:AG:293:VAL:HG11	1.82	0.56
18:D:657:U:C4'	34:T:28:GLN:HE22	2.15	0.56
18:D:1515:G:H2'	18:D:1516:G:H8	1.70	0.56
19:E:61:GLN:HE21	19:E:66:LEU:CD2	2.18	0.56
22:H:297:GLU:HG2	22:H:300:VAL:HG21	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:M:95:ARG:O	27:M:99:LEU:HG	2.05	0.56
29:O:5:GLN:NE2	29:O:20:PHE:CB	2.67	0.56
31:Q:15:GLN:OE1	31:Q:78:GLY:HA3	2.05	0.56
39:Y:58:ILE:HG23	39:Y:66:PHE:CE2	2.40	0.56
41:a:1442:U:H2'	41:a:1443:U:C6	2.40	0.56
41:a:1693:U:O2'	48:h:14:ARG:NH2	2.38	0.56
41:a:2291:U:H2'	41:a:2292:U:C6	2.40	0.56
42:b:45:PHE:CD2	42:b:80:ILE:HD11	2.40	0.56
52:l:148:ILE:HD13	52:l:187:VAL:HB	1.85	0.56
53:m:12:ARG:NE	53:m:44:VAL:HG21	2.19	0.56
58:r:54:LEU:HA	58:r:57:LYS:NZ	2.20	0.56
62:v:37:GLY:C	62:v:126:ILE:HD11	2.30	0.56
4:3:6:ARG:CZ	41:a:84:A:H2'	2.35	0.56
9:9:4:ASN:HB3	9:9:8:LYS:NZ	2.19	0.56
11:AA:856:ASN:OD1	16:AG:34:ALA:O	2.23	0.56
16:AG:49:ILE:HG22	16:AG:49:ILE:O	2.04	0.56
16:AG:221:ILE:HD13	16:AG:247:PRO:HA	1.85	0.56
18:D:280:C:O2	36:V:40:ARG:NH2	2.37	0.56
18:D:439:U:O2	18:D:440:C:C6	2.58	0.56
18:D:440:C:C4	18:D:441:A:N7	2.73	0.56
22:H:75:VAL:HG23	22:H:81:GLU:OE2	2.04	0.56
29:O:118:LEU:CD2	29:O:124:ARG:HA	2.36	0.56
34:T:21:ASP:OD1	34:T:22:THR:N	2.34	0.56
41:a:930:G:H1'	46:f:25:LEU:HD21	1.87	0.56
41:a:1266:G:OP1	49:i:16:ARG:NE	2.35	0.56
41:a:1568:G:N7	48:h:28:LYS:NZ	2.52	0.56
41:a:2191:A:H2'	41:a:2192:U:H6	1.71	0.56
41:a:2192:U:H2'	41:a:2193:G:C8	2.40	0.56
42:b:51:VAL:C	42:b:62:LYS:NZ	2.63	0.56
44:d:48:U:H2'	44:d:49:C:C6	2.40	0.56
62:v:49:ALA:C	62:v:53:MET:HE1	2.30	0.56
5:4:80:HIS:CD2	5:4:81:PRO:HD2	2.41	0.56
8:7:1:A:H2'	8:7:2:U:H6	1.70	0.56
11:AA:888:THR:O	16:AG:105:ILE:HD13	2.05	0.56
12:AB:38:ILE:HD13	12:AB:160:ARG:HH12	1.70	0.56
16:AG:457:GLU:HG2	16:AG:489:CYS:SG	2.45	0.56
21:G:5:SER:OG	21:G:8:ASP:OD2	2.24	0.56
21:G:131:LYS:HG3	21:G:132:LYS:HD2	1.87	0.56
24:J:121:LYS:HG2	24:J:131:ASN:ND2	2.20	0.56
24:J:196:ASN:ND2	24:J:199:LEU:HG	2.20	0.56
27:M:23:LEU:HD12	27:M:23:LEU:H	1.70	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:64:A:H2'	41:a:65:U:H6	1.71	0.56
41:a:589:U:H2'	41:a:590:A:H8	1.71	0.56
41:a:1198:U:H2'	41:a:1199:U:C6	2.40	0.56
41:a:1266:G:O2'	41:a:2012:G:O6	2.22	0.56
41:a:1570:A:H2'	41:a:1571:A:C8	2.41	0.56
41:a:2514:U:H2'	41:a:2515:C:C6	2.40	0.56
44:d:45:A:H5'	54:n:92:ARG:NH1	2.20	0.56
52:l:1:MET:HE1	52:l:20:GLY:N	2.21	0.56
54:n:62:GLY:C	54:n:95:ARG:NH2	2.63	0.56
56:p:85:LYS:HE3	56:p:164:TYR:CE1	2.41	0.56
63:w:35:LYS:NZ	63:w:112:TYR:HE2	2.04	0.56
5:4:53:LYS:HB2	5:4:56:PHE:CE2	2.40	0.56
5:4:80:HIS:CE1	5:4:83:LYS:HG3	2.41	0.56
11:AA:371:ARG:NH1	12:AB:67:THR:OG1	2.34	0.56
13:AC:45:ARG:HD3	13:AD:38:THR:HB	1.87	0.56
16:AG:173:PHE:N	16:AG:173:PHE:CD2	2.73	0.56
16:AG:434:LEU:HD23	16:AG:456:LEU:CG	2.35	0.56
18:D:521:G:O5'	32:R:70:GLU:OE1	2.23	0.56
21:G:35:ARG:HG2	21:G:40:ILE:HG13	1.86	0.56
22:H:100:LYS:O	22:H:104:ASP:N	2.38	0.56
22:H:163:ARG:CB	22:H:298:GLU:HG3	2.35	0.56
32:R:4:VAL:HG13	32:R:5:ASN:N	2.20	0.56
41:a:2315:G:H4'	54:n:127:ASN:HD22	1.71	0.56
45:e:20:ASN:O	45:e:23:ARG:HG2	2.05	0.56
62:v:41:LEU:CD2	62:v:45:GLN:HE22	2.18	0.56
9:9:43:LYS:HD3	9:9:98:GLU:CG	2.36	0.56
12:AB:130:PHE:CZ	12:AB:160:ARG:HG3	2.41	0.56
14:AE:888:CYS:SG	14:AE:889:ASP:N	2.74	0.56
10:B:75:C:P	41:a:2602:A:O2'	2.61	0.56
23:I:140:ASN:O	23:I:144:LEU:HD23	2.06	0.56
31:Q:61:PHE:O	31:Q:64:GLN:HG2	2.06	0.56
45:e:4:LYS:HZ2	45:e:7:ARG:HH12	1.54	0.56
52:l:17:THR:HA	52:l:21:ARG:NH2	2.19	0.56
54:n:4:LEU:HD11	54:n:101:GLU:HA	1.84	0.56
54:n:163:ASP:OD1	54:n:164:GLU:N	2.39	0.56
59:s:133:ALA:C	59:s:135:GLN:N	2.60	0.56
62:v:26:VAL:HG13	62:v:104:GLU:CD	2.30	0.56
2:1:7:HIS:HD2	2:1:10:ALA:HB2	1.68	0.56
5:4:77:VAL:HG11	5:4:79:ARG:NH2	2.21	0.56
16:AG:283:PHE:CE2	16:AG:331:LEU:O	2.58	0.56
18:D:324:G:N1	18:D:327:A:OP2	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:I:151:VAL:HG12	23:I:200:VAL:CG2	2.35	0.56
27:M:103:TRP:CD1	27:M:137:LYS:HZ1	2.22	0.56
31:Q:87:LYS:HG3	31:Q:113:VAL:CG2	2.36	0.56
41:a:84:A:N1	41:a:98:G:O2'	2.37	0.56
41:a:500:G:N1	41:a:503:A:OP2	2.37	0.56
41:a:659:G:H1'	52:l:100:MET:HE1	1.87	0.56
41:a:1365:A:O2'	43:c:11:ARG:NH1	2.38	0.56
41:a:2189:U:O2'	41:a:2190:G:C5'	2.54	0.56
41:a:2517:C:C2	41:a:2542:A:N6	2.74	0.56
47:g:5:ILE:CD1	54:n:64:LYS:HE3	2.36	0.56
3:2:2:ILE:HG21	3:2:42:GLU:OE1	2.06	0.56
4:3:94:ARG:HB3	4:3:103:ILE:HD12	1.86	0.56
12:AB:34:THR:O	12:AB:106:ASP:OD1	2.24	0.56
16:AG:334:TRP:N	16:AG:334:TRP:CD1	2.73	0.56
16:AG:422:GLU:CG	16:AG:426:GLY:HA3	2.35	0.56
10:B:18:G:H1'	10:B:58:A:C2	2.41	0.56
10:B:75:C:O2'	10:B:76:A:C8	2.58	0.56
18:D:1381:U:C2	18:D:1382:C:C6	2.94	0.56
24:J:62:ARG:HA	24:J:72:PHE:CE1	2.41	0.56
28:N:7:ILE:O	28:N:10:MET:HB3	2.05	0.56
34:T:35:GLN:CB	34:T:59:MET:HE1	2.36	0.56
41:a:923:G:O4'	42:b:29:GLU:OE2	2.24	0.56
49:i:40:ARG:HG3	49:i:41:HIS:ND1	2.21	0.56
60:t:59:LYS:CD	60:t:89:ASN:HA	2.36	0.56
18:D:280:C:H1'	36:V:40:ARG:NH1	2.21	0.56
18:D:671:G:O3'	26:L:79:ARG:NH2	2.39	0.56
18:D:946:A:H2'	18:D:947:G:C8	2.41	0.56
38:X:28:THR:HG22	38:X:29:ARG:NH1	2.21	0.56
41:a:839:U:H2'	41:a:840:C:C6	2.40	0.56
41:a:1289:C:C2	41:a:1290:C:C5	2.94	0.56
41:a:1980:G:O2'	41:a:1982:U:OP2	2.22	0.56
41:a:2849:U:P	65:y:93:ARG:NH2	2.78	0.56
46:f:7:ILE:CG2	46:f:55:VAL:HG21	2.35	0.56
54:n:114:PHE:O	54:n:115:ARG:NE	2.34	0.56
58:r:4:ILE:HD12	58:r:17:ASP:O	2.05	0.56
9:9:47:GLU:O	9:9:51:TYR:CE1	2.59	0.56
11:AA:818:VAL:HG22	11:AA:1096:ILE:HG12	1.87	0.56
11:AA:887:VAL:HB	16:AG:105:ILE:CB	2.36	0.56
16:AG:162:ILE:HG22	16:AG:162:ILE:O	2.06	0.56
16:AG:437:LEU:HD22	16:AG:488:ILE:HD12	1.88	0.56
18:D:827:U:O4	18:D:872:A:N1	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:14:GLN:OE1	26:L:83:ALA:HA	2.05	0.56
30:P:15:HIS:HA	30:P:18:ILE:HG22	1.87	0.56
34:T:3:LEU:HB2	34:T:35:GLN:CD	2.31	0.56
41:a:17:G:H2'	41:a:18:U:C6	2.41	0.56
41:a:608:A:H2'	41:a:609:A:C8	2.41	0.56
41:a:2720:U:OP1	65:y:53:ARG:NH2	2.38	0.56
61:u:125:LEU:HD23	61:u:125:LEU:N	2.21	0.56
63:w:24:MET:HA	63:w:24:MET:CE	2.33	0.56
64:x:36:TYR:CD2	64:x:52:SER:HB2	2.41	0.56
5:4:57:TYR:HB3	62:v:136:MET:HE3	1.87	0.56
8:7:6:U:C2	18:D:1493:A:H2	2.24	0.56
9:9:56:ARG:NH2	41:a:1084:A:C1'	2.69	0.56
9:9:73:LYS:HD2	9:9:73:LYS:O	2.06	0.56
11:AA:1070:HIS:NE2	11:AA:1114:GLU:OE1	2.36	0.56
14:AE:78:LEU:C	16:AG:144:LYS:HZ1	2.14	0.56
16:AG:285:ILE:O	16:AG:288:MET:CE	2.54	0.56
16:AG:362:TYR:CZ	16:AG:382:GLU:HG2	2.40	0.56
16:AG:387:VAL:CB	16:AG:388:PRO:CD	2.76	0.56
18:D:73:C:O2	18:D:73:C:H2'	2.06	0.56
22:H:116:VAL:HA	22:H:279:LYS:HD2	1.88	0.56
22:H:117:LYS:CB	22:H:279:LYS:O	2.54	0.56
23:I:150:LYS:HE3	23:I:201:TRP:CZ3	2.41	0.56
23:I:156:ARG:NH1	23:I:193:TYR:O	2.39	0.56
24:J:162:ALA:HA	24:J:165:ARG:CZ	2.36	0.56
25:K:99:ALA:HB2	25:K:124:LEU:HG	1.88	0.56
33:S:87:ALA:HB1	33:S:93:ILE:HG13	1.87	0.56
37:W:64:ASP:OD2	47:g:57:VAL:HG23	2.06	0.56
38:X:71:ARG:NE	47:g:50:ASP:O	2.39	0.56
41:a:372:G:O2'	41:a:400:G:O6	2.16	0.56
41:a:1799:G:C2	48:h:154:LEU:HD23	2.41	0.56
41:a:2063:C:O2	41:a:2063:C:H2'	2.06	0.56
45:e:2:LYS:HE2	45:e:49:ASP:HA	1.88	0.56
47:g:37:CYS:N	47:g:40:CYS:SG	2.71	0.56
50:j:3:GLY:N	50:j:204:LYS:HE2	2.20	0.56
50:j:91:THR:HB	50:j:94:GLN:HG2	1.87	0.56
56:p:94:TYR:CE1	56:p:107:LEU:O	2.59	0.56
4:3:47:LYS:HE3	41:a:483:A:OP1	2.07	0.55
9:9:24:SER:CB	9:9:86:MET:HE2	2.36	0.55
9:9:98:GLU:HA	9:9:101:LYS:HZ3	1.71	0.55
16:AG:354:ALA:HA	16:AG:357:ASP:H	1.70	0.55
16:AG:425:LEU:N	16:AG:429:LYS:HG3	2.20	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:588:G:C5'	28:N:3:MET:HE1	2.36	0.55
18:D:652:U:C3'	28:N:56:LYS:HZ1	2.18	0.55
25:K:112:ARG:HG3	25:K:116:GLU:OE2	2.06	0.55
35:U:14:ARG:HD3	35:U:42:ILE:HG21	1.86	0.55
41:a:1082:U:O4	41:a:1086:A:N1	2.39	0.55
41:a:2376:A:N3	64:x:111:ARG:NH1	2.54	0.55
41:a:2803:G:H2'	41:a:2804:U:C6	2.41	0.55
48:h:62:TYR:CD2	48:h:64:ILE:HD13	2.41	0.55
51:k:5:ILE:HG23	51:k:28:ARG:NH1	2.21	0.55
52:l:62:GLN:HA	52:l:69:ARG:NH2	2.20	0.55
65:y:106:LYS:HA	65:y:109:ARG:NH2	2.20	0.55
2:1:31:GLN:OE1	2:1:31:GLN:N	2.29	0.55
5:4:57:TYR:CB	62:v:136:MET:CE	2.84	0.55
10:A:16:C:O2	41:a:2181:U:H5'	2.06	0.55
11:AA:516:ASP:OD1	11:AA:516:ASP:O	2.22	0.55
11:AA:887:VAL:HG12	16:AG:105:ILE:HG23	1.88	0.55
16:AG:287:ALA:HB1	16:AG:331:LEU:CB	2.36	0.55
16:AG:320:ARG:HB2	16:AG:323:GLN:HB2	1.88	0.55
10:B:32:C:H5'	18:D:1340:A:O3'	2.06	0.55
18:D:502:A:O3'	32:R:116:LYS:HE2	2.06	0.55
22:H:118:GLY:O	22:H:133:GLY:HA2	2.06	0.55
22:H:268:VAL:HG23	22:H:269:ALA:H	1.72	0.55
26:L:42:TRP:CZ2	26:L:102:MET:CG	2.88	0.55
34:T:3:LEU:HD22	34:T:8:THR:HG22	1.88	0.55
41:a:2328:A:H2'	41:a:2329:U:H6	1.68	0.55
51:k:19:HIS:ND1	51:k:41:PRO:HD2	2.21	0.55
60:t:59:LYS:HD2	60:t:89:ASN:HA	1.87	0.55
60:t:112:PHE:CD2	60:t:115:ILE:HD12	2.39	0.55
62:v:70:ASP:C	62:v:92:TRP:HZ3	2.13	0.55
66:z:78:LYS:CD	66:z:117:LEU:CD2	2.84	0.55
6:5:9:DG:OP2	11:AA:473:ARG:HG2	2.05	0.55
8:7:12:U:H3'	8:7:12:U:H6	1.70	0.55
10:A:18:G:H1'	10:A:58:A:C2	2.42	0.55
10:A:22:G:O2'	10:A:23:C:P	2.65	0.55
16:AG:288:MET:HE2	16:AG:293:VAL:HB	1.89	0.55
16:AG:380:THR:C	16:AG:384:LEU:CD1	2.79	0.55
19:E:19:LYS:HG3	19:E:20:HIS:N	2.22	0.55
19:E:32:ILE:HG23	19:E:75:HIS:NE2	2.21	0.55
25:K:45:ARG:C	25:K:71:MET:HE2	2.32	0.55
25:K:74:VAL:HG11	25:K:144:LEU:HD12	1.88	0.55
27:M:71:PRO:HG3	27:M:103:TRP:CH2	2.42	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:728:G:C4	41:a:730:A:C8	2.95	0.55
41:a:1038:G:H2'	41:a:1039:A:C8	2.42	0.55
41:a:1349:C:C2	41:a:1350:C:C5	2.95	0.55
47:g:61:ASN:OD1	47:g:65:ASN:ND2	2.40	0.55
52:l:42:GLY:CA	52:l:92:HIS:HE1	2.18	0.55
56:p:109:PHE:CE1	56:p:152:ARG:NH1	2.74	0.55
59:s:37:ARG:HD3	59:s:39:LYS:HD2	1.88	0.55
63:w:67:PHE:O	63:w:71:ARG:N	2.39	0.55
4:3:26:LYS:HE2	4:3:26:LYS:HA	1.89	0.55
9:9:58:THR:HG21	9:9:81:LEU:CD2	2.31	0.55
9:9:112:ALA:O	9:9:123:ILE:CD1	2.54	0.55
12:AB:4:TRP:CD2	12:AB:27:ASN:ND2	2.75	0.55
13:AC:28:LEU:HD22	13:AC:201:LEU:HD23	1.88	0.55
16:AG:307:ILE:HD11	16:AG:336:LEU:HB2	1.87	0.55
18:D:1227:A:H5'	38:X:110:LYS:HZ1	1.71	0.55
18:D:1366:C:O2'	30:P:62:ARG:NH2	2.37	0.55
18:D:1370:G:C2	18:D:1371:G:C8	2.95	0.55
30:P:47:GLU:OE1	30:P:67:ILE:O	2.24	0.55
38:X:95:LEU:O	38:X:109:ARG:NH2	2.39	0.55
41:a:1469:A:H2'	41:a:1470:A:C8	2.42	0.55
45:e:23:ARG:HG3	45:e:24:GLU:N	2.21	0.55
50:j:55:LYS:NZ	50:j:60:VAL:N	2.55	0.55
54:n:5:HIS:HD2	54:n:97:TRP:CE2	2.24	0.55
55:o:16:LYS:HD2	55:o:65:ALA:HB1	1.88	0.55
59:s:73:VAL:HG12	59:s:74:TYR:N	2.20	0.55
8:7:11:U:O2	8:7:11:U:H2'	2.06	0.55
9:9:98:GLU:HA	9:9:101:LYS:NZ	2.22	0.55
9:9:122:GLN:HG2	9:9:123:ILE:H	1.70	0.55
16:AG:168:LEU:CD2	16:AG:231:GLY:HA2	2.09	0.55
16:AG:223:ILE:HD13	16:AG:223:ILE:C	2.32	0.55
10:B:5:G:H2'	10:B:6:G:C5'	2.35	0.55
17:C:54:GLN:HG3	17:C:55:LEU:N	2.21	0.55
18:D:276:G:H5'	36:V:17:MET:CE	2.35	0.55
18:D:1329:A:OP1	38:X:28:THR:HB	2.06	0.55
21:G:187:VAL:HG21	21:G:199:VAL:HG23	1.89	0.55
26:L:39:LEU:O	26:L:39:LEU:HD12	2.07	0.55
28:N:7:ILE:O	28:N:11:LEU:HD13	2.06	0.55
38:X:28:THR:O	38:X:31:LYS:HG2	2.07	0.55
39:Y:91:LYS:HZ1	39:Y:97:VAL:HG21	1.71	0.55
41:a:16:C:O4'	49:i:15:MET:CE	2.55	0.55
41:a:857:G:H5''	42:b:78:LYS:HZ2	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1818:U:O2'	48:h:153:GLN:O	2.23	0.55
50:j:156:PHE:CD2	59:s:81:ILE:HD13	2.42	0.55
50:j:180:VAL:O	50:j:180:VAL:HG13	2.05	0.55
55:o:30:ARG:NH1	61:u:62:PRO:HB3	2.21	0.55
55:o:55:LEU:HD21	55:o:59:ILE:HD11	1.87	0.55
59:s:13:ARG:CZ	59:s:49:ASP:O	2.54	0.55
5:4:55:GLU:O	5:4:59:GLU:OE1	2.24	0.55
16:AG:299:ASP:HB2	16:AG:303:HIS:HA	1.89	0.55
10:B:22:G:O2'	10:B:23:C:P	2.65	0.55
21:G:27:MET:CE	21:G:187:VAL:HG12	2.37	0.55
23:I:31:ASP:HA	33:S:65:ARG:HH22	1.72	0.55
24:J:138:SER:O	24:J:141:ASP:OD1	2.24	0.55
27:M:79:ARG:O	27:M:79:ARG:HD2	2.06	0.55
30:P:26:VAL:O	30:P:30:LYS:HG3	2.07	0.55
41:a:2314:A:P	54:n:88:LYS:HZ2	2.30	0.55
45:e:2:LYS:HE2	45:e:52:ARG:HE	1.71	0.55
45:e:28:LEU:HD23	45:e:43:LEU:HD22	1.89	0.55
45:e:53:VAL:O	45:e:57:LEU:HD23	2.06	0.55
55:o:30:ARG:HH12	61:u:62:PRO:HB3	1.70	0.55
56:p:149:ARG:NE	56:p:162:VAL:O	2.40	0.55
57:q:1:MET:HE1	57:q:35:GLN:C	2.32	0.55
66:z:92:ARG:HA	66:z:95:LEU:HD12	1.89	0.55
9:9:61:ARG:NH2	41:a:1045:C:O3'	2.40	0.55
16:AG:305:MET:HG3	16:AG:336:LEU:HB3	1.88	0.55
18:D:672:U:O2'	18:D:673:A:H5'	2.06	0.55
31:Q:84:VAL:CG2	31:Q:107:ILE:HD11	2.37	0.55
37:W:17:LYS:HB3	37:W:21:LYS:HZ3	1.71	0.55
39:Y:123:ALA:O	39:Y:126:ARG:HG3	2.07	0.55
40:Z:2:ILE:HB	40:Z:5:ASP:HB3	1.88	0.55
41:a:537:G:OP2	59:s:2:LYS:CE	2.55	0.55
41:a:888:C:N4	41:a:889:C:N4	2.55	0.55
41:a:1665:A:O3'	60:t:66:LYS:HD3	2.07	0.55
41:a:2557:G:H2'	41:a:2558:C:C6	2.42	0.55
41:a:2741:A:O3'	57:q:36:ARG:NH1	2.40	0.55
45:e:56:LEU:O	45:e:59:GLU:HG3	2.06	0.55
48:h:17:VAL:HB	48:h:204:VAL:HG12	1.89	0.55
48:h:67:PHE:CE1	48:h:105:LEU:HD11	2.42	0.55
48:h:107:PRO:HD2	48:h:110:LEU:HD22	1.89	0.55
54:n:134:GLU:N	54:n:134:GLU:OE1	2.40	0.55
65:y:27:GLU:HB3	65:y:87:LYS:NZ	2.22	0.55
2:1:43:ALA:O	2:1:47:VAL:HG12	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:7:1:A:C4	8:7:2:U:C5	2.95	0.55
8:7:8:U:H5'	18:D:1397:C:C2	2.41	0.55
10:A:22:G:H2'	10:A:23:C:H6	1.70	0.55
10:A:37:A:H2'	10:A:38:A:O4'	2.07	0.55
11:AA:633:LEU:HD13	11:AA:644:LEU:HD23	1.89	0.55
12:AB:138:ARG:CD	12:AB:155:LYS:CA	2.84	0.55
16:AG:130:PHE:O	16:AG:186:VAL:HG11	2.07	0.55
18:D:695:A:OP1	31:Q:53:ARG:HD3	2.07	0.55
22:H:9:PHE:O	22:H:13:LEU:HD23	2.06	0.55
23:I:164:ARG:CZ	23:I:166:GLU:OE2	2.54	0.55
24:J:175:ALA:O	24:J:178:MET:SD	2.65	0.55
25:K:77:ASN:HB2	25:K:82:GLN:HG3	1.89	0.55
27:M:23:LEU:O	27:M:27:VAL:HG13	2.07	0.55
38:X:96:PRO:HG3	38:X:102:THR:HG22	1.89	0.55
39:Y:44:LYS:O	39:Y:44:LYS:HD2	2.07	0.55
41:a:151:C:C2	41:a:152:A:C8	2.95	0.55
41:a:2335:A:OP2	64:x:13:ARG:HB3	2.06	0.55
41:a:2351:G:O6	55:o:39:LYS:CE	2.55	0.55
41:a:2483:C:O2	62:v:51:ARG:NH2	2.38	0.55
41:a:2838:G:C4	41:a:2839:G:C8	2.95	0.55
50:j:4:LEU:HD23	50:j:29:VAL:HG11	1.89	0.55
9:9:94:ARG:HB2	9:9:97:LYS:CE	2.36	0.55
11:AA:858:GLY:O	16:AG:38:LYS:NZ	2.39	0.55
12:AB:146:ILE:CG2	30:P:88:MET:O	2.55	0.55
16:AG:131:ARG:O	16:AG:134:GLU:CG	2.47	0.55
16:AG:131:ARG:CA	16:AG:186:VAL:HG11	2.35	0.55
16:AG:168:LEU:CD2	16:AG:230:PRO:O	2.55	0.55
16:AG:168:LEU:CD2	16:AG:230:PRO:C	2.74	0.55
16:AG:392:LEU:HD22	16:AG:395:ILE:HD11	1.87	0.55
10:B:22:G:H2'	10:B:23:C:H6	1.70	0.55
18:D:193:C:O2'	19:E:59:ASP:OD2	2.23	0.55
18:D:679:C:C2	18:D:680:C:C5	2.95	0.55
18:D:750:C:N3	18:D:751:U:C5	2.75	0.55
19:E:6:SER:O	19:E:10:ARG:HG2	2.07	0.55
28:N:7:ILE:O	28:N:11:LEU:CD1	2.55	0.55
28:N:78:VAL:HG12	28:N:85:ILE:HG21	1.89	0.55
33:S:27:LEU:HD21	33:S:48:LEU:N	2.21	0.55
39:Y:45:THR:HG21	39:Y:50:LYS:CD	2.36	0.55
41:a:784:G:H5'	41:a:785:G:OP1	2.06	0.55
41:a:851:C:H2'	41:a:852:U:H6	1.72	0.55
41:a:1386:C:H2'	41:a:1387:A:C8	2.42	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2591:C:H2'	41:a:2592:G:C8	2.42	0.55
41:a:2636:C:O2'	50:j:45:TYR:OH	1.99	0.55
48:h:157:SER:O	48:h:195:VAL:HG11	2.07	0.55
50:j:78:GLY:C	50:j:79:LEU:HD22	2.32	0.55
62:v:96:ILE:HA	62:v:100:LYS:NZ	2.21	0.55
9:9:23:LEU:HD21	9:9:96:PHE:CD1	2.42	0.55
14:AE:886:VAL:HG11	14:AE:1230:THR:HG21	1.89	0.55
16:AG:46:ARG:HH11	16:AG:61:ARG:HB3	1.71	0.55
16:AG:133:HIS:CE1	16:AG:136:GLU:OE1	2.59	0.55
16:AG:320:ARG:NH1	16:AG:320:ARG:HG2	2.22	0.55
16:AG:355:ALA:HB1	16:AG:382:GLU:CB	2.37	0.55
16:AG:448:LEU:HD22	16:AG:467:LEU:HD22	1.89	0.55
17:C:40:VAL:HG13	22:H:339:ARG:HD3	1.89	0.55
18:D:37:U:O4	18:D:397:A:N1	2.40	0.55
18:D:1078:U:O3'	25:K:138:ARG:NH2	2.40	0.55
20:F:26:ALA:HB3	20:F:28:VAL:HG23	1.88	0.55
23:I:132:ARG:HA	23:I:135:LYS:HE3	1.89	0.55
35:U:23:ASP:HB3	35:U:26:ASN:OD1	2.07	0.55
38:X:28:THR:CA	38:X:31:LYS:HE3	2.37	0.55
41:a:1059:G:H2'	41:a:1060:U:C5	2.41	0.55
41:a:1720:U:H2'	41:a:1721:G:O4'	2.06	0.55
46:f:10:THR:HB	46:f:56:LYS:NZ	2.22	0.55
60:t:12:ASP:CG	60:t:85:VAL:HG13	2.31	0.55
62:v:105:MET:HE2	62:v:108:VAL:HG21	1.87	0.55
2:1:62:ASP:OD1	2:1:62:ASP:O	2.24	0.54
6:5:16:DG:OP1	11:AA:175:ARG:NH2	2.39	0.54
8:7:6:U:C2	18:D:1493:A:C2	2.95	0.54
10:A:56:C:H1'	41:a:2112:G:C6	2.41	0.54
12:AB:8:TYR:OH	14:AE:278:ARG:NH2	2.40	0.54
12:AB:120:GLU:CB	12:AB:162:LEU:HD12	2.37	0.54
14:AE:287:ALA:CB	14:AE:292:VAL:HG22	2.37	0.54
10:B:37:A:H2'	10:B:38:A:O4'	2.07	0.54
18:D:50:A:O2'	18:D:360:G:N2	2.40	0.54
18:D:321:A:N7	18:D:328:C:O2'	2.38	0.54
18:D:1439:G:P	19:E:33:LYS:HZ1	2.28	0.54
26:L:99:ALA:HB3	26:L:104:LYS:NZ	2.22	0.54
34:T:80:GLN:O	34:T:84:ARG:HG2	2.05	0.54
41:a:1000:A:H2'	41:a:1001:A:C8	2.43	0.54
41:a:1708:C:C2	41:a:1709:U:C5	2.95	0.54
50:j:11:MET:SD	50:j:11:MET:C	2.90	0.54
59:s:15:TRP:HE3	59:s:135:GLN:O	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
61:u:122:VAL:HG13	61:u:125:LEU:CD1	2.34	0.54
62:v:45:GLN:NE2	62:v:125:PRO:HD2	2.22	0.54
2:1:31:GLN:O	2:1:35:ILE:HG12	2.07	0.54
4:3:7:ARG:HB3	41:a:85:G:OP2	2.08	0.54
4:3:48:PRO:HG3	4:3:56:GLY:HA3	1.89	0.54
10:A:41:C:O4'	27:M:143:ARG:NH1	2.40	0.54
10:A:76:A:N1	41:a:2422:C:C6	2.75	0.54
11:AA:29:SER:O	11:AA:33:ASP:HB2	2.07	0.54
11:AA:360:LEU:HD13	11:AA:378:ARG:HH11	1.72	0.54
12:AB:148:LYS:HD2	30:P:100:ILE:HD13	1.89	0.54
18:D:17:U:H2'	18:D:18:C:C6	2.42	0.54
22:H:5:PHE:O	22:H:9:PHE:HD1	1.90	0.54
22:H:71:ALA:C	22:H:72:LEU:HD12	2.32	0.54
23:I:66:VAL:HB	23:I:101:ILE:HD12	1.88	0.54
27:M:31:MET:SD	27:M:36:LYS:HA	2.47	0.54
31:Q:20:VAL:HB	31:Q:35:THR:HG23	1.89	0.54
41:a:151:C:H2'	41:a:152:A:H8	1.72	0.54
41:a:1353:A:H2'	41:a:1354:A:C8	2.42	0.54
41:a:2683:C:H4'	50:j:13:ARG:NH2	2.22	0.54
48:h:183:LYS:HE2	48:h:266:PHE:HA	1.88	0.54
52:l:188:MET:HE2	52:l:193:VAL:HA	1.89	0.54
60:t:18:ARG:HB3	60:t:45:GLU:HB3	1.89	0.54
5:4:6:ALA:HB3	5:4:65:VAL:CG2	2.37	0.54
9:9:52:MET:HE1	9:9:85:SER:CA	2.38	0.54
9:9:136:ILE:CD1	9:9:139:LEU:HD12	2.25	0.54
11:AA:846:GLY:C	16:AG:104:ARG:NH2	2.65	0.54
16:AG:220:VAL:HG12	16:AG:245:ILE:HG21	1.88	0.54
31:Q:20:VAL:HG13	31:Q:85:MET:HE1	1.89	0.54
41:a:1433:A:H2'	41:a:1434:A:O4'	2.07	0.54
41:a:1779:U:H5	41:a:1784:A:N7	2.06	0.54
41:a:2299:U:H2'	41:a:2300:C:H6	1.71	0.54
41:a:2454:G:C4	41:a:2455:G:C8	2.96	0.54
47:g:2:LYS:HB2	47:g:5:ILE:HD11	1.88	0.54
52:l:60:TRP:CD1	52:l:65:THR:HG21	2.43	0.54
54:n:46:ASP:CG	54:n:49:LEU:HD13	2.33	0.54
59:s:73:VAL:C	59:s:74:TYR:HD2	2.14	0.54
59:s:74:TYR:CE2	59:s:92:MET:HE1	2.42	0.54
62:v:45:GLN:HE21	62:v:124:LEU:HD23	1.72	0.54
1:0:10:LYS:HE2	41:a:996:A:OP1	2.07	0.54
1:0:13:ARG:NH2	66:z:97:ASP:OD1	2.40	0.54
6:5:10:DT:C7	11:AA:62:TYR:HB3	2.37	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:7:3:G:C3'	18:D:1500:A:N6	2.71	0.54
9:9:129:LEU:HD12	9:9:129:LEU:C	2.32	0.54
11:AA:243:PRO:HB2	11:AA:274:ILE:HG23	1.90	0.54
11:AA:985:GLU:HB3	11:AA:988:LYS:HB2	1.88	0.54
11:AA:1142:ARG:NH2	11:AA:1166:ASP:OD1	2.38	0.54
12:AB:134:ASP:OD1	12:AB:135:GLY:N	2.37	0.54
16:AG:285:ILE:HD12	16:AG:293:VAL:HG11	1.88	0.54
17:C:34:THR:HG22	17:C:36:SER:H	1.72	0.54
18:D:106:C:N4	19:E:10:ARG:NH2	2.56	0.54
20:F:63:GLU:HG2	20:F:67:ARG:NH2	2.22	0.54
23:I:155:GLY:O	23:I:196:ILE:HG12	2.07	0.54
28:N:87:LYS:HD2	28:N:91:GLU:CB	2.37	0.54
36:V:31:HIS:ND1	36:V:32:PRO:HD2	2.23	0.54
38:X:81:MET:CE	38:X:92:ARG:HD3	2.38	0.54
38:X:95:LEU:HB3	38:X:96:PRO:CD	2.37	0.54
41:a:3:U:H2'	41:a:4:U:C6	2.42	0.54
41:a:960:A:H61	62:v:82:MET:HE1	1.72	0.54
41:a:2020:A:C2	41:a:2022:U:O4'	2.61	0.54
41:a:2308:G:C5	54:n:77:PHE:HZ	2.26	0.54
52:l:69:ARG:HA	52:l:69:ARG:CZ	2.37	0.54
61:u:76:GLU:CG	61:u:111:ILE:HD13	2.37	0.54
65:y:15:GLN:NE2	65:y:16:ASP:OD1	2.40	0.54
2:1:25:ARG:NH2	41:a:519:U:O3'	2.40	0.54
3:2:56:GLU:OE1	3:2:87:LEU:CA	2.56	0.54
5:4:80:HIS:ND1	5:4:83:LYS:N	2.55	0.54
11:AA:811:ASN:HA	11:AA:815:SER:HB2	1.90	0.54
14:AE:679:TYR:OH	14:AE:754:ILE:O	2.23	0.54
17:C:60:LYS:HD2	18:D:734:G:O2'	2.07	0.54
21:G:19:GLN:CD	22:H:76:GLU:HB3	2.33	0.54
24:J:105:MET:HE1	24:J:171:LEU:HB3	1.89	0.54
30:P:5:ARG:HG3	30:P:7:ARG:NH1	2.23	0.54
41:a:1019:U:O4	41:a:1142:A:N1	2.40	0.54
41:a:1565:C:OP1	48:h:18:LYS:NZ	2.26	0.54
41:a:1656:C:OP1	50:j:141:ARG:CD	2.55	0.54
41:a:1790:C:H2'	41:a:1791:A:C5	2.43	0.54
41:a:2064:C:H2'	41:a:2065:C:C6	2.42	0.54
11:AA:143:ARG:NH2	11:AA:512:SER:O	2.40	0.54
14:AE:901:ARG:HH21	14:AE:906:GLY:HA2	1.73	0.54
14:AE:968:ASN:HA	14:AE:1117:SER:HB2	1.90	0.54
16:AG:63:LEU:HD23	16:AG:63:LEU:N	2.21	0.54
16:AG:358:THR:O	16:AG:358:THR:OG1	2.24	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1071:C:H2'	18:D:1072:G:H8	1.71	0.54
19:E:24:ARG:O	19:E:28:MET:HG2	2.08	0.54
22:H:279:LYS:HA	22:H:331:MET:HB3	1.89	0.54
24:J:67:VAL:HB	24:J:72:PHE:CE1	2.43	0.54
37:W:12:ASP:OD2	37:W:35:SER:OG	2.14	0.54
41:a:1348:C:C5	41:a:1349:C:C5	2.95	0.54
41:a:2356:U:H4'	42:b:20:ARG:HH11	1.73	0.54
41:a:2742:G:P	57:q:36:ARG:NH1	2.80	0.54
60:t:73:ASP:OD2	65:y:80:VAL:HG11	2.08	0.54
2:1:24:ILE:O	2:1:71:VAL:HG21	2.08	0.54
5:4:35:GLU:OE1	5:4:93:ARG:CZ	2.56	0.54
6:5:10:DT:O3'	12:AB:68:THR:C	2.47	0.54
12:AB:78:PHE:CD1	12:AB:85:PRO:HB3	2.42	0.54
16:AG:201:LYS:HB3	16:AG:201:LYS:HZ3	1.71	0.54
16:AG:327:LEU:HD12	16:AG:327:LEU:O	2.08	0.54
18:D:375:U:OP1	35:U:70:ARG:CG	2.54	0.54
18:D:399:G:H2'	18:D:400:C:C6	2.43	0.54
21:G:18:HIS:HA	22:H:43:LYS:HD2	1.90	0.54
30:P:6:ILE:HG21	30:P:100:ILE:HD11	1.89	0.54
39:Y:45:THR:HG21	39:Y:50:LYS:HZ2	1.72	0.54
41:a:1563:U:H2'	41:a:1564:C:C6	2.43	0.54
41:a:2766:A:C2	41:a:2767:C:C6	2.96	0.54
41:a:2822:G:O2'	41:a:2824:C:OP2	2.20	0.54
62:v:97:GLN:H	62:v:100:LYS:CE	2.21	0.54
1:0:10:LYS:CE	41:a:994:C:O2'	2.56	0.54
2:1:41:LYS:HZ2	41:a:2009:A:P	2.30	0.54
7:6:20:DA:C6	8:7:51:G:N2	2.76	0.54
10:A:60:U:P	10:A:61:C:H41	2.30	0.54
14:AE:1346:GLY:O	14:AE:1350:ASN:ND2	2.41	0.54
16:AG:448:LEU:HD11	16:AG:481:LEU:HD13	1.89	0.54
17:C:13:PHE:CZ	17:C:21:ILE:HG12	2.42	0.54
18:D:547:A:OP1	24:J:70:ARG:NH2	2.41	0.54
18:D:713:G:H2'	18:D:714:G:C8	2.42	0.54
18:D:1399:C:O2	18:D:1502:A:N6	2.40	0.54
19:E:25:ARG:HD3	19:E:66:LEU:HD21	1.90	0.54
20:F:19:PHE:CE2	31:Q:110:ILE:HG22	2.42	0.54
26:L:3:HIS:CG	26:L:65:GLU:OE1	2.61	0.54
27:M:79:ARG:HB2	27:M:84:THR:HG23	1.90	0.54
41:a:1084:A:N7	41:a:1085:A:C6	2.76	0.54
41:a:1321:A:C4	41:a:1322:A:C8	2.95	0.54
41:a:2259:U:C2	41:a:2260:C:C6	2.96	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:f:10:THR:HG23	46:f:11:ARG:HG2	1.90	0.54
50:j:124:ARG:NH1	50:j:161:MET:O	2.41	0.54
54:n:16:LEU:HD11	54:n:169:LEU:CD1	2.38	0.54
54:n:71:ARG:O	54:n:81:GLN:NE2	2.41	0.54
61:u:57:LEU:HA	61:u:60:ARG:HD2	1.89	0.54
63:w:24:MET:HE2	63:w:34:ILE:HD13	1.87	0.54
63:w:44:LEU:HD22	63:w:113:ILE:HD13	1.90	0.54
1:0:10:LYS:CE	41:a:996:A:OP1	2.56	0.54
3:2:19:LYS:NZ	41:a:1340:U:OP1	2.34	0.54
5:4:19:ARG:NH1	44:d:95:U:OP2	2.41	0.54
11:AA:377:THR:HB	12:AB:65:HIS:NE2	2.23	0.54
13:AC:100:LEU:HD23	13:AC:115:ILE:HG21	1.90	0.54
14:AE:1045:THR:O	14:AE:1067:ARG:NH2	2.41	0.54
16:AG:336:LEU:HD12	16:AG:336:LEU:N	2.01	0.54
18:D:110:C:O2'	35:U:25:ARG:O	2.21	0.54
18:D:339:C:H2'	18:D:340:U:H6	1.73	0.54
21:G:30:PHE:CD1	21:G:45:LYS:NZ	2.74	0.54
22:H:268:VAL:HG23	22:H:269:ALA:N	2.22	0.54
25:K:131:THR:O	25:K:131:THR:HG22	2.08	0.54
26:L:62:MET:HA	26:L:62:MET:HE3	1.90	0.54
27:M:38:THR:O	27:M:42:ILE:HG12	2.08	0.54
28:N:83:LEU:HD21	32:R:4:VAL:HG11	1.89	0.54
32:R:102:LEU:O	32:R:103:ASP:OD1	2.26	0.54
39:Y:77:VAL:HG13	39:Y:81:LYS:HZ3	1.73	0.54
41:a:150:U:H2'	41:a:151:C:H6	1.72	0.54
41:a:2038:G:H2'	41:a:2039:U:O4'	2.08	0.54
41:a:2537:U:H2'	41:a:2538:C:C6	2.43	0.54
52:l:147:LEU:HD11	52:l:170:ARG:HD2	1.89	0.54
56:p:33:LEU:HD11	56:p:136:ALA:HB1	1.89	0.54
62:v:20:LEU:H	62:v:20:LEU:HD23	1.73	0.54
3:2:56:GLU:CD	3:2:88:LYS:HA	2.32	0.54
6:5:9:DG:H2'	12:AB:71:ALA:HA	1.90	0.54
12:AB:112:PRO:HB3	12:AB:132:GLU:OE1	2.07	0.54
14:AE:1175:LEU:O	14:AE:1187:GLU:HA	2.08	0.54
10:B:60:U:P	10:B:61:C:H41	2.30	0.54
19:E:78:ASN:C	19:E:82:GLN:OE1	2.50	0.54
20:F:19:PHE:CE1	20:F:23:CYS:SG	3.00	0.54
28:N:50:LYS:HE2	28:N:50:LYS:HA	1.89	0.54
37:W:47:LEU:O	37:W:62:VAL:HG23	2.08	0.54
41:a:2747:G:O6	41:a:2755:C:H5''	2.08	0.54
62:v:1:MET:HA	62:v:1:MET:CE	2.38	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:7:13:U:H6	8:7:13:U:H5''	1.73	0.53
9:9:97:LYS:HD3	9:9:130:PRO:HG3	1.89	0.53
11:AA:684:ASN:OD1	11:AA:687:ARG:NH2	2.41	0.53
11:AA:860:ALA:HB3	16:AG:38:LYS:HE2	1.89	0.53
18:D:280:C:H1'	36:V:40:ARG:CZ	2.37	0.53
18:D:736:C:H2'	18:D:737:C:C6	2.42	0.53
18:D:1076:U:OP1	21:G:174:LYS:NZ	2.40	0.53
25:K:14:LYS:HD3	25:K:117:VAL:CG1	2.38	0.53
28:N:31:LYS:HE3	28:N:31:LYS:HA	1.89	0.53
29:O:5:GLN:NE2	29:O:22:LYS:NZ	2.56	0.53
30:P:27:GLU:HA	30:P:30:LYS:NZ	2.23	0.53
39:Y:80:LYS:CG	39:Y:86:LYS:O	2.56	0.53
41:a:134:G:H2'	41:a:135:U:C6	2.43	0.53
41:a:589:U:H2'	41:a:590:A:C8	2.42	0.53
41:a:2867:G:C6	65:y:21:ARG:NH1	2.76	0.53
54:n:72:LYS:HA	54:n:81:GLN:HE22	1.74	0.53
62:v:96:ILE:HA	62:v:100:LYS:CE	2.38	0.53
64:x:38:GLN:HG2	64:x:50:ALA:CB	2.38	0.53
65:y:31:TRP:CZ3	65:y:40:LEU:HD11	2.42	0.53
3:2:19:LYS:O	3:2:22:THR:OG1	2.23	0.53
3:2:53:VAL:HG21	3:2:87:LEU:HD23	1.91	0.53
6:5:15:DT:C2	11:AA:183:TRP:CD1	2.94	0.53
7:6:14:DC:H2'	7:6:15:DC:C6	2.43	0.53
9:9:36:ASP:N	9:9:36:ASP:OD1	2.40	0.53
12:AB:154:VAL:HG11	12:AB:162:LEU:HD21	1.90	0.53
16:AG:374:VAL:O	16:AG:374:VAL:HG12	2.08	0.53
18:D:76:G:C4	18:D:77:A:C8	2.95	0.53
18:D:750:C:C2	18:D:751:U:C5	2.97	0.53
19:E:32:ILE:CG2	19:E:75:HIS:NE2	2.71	0.53
21:G:148:LEU:HD22	21:G:151:ILE:HD11	1.89	0.53
22:H:45:GLU:OE1	22:H:45:GLU:N	2.42	0.53
25:K:86:LYS:N	25:K:86:LYS:HD2	2.23	0.53
25:K:157:ARG:NE	28:N:43:GLU:O	2.40	0.53
27:M:85:TYR:CD2	27:M:151:PHE:CZ	2.97	0.53
39:Y:33:ASN:HB3	39:Y:36:GLU:CG	2.30	0.53
41:a:839:U:H2'	41:a:840:C:H6	1.73	0.53
41:a:1817:G:C6	41:a:1818:U:C5	2.95	0.53
41:a:2287:A:C8	41:a:2289:G:C8	2.97	0.53
50:j:55:LYS:NZ	50:j:59:ARG:HB3	2.23	0.53
53:m:25:LYS:HG3	53:m:29:GLN:HE22	1.73	0.53
60:t:2:ILE:CG2	60:t:8:LEU:HD11	2.38	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:w:33:ILE:CD1	63:w:112:TYR:CD1	2.91	0.53
3:2:6:ARG:NH2	3:2:10:VAL:HG22	2.24	0.53
3:2:54:GLU:OE1	3:2:54:GLU:N	2.38	0.53
10:A:16:C:O2	41:a:2181:U:C5'	2.57	0.53
11:AA:855:PRO:HD3	16:AG:103:ASP:OD2	2.08	0.53
16:AG:398:LEU:HG	16:AG:398:LEU:O	2.09	0.53
17:C:38:LYS:HE2	17:C:38:LYS:CA	2.37	0.53
17:C:39:ILE:HD12	18:D:719:C:O2	2.07	0.53
21:G:53:ALA:CA	21:G:56:GLU:OE1	2.57	0.53
27:M:27:VAL:HG12	27:M:43:VAL:HG21	1.90	0.53
41:a:581:C:H2'	41:a:582:A:H8	1.73	0.53
41:a:588:U:H2'	41:a:589:U:C6	2.43	0.53
41:a:813:U:H2'	41:a:814:C:H6	1.74	0.53
41:a:1830:C:C2	41:a:1831:G:C8	2.95	0.53
41:a:2297:A:C2	41:a:2320:U:O4'	2.61	0.53
48:h:175:ARG:CG	48:h:181:MET:HE1	2.35	0.53
51:k:8:LYS:HZ1	51:k:54:ILE:HG12	1.72	0.53
60:t:1:MET:CE	60:t:32:TYR:HB3	2.38	0.53
60:t:41:ILE:HD11	60:t:86:LEU:HD22	1.89	0.53
63:w:94:TYR:C	63:w:116:VAL:HG23	2.32	0.53
5:4:80:HIS:ND1	5:4:83:LYS:HB2	2.23	0.53
8:7:9:U:C4	18:D:1196:A:C8	2.97	0.53
11:AA:628:HIS:HB3	11:AA:647:ARG:HH21	1.73	0.53
11:AA:855:PRO:HA	16:AG:35:THR:OG1	2.06	0.53
12:AB:120:GLU:N	12:AB:162:LEU:HG	2.24	0.53
13:AD:16:ILE:HG23	13:AD:26:VAL:HG22	1.91	0.53
14:AE:66:LYS:HB2	14:AE:69:GLU:CB	2.38	0.53
16:AG:235:LYS:HG2	16:AG:235:LYS:O	2.07	0.53
18:D:197:A:HI'	18:D:198:G:OP2	2.08	0.53
19:E:22:ALA:CA	19:E:25:ARG:HH21	2.21	0.53
23:I:85:GLU:HG3	23:I:89:LYS:NZ	2.23	0.53
24:J:61:VAL:HG21	24:J:200:ILE:HD11	1.90	0.53
26:L:39:LEU:HB3	26:L:62:MET:HE1	1.88	0.53
29:O:10:GLY:HA3	29:O:78:ALA:O	2.08	0.53
32:R:56:ARG:HA	32:R:62:GLU:OE2	2.08	0.53
32:R:74:LEU:HD11	32:R:80:ILE:HD11	1.90	0.53
39:Y:102:ARG:HG3	39:Y:140:GLU:C	2.33	0.53
41:a:15:G:H2'	49:i:15:MET:HE1	1.90	0.53
41:a:743:A:O2'	41:a:1659:G:OP1	2.25	0.53
41:a:1774:C:O2	41:a:1774:C:H2'	2.07	0.53
41:a:2849:U:O5'	65:y:93:ARG:NH2	2.42	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:62:GLN:O	52:l:69:ARG:CD	2.56	0.53
52:l:175:ILE:CG2	52:l:180:LEU:HD11	2.38	0.53
59:s:133:ALA:C	59:s:135:GLN:H	2.16	0.53
61:u:111:ILE:HD12	61:u:111:ILE:N	2.24	0.53
62:v:1:MET:CE	62:v:44:ARG:HA	2.38	0.53
64:x:48:LEU:CD2	64:x:87:ILE:HD11	2.38	0.53
66:z:78:LYS:HD3	66:z:117:LEU:HD23	1.88	0.53
2:1:33:LEU:HD23	2:1:51:LEU:HD23	1.89	0.53
2:1:41:LYS:HE3	41:a:2010:G:P	2.48	0.53
7:6:20:DA:N6	8:7:51:G:C2	2.76	0.53
9:9:3:LEU:HD12	9:9:4:ASN:CA	2.37	0.53
9:9:51:TYR:HB2	9:9:88:HIS:HB2	1.90	0.53
11:AA:103:VAL:HB	11:AA:114:VAL:HG11	1.89	0.53
16:AG:179:VAL:HB	16:AG:199:ARG:HH11	1.73	0.53
16:AG:363:LEU:CG	16:AG:410:ALA:CB	2.71	0.53
19:E:25:ARG:HD3	19:E:66:LEU:HD11	1.90	0.53
28:N:87:LYS:HG2	28:N:92:LEU:HA	1.90	0.53
29:O:112:GLU:HG3	29:O:115:LYS:HZ3	1.73	0.53
30:P:47:GLU:OE1	30:P:67:ILE:HG23	2.09	0.53
32:R:25:GLU:OE1	32:R:59:ASN:ND2	2.42	0.53
32:R:27:CYS:SG	32:R:30:LYS:NZ	2.68	0.53
32:R:35:THR:OG1	32:R:56:ARG:HG3	2.08	0.53
41:a:372:G:O2'	43:c:54:LYS:HE3	2.09	0.53
41:a:414:C:H2'	41:a:415:A:C8	2.44	0.53
41:a:1278:C:H5'	63:w:24:MET:SD	2.49	0.53
50:j:4:LEU:HG	50:j:32:ASN:OD1	2.08	0.53
51:k:8:LYS:NZ	51:k:54:ILE:HG12	2.23	0.53
52:l:60:TRP:NE1	52:l:69:ARG:HH12	2.07	0.53
60:t:14:SER:OG	60:t:86:LEU:HD12	2.08	0.53
11:AA:143:ARG:NH1	11:AA:507:GLY:O	2.32	0.53
16:AG:434:LEU:CG	16:AG:456:LEU:N	2.71	0.53
10:B:38:A:O2'	18:D:790:A:P	2.66	0.53
18:D:1329:A:P	38:X:28:THR:HB	2.48	0.53
18:D:1358:U:O4	18:D:1363:A:N1	2.41	0.53
22:H:321:VAL:HG13	22:H:322:VAL:HG23	1.91	0.53
33:S:42:TRP:CH2	47:g:66:ILE:HG23	2.42	0.53
36:V:7:THR:CB	36:V:60:GLU:OE2	2.56	0.53
39:Y:33:ASN:HB2	39:Y:36:GLU:OE2	2.08	0.53
41:a:1412:U:C4	41:a:1413:A:N7	2.77	0.53
41:a:1657:U:O2'	50:j:138:LEU:HD22	2.08	0.53
41:a:2847:U:H2'	41:a:2848:G:O4'	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:g:11:GLU:HA	47:g:25:ARG:HA	1.91	0.53
54:n:93:GLY:HA2	54:n:97:TRP:CZ3	2.44	0.53
56:p:89:LEU:HD22	56:p:162:VAL:HG22	1.91	0.53
62:v:97:GLN:N	62:v:100:LYS:CE	2.72	0.53
3:2:30:ILE:HG22	3:2:32:LEU:CD2	2.38	0.53
5:4:35:GLU:OE1	5:4:36:ALA:O	2.27	0.53
9:9:3:LEU:CG	9:9:4:ASN:H	2.21	0.53
12:AB:154:VAL:HB	12:AB:159:PHE:HB3	1.91	0.53
16:AG:131:ARG:HA	16:AG:186:VAL:CG1	2.38	0.53
16:AG:447:LYS:O	16:AG:470:ILE:HD11	2.05	0.53
10:B:31:G:O2'	18:D:1340:A:H4'	2.08	0.53
18:D:274:A:H4'	36:V:16:LYS:NZ	2.24	0.53
23:I:24:ALA:C	23:I:29:PHE:CE2	2.87	0.53
27:M:4:ARG:CG	27:M:5:ARG:NH1	2.72	0.53
29:O:67:VAL:C	29:O:68:LYS:HD2	2.33	0.53
41:a:78:U:H2'	41:a:79:C:C6	2.44	0.53
41:a:2308:G:C6	54:n:77:PHE:CZ	2.97	0.53
50:j:48:ILE:HG12	50:j:84:LEU:HD21	1.90	0.53
60:t:35:VAL:HG23	60:t:63:VAL:HA	1.89	0.53
65:y:9:GLU:OE2	65:y:55:LEU:N	2.37	0.53
1:0:5:PHE:HB3	1:0:59:ILE:HD12	1.91	0.53
11:AA:855:PRO:HD2	11:AA:887:VAL:HG11	1.91	0.53
12:AB:5:TYR:HA	12:AB:88:VAL:HG22	1.90	0.53
14:AE:514:THR:HG21	14:AE:596:LEU:HD12	1.90	0.53
16:AG:442:ARG:C	16:AG:446:PHE:CD2	2.85	0.53
27:M:103:TRP:HB3	27:M:137:LYS:HD3	1.90	0.53
32:R:110:ARG:NE	32:R:117:TYR:CD2	2.76	0.53
41:a:150:U:H2'	41:a:151:C:C6	2.44	0.53
41:a:1243:C:H1'	61:u:4:ASN:O	2.09	0.53
41:a:1379:U:H4'	41:a:1380:G:OP1	2.09	0.53
42:b:21:LEU:HD21	42:b:41:ARG:NE	2.21	0.53
47:g:47:LYS:HD3	54:n:178:ARG:HH22	1.74	0.53
50:j:131:ASP:OD2	50:j:132:ALA:N	2.42	0.53
56:p:9:VAL:HG22	56:p:69:ARG:NH1	2.24	0.53
60:t:99:ILE:HD13	60:t:118:LEU:CB	2.39	0.53
63:w:24:MET:HE3	63:w:24:MET:CA	2.34	0.53
4:3:26:LYS:HE3	4:3:37:GLU:HB2	1.90	0.53
6:5:17:DC:H5''	6:5:17:DC:H6	1.74	0.53
11:AA:1117:LEU:HD12	11:AA:1195:ILE:HG12	1.90	0.53
12:AB:89:PRO:HG2	14:AE:290:ILE:HG22	1.91	0.53
13:AD:28:LEU:HD12	13:AD:201:LEU:HD23	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1439:G:P	19:E:33:LYS:NZ	2.82	0.53
19:E:56:PRO:HG2	19:E:57:ILE:HD12	1.90	0.53
21:G:23:TRP:HB3	21:G:39:HIS:CE1	2.44	0.53
29:O:47:VAL:O	29:O:80:ARG:HD2	2.09	0.53
29:O:88:MET:HE1	29:O:95:ARG:HA	1.89	0.53
38:X:22:ILE:HB	38:X:25:VAL:CG1	2.39	0.53
41:a:995:C:N4	59:s:2:LYS:HG3	2.24	0.53
41:a:1818:U:C6	48:h:156:ARG:NH2	2.77	0.53
8:7:3:G:C4'	18:D:1500:A:H62	2.22	0.53
12:AB:148:LYS:HZ1	12:AB:150:ILE:CD1	2.19	0.53
16:AG:440:VAL:HG21	16:AG:481:LEU:HD22	1.88	0.53
17:C:12:ARG:NH2	17:C:13:PHE:CE1	2.77	0.53
18:D:62:U:H2'	18:D:63:C:C6	2.44	0.53
18:D:679:C:H2'	18:D:680:C:H6	1.74	0.53
18:D:1208:C:H2'	18:D:1209:C:C6	2.40	0.53
20:F:31:GLU:O	20:F:34:ARG:HG2	2.08	0.53
41:a:67:U:N3	41:a:74:A:C6	2.77	0.53
41:a:858:G:OP2	42:b:78:LYS:NZ	2.35	0.53
41:a:1567:G:OP2	48:h:83:TYR:OH	2.19	0.53
41:a:2387:U:O4'	42:b:41:ARG:NH1	2.41	0.53
54:n:42:GLU:HG3	54:n:49:LEU:HD23	1.91	0.53
54:n:132:VAL:CG2	54:n:152:LEU:HD23	2.39	0.53
8:7:3:G:H3'	18:D:1500:A:N6	2.24	0.52
9:9:93:ALA:O	9:9:129:LEU:O	2.27	0.52
9:9:136:ILE:HG13	9:9:136:ILE:O	2.09	0.52
12:AB:135:GLY:C	12:AB:137:ALA:H	2.17	0.52
15:AF:3:ARG:NH1	15:AF:55:GLU:OE2	2.40	0.52
16:AG:362:TYR:CE1	16:AG:414:LEU:HD21	2.43	0.52
10:B:38:A:O2'	18:D:790:A:O5'	2.27	0.52
33:S:79:LEU:HB2	33:S:84:VAL:HG22	1.91	0.52
34:T:11:ILE:HA	34:T:14:GLU:OE1	2.09	0.52
38:X:68:ASP:O	38:X:71:ARG:HG3	2.09	0.52
39:Y:9:LYS:CG	39:Y:55:PRO:HB3	2.39	0.52
39:Y:16:MET:HG2	39:Y:16:MET:O	2.09	0.52
41:a:1295:C:C2	41:a:1296:G:C8	2.97	0.52
41:a:2649:C:H2'	41:a:2650:U:H6	1.73	0.52
41:a:2833:U:O2	50:j:58:ASN:ND2	2.41	0.52
53:m:26:ASN:O	53:m:30:VAL:HG23	2.08	0.52
61:u:54:GLN:NE2	61:u:60:ARG:HH12	2.07	0.52
62:v:115:GLU:O	62:v:119:LEU:HD23	2.09	0.52
1:0:4:VAL:HG22	1:0:40:MET:HB3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1:61:ASN:OD1	41:a:496:G:H1'	2.09	0.52
3:2:33:LYS:HG3	3:2:80:TRP:CE3	2.45	0.52
5:4:51:GLN:OE1	5:4:57:TYR:OH	2.27	0.52
9:9:87:GLU:OE2	9:9:95:LEU:HG	2.09	0.52
12:AB:27:ASN:ND2	12:AB:58:GLU:HB2	2.24	0.52
14:AE:1000:GLY:HA3	14:AE:1026:PRO:HG2	1.90	0.52
16:AG:363:LEU:HG	16:AG:410:ALA:N	2.16	0.52
17:C:60:LYS:HZ2	18:D:734:G:C2'	2.19	0.52
18:D:769:G:H4'	18:D:1513:A:H4'	1.90	0.52
18:D:1240:U:C5	27:M:116:MET:HE2	2.44	0.52
25:K:111:MET:HE2	25:K:125:ALA:HB1	1.90	0.52
25:K:134:ILE:HG23	25:K:135:ASN:N	2.24	0.52
26:L:36:ILE:O	26:L:36:ILE:HG23	2.09	0.52
28:N:66:PHE:CD2	28:N:67:GLN:OE1	2.62	0.52
29:O:118:LEU:HD23	29:O:124:ARG:HA	1.91	0.52
37:W:66:MET:HG2	37:W:74:PHE:CZ	2.44	0.52
38:X:31:LYS:HG3	38:X:32:ALA:N	2.25	0.52
39:Y:77:VAL:O	39:Y:77:VAL:HG12	2.08	0.52
39:Y:126:ARG:NH1	41:a:1083:U:P	2.83	0.52
41:a:271:G:C4	41:a:272:A:N7	2.77	0.52
41:a:1801:A:OP2	48:h:146:MET:HE1	2.09	0.52
41:a:1820:U:C4	48:h:159:GLY:HA3	2.44	0.52
41:a:1857:G:O2'	41:a:1884:G:N2	2.38	0.52
41:a:2649:C:C2	41:a:2650:U:C5	2.97	0.52
48:h:34:LEU:HD12	48:h:61:ALA:HB1	1.91	0.52
52:l:62:GLN:HA	52:l:69:ARG:HH21	1.74	0.52
56:p:32:GLU:C	56:p:33:LEU:HD12	2.34	0.52
58:r:6:LEU:CD1	58:r:37:VAL:HG23	2.39	0.52
58:r:33:GLN:HB3	58:r:35:LYS:HZ3	1.74	0.52
61:u:58:TYR:CE2	61:u:59:ARG:HG2	2.44	0.52
9:9:59:LEU:HB2	9:9:62:ARG:HG3	1.92	0.52
9:9:106:PHE:CG	9:9:106:PHE:O	2.63	0.52
13:AC:43:LEU:HD13	13:AC:217:ILE:HD11	1.90	0.52
16:AG:213:VAL:HG22	16:AG:213:VAL:O	2.08	0.52
21:G:58:ASN:OD1	21:G:223:GLU:OE1	2.26	0.52
22:H:332:VAL:HG12	22:H:334:ASP:H	1.73	0.52
23:I:58:GLU:OE2	23:I:60:PRO:HD3	2.09	0.52
23:I:85:GLU:O	23:I:89:LYS:HE2	2.08	0.52
26:L:70:VAL:O	26:L:74:LEU:CD2	2.56	0.52
31:Q:84:VAL:HG21	31:Q:97:ILE:CD1	2.38	0.52
38:X:96:PRO:CG	38:X:102:THR:HG22	2.40	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:590:A:H2'	41:a:591:U:C6	2.44	0.52
41:a:1394:U:H4'	41:a:1603:A:H4'	1.92	0.52
41:a:1607:C:N4	41:a:1622:G:OP2	2.34	0.52
41:a:1656:C:OP1	50:j:141:ARG:HD3	2.10	0.52
60:t:1:MET:CE	60:t:32:TYR:CB	2.87	0.52
7:6:19:DC:H2'	7:6:20:DA:H8	1.74	0.52
12:AB:154:VAL:HG11	12:AB:162:LEU:CD2	2.39	0.52
14:AE:77:ARG:HD3	14:AE:78:LEU:H	1.73	0.52
16:AG:63:LEU:HA	16:AG:92:TYR:HA	1.91	0.52
16:AG:393:LEU:CB	16:AG:398:LEU:HD22	1.90	0.52
16:AG:434:LEU:HD22	16:AG:459:LEU:CG	2.31	0.52
31:Q:23:ILE:N	31:Q:87:LYS:HZ1	2.08	0.52
31:Q:79:ILE:O	31:Q:105:PHE:HE1	1.91	0.52
31:Q:86:VAL:O	31:Q:113:VAL:HG22	2.08	0.52
33:S:87:ALA:N	33:S:90:ARG:NH1	2.58	0.52
41:a:69:C:O2	41:a:73:A:O2'	2.26	0.52
41:a:1145:C:N3	41:a:1146:C:C5	2.77	0.52
41:a:2093:G:OP2	58:r:22:LYS:HE3	2.09	0.52
51:k:9:ILE:CG2	51:k:27:LYS:HZ1	2.23	0.52
52:l:6:LYS:HZ2	52:l:119:ILE:HA	1.73	0.52
52:l:60:TRP:NE1	52:l:69:ARG:NH1	2.56	0.52
54:n:29:PRO:HB2	54:n:169:LEU:HD22	1.91	0.52
59:s:25:LEU:CD1	59:s:101:ILE:HD13	2.39	0.52
60:t:71:ARG:NH2	60:t:123:LEU:O	2.42	0.52
2:1:27:LYS:O	2:1:71:VAL:HG22	2.10	0.52
8:7:8:U:OP2	18:D:1397:C:C2	2.62	0.52
11:AA:246:LEU:HB3	11:AA:269:ILE:HD13	1.91	0.52
14:AE:342:LEU:HD23	14:AE:1352:ILE:HG23	1.90	0.52
16:AG:9:VAL:HG22	16:AG:24:PHE:CE2	2.44	0.52
18:D:512:U:P	24:J:44:ARG:NH1	2.83	0.52
18:D:1158:C:O2'	21:G:132:LYS:HE2	2.10	0.52
19:E:75:HIS:CE1	19:E:79:LEU:CD1	2.93	0.52
20:F:17:ARG:HA	20:F:20:LYS:NZ	2.24	0.52
21:G:168:HIS:HB3	21:G:169:GLU:OE1	2.10	0.52
23:I:105:GLU:OE1	23:I:107:ARG:NE	2.43	0.52
24:J:50:ASP:O	24:J:53:VAL:HG22	2.09	0.52
32:R:24:LEU:C	32:R:25:GLU:OE1	2.52	0.52
33:S:54:ASP:HA	33:S:59:ARG:HD2	1.90	0.52
38:X:13:LYS:HE3	38:X:17:ILE:HG21	1.91	0.52
39:Y:32:VAL:HA	39:Y:60:VAL:HG21	1.90	0.52
39:Y:58:ILE:HG23	39:Y:66:PHE:CD2	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:Z:18:ASP:HB3	40:Z:22:LEU:CD1	2.39	0.52
41:a:1167:C:C2	41:a:1168:G:C8	2.97	0.52
41:a:2355:G:H2'	41:a:2356:U:H5'	1.91	0.52
60:t:34:GLY:N	60:t:37:ASP:OD2	2.39	0.52
61:u:56:PRO:O	61:u:60:ARG:HG3	2.09	0.52
4:3:69:ASN:ND2	41:a:329:G:OP2	2.37	0.52
7:6:17:DC:H42	8:7:53:G:H1	1.58	0.52
13:AD:100:LEU:HD21	13:AD:121:VAL:HG11	1.90	0.52
14:AE:526:VAL:HG12	14:AE:549:LYS:HB2	1.90	0.52
14:AE:959:LYS:HB3	14:AE:983:LYS:HB2	1.92	0.52
10:B:48:C:C5	10:B:59:A:C8	2.98	0.52
18:D:1308:U:H3'	38:X:98:ARG:CZ	2.39	0.52
23:I:23:PHE:HD2	30:P:97:ASP:HB2	1.74	0.52
23:I:142:MET:SD	23:I:170:GLU:OE1	2.67	0.52
26:L:88:MET:HE2	26:L:90:MET:HE3	1.91	0.52
39:Y:77:VAL:HA	39:Y:80:LYS:HD3	1.91	0.52
39:Y:78:LEU:HD13	39:Y:108:ILE:HG12	1.90	0.52
39:Y:129:GLU:CB	39:Y:133:ARG:HH12	2.16	0.52
41:a:632:A:H2'	41:a:633:A:C8	2.44	0.52
41:a:833:A:H2'	41:a:834:G:C8	2.44	0.52
41:a:1138:G:C2	59:s:108:MET:HE2	2.45	0.52
41:a:1145:C:C2	41:a:1146:C:C5	2.97	0.52
58:r:143:ILE:O	58:r:143:ILE:HG13	2.08	0.52
63:w:14:SER:HA	63:w:17:ARG:CZ	2.39	0.52
63:w:37:THR:OG1	63:w:40:LYS:CD	2.57	0.52
6:5:9:DG:OP2	11:AA:470:ARG:O	2.28	0.52
18:D:537:G:OP1	32:R:110:ARG:NH2	2.38	0.52
18:D:1371:G:O3'	29:O:71:GLY:HA3	2.10	0.52
21:G:9:MET:HE1	21:G:47:VAL:CG2	2.40	0.52
23:I:49:LYS:NZ	23:I:75:ILE:HD11	2.24	0.52
25:K:137:VAL:O	25:K:141:ILE:HG12	2.09	0.52
28:N:79:SER:HB2	28:N:85:ILE:HG22	1.92	0.52
29:O:21:ILE:HD11	29:O:87:LEU:HD22	1.90	0.52
29:O:88:MET:HE1	29:O:95:ARG:HG3	1.91	0.52
34:T:24:SER:O	34:T:28:GLN:HG2	2.09	0.52
39:Y:60:VAL:CG2	39:Y:66:PHE:HB3	2.28	0.52
39:Y:126:ARG:NH1	41:a:1082:U:C5'	2.73	0.52
41:a:963:U:C2	41:a:964:C:C5	2.98	0.52
41:a:1204:A:O4'	41:a:1206:G:C8	2.63	0.52
41:a:2356:U:C5'	42:b:20:ARG:HD2	2.36	0.52
41:a:2473:U:C4	41:a:2474:U:C5	2.97	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2758:A:H1'	56:p:38:ASN:HD21	1.74	0.52
48:h:74:ILE:CG2	48:h:96:TYR:HD1	2.21	0.52
54:n:94:GLU:HA	54:n:97:TRP:CE3	2.43	0.52
56:p:149:ARG:CZ	56:p:154:PRO:HD3	2.40	0.52
62:v:21:ALA:HB2	62:v:100:LYS:HG3	1.90	0.52
9:9:88:HIS:O	9:9:89:PRO:C	2.51	0.52
12:AB:155:LYS:HG3	12:AB:156:ASN:N	2.25	0.52
16:AG:353:HIS:HA	16:AG:356:ILE:HG23	1.79	0.52
10:B:31:G:H2'	10:B:32:C:O4'	2.10	0.52
18:D:122:G:C5	18:D:123:U:C5	2.98	0.52
18:D:452:A:H2'	18:D:453:G:O4'	2.10	0.52
29:O:57:MET:HA	29:O:60:LYS:CE	2.40	0.52
29:O:123:ARG:O	29:O:123:ARG:CG	2.57	0.52
30:P:22:THR:O	30:P:25:ILE:HG22	2.10	0.52
34:T:63:ARG:HD2	34:T:67:LEU:CD2	2.40	0.52
38:X:71:ARG:HH12	47:g:52:ALA:HB3	1.75	0.52
39:Y:15:GLY:HA2	39:Y:50:LYS:CG	2.33	0.52
41:a:537:G:P	59:s:2:LYS:HE2	2.49	0.52
41:a:998:C:OP1	66:z:84:LYS:NZ	2.39	0.52
41:a:1340:U:C5	41:a:1603:A:C8	2.97	0.52
41:a:1666:G:P	60:t:66:LYS:HD3	2.50	0.52
49:i:26:THR:HG23	49:i:27:SER:N	2.25	0.52
51:k:29:THR:HG23	51:k:30:LYS:HG2	1.90	0.52
57:q:25:VAL:CB	57:q:35:GLN:OE1	2.56	0.52
58:r:29:PHE:CE2	58:r:35:LYS:HE3	2.45	0.52
58:r:71:LYS:HZ2	58:r:72:ILE:HA	1.73	0.52
59:s:9:GLU:HG2	59:s:10:THR:HG23	1.90	0.52
60:t:2:ILE:HG23	60:t:8:LEU:HD11	1.91	0.52
61:u:14:LYS:HE2	61:u:14:LYS:HA	1.91	0.52
4:3:6:ARG:HG3	41:a:84:A:H3'	1.91	0.52
9:9:61:ARG:NE	41:a:1046:A:P	2.83	0.52
11:AA:714:VAL:HB	11:AA:787:PRO:HD2	1.92	0.52
14:AE:510:LEU:HD22	14:AE:601:ILE:HD12	1.91	0.52
16:AG:244:ARG:CD	16:AG:244:ARG:N	2.73	0.52
21:G:26:LYS:HE3	21:G:194:ASP:OD1	2.10	0.52
21:G:49:MET:CE	21:G:201:PRO:CD	2.87	0.52
24:J:74:ASN:O	24:J:77:LYS:HG2	2.10	0.52
30:P:7:ARG:HD3	30:P:75:ASP:OD1	2.10	0.52
30:P:73:LEU:HD21	30:P:75:ASP:OD2	2.10	0.52
41:a:645:C:H2'	41:a:647:G:C8	2.45	0.52
41:a:888:C:C4	41:a:889:C:C4	2.97	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:g:47:LYS:HE2	54:n:115:ARG:NE	2.25	0.52
50:j:55:LYS:HZ2	50:j:60:VAL:HB	1.74	0.52
52:l:6:LYS:CE	52:l:119:ILE:HG23	2.40	0.52
59:s:31:GLU:HG2	59:s:142:ILE:HG12	1.92	0.52
66:z:108:ALA:O	66:z:111:GLU:HG3	2.09	0.52
10:A:56:C:C5	41:a:2168:G:C6	2.98	0.52
11:AA:255:ILE:HB	11:AA:263:VAL:HB	1.92	0.52
11:AA:847:PRO:N	16:AG:104:ARG:NH2	2.57	0.52
12:AB:138:ARG:HD3	12:AB:155:LYS:CB	2.39	0.52
14:AE:67:ASP:C	14:AE:69:GLU:H	2.18	0.52
16:AG:240:THR:HG21	16:AG:247:PRO:HG3	1.92	0.52
18:D:58:C:O2	18:D:58:C:H2'	2.09	0.52
18:D:564:C:P	32:R:12:ARG:HH22	2.30	0.52
18:D:842:U:H3'	18:D:843:U:C5'	2.39	0.52
20:F:37:PHE:HB3	31:Q:124:PRO:O	2.10	0.52
34:T:5:THR:HG23	34:T:6:GLU:N	2.25	0.52
34:T:15:PHE:CD2	34:T:30:ALA:CB	2.92	0.52
39:Y:116:MET:HG2	39:Y:124:MET:HB3	1.91	0.52
41:a:560:C:O2	66:z:48:ARG:NH1	2.37	0.52
41:a:607:U:C5	41:a:620:G:C5	2.97	0.52
42:b:68:LYS:HZ2	42:b:70:GLU:CD	2.16	0.52
4:3:66:GLN:OE1	4:3:69:ASN:ND2	2.43	0.51
8:7:13:U:H5''	8:7:13:U:C6	2.45	0.51
9:9:43:LYS:O	9:9:47:GLU:OE1	2.28	0.51
12:AB:81:PHE:CB	14:AE:141:PHE:O	2.58	0.51
12:AB:138:ARG:HD3	12:AB:155:LYS:HA	1.91	0.51
14:AE:795:TYR:OH	14:AE:1326:GLN:NE2	2.42	0.51
16:AG:139:THR:HA	16:AG:180:ARG:HA	1.92	0.51
16:AG:233:ARG:C	16:AG:327:LEU:HD21	2.34	0.51
16:AG:285:ILE:HD12	16:AG:293:VAL:HG21	1.91	0.51
16:AG:302:LYS:HZ1	16:AG:302:LYS:C	2.11	0.51
16:AG:399:ASP:OD2	16:AG:399:ASP:N	2.40	0.51
16:AG:422:GLU:CA	16:AG:426:GLY:N	2.62	0.51
18:D:1175:G:N3	18:D:1176:A:C8	2.77	0.51
21:G:23:TRP:HZ3	21:G:25:PRO:CA	2.21	0.51
22:H:48:ILE:HG22	22:H:49:PRO:O	2.10	0.51
25:K:156:LYS:NZ	28:N:71:VAL:O	2.42	0.51
30:P:66:GLU:CG	33:S:99:ALA:HB2	2.41	0.51
34:T:3:LEU:HD12	34:T:35:GLN:OE1	2.10	0.51
41:a:139:U:OP2	41:a:140:C:N4	2.43	0.51
41:a:1681:G:H2'	41:a:1757:A:N1	2.25	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2049:G:N2	50:j:161:MET:HE1	2.24	0.51
41:a:2093:G:P	58:r:22:LYS:HZ2	2.24	0.51
41:a:2329:U:H2'	41:a:2330:G:C8	2.45	0.51
41:a:2466:C:N3	41:a:2467:C:C5	2.78	0.51
44:d:15:A:O4'	44:d:109:A:C8	2.63	0.51
44:d:42:C:P	54:n:64:LYS:NZ	2.83	0.51
54:n:15:LYS:O	54:n:18:THR:N	2.43	0.51
54:n:71:ARG:C	54:n:81:GLN:HE22	2.17	0.51
1:0:71:LYS:HG3	1:0:73:LYS:HE3	1.92	0.51
5:4:2:PHE:CE2	5:4:55:GLU:HG2	2.45	0.51
9:9:47:GLU:O	9:9:49:GLY:N	2.43	0.51
11:AA:125:GLY:HA2	11:AA:499:SER:HB2	1.92	0.51
11:AA:846:GLY:C	16:AG:104:ARG:HH22	2.18	0.51
11:AA:1287:LEU:HD13	14:AE:1357:ILE:HD11	1.91	0.51
10:B:76:A:H1'	41:a:2494:G:P	2.50	0.51
17:C:70:TYR:CE2	26:L:49:TYR:CZ	2.98	0.51
18:D:932:C:C5'	27:M:4:ARG:HH11	2.22	0.51
18:D:936:C:C2	18:D:937:A:C8	2.98	0.51
18:D:1086:U:C2	18:D:1087:G:C8	2.97	0.51
21:G:90:PHE:CD2	21:G:154:MET:CE	2.93	0.51
22:H:32:ASP:OD1	22:H:35:VAL:CG2	2.58	0.51
23:I:178:LEU:HD23	23:I:178:LEU:C	2.35	0.51
30:P:24:GLU:O	30:P:28:THR:HG23	2.10	0.51
31:Q:61:PHE:CD1	31:Q:64:GLN:NE2	2.78	0.51
36:V:62:ARG:CA	36:V:73:TRP:CZ3	2.93	0.51
39:Y:55:PRO:C	39:Y:71:LYS:HE2	2.36	0.51
39:Y:78:LEU:HD12	39:Y:112:LYS:NZ	2.25	0.51
41:a:125:A:OP2	53:m:19:ARG:NH2	2.39	0.51
41:a:1154:G:OP2	66:z:58:ARG:CZ	2.58	0.51
41:a:1348:C:C4	41:a:1349:C:C6	2.97	0.51
41:a:2473:U:O2	41:a:2473:U:H2'	2.09	0.51
50:j:131:ASP:OD1	50:j:134:HIS:HB2	2.10	0.51
58:r:2:GLN:HG3	58:r:39:ALA:CB	2.40	0.51
62:v:53:MET:HE3	62:v:120:ALA:CB	2.40	0.51
66:z:74:ILE:HG13	66:z:78:LYS:CE	2.40	0.51
8:7:46:U:O2	8:7:46:U:H2'	2.11	0.51
14:AE:71:LEU:HD22	14:AE:90:VAL:HG21	1.93	0.51
14:AE:79:LYS:CG	16:AG:144:LYS:HD2	2.39	0.51
14:AE:122:SER:HB2	14:AE:132:LEU:HD12	1.91	0.51
16:AG:21:GLU:HB3	16:AG:49:ILE:HD12	1.92	0.51
16:AG:402:THR:HA	16:AG:405:ALA:HB3	1.91	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1176:A:H2'	18:D:1177:G:C8	2.45	0.51
22:H:279:LYS:HA	22:H:331:MET:CB	2.41	0.51
22:H:335:ILE:HG12	22:H:342:ILE:HD12	1.92	0.51
41:a:65:U:H2'	41:a:66:C:H6	1.75	0.51
41:a:617:G:C4'	52:l:199:MET:HE1	2.40	0.51
41:a:2391:G:P	55:o:32:ILE:HD12	2.50	0.51
48:h:52:ARG:NE	48:h:53:HIS:HE1	2.07	0.51
60:t:17:ARG:NH2	60:t:47:ILE:HG23	2.25	0.51
62:v:45:GLN:HE21	62:v:124:LEU:HA	1.75	0.51
66:z:86:ALA:C	66:z:87:SER:HG	2.17	0.51
11:AA:93:SER:HA	11:AA:128:PRO:HA	1.93	0.51
16:AG:362:TYR:CE2	16:AG:382:GLU:HG2	2.45	0.51
18:D:73:C:O2	18:D:73:C:C2'	2.57	0.51
18:D:718:A:H5'	31:Q:119:ASN:HD22	1.76	0.51
20:F:25:LYS:HE3	22:H:336:ASP:OD2	2.11	0.51
23:I:85:GLU:OE2	23:I:88:ARG:NH1	2.26	0.51
23:I:108:LYS:N	23:I:108:LYS:HD2	2.26	0.51
23:I:183:ASP:HB3	23:I:204:LYS:NZ	2.25	0.51
25:K:86:LYS:HZ2	25:K:95:PHE:HE1	1.56	0.51
27:M:66:LEU:CD1	27:M:101:MET:CE	2.84	0.51
30:P:15:HIS:HB3	30:P:70:HIS:CE1	2.46	0.51
39:Y:78:LEU:HD12	39:Y:112:LYS:HZ1	1.76	0.51
40:Z:26:MET:HA	40:Z:30:PHE:CD1	2.45	0.51
41:a:872:U:H2'	41:a:873:C:H6	1.75	0.51
41:a:1666:G:OP1	60:t:66:LYS:HD3	2.10	0.51
41:a:2420:C:H5''	51:k:8:LYS:HE2	1.92	0.51
41:a:2723:C:H4'	63:w:1:MET:HE3	1.92	0.51
41:a:2812:G:H2'	41:a:2813:A:C8	2.45	0.51
48:h:76:ALA:HB1	48:h:94:VAL:CG1	2.39	0.51
50:j:68:PHE:CD2	50:j:75:ALA:HA	2.46	0.51
52:l:168:ASP:HB3	52:l:183:PHE:HE2	1.75	0.51
55:o:31:HIS:O	55:o:32:ILE:C	2.54	0.51
62:v:112:LEU:HA	62:v:115:GLU:OE1	2.10	0.51
66:z:65:ILE:HB	66:z:76:TYR:HE1	1.75	0.51
10:A:6:G:C4	10:A:7:G:C8	2.98	0.51
10:A:33:U:H5''	27:M:84:THR:HG21	1.93	0.51
16:AG:444:LEU:HD22	16:AG:473:LEU:HD21	1.92	0.51
10:B:6:G:C4	10:B:7:G:C8	2.98	0.51
18:D:107:G:C6	19:E:10:ARG:NH2	2.79	0.51
18:D:564:C:H5'	36:V:34:TYR:HE1	1.76	0.51
18:D:767:A:H2'	18:D:768:A:O4'	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:868:C:H2'	18:D:869:G:O4'	2.11	0.51
21:G:17:GLY:HA3	21:G:41:ILE:HD11	1.92	0.51
22:H:36:VAL:O	22:H:48:ILE:HB	2.10	0.51
27:M:111:ARG:N	27:M:119:ARG:NH2	2.58	0.51
36:V:15:ASP:OD1	36:V:55:ILE:HD11	2.11	0.51
41:a:626:A:N3	61:u:78:ARG:CZ	2.74	0.51
41:a:960:A:H61	62:v:82:MET:CE	2.24	0.51
41:a:1095:A:H2'	41:a:1096:A:C8	2.45	0.51
41:a:1314:C:C2	41:a:1315:C:C5	2.98	0.51
41:a:2093:G:OP2	58:r:22:LYS:HG2	2.11	0.51
41:a:2250:G:OP1	41:a:2275:C:O2'	2.25	0.51
41:a:2428:G:H21	61:u:60:ARG:CZ	2.23	0.51
41:a:2814:A:H2'	41:a:2815:C:H6	1.75	0.51
51:k:21:TYR:CE2	51:k:49:TYR:CD2	2.98	0.51
59:s:15:TRP:CE3	59:s:135:GLN:O	2.63	0.51
63:w:58:ASP:CG	63:w:80:PHE:CZ	2.89	0.51
5:4:48:MET:CE	5:4:86:LEU:HG	2.40	0.51
10:A:31:G:H2'	10:A:32:C:O4'	2.10	0.51
12:AB:10:LYS:HB2	12:AB:73:ARG:HB3	1.90	0.51
12:AB:148:LYS:HE2	30:P:100:ILE:CD1	2.40	0.51
13:AD:205:MET:HE3	13:AD:213:PRO:HB3	1.91	0.51
16:AG:353:HIS:O	16:AG:356:ILE:CB	2.58	0.51
18:D:107:G:C5	19:E:10:ARG:NH2	2.79	0.51
18:D:978:A:C4	18:D:1319:A:C2	2.98	0.51
21:G:35:ARG:HB3	22:H:14:LYS:NZ	2.26	0.51
21:G:165:ASP:O	21:G:169:GLU:OE1	2.28	0.51
22:H:304:VAL:HG11	22:H:346:LEU:HB2	1.93	0.51
26:L:42:TRP:CZ2	26:L:102:MET:SD	3.01	0.51
35:U:39:PHE:CD1	35:U:50:THR:HG22	2.45	0.51
37:W:41:PHE:O	37:W:44:MET:HG3	2.11	0.51
39:Y:86:LYS:HD3	39:Y:86:LYS:O	2.11	0.51
41:a:265:A:N1	41:a:427:U:O2'	2.40	0.51
41:a:910:A:C8	62:v:12:MET:CE	2.93	0.51
41:a:1198:U:C2	41:a:1199:U:C5	2.98	0.51
41:a:1598:A:C5	41:a:1599:U:C5	2.99	0.51
41:a:1956:U:C4	41:a:1957:C:C5	2.99	0.51
48:h:108:LYS:HD3	48:h:194:GLU:HB2	1.92	0.51
51:k:19:HIS:CE1	51:k:41:PRO:HD3	2.45	0.51
52:l:1:MET:CE	52:l:16:GLU:HA	2.26	0.51
60:t:103:VAL:HG21	60:t:107:LEU:CD2	2.41	0.51
62:v:115:GLU:HA	62:v:118:LYS:NZ	2.25	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:17:GLY:HA2	1:0:97:LYS:HE2	1.92	0.51
11:AA:858:GLY:HA2	16:AG:38:LYS:HB2	1.78	0.51
12:AB:78:PHE:CG	12:AB:85:PRO:HB3	2.46	0.51
13:AC:140:ILE:HD12	13:AC:142:MET:HE3	1.91	0.51
16:AG:329:SER:HA	16:AG:336:LEU:HG	1.92	0.51
18:D:501:C:H2'	18:D:502:A:C8	2.46	0.51
18:D:728:A:H2'	18:D:729:A:C8	2.46	0.51
18:D:754:C:OP1	34:T:72:ARG:NH1	2.44	0.51
18:D:925:G:C2	18:D:927:G:C8	2.98	0.51
18:D:934:C:C4	18:D:1345:U:C5	2.99	0.51
18:D:1226:C:C2'	38:X:102:THR:OG1	2.58	0.51
20:F:36:GLU:OE2	20:F:36:GLU:HA	2.11	0.51
23:I:142:MET:CE	23:I:170:GLU:OE1	2.58	0.51
25:K:36:LEU:HD13	25:K:134:ILE:CD1	2.41	0.51
29:O:9:THR:HB	29:O:85:ARG:NH1	2.25	0.51
29:O:30:ILE:HD13	29:O:39:PHE:HE2	1.75	0.51
31:Q:52:PHE:CZ	31:Q:65:VAL:HG21	2.45	0.51
34:T:11:ILE:HD11	34:T:30:ALA:C	2.36	0.51
41:a:851:C:H2'	41:a:852:U:C6	2.46	0.51
41:a:1318:U:C2	41:a:1319:C:C5	2.99	0.51
41:a:2311:A:C5	54:n:77:PHE:CD2	2.98	0.51
41:a:2483:C:N3	62:v:123:LYS:NZ	2.54	0.51
41:a:2588:G:O6	41:a:2607:G:C6	2.64	0.51
41:a:2698:U:H2'	41:a:2699:C:C6	2.46	0.51
41:a:2813:A:C4	41:a:2814:A:C8	2.99	0.51
48:h:124:ILE:C	48:h:125:LYS:HD3	2.36	0.51
48:h:130:LEU:HD21	48:h:192:LEU:HD21	1.93	0.51
52:l:119:ILE:O	52:l:188:MET:N	2.35	0.51
59:s:15:TRP:CZ3	59:s:135:GLN:CA	2.93	0.51
8:7:2:U:H2'	8:7:3:G:C8	2.45	0.51
9:9:10:ALA:O	9:9:14:GLU:HG3	2.11	0.51
9:9:78:GLY:H	9:9:79:PRO:HD2	1.75	0.51
10:A:48:C:C5	10:A:59:A:C8	2.98	0.51
11:AA:524:ILE:HG21	11:AA:708:VAL:HG13	1.92	0.51
16:AG:285:ILE:CD1	16:AG:285:ILE:N	2.73	0.51
17:C:73:ARG:HH12	31:Q:113:VAL:HB	1.76	0.51
18:D:500:G:H2'	18:D:501:C:C6	2.46	0.51
18:D:512:U:P	24:J:44:ARG:HH12	2.34	0.51
18:D:564:C:H5'	36:V:34:TYR:CE1	2.46	0.51
21:G:16:PHE:HB3	22:H:43:LYS:HA	1.93	0.51
21:G:187:VAL:HG13	21:G:191:SER:HB2	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:342:ILE:HG12	22:H:344:LEU:HD12	1.93	0.51
29:O:88:MET:SD	29:O:95:ARG:CZ	2.99	0.51
30:P:7:ARG:HD3	30:P:75:ASP:CG	2.36	0.51
30:P:24:GLU:CA	30:P:27:GLU:OE1	2.59	0.51
41:a:414:C:H2'	41:a:415:A:H8	1.75	0.51
41:a:1824:G:O2'	48:h:252:THR:HG21	2.11	0.51
41:a:1853:A:H2'	41:a:1854:A:C8	2.46	0.51
41:a:2006:C:C2	41:a:2007:U:C5	2.99	0.51
41:a:2133:G:O2'	41:a:2157:G:N2	2.44	0.51
41:a:2193:G:O2'	41:a:2194:U:OP1	2.28	0.51
41:a:2387:U:C4'	42:b:41:ARG:HH12	2.24	0.51
41:a:2727:A:O2'	60:t:70:ARG:NH2	2.43	0.51
48:h:124:ILE:O	48:h:125:LYS:HD3	2.10	0.51
50:j:122:VAL:HG21	50:j:141:ARG:HG3	1.92	0.51
59:s:98:GLU:O	59:s:99:ARG:C	2.53	0.51
1:0:37:GLU:O	1:0:37:GLU:HG3	2.11	0.51
8:7:12:U:H5''	25:K:56:VAL:HG21	1.93	0.51
10:A:23:C:H2'	10:A:24:U:C6	2.46	0.51
10:A:48:C:O2'	10:A:49:G:OP2	2.27	0.51
12:AB:38:ILE:CD1	12:AB:160:ARG:HH11	2.21	0.51
12:AB:118:ILE:CD1	12:AB:142:LEU:CD2	2.89	0.51
16:AG:365:ILE:HD11	16:AG:406:LEU:HD22	1.87	0.51
16:AG:453:VAL:HG11	16:AG:459:LEU:CD2	2.41	0.51
10:B:23:C:H2'	10:B:24:U:C6	2.45	0.51
17:C:60:LYS:CE	18:D:735:C:C4'	2.88	0.51
18:D:49:U:C2	18:D:361:G:N2	2.79	0.51
18:D:335:C:C2	18:D:336:A:C8	2.99	0.51
18:D:501:C:H2'	18:D:502:A:H8	1.75	0.51
18:D:975:A:H8	18:D:1357:A:HO2'	1.57	0.51
21:G:109:GLN:C	21:G:113:ARG:HE	2.19	0.51
21:G:127:ASP:O	21:G:128:LYS:HB3	2.09	0.51
21:G:169:GLU:OE1	21:G:169:GLU:N	2.44	0.51
25:K:111:MET:O	25:K:114:VAL:HG12	2.11	0.51
25:K:158:GLY:C	25:K:159:LYS:HD2	2.34	0.51
26:L:4:TYR:OH	26:L:68:GLN:HG2	2.11	0.51
26:L:11:HIS:HB3	26:L:14:GLN:OE1	2.11	0.51
41:a:247:G:O6	55:o:12:LYS:NZ	2.39	0.51
41:a:1346:G:C6	41:a:1601:G:C6	2.98	0.51
41:a:1454:C:H4'	63:w:63:ARG:NH1	2.26	0.51
41:a:1791:A:C2	41:a:1829:A:C4'	2.94	0.51
41:a:1996:C:H5''	50:j:128:ARG:NH1	2.25	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:c:71:LEU:HB3	43:c:76:GLU:CG	2.41	0.51
59:s:84:ILE:HG23	59:s:84:ILE:O	2.11	0.51
63:w:82:GLU:OE1	63:w:82:GLU:N	2.44	0.51
63:w:103:ARG:HA	63:w:110:MET:HE2	1.93	0.51
11:AA:232:ILE:HG12	11:AA:237:LEU:HG	1.93	0.51
12:AB:9:CYS:N	12:AB:53:ASN:HB3	2.24	0.51
16:AG:442:ARG:C	16:AG:446:PHE:HD2	2.15	0.51
16:AG:451:ARG:NH2	16:AG:470:ILE:CG1	2.67	0.51
18:D:21:G:C2	18:D:22:G:C5	2.99	0.51
25:K:156:LYS:HE3	28:N:71:VAL:CG1	2.41	0.51
26:L:33:GLU:HG3	26:L:33:GLU:O	2.10	0.51
32:R:54:ARG:HH11	32:R:62:GLU:HG3	1.76	0.51
34:T:3:LEU:HD23	34:T:3:LEU:C	2.36	0.51
41:a:271:G:H4'	41:a:272:A:OP1	2.10	0.51
41:a:624:C:O2'	41:a:657:U:OP1	2.29	0.51
41:a:672:C:O2'	52:l:77:ILE:HD11	2.11	0.51
41:a:813:U:H2'	41:a:814:C:C6	2.46	0.51
41:a:2063:C:C2	41:a:2064:C:C6	2.99	0.51
41:a:2086:U:H2'	41:a:2087:G:C8	2.46	0.51
41:a:2252:G:H2'	41:a:2253:G:H8	1.76	0.51
41:a:2315:G:O2'	54:n:125:ARG:HD3	2.09	0.51
42:b:55:ARG:HA	42:b:55:ARG:CZ	2.41	0.51
49:i:30:VAL:HG22	49:i:37:LYS:CD	2.41	0.51
63:w:58:ASP:OD1	63:w:63:ARG:NH2	2.44	0.51
66:z:65:ILE:HB	66:z:76:TYR:CE1	2.45	0.51
2:1:36:LEU:HD23	2:1:48:LYS:HB2	1.94	0.50
5:4:63:ILE:CD1	5:4:91:PHE:CD2	2.93	0.50
6:5:9:DG:P	11:AA:473:ARG:HG3	2.51	0.50
10:A:19:G:N2	41:a:2112:G:O4'	2.44	0.50
11:AA:400:VAL:HG11	11:AA:452:ARG:HD2	1.94	0.50
11:AA:853:ASP:O	16:AG:103:ASP:OD2	2.29	0.50
13:AD:182:ARG:NH1	14:AE:534:GLU:OE1	2.44	0.50
13:AD:288:GLU:C	13:AD:290:LEU:N	2.69	0.50
16:AG:233:ARG:NH1	16:AG:324:ASN:HB2	2.25	0.50
17:C:25:ASP:O	17:C:29:LEU:HD13	2.11	0.50
21:G:23:TRP:CZ3	21:G:25:PRO:HA	2.47	0.50
22:H:273:ARG:NH1	22:H:297:GLU:HB2	2.25	0.50
27:M:71:PRO:HG3	27:M:103:TRP:HH2	1.76	0.50
27:M:132:GLY:O	27:M:136:LYS:NZ	2.44	0.50
34:T:64:ARG:NH2	34:T:89:ARG:HH22	2.07	0.50
39:Y:21:PRO:CB	39:Y:22:PRO:HD3	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:413:C:C2	41:a:414:C:C5	2.99	0.50
41:a:703:U:H2'	41:a:704:G:O4'	2.10	0.50
41:a:1453:A:C8	63:w:73:ASN:OD1	2.64	0.50
41:a:1996:C:H4'	50:j:128:ARG:HH12	1.75	0.50
45:e:20:ASN:HA	45:e:23:ARG:HE	1.76	0.50
47:g:58:ASP:CG	47:g:62:LYS:NZ	2.68	0.50
48:h:18:LYS:HE3	48:h:20:VAL:CG2	2.41	0.50
55:o:25:LYS:HB3	61:u:62:PRO:CG	2.40	0.50
59:s:15:TRP:CD1	59:s:53:TYR:HB2	2.46	0.50
62:v:31:PHE:O	62:v:104:GLU:CD	2.54	0.50
63:w:73:ASN:HD22	63:w:73:ASN:C	2.17	0.50
64:x:53:THR:HG23	64:x:74:VAL:HG21	1.92	0.50
65:y:9:GLU:CG	65:y:55:LEU:HB2	2.41	0.50
2:1:25:ARG:NH1	41:a:519:U:O3'	2.43	0.50
4:3:85:PHE:CE1	4:3:94:ARG:HG2	2.46	0.50
11:AA:481:LEU:HD23	12:AB:16:ARG:HD3	1.79	0.50
11:AA:856:ASN:O	16:AG:37:LYS:HB2	2.11	0.50
12:AB:158:GLU:HB3	12:AB:162:LEU:CD2	2.41	0.50
14:AE:430:HIS:HB3	14:AE:925:GLU:HG3	1.93	0.50
17:C:41:PRO:HD2	22:H:339:ARG:NE	2.26	0.50
18:D:653:U:O5'	28:N:56:LYS:HE2	2.11	0.50
26:L:9:MET:O	26:L:84:VAL:HA	2.12	0.50
26:L:67:PRO:O	26:L:70:VAL:HG22	2.11	0.50
29:O:51:PRO:HA	29:O:103:PHE:HE2	1.76	0.50
30:P:37:ARG:HB3	30:P:75:ASP:HB2	1.94	0.50
33:S:10:GLU:OE2	33:S:63:ARG:HD2	2.11	0.50
39:Y:126:ARG:HD2	39:Y:126:ARG:C	2.36	0.50
39:Y:139:VAL:HG22	39:Y:140:GLU:N	2.25	0.50
41:a:628:G:C5	41:a:636:G:N2	2.79	0.50
41:a:1945:G:C4	41:a:1946:U:C5	2.99	0.50
41:a:2259:U:C6	41:a:2427:C:C4	2.99	0.50
51:k:7:GLU:CG	51:k:27:LYS:HZ3	2.25	0.50
54:n:5:HIS:HD2	54:n:97:TRP:CZ2	2.29	0.50
59:s:30:THR:HG23	59:s:31:GLU:N	2.26	0.50
60:t:38:ILE:HD11	60:t:112:PHE:CZ	2.43	0.50
60:t:114:LYS:O	60:t:118:LEU:HD23	2.12	0.50
62:v:39:GLY:O	62:v:96:ILE:N	2.33	0.50
62:v:105:MET:SD	62:v:113:ALA:HB1	2.51	0.50
62:v:105:MET:HE2	62:v:108:VAL:CG2	2.40	0.50
63:w:12:ARG:NE	63:w:20:MET:HE1	2.26	0.50
64:x:4:LYS:N	64:x:4:LYS:HD3	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:55:VAL:CG1	9:9:56:ARG:N	2.74	0.50
11:AA:974:ARG:HD2	11:AA:1014:LEU:HD21	1.92	0.50
12:AB:21:LEU:HB3	12:AB:26:VAL:CG1	2.38	0.50
16:AG:12:VAL:HG23	16:AG:16:LYS:HE2	1.92	0.50
16:AG:65:VAL:HG12	16:AG:66:ASP:H	1.76	0.50
16:AG:123:ARG:NH1	16:AG:123:ARG:HG2	2.26	0.50
10:B:68:C:C3'	10:B:69:C:H5''	2.42	0.50
17:C:51:TYR:O	17:C:54:GLN:HG3	2.11	0.50
18:D:739:C:O2	34:T:42:HIS:HE1	1.94	0.50
18:D:1145:A:HO2'	18:D:1146:A:P	2.35	0.50
21:G:26:LYS:HD2	21:G:193:PRO:CG	2.39	0.50
22:H:163:ARG:CA	22:H:298:GLU:HG3	2.41	0.50
25:K:62:LYS:HA	25:K:62:LYS:HE3	1.93	0.50
25:K:88:VAL:HG12	25:K:93:ARG:HA	1.94	0.50
32:R:3:THR:O	32:R:6:GLN:N	2.44	0.50
35:U:73:ALA:HA	35:U:76:LYS:CE	2.41	0.50
38:X:71:ARG:HD2	38:X:72:GLU:N	2.27	0.50
41:a:4:U:C2	41:a:5:A:C8	3.00	0.50
41:a:580:U:H2'	41:a:581:C:C6	2.46	0.50
41:a:1790:C:C3'	41:a:1791:A:C8	2.94	0.50
48:h:131:PRO:HD3	48:h:189:ARG:NH1	2.27	0.50
52:l:170:ARG:CZ	52:l:176:ASP:OD1	2.58	0.50
56:p:24:ILE:HG22	56:p:35:ARG:O	2.11	0.50
59:s:35:ARG:HA	59:s:40:HIS:CE1	2.46	0.50
60:t:59:LYS:HZ3	60:t:92:GLU:HG2	1.75	0.50
66:z:82:GLY:HA3	66:z:117:LEU:CD1	2.41	0.50
3:2:10:VAL:HG12	3:2:11:LEU:HD12	1.93	0.50
6:5:10:DT:N1	11:AA:62:TYR:HE2	2.07	0.50
8:7:3:G:C4'	18:D:1500:A:N6	2.74	0.50
11:AA:400:VAL:HG21	11:AA:452:ARG:HE	1.77	0.50
11:AA:562:GLU:OE1	11:AA:662:SER:OG	2.28	0.50
14:AE:412:LEU:HD22	14:AE:441:LEU:HD21	1.94	0.50
10:B:56:C:H2'	10:B:57:A:H5''	1.93	0.50
18:D:311:C:O5'	18:D:311:C:H6	1.94	0.50
18:D:580:C:H2'	18:D:581:G:O4'	2.11	0.50
18:D:958:A:O4'	37:W:55:ARG:NH1	2.45	0.50
18:D:1073:U:O2'	21:G:105:LYS:HE3	2.11	0.50
22:H:303:LEU:O	22:H:303:LEU:CG	2.60	0.50
23:I:142:MET:HE3	23:I:170:GLU:C	2.36	0.50
26:L:99:ALA:HB3	26:L:104:LYS:HZ1	1.76	0.50
31:Q:63:ALA:HB1	31:Q:96:THR:CG2	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:65:VAL:HG23	38:X:66:GLU:OE1	2.12	0.50
41:a:413:C:H2'	41:a:414:C:C6	2.46	0.50
41:a:1665:A:O3'	60:t:66:LYS:NZ	2.44	0.50
41:a:2266:A:H4'	41:a:2267:A:N3	2.26	0.50
41:a:2489:U:H2'	41:a:2490:G:O4'	2.12	0.50
41:a:2682:A:C2	50:j:23:PRO:HB3	2.46	0.50
47:g:16:CYS:CB	47:g:37:CYS:CB	2.90	0.50
54:n:30:ARG:HB2	54:n:30:ARG:NH2	2.25	0.50
54:n:136:ILE:HD12	54:n:141:ILE:HG23	1.92	0.50
55:o:19:LYS:HE2	55:o:19:LYS:HA	1.91	0.50
61:u:58:TYR:CD2	61:u:59:ARG:HG2	2.46	0.50
5:4:2:PHE:CE1	5:4:56:PHE:CE1	2.99	0.50
10:A:15:G:H2'	10:A:15:G:N3	2.27	0.50
14:AE:417:ARG:NH1	15:AF:43:ASN:O	2.45	0.50
16:AG:384:LEU:O	16:AG:387:VAL:HG23	2.11	0.50
17:C:60:LYS:NZ	18:D:735:C:C5'	2.75	0.50
18:D:36:C:C4	18:D:37:U:C5	3.00	0.50
18:D:280:C:H1'	36:V:40:ARG:HH22	1.75	0.50
34:T:64:ARG:HA	34:T:64:ARG:HH11	1.75	0.50
41:a:65:U:C2	41:a:66:C:C6	3.00	0.50
41:a:125:A:C5	53:m:10:LEU:HD13	2.46	0.50
41:a:993:G:C6	41:a:1162:G:C6	2.99	0.50
41:a:1999:C:O3'	41:a:2722:G:N2	2.44	0.50
46:f:4:THR:CB	46:f:37:GLU:OE2	2.58	0.50
52:l:170:ARG:HH12	52:l:176:ASP:N	2.09	0.50
54:n:8:TYR:HE2	54:n:30:ARG:HE	1.59	0.50
54:n:122:PHE:CE2	54:n:128:TYR:CD1	3.00	0.50
56:p:127:THR:HG22	56:p:128:GLN:N	2.26	0.50
59:s:98:GLU:HB3	59:s:102:GLU:OE2	2.12	0.50
3:2:3:ARG:NH2	3:2:4:GLU:HB2	2.26	0.50
5:4:6:ALA:HB3	5:4:65:VAL:HG22	1.93	0.50
5:4:50:MET:CA	5:4:56:PHE:CZ	2.94	0.50
9:9:55:VAL:HG12	9:9:56:ARG:N	2.25	0.50
11:AA:146:VAL:HG21	11:AA:513:GLN:HE21	1.77	0.50
12:AB:148:LYS:HD2	30:P:100:ILE:HG21	1.94	0.50
13:AD:285:THR:HA	13:AD:315:GLY:HA2	1.93	0.50
14:AE:77:ARG:HB3	14:AE:80:HIS:HB2	1.93	0.50
14:AE:157:GLN:HE22	14:AE:188:LEU:HD22	1.76	0.50
16:AG:5:ILE:HA	16:AG:8:VAL:HB	1.93	0.50
25:K:113:ALA:CA	25:K:116:GLU:OE1	2.60	0.50
27:M:35:LYS:HE3	27:M:38:THR:OG1	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:P:26:VAL:HG12	30:P:30:LYS:HZ2	1.76	0.50
33:S:87:ALA:HA	33:S:90:ARG:HG2	1.94	0.50
34:T:3:LEU:HD22	34:T:8:THR:HG23	1.92	0.50
39:Y:37:PHE:HE1	39:Y:58:ILE:HD11	1.77	0.50
40:Z:15:SER:O	40:Z:19:VAL:HG23	2.11	0.50
41:a:532:A:H4'	41:a:533:G:C8	2.46	0.50
41:a:1801:A:OP1	48:h:146:MET:HE1	2.11	0.50
41:a:2063:C:O2	41:a:2450:A:N1	2.45	0.50
41:a:2063:C:C4	41:a:2064:C:C5	3.00	0.50
41:a:2849:U:OP2	65:y:93:ARG:CZ	2.60	0.50
41:a:2884:U:H3	49:i:40:ARG:NH1	2.09	0.50
59:s:15:TRP:CH2	59:s:135:GLN:HG3	2.46	0.50
62:v:73:ILE:HD12	62:v:73:ILE:N	2.26	0.50
3:2:49:LYS:HE2	3:2:49:LYS:HA	1.93	0.50
10:A:68:C:C3'	10:A:69:C:H5''	2.42	0.50
12:AB:27:ASN:OD1	12:AB:58:GLU:HB2	2.11	0.50
12:AB:119:THR:HB	12:AB:123:PHE:CE2	2.46	0.50
16:AG:11:ALA:HB3	16:AG:12:VAL:HA	1.92	0.50
16:AG:216:ILE:HG12	16:AG:221:ILE:HB	1.93	0.50
16:AG:305:MET:HG2	16:AG:336:LEU:HA	1.94	0.50
16:AG:440:VAL:CG2	16:AG:481:LEU:HD21	2.42	0.50
18:D:10:A:OP2	25:K:131:THR:HG21	2.12	0.50
18:D:821:G:H2'	18:D:822:U:C6	2.46	0.50
18:D:864:A:H2'	18:D:865:A:C8	2.46	0.50
18:D:877:G:H5''	28:N:80:ARG:HD2	1.93	0.50
22:H:267:TRP:CE3	22:H:340:ARG:HG3	2.47	0.50
24:J:104:ARG:NE	24:J:104:ARG:HA	2.27	0.50
41:a:234:U:C2	41:a:235:U:C6	3.00	0.50
41:a:372:G:C8	43:c:61:LYS:HD2	2.46	0.50
41:a:594:U:H2'	41:a:595:C:C6	2.47	0.50
41:a:780:G:C2	41:a:782:A:C2	3.00	0.50
41:a:1707:G:C8	41:a:1756:G:C5	2.98	0.50
41:a:1799:G:N2	48:h:154:LEU:HD23	2.26	0.50
41:a:2290:G:H2'	41:a:2291:U:C6	2.47	0.50
41:a:2902:C:N4	41:a:2903:U:O4	2.45	0.50
46:f:41:THR:N	46:f:45:ARG:HH22	2.09	0.50
50:j:46:ARG:NH1	50:j:85:ALA:O	2.32	0.50
50:j:154:LYS:HE3	50:j:156:PHE:CE1	2.46	0.50
51:k:21:TYR:CE2	51:k:49:TYR:CE2	2.99	0.50
52:l:155:GLU:HG2	52:l:156:ASN:N	2.27	0.50
60:t:121:GLU:HG3	60:t:123:LEU:HD12	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:48:MET:HE2	5:4:86:LEU:HG	1.93	0.50
11:AA:18:ARG:HE	11:AA:620:ASN:HA	1.77	0.50
12:AB:6:LEU:CD1	14:AE:291:ILE:CD1	2.84	0.50
12:AB:19:GLU:HG2	12:AB:23:ARG:CZ	2.42	0.50
16:AG:433:ASP:O	16:AG:456:LEU:HD21	2.12	0.50
18:D:339:C:C2	18:D:340:U:C5	2.99	0.50
18:D:552:U:O2'	32:R:83:ARG:O	2.19	0.50
18:D:674:G:P	26:L:86:ARG:HH22	2.35	0.50
18:D:744:C:H2'	18:D:745:G:C8	2.47	0.50
20:F:6:VAL:HG12	31:Q:108:THR:O	2.11	0.50
20:F:25:LYS:NZ	22:H:336:ASP:CG	2.68	0.50
21:G:38:VAL:HG22	22:H:75:VAL:HG21	1.92	0.50
23:I:132:ARG:HG3	23:I:135:LYS:HE3	1.92	0.50
27:M:108:ALA:O	27:M:119:ARG:NH2	2.45	0.50
32:R:110:ARG:HE	32:R:117:TYR:HD2	1.57	0.50
41:a:1791:A:C2	41:a:1829:A:O4'	2.65	0.50
41:a:1800:C:HO2'	41:a:1818:U:H3	1.60	0.50
41:a:2159:G:H2'	41:a:2160:C:C6	2.47	0.50
41:a:2305:U:H4'	54:n:133:ARG:NH2	2.26	0.50
41:a:2314:A:C1'	54:n:155:THR:HG21	2.42	0.50
41:a:2682:A:C5'	50:j:11:MET:HE1	2.42	0.50
53:m:25:LYS:C	53:m:29:GLN:HE22	2.20	0.50
62:v:41:LEU:HD22	62:v:45:GLN:NE2	2.27	0.50
65:y:106:LYS:HD3	65:y:109:ARG:NH2	2.26	0.50
2:1:35:ILE:O	2:1:39:THR:HG23	2.12	0.50
5:4:2:PHE:HB2	5:4:61:LEU:HG	1.94	0.50
5:4:62:THR:OG1	5:4:71:LYS:HE3	2.11	0.50
11:AA:529:ARG:HH11	11:AA:572:ILE:HG22	1.77	0.50
12:AB:50:LEU:HD21	14:AE:285:LEU:HB3	1.94	0.50
14:AE:59:ALA:HB3	14:AE:71:LEU:CD2	2.42	0.50
16:AG:248:VAL:HG21	16:AG:275:LEU:HG	1.94	0.50
16:AG:422:GLU:O	16:AG:427:ASP:N	2.45	0.50
18:D:610:U:O2	18:D:610:U:H2'	2.12	0.50
18:D:1106:G:O3'	23:I:172:ARG:HG3	2.11	0.50
21:G:23:TRP:CD1	21:G:39:HIS:CE1	2.97	0.50
23:I:68:ILE:HG13	23:I:101:ILE:HD11	1.93	0.50
24:J:76:TYR:CE2	24:J:204:TYR:CB	2.95	0.50
32:R:62:GLU:OE2	32:R:62:GLU:HA	2.10	0.50
40:Z:18:ASP:HB3	40:Z:22:LEU:HD12	1.93	0.50
41:a:228:C:N4	41:a:2407:A:N3	2.52	0.50
41:a:658:U:O2'	52:l:97:ASN:OD1	2.28	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1069:A:O2'	41:a:1073:A:N6	2.38	0.50
41:a:1665:A:C3'	60:t:66:LYS:HZ2	2.24	0.50
41:a:2179:C:C2	41:a:2180:U:C5	3.00	0.50
41:a:2627:G:O2'	41:a:2781:A:N1	2.43	0.50
41:a:2740:A:H2'	41:a:2741:A:C8	2.47	0.50
48:h:261:LYS:HA	48:h:264:ASP:OD2	2.12	0.50
50:j:152:PRO:CG	50:j:156:PHE:CZ	2.91	0.50
57:q:9:LYS:HZ3	57:q:11:CYS:C	2.20	0.50
59:s:131:ASN:OD1	59:s:131:ASN:O	2.30	0.50
61:u:55:MET:HE1	61:u:60:ARG:HA	1.94	0.50
4:3:6:ARG:HG2	4:3:7:ARG:H	1.77	0.49
6:5:9:DG:H5''	11:AA:473:ARG:HB3	0.54	0.49
12:AB:29:LEU:HB3	12:AB:56:PHE:HD2	1.77	0.49
14:AE:693:VAL:HG21	14:AE:743:MET:HE3	1.93	0.49
16:AG:171:GLU:OE2	16:AG:267:GLY:C	2.54	0.49
10:B:47:U:O2'	10:B:50:U:P	2.70	0.49
17:C:12:ARG:HG3	22:H:264:GLU:CB	2.41	0.49
17:C:66:SER:HA	26:L:49:TYR:CE1	2.47	0.49
18:D:136:C:C2	18:D:137:U:C6	3.00	0.49
18:D:376:G:OP2	35:U:70:ARG:HD3	2.12	0.49
18:D:653:U:O4'	28:N:56:LYS:CE	2.60	0.49
18:D:678:U:H2'	18:D:679:C:H6	1.76	0.49
24:J:170:TRP:CD2	24:J:186:PRO:HB3	2.47	0.49
31:Q:27:PHE:CE2	31:Q:89:PRO:HG2	2.46	0.49
37:W:44:MET:SD	37:W:62:VAL:CG1	2.99	0.49
41:a:16:C:O4'	49:i:15:MET:HE3	2.12	0.49
41:a:537:G:P	59:s:2:LYS:CE	3.00	0.49
41:a:1205:A:C2	52:l:165:HIS:CE1	3.00	0.49
41:a:2230:G:C5'	43:c:30:LEU:HD23	2.42	0.49
47:g:7:PRO:HD2	54:n:62:GLY:O	2.11	0.49
56:p:137:ASP:OD2	56:p:140:VAL:HG23	2.12	0.49
58:r:90:LEU:HD21	58:r:94:ILE:CD1	2.42	0.49
59:s:74:TYR:N	59:s:74:TYR:HD2	2.10	0.49
63:w:82:GLU:O	63:w:86:ARG:HG2	2.13	0.49
9:9:87:GLU:OE2	9:9:95:LEU:HB2	2.12	0.49
11:AA:856:ASN:CA	16:AG:35:THR:CA	2.81	0.49
14:AE:429:LEU:HB3	14:AE:925:GLU:HG2	1.94	0.49
14:AE:1179:PRO:HB2	14:AE:1182:GLY:H	1.75	0.49
16:AG:183:LEU:HD23	16:AG:197:VAL:HG22	1.94	0.49
16:AG:429:LYS:HD2	16:AG:429:LYS:N	2.24	0.49
17:C:60:LYS:HE3	18:D:735:C:C4'	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:71:THR:H	17:C:74:HIS:HE1	1.58	0.49
18:D:160:A:H2'	18:D:161:A:O4'	2.12	0.49
18:D:276:G:C5'	36:V:17:MET:CE	2.89	0.49
18:D:404:G:O2'	18:D:498:A:N1	2.38	0.49
18:D:1492:A:O2'	18:D:1493:A:H5''	2.11	0.49
23:I:82:GLU:O	23:I:86:LYS:CD	2.60	0.49
25:K:157:ARG:CZ	28:N:43:GLU:HG3	2.43	0.49
32:R:7:LEU:HD23	36:V:34:TYR:OH	2.12	0.49
37:W:49:ILE:HG23	37:W:49:ILE:O	2.12	0.49
38:X:3:ARG:HH22	47:g:36:VAL:HG12	1.77	0.49
39:Y:126:ARG:CZ	41:a:1083:U:P	3.00	0.49
41:a:3:U:C2	41:a:4:U:C5	3.01	0.49
41:a:132:G:H2'	41:a:133:U:C6	2.47	0.49
41:a:235:U:C2	41:a:236:C:C5	3.00	0.49
41:a:636:G:N2	61:u:76:GLU:OE2	2.45	0.49
41:a:840:C:C2	41:a:841:G:C8	3.01	0.49
41:a:1036:G:C6	41:a:1120:G:C5	3.01	0.49
41:a:1199:U:H2'	41:a:1200:C:C6	2.47	0.49
41:a:1348:C:O2	41:a:1348:C:H2'	2.11	0.49
41:a:1823:G:O2'	41:a:1824:G:H5'	2.12	0.49
41:a:2331:G:O2'	41:a:2336:A:N1	2.40	0.49
59:s:123:LYS:CE	59:s:132:HIS:HD2	2.25	0.49
62:v:29:GLY:HA3	62:v:104:GLU:HG3	1.94	0.49
66:z:53:ARG:HG3	66:z:57:PHE:HE2	1.75	0.49
2:1:17:VAL:HG12	2:1:76:VAL:HG21	1.94	0.49
3:2:11:LEU:HD12	3:2:11:LEU:N	2.26	0.49
4:3:94:ARG:CB	4:3:103:ILE:HD12	2.43	0.49
11:AA:887:VAL:O	16:AG:105:ILE:CD1	2.48	0.49
14:AE:287:ALA:HB3	14:AE:292:VAL:HG22	1.94	0.49
18:D:1218:C:H2'	18:D:1219:A:C8	2.48	0.49
18:D:1359:C:OP2	33:S:75:ARG:NE	2.43	0.49
21:G:114:LEU:HD13	21:G:144:LEU:HB3	1.94	0.49
25:K:134:ILE:O	25:K:137:VAL:HG12	2.12	0.49
25:K:157:ARG:HH12	28:N:43:GLU:CD	2.20	0.49
26:L:66:ALA:HB1	26:L:67:PRO:HD2	1.94	0.49
29:O:7:TYR:H	29:O:89:GLU:CD	2.19	0.49
39:Y:19:PRO:HG2	39:Y:23:VAL:CG2	2.42	0.49
39:Y:91:LYS:HZ3	39:Y:97:VAL:HG21	1.76	0.49
41:a:15:G:C2'	49:i:15:MET:HE1	2.42	0.49
41:a:57:C:H2'	41:a:58:G:O4'	2.12	0.49
41:a:851:C:C2	41:a:852:U:C5	3.00	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1571:A:H2'	41:a:1572:A:C8	2.47	0.49
41:a:1843:C:C2	41:a:1844:C:C5	3.00	0.49
41:a:2298:A:P	54:n:72:LYS:HZ3	2.33	0.49
41:a:2387:U:H4'	42:b:41:ARG:HH12	1.76	0.49
62:v:70:ASP:C	62:v:92:TRP:CZ3	2.91	0.49
66:z:79:PHE:CE1	66:z:110:VAL:HA	2.46	0.49
3:2:11:LEU:HD11	3:2:46:ALA:HB3	1.95	0.49
6:5:16:DG:O6	11:AA:451:ARG:NH1	2.45	0.49
9:9:43:LYS:HA	9:9:46:ARG:NH1	2.27	0.49
10:A:32:C:H2'	10:A:33:U:H5'	1.94	0.49
11:AA:618:GLN:HG3	14:AE:770:LEU:HD13	1.95	0.49
11:AA:1340:GLU:HB2	14:AE:19:ALA:HB3	1.94	0.49
14:AE:814:CYS:SG	14:AE:815:GLY:N	2.85	0.49
16:AG:342:ASP:N	16:AG:342:ASP:OD1	2.45	0.49
10:B:15:G:H2'	10:B:15:G:N3	2.27	0.49
10:B:23:C:H2'	10:B:24:U:H6	1.77	0.49
18:D:612:C:C2	18:D:613:C:C5	2.99	0.49
18:D:953:G:O6	38:X:103:LYS:CE	2.60	0.49
21:G:111:ILE:HD13	21:G:148:LEU:HD13	1.95	0.49
23:I:188:GLU:OE1	23:I:188:GLU:N	2.45	0.49
28:N:110:VAL:C	28:N:111:MET:SD	2.95	0.49
39:Y:126:ARG:HD2	39:Y:127:SER:N	2.26	0.49
41:a:44:A:H2'	41:a:45:G:O4'	2.12	0.49
41:a:145:C:H2'	41:a:146:A:H8	1.78	0.49
41:a:583:G:OP1	66:z:11:ARG:NH2	2.46	0.49
41:a:756:A:H2'	41:a:757:G:O4'	2.11	0.49
41:a:1142:A:C4	41:a:1144:A:N7	2.80	0.49
41:a:2815:C:O2'	49:i:41:HIS:ND1	2.41	0.49
46:f:41:THR:CA	46:f:45:ARG:HH22	2.26	0.49
47:g:47:LYS:NZ	54:n:115:ARG:HA	2.27	0.49
48:h:141:VAL:HG23	48:h:162:VAL:HG12	1.94	0.49
50:j:184:ARG:HH12	65:y:7:GLN:NE2	2.09	0.49
59:s:54:ILE:O	59:s:123:LYS:HG2	2.12	0.49
65:y:25:THR:OG1	65:y:88:ARG:HB3	2.12	0.49
66:z:58:ARG:HA	66:z:61:TRP:CE3	2.47	0.49
3:2:2:ILE:HG12	3:2:42:GLU:CD	2.37	0.49
7:6:19:DC:H2'	7:6:20:DA:C8	2.48	0.49
9:9:3:LEU:CD1	9:9:4:ASN:N	2.63	0.49
9:9:67:THR:N	9:9:68:PRO:CD	2.75	0.49
10:A:47:U:O2'	10:A:50:U:P	2.71	0.49
10:A:56:C:H2'	10:A:57:A:H5''	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:406:G:C5	18:D:495:A:C5	3.01	0.49
18:D:500:G:H2'	18:D:501:C:H6	1.77	0.49
18:D:555:U:H2'	18:D:556:C:C6	2.47	0.49
18:D:922:G:H2'	18:D:923:A:C8	2.47	0.49
18:D:1215:G:C2	18:D:1216:A:C8	3.01	0.49
19:E:54:MET:CE	19:E:75:HIS:HE1	2.23	0.49
21:G:110:SER:CA	21:G:113:ARG:HE	2.25	0.49
26:L:45:ARG:HA	26:L:45:ARG:NE	2.26	0.49
27:M:18:PHE:CD2	27:M:59:LEU:HD21	2.46	0.49
32:R:38:TYR:HE2	32:R:40:THR:CG2	2.25	0.49
32:R:109:ASP:C	32:R:111:LYS:HZ3	2.21	0.49
36:V:42:THR:HG22	36:V:44:LEU:HD11	1.95	0.49
39:Y:99:LYS:CD	39:Y:138:VAL:O	2.61	0.49
41:a:272:A:N3	41:a:273:G:C8	2.80	0.49
41:a:1223:G:N2	41:a:1226:A:OP2	2.43	0.49
41:a:2298:A:O3'	54:n:72:LYS:HE2	2.11	0.49
41:a:2349:G:O6	41:a:2369:A:N6	2.46	0.49
43:c:65:ASP:CG	43:c:66:THR:N	2.70	0.49
46:f:3:LYS:HE2	46:f:3:LYS:HA	1.94	0.49
48:h:52:ARG:HG3	48:h:53:HIS:ND1	2.27	0.49
52:l:5:LEU:HD23	52:l:122:GLU:HG3	1.95	0.49
5:4:35:GLU:OE1	5:4:93:ARG:NH2	2.46	0.49
9:9:31:ARG:HH22	41:a:1106:G:N2	2.10	0.49
9:9:52:MET:CE	9:9:85:SER:HB2	2.42	0.49
13:AD:100:LEU:HB2	13:AD:144:ILE:HG23	1.94	0.49
16:AG:296:ILE:HD11	16:AG:307:ILE:HD13	1.93	0.49
16:AG:440:VAL:HG22	16:AG:481:LEU:HD21	1.95	0.49
18:D:736:C:H2'	18:D:737:C:H6	1.77	0.49
18:D:744:C:H2'	18:D:745:G:H8	1.78	0.49
22:H:112:ILE:HA	22:H:121:THR:O	2.12	0.49
28:N:106:THR:HB	28:N:121:LEU:CD1	2.41	0.49
34:T:33:THR:HG21	34:T:85:LEU:HD12	1.94	0.49
35:U:12:LYS:HD3	35:U:13:LYS:HG2	1.93	0.49
36:V:62:ARG:HA	36:V:73:TRP:CZ3	2.47	0.49
37:W:33:THR:HG21	37:W:49:ILE:HD11	1.94	0.49
37:W:45:ILE:HA	37:W:62:VAL:HG11	1.93	0.49
38:X:81:MET:HG2	41:a:888:C:C4	2.48	0.49
41:a:1199:U:H2'	41:a:1200:C:H6	1.77	0.49
41:a:1830:C:N3	41:a:1831:G:N7	2.61	0.49
41:a:2591:C:H2'	41:a:2592:G:H8	1.78	0.49
41:a:2684:U:C4	41:a:2685:G:N7	2.81	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:n:16:LEU:HD12	54:n:28:VAL:CG2	2.43	0.49
54:n:70:ALA:O	54:n:81:GLN:OE1	2.31	0.49
60:t:7:MET:SD	60:t:18:ARG:NE	2.85	0.49
64:x:90:VAL:HG22	64:x:115:LEU:HD11	1.94	0.49
66:z:74:ILE:HG13	66:z:78:LYS:HE3	1.94	0.49
2:1:95:ARG:NH1	2:1:97:LEU:HG	2.28	0.49
9:9:67:THR:O	9:9:72:LEU:HG	2.12	0.49
18:D:373:A:C2	18:D:374:A:C8	3.00	0.49
18:D:1017:U:O2'	18:D:1018:G:O4'	2.30	0.49
18:D:1082:A:OP1	25:K:23:LYS:NZ	2.34	0.49
21:G:109:GLN:O	21:G:113:ARG:NE	2.45	0.49
25:K:13:GLU:C	25:K:14:LYS:HD2	2.38	0.49
26:L:35:LYS:HG3	26:L:36:ILE:N	2.27	0.49
27:M:111:ARG:NH2	27:M:126:ASP:OD2	2.43	0.49
28:N:35:ALA:O	28:N:39:VAL:HG23	2.12	0.49
28:N:96:MET:HG2	28:N:96:MET:O	2.12	0.49
34:T:77:ARG:HH21	34:T:80:GLN:NE2	2.11	0.49
38:X:92:ARG:HD2	41:a:888:C:OP1	2.12	0.49
41:a:807:U:OP2	61:u:41:ARG:NH1	2.42	0.49
41:a:1790:C:H2'	41:a:1791:A:C8	2.48	0.49
41:a:1808:A:O2'	43:c:3:ARG:NH1	2.46	0.49
41:a:1932:A:H2'	41:a:1933:G:O4'	2.13	0.49
41:a:2082:A:H2'	41:a:2083:G:O4'	2.12	0.49
41:a:2362:C:O5'	55:o:40:ARG:NH2	2.45	0.49
43:c:6:GLN:CD	43:c:49:LEU:HD22	2.38	0.49
57:q:1:MET:CA	57:q:1:MET:HE3	2.43	0.49
57:q:10:LEU:HD23	57:q:33:HIS:CD2	2.47	0.49
63:w:35:LYS:NZ	63:w:112:TYR:CE2	2.76	0.49
65:y:9:GLU:HG3	65:y:55:LEU:CB	2.42	0.49
2:1:1:MET:CG	2:1:62:ASP:OD2	2.61	0.49
2:1:71:VAL:O	2:1:71:VAL:HG23	2.13	0.49
5:4:89:ILE:CG2	5:4:91:PHE:CE1	2.95	0.49
8:7:3:G:H5''	18:D:1501:C:N4	2.27	0.49
12:AB:6:LEU:O	12:AB:7:LEU:HD23	2.13	0.49
12:AB:92:VAL:HG13	12:AB:96:LEU:O	2.13	0.49
12:AB:136:GLU:OE1	12:AB:160:ARG:CZ	2.61	0.49
14:AE:678:ARG:NH1	14:AE:756:GLU:OE1	2.46	0.49
16:AG:283:PHE:CE2	16:AG:332:SER:HA	2.47	0.49
16:AG:361:LYS:HD2	16:AG:417:ILE:HG12	1.92	0.49
16:AG:448:LEU:CG	16:AG:473:LEU:HD11	2.42	0.49
17:C:24:LYS:HB3	26:L:102:MET:HE1	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:109:A:H2'	18:D:326:G:N2	2.27	0.49
18:D:829:G:O3'	21:G:23:TRP:HZ2	1.95	0.49
23:I:77:ILE:HA	23:I:84:VAL:HG13	1.94	0.49
34:T:33:THR:OG1	34:T:63:ARG:NH1	2.45	0.49
41:a:1921:G:C2'	41:a:1922:G:H5'	2.43	0.49
41:a:2065:C:H2'	41:a:2066:C:H6	1.77	0.49
47:g:36:VAL:HG21	47:g:41:HIS:ND1	2.27	0.49
47:g:63:ARG:HG3	47:g:64:PHE:CD1	2.48	0.49
49:i:40:ARG:NH1	49:i:41:HIS:CD2	2.80	0.49
50:j:55:LYS:HZ2	50:j:60:VAL:CA	2.26	0.49
51:k:34:LEU:CB	51:k:51:GLU:OE2	2.60	0.49
56:p:94:TYR:CE1	56:p:152:ARG:HD2	2.48	0.49
57:q:1:MET:HE1	57:q:35:GLN:HA	1.95	0.49
59:s:55:ILE:CG1	59:s:123:LYS:HE2	2.42	0.49
60:t:1:MET:HE3	60:t:32:TYR:HB3	1.94	0.49
63:w:87:PHE:HZ	63:w:115:LEU:HG	1.77	0.49
65:y:7:GLN:NE2	65:y:10:GLN:HE21	2.11	0.49
66:z:51:ARG:N	66:z:51:ARG:HD2	2.28	0.49
2:1:23:LEU:HD21	2:1:35:ILE:HG22	1.94	0.49
7:6:23:DC:H5'	14:AE:259:ARG:NH1	2.11	0.49
9:9:52:MET:HE2	9:9:85:SER:HB2	1.95	0.49
11:AA:732:ILE:HD11	11:AA:769:PRO:HB3	1.95	0.49
12:AB:2:GLN:HB2	12:AB:60:ASP:HB2	1.95	0.49
14:AE:977:SER:OG	14:AE:980:THR:OG1	2.31	0.49
16:AG:130:PHE:C	16:AG:186:VAL:CG1	2.86	0.49
16:AG:360:THR:HG22	16:AG:360:THR:O	2.10	0.49
17:C:45:THR:HA	22:H:340:ARG:HH21	1.77	0.49
18:D:331:G:H2'	19:E:4:ILE:HD11	1.95	0.49
18:D:429:U:N3	18:D:431:A:N6	2.61	0.49
18:D:925:G:C6	18:D:927:G:N7	2.80	0.49
20:F:63:GLU:O	20:F:67:ARG:HD2	2.12	0.49
25:K:34:THR:HG22	25:K:52:LYS:HD3	1.95	0.49
27:M:74:GLU:OE1	27:M:75:VAL:O	2.30	0.49
27:M:109:ARG:O	27:M:119:ARG:NH2	2.46	0.49
41:a:208:C:H2'	41:a:209:C:H6	1.78	0.49
41:a:2230:G:H5'	43:c:30:LEU:HD23	1.94	0.49
41:a:2245:U:O2'	41:a:2436:G:OP2	2.27	0.49
41:a:2259:U:C2	41:a:2260:C:C5	3.01	0.49
41:a:2392:A:C8	41:a:2429:G:C2	3.00	0.49
41:a:2455:G:H2'	41:a:2456:C:C6	2.48	0.49
49:i:40:ARG:NH1	49:i:41:HIS:CG	2.81	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:k:21:TYR:CE2	51:k:38:LYS:CG	2.96	0.49
52:l:97:ASN:ND2	52:l:100:MET:HE2	2.27	0.49
56:p:26:ILE:HG22	56:p:75:MET:HE2	1.93	0.49
59:s:13:ARG:HH22	59:s:49:ASP:C	2.19	0.49
59:s:73:VAL:CG1	59:s:74:TYR:N	2.75	0.49
61:u:59:ARG:HH11	61:u:59:ARG:HG3	1.78	0.49
61:u:117:THR:HG23	61:u:118:THR:N	2.28	0.49
66:z:72:ASN:ND2	66:z:106:PHE:CE2	2.81	0.49
1:0:73:LYS:HZ1	1:0:88:GLY:HA3	1.78	0.49
3:2:12:ARG:HH21	45:e:29:ARG:CZ	2.25	0.49
3:2:14:PRO:HA	3:2:32:LEU:HD13	1.94	0.49
12:AB:112:PRO:HA	12:AB:133:PRO:HG2	1.95	0.49
14:AE:209:ASN:HB3	14:AE:214:ARG:HH21	1.76	0.49
14:AE:749:LYS:HD3	14:AE:753:SER:HB2	1.94	0.49
16:AG:277:ASP:HB2	16:AG:282:GLN:HG3	1.95	0.49
16:AG:285:ILE:HD13	16:AG:285:ILE:H	1.76	0.49
16:AG:353:HIS:HA	16:AG:356:ILE:HG21	1.94	0.49
16:AG:453:VAL:HG11	16:AG:459:LEU:CG	2.42	0.49
17:C:13:PHE:CB	17:C:51:TYR:CD1	2.95	0.49
18:D:373:A:C2	18:D:482:A:C6	3.00	0.49
18:D:1071:C:H2'	18:D:1072:G:C8	2.47	0.49
21:G:126:PHE:C	21:G:127:ASP:O	2.55	0.49
23:I:164:ARG:HD2	23:I:166:GLU:OE2	2.12	0.49
26:L:65:GLU:OE1	26:L:65:GLU:HA	2.12	0.49
29:O:32:GLN:HG3	29:O:32:GLN:O	2.13	0.49
32:R:74:LEU:HD11	32:R:80:ILE:CD1	2.43	0.49
33:S:57:PRO:HA	33:S:60:GLN:OE1	2.13	0.49
36:V:17:MET:HG3	36:V:20:SER:HB2	1.95	0.49
38:X:9:ILE:HG13	38:X:10:PRO:HD2	1.95	0.49
52:l:40:ARG:HD2	52:l:92:HIS:CE1	2.48	0.49
52:l:194:LYS:O	52:l:197:GLU:N	2.45	0.49
1:0:61:ALA:HB1	1:0:96:VAL:CG2	2.43	0.48
9:9:31:ARG:NE	9:9:31:ARG:HA	2.28	0.48
11:AA:554:HIS:HD2	11:AA:558:VAL:HB	1.77	0.48
12:AB:145:LEU:HD22	12:AB:148:LYS:HG2	1.95	0.48
14:AE:520:ALA:HB3	14:AE:546:ALA:HB2	1.95	0.48
18:D:563:A:N1	18:D:884:U:O4	2.46	0.48
25:K:46:VAL:N	25:K:71:MET:HE2	2.27	0.48
35:U:18:GLN:NE2	35:U:35:ARG:HE	2.10	0.48
36:V:42:THR:HG22	36:V:44:LEU:HD12	1.95	0.48
39:Y:99:LYS:HZ3	39:Y:138:VAL:HG13	1.75	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:920:A:C5	41:a:921:C:C5	3.01	0.48
41:a:2014:A:H2'	41:a:2015:A:C8	2.47	0.48
41:a:2677:G:C6	41:a:2731:G:C6	3.01	0.48
48:h:123:ALA:HB1	48:h:125:LYS:HZ3	1.78	0.48
52:l:149:ILE:CG2	52:l:175:ILE:HD11	2.43	0.48
56:p:9:VAL:HG23	56:p:69:ARG:HD2	1.95	0.48
56:p:162:VAL:O	56:p:162:VAL:HG12	2.11	0.48
59:s:13:ARG:NH2	59:s:50:THR:HA	2.27	0.48
59:s:125:TYR:HE2	59:s:130:HIS:HB2	1.78	0.48
60:t:88:ASN:OD1	60:t:89:ASN:N	2.46	0.48
62:v:95:LEU:HD23	62:v:95:LEU:H	1.78	0.48
63:w:38:LEU:N	63:w:39:PRO:CD	2.76	0.48
65:y:40:LEU:HD12	65:y:40:LEU:N	2.28	0.48
9:9:30:SER:O	9:9:31:ARG:CZ	2.59	0.48
11:AA:861:ALA:CB	16:AG:38:LYS:HZ1	2.27	0.48
12:AB:118:ILE:CG1	12:AB:142:LEU:CD2	2.91	0.48
16:AG:36:LYS:HE3	16:AG:43:ILE:HG12	1.95	0.48
16:AG:323:GLN:CA	16:AG:323:GLN:NE2	2.75	0.48
16:AG:389:MET:HA	16:AG:407:ARG:NH1	2.29	0.48
16:AG:444:LEU:HD22	16:AG:473:LEU:CD2	2.43	0.48
18:D:56:U:H2'	18:D:57:G:C8	2.48	0.48
18:D:893:C:C2	18:D:894:G:C8	3.01	0.48
18:D:1319:A:C8	18:D:1323:G:C6	3.01	0.48
23:I:142:MET:HG3	23:I:170:GLU:OE1	2.13	0.48
27:M:16:PRO:CB	29:O:46:MET:HG3	2.42	0.48
27:M:138:ARG:HG2	27:M:142:HIS:CE1	2.48	0.48
38:X:16:VAL:HG23	38:X:17:ILE:N	2.29	0.48
39:Y:57:VAL:HG23	39:Y:71:LYS:HD3	1.93	0.48
41:a:26:G:H2'	41:a:27:G:O4'	2.13	0.48
41:a:1021:A:H3'	41:a:1021:A:N3	2.27	0.48
41:a:1792:G:H5''	48:h:204:VAL:HG23	1.95	0.48
41:a:2093:G:O2'	41:a:2198:A:N1	2.41	0.48
41:a:2297:A:N1	41:a:2321:U:O4	2.47	0.48
41:a:2485:G:O3'	62:v:125:PRO:HB3	2.12	0.48
46:f:41:THR:N	46:f:45:ARG:NH2	2.61	0.48
52:l:147:LEU:HD11	52:l:170:ARG:CD	2.43	0.48
55:o:62:LEU:HG	55:o:65:ALA:HA	1.95	0.48
59:s:88:THR:O	59:s:92:MET:HE2	2.13	0.48
62:v:58:LYS:O	62:v:59:ARG:HG3	2.13	0.48
9:9:95:LEU:HD12	9:9:99:PHE:CZ	2.48	0.48
10:A:22:G:HO2'	10:A:23:C:P	2.36	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:79:LYS:HZ3	16:AG:153:ASP:CG	2.20	0.48
14:AE:425:ARG:NH1	14:AE:458:ASN:O	2.45	0.48
16:AG:288:MET:O	16:AG:288:MET:HG2	2.13	0.48
18:D:123:U:C2	18:D:124:C:C5	3.01	0.48
18:D:123:U:H5''	18:D:311:C:O2'	2.13	0.48
21:G:112:LYS:HE2	21:G:112:LYS:HA	1.95	0.48
24:J:170:TRP:CZ3	24:J:190:ASP:HB3	2.48	0.48
25:K:12:GLN:HB3	25:K:14:LYS:HZ2	1.79	0.48
34:T:64:ARG:CZ	34:T:64:ARG:CB	2.91	0.48
38:X:86:TYR:HE2	38:X:90:ARG:NH2	2.11	0.48
41:a:39:G:H1'	52:l:43:THR:HG21	1.95	0.48
41:a:379:G:N1	41:a:396:G:C6	2.81	0.48
41:a:1405:U:H2'	41:a:1406:U:C5	2.49	0.48
41:a:1590:A:H2'	41:a:1591:A:C8	2.48	0.48
48:h:160:THR:O	48:h:195:VAL:HG12	2.13	0.48
56:p:94:TYR:C	56:p:95:ARG:HD2	2.38	0.48
59:s:15:TRP:HD1	59:s:53:TYR:HB2	1.78	0.48
59:s:39:LYS:HA	59:s:44:TYR:HD1	1.78	0.48
62:v:41:LEU:HG	62:v:96:ILE:HG13	1.95	0.48
62:v:57:VAL:HG22	62:v:60:GLN:O	2.13	0.48
64:x:115:LEU:HD23	64:x:117:PHE:CE1	2.49	0.48
11:AA:1280:ALA:HB1	14:AE:918:ILE:HG22	1.96	0.48
12:AB:32:MET:HE2	12:AB:32:MET:HA	1.95	0.48
16:AG:148:ASP:HA	16:AG:164:ARG:CD	2.44	0.48
16:AG:453:VAL:HG11	16:AG:459:LEU:HD23	1.95	0.48
16:AG:467:LEU:HD12	16:AG:482:ILE:HD11	1.95	0.48
10:B:12:G:C2'	10:B:13:C:OP1	2.61	0.48
10:B:21:A:H4'	10:B:21:A:OP1	2.13	0.48
17:C:60:LYS:NZ	18:D:734:G:C3'	2.76	0.48
18:D:718:A:C5	31:Q:118:HIS:ND1	2.81	0.48
23:I:134:MET:HE2	23:I:168:TYR:CG	2.49	0.48
24:J:113:GLU:O	24:J:117:LEU:HG	2.13	0.48
28:N:40:LEU:O	28:N:44:GLY:N	2.47	0.48
29:O:52:LEU:O	29:O:55:VAL:O	2.31	0.48
30:P:5:ARG:HG3	30:P:7:ARG:HH12	1.78	0.48
38:X:94:GLY:C	38:X:95:LEU:HD22	2.38	0.48
39:Y:74:PRO:O	39:Y:78:LEU:HG	2.13	0.48
40:Z:26:MET:O	40:Z:30:PHE:HB2	2.14	0.48
41:a:879:G:H2'	41:a:880:G:O4'	2.14	0.48
41:a:1664:A:C2'	41:a:1665:A:H5'	2.43	0.48
41:a:2012:G:H8	41:a:2012:G:O5'	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2296:U:H4'	41:a:2297:A:OP1	2.14	0.48
41:a:2298:A:C4	41:a:2321:U:H5	2.31	0.48
41:a:2547:A:H2'	41:a:2548:U:H6	1.77	0.48
48:h:138:GLY:H	48:h:164:ILE:HG23	1.79	0.48
50:j:55:LYS:NZ	50:j:59:ARG:C	2.70	0.48
63:w:114:GLU:N	63:w:114:GLU:OE1	2.46	0.48
66:z:79:PHE:CE1	66:z:110:VAL:HG22	2.48	0.48
11:AA:565:GLU:HA	11:AA:569:ILE:HG12	1.95	0.48
11:AA:860:ALA:N	16:AG:38:LYS:HE2	2.17	0.48
12:AB:151:LYS:HG3	12:AB:151:LYS:O	2.14	0.48
14:AE:640:GLY:N	14:AE:643:ASP:OD2	2.42	0.48
16:AG:59:PHE:CD1	16:AG:97:ILE:HG23	2.48	0.48
16:AG:64:VAL:HA	16:AG:75:ILE:HB	1.94	0.48
16:AG:369:PHE:O	16:AG:373:LEU:HG	2.12	0.48
17:C:60:LYS:CD	18:D:734:G:O2'	2.62	0.48
18:D:35:G:H2'	18:D:36:C:C6	2.48	0.48
20:F:62:ARG:HD2	20:F:63:GLU:N	2.29	0.48
24:J:124:MET:HE2	24:J:128:ARG:CA	2.43	0.48
26:L:38:ARG:HD2	26:L:40:GLU:OE2	2.13	0.48
26:L:62:MET:HE3	26:L:62:MET:CA	2.44	0.48
29:O:52:LEU:HB3	29:O:58:VAL:HG12	1.94	0.48
30:P:63:ASP:OD1	30:P:65:TYR:CZ	2.67	0.48
30:P:64:GLN:OE1	30:P:65:TYR:O	2.31	0.48
39:Y:126:ARG:HH12	41:a:1082:U:C3'	2.26	0.48
41:a:1141:U:H4'	41:a:1142:A:O5'	2.13	0.48
41:a:2515:C:H2'	41:a:2516:A:H8	1.78	0.48
41:a:2809:A:H2'	41:a:2810:A:C8	2.48	0.48
42:b:47:ALA:HB2	42:b:59:LEU:HD11	1.94	0.48
47:g:47:LYS:HE2	54:n:115:ARG:CZ	2.43	0.48
50:j:55:LYS:HZ1	50:j:59:ARG:HB3	1.77	0.48
50:j:124:ARG:HA	50:j:165:MET:CE	2.44	0.48
54:n:22:TYR:HE2	54:n:29:PRO:HD3	1.79	0.48
57:q:2:LYS:CE	57:q:35:GLN:HE21	2.25	0.48
57:q:14:CYS:SG	57:q:33:HIS:HB3	2.53	0.48
59:s:1:MET:HE2	66:z:97:ASP:OD2	2.13	0.48
59:s:98:GLU:OE1	59:s:124:VAL:HG13	2.13	0.48
60:t:20:MET:HB3	60:t:44:LYS:HE2	1.95	0.48
66:z:78:LYS:CD	66:z:117:LEU:HD21	2.43	0.48
2:1:25:ARG:HH12	41:a:520:G:P	2.37	0.48
4:3:6:ARG:NH1	41:a:84:A:H2'	2.27	0.48
9:9:24:SER:O	9:9:116:GLU:HB2	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:A:23:C:H2'	10:A:24:U:H6	1.77	0.48
11:AA:9:LYS:HG2	11:AA:1171:ARG:HH12	1.77	0.48
11:AA:1286:THR:O	11:AA:1290:MET:HB2	2.14	0.48
14:AE:79:LYS:CB	16:AG:159:GLU:OE2	2.60	0.48
10:B:72:A:H2'	10:B:73:A:H5''	1.95	0.48
18:D:43:C:P	35:U:12:LYS:NZ	2.86	0.48
18:D:626:G:O3'	35:U:51:ARG:NH1	2.45	0.48
18:D:718:A:C8	31:Q:118:HIS:HB3	2.48	0.48
18:D:1307:U:H2'	18:D:1308:U:C6	2.49	0.48
18:D:1314:C:H2'	18:D:1315:U:C6	2.48	0.48
18:D:1314:C:H2'	18:D:1315:U:H6	1.78	0.48
18:D:1439:G:OP1	19:E:33:LYS:CE	2.61	0.48
25:K:159:LYS:HZ1	28:N:44:GLY:HA3	1.77	0.48
26:L:36:ILE:HD12	26:L:63:ASN:O	2.14	0.48
27:M:142:HIS:O	27:M:146:GLU:HG2	2.13	0.48
31:Q:94:GLU:O	31:Q:97:ILE:HG22	2.13	0.48
34:T:35:GLN:HB3	34:T:59:MET:HE1	1.95	0.48
34:T:71:LYS:HE2	34:T:78:TYR:CE2	2.48	0.48
41:a:576:U:H2'	41:a:577:G:C8	2.48	0.48
41:a:581:C:C2	41:a:582:A:C8	3.01	0.48
41:a:1844:C:C2	41:a:1845:G:C8	3.02	0.48
41:a:2189:U:O2'	41:a:2190:G:O5'	2.31	0.48
41:a:2193:G:O2'	41:a:2194:U:P	2.72	0.48
41:a:2313:C:H5''	54:n:88:LYS:HD2	1.95	0.48
41:a:2327:A:H2'	41:a:2328:A:C8	2.49	0.48
41:a:2685:G:OP1	60:t:78:ARG:NH2	2.47	0.48
41:a:2754:U:O3'	57:q:19:ARG:NH2	2.46	0.48
49:i:28:LEU:O	49:i:37:LYS:NZ	2.46	0.48
54:n:171:ALA:O	54:n:174:ASP:N	2.46	0.48
59:s:56:VAL:O	59:s:124:VAL:HA	2.14	0.48
64:x:115:LEU:HD23	64:x:117:PHE:HE1	1.79	0.48
5:4:80:HIS:CE1	5:4:83:LYS:HZ3	2.31	0.48
6:5:10:DT:C4'	11:AA:62:TYR:OH	2.61	0.48
6:5:17:DC:C5	11:AA:542:ARG:HD3	2.48	0.48
14:AE:67:ASP:C	14:AE:69:GLU:N	2.71	0.48
10:B:32:C:H2'	10:B:33:U:H5'	1.94	0.48
18:D:378:G:H2'	18:D:379:C:C6	2.49	0.48
18:D:429:U:O2	18:D:430:A:C8	2.67	0.48
18:D:908:A:H2'	18:D:909:A:C8	2.48	0.48
18:D:1397:C:OP1	18:D:1397:C:H2'	2.13	0.48
18:D:1428:A:H2'	18:D:1429:A:O4'	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:E:19:LYS:CG	19:E:20:HIS:N	2.76	0.48
19:E:22:ALA:HA	19:E:25:ARG:HE	1.78	0.48
20:F:4:ILE:CD1	20:F:19:PHE:HA	2.43	0.48
23:I:134:MET:HG3	23:I:151:VAL:CG2	2.44	0.48
24:J:61:VAL:HA	24:J:64:ILE:HD12	1.96	0.48
39:Y:77:VAL:HG13	39:Y:81:LYS:NZ	2.27	0.48
41:a:880:G:C2	41:a:881:G:C8	3.02	0.48
41:a:2091:C:O2	43:c:34:HIS:CE1	2.67	0.48
41:a:2093:G:N7	41:a:2225:A:H2'	2.28	0.48
50:j:101:PHE:CZ	50:j:203:VAL:HG12	2.48	0.48
51:k:7:GLU:HG2	51:k:27:LYS:NZ	2.29	0.48
54:n:8:TYR:HA	54:n:12:VAL:HB	1.95	0.48
54:n:134:GLU:CG	54:n:137:ILE:HG23	2.41	0.48
2:l:95:ARG:NH1	2:l:97:LEU:CD2	2.77	0.48
5:4:1:MET:SD	5:4:3:THR:OG1	2.71	0.48
9:9:25:ALA:CB	9:9:99:PHE:HD2	2.25	0.48
10:A:21:A:H4'	10:A:21:A:OP1	2.13	0.48
12:AB:155:LYS:CG	12:AB:156:ASN:H	2.26	0.48
14:AE:107:LEU:HD22	14:AE:299:LEU:HD21	1.95	0.48
14:AE:1289:ASN:ND2	14:AE:1300:ALA:O	2.46	0.48
16:AG:18:LEU:HB3	16:AG:19:PRO:HD2	1.95	0.48
16:AG:87:LEU:HD22	16:AG:93:VAL:HG21	1.96	0.48
16:AG:220:VAL:CG1	16:AG:245:ILE:HG21	2.44	0.48
18:D:35:G:N3	32:R:115:SER:OG	2.47	0.48
18:D:71:A:N1	18:D:100:G:C8	2.81	0.48
18:D:109:A:C6	18:D:326:G:C6	3.02	0.48
18:D:562:U:O3'	32:R:14:ARG:NH1	2.46	0.48
18:D:1530:G:H2'	18:D:1531:A:C8	2.49	0.48
23:I:119:SER:O	23:I:123:GLN:HG2	2.13	0.48
23:I:156:ARG:HG2	23:I:193:TYR:HB2	1.95	0.48
25:K:148:ASN:ND2	25:K:153:VAL:CG1	2.77	0.48
27:M:71:PRO:O	27:M:96:ARG:NH1	2.47	0.48
36:V:50:ASN:HB2	36:V:52:GLU:OE2	2.13	0.48
40:Z:29:LYS:HG3	40:Z:30:PHE:CE2	2.49	0.48
41:a:871:U:H2'	41:a:872:U:H6	1.78	0.48
41:a:962:G:C5	41:a:963:U:C5	3.01	0.48
41:a:1042:G:C4	41:a:1043:C:C6	3.02	0.48
41:a:2373:G:H2'	41:a:2374:C:C6	2.49	0.48
41:a:2884:U:N3	49:i:40:ARG:NH1	2.62	0.48
52:l:195:GLN:HA	52:l:198:GLU:OE1	2.13	0.48
3:2:56:GLU:OE2	3:2:88:LYS:CA	2.61	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:55:GLU:OE1	5:4:55:GLU:N	2.42	0.48
5:4:69:GLU:H	5:4:69:GLU:CD	2.22	0.48
9:9:6:GLN:HA	9:9:9:GLN:HE22	1.74	0.48
9:9:16:SER:HB2	9:9:20:LYS:NZ	2.29	0.48
9:9:103:ASN:HA	9:9:107:GLU:HG3	1.96	0.48
10:A:12:G:C2'	10:A:13:C:OP1	2.61	0.48
10:A:70:G:C3'	10:A:71:C:H5''	2.44	0.48
13:AD:182:ARG:O	13:AD:205:MET:HA	2.13	0.48
16:AG:26:ALA:HB1	16:AG:117:LYS:HG2	1.96	0.48
16:AG:27:LEU:HD22	16:AG:114:ILE:HG12	1.96	0.48
10:B:26:G:H1	10:B:44:A:N6	2.12	0.48
18:D:977:A:O2'	18:D:979:C:OP2	2.25	0.48
19:E:9:LYS:O	19:E:12:ILE:HG12	2.14	0.48
22:H:5:PHE:CE1	22:H:9:PHE:HE1	2.31	0.48
24:J:62:ARG:CZ	24:J:69:GLU:HG2	2.44	0.48
28:N:83:LEU:HD11	32:R:4:VAL:HB	1.95	0.48
29:O:49:ARG:C	29:O:53:GLU:OE1	2.57	0.48
31:Q:31:ILE:HD12	31:Q:46:THR:HG22	1.94	0.48
32:R:80:ILE:HG22	32:R:81:LEU:N	2.27	0.48
41:a:6:A:O3'	59:s:132:HIS:CE1	2.67	0.48
41:a:1197:G:C4	41:a:1198:U:C5	3.02	0.48
41:a:1858:A:N6	41:a:1884:G:H1'	2.29	0.48
41:a:2061:G:C2	41:a:2063:C:C5	3.01	0.48
65:y:13:MET:HG3	65:y:77:HIS:CE1	2.49	0.48
1:0:6:GLN:OE1	1:0:11:GLN:HG2	2.14	0.48
2:1:20:VAL:O	2:1:23:LEU:HB3	2.13	0.48
11:AA:411:ARG:NH2	11:AA:427:ASP:OD2	2.44	0.48
12:AB:115:LYS:CD	12:AB:129:ILE:CB	2.92	0.48
16:AG:448:LEU:HD23	16:AG:473:LEU:HD12	1.95	0.48
18:D:330:C:O2	18:D:330:C:H2'	2.14	0.48
18:D:1127:G:C2	18:D:1128:C:C6	3.01	0.48
24:J:139:PRO:N	24:J:182:PHE:HE2	2.12	0.48
24:J:195:ILE:HG13	24:J:195:ILE:O	2.14	0.48
25:K:112:ARG:C	25:K:116:GLU:OE1	2.57	0.48
30:P:15:HIS:O	30:P:18:ILE:HG22	2.14	0.48
31:Q:85:MET:HB3	31:Q:113:VAL:HG11	1.96	0.48
33:S:47:LYS:O	33:S:50:THR:OG1	2.25	0.48
38:X:86:TYR:CE2	38:X:90:ARG:NE	2.82	0.48
39:Y:126:ARG:NH1	41:a:1083:U:OP2	2.47	0.48
41:a:760:G:H2'	41:a:761:A:O4'	2.14	0.48
41:a:984:A:N3	41:a:984:A:H2'	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1708:C:H2'	41:a:1709:U:H6	1.79	0.48
41:a:2070:A:H2'	41:a:2071:A:O4'	2.14	0.48
45:e:10:SER:OG	45:e:13:GLU:HG2	2.13	0.48
51:k:19:HIS:CD2	51:k:20:PHE:N	2.82	0.48
56:p:69:ARG:NH1	56:p:73:ASN:HD21	2.11	0.48
58:r:18:GLN:HE22	58:r:39:ALA:HB2	1.79	0.48
62:v:53:MET:SD	62:v:53:MET:N	2.86	0.48
62:v:111:GLU:O	62:v:112:LEU:C	2.57	0.48
63:w:33:ILE:HD11	63:w:112:TYR:CE1	2.48	0.48
65:y:9:GLU:HG3	65:y:55:LEU:HB2	1.96	0.48
66:z:53:ARG:HG2	66:z:57:PHE:CE2	2.49	0.48
2:1:28:LYS:HB2	2:1:31:GLN:OE1	2.14	0.47
8:7:7:U:C6	8:7:7:U:C3'	2.97	0.47
11:AA:1314:GLN:HB2	15:AF:28:ARG:HH12	1.79	0.47
12:AB:51:PHE:CZ	14:AE:288:PRO:CG	2.89	0.47
14:AE:438:GLU:OE2	14:AE:481:ARG:NH1	2.44	0.47
16:AG:207:GLU:HG3	16:AG:210:ARG:HB3	1.96	0.47
16:AG:209:PHE:CE1	16:AG:262:VAL:HG21	2.49	0.47
16:AG:434:LEU:HD11	16:AG:454:CYS:C	2.39	0.47
18:D:808:C:OP1	34:T:48:LYS:HE3	2.14	0.47
18:D:1227:A:C2	18:D:1228:C:C6	3.02	0.47
24:J:76:TYR:HE2	24:J:204:TYR:CB	2.25	0.47
25:K:16:ILE:H	25:K:16:ILE:HD12	1.79	0.47
29:O:49:ARG:O	29:O:53:GLU:OE1	2.32	0.47
29:O:118:LEU:HD22	29:O:123:ARG:C	2.39	0.47
32:R:74:LEU:HD11	32:R:80:ILE:HG13	1.96	0.47
33:S:74:LEU:HD23	33:S:77:PHE:CD2	2.49	0.47
35:U:26:ASN:ND2	35:U:31:ARG:HE	2.11	0.47
41:a:2043:C:C2	41:a:2044:C:C5	3.02	0.47
41:a:2300:C:H2'	41:a:2301:C:H6	1.79	0.47
44:d:42:C:C5	54:n:66:LEU:HD12	2.49	0.47
62:v:42:THR:HA	62:v:93:VAL:HG12	1.95	0.47
64:x:90:VAL:HG13	64:x:115:LEU:HD11	1.96	0.47
5:4:1:MET:O	5:4:1:MET:HE3	2.13	0.47
9:9:24:SER:HG	9:9:85:SER:C	2.22	0.47
10:A:22:G:O2'	10:A:23:C:O5'	2.31	0.47
10:A:26:G:H1	10:A:44:A:N6	2.12	0.47
12:AB:54:TYR:HE2	14:AE:291:ILE:CG2	2.13	0.47
12:AB:102:LYS:NZ	14:AE:285:LEU:HG	2.28	0.47
16:AG:62:TRP:CD1	16:AG:95:ASP:HB2	2.49	0.47
16:AG:299:ASP:CB	16:AG:303:HIS:HA	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:B:22:G:O2'	10:B:23:C:O5'	2.32	0.47
17:C:65:LEU:HD23	17:C:67:LEU:HD11	1.96	0.47
18:D:1149:C:H2'	18:D:1150:A:H8	1.79	0.47
18:D:1176:A:H2'	18:D:1177:G:O4'	2.15	0.47
18:D:1227:A:H5'	38:X:110:LYS:NZ	2.28	0.47
18:D:1324:A:H2'	18:D:1325:C:H6	1.78	0.47
21:G:110:SER:HA	21:G:113:ARG:HH21	1.79	0.47
31:Q:78:GLY:O	31:Q:80:LYS:NZ	2.42	0.47
37:W:62:VAL:HG13	37:W:66:MET:CE	2.44	0.47
39:Y:106:GLN:OE1	39:Y:125:THR:HG21	2.14	0.47
39:Y:126:ARG:NH1	41:a:1082:U:O5'	2.46	0.47
41:a:208:C:C2	41:a:209:C:C5	3.01	0.47
41:a:1048:A:C4	41:a:1049:C:C5	3.02	0.47
41:a:1321:A:C5	41:a:1322:A:C8	3.02	0.47
41:a:1790:C:H2'	41:a:1791:A:N7	2.29	0.47
41:a:1825:U:P	48:h:52:ARG:NH1	2.86	0.47
41:a:2788:C:H2'	41:a:2789:C:C6	2.50	0.47
43:c:67:VAL:HA	43:c:70:GLU:OE1	2.14	0.47
44:d:117:G:OP1	64:x:56:LYS:HD3	2.14	0.47
50:j:1:MET:O	50:j:204:LYS:HD2	2.15	0.47
52:l:112:LEU:CD1	52:l:117:ARG:HB2	2.39	0.47
54:n:67:ILE:HD13	54:n:87:CYS:HB3	1.95	0.47
54:n:93:GLY:C	54:n:97:TRP:CZ3	2.92	0.47
58:r:58:LEU:O	58:r:61:VAL:HG22	2.14	0.47
59:s:123:LYS:HD2	59:s:132:HIS:HD2	1.78	0.47
62:v:29:GLY:H	62:v:104:GLU:HG2	1.79	0.47
63:w:56:LYS:NZ	63:w:90:ARG:O	2.48	0.47
4:3:40:ASN:OD1	4:3:65:ILE:HB	2.14	0.47
4:3:84:GLY:HA3	4:3:95:PHE:CZ	2.49	0.47
5:4:59:GLU:OE1	5:4:59:GLU:N	2.47	0.47
7:6:10:DG:OP1	14:AE:311:ARG:NH2	2.47	0.47
9:9:3:LEU:O	9:9:4:ASN:OD1	2.32	0.47
11:AA:318:SER:OG	11:AA:320:ASP:OD1	2.31	0.47
11:AA:363:LEU:HB3	11:AA:381:ALA:HB1	1.96	0.47
12:AB:118:ILE:HG12	12:AB:130:PHE:CD2	2.49	0.47
16:AG:61:ARG:HD2	16:AG:73:LYS:CD	2.44	0.47
16:AG:316:GLN:H	16:AG:316:GLN:HG3	1.54	0.47
16:AG:316:GLN:HE21	16:AG:316:GLN:HB2	1.49	0.47
10:B:70:G:C3'	10:B:71:C:H5''	2.44	0.47
18:D:745:G:H2'	18:D:746:A:C8	2.50	0.47
22:H:302:GLY:HA3	22:H:344:LEU:HD13	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:J:62:ARG:NE	24:J:72:PHE:CE2	2.81	0.47
29:O:21:ILE:HD12	29:O:86:ALA:HB1	1.96	0.47
30:P:10:LEU:C	30:P:11:LYS:HD3	2.40	0.47
31:Q:98:ARG:HG2	31:Q:98:ARG:HH11	1.78	0.47
33:S:20:TYR:CD2	33:S:55:SER:OG	2.67	0.47
34:T:15:PHE:CD2	34:T:30:ALA:HB2	2.49	0.47
34:T:29:VAL:HG22	34:T:66:LEU:HD23	1.96	0.47
41:a:1477:A:N6	41:a:1514:G:O2'	2.45	0.47
41:a:1841:U:C2	41:a:1842:G:C8	3.02	0.47
41:a:2246:G:H2'	41:a:2247:A:C8	2.49	0.47
41:a:2351:G:O6	55:o:39:LYS:HE3	2.13	0.47
41:a:2638:G:O2'	41:a:2775:G:N2	2.37	0.47
45:e:14:LEU:HD13	45:e:57:LEU:HD22	1.95	0.47
46:f:27:LEU:O	46:f:38:ARG:NE	2.44	0.47
47:g:45:THR:O	47:g:49:ARG:CZ	2.62	0.47
56:p:8:PRO:C	56:p:69:ARG:HH21	2.22	0.47
56:p:52:PHE:CZ	56:p:72:LEU:HD22	2.50	0.47
62:v:57:VAL:HG22	62:v:57:VAL:O	2.14	0.47
1:0:68:ARG:NH2	1:0:90:ARG:HD3	2.29	0.47
2:1:86:MET:HE1	2:1:88:ARG:CD	2.42	0.47
5:4:53:LYS:O	5:4:56:PHE:HD2	1.97	0.47
9:9:52:MET:HG2	9:9:95:LEU:CD1	2.39	0.47
9:9:77:VAL:CG1	9:9:82:ILE:HG21	2.41	0.47
9:9:108:VAL:O	9:9:109:LYS:HB3	2.12	0.47
10:A:35:A:H2'	10:A:36:U:C6	2.49	0.47
11:AA:786:GLY:N	11:AA:789:THR:OG1	2.41	0.47
12:AB:111:TYR:HB3	12:AB:112:PRO:HD2	1.96	0.47
14:AE:218:THR:HA	14:AE:221:ILE:HG22	1.96	0.47
17:C:21:ILE:HG13	17:C:54:GLN:NE2	2.28	0.47
18:D:1458:G:H5''	19:E:26:SER:OG	2.14	0.47
22:H:37:LEU:HD11	22:H:47:ALA:HA	1.96	0.47
25:K:96:MET:HG2	25:K:125:ALA:HB2	1.97	0.47
27:M:109:ARG:C	27:M:119:ARG:HH21	2.23	0.47
28:N:34:VAL:HG22	28:N:59:LEU:HD21	1.96	0.47
29:O:12:ARG:HD3	29:O:13:LYS:H	1.78	0.47
29:O:63:LEU:HD13	29:O:65:ILE:HD11	1.97	0.47
32:R:99:ARG:NE	32:R:104:CYS:SG	2.87	0.47
33:S:12:LYS:HD2	33:S:16:LEU:CD2	2.44	0.47
36:V:62:ARG:C	36:V:73:TRP:CE3	2.92	0.47
41:a:67:U:N3	41:a:74:A:N6	2.62	0.47
41:a:111:A:O2'	45:e:58:ASN:ND2	2.46	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:600:G:H2'	41:a:601:C:C6	2.49	0.47
41:a:909:A:H2'	41:a:912:C:H5	1.80	0.47
41:a:1429:G:H2'	41:a:1430:G:H8	1.79	0.47
41:a:2064:C:H2'	41:a:2065:C:H6	1.80	0.47
41:a:2294:G:OP1	64:x:10:ARG:HD3	2.15	0.47
41:a:2345:G:C5	41:a:2347:C:C5	3.03	0.47
51:k:25:LYS:HD3	51:k:51:GLU:OE2	2.14	0.47
62:v:53:MET:HB2	62:v:116:ALA:HB1	1.96	0.47
4:3:28:VAL:HG23	4:3:34:VAL:HG12	1.96	0.47
9:9:38:MET:SD	39:Y:117:THR:CG2	3.02	0.47
9:9:88:HIS:CB	9:9:89:PRO:CD	2.81	0.47
10:A:48:C:C2'	10:A:49:G:OP2	2.62	0.47
11:AA:103:VAL:HG12	11:AA:117:ILE:HG22	1.97	0.47
16:AG:266:LEU:CD2	16:AG:266:LEU:N	2.73	0.47
16:AG:285:ILE:O	16:AG:288:MET:HE3	2.14	0.47
17:C:73:ARG:NH1	31:Q:113:VAL:CG1	2.76	0.47
18:D:126:G:OP1	18:D:605:U:O2'	2.19	0.47
18:D:375:U:H1'	35:U:28:ARG:HH22	1.79	0.47
18:D:1225:A:H2'	18:D:1226:C:C5	2.49	0.47
18:D:1308:U:P	38:X:98:ARG:CG	3.03	0.47
26:L:9:MET:HE2	26:L:51:ILE:CD1	2.44	0.47
27:M:129:GLU:O	27:M:130:ASN:HB2	2.13	0.47
29:O:89:GLU:O	29:O:92:GLU:OE1	2.32	0.47
37:W:63:THR:HG22	37:W:64:ASP:N	2.30	0.47
39:Y:19:PRO:HG2	39:Y:24:GLY:H	1.79	0.47
41:a:963:U:H2'	41:a:964:C:H6	1.79	0.47
41:a:1385:A:C4	41:a:1386:C:C5	3.02	0.47
41:a:1666:G:H1'	60:t:3:GLN:NE2	2.29	0.47
41:a:2297:A:C6	41:a:2321:U:O4	2.68	0.47
41:a:2299:U:H2'	41:a:2300:C:C6	2.49	0.47
41:a:2748:A:C2	41:a:2757:A:C5	3.03	0.47
41:a:2803:G:H2'	41:a:2804:U:C5	2.50	0.47
52:l:141:MET:O	52:l:142:ALA:HB3	2.13	0.47
62:v:30:SER:N	62:v:104:GLU:OE2	2.48	0.47
2:1:23:LEU:HD21	2:1:35:ILE:CG2	2.44	0.47
5:4:21:ARG:HA	5:4:25:LYS:O	2.14	0.47
5:4:77:VAL:HG13	5:4:77:VAL:O	2.13	0.47
11:AA:560:PRO:O	14:AE:780:ARG:NH2	2.47	0.47
14:AE:516:ASP:OD1	14:AE:516:ASP:N	2.47	0.47
17:C:12:ARG:CZ	17:C:13:PHE:CE1	2.97	0.47
17:C:16:GLU:OE1	17:C:18:VAL:HB	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:35:G:H2'	18:D:36:C:H6	1.80	0.47
18:D:436:C:C2	18:D:437:U:C5	3.02	0.47
18:D:642:A:H2'	18:D:643:C:H6	1.80	0.47
18:D:1238:A:H2	18:D:1241:G:N3	2.13	0.47
22:H:70:VAL:HA	22:H:85:SER:HA	1.95	0.47
24:J:62:ARG:HA	24:J:72:PHE:CZ	2.50	0.47
25:K:157:ARG:NH1	28:N:43:GLU:CD	2.72	0.47
28:N:3:MET:O	28:N:3:MET:HE3	2.14	0.47
39:Y:126:ARG:NH2	41:a:1082:U:O3'	2.47	0.47
41:a:1656:C:P	50:j:141:ARG:HD2	2.54	0.47
41:a:1770:G:C6	41:a:1983:G:C5	3.03	0.47
41:a:2040:G:OP1	59:s:106:LYS:NZ	2.38	0.47
41:a:2362:C:OP2	55:o:40:ARG:CZ	2.63	0.47
41:a:2814:A:C4	41:a:2815:C:C5	3.03	0.47
42:b:26:PHE:H	42:b:29:GLU:HG3	1.79	0.47
42:b:45:PHE:HD2	42:b:80:ILE:HD11	1.77	0.47
53:m:25:LYS:HE2	53:m:28:ARG:HH21	1.78	0.47
54:n:12:VAL:O	54:n:16:LEU:HG	2.14	0.47
54:n:38:MET:HE3	54:n:57:LEU:HG	1.96	0.47
61:u:77:ILE:HD11	61:u:108:ALA:HB1	1.95	0.47
64:x:41:ALA:HB2	64:x:48:LEU:HD11	1.95	0.47
2:1:98:LYS:HZ2	41:a:2012:G:P	2.38	0.47
10:A:6:G:O2'	10:A:7:G:H8	1.98	0.47
10:A:30:G:O2'	10:A:31:G:H5'	2.15	0.47
11:AA:207:THR:OG1	11:AA:354:ASP:OD2	2.32	0.47
11:AA:689:ALA:HB2	11:AA:1233:LEU:HD23	1.97	0.47
11:AA:855:PRO:HG2	16:AG:105:ILE:HG22	1.94	0.47
12:AB:50:LEU:HD21	14:AE:285:LEU:CB	2.44	0.47
14:AE:74:LYS:HB2	14:AE:74:LYS:HE2	1.59	0.47
14:AE:77:ARG:HA	14:AE:77:ARG:CZ	2.44	0.47
14:AE:290:ILE:O	14:AE:290:ILE:HG12	2.14	0.47
14:AE:291:ILE:O	14:AE:293:ARG:N	2.46	0.47
14:AE:616:PRO:HA	14:AE:619:ILE:HG22	1.96	0.47
14:AE:804:ALA:O	14:AE:806:ASP:N	2.47	0.47
14:AE:1350:ASN:HA	14:AE:1353:VAL:HG12	1.97	0.47
16:AG:200:SER:C	16:AG:230:PRO:HG2	2.38	0.47
16:AG:233:ARG:O	16:AG:327:LEU:CD2	2.62	0.47
16:AG:243:LYS:O	16:AG:243:LYS:HD3	2.14	0.47
16:AG:363:LEU:HD11	16:AG:410:ALA:HB2	1.92	0.47
16:AG:437:LEU:CG	16:AG:456:LEU:CD1	2.92	0.47
18:D:72:A:C5	18:D:73:C:C5	3.02	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:106:C:N4	19:E:10:ARG:HH22	2.12	0.47
18:D:110:C:H2'	18:D:111:G:O4'	2.14	0.47
18:D:197:A:C6	18:D:221:C:H4'	2.50	0.47
18:D:216:U:H2'	18:D:217:C:C6	2.50	0.47
18:D:520:A:O2'	32:R:70:GLU:CD	2.58	0.47
18:D:653:U:N1	28:N:56:LYS:HE3	2.29	0.47
18:D:657:U:C1'	34:T:28:GLN:HE22	2.24	0.47
18:D:687:A:C2	18:D:704:A:C4	3.02	0.47
18:D:864:A:H5'	25:K:90:THR:O	2.14	0.47
18:D:1149:C:C2	18:D:1150:A:C8	3.03	0.47
18:D:1172:C:H2'	18:D:1173:U:C6	2.49	0.47
18:D:1325:C:C2	18:D:1326:U:C5	3.02	0.47
18:D:1363:A:C5	18:D:1365:G:C6	3.03	0.47
21:G:26:LYS:O	21:G:26:LYS:HD3	2.14	0.47
21:G:207:ILE:HG23	21:G:208:ARG:N	2.30	0.47
23:I:25:ASN:O	23:I:27:LYS:N	2.47	0.47
27:M:116:MET:HA	27:M:119:ARG:HD3	1.97	0.47
27:M:135:VAL:HG23	27:M:136:LYS:HD2	1.96	0.47
29:O:7:TYR:O	29:O:89:GLU:CD	2.58	0.47
31:Q:116:ILE:HG13	31:Q:116:ILE:O	2.14	0.47
33:S:63:ARG:HD3	33:S:68:GLY:O	2.15	0.47
36:V:30:LYS:HD3	36:V:35:GLY:HA2	1.97	0.47
39:Y:102:ARG:HE	39:Y:139:VAL:HG21	1.79	0.47
41:a:78:U:OP1	45:e:4:LYS:CE	2.63	0.47
41:a:272:A:C2	41:a:273:G:C5	3.02	0.47
41:a:592:A:C2	55:o:4:ILE:HD11	2.50	0.47
41:a:689:A:H2'	41:a:690:G:C8	2.50	0.47
41:a:1278:C:H2'	41:a:1279:G:H8	1.80	0.47
41:a:1333:G:C2	41:a:1334:G:C8	3.03	0.47
41:a:1385:A:O2'	41:a:1396:U:O2	2.31	0.47
41:a:1441:G:H2'	41:a:1442:U:C6	2.50	0.47
41:a:1467:U:C4	41:a:1546:G:C2	3.03	0.47
41:a:1791:A:C2	41:a:1829:A:H4'	2.49	0.47
41:a:2236:U:H2'	41:a:2237:G:O4'	2.15	0.47
41:a:2243:U:H2'	41:a:2244:U:H6	1.80	0.47
41:a:2520:C:C6	41:a:2567:G:H1'	2.50	0.47
41:a:2771:C:C2	41:a:2772:C:C5	3.01	0.47
45:e:20:ASN:HA	45:e:23:ARG:NE	2.30	0.47
48:h:130:LEU:HG	48:h:135:ILE:HG13	1.96	0.47
50:j:91:THR:HB	50:j:94:GLN:CG	2.44	0.47
52:l:148:ILE:CD1	52:l:187:VAL:HB	2.45	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:l:168:ASP:HB3	52:l:183:PHE:CE2	2.49	0.47
56:p:18:LYS:HE2	56:p:18:LYS:HA	1.96	0.47
58:r:135:HIS:CE1	58:r:137:GLU:OE1	2.62	0.47
62:v:41:LEU:HG	62:v:96:ILE:CG1	2.45	0.47
63:w:44:LEU:C	63:w:44:LEU:HD23	2.40	0.47
66:z:69:ALA:HA	66:z:106:PHE:HZ	1.80	0.47
66:z:69:ALA:O	66:z:74:ILE:HG22	2.15	0.47
9:9:69:PHE:C	9:9:72:LEU:HD11	2.40	0.47
9:9:99:PHE:O	9:9:103:ASN:OD1	2.32	0.47
10:A:33:U:H4'	27:M:84:THR:HB	1.96	0.47
10:A:68:C:H3'	10:A:69:C:H5''	1.97	0.47
10:A:72:A:H2'	10:A:73:A:H5''	1.95	0.47
12:AB:145:LEU:HD22	12:AB:148:LYS:CD	2.44	0.47
13:AD:59:VAL:HG22	13:AD:144:ILE:HA	1.97	0.47
14:AE:1108:GLN:HE21	14:AE:1123:ARG:HH12	1.62	0.47
16:AG:1:MET:SD	16:AG:53:SER:HA	2.55	0.47
16:AG:314:LEU:HB3	16:AG:318:ILE:HD12	1.97	0.47
16:AG:479:GLY:O	16:AG:483:MET:HG2	2.15	0.47
10:B:30:G:O2'	10:B:31:G:H5'	2.15	0.47
18:D:983:A:H5'	18:D:984:C:OP2	2.15	0.47
21:G:19:GLN:NE2	22:H:75:VAL:HG13	2.30	0.47
22:H:115:LYS:HA	22:H:119:GLY:C	2.40	0.47
23:I:181:ASP:HB2	23:I:207:ILE:HD11	1.97	0.47
25:K:57:PRO:HA	25:K:60:ILE:HG12	1.96	0.47
27:M:111:ARG:O	27:M:119:ARG:CZ	2.63	0.47
28:N:112:THR:HG23	28:N:115:ALA:H	1.80	0.47
29:O:127:PHE:CD1	29:O:128:SER:N	2.83	0.47
32:R:2:ALA:CA	32:R:7:LEU:HD11	2.45	0.47
39:Y:52:LEU:O	39:Y:54:ILE:HG12	2.14	0.47
41:a:872:U:H2'	41:a:873:C:C6	2.50	0.47
41:a:1818:U:H6	48:h:156:ARG:NH2	2.13	0.47
41:a:2250:G:OP1	62:v:84:LYS:NZ	2.48	0.47
41:a:2294:G:OP1	64:x:98:GLN:OE1	2.33	0.47
41:a:2299:U:P	54:n:72:LYS:HE2	2.54	0.47
41:a:2513:A:H2'	41:a:2514:U:C6	2.50	0.47
41:a:2537:U:H2'	41:a:2538:C:H6	1.79	0.47
48:h:181:MET:HE2	48:h:181:MET:CA	2.45	0.47
49:i:55:ILE:HD13	63:w:112:TYR:CE1	2.50	0.47
52:l:42:GLY:HA3	52:l:92:HIS:CE1	2.50	0.47
52:l:130:LYS:HB2	52:l:133:LEU:HD12	1.97	0.47
62:v:31:PHE:O	62:v:104:GLU:OE1	2.33	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:73:LYS:HE2	1:0:73:LYS:HA	1.95	0.47
2:1:11:ARG:CZ	2:1:11:ARG:HB3	2.45	0.47
3:2:57:VAL:HG12	3:2:86:THR:OG1	2.15	0.47
5:4:11:GLU:OE2	5:4:19:ARG:NH2	2.47	0.47
10:A:19:G:C6	41:a:2112:G:O5'	2.67	0.47
11:AA:836:LEU:HD13	11:AA:1054:LEU:HD13	1.95	0.47
11:AA:848:GLU:HG2	11:AA:888:THR:HG22	1.96	0.47
11:AA:1311:GLY:O	15:AF:31:GLN:NE2	2.47	0.47
12:AB:7:LEU:HD22	12:AB:78:PHE:HA	1.96	0.47
12:AB:116:VAL:N	12:AB:129:ILE:CD1	2.78	0.47
14:AE:81:ARG:NH2	14:AE:81:ARG:CG	2.77	0.47
16:AG:434:LEU:CD1	16:AG:455:THR:O	2.62	0.47
10:B:35:A:H2'	10:B:36:U:C6	2.49	0.47
10:B:75:C:O2'	10:B:76:A:N7	2.48	0.47
18:D:61:G:O6	19:E:10:ARG:NH2	2.47	0.47
18:D:307:C:O2	18:D:307:C:H2'	2.15	0.47
18:D:309:A:O2'	18:D:607:A:N1	2.45	0.47
18:D:739:C:O2'	34:T:42:HIS:CE1	2.67	0.47
18:D:1308:U:H3'	38:X:98:ARG:NH2	2.29	0.47
22:H:277:GLY:CA	22:H:331:MET:CE	2.93	0.47
23:I:132:ARG:CA	23:I:135:LYS:HE3	2.44	0.47
27:M:95:ARG:CZ	27:M:99:LEU:HD21	2.44	0.47
34:T:48:LYS:O	34:T:50:HIS:N	2.48	0.47
37:W:9:PRO:HD3	47:g:64:PHE:CD2	2.49	0.47
38:X:5:ALA:HB3	38:X:66:GLU:OE2	2.14	0.47
39:Y:102:ARG:HH11	39:Y:141:ASP:CG	2.23	0.47
41:a:104:A:C5	41:a:105:C:C5	3.02	0.47
41:a:1421:G:C2	41:a:1422:G:C8	3.03	0.47
41:a:1794:A:H2'	41:a:1795:C:H6	1.79	0.47
41:a:1936:A:N7	41:a:1945:G:C8	2.83	0.47
41:a:2298:A:H5''	54:n:72:LYS:CE	2.44	0.47
41:a:2351:G:O6	55:o:39:LYS:NZ	2.48	0.47
41:a:2517:C:O2	41:a:2542:A:N6	2.48	0.47
41:a:2700:A:H2'	41:a:2701:U:C6	2.50	0.47
46:f:41:THR:O	46:f:42:PRO:C	2.58	0.47
50:j:9:VAL:O	50:j:197:THR:HG23	2.15	0.47
52:l:42:GLY:HA3	52:l:92:HIS:HE1	1.80	0.47
62:v:50:ARG:CA	62:v:53:MET:SD	3.00	0.47
8:7:8:U:OP2	18:D:1397:C:O2	2.33	0.47
9:9:18:VAL:CG1	9:9:71:CYS:HB2	2.45	0.47
9:9:40:GLU:CD	9:9:54:VAL:HG11	2.39	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:887:VAL:CA	16:AG:105:ILE:CD1	2.57	0.47
12:AB:140:MET:HG2	30:P:102:LEU:O	2.15	0.47
13:AC:102:LEU:HD12	13:AC:115:ILE:HG12	1.96	0.47
13:AD:182:ARG:HD2	14:AE:581:MET:HE1	1.97	0.47
16:AG:33:THR:CG2	16:AG:43:ILE:HD11	2.44	0.47
18:D:306:A:C5	18:D:307:C:C5	3.02	0.47
18:D:520:A:O3'	32:R:70:GLU:OE2	2.32	0.47
18:D:977:A:H1'	18:D:982:U:O4	2.15	0.47
20:F:12:PHE:CE1	31:Q:107:ILE:CG2	2.98	0.47
21:G:117:LEU:HD21	21:G:141:LEU:HB2	1.97	0.47
25:K:157:ARG:NH1	28:N:43:GLU:HA	2.29	0.47
25:K:165:LEU:HD12	28:N:114:ARG:NH1	2.30	0.47
32:R:2:ALA:N	32:R:7:LEU:HD11	2.30	0.47
32:R:5:ASN:HB3	32:R:9:ARG:HH22	1.80	0.47
33:S:63:ARG:NH2	33:S:70:PRO:HD3	2.30	0.47
34:T:77:ARG:NH2	34:T:80:GLN:NE2	2.62	0.47
38:X:13:LYS:HZ2	38:X:17:ILE:HG21	1.80	0.47
41:a:247:G:H4'	41:a:386:G:C5	2.49	0.47
41:a:396:G:C5	41:a:397:U:C5	3.03	0.47
41:a:654:A:N6	55:o:19:LYS:HD2	2.29	0.47
41:a:666:A:H4'	61:u:48:ARG:HD2	1.97	0.47
41:a:783:A:N3	41:a:783:A:H2'	2.29	0.47
41:a:1068:G:O6	41:a:1069:A:N6	2.48	0.47
41:a:1442:U:H2'	41:a:1443:U:H6	1.79	0.47
41:a:1794:A:H2'	41:a:1795:C:C6	2.50	0.47
41:a:2197:U:C5	41:a:2224:G:C6	3.04	0.47
41:a:2364:C:H2'	41:a:2365:G:O4'	2.15	0.47
41:a:2646:C:O5'	41:a:2646:C:H6	1.98	0.47
41:a:2725:A:C4	41:a:2727:A:N7	2.83	0.47
45:e:10:SER:OG	45:e:12:GLU:HG2	2.15	0.47
46:f:12:SER:C	46:f:54:MET:HE1	2.40	0.47
55:o:31:HIS:O	55:o:32:ILE:O	2.33	0.47
60:t:1:MET:CE	60:t:32:TYR:CG	2.98	0.47
2:1:11:ARG:NH1	2:1:11:ARG:HB3	2.30	0.46
3:2:1:MET:HE1	41:a:139:U:C4	2.50	0.46
3:2:5:GLU:HA	45:e:22:LEU:CD2	2.45	0.46
8:7:47:U:H3'	8:7:47:U:H6	1.80	0.46
11:AA:400:VAL:HG22	11:AA:584:TYR:HB3	1.96	0.46
12:AB:104:ILE:C	12:AB:106:ASP:N	2.72	0.46
12:AB:138:ARG:CD	12:AB:155:LYS:HB2	2.45	0.46
14:AE:81:ARG:CZ	16:AG:144:LYS:HE2	2.45	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:287:ALA:CB	14:AE:292:VAL:CG2	2.93	0.46
14:AE:437:PHE:HZ	14:AE:453:VAL:HG11	1.79	0.46
16:AG:233:ARG:HG2	16:AG:270:ARG:CB	2.45	0.46
16:AG:353:HIS:O	16:AG:356:ILE:CA	2.63	0.46
10:B:48:C:O2'	10:B:49:G:OP2	2.27	0.46
10:B:70:G:C2'	10:B:71:C:H5''	2.45	0.46
17:C:12:ARG:NH1	17:C:13:PHE:CE1	2.83	0.46
18:D:22:G:C5	18:D:23:C:C5	3.03	0.46
18:D:610:U:N3	18:D:611:C:C5	2.83	0.46
20:F:33:ARG:O	20:F:36:GLU:HG2	2.14	0.46
21:G:131:LYS:HE3	21:G:132:LYS:NZ	2.29	0.46
22:H:303:LEU:C	22:H:305:HIS:N	2.73	0.46
24:J:41:HIS:ND1	24:J:44:ARG:NH1	2.63	0.46
26:L:17:GLN:C	26:L:21:MET:HE1	2.40	0.46
29:O:86:ALA:HA	29:O:89:GLU:HG2	1.95	0.46
30:P:27:GLU:HA	30:P:30:LYS:HZ1	1.80	0.46
39:Y:20:SER:CB	39:Y:21:PRO:HD3	2.43	0.46
41:a:257:C:O2	61:u:104:GLN:NE2	2.48	0.46
41:a:609:A:H3'	41:a:610:C:H6	1.80	0.46
41:a:1200:C:C2	41:a:1201:U:C5	3.03	0.46
41:a:2702:G:C5	41:a:2703:C:C5	3.03	0.46
42:b:26:PHE:O	42:b:29:GLU:HG2	2.15	0.46
48:h:2:ALA:N	48:h:20:VAL:O	2.48	0.46
49:i:4:GLN:OE1	49:i:8:PRO:CD	2.63	0.46
50:j:74:GLU:HG2	50:j:75:ALA:N	2.31	0.46
52:l:6:LYS:HZ1	52:l:120:VAL:N	2.11	0.46
58:r:31:VAL:HB	58:r:32:PRO:HD3	1.97	0.46
59:s:72:LYS:HD3	59:s:74:TYR:CZ	2.43	0.46
62:v:59:ARG:CZ	62:v:59:ARG:O	2.63	0.46
66:z:108:ALA:O	66:z:112:LYS:HG2	2.14	0.46
1:0:71:LYS:HE2	1:0:90:ARG:HD2	1.97	0.46
2:1:105:VAL:HG13	2:1:105:VAL:O	2.14	0.46
9:9:24:SER:HB3	9:9:116:GLU:HG3	1.94	0.46
9:9:51:TYR:CD2	9:9:88:HIS:HB2	2.49	0.46
16:AG:36:LYS:HB2	16:AG:101:THR:HG21	1.97	0.46
16:AG:400:GLU:H	16:AG:400:GLU:HG2	1.40	0.46
10:B:24:U:H2'	10:B:25:C:C6	2.51	0.46
10:B:46:G:H1'	10:B:47:U:C5	2.51	0.46
10:B:48:C:C2'	10:B:49:G:OP2	2.62	0.46
17:C:31:ASN:OD1	22:H:268:VAL:HG12	2.15	0.46
18:D:1324:A:H2'	18:D:1325:C:C6	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1412:C:H2'	18:D:1413:A:C8	2.49	0.46
30:P:13:PHE:CZ	30:P:67:ILE:HD11	2.50	0.46
30:P:13:PHE:CE1	30:P:67:ILE:HD11	2.50	0.46
41:a:493:G:H2'	41:a:494:G:O4'	2.16	0.46
41:a:644:A:H2'	41:a:645:C:O4'	2.15	0.46
41:a:1006:C:C2	41:a:1138:G:N2	2.83	0.46
41:a:1432:G:H2'	41:a:1433:A:H8	1.80	0.46
41:a:2013:A:C2	41:a:2613:U:O4	2.68	0.46
41:a:2200:C:OP2	43:c:37:ARG:NH2	2.48	0.46
41:a:2314:A:P	54:n:88:LYS:NZ	2.87	0.46
41:a:2419:U:O2'	41:a:2420:C:H5'	2.14	0.46
41:a:2454:G:C6	41:a:2499:C:N4	2.83	0.46
41:a:2683:C:OP1	65:y:51:ARG:NH2	2.46	0.46
48:h:108:LYS:HZ3	48:h:194:GLU:N	2.13	0.46
52:l:175:ILE:HG21	52:l:180:LEU:HD11	1.97	0.46
56:p:155:GLU:O	56:p:159:GLY:HA2	2.15	0.46
59:s:123:LYS:HD2	59:s:132:HIS:CD2	2.51	0.46
1:0:4:VAL:HG23	1:0:39:LEU:HB2	1.97	0.46
2:1:50:VAL:HG12	2:1:105:VAL:CG1	2.45	0.46
3:2:64:LYS:HD2	41:a:1312:U:C4	2.51	0.46
3:2:69:ARG:HG3	3:2:74:ILE:HD13	1.96	0.46
4:3:96:PHE:CE1	4:3:103:ILE:HG12	2.50	0.46
5:4:48:MET:HE3	5:4:85:LYS:HA	1.97	0.46
10:A:19:G:O6	41:a:2112:G:O5'	2.34	0.46
10:A:24:U:H2'	10:A:25:C:C6	2.51	0.46
10:A:70:G:C2'	10:A:71:C:H5''	2.45	0.46
11:AA:889:PRO:HA	16:AG:104:ARG:O	2.15	0.46
12:AB:120:GLU:H	12:AB:162:LEU:HG	1.80	0.46
16:AG:413:ALA:C	16:AG:416:THR:HG1	2.15	0.46
18:D:611:C:C2	18:D:612:C:C5	3.02	0.46
18:D:636:U:H5'	36:V:6:ARG:NE	2.29	0.46
18:D:915:A:C5	18:D:916:U:C6	3.04	0.46
18:D:1382:C:N3	18:D:1383:C:C5	2.83	0.46
18:D:1435:G:H2'	18:D:1436:U:C6	2.51	0.46
21:G:19:GLN:OE1	21:G:21:ARG:CG	2.63	0.46
21:G:110:SER:CA	21:G:113:ARG:HH21	2.27	0.46
23:I:22:TRP:CZ2	23:I:32:ASN:HB2	2.50	0.46
23:I:107:ARG:HB3	23:I:108:LYS:HD2	1.96	0.46
23:I:134:MET:HE1	23:I:167:TRP:CA	2.43	0.46
26:L:38:ARG:NE	26:L:97:THR:O	2.49	0.46
27:M:74:GLU:CG	27:M:91:VAL:CG1	2.94	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:P:59:LYS:HE2	30:P:62:ARG:NH2	2.31	0.46
34:T:76:ALA:C	34:T:80:GLN:NE2	2.68	0.46
36:V:11:ARG:HD2	36:V:56:GLY:C	2.39	0.46
37:W:32:ARG:NH2	37:W:34:TRP:CZ3	2.84	0.46
39:Y:5:GLN:NE2	39:Y:60:VAL:C	2.74	0.46
41:a:626:A:C2	61:u:78:ARG:CZ	2.98	0.46
41:a:1349:C:N3	41:a:1350:C:C5	2.83	0.46
41:a:1688:U:O2'	41:a:1700:A:N7	2.44	0.46
41:a:2616:C:N3	41:a:2617:U:C5	2.83	0.46
41:a:2784:U:H2'	41:a:2785:C:H6	1.80	0.46
47:g:47:LYS:HE2	54:n:115:ARG:HA	1.95	0.46
52:l:1:MET:CE	52:l:20:GLY:N	2.79	0.46
52:l:23:PHE:CE2	52:l:25:GLU:HA	2.50	0.46
59:s:25:LEU:HD11	59:s:101:ILE:HD13	1.97	0.46
60:t:76:VAL:HG12	65:y:73:VAL:HB	1.97	0.46
62:v:41:LEU:HD22	62:v:45:GLN:HE22	1.78	0.46
1:0:4:VAL:CG2	1:0:40:MET:HB3	2.45	0.46
9:9:25:ALA:HB3	9:9:99:PHE:CE2	2.50	0.46
9:9:97:LYS:HZ1	9:9:130:PRO:HA	1.79	0.46
10:A:52:G:C6	10:A:63:G:O6	2.68	0.46
11:AA:28:LEU:HD21	11:AA:524:ILE:HG13	1.97	0.46
14:AE:66:LYS:HB2	14:AE:69:GLU:HB3	1.97	0.46
17:C:60:LYS:HE3	18:D:735:C:H4'	1.97	0.46
18:D:1189:U:OP1	33:S:98:LYS:NZ	2.48	0.46
20:F:19:PHE:HE2	31:Q:110:ILE:C	2.23	0.46
25:K:143:GLY:O	25:K:146:ASN:OD1	2.34	0.46
33:S:6:MET:HE3	33:S:63:ARG:NH1	2.31	0.46
34:T:15:PHE:CE2	34:T:30:ALA:HB1	2.51	0.46
36:V:61:ILE:C	36:V:73:TRP:HZ3	2.23	0.46
38:X:56:LEU:HA	38:X:59:GLU:OE1	2.16	0.46
41:a:374:A:C2	41:a:375:G:H1'	2.51	0.46
41:a:392:U:C2	41:a:393:C:C5	3.03	0.46
41:a:921:C:C2	41:a:922:C:C5	3.03	0.46
42:b:51:VAL:HG12	42:b:59:LEU:HD12	1.96	0.46
50:j:71:ALA:HB3	50:j:73:VAL:HG22	1.98	0.46
55:o:55:LEU:CD2	55:o:59:ILE:HD11	2.45	0.46
56:p:94:TYR:HD1	56:p:107:LEU:HD23	1.79	0.46
60:t:24:VAL:HG23	60:t:33:ALA:HB2	1.97	0.46
61:u:19:LEU:CD2	61:u:27:LEU:HD12	2.45	0.46
64:x:35:ILE:HD11	64:x:102:ARG:HH21	1.81	0.46
1:0:82:HIS:CD2	61:u:23:ILE:HG12	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4:23:ALA:O	5:4:24:ASN:OD1	2.33	0.46
9:9:59:LEU:HB2	9:9:62:ARG:CG	2.45	0.46
11:AA:1314:GLN:HA	15:AF:28:ARG:HH22	1.80	0.46
12:AB:148:LYS:HB3	30:P:100:ILE:HG23	1.96	0.46
17:C:65:LEU:HG	17:C:65:LEU:O	2.16	0.46
18:D:277:C:OP1	36:V:43:LYS:HD2	2.15	0.46
18:D:737:C:C2	18:D:738:C:C5	3.03	0.46
18:D:1132:C:O2	18:D:1132:C:H2'	2.16	0.46
18:D:1152:A:OP1	30:P:72:ARG:NH1	2.43	0.46
18:D:1512:U:H2'	18:D:1513:A:C8	2.51	0.46
22:H:288:THR:O	22:H:288:THR:HG22	2.16	0.46
25:K:86:LYS:HE3	25:K:95:PHE:CD1	2.50	0.46
26:L:96:VAL:O	26:L:96:VAL:HG23	2.16	0.46
26:L:100:SER:HB2	26:L:101:PRO:HD2	1.98	0.46
27:M:69:VAL:HG23	27:M:100:ALA:HB1	1.97	0.46
28:N:8:ALA:N	28:N:77:ARG:HH11	2.14	0.46
28:N:114:ARG:HD2	28:N:114:ARG:C	2.41	0.46
31:Q:20:VAL:HB	31:Q:35:THR:CG2	2.45	0.46
36:V:7:THR:CG2	36:V:60:GLU:OE2	2.63	0.46
39:Y:37:PHE:HZ	39:Y:56:VAL:HG11	1.79	0.46
39:Y:102:ARG:HD3	39:Y:141:ASP:OD1	2.15	0.46
41:a:6:A:N3	59:s:135:GLN:NE2	2.59	0.46
41:a:591:U:C2	41:a:592:A:C8	3.03	0.46
41:a:1585:C:H2'	41:a:1586:A:O4'	2.15	0.46
41:a:1598:A:C4	41:a:1599:U:C6	3.03	0.46
41:a:1790:C:O3'	41:a:1791:A:C8	2.68	0.46
41:a:2357:G:P	42:b:20:ARG:CZ	3.04	0.46
43:c:71:LEU:O	43:c:76:GLU:HG3	2.16	0.46
48:h:52:ARG:NE	48:h:53:HIS:CE1	2.83	0.46
49:i:52:ARG:HA	49:i:52:ARG:NE	2.31	0.46
52:l:108:ILE:HD11	52:l:180:LEU:HD13	1.97	0.46
58:r:137:GLU:CG	58:r:138:VAL:HG23	2.39	0.46
60:t:8:LEU:N	60:t:8:LEU:HD12	2.31	0.46
65:y:14:LYS:NZ	65:y:76:THR:O	2.49	0.46
66:z:113:ALA:O	66:z:117:LEU:HD13	2.15	0.46
4:3:47:LYS:NZ	41:a:482:A:O5'	2.48	0.46
5:4:2:PHE:CE1	5:4:56:PHE:CZ	3.03	0.46
9:9:94:ARG:CB	9:9:97:LYS:CE	2.94	0.46
10:A:47:U:H2'	10:A:48:C:H4'	1.97	0.46
10:A:65:C:C2'	10:A:66:C:H5'	2.44	0.46
10:A:75:C:O2'	10:A:76:A:N7	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:86:GLU:N	14:AE:86:GLU:CD	2.73	0.46
15:AF:25:ARG:NH1	15:AF:61:ASN:OD1	2.49	0.46
16:AG:183:LEU:CD2	16:AG:197:VAL:HG22	2.45	0.46
16:AG:228:ARG:HB3	16:AG:234:ALA:HA	1.98	0.46
10:B:52:G:C6	10:B:63:G:O6	2.68	0.46
18:D:280:C:O2	36:V:40:ARG:NE	2.49	0.46
18:D:413:G:N2	18:D:428:G:H1'	2.31	0.46
18:D:771:G:C6	18:D:809:G:C6	3.03	0.46
18:D:821:G:H2'	18:D:822:U:H6	1.79	0.46
18:D:829:G:H4'	21:G:23:TRP:CH2	2.51	0.46
18:D:1226:C:H4'	18:D:1227:A:OP1	2.16	0.46
18:D:1513:A:H2'	18:D:1514:G:C8	2.51	0.46
19:E:54:MET:HE1	19:E:75:HIS:HE1	1.77	0.46
21:G:110:SER:HA	21:G:113:ARG:NE	2.29	0.46
22:H:270:ILE:HG23	22:H:270:ILE:O	2.15	0.46
26:L:29:ILE:HG21	26:L:64:VAL:HG13	1.97	0.46
33:S:79:LEU:HB2	33:S:84:VAL:CG2	2.45	0.46
38:X:3:ARG:HB3	47:g:35:ASP:OD2	2.15	0.46
39:Y:79:LEU:O	39:Y:83:ALA:HB3	2.16	0.46
40:Z:2:ILE:CG2	40:Z:5:ASP:H	2.28	0.46
41:a:6:A:H5'	59:s:131:ASN:OD1	2.15	0.46
41:a:15:G:O2'	49:i:15:MET:HE1	2.16	0.46
41:a:306:U:H2'	41:a:307:G:O4'	2.16	0.46
41:a:391:A:C5	41:a:392:U:C5	3.04	0.46
41:a:753:A:C6	41:a:754:U:C5	3.04	0.46
41:a:1406:U:O2'	41:a:1407:G:P	2.74	0.46
41:a:1499:C:C2	41:a:1500:G:C8	3.03	0.46
41:a:2755:C:H3'	57:q:19:ARG:NH2	2.31	0.46
43:c:76:GLU:OE1	43:c:77:LYS:O	2.34	0.46
45:e:12:GLU:CG	45:e:13:GLU:N	2.78	0.46
47:g:5:ILE:CD1	54:n:64:LYS:CE	2.92	0.46
49:i:40:ARG:HH12	49:i:41:HIS:CD2	2.34	0.46
59:s:55:ILE:HG13	59:s:123:LYS:CE	2.46	0.46
3:2:3:ARG:CZ	3:2:5:GLU:OE1	2.64	0.46
4:3:66:GLN:OE1	41:a:329:G:OP2	2.34	0.46
5:4:53:LYS:HD2	5:4:56:PHE:HE2	1.77	0.46
9:9:38:MET:HA	9:9:41:LEU:HD12	1.96	0.46
11:AA:380:ALA:HA	12:AB:65:HIS:ND1	2.28	0.46
12:AB:6:LEU:HD13	14:AE:291:ILE:CG1	2.46	0.46
16:AG:362:TYR:HE1	16:AG:414:LEU:HD21	1.80	0.46
17:C:14:THR:HG21	17:C:48:ARG:CD	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:C:21:ILE:CD1	17:C:54:GLN:NE2	2.79	0.46
18:D:110:C:HI'	35:U:25:ARG:NH1	2.31	0.46
18:D:915:A:C6	18:D:916:U:C5	3.04	0.46
18:D:1209:C:C2	18:D:1210:C:C5	3.04	0.46
20:F:58:LYS:O	20:F:62:ARG:NE	2.45	0.46
24:J:138:SER:O	24:J:139:PRO:C	2.57	0.46
30:P:63:ASP:OD2	30:P:63:ASP:C	2.59	0.46
36:V:15:ASP:OD1	36:V:15:ASP:N	2.49	0.46
39:Y:59:THR:C	39:Y:66:PHE:HB2	2.41	0.46
41:a:2743:U:O2'	56:p:153:ARG:NH1	2.49	0.46
41:a:2840:C:C2	41:a:2841:C:C5	3.03	0.46
47:g:18:CYS:HB2	47:g:40:CYS:HB3	1.96	0.46
49:i:4:GLN:OE1	49:i:7:LYS:HA	2.16	0.46
52:l:59:PRO:HB2	52:l:60:TRP:CE3	2.50	0.46
52:l:60:TRP:CD1	52:l:69:ARG:HH12	2.33	0.46
52:l:195:GLN:CA	52:l:198:GLU:OE1	2.64	0.46
57:q:1:MET:HE3	57:q:2:LYS:N	2.31	0.46
63:w:24:MET:HG3	63:w:44:LEU:HD12	1.97	0.46
2:l:86:MET:SD	2:l:88:ARG:HD2	2.56	0.46
6:5:10:DT:OP1	11:AA:476:LYS:HD2	2.16	0.46
11:AA:230:PHE:HB2	11:AA:333:ILE:HB	1.97	0.46
11:AA:812:PHE:HZ	14:AE:503:SER:HB2	1.81	0.46
13:AC:64:VAL:HG11	13:AC:78:ILE:HG21	1.97	0.46
16:AG:61:ARG:HA	16:AG:94:GLU:HA	1.97	0.46
16:AG:138:ILE:HD12	16:AG:183:LEU:CD1	2.45	0.46
16:AG:430:PRO:CG	16:AG:435:LEU:CD2	2.85	0.46
10:B:32:C:C2'	10:B:33:U:H5'	2.46	0.46
17:C:12:ARG:NH2	17:C:28:THR:CG2	2.79	0.46
17:C:63:ARG:HB3	17:C:70:TYR:CE1	2.51	0.46
18:D:151:A:C5	18:D:152:A:C8	3.04	0.46
18:D:440:C:C2	18:D:441:A:C8	3.04	0.46
18:D:765:G:C6	18:D:812:G:C4	3.04	0.46
18:D:816:A:OP1	18:D:1526:G:O2'	2.27	0.46
21:G:187:VAL:N	21:G:200:ILE:O	2.40	0.46
27:M:134:ALA:HA	27:M:137:LYS:CD	2.46	0.46
29:O:58:VAL:HG23	29:O:59:GLU:N	2.31	0.46
34:T:3:LEU:HD23	34:T:4:SER:O	2.15	0.46
34:T:11:ILE:HG21	34:T:31:LEU:CD1	2.45	0.46
35:U:73:ALA:HA	35:U:76:LYS:HE3	1.98	0.46
37:W:77:THR:OG1	37:W:78:ARG:HD2	2.15	0.46
41:a:1019:U:O4	41:a:1142:A:C2	2.69	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1131:G:OP1	59:s:81:ILE:O	2.34	0.46
41:a:1444:G:C4	41:a:1445:G:C8	3.04	0.46
41:a:1613:G:O2'	53:m:3:ARG:CZ	2.64	0.46
41:a:2420:C:C5'	51:k:8:LYS:NZ	2.76	0.46
42:b:53:CYS:SG	42:b:57:HIS:HA	2.55	0.46
58:r:18:GLN:OE1	58:r:39:ALA:HB1	2.15	0.46
62:v:41:LEU:HD22	62:v:124:LEU:HD22	1.97	0.46
62:v:97:GLN:H	62:v:100:LYS:NZ	2.13	0.46
62:v:114:ARG:C	62:v:118:LYS:HZ3	2.23	0.46
1:0:1:MET:O	1:0:15:SER:OG	2.19	0.46
3:2:64:LYS:CE	41:a:1312:U:O4	2.64	0.46
5:4:53:LYS:HD2	5:4:56:PHE:CZ	2.50	0.46
9:9:11:ILE:HG12	9:9:65:GLU:HG2	1.98	0.46
9:9:61:ARG:HE	41:a:1046:A:P	2.39	0.46
12:AB:51:PHE:CZ	14:AE:288:PRO:HG3	2.47	0.46
16:AG:362:TYR:HA	16:AG:413:ALA:HB3	1.94	0.46
10:B:75:C:P	41:a:2602:A:H2'	2.55	0.46
18:D:13:U:C4	18:D:21:G:C2	3.03	0.46
18:D:13:U:O2	18:D:915:A:N7	2.49	0.46
18:D:371:A:H2'	18:D:372:C:O4'	2.15	0.46
18:D:377:G:OP1	35:U:5:ARG:CZ	2.60	0.46
18:D:1006:G:H2'	18:D:1007:U:H6	1.81	0.46
18:D:1368:A:OP1	29:O:113:ARG:NH2	2.48	0.46
19:E:54:MET:O	19:E:58:VAL:HG23	2.16	0.46
19:E:82:GLN:HA	19:E:85:LYS:NZ	2.31	0.46
21:G:197:ASP:OD2	21:G:198:PHE:CE2	2.68	0.46
22:H:297:GLU:OE1	22:H:297:GLU:N	2.49	0.46
23:I:142:MET:HE3	23:I:170:GLU:HB3	1.98	0.46
23:I:183:ASP:HB3	23:I:204:LYS:HZ2	1.80	0.46
27:M:87:VAL:HG12	27:M:151:PHE:HD2	1.81	0.46
33:S:77:PHE:CE1	33:S:93:ILE:HD13	2.51	0.46
35:U:45:GLU:HG3	35:U:46:LYS:CD	2.45	0.46
37:W:9:PRO:HB3	47:g:64:PHE:CZ	2.51	0.46
39:Y:99:LYS:NZ	39:Y:138:VAL:O	2.44	0.46
41:a:57:C:C2	41:a:58:G:C8	3.04	0.46
41:a:67:U:N3	41:a:68:G:C5	2.84	0.46
41:a:404:A:H4'	41:a:405:U:O5'	2.15	0.46
41:a:644:A:C2	41:a:2369:A:H1'	2.51	0.46
41:a:685:A:C8	41:a:773:U:C4	3.04	0.46
41:a:985:C:N3	41:a:986:C:C5	2.84	0.46
41:a:1209:U:O2'	41:a:1237:A:N1	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1406:U:O2	41:a:1407:G:C8	2.69	0.46
41:a:2289:G:C2	41:a:2290:G:C8	3.04	0.46
41:a:2700:A:H2'	41:a:2701:U:H6	1.81	0.46
41:a:2756:U:N3	41:a:2758:A:N6	2.64	0.46
41:a:2766:A:C5	41:a:2767:C:C5	3.03	0.46
42:b:51:VAL:N	42:b:62:LYS:HZ3	2.14	0.46
45:e:24:GLU:O	45:e:25:GLN:C	2.57	0.46
52:l:149:ILE:CD1	52:l:188:MET:HG2	2.46	0.46
52:l:194:LYS:HG3	52:l:198:GLU:OE2	2.16	0.46
54:n:4:LEU:HD21	54:n:101:GLU:HG3	1.98	0.46
3:2:3:ARG:HH22	3:2:4:GLU:HB2	1.81	0.46
4:3:46:GLN:NE2	4:3:54:GLN:HG2	2.31	0.46
5:4:50:MET:C	5:4:56:PHE:CZ	2.94	0.46
8:7:50:U:H4'	14:AE:322:ARG:CZ	2.46	0.46
9:9:122:GLN:OE1	9:9:122:GLN:N	2.49	0.46
11:AA:888:THR:N	16:AG:105:ILE:HD12	2.24	0.46
13:AD:35:PHE:HA	13:AD:38:THR:HG22	1.97	0.46
14:AE:388:ARG:NH2	14:AE:414:GLU:OE1	2.49	0.46
14:AE:848:VAL:HB	14:AE:858:VAL:HG22	1.98	0.46
10:B:65:C:C2'	10:B:66:C:H5'	2.44	0.46
17:C:34:THR:HB	17:C:38:LYS:H	1.81	0.46
18:D:1323:G:H2'	18:D:1324:A:H8	1.78	0.46
20:F:37:PHE:HA	31:Q:123:PRO:HB2	1.97	0.46
23:I:85:GLU:O	23:I:89:LYS:CE	2.64	0.46
27:M:17:LYS:HZ2	27:M:18:PHE:HD1	1.58	0.46
29:O:113:ARG:HG3	29:O:115:LYS:HZ2	1.81	0.46
30:P:5:ARG:HE	30:P:7:ARG:CZ	2.29	0.46
34:T:32:LEU:HD12	34:T:63:ARG:HA	1.98	0.46
34:T:75:VAL:HG13	34:T:76:ALA:N	2.31	0.46
39:Y:59:THR:O	39:Y:66:PHE:HB2	2.16	0.46
41:a:152:A:H2'	41:a:153:U:C6	2.51	0.46
41:a:235:U:N3	41:a:236:C:C5	2.84	0.46
41:a:1007:C:H5''	59:s:37:ARG:NE	2.31	0.46
41:a:1064:C:O2'	41:a:1065:U:C6	2.62	0.46
41:a:1786:A:H1'	41:a:1938:A:N6	2.31	0.46
44:d:44:G:H3'	54:n:92:ARG:HH22	1.81	0.46
44:d:82:U:O2'	46:f:53:PHE:HE2	1.98	0.46
44:d:83:G:H1'	46:f:53:PHE:CE1	2.51	0.46
46:f:12:SER:CA	46:f:32:ILE:HD11	2.46	0.46
50:j:5:VAL:HG22	50:j:202:ILE:CD1	2.45	0.46
52:l:35:TYR:CD2	52:l:178:VAL:HG11	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:73:VAL:CG1	59:s:86:GLN:HG2	2.46	0.46
61:u:110:VAL:HG23	61:u:125:LEU:HD12	1.97	0.46
63:w:37:THR:HG21	63:w:40:LYS:HZ2	1.81	0.46
63:w:115:LEU:HD23	63:w:117:ASP:N	2.30	0.46
64:x:15:ARG:C	64:x:19:GLN:HE22	2.24	0.46
64:x:88:LYS:O	64:x:115:LEU:HD12	2.16	0.46
65:y:103:ARG:HD3	65:y:107:ALA:C	2.41	0.46
3:2:18:GLU:O	3:2:22:THR:HG23	2.15	0.45
7:6:25:DT:H2'	11:AA:496:LYS:CG	2.44	0.45
9:9:87:GLU:OE2	9:9:95:LEU:CG	2.64	0.45
11:AA:176:ILE:HD11	11:AA:428:VAL:HG21	1.98	0.45
12:AB:6:LEU:CD2	14:AE:291:ILE:HD11	2.47	0.45
12:AB:76:SER:O	12:AB:77:HIS:HB3	2.16	0.45
16:AG:305:MET:HB3	16:AG:305:MET:HE3	1.77	0.45
10:B:6:G:O2'	10:B:7:G:H8	1.98	0.45
17:C:33:ILE:O	22:H:338:GLU:CG	2.64	0.45
18:D:269:C:H2'	18:D:270:A:C8	2.51	0.45
18:D:337:G:H2'	18:D:338:A:H8	1.80	0.45
18:D:696:A:H2'	18:D:697:U:H6	1.81	0.45
18:D:723:U:O2	20:F:49:LYS:HE3	2.16	0.45
18:D:878:A:P	28:N:80:ARG:NH1	2.89	0.45
18:D:1287:A:H2'	18:D:1288:A:C8	2.51	0.45
18:D:1437:A:C2	18:D:1438:G:C8	3.04	0.45
20:F:12:PHE:CD1	31:Q:101:ASN:ND2	2.84	0.45
22:H:32:ASP:OD1	22:H:35:VAL:HG23	2.16	0.45
22:H:287:LEU:HG	22:H:292:CYS:HA	1.97	0.45
32:R:30:LYS:O	32:R:81:LEU:HD12	2.16	0.45
32:R:68:GLY:O	32:R:99:ARG:NH1	2.49	0.45
33:S:12:LYS:NZ	33:S:16:LEU:HD21	2.31	0.45
33:S:82:ILE:HG23	33:S:83:LYS:N	2.31	0.45
34:T:47:LYS:HD3	34:T:47:LYS:N	2.30	0.45
41:a:132:G:H2'	41:a:133:U:H6	1.81	0.45
41:a:584:C:N4	41:a:585:G:C6	2.83	0.45
41:a:669:G:H2'	41:a:670:A:N7	2.31	0.45
41:a:783:A:O2'	41:a:785:G:OP1	2.35	0.45
41:a:1107:G:C5	41:a:1108:U:C5	3.04	0.45
41:a:1433:A:H2'	41:a:1434:A:C1'	2.46	0.45
41:a:1707:G:C4	41:a:1708:C:C5	3.04	0.45
41:a:2190:G:H2'	41:a:2191:A:C8	2.49	0.45
41:a:2314:A:O2'	54:n:155:THR:HG21	2.16	0.45
41:a:2373:G:H2'	41:a:2374:C:H6	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:d:57:A:C5	54:n:26:MET:CE	2.99	0.45
46:f:41:THR:C	46:f:45:ARG:NH2	2.75	0.45
47:g:1:MET:SD	47:g:3:LYS:HE2	2.56	0.45
47:g:16:CYS:SG	47:g:37:CYS:N	2.89	0.45
52:l:21:ARG:HA	52:l:21:ARG:NE	2.31	0.45
56:p:149:ARG:CZ	56:p:162:VAL:C	2.89	0.45
57:q:2:LYS:HE2	57:q:35:GLN:HG3	1.98	0.45
60:t:48:PRO:HB2	60:t:49:ARG:HE	1.81	0.45
63:w:48:VAL:CG1	63:w:49:GLU:OE1	2.64	0.45
63:w:60:VAL:CG2	63:w:63:ARG:NH1	2.78	0.45
65:y:106:LYS:CD	65:y:109:ARG:NH2	2.79	0.45
9:9:59:LEU:HD22	9:9:62:ARG:HE	1.81	0.45
9:9:60:LEU:O	9:9:64:VAL:HB	2.15	0.45
10:A:46:G:H1'	10:A:47:U:C5	2.51	0.45
10:A:50:U:H2'	10:A:51:C:C6	2.51	0.45
12:AB:17:ALA:O	12:AB:20:HIS:HB2	2.17	0.45
14:AE:283:LEU:HD23	14:AE:292:VAL:HG11	1.99	0.45
14:AE:287:ALA:HB3	14:AE:292:VAL:CG2	2.46	0.45
16:AG:102:PHE:HB3	16:AG:103:ASP:CG	2.41	0.45
16:AG:168:LEU:CB	16:AG:171:GLU:HB2	2.46	0.45
16:AG:434:LEU:HD22	16:AG:459:LEU:HD11	1.80	0.45
16:AG:434:LEU:HD11	16:AG:455:THR:CA	2.46	0.45
18:D:673:A:H2'	18:D:674:G:H8	1.79	0.45
18:D:928:G:H2'	18:D:929:G:H8	1.81	0.45
18:D:1055:A:H1'	23:I:156:ARG:NH2	2.31	0.45
18:D:1086:U:O2	18:D:1086:U:H2'	2.16	0.45
18:D:1436:U:H2'	18:D:1437:A:H8	1.82	0.45
20:F:63:GLU:HG2	20:F:67:ARG:CZ	2.46	0.45
20:F:64:ASN:HA	20:F:67:ARG:HD3	1.97	0.45
21:G:18:HIS:CE1	21:G:205:ASP:OD1	2.68	0.45
21:G:101:LEU:HD22	21:G:157:LEU:HD12	1.99	0.45
23:I:135:LYS:O	23:I:138:VAL:HG12	2.17	0.45
26:L:43:GLY:HA2	26:L:58:HIS:CE1	2.51	0.45
32:R:18:LYS:HD3	32:R:19:SER:O	2.16	0.45
37:W:32:ARG:NH2	37:W:34:TRP:HZ3	2.14	0.45
37:W:44:MET:HE1	37:W:66:MET:HE1	1.98	0.45
38:X:104:THR:O	38:X:104:THR:HG22	2.16	0.45
41:a:642:U:O2'	41:a:644:A:N7	2.43	0.45
41:a:1013:C:C2	41:a:1014:A:C8	3.04	0.45
41:a:1408:G:N3	41:a:1408:G:H2'	2.31	0.45
41:a:1588:G:C6	41:a:1589:U:O4	2.69	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1796:U:H2'	41:a:1797:G:C8	2.50	0.45
41:a:2029:G:N1	41:a:2033:A:OP2	2.35	0.45
41:a:2309:A:H2'	41:a:2310:C:O4'	2.16	0.45
41:a:2649:C:H2'	41:a:2650:U:C6	2.51	0.45
48:h:33:LEU:O	48:h:64:ILE:HG12	2.16	0.45
49:i:30:VAL:HG22	49:i:37:LYS:HD2	1.99	0.45
50:j:5:VAL:HB	50:j:32:ASN:ND2	2.31	0.45
52:l:112:LEU:HD21	52:l:186:VAL:HB	1.97	0.45
56:p:69:ARG:NH1	56:p:73:ASN:CG	2.74	0.45
5:4:53:LYS:HB3	5:4:55:GLU:OE1	2.16	0.45
9:9:34:THR:HA	9:9:37:LYS:NZ	2.32	0.45
9:9:68:PRO:O	9:9:72:LEU:HD21	2.16	0.45
13:AD:15:ASP:OD1	13:AD:27:THR:OG1	2.30	0.45
13:AD:286:GLU:N	13:AD:315:GLY:HA2	2.18	0.45
14:AE:797:THR:HG22	14:AE:924:GLY:HA3	1.97	0.45
16:AG:271:ILE:H	16:AG:271:ILE:HG12	1.49	0.45
18:D:110:C:O4'	35:U:25:ARG:NH2	2.49	0.45
18:D:311:C:P	35:U:31:ARG:NH2	2.86	0.45
18:D:356:A:N3	18:D:368:U:O2'	2.37	0.45
18:D:511:C:C2	18:D:512:U:C5	3.05	0.45
18:D:918:A:H2'	18:D:919:A:C8	2.51	0.45
18:D:984:C:C2	18:D:985:C:C5	3.05	0.45
18:D:1173:U:H2'	18:D:1174:G:O4'	2.16	0.45
19:E:54:MET:C	19:E:54:MET:SD	3.00	0.45
21:G:100:MET:SD	21:G:101:LEU:HD23	2.57	0.45
24:J:95:GLU:HG3	24:J:186:PRO:HG3	1.98	0.45
26:L:18:VAL:N	26:L:19:PRO:CD	2.80	0.45
26:L:94:HIS:CG	26:L:95:ALA:H	2.34	0.45
27:M:44:TYR:C	27:M:44:TYR:CD2	2.94	0.45
32:R:18:LYS:NZ	32:R:19:SER:O	2.47	0.45
36:V:64:CYS:SG	36:V:74:THR:OG1	2.69	0.45
39:Y:96:LYS:HD2	39:Y:136:GLY:O	2.15	0.45
39:Y:102:ARG:HH21	39:Y:105:LEU:HD12	1.81	0.45
41:a:152:A:H2'	41:a:153:U:H6	1.82	0.45
41:a:182:A:H2'	41:a:183:C:H6	1.81	0.45
41:a:590:A:H2'	41:a:591:U:H6	1.80	0.45
41:a:923:G:H4'	42:b:29:GLU:OE1	2.16	0.45
41:a:1065:U:HO2'	41:a:1066:U:C5'	2.27	0.45
41:a:1083:U:O2'	41:a:1085:A:OP2	2.31	0.45
41:a:1796:U:H2'	41:a:1797:G:H8	1.82	0.45
41:a:1797:G:O2'	48:h:257:THR:HG22	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2061:G:N2	41:a:2063:C:C6	2.84	0.45
41:a:2572:A:OP2	50:j:149:ASN:O	2.35	0.45
44:d:45:A:C5'	54:n:92:ARG:NH1	2.79	0.45
47:g:56:ARG:HH21	47:g:59:ARG:HH21	1.62	0.45
49:i:28:LEU:HB2	49:i:37:LYS:HE3	1.98	0.45
51:k:19:HIS:CE1	51:k:41:PRO:CD	3.00	0.45
52:l:198:GLU:OE1	52:l:198:GLU:N	2.43	0.45
54:n:22:TYR:CE2	54:n:28:VAL:HA	2.52	0.45
63:w:60:VAL:HG22	63:w:63:ARG:HH12	1.78	0.45
64:x:83:LEU:HD23	64:x:88:LYS:HE3	1.97	0.45
65:y:7:GLN:HE21	65:y:10:GLN:HE21	1.65	0.45
66:z:89:GLU:N	66:z:89:GLU:OE1	2.50	0.45
9:9:23:LEU:HA	9:9:118:ILE:HD11	1.99	0.45
9:9:129:LEU:HD12	9:9:130:PRO:CD	2.43	0.45
10:A:32:C:C2'	10:A:33:U:H5'	2.46	0.45
11:AA:855:PRO:HG3	16:AG:105:ILE:CG2	2.40	0.45
14:AE:79:LYS:N	16:AG:144:LYS:NZ	2.58	0.45
14:AE:781:LYS:HD2	14:AE:933:ARG:HH12	1.81	0.45
14:AE:1341:ARG:NH1	14:AE:1343:GLU:OE2	2.49	0.45
16:AG:227:ALA:CB	16:AG:331:LEU:HA	2.46	0.45
16:AG:434:LEU:CD2	16:AG:455:THR:O	2.51	0.45
10:B:68:C:H3'	10:B:69:C:H5''	1.97	0.45
18:D:735:C:C2	18:D:736:C:C5	3.05	0.45
18:D:915:A:C5	18:D:916:U:C5	3.04	0.45
21:G:77:SER:O	21:G:80:VAL:HG12	2.17	0.45
22:H:304:VAL:O	22:H:304:VAL:CG1	2.63	0.45
23:I:134:MET:HE2	23:I:168:TYR:CB	2.46	0.45
23:I:165:THR:C	23:I:166:GLU:OE1	2.60	0.45
25:K:13:GLU:O	25:K:13:GLU:HG2	2.16	0.45
27:M:78:ARG:CG	27:M:80:VAL:HG13	2.46	0.45
28:N:26:THR:OG1	28:N:58:GLU:CG	2.65	0.45
32:R:56:ARG:HG2	32:R:62:GLU:CD	2.41	0.45
33:S:86:GLU:OE2	33:S:90:ARG:HD3	2.16	0.45
37:W:5:LEU:HD12	37:W:6:LYS:N	2.32	0.45
37:W:44:MET:HE1	37:W:62:VAL:CG2	2.46	0.45
39:Y:126:ARG:HH12	41:a:1082:U:C5'	2.29	0.45
41:a:136:G:C6	41:a:144:A:C6	3.04	0.45
41:a:396:G:C6	41:a:397:U:C4	3.04	0.45
41:a:580:U:O3'	66:z:31:VAL:HG13	2.17	0.45
41:a:597:G:H2'	41:a:598:U:C6	2.52	0.45
41:a:607:U:C6	41:a:620:G:C5	3.05	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1398:C:C2	41:a:1399:C:C5	3.04	0.45
41:a:1665:A:C5'	60:t:66:LYS:NZ	2.79	0.45
41:a:2356:U:H4'	42:b:20:ARG:HD2	1.99	0.45
41:a:2766:A:C6	41:a:2767:C:C5	3.05	0.45
48:h:52:ARG:CD	48:h:53:HIS:CE1	3.00	0.45
54:n:10:ASP:OD1	54:n:11:GLU:HG3	2.17	0.45
56:p:11:VAL:HG13	56:p:11:VAL:O	2.17	0.45
60:t:59:LYS:HZ3	60:t:92:GLU:CG	2.28	0.45
1:0:71:LYS:CG	1:0:73:LYS:HE3	2.47	0.45
4:3:49:VAL:HG23	4:3:51:ALA:O	2.16	0.45
8:7:13:U:H3	24:J:47:ARG:NH1	2.10	0.45
12:AB:118:ILE:CG1	12:AB:142:LEU:HD23	2.46	0.45
12:AB:126:PHE:CD2	12:AB:126:PHE:O	2.70	0.45
14:AE:482:ALA:HA	15:AF:6:VAL:HG21	1.98	0.45
14:AE:1321:SER:OG	14:AE:1349:GLU:OE2	2.32	0.45
16:AG:123:ARG:HG2	16:AG:123:ARG:HH11	1.82	0.45
16:AG:127:VAL:HG22	16:AG:192:GLY:CA	2.46	0.45
16:AG:197:VAL:HG11	16:AG:199:ARG:NH2	2.30	0.45
16:AG:288:MET:HE3	16:AG:289:ALA:N	2.32	0.45
16:AG:314:LEU:HD22	16:AG:318:ILE:HD11	1.98	0.45
16:AG:349:GLN:C	16:AG:351:GLU:N	2.71	0.45
16:AG:429:LYS:HB2	16:AG:430:PRO:HD2	1.99	0.45
17:C:70:TYR:C	17:C:71:THR:HG23	2.42	0.45
18:D:827:U:O4	18:D:872:A:C2	2.69	0.45
18:D:1109:C:C2	18:D:1110:A:C8	3.04	0.45
20:F:38:TYR:O	31:Q:126:LYS:HE3	2.17	0.45
21:G:52:GLU:C	21:G:56:GLU:OE1	2.60	0.45
23:I:24:ALA:HB3	23:I:29:PHE:CD2	2.51	0.45
24:J:172:GLU:CD	24:J:183:LYS:HD2	2.42	0.45
25:K:13:GLU:OE1	25:K:13:GLU:N	2.50	0.45
27:M:69:VAL:HA	27:M:138:ARG:HD3	1.97	0.45
29:O:51:PRO:CA	29:O:103:PHE:HE2	2.30	0.45
29:O:80:ARG:NH2	29:O:103:PHE:CE1	2.85	0.45
31:Q:88:GLY:H	31:Q:114:THR:HG22	1.81	0.45
32:R:122:PRO:C	32:R:123:LYS:HE2	2.42	0.45
38:X:55:THR:HG22	38:X:59:GLU:OE2	2.17	0.45
39:Y:126:ARG:HH12	41:a:1083:U:P	2.40	0.45
41:a:15:G:O2'	49:i:15:MET:HE2	2.15	0.45
41:a:78:U:P	45:e:4:LYS:NZ	2.89	0.45
41:a:400:G:OP2	43:c:57:ARG:NH1	2.46	0.45
41:a:1042:G:C5	41:a:1043:C:C5	3.05	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1889:A:H2'	41:a:1890:A:C8	2.51	0.45
41:a:2065:C:C2	41:a:2066:C:C5	3.04	0.45
41:a:2262:U:N3	41:a:2263:C:C5	2.85	0.45
41:a:2814:A:C5	41:a:2815:C:C5	3.05	0.45
43:c:43:GLU:OE1	43:c:45:ARG:CB	2.65	0.45
45:e:2:LYS:HZ1	45:e:49:ASP:C	2.23	0.45
45:e:12:GLU:HG2	45:e:13:GLU:H	1.80	0.45
46:f:51:VAL:O	46:f:55:VAL:HG12	2.17	0.45
47:g:8:LYS:NZ	47:g:10:GLU:HB2	2.30	0.45
48:h:200:HIS:O	48:h:203:ARG:HG2	2.17	0.45
51:k:7:GLU:OE2	51:k:27:LYS:HG3	2.16	0.45
60:t:17:ARG:NH2	60:t:47:ILE:CG2	2.80	0.45
65:y:31:TRP:CH2	65:y:40:LEU:HD11	2.52	0.45
1:0:74:ILE:HD12	1:0:74:ILE:N	2.32	0.45
4:3:4:LYS:O	4:3:94:ARG:NH2	2.40	0.45
7:6:25:DT:OP1	11:AA:496:LYS:O	2.34	0.45
9:9:61:ARG:CD	41:a:1046:A:O5'	2.64	0.45
10:A:18:G:N7	10:A:57:A:N6	2.65	0.45
11:AA:694:ARG:HH22	13:AC:80:GLU:HG3	1.82	0.45
11:AA:855:PRO:HG2	16:AG:35:THR:HG21	1.97	0.45
14:AE:210:SER:O	14:AE:214:ARG:N	2.43	0.45
14:AE:420:PRO:HA	14:AE:437:PHE:O	2.16	0.45
10:B:47:U:H2'	10:B:48:C:H4'	1.97	0.45
10:B:50:U:H2'	10:B:51:C:C6	2.51	0.45
17:C:65:LEU:HD23	17:C:67:LEU:CD1	2.47	0.45
18:D:198:G:OP2	18:D:198:G:C8	2.70	0.45
18:D:278:G:OP2	36:V:43:LYS:NZ	2.49	0.45
18:D:826:C:O2	28:N:16:ASN:ND2	2.49	0.45
18:D:878:A:P	28:N:80:ARG:HH11	2.39	0.45
18:D:1308:U:H3'	38:X:98:ARG:NE	2.32	0.45
18:D:1328:C:O3'	38:X:28:THR:HB	2.16	0.45
19:E:49:LYS:O	19:E:53:GLU:HG2	2.16	0.45
22:H:54:LYS:HD3	22:H:58:GLY:C	2.42	0.45
22:H:303:LEU:O	22:H:304:VAL:C	2.59	0.45
23:I:39:VAL:HG23	23:I:40:ARG:N	2.32	0.45
24:J:35:GLU:HG2	24:J:36:GLN:N	2.32	0.45
25:K:145:GLU:CD	25:K:146:ASN:N	2.74	0.45
25:K:150:PRO:O	25:K:153:VAL:HG22	2.16	0.45
26:L:98:GLU:OE1	26:L:98:GLU:HA	2.16	0.45
28:N:89:LYS:HG3	28:N:90:ASP:N	2.32	0.45
29:O:86:ALA:O	29:O:89:GLU:CG	2.65	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Q:51:GLY:O	31:Q:56:ARG:NH1	2.50	0.45
35:U:8:ARG:H	35:U:28:ARG:HD3	1.81	0.45
41:a:581:C:H2'	41:a:582:A:C8	2.50	0.45
41:a:893:C:H2'	41:a:894:U:C5	2.51	0.45
41:a:1310:G:N2	41:a:1313:U:C4	2.85	0.45
41:a:2650:U:H2'	41:a:2651:C:H6	1.81	0.45
41:a:2683:C:H4'	50:j:13:ARG:NH1	2.31	0.45
47:g:47:LYS:CD	54:n:178:ARG:HH22	2.29	0.45
54:n:92:ARG:CZ	54:n:92:ARG:HB3	2.47	0.45
55:o:30:ARG:HH12	61:u:62:PRO:N	2.15	0.45
59:s:17:VAL:HG23	59:s:137:PRO:HB2	1.98	0.45
59:s:136:GLN:N	59:s:137:PRO:CD	2.78	0.45
11:AA:800:MET:HB2	11:AA:800:MET:HE3	1.69	0.45
13:AD:212:ASP:OD1	13:AD:212:ASP:N	2.48	0.45
14:AE:230:SER:OG	14:AE:231:GLY:N	2.49	0.45
14:AE:774:ILE:HA	14:AE:777:HIS:HD2	1.82	0.45
16:AG:239:LYS:CG	16:AG:276:TRP:HB3	2.46	0.45
18:D:675:A:C1'	31:Q:118:HIS:CD2	2.99	0.45
18:D:958:A:C4	37:W:55:ARG:HD3	2.52	0.45
18:D:980:C:C5	18:D:981:U:C2	3.04	0.45
18:D:1060:U:OP1	33:S:85:ARG:NH1	2.48	0.45
18:D:1060:U:H5''	30:P:53:ILE:HD12	1.99	0.45
18:D:1333:A:H2'	18:D:1334:G:O4'	2.17	0.45
18:D:1436:U:H2'	18:D:1437:A:C8	2.52	0.45
21:G:163:VAL:CG1	21:G:185:ALA:HA	2.47	0.45
22:H:24:VAL:HG23	22:H:24:VAL:O	2.16	0.45
22:H:32:ASP:OD1	22:H:35:VAL:HG22	2.17	0.45
22:H:332:VAL:O	22:H:344:LEU:HA	2.17	0.45
23:I:114:LYS:HB2	23:I:185:ASN:ND2	2.32	0.45
27:M:31:MET:HE1	27:M:36:LYS:CD	2.46	0.45
32:R:30:LYS:C	32:R:81:LEU:HD12	2.42	0.45
34:T:10:LYS:CD	34:T:14:GLU:OE2	2.65	0.45
39:Y:64:ARG:O	39:Y:64:ARG:HG3	2.17	0.45
41:a:6:A:O2'	59:s:132:HIS:ND1	2.50	0.45
41:a:535:G:H2'	41:a:536:G:O4'	2.17	0.45
41:a:536:G:H4'	66:z:57:PHE:HZ	1.82	0.45
41:a:1753:G:N2	41:a:1755:A:H3'	2.32	0.45
41:a:2297:A:C2	41:a:2320:U:C1'	3.00	0.45
41:a:2313:C:O3'	54:n:88:LYS:NZ	2.48	0.45
41:a:2727:A:O3'	60:t:70:ARG:NH2	2.50	0.45
52:l:112:LEU:HD23	52:l:118:LEU:HD13	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:m:12:ARG:CG	53:m:44:VAL:HG21	2.47	0.45
54:n:15:LYS:O	54:n:19:GLU:OE1	2.35	0.45
57:q:30:GLU:HB3	57:q:33:HIS:ND1	2.31	0.45
58:r:72:ILE:HG22	58:r:72:ILE:O	2.16	0.45
65:y:31:TRP:CZ3	65:y:40:LEU:CD1	3.00	0.45
5:4:16:ALA:O	5:4:20:LEU:HG	2.17	0.45
9:9:14:GLU:OE1	9:9:63:ALA:HA	2.17	0.45
10:A:11:A:H2'	10:A:12:G:O4'	2.17	0.45
11:AA:24:VAL:HG11	11:AA:704:MET:HE1	1.99	0.45
11:AA:674:ASP:OD2	11:AA:1070:HIS:ND1	2.50	0.45
12:AB:6:LEU:HD22	14:AE:291:ILE:HD11	1.99	0.45
16:AG:233:ARG:HB2	16:AG:327:LEU:CD2	2.42	0.45
10:B:18:G:N7	10:B:57:A:N6	2.65	0.45
18:D:13:U:O4	18:D:20:U:O4	2.34	0.45
18:D:611:C:C2	18:D:612:C:C6	3.04	0.45
20:F:58:LYS:HD2	20:F:58:LYS:C	2.42	0.45
31:Q:52:PHE:CE1	31:Q:65:VAL:HG21	2.52	0.45
41:a:1370:C:H2'	41:a:1371:G:O4'	2.16	0.45
41:a:1709:U:H2'	41:a:1710:G:H8	1.82	0.45
41:a:2016:U:H2'	41:a:2017:U:C6	2.52	0.45
41:a:2347:C:C2	41:a:2348:U:C5	3.05	0.45
43:c:41:GLU:O	43:c:44:LYS:HD2	2.16	0.45
48:h:130:LEU:CA	48:h:189:ARG:HH11	2.24	0.45
52:l:117:ARG:NH1	61:u:1:MET:HE2	2.31	0.45
54:n:63:GLN:HG3	54:n:95:ARG:HD3	1.99	0.45
60:t:42:THR:HG22	60:t:44:LYS:HZ1	1.80	0.45
60:t:108:ARG:CG	60:t:116:ILE:HD13	2.46	0.45
2:1:20:VAL:CG1	2:1:47:VAL:HG11	2.46	0.45
5:4:29:ILE:HD13	44:d:74:U:O2	2.17	0.45
5:4:31:TYR:O	5:4:92:VAL:HA	2.17	0.45
8:7:4:U:C5	18:D:1403:C:H1'	2.52	0.45
9:9:23:LEU:O	9:9:87:GLU:CD	2.60	0.45
11:AA:735:LYS:HA	11:AA:748:ILE:HG22	1.98	0.45
12:AB:16:ARG:C	12:AB:19:GLU:H	2.24	0.45
12:AB:52:PRO:O	12:AB:54:TYR:N	2.50	0.45
12:AB:148:LYS:HZ2	12:AB:150:ILE:CG1	2.28	0.45
14:AE:513:MET:HB3	14:AE:513:MET:HE2	1.79	0.45
16:AG:3:LYS:HB3	16:AG:6:LEU:HD12	1.99	0.45
16:AG:142:VAL:HB	16:AG:175:PRO:HA	1.99	0.45
17:C:12:ARG:HB2	22:H:264:GLU:CB	2.46	0.45
18:D:824:G:C6	18:D:877:G:C6	3.05	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:982:U:P	33:S:6:MET:HE2	2.56	0.45
18:D:1202:U:C5	18:D:1203:C:C5	3.04	0.45
19:E:16:LYS:HG3	19:E:19:LYS:HE3	1.98	0.45
21:G:27:MET:HG3	21:G:27:MET:O	2.16	0.45
25:K:16:ILE:HD11	25:K:38:VAL:CG1	2.46	0.45
25:K:93:ARG:NH2	25:K:93:ARG:CB	2.80	0.45
27:M:4:ARG:HG2	27:M:5:ARG:NH2	2.32	0.45
29:O:113:ARG:HD2	33:S:101:TRP:CD1	2.52	0.45
30:P:26:VAL:C	30:P:30:LYS:HZ3	2.24	0.45
30:P:64:GLN:HG2	33:S:101:TRP:CH2	2.52	0.45
31:Q:21:ALA:HB3	31:Q:84:VAL:HG22	1.99	0.45
33:S:86:GLU:O	33:S:90:ARG:HG2	2.17	0.45
39:Y:102:ARG:NE	39:Y:139:VAL:CG2	2.78	0.45
39:Y:139:VAL:HG22	39:Y:140:GLU:H	1.82	0.45
41:a:219:A:H2'	41:a:220:G:C8	2.52	0.45
41:a:598:U:H2'	41:a:599:A:C8	2.52	0.45
41:a:841:G:H2'	41:a:842:U:C6	2.52	0.45
41:a:1803:A:O2'	48:h:257:THR:HG21	2.16	0.45
41:a:2049:G:N2	50:j:161:MET:CE	2.80	0.45
41:a:2743:U:H2'	41:a:2744:G:O4'	2.17	0.45
41:a:2747:G:C2	41:a:2756:U:C5	3.05	0.45
52:l:23:PHE:CD1	52:l:111:GLU:CD	2.94	0.45
52:l:90:GLN:OE1	52:l:92:HIS:CD2	2.70	0.45
52:l:95:LYS:HE2	52:l:97:ASN:OD1	2.17	0.45
54:n:127:ASN:OD1	54:n:157:THR:HA	2.16	0.45
56:p:91:GLY:HA3	56:p:94:TYR:CD2	2.52	0.45
59:s:98:GLU:O	59:s:100:VAL:N	2.50	0.45
61:u:95:LEU:HD12	61:u:100:ILE:HG21	1.99	0.45
64:x:90:VAL:CG2	64:x:115:LEU:HD21	2.46	0.45
4:3:5:ILE:HG13	4:3:72:ILE:HD11	1.99	0.45
4:3:47:LYS:HZ3	41:a:482:A:P	2.40	0.45
5:4:50:MET:CB	5:4:56:PHE:CE1	2.98	0.45
8:7:2:U:OP2	18:D:926:G:N2	2.50	0.45
10:A:18:G:N2	10:A:58:A:N7	2.65	0.45
10:A:71:C:H2'	10:A:72:A:N9	2.32	0.45
11:AA:1246:ARG:HH11	11:AA:1266:GLY:HA2	1.82	0.45
16:AG:226:ALA:HB1	16:AG:228:ARG:HH11	1.81	0.45
10:B:18:G:C5	10:B:57:A:N6	2.85	0.45
17:C:41:PRO:HD2	22:H:339:ARG:CZ	2.47	0.45
17:C:60:LYS:CE	18:D:735:C:O4'	2.64	0.45
18:D:633:G:H2'	18:D:634:C:H6	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:676:A:H2'	18:D:677:U:H6	1.82	0.45
18:D:753:A:OP1	34:T:73:LYS:NZ	2.50	0.45
18:D:1108:G:C5	18:D:1109:C:C5	3.04	0.45
20:F:12:PHE:CD1	31:Q:101:ASN:CG	2.95	0.45
20:F:26:ALA:CB	20:F:28:VAL:HG23	2.47	0.45
28:N:6:PRO:O	28:N:33:LYS:NZ	2.48	0.45
29:O:88:MET:CE	29:O:95:ARG:HG3	2.47	0.45
34:T:8:THR:O	34:T:11:ILE:HG22	2.16	0.45
38:X:55:THR:C	38:X:59:GLU:OE1	2.60	0.45
41:a:221:A:C4	41:a:266:G:N7	2.85	0.45
41:a:372:G:O2'	43:c:54:LYS:CE	2.64	0.45
41:a:607:U:C6	41:a:620:G:N7	2.85	0.45
41:a:952:G:C6	41:a:966:G:C6	3.04	0.45
41:a:1182:G:H2'	41:a:1183:U:O4'	2.17	0.45
41:a:2848:G:C2	41:a:2867:G:C4	3.05	0.45
48:h:108:LYS:HZ3	48:h:194:GLU:H	1.65	0.45
48:h:146:MET:HE2	48:h:153:GLN:OE1	2.17	0.45
48:h:164:ILE:O	48:h:164:ILE:CG2	2.66	0.45
50:j:181:ASP:HB3	50:j:186:LEU:HB2	1.99	0.45
52:l:23:PHE:CE2	52:l:25:GLU:HB2	2.52	0.45
54:n:16:LEU:HD11	54:n:169:LEU:HD13	1.99	0.45
54:n:175:PHE:CG	54:n:176:PRO:HD2	2.52	0.45
59:s:28:LEU:HD12	59:s:142:ILE:CG2	2.47	0.45
59:s:31:GLU:HG2	59:s:142:ILE:HG23	1.98	0.45
59:s:118:MET:HB3	59:s:118:MET:HE2	1.81	0.45
63:w:60:VAL:HA	63:w:63:ARG:CZ	2.47	0.45
9:9:43:LYS:HE3	9:9:102:ALA:HB2	1.97	0.44
10:A:18:G:C5	10:A:57:A:N6	2.85	0.44
10:A:53:G:N3	10:A:54:U:C5	2.85	0.44
11:AA:1002:LEU:HD21	11:AA:1007:LYS:HB2	1.98	0.44
12:AB:31:PRO:C	12:AB:50:LEU:H	2.22	0.44
12:AB:118:ILE:HD12	12:AB:142:LEU:HD22	1.99	0.44
14:AE:68:TYR:CA	14:AE:92:VAL:HG23	2.45	0.44
10:B:18:G:N2	10:B:58:A:N7	2.65	0.44
10:B:53:G:N3	10:B:54:U:C5	2.85	0.44
10:B:71:C:H2'	10:B:72:A:N9	2.32	0.44
17:C:34:THR:OG1	17:C:38:LYS:HB2	2.17	0.44
17:C:60:LYS:HZ3	18:D:735:C:P	2.37	0.44
18:D:339:C:H2'	18:D:340:U:C6	2.50	0.44
18:D:983:A:H2	18:D:984:C:C6	2.34	0.44
18:D:1172:C:H2'	18:D:1173:U:H6	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:E:25:ARG:O	19:E:28:MET:HB2	2.18	0.44
20:F:16:LEU:HD12	31:Q:98:ARG:HH21	1.81	0.44
21:G:96:TRP:CZ2	21:G:100:MET:HE3	2.52	0.44
21:G:109:GLN:C	21:G:113:ARG:NE	2.75	0.44
22:H:49:PRO:CD	22:H:84:LEU:HD21	2.47	0.44
22:H:332:VAL:CG1	22:H:335:ILE:HG13	2.48	0.44
26:L:47:LEU:HD13	26:L:51:ILE:HG12	1.98	0.44
27:M:134:ALA:O	27:M:137:LYS:CG	2.65	0.44
32:R:31:ARG:O	32:R:57:LEU:HD12	2.17	0.44
34:T:11:ILE:HD11	34:T:30:ALA:HB1	1.99	0.44
34:T:15:PHE:CE2	34:T:30:ALA:CB	3.00	0.44
41:a:1146:C:C2	41:a:1147:A:C8	3.05	0.44
41:a:2478:A:C8	41:a:2529:G:C5	3.04	0.44
41:a:2646:C:OP2	41:a:2732:G:O2'	2.25	0.44
41:a:2783:U:H2'	41:a:2784:U:C6	2.52	0.44
50:j:48:ILE:CD1	50:j:84:LEU:HD11	2.46	0.44
54:n:122:PHE:HZ	54:n:170:LEU:HD12	1.81	0.44
55:o:26:HIS:CE1	55:o:48:ALA:HB2	2.51	0.44
56:p:85:LYS:HE3	56:p:164:TYR:HD1	1.67	0.44
57:q:1:MET:HE3	57:q:1:MET:HA	1.97	0.44
1:0:73:LYS:HD3	1:0:86:GLN:HE21	1.82	0.44
6:5:10:DT:H2''	12:AB:71:ALA:H	1.79	0.44
8:7:54:A:O3'	11:AA:688:GLN:NE2	2.51	0.44
9:9:3:LEU:HD12	9:9:5:LEU:H	1.81	0.44
9:9:58:THR:HG23	41:a:1108:U:OP1	2.17	0.44
13:AD:185:TYR:HA	13:AD:202:VAL:O	2.16	0.44
14:AE:79:LYS:CD	16:AG:159:GLU:HB3	2.44	0.44
16:AG:331:LEU:HA	16:AG:331:LEU:HD23	1.79	0.44
16:AG:403:VAL:C	16:AG:405:ALA:N	2.73	0.44
10:B:3:C:H2'	10:B:4:G:C8	2.52	0.44
18:D:219:U:C2	18:D:220:G:C8	3.05	0.44
18:D:524:G:H2'	18:D:525:C:C6	2.52	0.44
18:D:1250:A:H2'	18:D:1251:A:C8	2.52	0.44
21:G:110:SER:OG	21:G:113:ARG:NH2	2.47	0.44
22:H:12:SER:O	22:H:16:ILE:HG13	2.17	0.44
22:H:53:PHE:CE1	22:H:68:VAL:HG21	2.52	0.44
22:H:294:VAL:HG11	22:H:330:VAL:HG21	2.00	0.44
23:I:55:ILE:HD12	23:I:68:ILE:HG12	1.99	0.44
23:I:84:VAL:HG23	23:I:85:GLU:N	2.31	0.44
23:I:156:ARG:CG	23:I:193:TYR:HB2	2.47	0.44
25:K:50:TYR:O	25:K:66:LYS:HE3	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:72:ASP:O	26:L:76:THR:HG23	2.17	0.44
29:O:94:LEU:O	29:O:97:GLU:CD	2.60	0.44
32:R:27:CYS:CB	32:R:30:LYS:HE2	2.47	0.44
32:R:110:ARG:N	32:R:111:LYS:HZ3	2.14	0.44
33:S:6:MET:HE3	33:S:63:ARG:HH11	1.82	0.44
37:W:13:LEU:O	37:W:17:LYS:HG2	2.17	0.44
38:X:90:ARG:HD3	38:X:96:PRO:O	2.17	0.44
41:a:900:A:C5	41:a:901:C:C5	3.06	0.44
41:a:1144:A:C4	41:a:1145:C:C5	3.05	0.44
41:a:2095:A:O5'	58:r:11:ASN:OD1	2.35	0.44
41:a:2450:A:C2	41:a:2451:A:C4	3.06	0.44
41:a:2646:C:H2'	41:a:2647:U:O4'	2.17	0.44
41:a:2723:C:H2'	41:a:2724:U:O4'	2.16	0.44
46:f:4:THR:CG2	46:f:37:GLU:OE2	2.65	0.44
48:h:220:VAL:HG23	48:h:225:MET:HE3	2.00	0.44
50:j:3:GLY:O	50:j:82:PHE:CE2	2.69	0.44
51:k:34:LEU:HB2	51:k:51:GLU:OE2	2.16	0.44
55:o:45:ARG:HB3	55:o:46:PRO:CD	2.48	0.44
56:p:141:ILE:HG13	56:p:142:GLY:N	2.32	0.44
3:2:6:ARG:NE	3:2:42:GLU:CD	2.76	0.44
10:A:3:C:H2'	10:A:4:G:C8	2.53	0.44
12:AB:117:ILE:HG21	12:AB:161:LYS:HE2	1.99	0.44
12:AB:120:GLU:CB	12:AB:162:LEU:HB2	2.46	0.44
12:AB:130:PHE:HZ	12:AB:160:ARG:HG3	1.81	0.44
16:AG:339:MET:HE2	16:AG:343:ASP:HB3	1.98	0.44
18:D:682:G:C2	18:D:683:G:C8	3.06	0.44
19:E:58:VAL:HG12	19:E:72:ALA:HB1	1.99	0.44
21:G:128:LYS:C	21:G:129:LEU:HD12	2.42	0.44
23:I:14:ILE:HG13	23:I:178:LEU:HD21	1.99	0.44
23:I:147:LYS:HE2	23:I:204:LYS:C	2.42	0.44
25:K:134:ILE:CG2	25:K:135:ASN:N	2.81	0.44
26:L:9:MET:C	26:L:9:MET:SD	3.00	0.44
33:S:77:PHE:HE1	33:S:93:ILE:CG2	2.30	0.44
37:W:79:THR:HG23	37:W:79:THR:O	2.17	0.44
38:X:2:ALA:HB1	38:X:9:ILE:CG2	2.47	0.44
39:Y:19:PRO:CG	39:Y:23:VAL:CG2	2.95	0.44
41:a:65:U:N3	41:a:66:C:C5	2.86	0.44
41:a:68:G:N2	41:a:74:A:OP2	2.51	0.44
41:a:150:U:C2	41:a:151:C:C5	3.06	0.44
41:a:598:U:H2'	41:a:599:A:H8	1.82	0.44
41:a:730:A:C2	41:a:731:C:C6	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:782:A:N1	48:h:225:MET:HE2	2.32	0.44
41:a:957:C:H5'	62:v:75:GLU:OE1	2.18	0.44
41:a:1414:C:H2'	41:a:1415:U:C6	2.52	0.44
41:a:1443:U:H2'	41:a:1444:G:H8	1.83	0.44
41:a:1656:C:C2	41:a:1657:U:C5	3.05	0.44
41:a:2492:U:C2'	41:a:2493:U:H5'	2.48	0.44
44:d:113:C:HO2'	64:x:46:GLU:CD	2.26	0.44
45:e:24:GLU:O	45:e:28:LEU:HD13	2.17	0.44
47:g:58:ASP:C	47:g:62:LYS:HZ1	2.25	0.44
48:h:240:PHE:CD2	48:h:240:PHE:O	2.71	0.44
50:j:55:LYS:HZ2	50:j:60:VAL:N	2.15	0.44
52:l:1:MET:HE3	52:l:19:PHE:C	2.42	0.44
52:l:6:LYS:HZ1	52:l:119:ILE:C	2.25	0.44
56:p:18:LYS:HG3	56:p:27:LYS:NZ	2.32	0.44
56:p:69:ARG:NH1	56:p:73:ASN:ND2	2.66	0.44
61:u:101:ILE:HG13	61:u:102:GLY:H	1.82	0.44
65:y:15:GLN:NE2	65:y:16:ASP:CG	2.75	0.44
5:4:31:TYR:HE2	44:d:104:A:O4'	2.01	0.44
6:5:11:DA:C5'	12:AB:67:THR:O	2.66	0.44
8:7:12:U:C3'	8:7:12:U:C6	3.00	0.44
9:9:4:ASN:OD1	41:a:1046:A:N7	2.50	0.44
11:AA:560:PRO:HB2	14:AE:776:THR:HG21	1.99	0.44
12:AB:103:ASP:CB	12:AB:105:VAL:HG23	2.34	0.44
12:AB:126:PHE:O	12:AB:126:PHE:CG	2.70	0.44
14:AE:53:ARG:HA	14:AE:54:ASP:HA	1.59	0.44
14:AE:1037:PHE:HB3	14:AE:1040:MET:HB2	1.97	0.44
16:AG:320:ARG:HH11	16:AG:320:ARG:HG2	1.80	0.44
16:AG:355:ALA:CB	16:AG:382:GLU:HG3	2.30	0.44
16:AG:429:LYS:CB	16:AG:430:PRO:HD2	2.48	0.44
17:C:34:THR:HG22	17:C:35:GLU:N	2.33	0.44
17:C:54:GLN:HA	17:C:57:ARG:NH1	2.33	0.44
18:D:511:C:C2	18:D:512:U:C6	3.05	0.44
18:D:622:A:C8	18:D:623:C:C6	3.06	0.44
18:D:653:U:O4'	28:N:56:LYS:HE3	2.16	0.44
18:D:1408:A:H2'	18:D:1409:C:C6	2.53	0.44
19:E:22:ALA:CA	19:E:25:ARG:HE	2.31	0.44
21:G:19:GLN:HE21	22:H:75:VAL:C	2.24	0.44
22:H:37:LEU:HG	22:H:46:SER:O	2.17	0.44
22:H:53:PHE:HE1	22:H:68:VAL:HG21	1.82	0.44
22:H:76:GLU:OE1	22:H:79:PHE:HA	2.18	0.44
26:L:17:GLN:CB	26:L:21:MET:HE1	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:78:PHE:HD1	26:L:87:SER:HB3	1.83	0.44
27:M:138:ARG:HG2	27:M:142:HIS:HE1	1.80	0.44
30:P:47:GLU:HB2	30:P:67:ILE:HG22	1.99	0.44
35:U:6:LEU:HD23	35:U:17:TYR:CB	2.47	0.44
36:V:30:LYS:HB2	36:V:37:PHE:CE1	2.52	0.44
38:X:14:HIS:HB2	38:X:17:ILE:HD13	1.99	0.44
39:Y:44:LYS:HG2	39:Y:68:PHE:HE2	1.83	0.44
39:Y:76:ALA:O	39:Y:80:LYS:NZ	2.33	0.44
41:a:66:C:N4	41:a:74:A:N6	2.66	0.44
41:a:251:A:OP1	61:u:58:TYR:OH	2.23	0.44
41:a:2514:U:H2'	41:a:2515:C:H6	1.81	0.44
41:a:2839:G:C5	41:a:2840:C:C5	3.05	0.44
46:f:31:ARG:HG2	46:f:34:HIS:HB2	2.00	0.44
49:i:17:ARG:CA	49:i:20:ASP:OD1	2.66	0.44
54:n:40:VAL:HG13	54:n:42:GLU:CD	2.43	0.44
55:o:8:ARG:HA	55:o:8:ARG:HE	1.83	0.44
56:p:11:VAL:HG12	56:p:48:ASN:O	2.17	0.44
56:p:121:ILE:HD12	56:p:141:ILE:CG2	2.39	0.44
63:w:13:ASN:O	63:w:17:ARG:HG3	2.17	0.44
65:y:106:LYS:CD	65:y:109:ARG:HH22	2.29	0.44
66:z:66:ASN:CG	66:z:76:TYR:HB2	2.42	0.44
1:0:46:GLU:HG2	1:0:48:LYS:NZ	2.32	0.44
1:0:77:PHE:CD2	66:z:40:ILE:HG21	2.53	0.44
2:1:36:LEU:N	2:1:36:LEU:HD12	2.33	0.44
11:AA:812:PHE:O	14:AE:504:GLN:NE2	2.40	0.44
11:AA:1244:HIS:NE2	11:AA:1266:GLY:O	2.44	0.44
16:AG:282:GLN:NE2	16:AG:282:GLN:CA	2.73	0.44
10:B:11:A:H2'	10:B:12:G:O4'	2.17	0.44
18:D:604:G:H2'	18:D:605:U:O4'	2.17	0.44
20:F:32:VAL:HG11	31:Q:89:PRO:HG3	1.99	0.44
21:G:117:LEU:HD23	21:G:141:LEU:HA	1.98	0.44
23:I:22:TRP:CZ2	23:I:32:ASN:CB	3.01	0.44
26:L:9:MET:HE3	26:L:9:MET:HB3	1.85	0.44
27:M:78:ARG:HG3	27:M:80:VAL:HG13	1.99	0.44
28:N:25:VAL:HG23	28:N:63:LEU:CD1	2.46	0.44
28:N:35:ALA:HB1	28:N:110:VAL:HG11	2.00	0.44
29:O:46:MET:HB3	29:O:50:GLN:NE2	2.31	0.44
30:P:40:ILE:HD12	30:P:73:LEU:CD2	2.48	0.44
31:Q:85:MET:HG3	31:Q:111:THR:HB	2.00	0.44
32:R:38:TYR:CE2	32:R:40:THR:CG2	3.01	0.44
34:T:3:LEU:CD2	34:T:8:THR:HG23	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:W:39:THR:HG22	37:W:40:ILE:N	2.33	0.44
39:Y:12:VAL:HG22	39:Y:54:ILE:O	2.17	0.44
39:Y:82:ALA:HB3	39:Y:108:ILE:CD1	2.48	0.44
41:a:234:U:C4	41:a:235:U:C5	3.05	0.44
41:a:575:A:C2	41:a:576:U:C6	3.05	0.44
41:a:1020:A:C2	41:a:1141:U:C2	3.06	0.44
41:a:1084:A:N6	41:a:1085:A:N1	2.66	0.44
41:a:1774:C:O2	41:a:1774:C:C2'	2.65	0.44
41:a:1820:U:C2	48:h:201:MET:HE2	2.53	0.44
41:a:1936:A:C8	41:a:1945:G:N7	2.86	0.44
41:a:2063:C:N3	41:a:2064:C:C5	2.86	0.44
41:a:2617:U:C2	41:a:2618:G:C8	3.06	0.44
41:a:2718:G:O2'	41:a:2847:U:H5''	2.18	0.44
43:c:33:LEU:O	43:c:34:HIS:CG	2.70	0.44
47:g:30:HIS:CG	47:g:31:ASP:N	2.85	0.44
50:j:61:THR:OG1	50:j:63:PRO:HD2	2.18	0.44
55:o:55:LEU:C	55:o:55:LEU:HD23	2.42	0.44
56:p:8:PRO:C	56:p:69:ARG:NH2	2.75	0.44
60:t:18:ARG:NE	60:t:18:ARG:HA	2.33	0.44
61:u:91:ASP:HB2	61:u:93:ASN:OD1	2.17	0.44
2:1:20:VAL:HG11	2:1:47:VAL:HG11	1.99	0.44
3:2:65:GLY:H	3:2:76:ARG:HH12	1.64	0.44
8:7:9:U:OP2	8:7:9:U:C6	2.71	0.44
9:9:25:ALA:HB2	9:9:96:PHE:HE1	1.82	0.44
11:AA:148:GLN:NE2	11:AA:535:PRO:O	2.40	0.44
11:AA:712:SER:OG	11:AA:713:GLY:N	2.49	0.44
11:AA:856:ASN:CA	16:AG:35:THR:CB	2.89	0.44
12:AB:31:PRO:HG2	12:AB:51:PHE:CE2	2.53	0.44
14:AE:41:PRO:HB2	14:AE:270:ARG:HG3	1.99	0.44
16:AG:80:ALA:HB1	16:AG:87:LEU:HB2	1.98	0.44
10:B:71:C:H2'	10:B:72:A:C1'	2.47	0.44
10:B:71:C:C3'	10:B:72:A:C8	2.99	0.44
18:D:375:U:OP2	35:U:70:ARG:NE	2.51	0.44
18:D:377:G:H2'	18:D:378:G:H8	1.81	0.44
18:D:414:A:C4	18:D:415:A:C8	3.05	0.44
18:D:715:A:H2'	18:D:716:A:C8	2.52	0.44
18:D:1486:G:H2'	18:D:1487:G:O4'	2.18	0.44
21:G:206:ALA:O	21:G:210:VAL:HG23	2.18	0.44
22:H:49:PRO:HD3	22:H:84:LEU:HD21	1.99	0.44
22:H:163:ARG:N	22:H:298:GLU:CG	2.81	0.44
23:I:72:ARG:NE	23:I:75:ILE:HB	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:O:63:LEU:CD1	29:O:65:ILE:HD11	2.48	0.44
31:Q:83:GLU:OE1	31:Q:83:GLU:N	2.51	0.44
34:T:26:GLU:HG3	34:T:27:VAL:N	2.33	0.44
38:X:106:ALA:HB3	38:X:110:LYS:CD	2.48	0.44
41:a:30:G:H2'	41:a:31:C:H6	1.82	0.44
41:a:564:C:H1'	66:z:37:GLN:NE2	2.32	0.44
41:a:980:A:N7	41:a:1136:G:H5'	2.32	0.44
41:a:1170:C:O2	41:a:1170:C:H2'	2.18	0.44
41:a:1656:C:H2'	41:a:1657:U:H6	1.81	0.44
41:a:2702:G:C4	41:a:2703:C:C5	3.06	0.44
48:h:123:ALA:HB1	48:h:125:LYS:CE	2.47	0.44
49:i:23:THR:O	49:i:23:THR:HG23	2.17	0.44
50:j:131:ASP:OD1	50:j:133:THR:O	2.36	0.44
52:l:6:LYS:CE	52:l:119:ILE:CG2	2.94	0.44
53:m:41:ARG:NE	53:m:41:ARG:HA	2.32	0.44
61:u:55:MET:SD	61:u:60:ARG:N	2.90	0.44
61:u:101:ILE:C	61:u:105:ILE:HD12	2.42	0.44
62:v:3:GLN:OE1	62:v:4:PRO:O	2.36	0.44
62:v:38:ARG:HB2	62:v:98:PRO:HD3	2.00	0.44
66:z:76:TYR:O	66:z:80:ILE:HG12	2.16	0.44
2:1:78:GLU:OE2	41:a:1262:A:OP1	2.35	0.44
8:7:1:A:C6	10:B:37:A:N1	2.86	0.44
11:AA:557:ARG:NH2	11:AA:607:SER:O	2.49	0.44
13:AD:68:TYR:HE1	13:AD:79:LEU:HD13	1.82	0.44
14:AE:708:ASN:OD1	14:AE:708:ASN:N	2.50	0.44
16:AG:167:MET:HE3	16:AG:173:PHE:CZ	2.53	0.44
18:D:76:G:C5	18:D:77:A:C8	3.06	0.44
18:D:153:C:C2	18:D:154:U:C5	3.06	0.44
18:D:376:G:C2	18:D:389:A:C2	3.06	0.44
18:D:415:A:C6	18:D:416:G:C5	3.06	0.44
18:D:1062:U:H2'	18:D:1063:C:C6	2.53	0.44
21:G:48:PRO:O	21:G:52:GLU:HG2	2.18	0.44
27:M:23:LEU:H	27:M:23:LEU:CD1	2.30	0.44
27:M:106:GLU:O	27:M:110:LYS:HG2	2.18	0.44
30:P:24:GLU:OE2	30:P:92:LEU:CD2	2.66	0.44
30:P:73:LEU:C	30:P:73:LEU:HD23	2.42	0.44
30:P:85:ASP:HA	30:P:88:MET:HE3	2.00	0.44
31:Q:14:LYS:HD3	31:Q:15:GLN:N	2.32	0.44
31:Q:84:VAL:HG21	31:Q:97:ILE:HD11	1.99	0.44
31:Q:91:PRO:HD2	31:Q:92:GLY:H	1.83	0.44
32:R:23:ALA:O	32:R:25:GLU:OE1	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:T:71:LYS:HE2	34:T:78:TYR:CD2	2.52	0.44
36:V:27:ARG:HH12	36:V:42:THR:CB	2.26	0.44
38:X:3:ARG:NH2	54:n:110:ARG:HD2	2.33	0.44
41:a:27:G:C4	41:a:512:G:N2	2.86	0.44
41:a:536:G:H4'	66:z:57:PHE:CZ	2.53	0.44
41:a:582:A:H5''	66:z:11:ARG:HH12	1.81	0.44
41:a:833:A:OP2	61:u:39:LYS:NZ	2.49	0.44
41:a:837:C:N3	41:a:941:A:N6	2.65	0.44
41:a:910:A:H8	62:v:12:MET:CE	2.31	0.44
41:a:1792:G:C5'	48:h:204:VAL:HG23	2.48	0.44
41:a:1856:U:H2'	41:a:1857:G:O4'	2.17	0.44
41:a:1997:C:OP1	50:j:128:ARG:NH1	2.51	0.44
41:a:2636:C:O2'	50:j:45:TYR:CZ	2.70	0.44
52:l:23:PHE:CD1	52:l:111:GLU:CG	3.00	0.44
54:n:101:GLU:O	54:n:105:THR:OG1	2.23	0.44
55:o:49:MET:H	55:o:49:MET:HE2	1.81	0.44
56:p:137:ASP:OD2	56:p:140:VAL:N	2.45	0.44
59:s:35:ARG:HA	59:s:40:HIS:ND1	2.32	0.44
62:v:96:ILE:HA	62:v:100:LYS:HE3	2.00	0.44
62:v:111:GLU:HG2	62:v:112:LEU:N	2.33	0.44
1:0:93:PHE:CD1	1:0:93:PHE:C	2.96	0.44
9:9:118:ILE:HB	9:9:119:PRO:CD	2.42	0.44
10:A:3:C:H2'	10:A:4:G:H8	1.82	0.44
12:AB:38:ILE:HA	12:AB:42:LYS:O	2.18	0.44
13:AD:192:VAL:HG12	13:AD:193:GLU:H	1.83	0.44
14:AE:596:LEU:HD23	14:AE:596:LEU:HA	1.89	0.44
16:AG:19:PRO:HG2	16:AG:19:PRO:O	2.17	0.44
16:AG:227:ALA:CB	16:AG:330:GLN:C	2.90	0.44
16:AG:320:ARG:HB2	16:AG:323:GLN:CB	2.48	0.44
10:B:3:C:H2'	10:B:4:G:H8	1.82	0.44
10:B:72:A:C3'	10:B:73:A:H5''	2.47	0.44
17:C:38:LYS:NZ	18:D:718:A:C6	2.79	0.44
18:D:108:G:H5''	18:D:108:G:N3	2.33	0.44
18:D:429:U:O2	18:D:430:A:N7	2.50	0.44
18:D:598:U:C2	18:D:599:C:C5	3.06	0.44
18:D:610:U:C2	18:D:611:C:C6	3.06	0.44
18:D:704:A:C4	18:D:705:G:C8	3.05	0.44
18:D:1104:G:H4'	21:G:113:ARG:HH22	1.83	0.44
18:D:1120:C:C2	18:D:1121:U:C5	3.06	0.44
18:D:1146:A:C5	18:D:1147:C:C5	3.06	0.44
18:D:1477:U:H2'	18:D:1478:U:C6	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:8:LEU:HD12	22:H:11:GLU:OE2	2.18	0.44
22:H:19:ARG:O	22:H:72:LEU:HD13	2.18	0.44
22:H:277:GLY:HA2	22:H:331:MET:HE3	1.97	0.44
23:I:34:ASP:HB2	33:S:65:ARG:CZ	2.48	0.44
23:I:72:ARG:HD3	23:I:75:ILE:HG22	1.99	0.44
24:J:74:ASN:CA	24:J:77:LYS:HZ3	2.31	0.44
27:M:132:GLY:O	27:M:136:LYS:HD3	2.17	0.44
28:N:110:VAL:O	28:N:111:MET:SD	2.75	0.44
33:S:13:ARG:HB3	33:S:60:GLN:HE21	1.81	0.44
35:U:14:ARG:CD	35:U:42:ILE:HG21	2.47	0.44
39:Y:14:ALA:O	39:Y:16:MET:N	2.51	0.44
39:Y:45:THR:HG22	39:Y:45:THR:O	2.16	0.44
39:Y:60:VAL:HG22	39:Y:66:PHE:CD1	2.52	0.44
41:a:39:G:H2'	41:a:40:U:C6	2.53	0.44
41:a:579:G:H2'	41:a:580:U:C6	2.53	0.44
41:a:594:U:C2	41:a:595:C:C5	3.05	0.44
41:a:730:A:C2	41:a:731:C:C5	3.05	0.44
41:a:1177:G:O2'	41:a:1178:C:O5'	2.28	0.44
41:a:1754:A:N1	41:a:2716:C:O2'	2.34	0.44
41:a:2000:C:OP1	63:w:5:LYS:NZ	2.38	0.44
41:a:2025:C:H2'	41:a:2026:U:C6	2.53	0.44
41:a:2179:C:H2'	41:a:2180:U:H6	1.83	0.44
41:a:2391:G:P	55:o:32:ILE:CD1	3.06	0.44
41:a:2484:G:OP1	62:v:44:ARG:HD2	2.18	0.44
41:a:2548:U:C2	41:a:2549:G:C8	3.06	0.44
41:a:2850:A:N7	41:a:2868:A:O2'	2.41	0.44
43:c:4:VAL:HG22	43:c:11:ARG:HG2	2.00	0.44
45:e:59:GLU:CD	45:e:60:LYS:N	2.76	0.44
48:h:45:ASN:OD1	48:h:46:ASN:N	2.51	0.44
48:h:205:LEU:HD12	48:h:210:ALA:CB	2.44	0.44
50:j:104:VAL:HG12	50:j:105:LYS:N	2.33	0.44
50:j:172:VAL:HG11	50:j:175:LEU:HD11	1.99	0.44
54:n:79:ILE:O	54:n:79:ILE:HG13	2.17	0.44
56:p:27:LYS:HG2	56:p:32:GLU:HG3	1.99	0.44
56:p:83:PHE:HB2	56:p:141:ILE:HD13	2.00	0.44
58:r:127:GLU:OE2	58:r:143:ILE:HB	2.17	0.44
64:x:14:ALA:O	64:x:15:ARG:C	2.61	0.44
64:x:50:ALA:O	64:x:81:ARG:NH2	2.51	0.44
5:4:57:TYR:HD2	62:v:136:MET:HE2	1.81	0.44
9:9:31:ARG:HD2	41:a:1053:C:HO2'	1.75	0.44
10:A:35:A:H2'	10:A:36:U:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:557:ARG:HB3	11:AA:587:LEU:HD13	2.00	0.44
11:AA:746:ALA:O	11:AA:974:ARG:NH2	2.45	0.44
12:AB:33:ILE:HD11	12:AB:104:ILE:HA	2.00	0.44
12:AB:37:LYS:HE3	12:AB:109:THR:HG21	1.99	0.44
13:AD:78:ILE:HA	13:AD:81:ILE:HG22	2.00	0.44
14:AE:78:LEU:HD23	14:AE:78:LEU:HA	1.81	0.44
14:AE:658:GLU:HA	14:AE:661:VAL:HG22	2.00	0.44
15:AF:58:LEU:HD12	15:AF:59:ILE:HG12	2.00	0.44
16:AG:227:ALA:HB2	16:AG:330:GLN:O	2.18	0.44
17:C:18:VAL:HG13	17:C:51:TYR:HE1	1.83	0.44
17:C:70:TYR:CE2	26:L:49:TYR:CE1	3.05	0.44
18:D:323:U:H4'	19:E:17:ALA:HB3	1.99	0.44
18:D:526:C:P	32:R:88:LYS:HZ3	2.41	0.44
18:D:754:C:O5'	34:T:72:ARG:NH1	2.51	0.44
18:D:1125:U:N3	18:D:1127:G:C8	2.86	0.44
18:D:1492:A:HO2'	18:D:1493:A:H5''	1.83	0.44
19:E:78:ASN:O	19:E:82:GLN:CD	2.61	0.44
20:F:66:ARG:NH2	20:F:67:ARG:NH2	2.66	0.44
21:G:23:TRP:CZ3	21:G:25:PRO:CA	3.01	0.44
22:H:71:ALA:HB3	22:H:83:LEU:O	2.18	0.44
23:I:134:MET:HG3	23:I:151:VAL:HG21	2.00	0.44
26:L:16:GLU:H	26:L:16:GLU:CD	2.26	0.44
31:Q:63:ALA:CB	31:Q:92:GLY:HA3	2.48	0.44
31:Q:84:VAL:HG11	31:Q:97:ILE:HD13	2.00	0.44
34:T:11:ILE:HD13	34:T:31:LEU:HA	1.99	0.44
34:T:15:PHE:HE2	34:T:30:ALA:HB1	1.82	0.44
34:T:71:LYS:HZ3	34:T:78:TYR:HD2	1.62	0.44
39:Y:33:ASN:OD1	39:Y:35:MET:HE2	2.17	0.44
41:a:128:C:C2	41:a:129:C:C5	3.06	0.44
41:a:145:C:H2'	41:a:146:A:C8	2.53	0.44
41:a:782:A:C6	48:h:225:MET:HE2	2.53	0.44
41:a:1824:G:P	48:h:52:ARG:CZ	3.06	0.44
41:a:1913:A:OP2	41:a:1913:A:H3'	2.18	0.44
41:a:1996:C:OP1	60:t:31:ARG:NH2	2.51	0.44
41:a:2021:C:OP1	49:i:9:THR:HG21	2.18	0.44
41:a:2098:U:H2'	41:a:2099:U:O4'	2.18	0.44
41:a:2149:U:H2'	41:a:2150:C:C6	2.52	0.44
44:d:46:A:C5	44:d:47:C:C5	3.06	0.44
47:g:59:ARG:HH11	47:g:63:ARG:CD	2.31	0.44
51:k:7:GLU:HG3	51:k:27:LYS:HZ3	1.83	0.44
54:n:117:LEU:N	54:n:177:PHE:O	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
59:s:132:HIS:HA	59:s:135:GLN:OE1	2.17	0.44
63:w:79:LEU:HD23	63:w:83:LEU:HD12	1.99	0.44
1:0:1:MET:HE3	1:0:43:ASN:OD1	2.18	0.43
2:1:36:LEU:HD23	2:1:48:LYS:CA	2.48	0.43
5:4:80:HIS:CD2	5:4:83:LYS:HZ3	2.34	0.43
11:AA:11:ILE:HG22	11:AA:1172:LEU:HD11	1.99	0.43
11:AA:813:GLU:HB2	14:AE:461:PHE:HD2	1.83	0.43
11:AA:998:LEU:HG	11:AA:1011:LEU:HB3	1.99	0.43
13:AD:211:ILE:HD12	13:AD:211:ILE:HA	1.92	0.43
16:AG:104:ARG:N	16:AG:105:ILE:HG12	2.32	0.43
16:AG:363:LEU:CD2	16:AG:409:ARG:N	2.81	0.43
18:D:284:C:C2	18:D:285:C:C5	3.06	0.43
18:D:642:A:C5	18:D:643:C:C5	3.06	0.43
18:D:882:C:O2'	18:D:883:C:H5'	2.18	0.43
18:D:1240:U:OP1	27:M:116:MET:HB2	2.17	0.43
18:D:1315:U:O2	18:D:1360:A:H2	2.00	0.43
22:H:5:PHE:CZ	22:H:9:PHE:CE1	3.06	0.43
23:I:151:VAL:CG1	23:I:200:VAL:HG23	2.47	0.43
25:K:76:LEU:HD12	25:K:76:LEU:O	2.17	0.43
26:L:5:GLU:CD	26:L:61:LEU:HD11	2.42	0.43
26:L:27:ALA:O	26:L:30:THR:OG1	2.29	0.43
29:O:91:ASP:HB3	29:O:94:LEU:HG	2.00	0.43
37:W:68:GLY:HA3	47:g:59:ARG:HH22	1.82	0.43
41:a:607:U:N3	41:a:608:A:N7	2.66	0.43
41:a:858:G:OP1	42:b:78:LYS:HE3	2.18	0.43
41:a:1282:U:H2'	41:a:1283:G:O4'	2.18	0.43
41:a:1349:C:O2	41:a:1349:C:H2'	2.17	0.43
41:a:1790:C:C2'	41:a:1791:A:C8	3.00	0.43
41:a:2175:C:C2'	41:a:2176:A:C8	3.00	0.43
41:a:2590:A:H2'	41:a:2591:C:H6	1.83	0.43
41:a:2615:U:H2'	41:a:2616:C:H6	1.83	0.43
41:a:2848:G:O2'	41:a:2867:G:N2	2.36	0.43
46:f:41:THR:O	46:f:43:ALA:N	2.51	0.43
50:j:107:VAL:HG13	50:j:177:VAL:HG11	2.00	0.43
56:p:94:TYR:CE1	56:p:152:ARG:CD	3.01	0.43
59:s:124:VAL:O	59:s:125:TYR:HD1	2.01	0.43
61:u:95:LEU:HD13	61:u:100:ILE:HD12	2.00	0.43
66:z:14:HIS:O	66:z:18:LEU:HD23	2.18	0.43
4:3:6:ARG:CG	4:3:7:ARG:H	2.31	0.43
5:4:53:LYS:HB2	5:4:56:PHE:HE2	1.84	0.43
9:9:19:ALA:CB	9:9:69:PHE:HE2	2.31	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:A:72:A:C3'	10:A:73:A:H5''	2.47	0.43
12:AB:160:ARG:HE	12:AB:160:ARG:HB2	1.47	0.43
14:AE:708:ASN:HA	14:AE:713:GLU:HA	2.00	0.43
16:AG:143:LYS:HG3	16:AG:153:ASP:HB2	1.99	0.43
10:B:22:G:C8	10:B:22:G:OP2	2.71	0.43
10:B:75:C:P	41:a:2602:A:C2'	3.06	0.43
18:D:636:U:H2'	18:D:637:C:C6	2.52	0.43
18:D:671:G:H4'	26:L:79:ARG:HH12	1.83	0.43
18:D:1389:C:H2'	18:D:1390:U:O4'	2.17	0.43
18:D:1434:A:H2'	18:D:1435:G:O4'	2.18	0.43
19:E:60:ARG:O	19:E:64:LYS:HD3	2.18	0.43
20:F:3:VAL:HG22	31:Q:111:THR:HG23	2.01	0.43
21:G:9:MET:HE1	21:G:47:VAL:HG22	1.99	0.43
23:I:54:ARG:NH1	23:I:69:HIS:NE2	2.66	0.43
23:I:152:GLU:OE2	23:I:165:THR:HG23	2.18	0.43
26:L:35:LYS:HD2	26:L:36:ILE:H	1.82	0.43
28:N:84:ARG:HD2	28:N:124:GLU:OE1	2.17	0.43
29:O:51:PRO:HB3	29:O:103:PHE:HD2	1.83	0.43
30:P:10:LEU:HD22	30:P:22:THR:HG22	1.98	0.43
30:P:52:LEU:HB2	33:S:81:ARG:NH1	2.33	0.43
32:R:57:LEU:HD23	32:R:59:ASN:CG	2.43	0.43
33:S:13:ARG:HB3	33:S:60:GLN:CG	2.46	0.43
38:X:11:ASP:HA	38:X:45:ILE:HB	1.99	0.43
39:Y:93:ASN:O	39:Y:94:LYS:HD2	2.16	0.43
41:a:861:A:H2'	41:a:862:G:O4'	2.17	0.43
41:a:1434:A:H2'	41:a:1435:G:C8	2.53	0.43
41:a:2092:U:C2	41:a:2225:A:O2'	2.71	0.43
41:a:2465:C:O2	41:a:2465:C:H2'	2.18	0.43
41:a:2477:U:O2	57:q:4:ARG:NH2	2.51	0.43
41:a:2747:G:N2	41:a:2756:U:C6	2.85	0.43
48:h:251:GLN:OE1	48:h:255:LYS:CB	2.65	0.43
51:k:7:GLU:OE2	51:k:27:LYS:N	2.50	0.43
52:l:73:ILE:HG13	52:l:73:ILE:O	2.17	0.43
53:m:26:ASN:HA	53:m:29:GLN:HE21	1.83	0.43
54:n:15:LYS:C	54:n:19:GLU:OE1	2.61	0.43
54:n:132:VAL:CG2	54:n:152:LEU:CD2	2.97	0.43
55:o:13:ARG:HG2	61:u:62:PRO:O	2.18	0.43
57:q:3:VAL:HG12	57:q:36:ARG:HB3	2.01	0.43
63:w:87:PHE:HE2	63:w:115:LEU:HD11	1.83	0.43
65:y:106:LYS:CB	65:y:109:ARG:HH22	2.31	0.43
1:0:13:ARG:CZ	66:z:97:ASP:CG	2.91	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:0:60:LYS:N	1:0:60:LYS:HD3	2.33	0.43
2:1:36:LEU:CD2	2:1:48:LYS:HA	2.49	0.43
4:3:7:ARG:NH2	41:a:84:A:O2'	2.51	0.43
9:9:92:ALA:HB3	9:9:129:LEU:HB2	2.00	0.43
9:9:138:ARG:NH1	40:Z:10:ALA:O	2.51	0.43
11:AA:27:LEU:O	11:AA:528:ARG:NH1	2.43	0.43
12:AB:143:LEU:HD23	12:AB:143:LEU:C	2.42	0.43
13:AD:22:THR:OG1	13:AD:207:THR:OG1	2.36	0.43
14:AE:362:ARG:H	14:AE:365:GLN:HE21	1.66	0.43
17:C:40:VAL:HG13	22:H:339:ARG:CD	2.49	0.43
17:C:40:VAL:HG22	22:H:339:ARG:HE	1.83	0.43
18:D:262:A:H2'	18:D:263:A:C8	2.53	0.43
18:D:746:A:H2'	18:D:747:A:C8	2.53	0.43
18:D:986:U:H2'	18:D:987:G:C8	2.53	0.43
18:D:1103:C:O2	21:G:106:THR:HG21	2.18	0.43
18:D:1315:U:H2'	18:D:1316:G:O4'	2.19	0.43
21:G:64:LYS:HE2	21:G:64:LYS:HA	1.99	0.43
21:G:66:LYS:HB3	21:G:154:MET:HE3	1.99	0.43
21:G:117:LEU:HD21	21:G:141:LEU:CA	2.48	0.43
23:I:58:GLU:OE2	30:P:94:ALA:HB1	2.16	0.43
24:J:49:SER:O	24:J:53:VAL:HG13	2.18	0.43
24:J:105:MET:HE3	24:J:171:LEU:HD13	1.97	0.43
25:K:50:TYR:C	25:K:66:LYS:HE3	2.43	0.43
28:N:35:ALA:HB1	28:N:110:VAL:CG1	2.48	0.43
29:O:21:ILE:CD1	29:O:86:ALA:CB	2.97	0.43
29:O:89:GLU:HG3	29:O:90:TYR:N	2.33	0.43
32:R:25:GLU:OE2	32:R:30:LYS:HE3	2.18	0.43
34:T:5:THR:O	34:T:8:THR:OG1	2.27	0.43
35:U:9:HIS:O	35:U:16:PHE:HB3	2.19	0.43
38:X:71:ARG:NH1	38:X:72:GLU:HG3	2.33	0.43
38:X:90:ARG:NH1	38:X:96:PRO:O	2.51	0.43
41:a:208:C:H2'	41:a:209:C:C6	2.53	0.43
41:a:870:U:N3	41:a:871:U:C5	2.86	0.43
41:a:963:U:H2'	41:a:964:C:C6	2.53	0.43
41:a:966:G:O4'	41:a:2267:A:N6	2.51	0.43
41:a:1409:U:C2	41:a:1410:G:C8	3.06	0.43
41:a:1657:U:C2	41:a:1658:C:C5	3.06	0.43
41:a:2615:U:C2	49:i:4:GLN:HA	2.53	0.43
49:i:55:ILE:CD1	63:w:112:TYR:CD2	3.02	0.43
54:n:6:ASP:O	54:n:10:ASP:OD2	2.36	0.43
54:n:66:LEU:HD23	54:n:67:ILE:N	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:r:6:LEU:C	58:r:15:LEU:HD13	2.43	0.43
66:z:78:LYS:HG2	66:z:117:LEU:HD21	2.00	0.43
2:1:99:ARG:CZ	2:1:99:ARG:HB2	2.49	0.43
6:5:18:DG:C4	6:5:19:DC:C5	3.06	0.43
8:7:1:A:H2'	8:7:2:U:C6	2.50	0.43
10:A:19:G:N3	10:A:57:A:C4	2.87	0.43
11:AA:855:PRO:HD2	16:AG:105:ILE:CG2	2.41	0.43
13:AD:29:GLU:HB3	13:AD:200:LYS:HG3	1.99	0.43
14:AE:412:LEU:HA	14:AE:415:VAL:HG22	2.01	0.43
16:AG:353:HIS:C	16:AG:357:ASP:HB2	2.43	0.43
10:B:31:G:H5'	18:D:1339:A:C2	2.54	0.43
18:D:283:U:C2	18:D:284:C:C6	3.06	0.43
18:D:316:C:C2	18:D:317:U:C5	3.06	0.43
18:D:564:C:OP1	32:R:12:ARG:CZ	2.66	0.43
18:D:575:G:O2'	18:D:821:G:OP2	2.34	0.43
18:D:1160:G:C6	18:D:1161:C:C5	3.06	0.43
19:E:54:MET:HB2	19:E:54:MET:HE2	1.86	0.43
22:H:123:GLU:HA	22:H:128:ARG:HA	2.01	0.43
26:L:36:ILE:HD12	26:L:36:ILE:HA	1.81	0.43
28:N:10:MET:HB2	28:N:33:LYS:HZ2	1.84	0.43
29:O:86:ALA:C	29:O:89:GLU:HG2	2.43	0.43
32:R:24:LEU:HD23	32:R:95:TYR:HE1	1.83	0.43
36:V:31:HIS:NE2	36:V:34:TYR:HD2	2.15	0.43
36:V:63:GLU:HB2	36:V:73:TRP:CZ2	2.53	0.43
38:X:22:ILE:HG23	38:X:66:GLU:CD	2.44	0.43
41:a:39:G:H2'	41:a:40:U:H6	1.83	0.43
41:a:153:U:H2'	41:a:154:U:C6	2.53	0.43
41:a:532:A:N7	41:a:2021:C:O2'	2.44	0.43
41:a:580:U:C2	41:a:581:C:C5	3.07	0.43
41:a:997:G:H5''	66:z:92:ARG:NH1	2.34	0.43
41:a:1021:A:C2	41:a:1141:U:O4	2.71	0.43
41:a:1269:A:H2'	41:a:1270:C:C6	2.54	0.43
41:a:1506:U:H2'	41:a:1507:C:C6	2.54	0.43
41:a:1900:A:N1	41:a:1970:A:C6	2.87	0.43
41:a:2345:G:N3	41:a:2381:A:H2'	2.33	0.43
41:a:2393:U:C5'	55:o:30:ARG:NH1	2.76	0.43
41:a:2665:A:C2	41:a:2666:C:C6	3.07	0.43
41:a:2846:G:H2'	41:a:2847:U:O4'	2.19	0.43
44:d:39:A:C2	44:d:44:G:C2	3.07	0.43
50:j:146:ILE:HA	50:j:159:LYS:HZ1	1.82	0.43
50:j:204:LYS:HD3	50:j:205:PRO:HD2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:r:90:LEU:HD21	58:r:94:ILE:HD11	2.00	0.43
62:v:104:GLU:OE1	62:v:104:GLU:HA	2.17	0.43
63:w:12:ARG:HE	63:w:20:MET:HE1	1.82	0.43
65:y:55:LEU:HA	65:y:77:HIS:CD2	2.54	0.43
2:1:19:LEU:HD23	2:1:19:LEU:C	2.43	0.43
3:2:56:GLU:OE1	3:2:86:THR:C	2.61	0.43
6:5:9:DG:H5'	11:AA:473:ARG:CD	2.45	0.43
9:9:70:GLU:HG3	9:9:71:CYS:N	2.34	0.43
11:AA:914:LYS:N	16:AG:105:ILE:C	2.76	0.43
11:AA:1157:GLN:HG3	11:AA:1159:VAL:HG13	2.00	0.43
16:AG:162:ILE:HD13	16:AG:167:MET:CE	2.49	0.43
16:AG:183:LEU:HD22	16:AG:195:LEU:HD13	2.00	0.43
10:B:49:G:O5'	10:B:49:G:H8	2.01	0.43
10:B:75:C:OP1	41:a:2602:A:C8	2.71	0.43
17:C:20:GLU:OE1	17:C:21:ILE:O	2.37	0.43
18:D:13:U:N3	18:D:915:A:C6	2.87	0.43
18:D:674:G:H2'	18:D:675:A:H8	1.84	0.43
18:D:1125:U:O2'	18:D:1126:U:O5'	2.32	0.43
18:D:1157:A:C2	18:D:1181:G:C5	3.06	0.43
18:D:1208:C:C2	18:D:1209:C:C6	3.07	0.43
18:D:1325:C:N3	18:D:1326:U:C5	2.87	0.43
18:D:1381:U:C4	18:D:1382:C:C5	3.07	0.43
18:D:1437:A:N3	18:D:1438:G:C8	2.86	0.43
19:E:16:LYS:HA	19:E:19:LYS:HE3	2.00	0.43
19:E:21:ASN:HB3	19:E:25:ARG:CZ	2.48	0.43
21:G:134:ALA:HA	21:G:137:ARG:HE	1.83	0.43
22:H:76:GLU:OE2	22:H:78:GLY:C	2.61	0.43
22:H:303:LEU:CD2	22:H:305:HIS:HB3	2.48	0.43
23:I:58:GLU:OE1	23:I:65:ARG:CZ	2.65	0.43
23:I:149:ILE:HG13	23:I:201:TRP:O	2.18	0.43
24:J:76:TYR:HE1	24:J:86:THR:HG21	1.83	0.43
26:L:2:ARG:HB2	26:L:4:TYR:CE1	2.53	0.43
27:M:17:LYS:NZ	27:M:18:PHE:CE1	2.78	0.43
27:M:23:LEU:N	27:M:23:LEU:CD1	2.81	0.43
35:U:7:ALA:HA	35:U:28:ARG:HD3	2.01	0.43
38:X:79:ARG:NH1	47:g:56:ARG:NH1	2.66	0.43
38:X:81:MET:HE1	38:X:92:ARG:CD	2.48	0.43
39:Y:99:LYS:CE	39:Y:140:GLU:OE2	2.66	0.43
41:a:61:C:C2	41:a:94:A:C2	3.06	0.43
41:a:307:G:N1	41:a:310:A:OP2	2.47	0.43
41:a:677:A:O2'	41:a:2071:A:H5'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:841:G:C6	41:a:938:G:C6	3.06	0.43
41:a:969:G:H2'	41:a:970:U:C6	2.54	0.43
41:a:1074:G:C6	41:a:1075:C:N4	2.87	0.43
41:a:1169:A:N3	41:a:1169:A:O4'	2.51	0.43
41:a:1657:U:H2'	41:a:1658:C:H6	1.83	0.43
41:a:2466:C:C2	41:a:2467:C:C5	3.06	0.43
41:a:2529:G:N7	57:q:32:LYS:NZ	2.63	0.43
41:a:2616:C:C2	41:a:2617:U:C6	3.06	0.43
44:d:48:U:P	64:x:30:ARG:HH22	2.41	0.43
46:f:6:LYS:HD3	46:f:6:LYS:N	2.33	0.43
52:l:23:PHE:CE2	52:l:25:GLU:CB	3.01	0.43
52:l:131:THR:HG22	52:l:160:ALA:O	2.18	0.43
54:n:33:LYS:CA	54:n:96:MET:SD	3.07	0.43
61:u:122:VAL:CG1	61:u:125:LEU:CD2	2.90	0.43
62:v:74:THR:HA	62:v:88:ASN:O	2.19	0.43
65:y:106:LYS:CA	65:y:109:ARG:NH2	2.81	0.43
9:9:6:GLN:CA	9:9:9:GLN:NE2	2.78	0.43
9:9:31:ARG:NH1	41:a:1054:A:C1'	2.80	0.43
11:AA:811:ASN:ND2	11:AA:1098:LEU:O	2.51	0.43
14:AE:859:PRO:HD2	14:AE:862:THR:HG21	2.01	0.43
16:AG:252:VAL:HG13	16:AG:256:GLY:HA2	1.99	0.43
16:AG:402:THR:C	16:AG:405:ALA:H	2.26	0.43
17:C:60:LYS:NZ	18:D:735:C:C4'	2.74	0.43
18:D:414:A:C2	18:D:415:A:H1'	2.54	0.43
18:D:679:C:N3	18:D:680:C:C5	2.87	0.43
18:D:792:A:H1'	18:D:794:A:N7	2.34	0.43
18:D:1410:A:H2'	18:D:1411:C:C6	2.54	0.43
21:G:131:LYS:O	21:G:135:LEU:HD13	2.19	0.43
21:G:171:ILE:H	21:G:171:ILE:HD12	1.82	0.43
22:H:19:ARG:HD3	22:H:19:ARG:N	2.32	0.43
23:I:69:HIS:CD2	23:I:104:ALA:HB3	2.54	0.43
23:I:132:ARG:CB	23:I:135:LYS:HE3	2.49	0.43
23:I:157:LEU:CD1	23:I:166:GLU:OE1	2.66	0.43
26:L:53:LYS:HD2	26:L:53:LYS:O	2.18	0.43
28:N:31:LYS:N	28:N:31:LYS:HD2	2.34	0.43
34:T:3:LEU:HD23	34:T:4:SER:N	2.33	0.43
36:V:27:ARG:HH22	36:V:42:THR:N	2.16	0.43
40:Z:29:LYS:O	40:Z:30:PHE:HD2	2.02	0.43
41:a:196:A:C5	41:a:805:G:C6	3.07	0.43
41:a:479:A:H4'	41:a:480:A:OP1	2.18	0.43
41:a:1095:A:O2'	41:a:1096:A:O4'	2.36	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1141:U:H4'	41:a:1142:A:O4'	2.18	0.43
41:a:1441:G:H2'	41:a:1442:U:H6	1.83	0.43
41:a:1565:C:OP1	48:h:18:LYS:HD2	2.19	0.43
41:a:1683:U:H2'	41:a:1684:G:H8	1.83	0.43
41:a:1879:C:H2'	41:a:1880:U:O4'	2.18	0.43
41:a:2462:C:C2	41:a:2463:C:C5	3.07	0.43
41:a:2768:U:OP1	59:s:85:LYS:HE3	2.19	0.43
44:d:45:A:C4	44:d:46:A:C8	3.05	0.43
44:d:69:G:C5	44:d:70:C:C6	3.06	0.43
44:d:89:U:O2'	44:d:90:C:P	2.76	0.43
48:h:183:LYS:CE	48:h:265:LYS:O	2.67	0.43
51:k:34:LEU:HB3	51:k:51:GLU:OE2	2.19	0.43
55:o:31:HIS:C	55:o:33:LEU:HG	2.43	0.43
55:o:39:LYS:HG2	55:o:42:ARG:HH12	1.82	0.43
60:t:50:GLY:O	60:t:53:LYS:NZ	2.47	0.43
61:u:103:ILE:HG23	61:u:104:GLN:HG2	1.99	0.43
62:v:1:MET:SD	62:v:44:ARG:HA	2.59	0.43
62:v:53:MET:O	62:v:57:VAL:HG12	2.18	0.43
2:1:74:ILE:HD12	2:1:74:ILE:N	2.34	0.43
3:2:56:GLU:CD	3:2:88:LYS:N	2.76	0.43
4:3:7:ARG:HG3	4:3:8:ASP:N	2.33	0.43
9:9:35:VAL:HG22	9:9:35:VAL:O	2.19	0.43
10:A:71:C:H2'	10:A:72:A:C1'	2.48	0.43
11:AA:444:ASP:OD1	11:AA:444:ASP:N	2.51	0.43
11:AA:1275:VAL:HG13	11:AA:1287:LEU:HD11	2.01	0.43
11:AA:1313:HIS:CD2	15:AF:31:GLN:HE22	2.37	0.43
12:AB:6:LEU:N	12:AB:88:VAL:HG21	2.31	0.43
12:AB:112:PRO:HA	12:AB:133:PRO:CG	2.49	0.43
12:AB:138:ARG:HG3	12:AB:154:VAL:O	2.19	0.43
14:AE:288:PRO:HD2	14:AE:291:ILE:CB	2.42	0.43
16:AG:402:THR:C	16:AG:405:ALA:HB3	2.41	0.43
17:C:60:LYS:NZ	18:D:735:C:P	2.92	0.43
18:D:34:C:H2'	18:D:35:G:H8	1.83	0.43
18:D:235:C:H2'	18:D:236:A:H8	1.84	0.43
18:D:553:A:H2'	18:D:554:A:H8	1.84	0.43
18:D:622:A:C8	18:D:623:C:C5	3.06	0.43
18:D:690:G:H2'	18:D:691:G:C8	2.54	0.43
18:D:1184:G:C2	18:D:1185:G:C8	3.07	0.43
22:H:74:ALA:HB3	22:H:83:LEU:HD23	1.99	0.43
24:J:41:HIS:ND1	24:J:44:ARG:CZ	2.82	0.43
26:L:9:MET:HE3	26:L:86:ARG:N	2.33	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:L:85:ILE:HG22	26:L:86:ARG:HG2	1.99	0.43
28:N:47:GLU:OE1	28:N:63:LEU:C	2.62	0.43
29:O:56:ASP:OD1	29:O:56:ASP:N	2.50	0.43
34:T:15:PHE:HD2	34:T:30:ALA:CB	2.31	0.43
34:T:63:ARG:HD2	34:T:67:LEU:HD21	2.00	0.43
36:V:28:PHE:CZ	36:V:37:PHE:HB3	2.53	0.43
41:a:66:C:H42	41:a:74:A:H62	1.67	0.43
41:a:74:A:N7	41:a:88:G:C5	2.87	0.43
41:a:197:A:N6	41:a:2430:A:H2'	2.34	0.43
41:a:573:U:O2'	41:a:574:A:H3'	2.19	0.43
41:a:587:C:O2	61:u:33:ARG:NH2	2.52	0.43
41:a:654:A:H61	55:o:19:LYS:CD	2.31	0.43
41:a:1028:A:N6	41:a:1125:G:H2'	2.34	0.43
41:a:1406:U:O2'	41:a:1407:G:C5'	2.66	0.43
41:a:1956:U:C2	41:a:1957:C:C6	3.06	0.43
41:a:2093:G:P	58:r:22:LYS:HG2	2.58	0.43
41:a:2884:U:O2	49:i:40:ARG:NH2	2.51	0.43
44:d:31:C:C2	44:d:32:U:C5	3.06	0.43
45:e:13:GLU:C	45:e:17:GLU:OE1	2.61	0.43
47:g:47:LYS:CE	54:n:115:ARG:HA	2.49	0.43
48:h:74:ILE:HB	48:h:96:TYR:CD1	2.53	0.43
48:h:265:LYS:HG3	48:h:266:PHE:CD2	2.52	0.43
49:i:17:ARG:CB	49:i:20:ASP:OD1	2.67	0.43
51:k:21:TYR:CE2	51:k:38:LYS:HB3	2.54	0.43
60:t:44:LYS:N	60:t:44:LYS:HD2	2.33	0.43
63:w:73:ASN:C	63:w:73:ASN:ND2	2.76	0.43
66:z:82:GLY:CA	66:z:117:LEU:HD11	2.48	0.43
3:2:6:ARG:HE	3:2:42:GLU:CD	2.27	0.43
9:9:6:GLN:CA	9:9:9:GLN:HE22	2.30	0.43
11:AA:829:THR:HG23	11:AA:1059:ARG:HG2	2.00	0.43
12:AB:90:SER:HB2	12:AB:94:HIS:CE1	2.54	0.43
12:AB:115:LYS:CD	12:AB:129:ILE:CD1	2.63	0.43
12:AB:126:PHE:CE1	12:AB:128:ALA:HB2	2.53	0.43
12:AB:138:ARG:HD2	12:AB:154:VAL:C	2.41	0.43
14:AE:975:ILE:HG22	14:AE:977:SER:H	1.83	0.43
16:AG:15:GLU:HA	16:AG:18:LEU:HB2	2.00	0.43
16:AG:126:VAL:HG13	16:AG:130:PHE:CE2	2.53	0.43
16:AG:277:ASP:HB2	16:AG:282:GLN:CB	2.49	0.43
17:C:27:ALA:HB1	17:C:30:LYS:HE3	2.00	0.43
17:C:27:ALA:CA	17:C:30:LYS:HE3	2.49	0.43
18:D:33:A:H2'	18:D:34:C:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:246:A:C2	18:D:282:A:C4	3.06	0.43
18:D:360:G:H2'	18:D:361:G:C8	2.53	0.43
18:D:562:U:C3'	32:R:14:ARG:NH1	2.81	0.43
18:D:611:C:N3	18:D:612:C:C5	2.87	0.43
18:D:621:A:H2'	18:D:622:A:C8	2.54	0.43
18:D:689:C:H2'	18:D:690:G:O4'	2.18	0.43
18:D:723:U:C2	20:F:49:LYS:HE3	2.54	0.43
18:D:1120:C:H2'	18:D:1121:U:H6	1.83	0.43
18:D:1194:U:H2'	18:D:1195:C:C6	2.54	0.43
18:D:1276:G:O5'	18:D:1276:G:H8	2.02	0.43
18:D:1328:C:H5''	38:X:28:THR:HG21	2.01	0.43
19:E:60:ARG:HG3	19:E:64:LYS:NZ	2.34	0.43
20:F:51:SER:OG	20:F:55:ARG:NH2	2.52	0.43
22:H:135:LEU:CB	22:H:157:ILE:CB	2.97	0.43
22:H:332:VAL:O	22:H:333:LEU:HB2	2.17	0.43
25:K:157:ARG:HG2	25:K:159:LYS:HD3	2.01	0.43
29:O:5:GLN:NE2	29:O:22:LYS:HZ2	2.15	0.43
29:O:39:PHE:CD1	29:O:44:ALA:HB1	2.53	0.43
31:Q:85:MET:HB3	31:Q:113:VAL:HG21	2.00	0.43
32:R:74:LEU:HD11	32:R:80:ILE:CG1	2.48	0.43
41:a:30:G:P	66:z:5:LYS:HZ3	2.28	0.43
41:a:271:G:C2	41:a:272:A:C5	3.07	0.43
41:a:996:A:N6	41:a:1160:G:C6	2.87	0.43
41:a:1141:U:O2	41:a:1142:A:N6	2.52	0.43
41:a:1360:G:C8	41:a:1361:G:C8	3.06	0.43
41:a:1386:C:H2'	41:a:1387:A:H8	1.82	0.43
41:a:1839:G:C4	41:a:1840:G:C8	3.06	0.43
41:a:2332:C:OP1	42:b:77:ARG:NH2	2.52	0.43
44:d:55:U:O2'	54:n:26:MET:CE	2.67	0.43
52:l:47:LYS:HA	52:l:51:GLU:CD	2.44	0.43
52:l:170:ARG:NH1	52:l:175:ILE:HA	2.34	0.43
59:s:88:THR:OG1	59:s:91:GLU:HG3	2.18	0.43
62:v:105:MET:HE1	62:v:113:ALA:HA	2.01	0.43
9:9:80:THR:O	41:a:1108:U:OP1	2.37	0.43
11:AA:300:ASP:OD1	11:AA:313:ALA:N	2.52	0.43
11:AA:853:ASP:HA	16:AG:102:PHE:CE1	2.54	0.43
11:AA:858:GLY:H	16:AG:38:LYS:HE3	1.84	0.43
11:AA:887:VAL:HG23	16:AG:105:ILE:CG1	2.41	0.43
13:AD:288:GLU:C	13:AD:290:LEU:H	2.26	0.43
14:AE:117:LEU:HG	14:AE:118:LYS:HG3	2.00	0.43
16:AG:10:GLU:C	16:AG:12:VAL:HG22	2.44	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:AG:398:LEU:C	16:AG:398:LEU:HD12	2.44	0.43
18:D:57:G:H2'	18:D:58:C:C6	2.54	0.43
18:D:68:G:H3'	18:D:69:G:H5''	2.01	0.43
18:D:736:C:H4'	26:L:88:MET:HE2	1.99	0.43
18:D:1244:G:C6	18:D:1294:G:C6	3.06	0.43
18:D:1464:U:C2	18:D:1465:A:C8	3.07	0.43
18:D:1513:A:H2'	18:D:1514:G:H8	1.84	0.43
21:G:30:PHE:CG	21:G:45:LYS:NZ	2.87	0.43
23:I:54:ARG:NH2	23:I:56:VAL:CG2	2.82	0.43
31:Q:20:VAL:CG1	31:Q:21:ALA:N	2.82	0.43
31:Q:47:ALA:O	31:Q:52:PHE:HB2	2.19	0.43
31:Q:87:LYS:CG	31:Q:113:VAL:CG2	2.89	0.43
35:U:12:LYS:HD2	35:U:13:LYS:HG2	2.00	0.43
41:a:125:A:C6	53:m:10:LEU:HD13	2.54	0.43
41:a:531:C:C5	41:a:2035:G:C2	3.07	0.43
41:a:1065:U:O2'	41:a:1066:U:P	2.77	0.43
41:a:1319:C:O2'	41:a:1320:C:H5'	2.18	0.43
41:a:2037:A:H2'	41:a:2038:G:C8	2.54	0.43
41:a:2061:G:N3	41:a:2063:C:C5	2.87	0.43
41:a:2262:U:OP1	41:a:2387:U:O2'	2.33	0.43
41:a:2297:A:C2	41:a:2321:U:C5	3.07	0.43
50:j:37:VAL:HG13	50:j:48:ILE:CD1	2.49	0.43
52:l:105:LEU:O	52:l:109:LEU:HG	2.18	0.43
54:n:126:GLY:HA2	54:n:163:ASP:HA	2.01	0.43
58:r:137:GLU:OE1	58:r:137:GLU:N	2.41	0.43
2:1:1:MET:SD	2:1:62:ASP:OD2	2.76	0.43
2:1:85:ILE:O	2:1:85:ILE:HG13	2.19	0.43
3:2:37:ASP:OD2	3:2:37:ASP:C	2.62	0.43
3:2:76:ARG:CZ	3:2:77:ARG:O	2.67	0.43
9:9:3:LEU:CD1	9:9:5:LEU:H	2.31	0.43
9:9:18:VAL:HG13	9:9:71:CYS:HB2	2.00	0.43
11:AA:699:LEU:HG	11:AA:799:ASN:HD22	1.84	0.43
12:AB:2:GLN:HA	12:AB:59:PHE:C	2.43	0.43
12:AB:148:LYS:HB2	12:AB:149:GLU:H	1.72	0.43
14:AE:694:SER:OG	14:AE:738:ARG:NE	2.52	0.43
10:B:19:G:N3	10:B:57:A:C4	2.87	0.43
10:B:35:A:H2'	10:B:36:U:O4'	2.18	0.43
17:C:29:LEU:CD2	17:C:59:ILE:HD13	2.48	0.43
18:D:381:C:H2'	18:D:382:A:O4'	2.19	0.43
18:D:1073:U:O2	21:G:103:ASN:ND2	2.52	0.43
18:D:1127:G:C6	18:D:1128:C:C5	3.07	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:1261:A:C5	18:D:1262:C:C6	3.07	0.43
21:G:208:ARG:NH1	21:G:211:THR:CG2	2.82	0.43
22:H:124:LEU:O	22:H:127:ILE:O	2.36	0.43
22:H:305:HIS:CG	22:H:306:VAL:H	2.36	0.43
23:I:22:TRP:CH2	23:I:32:ASN:HB2	2.53	0.43
23:I:72:ARG:O	23:I:75:ILE:HG22	2.19	0.43
25:K:45:ARG:CG	25:K:45:ARG:NE	2.72	0.43
25:K:97:GLN:NE2	25:K:98:PRO:O	2.52	0.43
25:K:157:ARG:NH2	28:N:43:GLU:OE2	2.46	0.43
26:L:1:MET:HE2	26:L:67:PRO:N	2.34	0.43
26:L:2:ARG:C	26:L:65:GLU:OE1	2.62	0.43
26:L:25:TYR:O	26:L:26:THR:C	2.61	0.43
30:P:45:ARG:HD3	30:P:47:GLU:OE2	2.19	0.43
30:P:64:GLN:HG2	33:S:101:TRP:HH2	1.84	0.43
32:R:18:LYS:HD3	32:R:19:SER:N	2.33	0.43
34:T:63:ARG:HD2	34:T:67:LEU:HD23	2.01	0.43
34:T:70:LEU:CD2	34:T:78:TYR:HA	2.49	0.43
37:W:17:LYS:O	37:W:21:LYS:HD3	2.19	0.43
37:W:44:MET:HE1	37:W:62:VAL:HG22	2.01	0.43
38:X:71:ARG:CZ	47:g:50:ASP:O	2.67	0.43
41:a:133:U:H2'	41:a:134:G:C8	2.54	0.43
41:a:192:C:C5	41:a:193:U:C6	3.07	0.43
41:a:310:A:C2	41:a:330:A:C4	3.07	0.43
41:a:353:C:C2	41:a:354:A:C8	3.07	0.43
41:a:407:G:C2	41:a:408:G:C8	3.07	0.43
41:a:685:A:N1	41:a:787:C:H1'	2.33	0.43
41:a:814:C:C2	41:a:815:C:C5	3.07	0.43
41:a:1076:C:H2'	41:a:1077:A:C8	2.54	0.43
41:a:1139:G:O3'	59:s:26:GLY:HA3	2.19	0.43
41:a:1144:A:C5	41:a:1145:C:C5	3.07	0.43
41:a:1145:C:H2'	41:a:1146:C:H6	1.84	0.43
41:a:1996:C:OP1	60:t:31:ARG:NE	2.52	0.43
41:a:2756:U:N3	41:a:2758:A:C6	2.87	0.43
41:a:2806:C:H2'	41:a:2807:U:O4'	2.18	0.43
41:a:2884:U:C2	49:i:40:ARG:CZ	3.02	0.43
43:c:76:GLU:CD	43:c:76:GLU:C	2.87	0.43
52:l:6:LYS:NZ	52:l:119:ILE:C	2.77	0.43
62:v:31:PHE:C	62:v:104:GLU:OE1	2.62	0.43
62:v:105:MET:HG2	62:v:117:PHE:CZ	2.54	0.43
63:w:33:ILE:HG13	63:w:112:TYR:HD1	1.83	0.43
66:z:76:TYR:OH	66:z:92:ARG:HG2	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1:24:ILE:HG21	2:1:36:LEU:CD1	2.42	0.42
2:1:31:GLN:H	2:1:31:GLN:CD	2.22	0.42
2:1:75:PHE:HE2	2:1:77:ASP:OD2	2.02	0.42
2:1:95:ARG:C	2:1:95:ARG:HD2	2.44	0.42
3:2:3:ARG:HG3	3:2:5:GLU:OE1	2.19	0.42
4:3:7:ARG:O	4:3:25:VAL:O	2.36	0.42
10:A:71:C:C3'	10:A:72:A:C8	2.99	0.42
11:AA:646:SER:OG	11:AA:647:ARG:N	2.52	0.42
12:AB:140:MET:CE	30:P:102:LEU:CG	2.85	0.42
14:AE:66:LYS:HB2	14:AE:69:GLU:HB2	2.01	0.42
16:AG:224:LYS:HB2	16:AG:276:TRP:CH2	2.54	0.42
16:AG:226:ALA:O	16:AG:228:ARG:N	2.52	0.42
16:AG:234:ALA:HB3	16:AG:271:ILE:HD13	2.01	0.42
16:AG:431:ALA:O	16:AG:435:LEU:HG	2.18	0.42
17:C:33:ILE:O	22:H:338:GLU:HB3	2.19	0.42
18:D:562:U:C2'	32:R:14:ARG:NH1	2.78	0.42
18:D:687:A:C2	18:D:704:A:C6	3.07	0.42
18:D:964:A:C2	18:D:969:A:N3	2.87	0.42
18:D:1107:C:OP1	23:I:172:ARG:HG3	2.19	0.42
18:D:1408:A:H2'	18:D:1409:C:H6	1.84	0.42
18:D:1437:A:C4	18:D:1438:G:C8	3.08	0.42
22:H:31:ILE:O	22:H:31:ILE:HG13	2.18	0.42
23:I:149:ILE:HG13	23:I:202:ILE:HG12	2.01	0.42
27:M:18:PHE:HD2	27:M:59:LEU:CD2	2.31	0.42
32:R:59:ASN:OD1	32:R:59:ASN:C	2.62	0.42
41:a:78:U:C5'	45:e:4:LYS:NZ	2.81	0.42
41:a:320:A:N3	52:l:163:ASN:ND2	2.67	0.42
41:a:538:A:H2'	41:a:539:G:O4'	2.18	0.42
41:a:858:G:P	42:b:78:LYS:HZ1	2.39	0.42
41:a:1009:A:OP2	59:s:39:LYS:NZ	2.37	0.42
41:a:1114:C:H2'	41:a:1115:G:C8	2.54	0.42
41:a:1511:G:H2'	41:a:1512:C:H6	1.84	0.42
41:a:1695:G:N7	48:h:14:ARG:NH2	2.66	0.42
41:a:1707:G:C5	41:a:1708:C:C5	3.07	0.42
41:a:2216:G:H2'	41:a:2217:G:H8	1.83	0.42
41:a:2332:C:P	42:b:77:ARG:HH21	2.42	0.42
41:a:2467:C:OP1	57:q:8:LYS:NZ	2.45	0.42
41:a:2516:A:O2'	41:a:2517:C:H5'	2.18	0.42
41:a:2845:U:H5''	65:y:52:ASN:O	2.19	0.42
44:d:57:A:N7	54:n:26:MET:CE	2.82	0.42
46:f:10:THR:CB	46:f:56:LYS:NZ	2.81	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:h:94:VAL:HG12	48:h:95:LEU:N	2.34	0.42
48:h:130:LEU:CD2	48:h:192:LEU:HD21	2.48	0.42
54:n:20:PHE:HB3	54:n:22:TYR:CE1	2.54	0.42
56:p:94:TYR:HE1	56:p:152:ARG:HD2	1.83	0.42
56:p:149:ARG:HD2	56:p:162:VAL:O	2.18	0.42
59:s:1:MET:SD	59:s:2:LYS:C	3.02	0.42
60:t:7:MET:SD	60:t:18:ARG:HG3	2.59	0.42
60:t:59:LYS:NZ	60:t:92:GLU:CG	2.82	0.42
63:w:12:ARG:NH1	63:w:20:MET:HE1	2.33	0.42
66:z:76:TYR:CZ	66:z:80:ILE:HD11	2.54	0.42
5:4:2:PHE:CD1	5:4:56:PHE:CE1	3.08	0.42
6:5:10:DT:C4	11:AA:62:TYR:HE2	2.27	0.42
8:7:49:G:C2'	8:7:50:U:H5'	2.49	0.42
10:A:11:A:H2'	10:A:12:G:C8	2.54	0.42
10:A:19:G:N2	41:a:2112:G:C1'	2.82	0.42
10:A:19:G:O6	41:a:2111:U:H3'	2.18	0.42
10:A:47:U:O2'	10:A:50:U:OP1	2.37	0.42
10:A:49:G:H8	10:A:49:G:O5'	2.01	0.42
11:AA:22:LEU:HB3	11:AA:655:VAL:HG11	2.01	0.42
11:AA:125:GLY:H	11:AA:495:ALA:HB1	1.84	0.42
12:AB:4:TRP:CE2	12:AB:27:ASN:ND2	2.87	0.42
13:AC:48:LEU:HD23	13:AC:48:LEU:HA	1.87	0.42
14:AE:513:MET:HE1	14:AE:579:LEU:HD13	2.01	0.42
16:AG:288:MET:HE3	16:AG:288:MET:C	2.45	0.42
10:B:11:A:H2'	10:B:12:G:C8	2.54	0.42
10:B:12:G:H2'	10:B:13:C:OP1	2.19	0.42
10:B:47:U:O2'	10:B:50:U:OP1	2.37	0.42
10:B:57:A:O5'	10:B:58:A:P	2.77	0.42
18:D:119:A:C2	18:D:240:G:C8	3.07	0.42
18:D:320:A:H2'	18:D:321:A:O4'	2.19	0.42
18:D:384:G:H2'	18:D:385:C:C6	2.54	0.42
18:D:393:A:C2	18:D:394:G:C8	3.07	0.42
18:D:1526:G:P	20:F:42:THR:HG23	2.59	0.42
19:E:48:GLN:CD	19:E:49:LYS:N	2.77	0.42
20:F:39:GLU:OE2	20:F:44:GLU:HA	2.18	0.42
21:G:26:LYS:CE	21:G:193:PRO:HG2	2.49	0.42
23:I:112:ASP:O	23:I:116:VAL:HG23	2.18	0.42
26:L:29:ILE:HD12	26:L:29:ILE:H	1.83	0.42
30:P:45:ARG:CD	30:P:47:GLU:OE2	2.67	0.42
31:Q:35:THR:HA	31:Q:42:LEU:HG	2.01	0.42
32:R:80:ILE:CG2	32:R:81:LEU:N	2.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:W:12:ASP:OD2	37:W:35:SER:CB	2.66	0.42
37:W:40:ILE:HG22	37:W:67:VAL:HA	2.01	0.42
38:X:64:VAL:O	38:X:64:VAL:HG13	2.19	0.42
38:X:101:ARG:C	38:X:103:LYS:H	2.26	0.42
39:Y:14:ALA:C	39:Y:50:LYS:HD2	2.44	0.42
39:Y:23:VAL:HG23	39:Y:27:LEU:HD23	2.01	0.42
41:a:533:G:H2'	41:a:534:U:C6	2.54	0.42
41:a:838:C:C2	41:a:839:U:C6	3.07	0.42
41:a:1212:G:O2'	41:a:1236:G:N2	2.47	0.42
41:a:1365:A:O5'	43:c:28:ARG:NH2	2.51	0.42
41:a:1509:A:N3	41:a:1510:G:C8	2.87	0.42
41:a:1778:U:C4	41:a:1784:A:C4	3.07	0.42
41:a:2557:G:H2'	41:a:2558:C:H6	1.84	0.42
41:a:2579:C:O3'	50:j:137:SER:OG	2.32	0.42
45:e:37:LEU:HD21	45:e:43:LEU:CD2	2.49	0.42
50:j:81:GLU:O	50:j:82:PHE:HD1	2.01	0.42
50:j:91:THR:HG22	50:j:92:VAL:N	2.34	0.42
50:j:156:PHE:CE2	59:s:81:ILE:HD13	2.54	0.42
54:n:12:VAL:HG13	54:n:172:ALA:CB	2.49	0.42
54:n:38:MET:HE3	54:n:57:LEU:CG	2.49	0.42
54:n:73:SER:OG	54:n:80:ARG:HA	2.19	0.42
56:p:45:HIS:HA	56:p:50:LEU:HD13	2.01	0.42
59:s:125:TYR:HE2	59:s:130:HIS:CB	2.32	0.42
59:s:125:TYR:CD2	59:s:130:HIS:HB3	2.54	0.42
60:t:7:MET:C	60:t:18:ARG:NH1	2.77	0.42
65:y:27:GLU:HB3	65:y:87:LYS:HZ1	1.83	0.42
6:5:20:DC:H1'	6:5:21:DA:H5'	2.00	0.42
9:9:15:VAL:HG22	9:9:66:GLY:HA3	2.00	0.42
10:A:18:G:H2'	10:A:57:A:C2	2.55	0.42
10:A:22:G:OP2	10:A:22:G:C8	2.72	0.42
10:A:32:C:O3'	27:M:86:GLN:NE2	2.51	0.42
14:AE:1036:ARG:HE	14:AE:1081:VAL:HG21	1.83	0.42
16:AG:216:ILE:HG13	16:AG:216:ILE:H	1.63	0.42
17:C:44:ILE:CG2	22:H:340:ARG:HB2	2.49	0.42
18:D:120:A:O2'	18:D:122:G:N7	2.48	0.42
18:D:404:G:N7	24:J:2:ALA:HB3	2.33	0.42
18:D:1149:C:N3	18:D:1150:A:N7	2.67	0.42
18:D:1317:C:OP2	33:S:28:LYS:CE	2.66	0.42
20:F:14:VAL:O	20:F:18:ARG:HG3	2.19	0.42
21:G:55:ALA:O	21:G:59:LYS:HG2	2.19	0.42
28:N:66:PHE:C	28:N:67:GLN:OE1	2.62	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:X:71:ARG:NH1	47:g:51:VAL:O	2.52	0.42
39:Y:14:ALA:O	39:Y:15:GLY:C	2.61	0.42
41:a:1198:U:H2'	41:a:1199:U:H6	1.83	0.42
41:a:1665:A:C5'	60:t:66:LYS:HZ2	2.32	0.42
41:a:1827:U:O2'	41:a:1828:G:H5'	2.19	0.42
41:a:1863:G:H2'	41:a:1864:U:H6	1.83	0.42
41:a:2002:G:OP1	63:w:13:ASN:HA	2.19	0.42
41:a:2215:C:H2'	41:a:2216:G:C8	2.54	0.42
41:a:2305:U:H5''	54:n:131:GLY:HA3	2.00	0.42
41:a:2387:U:C4'	42:b:41:ARG:NH1	2.81	0.42
41:a:2547:A:H4'	60:t:29:HIS:NE2	2.34	0.42
41:a:2547:A:H5'	60:t:29:HIS:NE2	2.34	0.42
48:h:71:LYS:CB	48:h:96:TYR:CZ	3.03	0.42
52:l:170:ARG:CZ	52:l:174:GLY:O	2.66	0.42
55:o:45:ARG:HD2	55:o:45:ARG:C	2.43	0.42
58:r:93:SER:OG	58:r:121:VAL:HG13	2.18	0.42
62:v:49:ALA:O	62:v:50:ARG:C	2.62	0.42
66:z:91:ASP:O	66:z:95:LEU:HG	2.20	0.42
10:A:57:A:O5'	10:A:58:A:P	2.77	0.42
12:AB:26:VAL:HG13	12:AB:58:GLU:H	1.84	0.42
13:AC:228:LEU:HD21	13:AD:224:LEU:HD23	2.01	0.42
13:AD:86:LYS:NZ	14:AE:532:GLU:OE2	2.44	0.42
14:AE:609:TYR:HD2	14:AE:610:ARG:HD2	1.84	0.42
16:AG:343:ASP:HB3	16:AG:347:LYS:HE3	2.01	0.42
16:AG:440:VAL:HG13	16:AG:444:LEU:HB2	2.00	0.42
17:C:34:THR:HB	17:C:38:LYS:N	2.35	0.42
18:D:43:C:P	35:U:12:LYS:HE3	2.59	0.42
18:D:124:C:H2'	18:D:125:U:H6	1.84	0.42
18:D:195:A:H5''	19:E:64:LYS:HZ1	1.84	0.42
18:D:256:U:H2'	18:D:257:G:C8	2.55	0.42
18:D:505:G:C2	18:D:506:G:C5	3.07	0.42
18:D:520:A:C2'	32:R:70:GLU:OE1	2.67	0.42
18:D:619:U:H1'	24:J:128:ARG:NH1	2.34	0.42
18:D:619:U:O4'	24:J:128:ARG:CZ	2.67	0.42
18:D:796:C:N3	18:D:797:C:C5	2.87	0.42
18:D:1073:U:C2	18:D:1074:G:C8	3.07	0.42
18:D:1152:A:P	30:P:72:ARG:HH22	2.43	0.42
18:D:1318:A:OP1	37:W:7:LYS:NZ	2.52	0.42
18:D:1329:A:OP1	38:X:28:THR:CB	2.66	0.42
18:D:1382:C:O2	18:D:1382:C:H2'	2.19	0.42
19:E:9:LYS:CG	19:E:13:GLN:NE2	2.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:H:74:ALA:HB3	22:H:83:LEU:CD2	2.50	0.42
23:I:23:PHE:CD2	30:P:97:ASP:HB2	2.53	0.42
23:I:156:ARG:O	23:I:156:ARG:HG3	2.20	0.42
25:K:68:ARG:O	25:K:69:ARG:NH1	2.52	0.42
25:K:100:SER:O	25:K:103:THR:HG22	2.20	0.42
25:K:132:ASN:OD1	25:K:134:ILE:HG22	2.18	0.42
27:M:99:LEU:O	27:M:103:TRP:CG	2.73	0.42
29:O:35:LEU:CD2	29:O:45:ARG:HG2	2.50	0.42
29:O:80:ARG:O	29:O:84:THR:HG23	2.18	0.42
30:P:7:ARG:HA	30:P:75:ASP:OD1	2.20	0.42
30:P:56:HIS:C	30:P:57:VAL:HG23	2.44	0.42
32:R:116:LYS:O	32:R:117:TYR:HD1	2.02	0.42
35:U:5:ARG:HA	35:U:71:VAL:HG21	2.01	0.42
36:V:43:LYS:C	36:V:44:LEU:HD12	2.45	0.42
38:X:25:VAL:O	38:X:25:VAL:HG13	2.19	0.42
39:Y:32:VAL:HG13	39:Y:58:ILE:HG21	2.00	0.42
39:Y:99:LYS:HD2	39:Y:140:GLU:OE2	2.19	0.42
39:Y:102:ARG:NH1	39:Y:139:VAL:HG11	2.33	0.42
41:a:136:G:H2'	41:a:137:U:H6	1.85	0.42
41:a:471:A:H2'	41:a:472:A:O4'	2.19	0.42
41:a:973:A:O4'	41:a:1188:U:C6	2.72	0.42
41:a:1064:C:H2'	41:a:1065:U:C6	2.55	0.42
41:a:1665:A:O3'	60:t:66:LYS:CD	2.67	0.42
41:a:1792:G:H5'	48:h:204:VAL:CG2	2.49	0.42
41:a:1906:G:C8	41:a:1929:G:C5	3.08	0.42
41:a:2093:G:P	58:r:22:LYS:CG	3.07	0.42
41:a:2228:G:H2'	41:a:2229:U:C6	2.54	0.42
41:a:2249:U:H3'	41:a:2250:G:C5'	2.50	0.42
41:a:2273:A:H2'	41:a:2274:A:C8	2.54	0.42
41:a:2287:A:N7	41:a:2289:G:C8	2.87	0.42
41:a:2554:U:H2'	41:a:2555:U:C6	2.55	0.42
41:a:2821:A:C2	41:a:2822:G:C4	3.07	0.42
45:e:8:GLU:O	45:e:9:LYS:HD3	2.19	0.42
49:i:4:GLN:OE1	49:i:8:PRO:HD3	2.19	0.42
50:j:16:THR:HG23	50:j:20:VAL:O	2.19	0.42
50:j:61:THR:HG23	50:j:63:PRO:HD2	2.00	0.42
54:n:104:ILE:HD11	54:n:174:ASP:O	2.19	0.42
55:o:30:ARG:CZ	61:u:62:PRO:HB3	2.50	0.42
56:p:70:ALA:HA	56:p:73:ASN:HD22	1.84	0.42
58:r:8:LYS:NZ	58:r:9:VAL:O	2.52	0.42
58:r:114:GLU:OE2	58:r:133:GLN:N	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:t:110:GLU:OE1	60:t:110:GLU:HA	2.19	0.42
62:v:112:LEU:HD12	62:v:112:LEU:N	2.34	0.42
63:w:70:THR:O	63:w:71:ARG:HB2	2.20	0.42
66:z:111:GLU:OE2	66:z:112:LYS:HD2	2.19	0.42
1:0:77:PHE:CD1	1:0:77:PHE:C	2.98	0.42
4:3:35:ILE:HD11	4:3:62:GLU:C	2.44	0.42
9:9:87:GLU:OE1	9:9:91:ALA:O	2.38	0.42
11:AA:478:ARG:HD2	11:AA:492:MET:HA	2.02	0.42
11:AA:861:ALA:HB1	11:AA:882:ILE:HD13	2.01	0.42
12:AB:81:PHE:CD1	14:AE:141:PHE:O	2.72	0.42
13:AD:46:ILE:HD11	13:AD:224:LEU:HD13	2.02	0.42
14:AE:287:ALA:HB2	14:AE:292:VAL:HG11	1.85	0.42
18:D:250:A:O4'	18:D:252:U:C6	2.72	0.42
18:D:427:U:OP2	18:D:428:G:O2'	2.31	0.42
18:D:563:A:C2	18:D:567:G:C6	3.07	0.42
18:D:662:U:H2'	18:D:663:A:C8	2.55	0.42
18:D:768:A:C5	18:D:769:G:C8	3.08	0.42
18:D:877:G:H5''	28:N:80:ARG:CD	2.49	0.42
18:D:1302:C:OP1	38:X:13:LYS:NZ	2.30	0.42
19:E:50:ALA:O	19:E:54:MET:HG3	2.20	0.42
21:G:134:ALA:HA	21:G:137:ARG:NE	2.35	0.42
22:H:308:GLU:OE1	22:H:308:GLU:N	2.52	0.42
24:J:67:VAL:CG1	24:J:71:GLN:HB3	2.50	0.42
24:J:67:VAL:C	24:J:72:PHE:CE2	2.98	0.42
24:J:117:LEU:HD23	24:J:154:ARG:HH21	1.84	0.42
25:K:133:PRO:HA	25:K:136:VAL:HG12	2.02	0.42
26:L:18:VAL:HA	26:L:21:MET:CE	2.47	0.42
30:P:40:ILE:HD12	30:P:73:LEU:HD22	2.02	0.42
35:U:19:VAL:HG11	35:U:52:LEU:CD2	2.50	0.42
37:W:66:MET:SD	37:W:66:MET:C	3.03	0.42
39:Y:99:LYS:CE	39:Y:138:VAL:O	2.67	0.42
41:a:6:A:O3'	59:s:132:HIS:HE1	2.03	0.42
41:a:136:G:H2'	41:a:137:U:C6	2.54	0.42
41:a:209:C:N3	41:a:210:C:C5	2.88	0.42
41:a:780:G:O2'	41:a:783:A:N6	2.53	0.42
41:a:1588:G:N1	41:a:1589:U:O4	2.53	0.42
41:a:1589:U:O2'	41:a:1590:A:O5'	2.29	0.42
41:a:1709:U:H2'	41:a:1710:G:C8	2.55	0.42
41:a:2052:A:O2'	50:j:148:GLN:O	2.37	0.42
41:a:2813:A:H2'	41:a:2814:A:H8	1.84	0.42
41:a:2859:G:H2'	41:a:2860:A:C8	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:d:39:A:C2	44:d:44:G:C4	3.07	0.42
48:h:91:ILE:HG22	48:h:92:ALA:N	2.33	0.42
52:l:6:LYS:NZ	52:l:119:ILE:CG2	2.83	0.42
53:m:25:LYS:HG3	53:m:29:GLN:NE2	2.34	0.42
60:t:1:MET:HE1	60:t:32:TYR:CB	2.49	0.42
62:v:95:LEU:C	62:v:96:ILE:HD13	2.45	0.42
64:x:83:LEU:HD23	64:x:88:LYS:HD3	2.02	0.42
65:y:106:LYS:HB3	65:y:109:ARG:HH22	1.84	0.42
66:z:12:ALA:O	66:z:16:LYS:HG2	2.19	0.42
2:1:98:LYS:CD	41:a:2012:G:OP1	2.63	0.42
3:2:11:LEU:HD11	3:2:46:ALA:CB	2.50	0.42
3:2:42:GLU:OE1	3:2:42:GLU:HA	2.19	0.42
5:4:80:HIS:HE1	5:4:82:TYR:CG	2.37	0.42
6:5:16:DG:N1	11:AA:451:ARG:NH2	2.66	0.42
9:9:105:LYS:O	9:9:106:PHE:HB3	2.18	0.42
12:AB:120:GLU:H	12:AB:162:LEU:CD1	2.33	0.42
13:AD:31:LEU:HD21	13:AD:39:LEU:HD12	2.00	0.42
14:AE:97:VAL:HG12	14:AE:101:ARG:HG3	2.01	0.42
16:AG:10:GLU:CA	16:AG:12:VAL:HG22	2.50	0.42
16:AG:161:VAL:HB	16:AG:196:PHE:CE1	2.54	0.42
16:AG:208:LEU:HA	16:AG:208:LEU:HD23	1.78	0.42
16:AG:288:MET:HE2	16:AG:288:MET:HB3	1.75	0.42
16:AG:318:ILE:CG1	16:AG:338:VAL:HG11	2.48	0.42
10:B:18:G:H2'	10:B:57:A:C2	2.55	0.42
18:D:119:A:C4	18:D:240:G:N7	2.88	0.42
18:D:306:A:C4	18:D:307:C:C6	3.08	0.42
18:D:552:U:H2'	18:D:553:A:H8	1.84	0.42
18:D:953:G:O6	38:X:103:LYS:HE3	2.18	0.42
20:F:31:GLU:CD	20:F:34:ARG:NH2	2.77	0.42
21:G:101:LEU:HD22	21:G:157:LEU:HD11	1.99	0.42
22:H:34:ASP:OD1	22:H:50:ALA:HB3	2.20	0.42
22:H:308:GLU:C	22:H:310:ASP:H	2.25	0.42
22:H:310:ASP:HB2	22:H:316:ILE:HD11	2.00	0.42
22:H:330:VAL:HB	22:H:344:LEU:HD23	2.01	0.42
22:H:331:MET:SD	22:H:348:GLN:HG2	2.60	0.42
24:J:139:PRO:N	24:J:182:PHE:CE2	2.88	0.42
24:J:153:SER:OG	24:J:156:LYS:NZ	2.53	0.42
24:J:170:TRP:CD1	24:J:171:LEU:HG	2.54	0.42
26:L:88:MET:SD	26:L:90:MET:HE3	2.60	0.42
29:O:50:GLN:OE1	29:O:50:GLN:HA	2.20	0.42
32:R:57:LEU:HG	32:R:58:THR:N	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:T:10:LYS:CE	34:T:10:LYS:HA	2.49	0.42
34:T:10:LYS:HA	34:T:10:LYS:HE2	2.00	0.42
34:T:70:LEU:HG	34:T:78:TYR:HB2	2.02	0.42
35:U:8:ARG:HD3	35:U:17:TYR:CE1	2.54	0.42
41:a:340:A:H2'	41:a:341:C:O4'	2.19	0.42
41:a:532:A:OP1	66:z:45:TYR:OH	2.32	0.42
41:a:579:G:H2'	41:a:580:U:H6	1.85	0.42
41:a:1443:U:H2'	41:a:1444:G:C8	2.54	0.42
41:a:1587:G:H2'	41:a:1588:G:C8	2.54	0.42
41:a:1770:G:C5	41:a:1983:G:C6	3.07	0.42
41:a:1773:A:C8	41:a:1829:A:C8	3.08	0.42
41:a:2175:C:H2'	41:a:2176:A:N7	2.33	0.42
41:a:2571:U:O2'	50:j:151:THR:HG23	2.20	0.42
41:a:2759:G:O2'	56:p:35:ARG:NH1	2.53	0.42
42:b:40:GLN:NE2	42:b:59:LEU:HD23	2.35	0.42
44:d:82:U:O2	46:f:53:PHE:HZ	2.02	0.42
45:e:56:LEU:C	45:e:59:GLU:HG3	2.44	0.42
52:l:77:ILE:O	52:l:77:ILE:HG22	2.19	0.42
54:n:66:LEU:HD23	54:n:67:ILE:C	2.45	0.42
59:s:15:TRP:CE3	59:s:135:GLN:HB3	2.55	0.42
59:s:92:MET:HB2	59:s:92:MET:HE3	1.13	0.42
62:v:115:GLU:HA	62:v:118:LYS:HE2	2.02	0.42
63:w:18:GLN:HE22	63:w:22:ARG:NE	2.18	0.42
63:w:58:ASP:HB2	63:w:80:PHE:CE1	2.54	0.42
65:y:37:LYS:CA	65:y:37:LYS:HE3	2.49	0.42
66:z:47:TYR:CZ	66:z:51:ARG:NH2	2.87	0.42
10:A:5:G:C2'	10:A:6:G:C5'	2.98	0.42
12:AB:120:GLU:H	12:AB:162:LEU:CG	2.31	0.42
12:AB:145:LEU:HG	30:P:88:MET:HE2	2.01	0.42
12:AB:161:LYS:HD3	12:AB:161:LYS:HA	1.55	0.42
16:AG:36:LYS:HB3	16:AG:36:LYS:HE2	1.87	0.42
16:AG:244:ARG:N	16:AG:244:ARG:HE	2.18	0.42
16:AG:277:ASP:HB2	16:AG:282:GLN:CG	2.50	0.42
17:C:51:TYR:HD1	17:C:54:GLN:HE22	1.58	0.42
18:D:21:G:H2'	18:D:22:G:C8	2.54	0.42
18:D:197:A:N6	18:D:221:C:H4'	2.35	0.42
18:D:323:U:H2'	18:D:324:G:O4'	2.19	0.42
18:D:374:A:C6	18:D:375:U:C4	3.08	0.42
18:D:416:G:C4	18:D:417:G:C8	3.07	0.42
18:D:501:C:H1'	18:D:549:C:H1'	2.02	0.42
18:D:684:U:H2'	18:D:685:G:O4'	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:951:G:C6	18:D:1231:G:C6	3.08	0.42
18:D:986:U:H2'	18:D:987:G:O4'	2.19	0.42
18:D:1035:A:C6	18:D:1036:A:N7	2.88	0.42
18:D:1202:U:N3	33:S:82:ILE:HD13	2.35	0.42
18:D:1381:U:N3	18:D:1382:C:C5	2.88	0.42
23:I:72:ARG:HD3	23:I:75:ILE:H	1.84	0.42
25:K:88:VAL:HG12	25:K:93:ARG:HG2	2.02	0.42
26:L:1:MET:CE	26:L:67:PRO:HD3	2.50	0.42
29:O:118:LEU:HD23	29:O:118:LEU:HA	1.87	0.42
35:U:69:ASP:C	35:U:69:ASP:OD1	2.63	0.42
36:V:31:HIS:HD2	36:V:34:TYR:H	1.62	0.42
41:a:235:U:C2	41:a:236:C:C6	3.08	0.42
41:a:250:G:H2'	41:a:251:A:C8	2.54	0.42
41:a:367:G:C4	41:a:368:A:C8	3.07	0.42
41:a:704:G:O2'	41:a:726:G:N2	2.45	0.42
41:a:751:A:C6	41:a:789:A:C5	3.08	0.42
41:a:826:U:O2'	61:u:53:GLY:HA3	2.20	0.42
41:a:1385:A:C4	41:a:1403:A:C2	3.06	0.42
41:a:1789:A:OP2	48:h:221:ARG:NH1	2.52	0.42
41:a:2049:G:H21	50:j:161:MET:HE2	1.84	0.42
41:a:2066:C:O2'	41:a:2067:G:H5'	2.20	0.42
41:a:2097:A:H2'	41:a:2098:U:C6	2.55	0.42
41:a:2226:C:H2'	41:a:2227:A:O4'	2.20	0.42
41:a:2708:G:O4'	63:w:71:ARG:NH1	2.52	0.42
50:j:55:LYS:NZ	50:j:56:LYS:O	2.49	0.42
52:l:31:VAL:HG21	52:l:104:ALA:CB	2.50	0.42
56:p:109:PHE:CD1	56:p:152:ARG:NH1	2.88	0.42
62:v:71:LYS:C	62:v:92:TRP:CE3	2.97	0.42
63:w:45:ARG:O	63:w:49:GLU:CD	2.62	0.42
9:9:69:PHE:O	9:9:72:LEU:HD11	2.20	0.42
9:9:94:ARG:N	9:9:94:ARG:HD3	2.34	0.42
12:AB:3:SER:HB3	12:AB:5:TYR:CE2	2.55	0.42
12:AB:104:ILE:C	12:AB:106:ASP:H	2.27	0.42
12:AB:130:PHE:HB3	12:AB:133:PRO:HD3	2.00	0.42
12:AB:136:GLU:HG3	12:AB:160:ARG:HH12	1.84	0.42
14:AE:211:GLU:HA	14:AE:214:ARG:HD2	2.01	0.42
14:AE:907:HIS:ND1	14:AE:908:ILE:O	2.48	0.42
10:B:31:G:H2'	10:B:32:C:C6	2.55	0.42
18:D:299:G:H2'	18:D:300:A:C8	2.55	0.42
18:D:407:U:C2	18:D:408:A:C8	3.08	0.42
18:D:415:A:C4	18:D:416:G:C8	3.07	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:539:A:H2'	18:D:540:G:C8	2.54	0.42
18:D:611:C:H2'	18:D:611:C:O2	2.18	0.42
18:D:946:A:C2	18:D:947:G:C5	3.08	0.42
18:D:1319:A:C8	18:D:1323:G:C5	3.08	0.42
18:D:1499:A:H2'	18:D:1500:A:H8	1.84	0.42
21:G:115:LYS:C	21:G:115:LYS:HD3	2.45	0.42
23:I:14:ILE:HB	23:I:178:LEU:HD21	2.01	0.42
24:J:69:GLU:HA	24:J:72:PHE:HD2	1.83	0.42
25:K:111:MET:O	25:K:112:ARG:C	2.63	0.42
26:L:94:HIS:CG	26:L:95:ALA:N	2.88	0.42
29:O:116:VAL:HG11	30:P:62:ARG:HD3	2.02	0.42
31:Q:21:ALA:HA	31:Q:34:ILE:HD13	2.01	0.42
32:R:79:VAL:O	32:R:103:ASP:HB2	2.20	0.42
34:T:77:ARG:HA	34:T:77:ARG:NE	2.35	0.42
35:U:67:ILE:O	35:U:67:ILE:HG13	2.20	0.42
39:Y:45:THR:HG21	39:Y:50:LYS:NZ	2.34	0.42
39:Y:82:ALA:HB3	39:Y:108:ILE:HD12	2.01	0.42
39:Y:102:ARG:CZ	39:Y:139:VAL:HG21	2.50	0.42
41:a:299:A:C5	41:a:322:A:C2	3.08	0.42
41:a:301:G:C4	41:a:302:C:C5	3.07	0.42
41:a:826:U:OP1	41:a:2428:G:H3'	2.20	0.42
41:a:1080:A:C6	41:a:1081:U:C4	3.07	0.42
48:h:71:LYS:HB2	48:h:96:TYR:CZ	2.54	0.42
49:i:55:ILE:HD13	63:w:112:TYR:CE2	2.54	0.42
50:j:57:ALA:O	50:j:60:VAL:HG12	2.19	0.42
50:j:100:LEU:O	50:j:100:LEU:HD12	2.20	0.42
50:j:170:VAL:HG13	50:j:194:PRO:CG	2.50	0.42
52:l:108:ILE:HD11	52:l:180:LEU:CD1	2.50	0.42
54:n:93:GLY:O	54:n:94:GLU:C	2.61	0.42
54:n:132:VAL:HG11	54:n:137:ILE:HD11	2.01	0.42
55:o:29:LEU:O	55:o:29:LEU:HG	2.19	0.42
60:t:108:ARG:HG2	60:t:116:ILE:HD13	2.01	0.42
63:w:96:ARG:O	63:w:113:ILE:HA	2.19	0.42
64:x:36:TYR:CD2	64:x:37:ALA:N	2.88	0.42
64:x:79:ALA:CB	64:x:113:ALA:HB3	2.49	0.42
1:0:52:PRO:HB3	66:z:89:GLU:CD	2.44	0.42
4:3:6:ARG:HG2	4:3:7:ARG:N	2.35	0.42
5:4:77:VAL:HG21	5:4:79:ARG:HE	1.84	0.42
6:5:25:DA:N3	7:6:5:DG:N2	2.68	0.42
8:7:7:U:H6	8:7:7:U:H3'	1.84	0.42
9:9:2:ALA:HB3	9:9:6:GLN:HB2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:9:17:GLU:OE2	9:9:53:ARG:NH2	2.53	0.42
9:9:24:SER:O	9:9:96:PHE:CE1	2.72	0.42
11:AA:144:VAL:HG23	11:AA:515:MET:HB2	2.02	0.42
11:AA:616:ILE:HG13	11:AA:652:TYR:HB2	2.02	0.42
12:AB:120:GLU:HB3	12:AB:162:LEU:HD12	2.02	0.42
12:AB:148:LYS:CD	30:P:100:ILE:HD13	2.49	0.42
14:AE:800:LEU:HB3	14:AE:920:ALA:HB1	2.02	0.42
14:AE:975:ILE:HD11	14:AE:1003:LEU:HG	2.02	0.42
16:AG:185:SER:O	16:AG:185:SER:OG	2.37	0.42
16:AG:314:LEU:HA	16:AG:314:LEU:HD23	1.85	0.42
16:AG:434:LEU:HD23	16:AG:456:LEU:CB	2.49	0.42
17:C:38:LYS:HA	17:C:38:LYS:CE	2.37	0.42
17:C:44:ILE:HG22	22:H:340:ARG:HB2	2.02	0.42
17:C:70:TYR:HE2	26:L:49:TYR:CZ	2.38	0.42
18:D:643:C:C2	18:D:644:U:C5	3.08	0.42
18:D:745:G:H2'	18:D:746:A:H8	1.84	0.42
18:D:771:G:H2'	18:D:772:U:C6	2.54	0.42
18:D:1368:A:OP1	29:O:116:VAL:HG23	2.20	0.42
18:D:1436:U:C2	18:D:1437:A:C8	3.07	0.42
18:D:1499:A:O2'	18:D:1500:A:H5'	2.20	0.42
22:H:33:LYS:HA	22:H:34:ASP:HA	1.80	0.42
22:H:52:GLN:OE1	22:H:52:GLN:HA	2.18	0.42
22:H:322:VAL:HG12	22:H:323:ASN:HB2	2.02	0.42
25:K:15:LEU:HD23	25:K:15:LEU:H	1.83	0.42
25:K:77:ASN:O	25:K:80:THR:O	2.37	0.42
27:M:66:LEU:HD13	27:M:101:MET:HE3	1.93	0.42
32:R:30:LYS:CB	32:R:57:LEU:HD11	2.47	0.42
35:U:26:ASN:ND2	35:U:31:ARG:CD	2.83	0.42
41:a:126:A:OP1	53:m:45:SER:OG	2.37	0.42
41:a:553:G:C5	41:a:554:U:C5	3.07	0.42
41:a:635:C:OP1	61:u:126:ARG:CZ	2.68	0.42
41:a:667:U:H2'	41:a:668:A:O4'	2.20	0.42
41:a:749:A:H2'	41:a:749:A:N3	2.35	0.42
41:a:958:U:C4	62:v:40:ARG:NH1	2.88	0.42
41:a:1008:A:OP2	59:s:37:ARG:NH1	2.53	0.42
41:a:1361:G:H2'	41:a:1362:C:C6	2.54	0.42
41:a:1839:G:O4'	41:a:1927:A:O4'	2.37	0.42
41:a:1874:C:H2'	41:a:1875:G:O4'	2.20	0.42
41:a:1906:G:C8	41:a:1929:G:C4	3.08	0.42
41:a:2585:U:O2	41:a:2585:U:O4'	2.37	0.42
41:a:2884:U:P	49:i:40:ARG:NH2	2.93	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:c:3:ARG:CD	43:c:30:LEU:HD11	2.50	0.42
44:d:83:G:O4'	46:f:53:PHE:CE2	2.72	0.42
48:h:108:LYS:CG	48:h:109:GLY:N	2.83	0.42
50:j:184:ARG:NH2	65:y:7:GLN:CD	2.72	0.42
51:k:9:ILE:HG22	51:k:27:LYS:HZ1	1.84	0.42
51:k:9:ILE:HD12	51:k:51:GLU:HG2	2.01	0.42
53:m:25:LYS:O	53:m:28:ARG:N	2.52	0.42
54:n:38:MET:HE3	54:n:57:LEU:CD2	2.50	0.42
60:t:118:LEU:O	60:t:119:ALA:C	2.62	0.42
62:v:72:PRO:C	62:v:73:ILE:HD12	2.43	0.42
63:w:45:ARG:CZ	63:w:45:ARG:HB3	2.50	0.42
66:z:111:GLU:OE2	66:z:112:LYS:HG2	2.19	0.42
1:0:47:VAL:C	1:0:48:LYS:HD3	2.45	0.42
9:9:41:LEU:CD1	39:Y:117:THR:HG22	2.50	0.42
9:9:47:GLU:C	9:9:49:GLY:H	2.28	0.42
12:AB:6:LEU:HB2	12:AB:56:PHE:CE1	2.55	0.42
12:AB:29:LEU:H	12:AB:56:PHE:HB2	1.84	0.42
12:AB:104:ILE:O	12:AB:106:ASP:N	2.53	0.42
12:AB:116:VAL:H	12:AB:129:ILE:CD1	2.32	0.42
16:AG:56:PHE:CD1	16:AG:56:PHE:N	2.88	0.42
16:AG:63:LEU:HD22	16:AG:92:TYR:CD1	2.48	0.42
16:AG:206:ILE:HD11	16:AG:226:ALA:HB2	2.02	0.42
16:AG:226:ALA:HB1	16:AG:228:ARG:NH1	2.34	0.42
16:AG:285:ILE:O	16:AG:288:MET:HE2	2.19	0.42
16:AG:320:ARG:HD3	16:AG:320:ARG:HA	1.83	0.42
16:AG:365:ILE:CD1	16:AG:406:LEU:HD21	2.37	0.42
16:AG:433:ASP:HB3	16:AG:456:LEU:CD1	2.47	0.42
10:B:59:A:C5	10:B:60:U:C5	3.08	0.42
17:C:44:ILE:HG21	22:H:339:ARG:C	2.45	0.42
18:D:36:C:N4	18:D:37:U:C4	2.88	0.42
18:D:76:G:C5	18:D:77:A:N7	2.88	0.42
18:D:429:U:H3	18:D:431:A:N6	2.18	0.42
18:D:860:A:H2'	18:D:861:G:O4'	2.20	0.42
18:D:960:U:O2'	18:D:1223:C:H4'	2.20	0.42
18:D:1186:G:O2'	29:O:112:GLU:OE2	2.33	0.42
20:F:5:LYS:HD3	31:Q:109:ASN:OD1	2.19	0.42
21:G:30:PHE:CE1	21:G:201:PRO:HG3	2.53	0.42
25:K:123:VAL:O	25:K:124:LEU:HD23	2.20	0.42
28:N:83:LEU:HD21	32:R:4:VAL:HG21	2.02	0.42
30:P:59:LYS:HE2	30:P:62:ARG:CZ	2.50	0.42
30:P:66:GLU:HG3	33:S:99:ALA:HB2	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:U:4:ILE:N	35:U:4:ILE:HD12	2.34	0.42
36:V:48:ASP:HB3	36:V:75:LEU:CD2	2.49	0.42
37:W:69:HIS:ND1	37:W:74:PHE:HZ	2.17	0.42
41:a:62:U:O2	41:a:62:U:H2'	2.20	0.42
41:a:1343:G:C8	41:a:1597:A:C6	3.08	0.42
41:a:1560:G:C5	41:a:1561:C:C5	3.08	0.42
41:a:1597:A:H5''	41:a:1598:A:H5'	2.02	0.42
41:a:2259:U:C5	41:a:2427:C:N4	2.88	0.42
41:a:2507:C:C2	41:a:2583:G:C2	3.07	0.42
48:h:18:LYS:HE3	48:h:20:VAL:HG21	2.02	0.42
50:j:175:LEU:HB3	50:j:189:VAL:CG1	2.50	0.42
52:l:6:LYS:HB2	52:l:6:LYS:HE2	1.75	0.42
52:l:147:LEU:HB2	52:l:183:PHE:CD2	2.48	0.42
53:m:4:THR:HG23	53:m:5:PHE:N	2.34	0.42
59:s:37:ARG:HA	59:s:118:MET:HE1	2.00	0.42
62:v:135:VAL:HG12	62:v:136:MET:CE	2.50	0.42
63:w:38:LEU:HB3	63:w:39:PRO:HD3	2.02	0.42
1:0:40:MET:HE1	66:z:105:ALA:HB1	2.01	0.41
1:0:77:PHE:HE1	1:0:79:ARG:HA	1.85	0.41
3:2:26:LYS:HE3	3:2:26:LYS:HA	2.02	0.41
9:9:31:ARG:HH11	41:a:1054:A:C4'	2.31	0.41
9:9:77:VAL:O	9:9:77:VAL:CG1	2.67	0.41
10:A:5:G:C2'	10:A:6:G:H5'	2.50	0.41
11:AA:590:PRO:HB2	11:AA:655:VAL:HG21	2.01	0.41
11:AA:856:ASN:CB	16:AG:35:THR:HB	2.48	0.41
12:AB:123:PHE:CD2	12:AB:123:PHE:N	2.79	0.41
14:AE:282:LEU:HD23	14:AE:282:LEU:HA	1.92	0.41
16:AG:131:ARG:HG2	16:AG:186:VAL:CG1	2.47	0.41
16:AG:152:LEU:HD11	16:AG:162:ILE:HD12	2.02	0.41
16:AG:207:GLU:HG3	16:AG:210:ARG:CB	2.50	0.41
16:AG:296:ILE:CD1	16:AG:307:ILE:HA	2.49	0.41
16:AG:354:ALA:O	16:AG:358:THR:HG22	2.19	0.41
10:B:5:G:C2'	10:B:6:G:C5'	2.98	0.41
17:C:12:ARG:CB	22:H:264:GLU:CB	2.98	0.41
18:D:105:G:C5	18:D:106:C:C4	3.08	0.41
18:D:122:G:C4	18:D:123:U:C6	3.08	0.41
18:D:255:G:H2'	18:D:256:U:C6	2.55	0.41
18:D:519:C:H2'	18:D:520:A:O4'	2.20	0.41
18:D:603:U:C2	18:D:604:G:C8	3.08	0.41
18:D:678:U:C2	18:D:679:C:C5	3.08	0.41
18:D:892:A:H2'	18:D:893:C:H6	1.83	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:908:A:H2'	18:D:909:A:H8	1.85	0.41
18:D:921:U:H4'	25:K:23:LYS:HZ3	1.83	0.41
18:D:1064:G:O2'	18:D:1190:G:N2	2.53	0.41
18:D:1172:C:C2	18:D:1173:U:C5	3.08	0.41
18:D:1227:A:OP2	38:X:110:LYS:HE3	2.20	0.41
18:D:1241:G:H2'	18:D:1242:G:H8	1.85	0.41
18:D:1309:G:C8	38:X:98:ARG:NH2	2.88	0.41
19:E:22:ALA:N	19:E:25:ARG:HE	2.17	0.41
19:E:53:GLU:O	19:E:57:ILE:CD1	2.68	0.41
20:F:5:LYS:CD	31:Q:109:ASN:OD1	2.68	0.41
21:G:9:MET:HE2	21:G:43:LEU:HD22	2.00	0.41
21:G:47:VAL:N	21:G:48:PRO:HD2	2.35	0.41
23:I:184:TYR:O	23:I:185:ASN:CG	2.63	0.41
24:J:65:TYR:CE1	24:J:100:ASN:CG	2.98	0.41
24:J:76:TYR:O	24:J:76:TYR:CD1	2.73	0.41
27:M:18:PHE:CD2	27:M:59:LEU:HG	2.55	0.41
27:M:37:SER:HA	29:O:41:ARG:HH22	1.85	0.41
27:M:66:LEU:HA	27:M:66:LEU:HD12	1.84	0.41
28:N:27:MET:CA	28:N:58:GLU:OE2	2.67	0.41
29:O:97:GLU:HG2	29:O:98:LEU:N	2.35	0.41
29:O:112:GLU:HG3	29:O:115:LYS:NZ	2.35	0.41
31:Q:79:ILE:O	31:Q:105:PHE:CE1	2.72	0.41
33:S:87:ALA:CA	33:S:90:ARG:CZ	2.96	0.41
38:X:3:ARG:CZ	54:n:110:ARG:HD2	2.49	0.41
38:X:16:VAL:HG23	38:X:17:ILE:CD1	2.47	0.41
40:Z:20:VAL:O	40:Z:20:VAL:CG1	2.65	0.41
41:a:16:C:O4'	49:i:15:MET:HE1	2.20	0.41
41:a:30:G:H2'	41:a:31:C:C6	2.56	0.41
41:a:104:A:C4	41:a:105:C:C6	3.08	0.41
41:a:729:G:C8	48:h:207:LYS:HD2	2.54	0.41
41:a:920:A:H2'	41:a:921:C:H6	1.85	0.41
41:a:938:G:C2	41:a:939:G:C8	3.08	0.41
41:a:1069:A:C2	41:a:1073:A:C8	3.08	0.41
41:a:1246:A:H2'	41:a:1247:A:O4'	2.20	0.41
41:a:1365:A:H4'	43:c:11:ARG:HH12	1.85	0.41
41:a:1824:G:H3'	48:h:52:ARG:HH12	1.84	0.41
41:a:1829:A:C8	41:a:1830:C:C5	3.08	0.41
41:a:1946:U:H2'	41:a:1947:C:H6	1.84	0.41
41:a:2072:C:C2	41:a:2073:C:C5	3.08	0.41
41:a:2218:G:C5'	43:c:45:ARG:HH12	2.33	0.41
41:a:2357:G:OP1	42:b:20:ARG:CZ	2.68	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:2392:A:N3	61:u:55:MET:HE2	2.34	0.41
41:a:2673:G:C2	41:a:2674:G:C8	3.08	0.41
41:a:2856:A:C6	41:a:2862:G:C6	3.08	0.41
42:b:21:LEU:HD23	42:b:40:GLN:HA	2.00	0.41
44:d:57:A:C4	44:d:58:A:C8	3.08	0.41
45:e:20:ASN:O	45:e:24:GLU:HG2	2.20	0.41
48:h:245:VAL:HA	48:h:250:VAL:O	2.20	0.41
50:j:68:PHE:CE1	50:j:79:LEU:HD21	2.54	0.41
50:j:151:THR:HG23	50:j:151:THR:O	2.19	0.41
52:l:23:PHE:CD1	52:l:111:GLU:HG3	2.55	0.41
52:l:46:GLN:O	52:l:88:ARG:NH2	2.53	0.41
52:l:149:ILE:HG21	52:l:175:ILE:HD11	2.01	0.41
54:n:141:ILE:HD11	54:n:146:VAL:CG1	2.50	0.41
54:n:142:ASP:CB	54:n:145:LYS:HE2	2.40	0.41
59:s:28:LEU:HD12	59:s:142:ILE:HG21	2.01	0.41
59:s:99:ARG:O	59:s:100:VAL:C	2.63	0.41
60:t:43:ILE:HD12	60:t:56:ASP:HB2	2.02	0.41
62:v:115:GLU:O	62:v:116:ALA:C	2.61	0.41
63:w:38:LEU:HA	63:w:38:LEU:HD12	1.72	0.41
1:0:74:ILE:HD13	1:0:87:GLN:OE1	2.20	0.41
3:2:5:GLU:HA	45:e:22:LEU:HD21	2.02	0.41
5:4:18:ARG:NH1	44:d:93:C:OP2	2.53	0.41
8:7:12:U:H4'	25:K:56:VAL:HG11	2.02	0.41
9:9:95:LEU:O	9:9:98:GLU:HG2	2.20	0.41
10:A:31:G:H2'	10:A:32:C:C6	2.55	0.41
10:A:68:C:C4	10:A:69:C:C5	3.08	0.41
12:AB:115:LYS:HD2	12:AB:129:ILE:CB	2.50	0.41
12:AB:143:LEU:HD23	12:AB:144:ASN:N	2.35	0.41
16:AG:105:ILE:HD13	16:AG:105:ILE:H	1.85	0.41
16:AG:363:LEU:HD21	16:AG:407:ARG:C	2.46	0.41
16:AG:434:LEU:HG	16:AG:455:THR:CA	2.50	0.41
16:AG:434:LEU:CD1	16:AG:455:THR:CA	2.98	0.41
17:C:71:THR:H	17:C:74:HIS:CE1	2.38	0.41
18:D:124:C:H2'	18:D:125:U:C6	2.56	0.41
18:D:997:U:O2'	18:D:998:C:H5'	2.21	0.41
18:D:1251:A:H2'	18:D:1252:A:O4'	2.20	0.41
22:H:309:MET:HE1	22:H:318:PRO:HG3	2.02	0.41
26:L:14:GLN:NE2	26:L:83:ALA:HB2	2.35	0.41
32:R:7:LEU:HD21	32:R:12:ARG:HH21	1.82	0.41
39:Y:96:LYS:CD	39:Y:136:GLY:HA3	2.50	0.41
41:a:138:U:C5	41:a:142:A:C6	3.08	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:888:C:C4	41:a:889:C:N4	2.89	0.41
41:a:975:A:H1'	41:a:990:A:C2	2.55	0.41
41:a:2072:C:C2	41:a:2073:C:C6	3.08	0.41
41:a:2461:A:H2'	41:a:2462:C:C6	2.56	0.41
41:a:2756:U:C4	41:a:2759:G:O6	2.73	0.41
45:e:14:LEU:HA	45:e:17:GLU:OE1	2.20	0.41
54:n:38:MET:HE3	54:n:57:LEU:HD23	2.01	0.41
61:u:96:LYS:HG2	61:u:102:GLY:O	2.21	0.41
64:x:15:ARG:C	64:x:19:GLN:NE2	2.78	0.41
64:x:88:LYS:HB3	64:x:116:GLN:OE1	2.20	0.41
6:5:10:DT:C2	11:AA:62:TYR:HE2	2.38	0.41
9:9:23:LEU:HA	9:9:118:ILE:HG13	2.01	0.41
9:9:33:VAL:O	9:9:36:ASP:OD1	2.39	0.41
9:9:117:LEU:O	9:9:117:LEU:CD2	2.67	0.41
10:A:9:G:O2'	10:A:45:G:H2'	2.20	0.41
10:A:19:G:O6	41:a:2112:G:P	2.79	0.41
11:AA:889:PRO:HG2	16:AG:104:ARG:NH1	2.34	0.41
12:AB:90:SER:HB2	12:AB:94:HIS:NE2	2.35	0.41
12:AB:118:ILE:CG2	12:AB:128:ALA:HB3	2.50	0.41
14:AE:836:ARG:HG3	14:AE:869:CYS:HB3	2.02	0.41
10:B:68:C:C4	10:B:69:C:C5	3.08	0.41
17:C:12:ARG:NH1	17:C:13:PHE:CD1	2.89	0.41
17:C:13:PHE:CG	17:C:51:TYR:CD1	3.08	0.41
18:D:13:U:C2	18:D:914:A:C8	3.08	0.41
18:D:562:U:O3'	32:R:12:ARG:HD2	2.20	0.41
18:D:1113:C:C2	18:D:1114:C:C5	3.08	0.41
18:D:1202:U:C2	33:S:82:ILE:HD13	2.55	0.41
18:D:1439:G:H2'	18:D:1440:U:O4'	2.20	0.41
18:D:1493:A:O2'	18:D:1494:G:P	2.78	0.41
22:H:267:TRP:CD2	22:H:340:ARG:HG3	2.56	0.41
22:H:293:PHE:HA	22:H:302:GLY:O	2.20	0.41
26:L:9:MET:SD	26:L:57:ALA:HB1	2.60	0.41
26:L:42:TRP:HB2	26:L:59:TYR:O	2.20	0.41
30:P:79:PRO:HG2	30:P:102:LEU:HD11	2.02	0.41
31:Q:64:GLN:HB2	31:Q:99:ALA:HB2	2.01	0.41
32:R:21:VAL:HG23	32:R:21:VAL:O	2.19	0.41
32:R:66:TYR:CE2	32:R:68:GLY:HA2	2.55	0.41
41:a:241:A:N1	41:a:255:A:H5''	2.36	0.41
41:a:271:G:C2	41:a:272:A:C4	3.08	0.41
41:a:592:A:N3	55:o:4:ILE:HD11	2.35	0.41
41:a:985:C:O2	41:a:986:C:C6	2.73	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:1248:G:O2'	66:z:3:ARG:HA	2.21	0.41
41:a:1580:A:C8	41:a:1581:G:C8	3.08	0.41
41:a:1953:A:H2	41:a:2549:G:N3	2.18	0.41
41:a:2049:G:H21	50:j:161:MET:CE	2.33	0.41
41:a:2193:G:HO2'	41:a:2194:U:P	2.41	0.41
41:a:2586:U:H2'	41:a:2587:A:O4'	2.20	0.41
41:a:2838:G:C6	41:a:2839:G:C5	3.08	0.41
46:f:4:THR:O	46:f:5:ILE:HD13	2.19	0.41
61:u:123:ARG:C	61:u:125:LEU:HD22	2.45	0.41
64:x:88:LYS:O	64:x:116:GLN:OE1	2.38	0.41
2:1:49:LYS:NZ	41:a:491:G:O6	2.50	0.41
4:3:85:PHE:HE1	4:3:94:ARG:HG2	1.86	0.41
6:5:16:DG:N2	11:AA:445:ILE:HG21	2.35	0.41
9:9:4:ASN:HB3	9:9:8:LYS:HZ2	1.82	0.41
11:AA:22:LEU:HD22	11:AA:603:ILE:HG21	2.02	0.41
11:AA:855:PRO:CB	16:AG:35:THR:HG1	2.22	0.41
12:AB:148:LYS:HB3	30:P:100:ILE:CG2	2.50	0.41
14:AE:319:SER:HA	14:AE:320:ASN:HA	1.73	0.41
16:AG:26:ALA:CA	16:AG:117:LYS:HG2	2.51	0.41
16:AG:87:LEU:CD2	16:AG:93:VAL:HG21	2.50	0.41
16:AG:339:MET:SD	16:AG:347:LYS:CD	2.94	0.41
10:B:38:A:H4'	18:D:790:A:C1'	2.49	0.41
18:D:407:U:O4'	24:J:116:GLN:OE1	2.39	0.41
18:D:585:G:P	36:V:39:LYS:HZ1	2.44	0.41
18:D:671:G:O2'	26:L:79:ARG:NH1	2.54	0.41
18:D:958:A:H5'	37:W:55:ARG:HH12	1.86	0.41
18:D:980:C:H2'	18:D:981:U:H5'	2.01	0.41
18:D:1071:C:C5'	25:K:54:ARG:HH21	2.33	0.41
18:D:1411:C:H2'	18:D:1412:C:C6	2.56	0.41
18:D:1524:C:H2'	18:D:1525:G:C8	2.55	0.41
19:E:66:LEU:HA	19:E:66:LEU:HD12	1.85	0.41
22:H:336:ASP:HB2	22:H:341:ARG:H	1.84	0.41
23:I:47:LEU:HD22	23:I:76:VAL:HG22	2.03	0.41
23:I:105:GLU:CD	23:I:107:ARG:HH21	2.28	0.41
25:K:82:GLN:OE1	25:K:147:MET:SD	2.78	0.41
27:M:16:PRO:HB3	29:O:46:MET:CG	2.47	0.41
27:M:108:ALA:O	27:M:119:ARG:NE	2.51	0.41
29:O:44:ALA:HA	29:O:47:VAL:HG22	2.03	0.41
31:Q:94:GLU:O	31:Q:94:GLU:OE2	2.39	0.41
32:R:3:THR:O	32:R:4:VAL:C	2.62	0.41
41:a:244:A:H2'	41:a:245:G:O4'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:319:G:H2'	41:a:320:A:O4'	2.21	0.41
41:a:1820:U:C2	48:h:201:MET:CE	3.03	0.41
41:a:1824:G:O5'	48:h:52:ARG:NH1	2.53	0.41
41:a:2244:U:O2'	41:a:2245:U:H5'	2.20	0.41
41:a:2708:G:C2	41:a:2709:G:C8	3.08	0.41
41:a:2722:G:C2'	41:a:2723:C:H5'	2.50	0.41
42:b:30:SER:HA	42:b:66:LYS:HD2	2.03	0.41
48:h:53:HIS:NE2	48:h:219:THR:HG23	2.36	0.41
48:h:124:ILE:O	48:h:124:ILE:HG22	2.21	0.41
48:h:220:VAL:CG2	48:h:225:MET:HE3	2.51	0.41
49:i:34:SER:OG	49:i:36:GLU:HG2	2.20	0.41
50:j:47:ALA:O	50:j:48:ILE:HD13	2.20	0.41
51:k:8:LYS:HA	51:k:24:THR:HA	2.02	0.41
51:k:26:ASN:OD1	51:k:28:ARG:CZ	2.69	0.41
54:n:4:LEU:HD11	54:n:101:GLU:CB	2.50	0.41
60:t:113:MET:SD	60:t:113:MET:C	3.03	0.41
60:t:116:ILE:HG13	60:t:117:SER:N	2.34	0.41
61:u:98:ALA:O	61:u:100:ILE:HG13	2.20	0.41
62:v:57:VAL:CG2	62:v:60:GLN:O	2.69	0.41
62:v:104:GLU:CD	62:v:105:MET:H	2.28	0.41
63:w:98:LEU:CD1	63:w:114:GLU:OE2	2.68	0.41
2:1:23:LEU:CD2	2:1:39:THR:HG21	2.50	0.41
3:2:22:THR:O	3:2:25:GLU:HG2	2.20	0.41
6:5:10:DT:C2'	12:AB:69:ILE:C	2.89	0.41
7:6:14:DC:H2'	7:6:15:DC:H6	1.85	0.41
16:AG:204:MET:SD	16:AG:204:MET:O	2.79	0.41
16:AG:393:LEU:CG	16:AG:398:LEU:HD22	2.22	0.41
16:AG:437:LEU:CD2	16:AG:488:ILE:CD1	2.98	0.41
10:B:5:G:C2'	10:B:6:G:H5'	2.50	0.41
10:B:9:G:O2'	10:B:45:G:H2'	2.20	0.41
17:C:30:LYS:HA	17:C:33:ILE:HG12	2.01	0.41
18:D:124:C:C2	18:D:125:U:C5	3.09	0.41
18:D:545:C:OP2	24:J:62:ARG:NH1	2.53	0.41
18:D:1061:G:H5'	30:P:61:ALA:HB2	2.03	0.41
18:D:1140:C:O2'	18:D:1141:C:O5'	2.39	0.41
24:J:76:TYR:HE1	24:J:86:THR:CG2	2.33	0.41
25:K:123:VAL:C	25:K:124:LEU:HD23	2.45	0.41
27:M:17:LYS:HZ2	27:M:18:PHE:HE1	1.56	0.41
27:M:74:GLU:HG3	27:M:91:VAL:CG1	2.50	0.41
29:O:50:GLN:CA	29:O:53:GLU:OE1	2.69	0.41
32:R:110:ARG:O	32:R:111:LYS:NZ	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Y:24:GLY:HA3	39:Y:25:PRO:HD3	1.86	0.41
39:Y:102:ARG:NH2	39:Y:139:VAL:HG11	2.36	0.41
40:Z:6:GLN:CD	40:Z:10:ALA:HB2	2.45	0.41
41:a:152:A:C6	41:a:153:U:C4	3.09	0.41
41:a:485:C:N3	41:a:486:C:C5	2.89	0.41
41:a:666:A:H1'	55:o:4:ILE:CD1	2.47	0.41
41:a:700:G:H2'	41:a:701:G:O4'	2.20	0.41
41:a:874:G:C6	41:a:904:G:C6	3.08	0.41
41:a:1026:G:C8	41:a:1134:A:C4	3.09	0.41
41:a:1281:G:H2'	41:a:1282:U:C6	2.55	0.41
41:a:1289:C:H2'	41:a:1290:C:H6	1.84	0.41
41:a:1399:C:C2	41:a:1400:U:C5	3.08	0.41
41:a:1494:A:H2'	41:a:1495:A:C8	2.56	0.41
41:a:1542:U:H2'	41:a:1543:G:O4'	2.20	0.41
41:a:1664:A:N3	60:t:67:LYS:NZ	2.59	0.41
41:a:1951:U:OP1	60:t:54:LYS:NZ	2.52	0.41
41:a:1996:C:H4'	50:j:128:ARG:NH1	2.35	0.41
41:a:2469:A:C6	41:a:2482:A:C8	3.07	0.41
41:a:2520:C:O2'	41:a:2521:C:H5'	2.20	0.41
41:a:2784:U:H2'	41:a:2785:C:C6	2.55	0.41
41:a:2808:G:N1	41:a:2891:U:C5	2.89	0.41
45:e:25:GLN:OE1	45:e:50:VAL:HG21	2.19	0.41
48:h:161:TYR:HB3	48:h:194:GLU:HG2	2.03	0.41
48:h:180:GLU:O	48:h:181:MET:HE2	2.21	0.41
50:j:68:PHE:CZ	50:j:79:LEU:HD21	2.56	0.41
52:l:97:ASN:CG	52:l:100:MET:HE2	2.46	0.41
54:n:8:TYR:O	54:n:13:VAL:HG23	2.20	0.41
56:p:52:PHE:HZ	56:p:72:LEU:HD22	1.85	0.41
56:p:91:GLY:HA2	56:p:160:LYS:CD	2.51	0.41
60:t:1:MET:HE3	60:t:32:TYR:CB	2.50	0.41
62:v:45:GLN:HE22	62:v:125:PRO:HD2	1.84	0.41
62:v:105:MET:HG2	62:v:117:PHE:HZ	1.85	0.41
63:w:24:MET:CG	63:w:44:LEU:HD12	2.51	0.41
63:w:33:ILE:CD1	63:w:112:TYR:HD1	2.32	0.41
1:0:4:VAL:HG23	1:0:4:VAL:O	2.20	0.41
3:2:26:LYS:HA	3:2:26:LYS:CE	2.50	0.41
9:9:96:PHE:HB3	9:9:125:ARG:HH11	1.86	0.41
9:9:118:ILE:CB	9:9:119:PRO:HD3	2.44	0.41
10:A:59:A:C5	10:A:60:U:C5	3.08	0.41
11:AA:310:ILE:HG21	11:AA:325:LEU:HB3	2.02	0.41
12:AB:6:LEU:HB2	12:AB:56:PHE:HE1	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AE:134:ASP:HB3	14:AE:159:ILE:HG21	2.02	0.41
14:AE:641:ILE:HD12	14:AE:641:ILE:HA	1.89	0.41
16:AG:284:VAL:HG22	16:AG:305:MET:CE	2.50	0.41
16:AG:296:ILE:HD11	16:AG:307:ILE:CD1	2.50	0.41
17:C:60:LYS:NZ	18:D:735:C:O5'	2.52	0.41
18:D:21:G:N2	18:D:22:G:C6	2.88	0.41
18:D:511:C:O3'	24:J:44:ARG:NH1	2.53	0.41
18:D:680:C:H2'	18:D:681:A:H8	1.85	0.41
18:D:687:A:C8	18:D:701:U:C4	3.08	0.41
18:D:820:U:H4'	18:D:821:G:OP2	2.20	0.41
18:D:1174:G:H2'	18:D:1175:G:H5'	2.02	0.41
18:D:1411:C:H2'	18:D:1412:C:H6	1.85	0.41
21:G:187:VAL:HG11	21:G:193:PRO:HB3	2.02	0.41
22:H:70:VAL:O	22:H:71:ALA:HB2	2.20	0.41
23:I:6:HIS:CD2	33:S:89:MET:HB3	2.55	0.41
23:I:123:GLN:HB3	23:I:128:VAL:HG21	2.02	0.41
23:I:147:LYS:CE	23:I:204:LYS:C	2.94	0.41
25:K:147:MET:HE3	25:K:147:MET:O	2.21	0.41
26:L:3:HIS:CD2	26:L:95:ALA:HA	2.56	0.41
26:L:51:ILE:CG2	26:L:86:ARG:HH21	2.34	0.41
29:O:35:LEU:HD21	29:O:45:ARG:HG2	2.03	0.41
29:O:50:GLN:HA	29:O:53:GLU:OE1	2.20	0.41
31:Q:20:VAL:HG12	31:Q:21:ALA:N	2.35	0.41
34:T:18:ASP:O	34:T:19:ALA:HB3	2.20	0.41
39:Y:57:VAL:CG2	39:Y:71:LYS:HD3	2.50	0.41
41:a:66:C:H42	41:a:74:A:N6	2.19	0.41
41:a:690:G:H2'	41:a:691:C:H6	1.85	0.41
41:a:832:U:H2'	41:a:833:A:C8	2.50	0.41
41:a:1322:A:C4	41:a:1323:C:C6	3.08	0.41
41:a:2024:G:C5	41:a:2040:G:C2	3.09	0.41
41:a:2040:G:H2'	41:a:2041:U:O4'	2.20	0.41
41:a:2291:U:H2'	41:a:2292:U:H6	1.85	0.41
41:a:2454:G:C5	41:a:2455:G:C8	3.09	0.41
41:a:2539:C:O2'	57:q:36:ARG:NH2	2.54	0.41
41:a:2686:G:H2'	41:a:2687:U:C6	2.56	0.41
48:h:196:GLY:O	48:h:198:ALA:N	2.53	0.41
51:k:26:ASN:OD1	51:k:28:ARG:NH1	2.54	0.41
55:o:22:PHE:CE1	55:o:62:LEU:HD23	2.55	0.41
62:v:112:LEU:CA	62:v:115:GLU:OE1	2.68	0.41
64:x:21:LEU:HD23	64:x:21:LEU:C	2.46	0.41
64:x:86:GLY:HA2	64:x:88:LYS:HZ3	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2:50:LEU:HD23	45:e:26:PHE:CE1	2.56	0.41
8:7:50:U:H4'	14:AE:322:ARG:NH1	2.36	0.41
9:9:23:LEU:CA	9:9:118:ILE:HD11	2.50	0.41
11:AA:849:GLU:HB2	11:AA:887:VAL:HG23	2.02	0.41
12:AB:120:GLU:C	12:AB:162:LEU:HD12	2.46	0.41
12:AB:155:LYS:HB3	12:AB:158:GLU:CD	2.45	0.41
13:AD:205:MET:HE1	13:AD:217:ILE:HG13	2.03	0.41
14:AE:138:VAL:HG21	14:AE:145:VAL:HB	2.03	0.41
14:AE:1272:SER:HB2	14:AE:1292:LEU:HD11	2.02	0.41
16:AG:323:GLN:HE21	16:AG:323:GLN:C	2.28	0.41
10:B:9:G:H5''	10:B:11:A:OP2	2.21	0.41
18:D:71:A:C6	18:D:100:G:N7	2.89	0.41
18:D:104:G:C2	18:D:105:G:C5	3.09	0.41
18:D:107:G:C6	18:D:108:G:C4	3.09	0.41
18:D:296:U:O2'	18:D:556:C:O2	2.34	0.41
18:D:375:U:H1'	35:U:28:ARG:NH2	2.36	0.41
18:D:567:G:H2'	18:D:568:G:O4'	2.21	0.41
18:D:602:A:H2'	18:D:603:U:H6	1.86	0.41
18:D:982:U:H4'	18:D:983:A:O4'	2.21	0.41
18:D:1240:U:H5	27:M:116:MET:HE2	1.83	0.41
23:I:66:VAL:HB	23:I:101:ILE:CD1	2.50	0.41
23:I:135:LYS:HG3	23:I:136:ARG:N	2.35	0.41
25:K:80:THR:OG1	25:K:81:LEU:N	2.53	0.41
30:P:73:LEU:HD23	30:P:73:LEU:O	2.21	0.41
31:Q:75:LYS:HG3	31:Q:76:GLU:N	2.36	0.41
32:R:25:GLU:CD	32:R:30:LYS:HE3	2.46	0.41
38:X:13:LYS:CE	38:X:17:ILE:HG21	2.51	0.41
39:Y:34:ILE:HG22	39:Y:34:ILE:O	2.20	0.41
40:Z:14:MET:CG	40:Z:15:SER:H	2.25	0.41
41:a:1252:G:H1	66:z:37:GLN:HE21	1.68	0.41
41:a:1589:U:O2'	41:a:1590:A:P	2.79	0.41
41:a:1651:G:P	63:w:40:LYS:NZ	2.94	0.41
41:a:1822:C:O2'	41:a:1823:G:H5'	2.21	0.41
41:a:2065:C:C2	41:a:2066:C:C6	3.09	0.41
41:a:2210:U:H4'	41:a:2211:A:OP1	2.20	0.41
41:a:2393:U:H2'	41:a:2394:C:C6	2.56	0.41
41:a:2483:C:H1'	62:v:51:ARG:CZ	2.51	0.41
41:a:2590:A:C2	41:a:2591:C:C5	3.08	0.41
41:a:2649:C:C2	41:a:2650:U:C6	3.09	0.41
41:a:2783:U:H2'	41:a:2784:U:H6	1.85	0.41
41:a:2848:G:HO2'	41:a:2867:G:H22	1.63	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:b:17:GLU:OE1	42:b:19:LYS:HD3	2.21	0.41
42:b:70:GLU:HG3	42:b:72:LYS:HG2	2.03	0.41
43:c:66:THR:O	43:c:69:ALA:N	2.53	0.41
44:d:57:A:N7	54:n:26:MET:HE3	2.36	0.41
50:j:34:VAL:HG13	50:j:48:ILE:CG2	2.50	0.41
52:l:170:ARG:HH22	52:l:176:ASP:H	1.68	0.41
52:l:171:ASP:OD1	52:l:172:ALA:N	2.53	0.41
53:m:25:LYS:O	53:m:26:ASN:C	2.64	0.41
53:m:44:VAL:O	53:m:45:SER:C	2.63	0.41
56:p:85:LYS:HD2	56:p:85:LYS:HA	1.77	0.41
56:p:149:ARG:CZ	56:p:154:PRO:CD	2.98	0.41
58:r:54:LEU:HD12	58:r:57:LYS:HE2	2.03	0.41
58:r:66:ASN:CG	58:r:135:HIS:HB2	2.45	0.41
60:t:24:VAL:O	60:t:24:VAL:HG13	2.20	0.41
60:t:61:VAL:HG11	60:t:112:PHE:CE2	2.56	0.41
60:t:102:PRO:HD3	65:y:66:ASN:HB2	2.03	0.41
63:w:28:LEU:HD23	63:w:48:VAL:HG21	2.03	0.41
64:x:51:ALA:HB3	64:x:78:VAL:HB	2.02	0.41
65:y:110:ILE:O	65:y:110:ILE:HG13	2.19	0.41
3:2:6:ARG:NH2	3:2:10:VAL:CG2	2.84	0.41
8:7:1:A:H61	10:B:37:A:N6	2.19	0.41
9:9:33:VAL:H	9:9:36:ASP:CG	2.28	0.41
9:9:129:LEU:CD1	9:9:130:PRO:HD3	2.45	0.41
10:A:33:U:H5''	27:M:84:THR:CG2	2.50	0.41
17:C:70:TYR:CD2	18:D:674:G:H4'	2.55	0.41
18:D:5:U:O2	18:D:5:U:O4'	2.37	0.41
18:D:545:C:C5'	24:J:69:GLU:HG3	2.38	0.41
18:D:678:U:H2'	18:D:679:C:C6	2.53	0.41
18:D:1512:U:H2'	18:D:1513:A:H8	1.85	0.41
21:G:96:TRP:CH2	21:G:100:MET:CE	3.04	0.41
23:I:8:ASN:O	23:I:12:LEU:CD2	2.69	0.41
23:I:156:ARG:HD3	23:I:160:ALA:O	2.21	0.41
24:J:76:TYR:CD1	24:J:76:TYR:C	2.98	0.41
26:L:24:ARG:HA	26:L:24:ARG:NE	2.36	0.41
29:O:5:GLN:HE21	29:O:20:PHE:HB3	1.81	0.41
30:P:6:ILE:CG2	30:P:100:ILE:HG13	2.51	0.41
30:P:56:HIS:ND1	30:P:57:VAL:HG23	2.36	0.41
31:Q:20:VAL:HG13	31:Q:85:MET:CE	2.51	0.41
36:V:11:ARG:HD3	36:V:57:ASP:C	2.46	0.41
41:a:523:C:O2	41:a:554:U:O2'	2.37	0.41
41:a:529:A:C4	41:a:2023:C:C5	3.09	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:945:A:C4	41:a:2448:A:C2	3.09	0.41
41:a:1048:A:C5	41:a:1049:C:C5	3.09	0.41
41:a:1223:G:N1	41:a:1226:A:OP2	2.50	0.41
41:a:1746:A:H2'	41:a:1747:U:C6	2.56	0.41
44:d:47:C:C4	44:d:48:U:C5	3.09	0.41
48:h:163:GLN:HB3	48:h:175:ARG:HB3	2.02	0.41
48:h:165:VAL:HG12	48:h:175:ARG:NH2	2.35	0.41
50:j:146:ILE:HA	50:j:159:LYS:NZ	2.36	0.41
51:k:21:TYR:CZ	51:k:38:LYS:CG	3.04	0.41
52:l:193:VAL:O	52:l:194:LYS:C	2.62	0.41
55:o:45:ARG:HB3	55:o:46:PRO:HD3	2.02	0.41
56:p:52:PHE:CE1	56:p:69:ARG:HD3	2.55	0.41
56:p:94:TYR:HE1	56:p:152:ARG:CD	2.34	0.41
63:w:60:VAL:HG22	63:w:63:ARG:HH11	1.80	0.41
63:w:78:LYS:HA	63:w:82:GLU:OE1	2.21	0.41
2:1:93:ALA:HB2	41:a:1614:A:C2	2.56	0.41
3:2:30:ILE:HG22	3:2:32:LEU:HD22	2.03	0.41
3:2:57:VAL:HG22	3:2:58:VAL:N	2.35	0.41
5:4:77:VAL:HG21	5:4:79:ARG:HH21	1.86	0.41
6:5:23:DA:H2''	6:5:24:DC:C6	2.56	0.41
8:7:13:U:C6	8:7:13:U:C3'	3.03	0.41
9:9:129:LEU:N	9:9:130:PRO:CD	2.81	0.41
10:A:9:G:H5''	10:A:11:A:OP2	2.21	0.41
10:A:12:G:H2'	10:A:13:C:OP1	2.20	0.41
11:AA:135:THR:HG22	11:AA:144:VAL:HG22	2.02	0.41
11:AA:836:LEU:HD21	11:AA:921:PRO:HD3	2.02	0.41
11:AA:936:ARG:HB2	11:AA:1042:LEU:HD12	2.03	0.41
11:AA:979:LEU:HD11	11:AA:1011:LEU:HD11	2.03	0.41
12:AB:2:GLN:HA	12:AB:59:PHE:O	2.21	0.41
12:AB:5:TYR:HA	12:AB:88:VAL:CG2	2.49	0.41
12:AB:72:THR:HG22	12:AB:73:ARG:H	1.86	0.41
12:AB:151:LYS:HB2	12:AB:151:LYS:HE2	1.49	0.41
14:AE:288:PRO:CG	14:AE:291:ILE:HG21	2.50	0.41
14:AE:1022:PRO:O	14:AE:1126:GLN:NE2	2.45	0.41
16:AG:205:LEU:HD12	16:AG:205:LEU:H	1.86	0.41
16:AG:258:ARG:HH11	16:AG:258:ARG:CG	2.13	0.41
17:C:21:ILE:HD12	17:C:54:GLN:HG3	2.03	0.41
18:D:338:A:C5	18:D:339:C:C5	3.08	0.41
18:D:437:U:H5''	24:J:152:GLN:CD	2.46	0.41
18:D:570:G:H1'	18:D:820:U:C4	2.56	0.41
18:D:588:G:C4'	28:N:3:MET:HE1	2.48	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:680:C:C2	18:D:681:A:C8	3.09	0.41
18:D:915:A:H8	18:D:915:A:O5'	2.04	0.41
18:D:920:U:H2'	18:D:921:U:C6	2.56	0.41
18:D:963:G:C2	18:D:964:A:C8	3.09	0.41
18:D:985:C:C2	18:D:986:U:C5	3.08	0.41
18:D:1119:C:C2	18:D:1120:C:C5	3.09	0.41
18:D:1189:U:H4'	23:I:10:ILE:CD1	2.50	0.41
18:D:1217:C:P	33:S:9:ARG:HE	2.41	0.41
18:D:1227:A:C8	38:X:116:ILE:HD11	2.56	0.41
18:D:1228:C:OP1	38:X:107:ARG:CZ	2.69	0.41
20:F:39:GLU:OE1	20:F:44:GLU:OE1	2.38	0.41
22:H:86:ARG:O	22:H:89:ALA:HB3	2.20	0.41
22:H:277:GLY:C	22:H:331:MET:HE3	2.45	0.41
22:H:308:GLU:HB2	22:H:347:LYS:HB3	2.02	0.41
23:I:36:ASP:OD1	23:I:57:ILE:HG21	2.20	0.41
24:J:62:ARG:CG	24:J:72:PHE:CZ	3.02	0.41
24:J:197:GLU:HA	24:J:200:ILE:HD12	2.03	0.41
25:K:58:ALA:O	25:K:62:LYS:HG2	2.21	0.41
25:K:115:LEU:O	25:K:120:VAL:HG12	2.20	0.41
26:L:24:ARG:O	26:L:25:TYR:C	2.62	0.41
26:L:42:TRP:HB3	26:L:59:TYR:HB2	2.03	0.41
27:M:31:MET:HE3	27:M:31:MET:HB2	1.76	0.41
27:M:138:ARG:O	27:M:142:HIS:ND1	2.46	0.41
28:N:28:PRO:HD3	28:N:58:GLU:OE1	2.21	0.41
29:O:21:ILE:HD12	29:O:86:ALA:CB	2.50	0.41
32:R:4:VAL:CG1	32:R:5:ASN:N	2.83	0.41
32:R:25:GLU:HG2	32:R:25:GLU:O	2.20	0.41
32:R:57:LEU:HD23	32:R:59:ASN:OD1	2.20	0.41
34:T:32:LEU:HD13	34:T:62:GLN:OE1	2.21	0.41
34:T:64:ARG:CZ	34:T:89:ARG:HH12	2.33	0.41
34:T:70:LEU:HD23	34:T:78:TYR:HD1	1.85	0.41
36:V:7:THR:C	36:V:8:LEU:HD12	2.46	0.41
37:W:45:ILE:HA	37:W:62:VAL:HG12	2.01	0.41
38:X:3:ARG:NH2	47:g:35:ASP:HB2	2.36	0.41
38:X:55:THR:CG2	38:X:59:GLU:OE2	2.69	0.41
38:X:71:ARG:NH1	38:X:72:GLU:CG	2.84	0.41
39:Y:42:ASN:OD1	39:Y:50:LYS:NZ	2.46	0.41
39:Y:63:ASP:C	39:Y:65:SER:H	2.28	0.41
39:Y:79:LEU:HD11	39:Y:132:ALA:HA	2.02	0.41
39:Y:82:ALA:CB	39:Y:108:ILE:HD12	2.50	0.41
39:Y:100:ILE:HG13	39:Y:139:VAL:CB	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:184:C:C2	41:a:213:A:N1	2.89	0.41
41:a:209:C:C2	41:a:210:C:C6	3.09	0.41
41:a:272:A:C2	41:a:273:G:C8	3.09	0.41
41:a:397:U:H2'	41:a:398:C:C6	2.56	0.41
41:a:465:G:H2'	41:a:466:A:C8	2.56	0.41
41:a:829:A:N7	41:a:2247:A:O2'	2.52	0.41
41:a:967:U:H2'	41:a:968:C:C6	2.56	0.41
41:a:1057:A:C2	41:a:1082:U:N3	2.89	0.41
41:a:1355:G:C2	41:a:1356:G:C8	3.09	0.41
41:a:1398:C:H2'	41:a:1399:C:H6	1.86	0.41
41:a:1427:A:H4'	41:a:1428:C:O4'	2.21	0.41
41:a:1565:C:O2'	41:a:1567:G:N7	2.47	0.41
41:a:1607:C:H4'	41:a:1608:A:O5'	2.21	0.41
41:a:1827:U:H2'	41:a:1828:G:O4'	2.20	0.41
41:a:1945:G:H2'	41:a:1946:U:C6	2.56	0.41
41:a:2366:A:H2'	41:a:2367:G:O4'	2.21	0.41
41:a:2526:G:O2'	57:q:1:MET:HG2	2.21	0.41
41:a:2569:G:C2	41:a:2570:G:C8	3.09	0.41
44:d:8:C:O2'	64:x:40:ILE:HD13	2.21	0.41
44:d:55:U:H2'	44:d:56:G:C8	2.56	0.41
47:g:7:PRO:CG	54:n:62:GLY:O	2.69	0.41
47:g:57:VAL:O	47:g:58:ASP:C	2.62	0.41
48:h:35:GLU:CD	48:h:64:ILE:HD11	2.46	0.41
48:h:189:ARG:HG3	48:h:190:ALA:N	2.36	0.41
49:i:14:GLY:HA2	49:i:17:ARG:HG2	2.02	0.41
50:j:5:VAL:HB	50:j:32:ASN:HD21	1.86	0.41
50:j:11:MET:SD	50:j:12:THR:N	2.94	0.41
50:j:37:VAL:HG22	50:j:48:ILE:HD12	2.02	0.41
50:j:55:LYS:HD3	50:j:55:LYS:C	2.45	0.41
51:k:7:GLU:CG	51:k:27:LYS:NZ	2.84	0.41
52:l:148:ILE:O	52:l:169:VAL:HA	2.21	0.41
52:l:194:LYS:O	52:l:195:GLN:C	2.64	0.41
52:l:194:LYS:C	52:l:198:GLU:OE1	2.63	0.41
54:n:26:MET:HE3	54:n:26:MET:HB2	1.92	0.41
56:p:17:VAL:O	56:p:18:LYS:CE	2.65	0.41
56:p:89:LEU:HD22	56:p:162:VAL:HA	2.02	0.41
56:p:121:ILE:CD1	56:p:141:ILE:HG22	2.43	0.41
56:p:149:ARG:HH12	56:p:154:PRO:HB3	1.85	0.41
58:r:110:VAL:HG21	58:r:114:GLU:OE1	2.21	0.41
59:s:123:LYS:CD	59:s:132:HIS:HD2	2.33	0.41
60:t:103:VAL:HG21	60:t:107:LEU:HD22	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:w:18:GLN:CD	63:w:18:GLN:C	2.88	0.41
64:x:48:LEU:CD2	64:x:87:ILE:CD1	2.98	0.41
65:y:48:ILE:HD11	65:y:101:ARG:NH1	2.36	0.41
1:0:13:ARG:C	1:0:14:VAL:HG13	2.46	0.41
3:2:9:LYS:O	3:2:12:ARG:NH2	2.52	0.41
4:3:6:ARG:NH1	41:a:84:A:C2'	2.84	0.41
9:9:38:MET:SD	9:9:41:LEU:HD12	2.61	0.41
10:A:25:C:C4	10:A:26:G:C8	3.09	0.41
11:AA:657:THR:HG21	11:AA:1188:ASP:HB2	2.03	0.41
13:AC:42:ALA:HB1	13:AC:224:LEU:HD11	2.03	0.41
13:AD:305:ASP:C	13:AD:307:LEU:H	2.28	0.41
14:AE:1155:ILE:HG13	14:AE:1210:ILE:HB	2.02	0.41
16:AG:43:ILE:HB	16:AG:44:ASP:H	1.54	0.41
16:AG:323:GLN:NE2	16:AG:323:GLN:C	2.79	0.41
16:AG:442:ARG:O	16:AG:446:PHE:CE2	2.63	0.41
18:D:105:G:H2'	18:D:106:C:C6	2.56	0.41
18:D:218:U:H2'	18:D:219:U:O4'	2.20	0.41
18:D:338:A:H2'	18:D:339:C:H6	1.86	0.41
18:D:517:G:O2'	18:D:530:G:H4'	2.21	0.41
18:D:1080:A:OP1	25:K:50:TYR:OH	2.29	0.41
18:D:1186:G:H4'	29:O:112:GLU:OE2	2.21	0.41
20:F:66:ARG:NH2	20:F:67:ARG:HH22	2.19	0.41
21:G:20:THR:O	21:G:23:TRP:CD1	2.74	0.41
21:G:56:GLU:HG2	21:G:198:PHE:CZ	2.55	0.41
21:G:142:GLU:OE2	21:G:142:GLU:HA	2.21	0.41
21:G:185:ALA:O	21:G:200:ILE:N	2.54	0.41
23:I:6:HIS:CD2	23:I:9:GLY:H	2.39	0.41
23:I:113:ALA:HB1	23:I:200:VAL:HG13	2.02	0.41
24:J:188:ARG:NH2	24:J:195:ILE:HG13	2.36	0.41
25:K:46:VAL:HG12	25:K:47:GLY:N	2.36	0.41
26:L:35:LYS:CG	26:L:36:ILE:N	2.84	0.41
29:O:56:ASP:O	29:O:60:LYS:CE	2.69	0.41
33:S:86:GLU:CD	33:S:90:ARG:HD3	2.45	0.41
35:U:4:ILE:HG22	35:U:71:VAL:HG11	2.03	0.41
37:W:34:TRP:O	37:W:36:ARG:N	2.54	0.41
39:Y:9:LYS:O	39:Y:10:LEU:HG	2.21	0.41
39:Y:60:VAL:CG2	39:Y:66:PHE:CD1	3.03	0.41
41:a:181:A:H2'	41:a:182:A:C8	2.56	0.41
41:a:205:G:O2'	41:a:206:U:OP2	2.39	0.41
41:a:320:A:H2'	52:l:131:THR:HG21	2.03	0.41
41:a:409:G:H2'	41:a:410:G:C8	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:a:451:U:C5'	52:l:47:LYS:HZ3	2.34	0.41
41:a:910:A:N1	41:a:2277:G:H1'	2.36	0.41
41:a:1128:G:C6	41:a:2518:A:C6	3.08	0.41
41:a:1172:C:C2'	41:a:1173:U:O4'	2.68	0.41
41:a:1430:G:H2'	41:a:1431:A:C8	2.56	0.41
41:a:1589:U:C2	41:a:1590:A:C8	3.09	0.41
41:a:1676:A:H2'	41:a:1677:A:C8	2.56	0.41
41:a:1792:G:C5'	48:h:204:VAL:CG2	2.99	0.41
41:a:1881:C:C4	41:a:1882:U:C5	3.09	0.41
41:a:1957:C:C2	41:a:1958:C:C5	3.09	0.41
41:a:2531:A:C4	41:a:2532:G:C8	3.08	0.41
41:a:2619:C:O2'	41:a:2620:C:H5'	2.21	0.41
42:b:39:ARG:HA	42:b:39:ARG:HD2	1.92	0.41
44:d:8:C:O3'	64:x:25:ARG:NH1	2.48	0.41
45:e:13:GLU:O	45:e:14:LEU:C	2.63	0.41
54:n:61:SER:O	54:n:63:GLN:N	2.54	0.41
54:n:94:GLU:CA	54:n:97:TRP:HE3	2.30	0.41
54:n:136:ILE:HA	54:n:141:ILE:HG21	2.01	0.41
59:s:53:TYR:HD1	59:s:121:LYS:HB3	1.86	0.41
60:t:80:ASP:OD1	65:y:68:GLU:OE2	2.39	0.41
60:t:98:ARG:NH2	60:t:118:LEU:C	2.79	0.41
61:u:101:ILE:HG13	61:u:102:GLY:N	2.36	0.41
62:v:112:LEU:HD12	62:v:112:LEU:H	1.86	0.41
63:w:49:GLU:OE2	63:w:95:THR:CG2	2.69	0.41
2:1:10:ALA:CB	2:1:103:ILE:CD1	2.99	0.40
3:2:19:LYS:HE2	3:2:84:TYR:OH	2.21	0.40
5:4:50:MET:O	5:4:56:PHE:CZ	2.74	0.40
8:7:4:U:C6	18:D:1403:C:H1'	2.57	0.40
9:9:11:ILE:O	9:9:15:VAL:HG23	2.20	0.40
10:A:1:C:C5	10:A:2:G:N7	2.89	0.40
11:AA:469:VAL:HA	11:AA:472:GLU:HG2	2.01	0.40
11:AA:856:ASN:CG	16:AG:35:THR:HB	2.43	0.40
11:AA:1105:SER:HB2	14:AE:731:ARG:HB3	2.03	0.40
12:AB:32:MET:O	12:AB:50:LEU:HD12	2.21	0.40
12:AB:36:GLU:HA	12:AB:44:THR:O	2.22	0.40
12:AB:72:THR:HG22	12:AB:73:ARG:N	2.36	0.40
13:AD:257:VAL:N	13:AD:276:HIS:O	2.34	0.40
14:AE:79:LYS:N	16:AG:144:LYS:HZ1	2.16	0.40
14:AE:1003:LEU:HD23	14:AE:1018:ALA:HB2	2.02	0.40
16:AG:125:MET:O	16:AG:125:MET:SD	2.79	0.40
10:B:1:C:C5	10:B:2:G:N7	2.89	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:356:A:C5	18:D:357:G:C8	3.08	0.40
18:D:414:A:C2	18:D:415:A:N9	2.89	0.40
18:D:718:A:C2	31:Q:118:HIS:CE1	3.09	0.40
18:D:1141:C:O2'	18:D:1142:G:P	2.79	0.40
19:E:9:LYS:CG	19:E:13:GLN:HE22	2.34	0.40
20:F:12:PHE:O	20:F:16:LEU:HG	2.21	0.40
21:G:6:MET:HA	21:G:9:MET:SD	2.61	0.40
21:G:36:ASN:ND2	22:H:19:ARG:HD2	2.36	0.40
21:G:58:ASN:HB2	21:G:220:THR:OG1	2.21	0.40
22:H:308:GLU:HG2	22:H:309:MET:N	2.31	0.40
23:I:8:ASN:O	23:I:12:LEU:HD23	2.21	0.40
23:I:172:ARG:NH1	23:I:174:PRO:HG3	2.35	0.40
24:J:74:ASN:OD1	24:J:74:ASN:C	2.64	0.40
27:M:35:LYS:CE	27:M:38:THR:OG1	2.69	0.40
28:N:8:ALA:N	28:N:77:ARG:NH1	2.69	0.40
30:P:47:GLU:N	30:P:47:GLU:CD	2.79	0.40
31:Q:20:VAL:CG1	31:Q:85:MET:CE	2.99	0.40
31:Q:36:ASP:OD1	31:Q:40:ASN:N	2.53	0.40
33:S:37:SER:O	33:S:41:ARG:HG3	2.22	0.40
33:S:63:ARG:HH22	33:S:70:PRO:HD3	1.85	0.40
34:T:26:GLU:CG	34:T:27:VAL:N	2.83	0.40
40:Z:17:MET:HE3	40:Z:17:MET:HB2	1.90	0.40
41:a:67:U:C4	41:a:74:A:N6	2.89	0.40
41:a:151:C:C2	41:a:152:A:N7	2.89	0.40
41:a:443:A:N1	52:l:40:ARG:NH2	2.69	0.40
41:a:593:U:H2'	41:a:594:U:H6	1.85	0.40
41:a:708:G:H2'	41:a:709:U:H6	1.86	0.40
41:a:841:G:H2'	41:a:842:U:H6	1.86	0.40
41:a:1177:G:O2'	41:a:1178:C:O4'	2.39	0.40
41:a:1858:A:C2	41:a:1885:A:C1'	3.04	0.40
41:a:1923:U:OP2	41:a:1923:U:H2'	2.21	0.40
41:a:1946:U:H2'	41:a:1947:C:C6	2.56	0.40
41:a:2310:C:O2'	54:n:74:VAL:HG12	2.20	0.40
41:a:2397:G:H2'	41:a:2398:U:H6	1.86	0.40
41:a:2422:C:C4	41:a:2424:C:C4	3.09	0.40
41:a:2839:G:C4	41:a:2840:C:C6	3.09	0.40
41:a:2848:G:N2	41:a:2867:G:C4	2.89	0.40
44:d:113:C:O2'	64:x:46:GLU:CD	2.64	0.40
48:h:144:VAL:HG21	48:h:162:VAL:HG11	2.01	0.40
53:m:42:LEU:N	53:m:42:LEU:HD12	2.36	0.40
58:r:8:LYS:HD2	58:r:13:GLY:O	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:t:20:MET:CB	60:t:44:LYS:HE2	2.51	0.40
62:v:31:PHE:CZ	62:v:110:GLU:HG3	2.56	0.40
63:w:12:ARG:CZ	63:w:20:MET:HE1	2.51	0.40
63:w:87:PHE:O	63:w:88:ALA:C	2.64	0.40
65:y:6:LYS:C	65:y:6:LYS:HD2	2.46	0.40
1:0:13:ARG:NH2	66:z:97:ASP:CG	2.79	0.40
4:3:86:ARG:CZ	4:3:95:PHE:CD2	3.04	0.40
4:3:86:ARG:NH2	4:3:95:PHE:CB	2.84	0.40
5:4:48:MET:CE	5:4:85:LYS:HA	2.51	0.40
6:5:10:DT:H2''	12:AB:69:ILE:CA	2.50	0.40
8:7:47:U:C3'	8:7:47:U:C6	3.04	0.40
9:9:94:ARG:HB3	9:9:97:LYS:NZ	2.32	0.40
9:9:94:ARG:HA	9:9:97:LYS:HG2	2.03	0.40
11:AA:998:LEU:HD21	11:AA:1015:ALA:HB2	2.02	0.40
12:AB:2:GLN:N	12:AB:60:ASP:OD1	2.54	0.40
12:AB:8:TYR:C	12:AB:76:SER:HB3	2.47	0.40
13:AD:76:GLU:HB3	13:AD:80:GLU:HB3	2.01	0.40
14:AE:79:LYS:HB3	14:AE:79:LYS:NZ	2.36	0.40
14:AE:438:GLU:HG3	14:AE:485:MET:HE1	2.03	0.40
14:AE:706:VAL:HG12	14:AE:715:LYS:HB3	2.04	0.40
14:AE:978:ARG:HB2	14:AE:999:TYR:HB2	2.04	0.40
16:AG:21:GLU:HB3	16:AG:49:ILE:CD1	2.49	0.40
16:AG:216:ILE:HG12	16:AG:221:ILE:HG13	2.03	0.40
16:AG:374:VAL:O	16:AG:374:VAL:CG1	2.68	0.40
10:B:16:C:O2	10:B:16:C:H2'	2.21	0.40
18:D:19:A:C6	18:D:20:U:C4	3.10	0.40
18:D:112:G:N1	18:D:330:C:C5	2.88	0.40
18:D:174:A:C5	18:D:175:C:C5	3.10	0.40
18:D:262:A:C4	18:D:263:A:C8	3.10	0.40
18:D:436:C:N3	18:D:437:U:C5	2.89	0.40
18:D:695:A:O5'	31:Q:53:ARG:CZ	2.69	0.40
18:D:1061:G:C6	18:D:1197:A:C6	3.09	0.40
21:G:17:GLY:C	22:H:43:LYS:HD2	2.46	0.40
22:H:170:ARG:O	22:H:171:ARG:CB	2.69	0.40
24:J:196:ASN:OD1	24:J:198:HIS:ND1	2.54	0.40
26:L:4:TYR:CD2	26:L:91:ARG:HD2	2.57	0.40
26:L:9:MET:CE	26:L:86:ARG:HG2	2.52	0.40
26:L:38:ARG:CZ	26:L:99:ALA:HB2	2.51	0.40
28:N:83:LEU:HD11	32:R:4:VAL:CB	2.51	0.40
28:N:83:LEU:HD23	36:V:36:LYS:CD	2.52	0.40
29:O:118:LEU:HB3	29:O:120:LYS:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Q:49:GLY:C	31:Q:51:GLY:H	2.29	0.40
32:R:3:THR:O	32:R:6:GLN:HB2	2.20	0.40
34:T:21:ASP:C	34:T:22:THR:HG1	2.16	0.40
34:T:67:LEU:CD1	34:T:78:TYR:CE1	3.04	0.40
36:V:61:ILE:HA	36:V:75:LEU:HA	2.02	0.40
39:Y:81:LYS:N	39:Y:81:LYS:CD	2.79	0.40
40:Z:6:GLN:HG3	40:Z:6:GLN:O	2.21	0.40
41:a:151:C:H2'	41:a:152:A:C8	2.53	0.40
41:a:458:G:N2	41:a:469:G:H2'	2.36	0.40
41:a:638:G:H2'	41:a:639:U:C6	2.56	0.40
41:a:966:G:C1'	41:a:2267:A:H62	2.35	0.40
41:a:1289:C:H2'	41:a:1290:C:C6	2.56	0.40
41:a:1843:C:H2'	41:a:1844:C:H6	1.85	0.40
41:a:1958:C:O2'	41:a:1959:G:H5'	2.20	0.40
41:a:1964:G:H4'	41:a:1965:C:OP2	2.19	0.40
41:a:2216:G:H2'	41:a:2217:G:C8	2.57	0.40
41:a:2246:G:H2'	41:a:2247:A:H8	1.86	0.40
41:a:2511:U:C5'	50:j:130:GLN:OE1	2.68	0.40
41:a:2728:U:H5'	60:t:70:ARG:HH22	1.85	0.40
42:b:55:ARG:CZ	42:b:55:ARG:CA	2.99	0.40
48:h:210:ALA:HA	48:h:213:TRP:CE3	2.56	0.40
62:v:45:GLN:NE2	62:v:124:LEU:HD23	2.37	0.40
66:z:66:ASN:OD1	66:z:70:ARG:NE	2.54	0.40
2:1:25:ARG:NH2	41:a:520:G:P	2.87	0.40
5:4:2:PHE:CZ	5:4:56:PHE:CZ	3.08	0.40
6:5:27:DG:N2	7:6:3:DC:O2	2.53	0.40
11:AA:524:ILE:HD12	11:AA:712:SER:HB2	2.02	0.40
12:AB:31:PRO:HG2	12:AB:51:PHE:CD2	2.55	0.40
13:AC:57:THR:HG22	13:AC:58:GLU:HG3	2.04	0.40
14:AE:291:ILE:HD13	14:AE:291:ILE:HA	1.85	0.40
14:AE:1175:LEU:HD22	14:AE:1190:ILE:HD11	2.03	0.40
16:AG:307:ILE:HG13	16:AG:338:VAL:HG23	2.02	0.40
18:D:18:C:C2	18:D:19:A:C8	3.09	0.40
18:D:74:A:C4	18:D:75:G:C8	3.09	0.40
18:D:135:C:H2'	35:U:1:MET:CE	2.51	0.40
18:D:553:A:H2'	18:D:554:A:C8	2.57	0.40
18:D:953:G:O6	38:X:103:LYS:HE2	2.22	0.40
18:D:983:A:C2	18:D:984:C:C6	3.10	0.40
20:F:4:ILE:HD12	20:F:19:PHE:HA	2.03	0.40
21:G:85:LEU:HD23	21:G:85:LEU:C	2.46	0.40
21:G:90:PHE:CD2	21:G:154:MET:SD	3.14	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:G:137:ARG:HG3	21:G:138:THR:N	2.36	0.40
22:H:116:VAL:O	22:H:279:LYS:HG2	2.21	0.40
22:H:309:MET:HE1	22:H:318:PRO:CA	2.51	0.40
24:J:74:ASN:CG	24:J:77:LYS:HZ3	2.29	0.40
27:M:109:ARG:CD	27:M:116:MET:HE1	2.49	0.40
33:S:10:GLU:CD	33:S:63:ARG:HD2	2.45	0.40
33:S:10:GLU:OE1	33:S:63:ARG:HG3	2.20	0.40
34:T:3:LEU:HB2	34:T:35:GLN:OE1	2.21	0.40
35:U:25:ARG:HA	35:U:25:ARG:HD2	1.95	0.40
38:X:36:ALA:HB3	38:X:59:GLU:HG2	2.03	0.40
39:Y:55:PRO:C	39:Y:71:LYS:CE	2.95	0.40
41:a:18:U:H2'	41:a:19:A:H8	1.86	0.40
41:a:404:A:O2'	41:a:405:U:P	2.78	0.40
41:a:789:A:N1	53:m:3:ARG:NH2	2.68	0.40
41:a:1239:G:H2'	41:a:1240:U:O4'	2.21	0.40
41:a:1449:G:C6	41:a:1450:G:N7	2.89	0.40
41:a:1589:U:HO2'	41:a:1590:A:P	2.45	0.40
41:a:2589:A:H2'	41:a:2590:A:H8	1.86	0.40
43:c:67:VAL:CA	43:c:70:GLU:OE1	2.69	0.40
44:d:31:C:O2	44:d:31:C:H2'	2.21	0.40
48:h:53:HIS:CD2	48:h:218:PRO:O	2.74	0.40
48:h:246:THR:HB	48:h:247:PRO:HD2	2.04	0.40
50:j:152:PRO:HG3	50:j:156:PHE:CE1	2.53	0.40
54:n:138:PHE:O	54:n:141:ILE:HG22	2.20	0.40
54:n:150:ARG:HG2	54:n:151:GLY:N	2.36	0.40
58:r:5:LEU:CD1	58:r:17:ASP:HB3	2.50	0.40
59:s:133:ALA:O	59:s:134:ALA:C	2.62	0.40
62:v:106:ASP:OD1	62:v:106:ASP:C	2.64	0.40
65:y:48:ILE:HD11	65:y:101:ARG:HH11	1.86	0.40
66:z:74:ILE:CD1	66:z:79:PHE:HA	2.40	0.40
2:1:32:ALA:O	2:1:36:LEU:CD1	2.70	0.40
9:9:43:LYS:HA	9:9:46:ARG:NH2	2.36	0.40
9:9:122:GLN:HG2	9:9:123:ILE:HG22	2.04	0.40
11:AA:213:LEU:HD13	11:AA:422:LYS:HG2	2.03	0.40
11:AA:741:MET:HE3	11:AA:974:ARG:HH12	1.85	0.40
12:AB:35:LEU:HA	12:AB:106:ASP:CG	2.33	0.40
16:AG:27:LEU:CD2	16:AG:114:ILE:HG23	2.51	0.40
16:AG:282:GLN:HA	16:AG:282:GLN:HE21	1.82	0.40
16:AG:433:ASP:O	16:AG:456:LEU:CD2	2.69	0.40
10:B:16:C:H3'	10:B:17:C:H5''	2.04	0.40
10:B:25:C:C4	10:B:26:G:C8	3.09	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:D:334:C:N3	18:D:335:C:C5	2.90	0.40
18:D:376:G:O3'	35:U:5:ARG:NE	2.54	0.40
18:D:597:G:N7	18:D:598:U:C5	2.89	0.40
18:D:1216:A:C2	18:D:1217:C:C5	3.10	0.40
18:D:1326:U:H2'	18:D:1327:C:C6	2.56	0.40
19:E:43:ASP:CG	19:E:46:ALA:HB3	2.46	0.40
21:G:30:PHE:HA	21:G:45:LYS:NZ	2.35	0.40
22:H:287:LEU:HD23	22:H:318:PRO:HB2	2.02	0.40
23:I:30:ALA:O	33:S:65:ARG:NH2	2.55	0.40
23:I:34:ASP:O	23:I:38:LYS:HG3	2.21	0.40
23:I:147:LYS:CE	23:I:204:LYS:O	2.69	0.40
24:J:138:SER:C	24:J:182:PHE:CE2	3.00	0.40
27:M:93:PRO:O	27:M:96:ARG:HG2	2.21	0.40
29:O:25:ASN:HB2	29:O:27:LYS:NZ	2.36	0.40
35:U:6:LEU:HD12	35:U:6:LEU:N	2.36	0.40
35:U:34:GLU:OE1	35:U:60:TRP:NE1	2.54	0.40
36:V:26:GLU:OE2	36:V:39:LYS:C	2.65	0.40
36:V:31:HIS:HD2	36:V:34:TYR:HB2	1.85	0.40
40:Z:29:LYS:C	40:Z:30:PHE:HD2	2.29	0.40
41:a:216:A:C8	41:a:432:A:C6	3.10	0.40
41:a:379:G:C6	41:a:396:G:O6	2.75	0.40
41:a:672:C:O2'	41:a:673:C:H5'	2.22	0.40
41:a:752:A:P	53:m:1:MET:HE1	2.62	0.40
41:a:1709:U:C2	41:a:1710:G:C8	3.10	0.40
41:a:2205:A:H2'	41:a:2206:C:C6	2.56	0.40
41:a:2223:G:OP1	48:h:171:TYR:OH	2.38	0.40
41:a:2812:G:H2'	41:a:2813:A:H8	1.87	0.40
46:f:25:LEU:C	46:f:25:LEU:HD23	2.46	0.40
48:h:84:ASP:OD2	48:h:91:ILE:HD11	2.21	0.40
50:j:45:TYR:CE2	50:j:81:GLU:OE2	2.75	0.40
50:j:175:LEU:HB3	50:j:189:VAL:HG12	2.02	0.40
55:o:33:LEU:O	55:o:41:LYS:NZ	2.45	0.40
59:s:4:PHE:CG	66:z:100:VAL:HG11	2.56	0.40
60:t:7:MET:CE	60:t:20:MET:HB2	2.51	0.40
60:t:58:LEU:N	60:t:58:LEU:HD23	2.36	0.40
61:u:123:ARG:HG3	61:u:143:GLU:OE2	2.21	0.40
4:3:4:LYS:O	4:3:72:ILE:HD11	2.22	0.40
5:4:78:GLN:OE1	5:4:88:HIS:HB3	2.21	0.40
8:7:9:U:O4	18:D:1196:A:C1'	2.70	0.40
9:9:43:LYS:HG2	9:9:98:GLU:CD	2.47	0.40
10:A:16:C:O2	41:a:2181:U:H5''	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AA:1247:SER:OG	11:AA:1248:THR:N	2.55	0.40
12:AB:147:ASN:C	12:AB:147:ASN:ND2	2.79	0.40
12:AB:158:GLU:HA	12:AB:162:LEU:HA	2.04	0.40
14:AE:144:TYR:HH	14:AE:293:ARG:HH22	1.63	0.40
14:AE:247:PRO:HA	14:AE:250:ARG:HE	1.86	0.40
14:AE:1178:THR:HG23	14:AE:1184:ASP:HB3	2.04	0.40
16:AG:425:LEU:HA	16:AG:429:LYS:HE3	0.95	0.40
18:D:72:A:C4	18:D:73:C:C6	3.10	0.40
18:D:228:A:C4'	35:U:63:GLN:CD	2.95	0.40
18:D:280:C:H1'	36:V:40:ARG:HH12	1.86	0.40
18:D:575:G:H4'	18:D:576:C:OP1	2.22	0.40
18:D:977:A:N6	18:D:1224:U:O5'	2.55	0.40
18:D:1095:U:H2'	18:D:1096:C:C6	2.57	0.40
18:D:1288:A:H2'	18:D:1289:A:O4'	2.21	0.40
19:E:9:LYS:HG3	19:E:13:GLN:HE22	1.87	0.40
19:E:15:GLU:CD	19:E:16:LYS:N	2.80	0.40
19:E:59:ASP:OD1	19:E:76:LYS:HE2	2.21	0.40
21:G:94:HIS:O	21:G:95:ARG:C	2.64	0.40
21:G:126:PHE:O	21:G:127:ASP:C	2.65	0.40
22:H:16:ILE:O	22:H:25:ARG:NH1	2.55	0.40
22:H:303:LEU:O	22:H:305:HIS:HA	2.21	0.40
24:J:71:GLN:O	24:J:74:ASN:HB3	2.22	0.40
30:P:21:ALA:HB1	30:P:92:LEU:HD23	2.03	0.40
36:V:77:ARG:HH21	36:V:78:VAL:H	1.69	0.40
41:a:134:G:H2'	41:a:135:U:H6	1.85	0.40
41:a:239:C:O2'	41:a:622:G:O2'	2.32	0.40
41:a:838:C:H2'	41:a:839:U:H6	1.87	0.40
41:a:849:A:H2'	41:a:850:U:H6	1.87	0.40
41:a:873:C:C2	41:a:874:G:C8	3.09	0.40
41:a:1144:A:C6	41:a:1145:C:C4	3.09	0.40
41:a:1944:U:C5	41:a:1955:U:C2	3.09	0.40
41:a:2072:C:N3	41:a:2073:C:C5	2.90	0.40
41:a:2314:A:O2'	54:n:155:THR:CG2	2.69	0.40
41:a:2639:A:H2'	41:a:2640:G:O4'	2.21	0.40
41:a:2650:U:C2	41:a:2651:C:C5	3.08	0.40
42:b:55:ARG:NE	42:b:55:ARG:CA	2.84	0.40
42:b:58:THR:O	42:b:58:THR:HG23	2.21	0.40
46:f:34:HIS:CD2	46:f:36:VAL:CG2	3.05	0.40
47:g:58:ASP:O	47:g:59:ARG:C	2.64	0.40
50:j:55:LYS:HD3	50:j:56:LYS:N	2.36	0.40
50:j:61:THR:CG2	50:j:64:GLU:HG2	2.52	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:j:171:THR:CG2	50:j:173:GLN:HE21	2.35	0.40
54:n:16:LEU:HD11	54:n:169:LEU:HD12	2.03	0.40
59:s:1:MET:SD	59:s:2:LYS:O	2.79	0.40
61:u:95:LEU:HD13	61:u:100:ILE:CD1	2.51	0.40
62:v:50:ARG:O	62:v:53:MET:SD	2.80	0.40
65:y:27:GLU:OE2	65:y:29:LYS:HG2	2.22	0.40

There are no symmetry-related clashes.

## 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	0	101/103 (98%)	97 (96%)	4 (4%)	0	100	100
2	1	108/110 (98%)	107 (99%)	1 (1%)	0	100	100
3	2	92/100 (92%)	87 (95%)	5 (5%)	0	100	100
4	3	101/104 (97%)	98 (97%)	3 (3%)	0	100	100
5	4	92/94 (98%)	89 (97%)	3 (3%)	0	100	100
9	9	146/165 (88%)	100 (68%)	43 (30%)	3 (2%)	5	24
11	AA	1312/1342 (98%)	1198 (91%)	113 (9%)	1 (0%)	48	78
12	AB	159/162 (98%)	103 (65%)	44 (28%)	12 (8%)	1	5
13	AC	217/329 (66%)	203 (94%)	12 (6%)	2 (1%)	14	41
13	AD	293/329 (89%)	263 (90%)	27 (9%)	3 (1%)	13	39
14	AE	1331/1407 (95%)	1211 (91%)	114 (9%)	6 (0%)	25	54
15	AF	80/91 (88%)	77 (96%)	3 (4%)	0	100	100
16	AG	493/495 (100%)	375 (76%)	86 (17%)	32 (6%)	1	7
17	C	64/75 (85%)	63 (98%)	1 (2%)	0	100	100
19	E	84/87 (97%)	83 (99%)	1 (1%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
20	F	68/71 (96%)	68 (100%)	0	0	100	100
21	G	223/241 (92%)	212 (95%)	10 (4%)	1 (0%)	30	60
22	H	255/557 (46%)	182 (71%)	66 (26%)	7 (3%)	4	21
23	I	206/233 (88%)	193 (94%)	13 (6%)	0	100	100
24	J	203/206 (98%)	201 (99%)	2 (1%)	0	100	100
25	K	154/167 (92%)	145 (94%)	8 (5%)	1 (1%)	22	50
26	L	102/135 (76%)	97 (95%)	4 (4%)	1 (1%)	13	39
27	M	149/179 (83%)	140 (94%)	8 (5%)	1 (1%)	19	47
28	N	127/130 (98%)	123 (97%)	4 (3%)	0	100	100
29	O	125/130 (96%)	116 (93%)	8 (6%)	1 (1%)	16	44
30	P	97/103 (94%)	89 (92%)	8 (8%)	0	100	100
31	Q	115/129 (89%)	107 (93%)	8 (7%)	0	100	100
32	R	117/124 (94%)	112 (96%)	5 (4%)	0	100	100
33	S	98/101 (97%)	97 (99%)	1 (1%)	0	100	100
34	T	86/89 (97%)	83 (96%)	3 (4%)	0	100	100
35	U	80/82 (98%)	76 (95%)	4 (5%)	0	100	100
36	V	78/84 (93%)	74 (95%)	4 (5%)	0	100	100
37	W	81/92 (88%)	80 (99%)	1 (1%)	0	100	100
38	X	114/118 (97%)	103 (90%)	9 (8%)	2 (2%)	7	27
39	Y	139/142 (98%)	101 (73%)	38 (27%)	0	100	100
40	Z	28/121 (23%)	22 (79%)	6 (21%)	0	100	100
42	b	74/85 (87%)	73 (99%)	1 (1%)	0	100	100
43	c	75/78 (96%)	72 (96%)	3 (4%)	0	100	100
45	e	60/63 (95%)	57 (95%)	3 (5%)	0	100	100
46	f	56/59 (95%)	52 (93%)	4 (7%)	0	100	100
47	g	64/70 (91%)	62 (97%)	2 (3%)	0	100	100
48	h	269/273 (98%)	255 (95%)	14 (5%)	0	100	100
49	i	54/57 (95%)	49 (91%)	5 (9%)	0	100	100
50	j	207/209 (99%)	198 (96%)	9 (4%)	0	100	100
51	k	50/55 (91%)	50 (100%)	0	0	100	100
52	l	199/201 (99%)	188 (94%)	11 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
53	m	44/46 (96%)	43 (98%)	1 (2%)	0	100	100
54	n	175/179 (98%)	160 (91%)	14 (8%)	1 (1%)	22	50
55	o	62/65 (95%)	57 (92%)	4 (6%)	1 (2%)	8	29
56	p	173/177 (98%)	162 (94%)	11 (6%)	0	100	100
57	q	36/38 (95%)	35 (97%)	1 (3%)	0	100	100
58	r	147/149 (99%)	139 (95%)	8 (5%)	0	100	100
59	s	140/142 (99%)	133 (95%)	7 (5%)	0	100	100
60	t	121/123 (98%)	114 (94%)	7 (6%)	0	100	100
61	u	142/144 (99%)	134 (94%)	8 (6%)	0	100	100
62	v	134/136 (98%)	129 (96%)	5 (4%)	0	100	100
63	w	117/127 (92%)	112 (96%)	5 (4%)	0	100	100
64	x	114/117 (97%)	107 (94%)	7 (6%)	0	100	100
65	y	112/115 (97%)	105 (94%)	7 (6%)	0	100	100
66	z	115/118 (98%)	111 (96%)	4 (4%)	0	100	100
All	All	10058/11053 (91%)	9172 (91%)	811 (8%)	75 (1%)	21	47

All (75) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
12	AB	107	PRO
12	AB	130	PHE
12	AB	137	ALA
12	AB	144	ASN
12	AB	160	ARG
13	AD	286	GLU
16	AG	34	ALA
16	AG	105	ILE
16	AG	187	ARG
16	AG	227	ALA
21	G	127	ASP
22	H	304	VAL
27	M	56	LYS
55	o	32	ILE
12	AB	105	VAL
12	AB	127	GLN
14	AE	92	VAL
14	AE	175	GLU

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Mol	Chain	Res	Type
16	AG	63	LEU
16	AG	104	ARG
16	AG	242	ASP
16	AG	268	GLY
16	AG	279	ASN
16	AG	350	ALA
16	AG	365	ILE
22	H	171	ARG
22	H	305	HIS
22	H	309	MET
12	AB	141	LEU
12	AB	152	HIS
16	AG	33	THR
16	AG	102	PHE
16	AG	200	SER
16	AG	305	MET
16	AG	319	GLY
16	AG	355	ALA
16	AG	396	GLU
16	AG	425	LEU
25	K	78	ASN
26	L	96	VAL
29	O	13	LYS
54	n	177	PHE
13	AC	193	GLU
16	AG	48	GLN
16	AG	265	GLU
16	AG	363	LEU
16	AG	401	PRO
22	H	82	THR
12	AB	53	ASN
13	AD	289	LEU
14	AE	74	LYS
14	AE	286	ALA
16	AG	224	LYS
16	AG	399	ASP
22	H	143	ASP
38	X	6	GLY
38	X	66	GLU
9	9	79	PRO
9	9	88	HIS
13	AC	192	VAL

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Mol	Chain	Res	Type
13	AD	306	VAL
14	AE	290	ILE
16	AG	43	ILE
16	AG	70	GLN
16	AG	169	PRO
16	AG	171	GLU
16	AG	323	GLN
16	AG	364	ASP
16	AG	382	GLU
22	H	71	ALA
14	AE	292	VAL
9	9	129	LEU
11	AA	1317	PRO
12	AB	150	ILE
12	AB	116	VAL

### 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.

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	0	84/84 (100%)	84 (100%)	0	100	100
2	1	93/93 (100%)	93 (100%)	0	100	100
3	2	81/84 (96%)	81 (100%)	0	100	100
4	3	84/85 (99%)	84 (100%)	0	100	100
5	4	78/78 (100%)	78 (100%)	0	100	100
9	9	112/123 (91%)	112 (100%)	0	100	100
11	AA	1135/1157 (98%)	1130 (100%)	5 (0%)	89	93
12	AB	141/142 (99%)	117 (83%)	24 (17%)	1	6
13	AC	186/286 (65%)	186 (100%)	0	100	100
13	AD	185/286 (65%)	185 (100%)	0	100	100
14	AE	1122/1168 (96%)	1104 (98%)	18 (2%)	58	75
15	AF	70/75 (93%)	69 (99%)	1 (1%)	62	77

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
16	AG	409/409 (100%)	288 (70%)	121 (30%)	0	1
17	C	57/65 (88%)	57 (100%)	0	100	100
19	E	65/66 (98%)	65 (100%)	0	100	100
20	F	60/61 (98%)	60 (100%)	0	100	100
21	G	187/199 (94%)	187 (100%)	0	100	100
22	H	137/461 (30%)	137 (100%)	0	100	100
23	I	171/190 (90%)	171 (100%)	0	100	100
24	J	172/173 (99%)	171 (99%)	1 (1%)	84	90
25	K	119/126 (94%)	119 (100%)	0	100	100
26	L	91/116 (78%)	91 (100%)	0	100	100
27	M	124/147 (84%)	124 (100%)	0	100	100
28	N	104/105 (99%)	104 (100%)	0	100	100
29	O	105/107 (98%)	105 (100%)	0	100	100
30	P	86/90 (96%)	86 (100%)	0	100	100
31	Q	90/99 (91%)	90 (100%)	0	100	100
32	R	101/104 (97%)	101 (100%)	0	100	100
33	S	83/84 (99%)	83 (100%)	0	100	100
34	T	76/77 (99%)	76 (100%)	0	100	100
35	U	65/65 (100%)	65 (100%)	0	100	100
36	V	74/78 (95%)	74 (100%)	0	100	100
37	W	72/79 (91%)	72 (100%)	0	100	100
38	X	94/96 (98%)	94 (100%)	0	100	100
39	Y	109/110 (99%)	109 (100%)	0	100	100
40	Z	26/85 (31%)	26 (100%)	0	100	100
42	b	58/63 (92%)	58 (100%)	0	100	100
43	c	67/68 (98%)	67 (100%)	0	100	100
45	e	54/55 (98%)	54 (100%)	0	100	100
46	f	48/49 (98%)	48 (100%)	0	100	100
47	g	59/62 (95%)	59 (100%)	0	100	100
48	h	216/218 (99%)	216 (100%)	0	100	100
49	i	47/48 (98%)	47 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
50	j	164/164 (100%)	164 (100%)	0	100	100
51	k	47/49 (96%)	47 (100%)	0	100	100
52	l	165/165 (100%)	165 (100%)	0	100	100
53	m	38/38 (100%)	38 (100%)	0	100	100
54	n	148/150 (99%)	148 (100%)	0	100	100
55	o	51/52 (98%)	51 (100%)	0	100	100
56	p	136/138 (99%)	136 (100%)	0	100	100
57	q	34/34 (100%)	34 (100%)	0	100	100
58	r	114/114 (100%)	114 (100%)	0	100	100
59	s	116/116 (100%)	116 (100%)	0	100	100
60	t	104/104 (100%)	104 (100%)	0	100	100
61	u	103/103 (100%)	103 (100%)	0	100	100
62	v	109/109 (100%)	109 (100%)	0	100	100
63	w	99/103 (96%)	99 (100%)	0	100	100
64	x	86/87 (99%)	86 (100%)	0	100	100
65	y	99/100 (99%)	99 (100%)	0	100	100
66	z	89/90 (99%)	89 (100%)	0	100	100
All	All	8299/9132 (91%)	8129 (98%)	170 (2%)	50	70

All (170) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
11	AA	145	ILE
11	AA	615	VAL
11	AA	914	LYS
11	AA	1076	ILE
11	AA	1151	LEU
12	AB	115	LYS
12	AB	116	VAL
12	AB	117	ILE
12	AB	118	ILE
12	AB	119	THR
12	AB	120	GLU
12	AB	123	PHE
12	AB	124	GLU
12	AB	126	PHE

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Mol	Chain	Res	Type
12	AB	127	GLN
12	AB	129	ILE
12	AB	138	ARG
12	AB	141	LEU
12	AB	145	LEU
12	AB	146	ILE
12	AB	147	ASN
12	AB	148	LYS
12	AB	149	GLU
12	AB	150	ILE
12	AB	151	LYS
12	AB	154	VAL
12	AB	159	PHE
12	AB	160	ARG
12	AB	161	LYS
14	AE	69	GLU
14	AE	70	CYS
14	AE	71	LEU
14	AE	74	LYS
14	AE	76	LYS
14	AE	77	ARG
14	AE	78	LEU
14	AE	79	LYS
14	AE	81	ARG
14	AE	84	ILE
14	AE	85	CYS
14	AE	92	VAL
14	AE	93	THR
14	AE	290	ILE
14	AE	291	ILE
14	AE	292	VAL
14	AE	514	THR
14	AE	1261	LEU
15	AF	63	ILE
16	AG	5	ILE
16	AG	12	VAL
16	AG	16	LYS
16	AG	36	LYS
16	AG	37	LYS
16	AG	40	GLU
16	AG	43	ILE
16	AG	47	VAL

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Mol	Chain	Res	Type
16	AG	49	ILE
16	AG	50	ASP
16	AG	56	PHE
16	AG	57	ASP
16	AG	61	ARG
16	AG	63	LEU
16	AG	75	ILE
16	AG	97	ILE
16	AG	103	ASP
16	AG	104	ARG
16	AG	114	ILE
16	AG	117	LYS
16	AG	123	ARG
16	AG	125	MET
16	AG	126	VAL
16	AG	131	ARG
16	AG	138	ILE
16	AG	139	THR
16	AG	144	LYS
16	AG	150	ILE
16	AG	154	LEU
16	AG	157	ASN
16	AG	162	ILE
16	AG	167	MET
16	AG	173	PHE
16	AG	174	ARG
16	AG	179	VAL
16	AG	180	ARG
16	AG	182	VAL
16	AG	187	ARG
16	AG	189	GLU
16	AG	195	LEU
16	AG	197	VAL
16	AG	199	ARG
16	AG	201	LYS
16	AG	203	GLU
16	AG	204	MET
16	AG	205	LEU
16	AG	206	ILE
16	AG	207	GLU
16	AG	210	ARG
16	AG	211	ILE

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Mol	Chain	Res	Type
16	AG	213	VAL
16	AG	218	GLU
16	AG	219	GLU
16	AG	221	ILE
16	AG	223	ILE
16	AG	232	SER
16	AG	235	LYS
16	AG	236	ILE
16	AG	238	VAL
16	AG	239	LYS
16	AG	240	THR
16	AG	241	ASN
16	AG	243	LYS
16	AG	244	ARG
16	AG	245	ILE
16	AG	254	MET
16	AG	255	ARG
16	AG	258	ARG
16	AG	259	VAL
16	AG	260	GLN
16	AG	262	VAL
16	AG	265	GLU
16	AG	270	ARG
16	AG	271	ILE
16	AG	273	ILE
16	AG	274	VAL
16	AG	276	TRP
16	AG	278	ASP
16	AG	282	GLN
16	AG	285	ILE
16	AG	288	MET
16	AG	296	ILE
16	AG	297	VAL
16	AG	299	ASP
16	AG	300	GLU
16	AG	301	ASP
16	AG	302	LYS
16	AG	303	HIS
16	AG	305	MET
16	AG	309	VAL
16	AG	310	GLU
16	AG	313	ASN

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Mol	Chain	Res	Type
16	AG	316	GLN
16	AG	320	ARG
16	AG	321	ASN
16	AG	323	GLN
16	AG	327	LEU
16	AG	331	LEU
16	AG	335	GLU
16	AG	336	LEU
16	AG	337	ASN
16	AG	338	VAL
16	AG	340	THR
16	AG	341	VAL
16	AG	342	ASP
16	AG	344	LEU
16	AG	356	ILE
16	AG	358	THR
16	AG	359	PHE
16	AG	361	LYS
16	AG	365	ILE
16	AG	380	THR
16	AG	387	VAL
16	AG	390	LYS
16	AG	396	GLU
16	AG	398	LEU
16	AG	399	ASP
16	AG	400	GLU
16	AG	425	LEU
16	AG	428	ASN
16	AG	429	LYS
24	J	47	ARG

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (116) such sidechains are listed below:

Mol	Chain	Res	Type
1	0	6	GLN
1	0	82	HIS
2	1	7	HIS
2	1	9	HIS
3	2	15	HIS
9	9	9	GLN
11	AA	69	GLN
11	AA	314	ASN

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Mol	Chain	Res	Type
11	AA	330	HIS
11	AA	343	HIS
11	AA	387	ASN
11	AA	513	GLN
11	AA	554	HIS
11	AA	688	GLN
11	AA	808	ASN
11	AA	856	ASN
11	AA	1237	HIS
11	AA	1268	GLN
11	AA	1288	GLN
11	AA	1313	HIS
12	AB	27	ASN
12	AB	65	HIS
12	AB	147	ASN
13	AC	23	HIS
13	AD	66	HIS
13	AD	84	ASN
13	AD	117	HIS
13	AD	227	GLN
14	AE	80	HIS
14	AE	157	GLN
14	AE	164	GLN
14	AE	232	ASN
14	AE	365	GLN
14	AE	424	ASN
14	AE	450	HIS
14	AE	469	HIS
14	AE	777	HIS
14	AE	910	ASN
14	AE	1108	GLN
14	AE	1235	ASN
14	AE	1238	GLN
14	AE	1326	GLN
15	AF	62	GLN
16	AG	108	GLN
16	AG	194	GLN
16	AG	260	GLN
16	AG	282	GLN
16	AG	316	GLN
16	AG	323	GLN
16	AG	324	ASN

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Mol	Chain	Res	Type
16	AG	337	ASN
16	AG	428	ASN
17	C	54	GLN
19	E	13	GLN
19	E	61	GLN
20	F	9	ASN
21	G	18	HIS
21	G	36	ASN
21	G	39	HIS
21	G	94	HIS
23	I	3	GLN
23	I	6	HIS
23	I	8	ASN
23	I	190	HIS
24	J	36	GLN
24	J	131	ASN
24	J	198	HIS
25	K	89	HIS
26	L	3	HIS
26	L	68	GLN
27	M	68	ASN
27	M	86	GLN
29	O	5	GLN
29	O	81	HIS
29	O	126	GLN
32	R	77	HIS
33	S	4	GLN
34	T	80	GLN
35	U	26	ASN
35	U	59	HIS
39	Y	5	GLN
45	e	25	GLN
45	e	31	GLN
46	f	20	HIS
46	f	49	ASN
47	g	33	ASN
48	h	53	HIS
48	h	90	ASN
49	i	6	ASN
50	j	173	GLN
51	k	19	HIS
52	l	92	HIS

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Mol	Chain	Res	Type
52	l	94	GLN
52	l	156	ASN
52	l	163	ASN
53	m	29	GLN
54	n	5	HIS
54	n	81	GLN
55	o	24	HIS
55	o	28	ASN
57	q	13	ASN
58	r	133	GLN
59	s	86	GLN
59	s	132	HIS
62	v	13	HIS
62	v	45	GLN
62	v	97	GLN
63	w	18	GLN
64	x	19	GLN
64	x	29	HIS
64	x	38	GLN
65	y	10	GLN
65	y	77	HIS
66	z	37	GLN
66	z	72	ASN
66	z	81	ASN

### 5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
10	A	75/76 (98%)	29 (38%)	8 (10%)
10	B	75/76 (98%)	29 (38%)	8 (10%)
18	D	1514/1542 (98%)	289 (19%)	20 (1%)
41	a	2859/2904 (98%)	510 (17%)	0
44	d	119/120 (99%)	15 (12%)	0
8	7	23/56 (41%)	14 (60%)	3 (13%)
All	All	4665/4774 (97%)	886 (18%)	39 (0%)

All (886) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
8	7	3	G
8	7	4	U

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Mol	Chain	Res	Type
8	7	5	U
8	7	7	U
8	7	8	U
8	7	9	U
8	7	10	U
8	7	11	U
8	7	12	U
8	7	13	U
8	7	14	U
8	7	47	U
8	7	50	U
8	7	52	A
10	A	2	G
10	A	6	G
10	A	7	G
10	A	8	U
10	A	10	G
10	A	13	C
10	A	14	A
10	A	16	C
10	A	17	C
10	A	18	G
10	A	19	G
10	A	20	U
10	A	21	A
10	A	22	G
10	A	23	C
10	A	30	G
10	A	46	G
10	A	47	U
10	A	48	C
10	A	49	G
10	A	52	G
10	A	57	A
10	A	58	A
10	A	59	A
10	A	61	C
10	A	66	C
10	A	69	C
10	A	71	C
10	A	73	A
10	B	2	G

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Mol	Chain	Res	Type
10	B	6	G
10	B	7	G
10	B	8	U
10	B	10	G
10	B	13	C
10	B	14	A
10	B	16	C
10	B	17	C
10	B	18	G
10	B	19	G
10	B	20	U
10	B	21	A
10	B	22	G
10	B	23	C
10	B	30	G
10	B	46	G
10	B	47	U
10	B	48	C
10	B	49	G
10	B	52	G
10	B	57	A
10	B	58	A
10	B	59	A
10	B	61	C
10	B	66	C
10	B	69	C
10	B	71	C
10	B	73	A
18	D	4	U
18	D	5	U
18	D	9	G
18	D	22	G
18	D	29	U
18	D	32	A
18	D	39	G
18	D	47	C
18	D	48	C
18	D	50	A
18	D	51	A
18	D	52	C
18	D	54	C
18	D	69	G

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Mol	Chain	Res	Type
18	D	71	A
18	D	72	A
18	D	74	A
18	D	76	G
18	D	82	G
18	D	83	C
18	D	84	U
18	D	87	C
18	D	90	C
18	D	94	G
18	D	95	C
18	D	96	U
18	D	108	G
18	D	120	A
18	D	121	U
18	D	122	G
18	D	131	A
18	D	141	G
18	D	144	G
18	D	148	G
18	D	149	A
18	D	160	A
18	D	164	G
18	D	173	U
18	D	181	A
18	D	182	A
18	D	183	C
18	D	184	G
18	D	197	A
18	D	198	G
18	D	204	G
18	D	208	U
18	D	209	U
18	D	210	C
18	D	211	G
18	D	212	G
18	D	216	U
18	D	226	G
18	D	245	U
18	D	247	G
18	D	251	G
18	D	253	A

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Mol	Chain	Res	Type
18	D	258	G
18	D	262	A
18	D	266	G
18	D	267	C
18	D	271	C
18	D	279	A
18	D	289	G
18	D	299	G
18	D	306	A
18	D	321	A
18	D	328	C
18	D	329	A
18	D	332	G
18	D	347	G
18	D	352	C
18	D	353	A
18	D	354	G
18	D	355	C
18	D	367	U
18	D	372	C
18	D	373	A
18	D	376	G
18	D	382	A
18	D	384	G
18	D	392	C
18	D	397	A
18	D	398	U
18	D	406	G
18	D	411	A
18	D	412	A
18	D	413	G
18	D	414	A
18	D	421	U
18	D	422	C
18	D	424	G
18	D	429	U
18	D	446	G
18	D	451	A
18	D	457	G
18	D	458	U
18	D	463	U
18	D	464	U

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Mol	Chain	Res	Type
18	D	467	U
18	D	468	A
18	D	469	C
18	D	478	A
18	D	479	U
18	D	481	G
18	D	484	G
18	D	485	U
18	D	486	U
18	D	496	A
18	D	497	G
18	D	505	G
18	D	511	C
18	D	518	C
18	D	519	C
18	D	526	C
18	D	531	U
18	D	532	A
18	D	533	A
18	D	542	G
18	D	547	A
18	D	559	A
18	D	564	C
18	D	568	G
18	D	572	A
18	D	573	A
18	D	576	C
18	D	577	G
18	D	579	A
18	D	596	A
18	D	628	G
18	D	633	G
18	D	649	A
18	D	650	G
18	D	653	U
18	D	665	A
18	D	695	A
18	D	700	G
18	D	723	U
18	D	724	G
18	D	731	G
18	D	734	G

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Mol	Chain	Res	Type
18	D	735	C
18	D	747	A
18	D	748	G
18	D	755	G
18	D	760	G
18	D	777	A
18	D	793	U
18	D	794	A
18	D	815	A
18	D	817	C
18	D	828	U
18	D	829	G
18	D	832	G
18	D	841	C
18	D	843	U
18	D	844	G
18	D	845	A
18	D	846	G
18	D	849	G
18	D	874	G
18	D	887	G
18	D	902	G
18	D	914	A
18	D	916	U
18	D	926	G
18	D	934	C
18	D	935	A
18	D	960	U
18	D	963	G
18	D	965	U
18	D	969	A
18	D	972	C
18	D	975	A
18	D	976	G
18	D	987	G
18	D	991	U
18	D	992	U
18	D	993	G
18	D	996	A
18	D	1004	A
18	D	1008	U
18	D	1009	U

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Mol	Chain	Res	Type
18	D	1017	U
18	D	1018	G
18	D	1021	A
18	D	1024	G
18	D	1026	G
18	D	1028	C
18	D	1030	U
18	D	1031	C
18	D	1037	C
18	D	1043	G
18	D	1044	A
18	D	1046	A
18	D	1065	U
18	D	1085	U
18	D	1094	G
18	D	1095	U
18	D	1099	G
18	D	1101	A
18	D	1108	G
18	D	1124	G
18	D	1133	G
18	D	1135	U
18	D	1136	C
18	D	1137	C
18	D	1139	G
18	D	1140	C
18	D	1141	C
18	D	1142	G
18	D	1143	G
18	D	1145	A
18	D	1146	A
18	D	1151	A
18	D	1152	A
18	D	1159	U
18	D	1167	A
18	D	1171	A
18	D	1174	G
18	D	1175	G
18	D	1176	A
18	D	1184	G
18	D	1193	G
18	D	1196	A

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Mol	Chain	Res	Type
18	D	1197	A
18	D	1206	G
18	D	1211	U
18	D	1212	U
18	D	1213	A
18	D	1214	C
18	D	1215	G
18	D	1226	C
18	D	1227	A
18	D	1228	C
18	D	1238	A
18	D	1257	A
18	D	1260	G
18	D	1275	A
18	D	1276	G
18	D	1278	G
18	D	1279	G
18	D	1280	A
18	D	1285	A
18	D	1286	U
18	D	1287	A
18	D	1299	A
18	D	1300	G
18	D	1302	C
18	D	1305	G
18	D	1312	G
18	D	1317	C
18	D	1320	C
18	D	1323	G
18	D	1338	G
18	D	1340	A
18	D	1346	A
18	D	1347	G
18	D	1353	G
18	D	1363	A
18	D	1370	G
18	D	1378	C
18	D	1379	G
18	D	1381	U
18	D	1391	U
18	D	1396	A
18	D	1397	C

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Mol	Chain	Res	Type
18	D	1398	A
18	D	1404	C
18	D	1419	G
18	D	1429	A
18	D	1441	A
18	D	1446	A
18	D	1447	A
18	D	1448	C
18	D	1452	C
18	D	1453	G
18	D	1475	G
18	D	1487	G
18	D	1491	G
18	D	1492	A
18	D	1493	A
18	D	1494	G
18	D	1497	G
18	D	1503	A
18	D	1506	U
18	D	1517	G
18	D	1529	G
18	D	1530	G
18	D	1534	A
41	a	4	U
41	a	10	A
41	a	15	G
41	a	23	G
41	a	34	U
41	a	35	G
41	a	46	G
41	a	58	G
41	a	60	G
41	a	62	U
41	a	63	A
41	a	71	A
41	a	74	A
41	a	75	G
41	a	83	A
41	a	84	A
41	a	85	G
41	a	96	C
41	a	101	A

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Mol	Chain	Res	Type
41	a	102	U
41	a	103	A
41	a	110	G
41	a	118	A
41	a	119	A
41	a	120	U
41	a	131	A
41	a	136	G
41	a	139	U
41	a	140	C
41	a	141	G
41	a	163	C
41	a	165	A
41	a	181	A
41	a	196	A
41	a	215	G
41	a	216	A
41	a	222	A
41	a	225	C
41	a	248	G
41	a	249	C
41	a	261	G
41	a	264	C
41	a	265	A
41	a	266	G
41	a	267	C
41	a	271	G
41	a	272	A
41	a	275	C
41	a	276	U
41	a	278	A
41	a	285	G
41	a	291	G
41	a	311	A
41	a	329	G
41	a	330	A
41	a	353	C
41	a	361	G
41	a	362	A
41	a	371	A
41	a	372	G
41	a	373	U

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Mol	Chain	Res	Type
41	a	375	G
41	a	383	C
41	a	386	G
41	a	396	G
41	a	405	U
41	a	411	G
41	a	412	A
41	a	420	C
41	a	424	G
41	a	451	U
41	a	457	A
41	a	477	A
41	a	481	G
41	a	491	G
41	a	501	A
41	a	503	A
41	a	504	A
41	a	505	A
41	a	509	C
41	a	522	A
41	a	529	A
41	a	531	C
41	a	532	A
41	a	537	G
41	a	543	G
41	a	545	U
41	a	546	U
41	a	547	A
41	a	549	G
41	a	551	G
41	a	563	A
41	a	569	U
41	a	573	U
41	a	575	A
41	a	588	U
41	a	603	A
41	a	609	A
41	a	613	A
41	a	614	A
41	a	615	U
41	a	616	A
41	a	627	A

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Mol	Chain	Res	Type
41	a	637	A
41	a	645	C
41	a	647	G
41	a	654	A
41	a	668	A
41	a	686	U
41	a	710	U
41	a	717	C
41	a	730	A
41	a	738	G
41	a	757	G
41	a	764	A
41	a	765	C
41	a	775	G
41	a	776	G
41	a	782	A
41	a	784	G
41	a	785	G
41	a	800	A
41	a	805	G
41	a	812	C
41	a	819	A
41	a	827	U
41	a	828	U
41	a	845	A
41	a	846	U
41	a	858	G
41	a	859	G
41	a	869	G
41	a	878	A
41	a	881	G
41	a	884	U
41	a	885	C
41	a	888	C
41	a	891	G
41	a	895	U
41	a	896	A
41	a	897	C
41	a	899	A
41	a	907	G
41	a	910	A
41	a	914	G

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Mol	Chain	Res	Type
41	a	915	C
41	a	931	U
41	a	941	A
41	a	945	A
41	a	946	C
41	a	953	G
41	a	961	C
41	a	974	G
41	a	983	A
41	a	995	C
41	a	996	A
41	a	999	U
41	a	1005	C
41	a	1012	U
41	a	1013	C
41	a	1022	G
41	a	1023	U
41	a	1026	G
41	a	1033	U
41	a	1041	G
41	a	1045	C
41	a	1046	A
41	a	1047	G
41	a	1060	U
41	a	1061	U
41	a	1064	C
41	a	1066	U
41	a	1067	A
41	a	1068	G
41	a	1070	A
41	a	1071	G
41	a	1073	A
41	a	1074	G
41	a	1079	C
41	a	1080	A
41	a	1081	U
41	a	1082	U
41	a	1083	U
41	a	1084	A
41	a	1087	G
41	a	1088	A
41	a	1090	A

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Mol	Chain	Res	Type
41	a	1095	A
41	a	1101	U
41	a	1107	G
41	a	1110	G
41	a	1111	A
41	a	1112	G
41	a	1119	U
41	a	1122	G
41	a	1128	G
41	a	1132	U
41	a	1134	A
41	a	1135	C
41	a	1142	A
41	a	1169	A
41	a	1170	C
41	a	1173	U
41	a	1174	U
41	a	1175	A
41	a	1176	U
41	a	1177	G
41	a	1178	C
41	a	1179	G
41	a	1180	U
41	a	1186	G
41	a	1238	G
41	a	1248	G
41	a	1253	A
41	a	1256	G
41	a	1266	G
41	a	1271	G
41	a	1272	A
41	a	1273	U
41	a	1300	G
41	a	1301	A
41	a	1321	A
41	a	1345	C
41	a	1352	U
41	a	1365	A
41	a	1368	G
41	a	1378	A
41	a	1379	U
41	a	1380	G

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Mol	Chain	Res	Type
41	a	1383	A
41	a	1392	A
41	a	1395	A
41	a	1406	U
41	a	1408	G
41	a	1409	U
41	a	1414	C
41	a	1416	G
41	a	1417	C
41	a	1419	A
41	a	1420	A
41	a	1428	C
41	a	1452	G
41	a	1453	A
41	a	1460	U
41	a	1482	G
41	a	1490	A
41	a	1497	U
41	a	1503	A
41	a	1508	A
41	a	1509	A
41	a	1510	G
41	a	1515	A
41	a	1529	G
41	a	1534	U
41	a	1535	A
41	a	1536	C
41	a	1537	G
41	a	1554	U
41	a	1559	U
41	a	1566	A
41	a	1569	A
41	a	1578	U
41	a	1580	A
41	a	1581	G
41	a	1582	C
41	a	1583	A
41	a	1584	U
41	a	1585	C
41	a	1589	U
41	a	1590	A
41	a	1608	A

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Mol	Chain	Res	Type
41	a	1610	A
41	a	1613	G
41	a	1647	U
41	a	1648	U
41	a	1649	G
41	a	1651	G
41	a	1665	A
41	a	1674	G
41	a	1677	A
41	a	1703	G
41	a	1714	U
41	a	1715	G
41	a	1729	U
41	a	1730	C
41	a	1732	C
41	a	1738	G
41	a	1750	G
41	a	1758	U
41	a	1764	C
41	a	1773	A
41	a	1791	A
41	a	1800	C
41	a	1808	A
41	a	1811	G
41	a	1816	C
41	a	1829	A
41	a	1833	C
41	a	1847	A
41	a	1848	A
41	a	1858	A
41	a	1859	U
41	a	1862	G
41	a	1864	U
41	a	1869	G
41	a	1870	C
41	a	1872	A
41	a	1873	G
41	a	1906	G
41	a	1907	G
41	a	1913	A
41	a	1914	C
41	a	1919	A

*Continued on next page...*

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Mol	Chain	Res	Type
41	a	1920	C
41	a	1922	G
41	a	1923	U
41	a	1924	C
41	a	1925	C
41	a	1926	U
41	a	1929	G
41	a	1930	G
41	a	1936	A
41	a	1938	A
41	a	1955	U
41	a	1965	C
41	a	1967	C
41	a	1970	A
41	a	1971	U
41	a	1972	G
41	a	1987	A
41	a	1991	U
41	a	1992	G
41	a	1993	U
41	a	1997	C
41	a	2002	G
41	a	2020	A
41	a	2022	U
41	a	2023	C
41	a	2027	G
41	a	2033	A
41	a	2043	C
41	a	2051	A
41	a	2052	A
41	a	2055	C
41	a	2056	G
41	a	2060	A
41	a	2061	G
41	a	2062	A
41	a	2063	C
41	a	2097	A
41	a	2099	U
41	a	2100	G
41	a	2108	A
41	a	2110	G
41	a	2111	U

*Continued on next page...*



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Mol	Chain	Res	Type
41	a	2113	U
41	a	2115	G
41	a	2116	G
41	a	2117	A
41	a	2118	U
41	a	2121	G
41	a	2122	U
41	a	2124	G
41	a	2125	G
41	a	2126	A
41	a	2127	G
41	a	2128	G
41	a	2131	U
41	a	2132	U
41	a	2133	G
41	a	2134	A
41	a	2139	U
41	a	2141	G
41	a	2146	C
41	a	2147	A
41	a	2154	A
41	a	2157	G
41	a	2158	A
41	a	2159	G
41	a	2162	G
41	a	2163	A
41	a	2164	C
41	a	2165	C
41	a	2169	A
41	a	2171	A
41	a	2172	U
41	a	2178	C
41	a	2182	U
41	a	2183	A
41	a	2185	U
41	a	2189	U
41	a	2190	G
41	a	2193	G
41	a	2194	U
41	a	2198	A
41	a	2204	G
41	a	2211	A

*Continued on next page...*

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Mol	Chain	Res	Type
41	a	2212	A
41	a	2225	A
41	a	2226	C
41	a	2229	U
41	a	2238	G
41	a	2239	G
41	a	2250	G
41	a	2268	A
41	a	2278	A
41	a	2283	C
41	a	2287	A
41	a	2288	A
41	a	2297	A
41	a	2305	U
41	a	2308	G
41	a	2309	A
41	a	2322	A
41	a	2325	G
41	a	2327	A
41	a	2333	A
41	a	2335	A
41	a	2339	C
41	a	2345	G
41	a	2347	C
41	a	2350	C
41	a	2361	G
41	a	2372	U
41	a	2376	A
41	a	2383	G
41	a	2385	C
41	a	2402	U
41	a	2403	C
41	a	2406	A
41	a	2423	U
41	a	2424	C
41	a	2425	A
41	a	2426	A
41	a	2429	G
41	a	2430	A
41	a	2431	U
41	a	2434	A
41	a	2435	A

*Continued on next page...*

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Mol	Chain	Res	Type
41	a	2441	U
41	a	2447	G
41	a	2448	A
41	a	2470	G
41	a	2474	U
41	a	2476	A
41	a	2478	A
41	a	2484	G
41	a	2491	U
41	a	2502	G
41	a	2506	U
41	a	2512	C
41	a	2513	A
41	a	2518	A
41	a	2520	C
41	a	2525	G
41	a	2529	G
41	a	2535	G
41	a	2554	U
41	a	2566	A
41	a	2567	G
41	a	2573	C
41	a	2574	G
41	a	2585	U
41	a	2586	U
41	a	2602	A
41	a	2603	G
41	a	2609	U
41	a	2610	C
41	a	2611	C
41	a	2613	U
41	a	2629	U
41	a	2630	G
41	a	2663	G
41	a	2669	G
41	a	2671	G
41	a	2689	U
41	a	2690	U
41	a	2714	G
41	a	2716	C
41	a	2726	A
41	a	2744	G

*Continued on next page...*

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Mol	Chain	Res	Type
41	a	2748	A
41	a	2758	A
41	a	2765	A
41	a	2777	G
41	a	2778	A
41	a	2791	G
41	a	2793	C
41	a	2796	U
41	a	2797	U
41	a	2798	U
41	a	2799	A
41	a	2801	G
41	a	2818	U
41	a	2820	A
41	a	2823	A
41	a	2825	G
41	a	2835	A
41	a	2849	U
41	a	2859	G
41	a	2861	U
41	a	2867	G
41	a	2873	A
41	a	2874	C
41	a	2880	C
41	a	2883	A
41	a	2884	U
41	a	2885	G
41	a	2891	U
41	a	2902	C
44	d	2	G
44	d	13	G
44	d	16	G
44	d	17	C
44	d	35	C
44	d	45	A
44	d	51	G
44	d	56	G
44	d	57	A
44	d	66	A
44	d	88	C
44	d	89	U
44	d	90	C

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Mol	Chain	Res	Type
44	d	99	A
44	d	109	A

All (39) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
8	7	4	U
8	7	7	U
8	7	10	U
10	A	6	G
10	A	7	G
10	A	9	G
10	A	12	G
10	A	21	A
10	A	22	G
10	A	57	A
10	A	60	U
10	B	6	G
10	B	7	G
10	B	9	G
10	B	12	G
10	B	21	A
10	B	22	G
10	B	57	A
10	B	60	U
18	D	121	U
18	D	181	A
18	D	183	C
18	D	197	A
18	D	209	U
18	D	428	G
18	D	496	A
18	D	517	G
18	D	991	U
18	D	992	U
18	D	1145	A
18	D	1196	A
18	D	1211	U
18	D	1212	U
18	D	1213	A
18	D	1214	C
18	D	1447	A

*Continued on next page...*

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Mol	Chain	Res	Type
18	D	1491	G
18	D	1492	A
18	D	1493	A

## 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](#)

Of 3 ligands modelled in this entry, 3 are monoatomic - leaving 0 for Mogul analysis.

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

## 5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

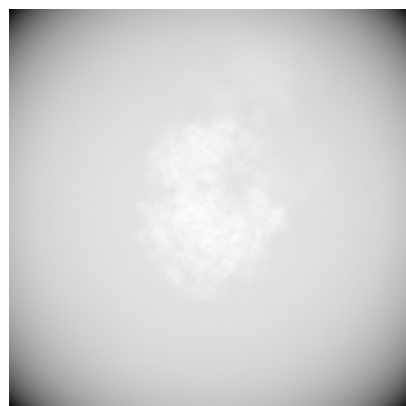
## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-43387. These allow visual inspection of the internal detail of the map and identification of artifacts.

Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

### 6.1 Orthogonal projections [i](#)

#### 6.1.1 Primary map



X

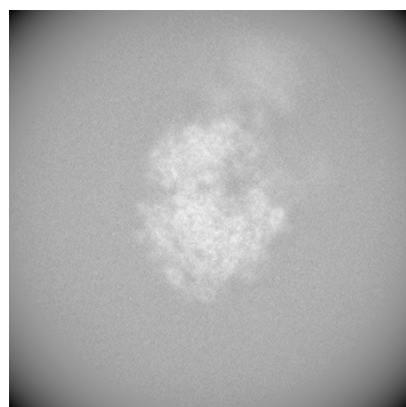


Y

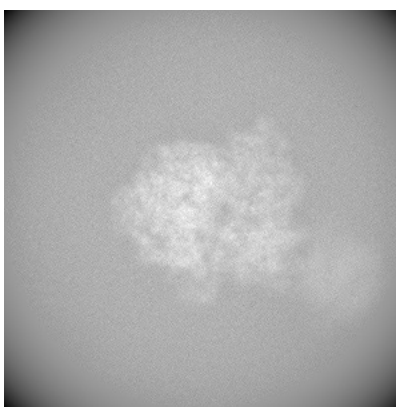


Z

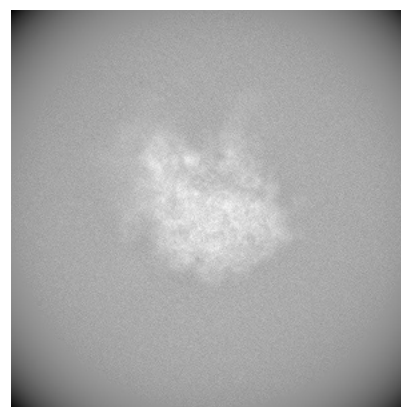
#### 6.1.2 Raw map



X



Y

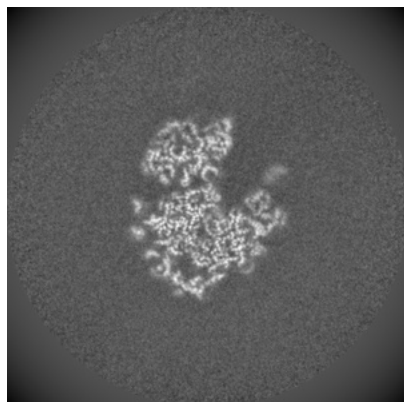


Z

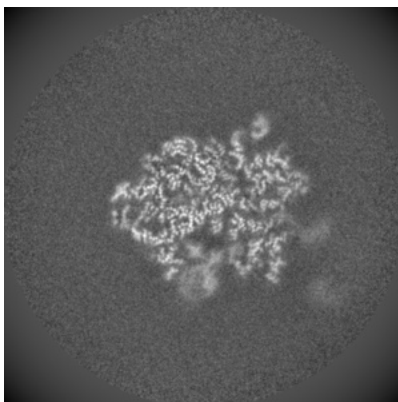
The images above show the map projected in three orthogonal directions.

## 6.2 Central slices [i](#)

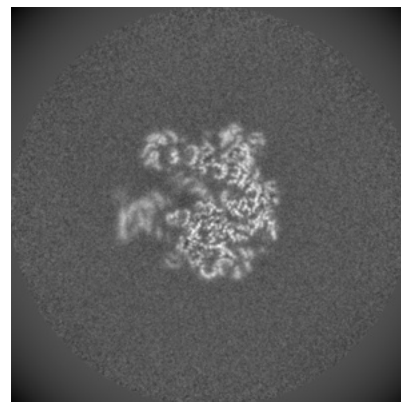
### 6.2.1 Primary map



X Index: 310

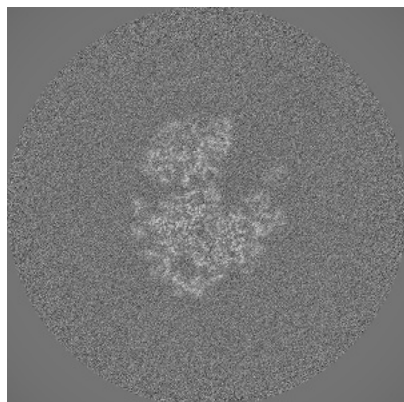


Y Index: 310

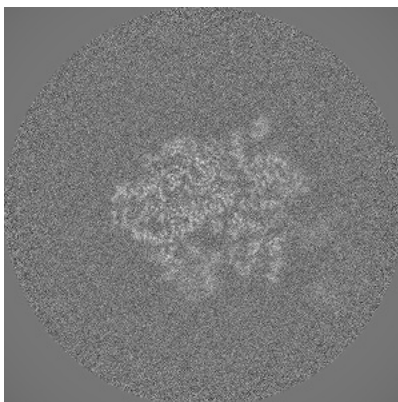


Z Index: 310

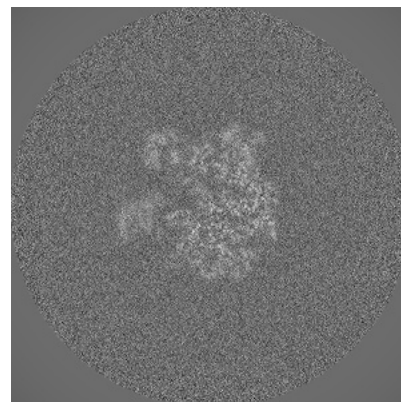
### 6.2.2 Raw map



X Index: 310



Y Index: 310



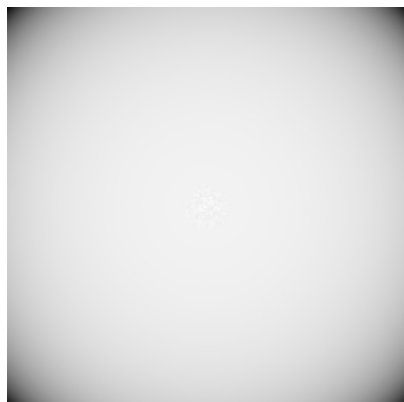
Z Index: 310

The images above show central slices of the map in three orthogonal directions.

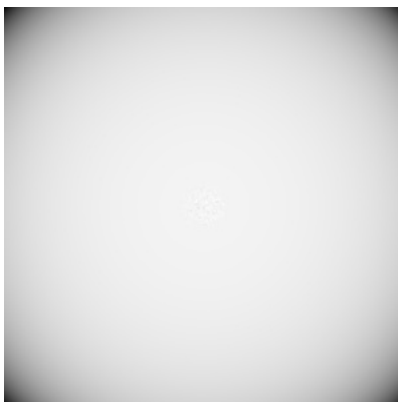


## 6.3 Largest variance slices [i](#)

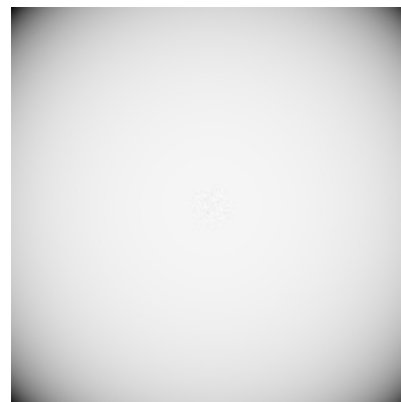
### 6.3.1 Primary map



X Index: 0

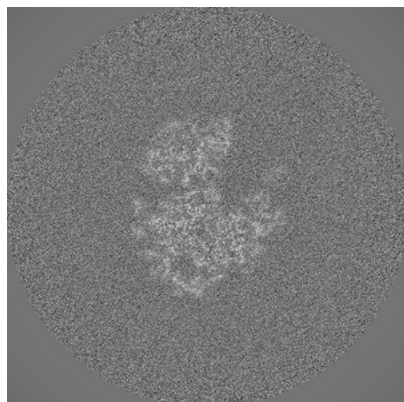


Y Index: 0

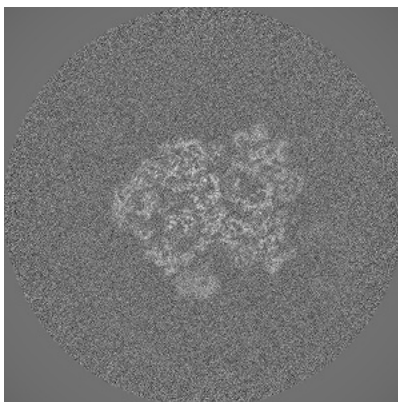


Z Index: 0

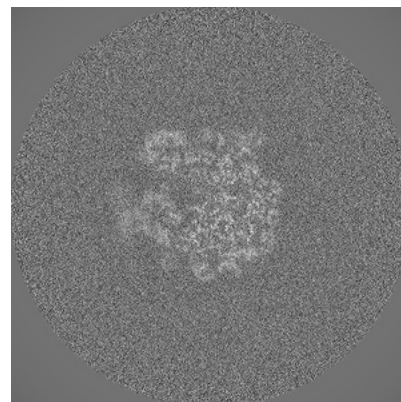
### 6.3.2 Raw map



X Index: 310



Y Index: 299

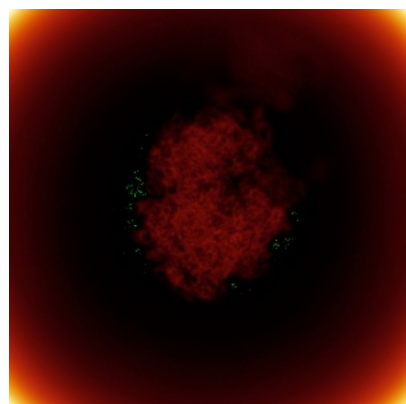


Z Index: 302

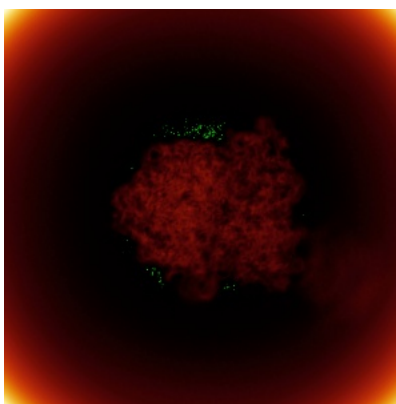
The images above show the largest variance slices of the map in three orthogonal directions.

## 6.4 Orthogonal standard-deviation projections (False-color) [i](#)

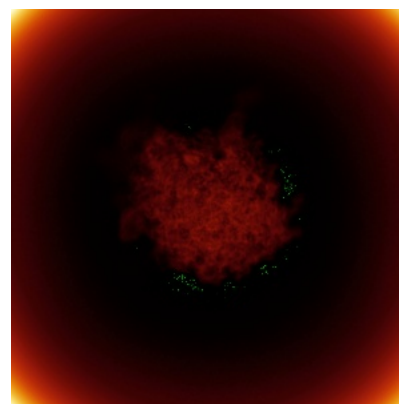
### 6.4.1 Primary map



X

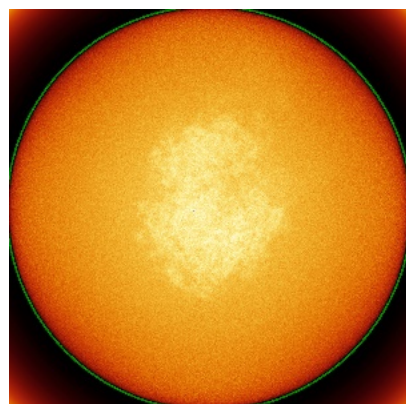


Y

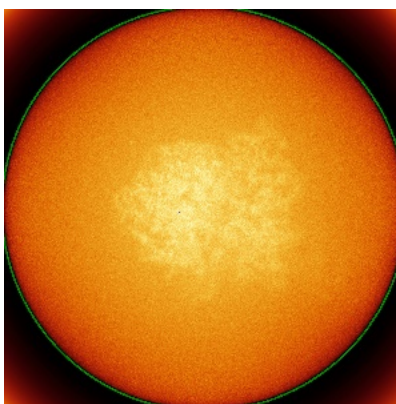


Z

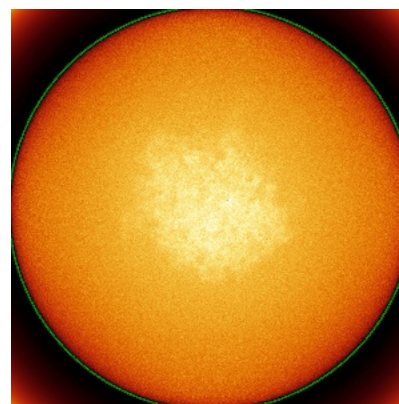
### 6.4.2 Raw map



X



Y

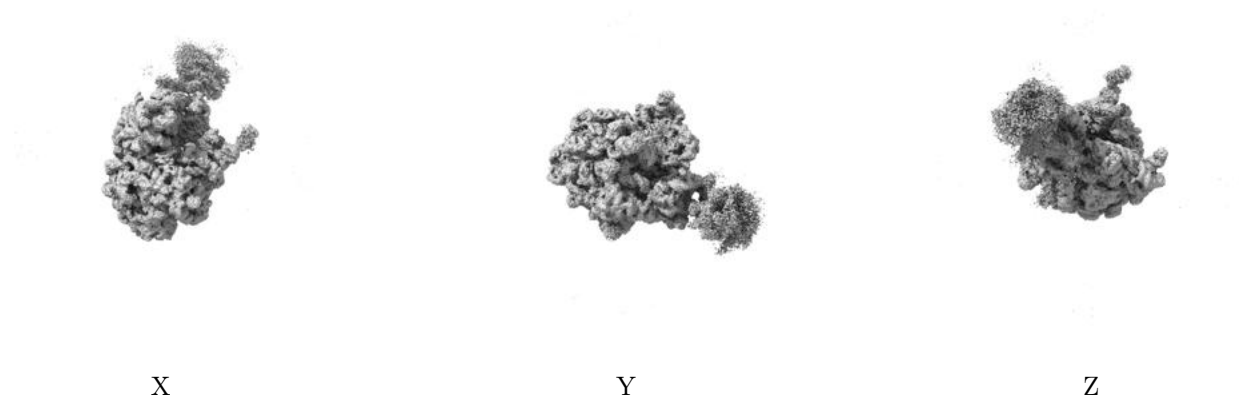


Z

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.

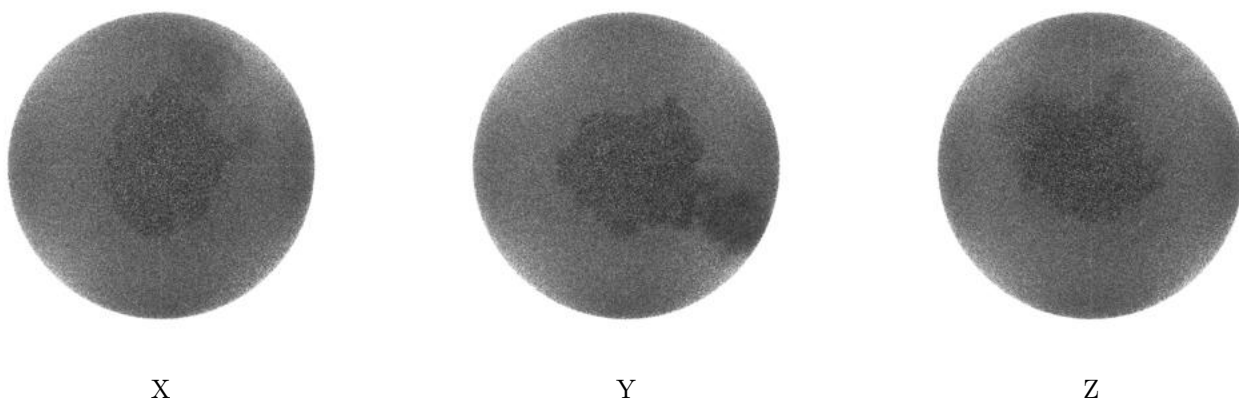
## 6.5 Orthogonal surface views [i](#)

### 6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 1e-06. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

### 6.5.2 Raw map



These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

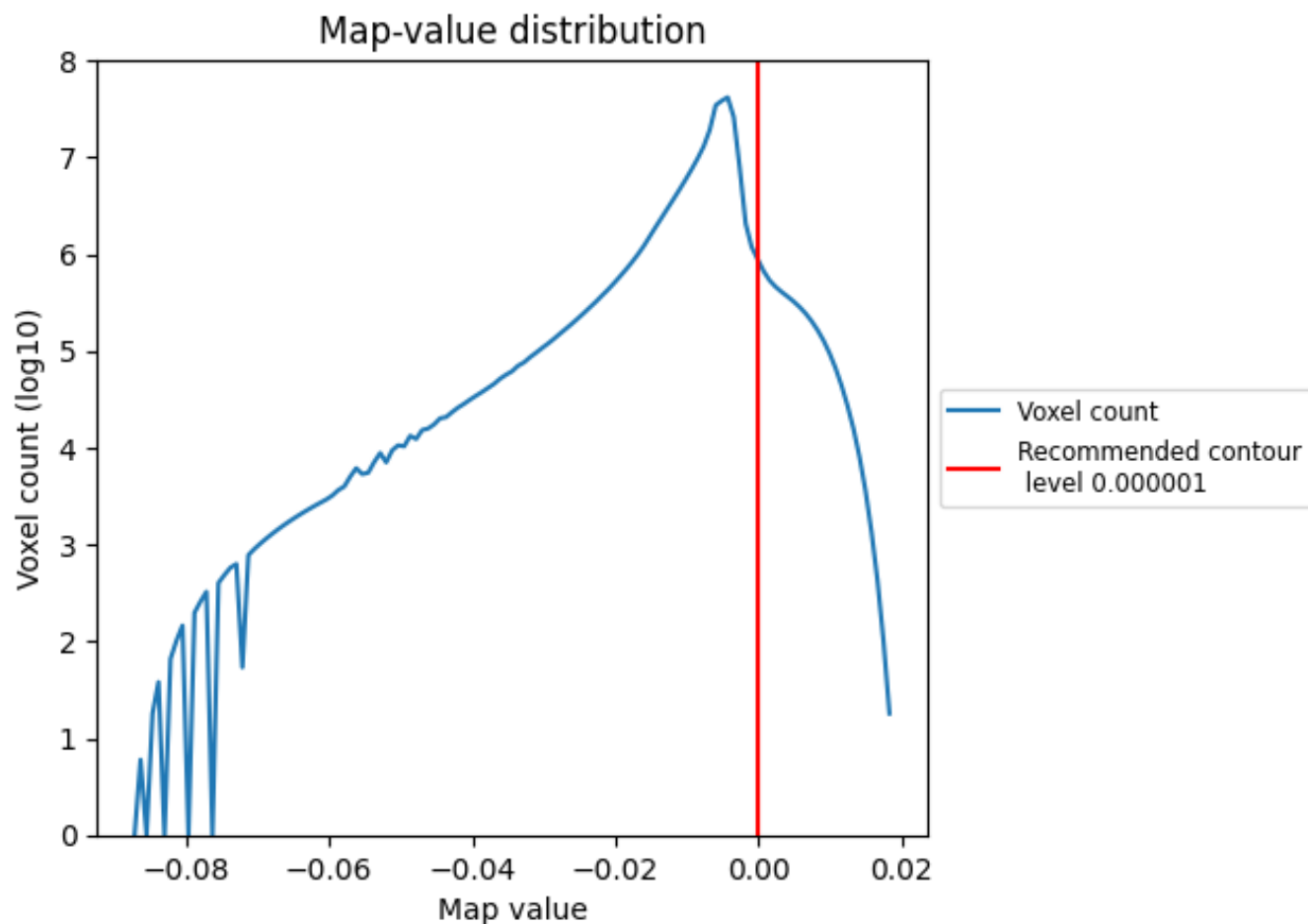
## 6.6 Mask visualisation [i](#)

This section was not generated. No masks/segmentation were deposited.

## 7 Map analysis [i](#)

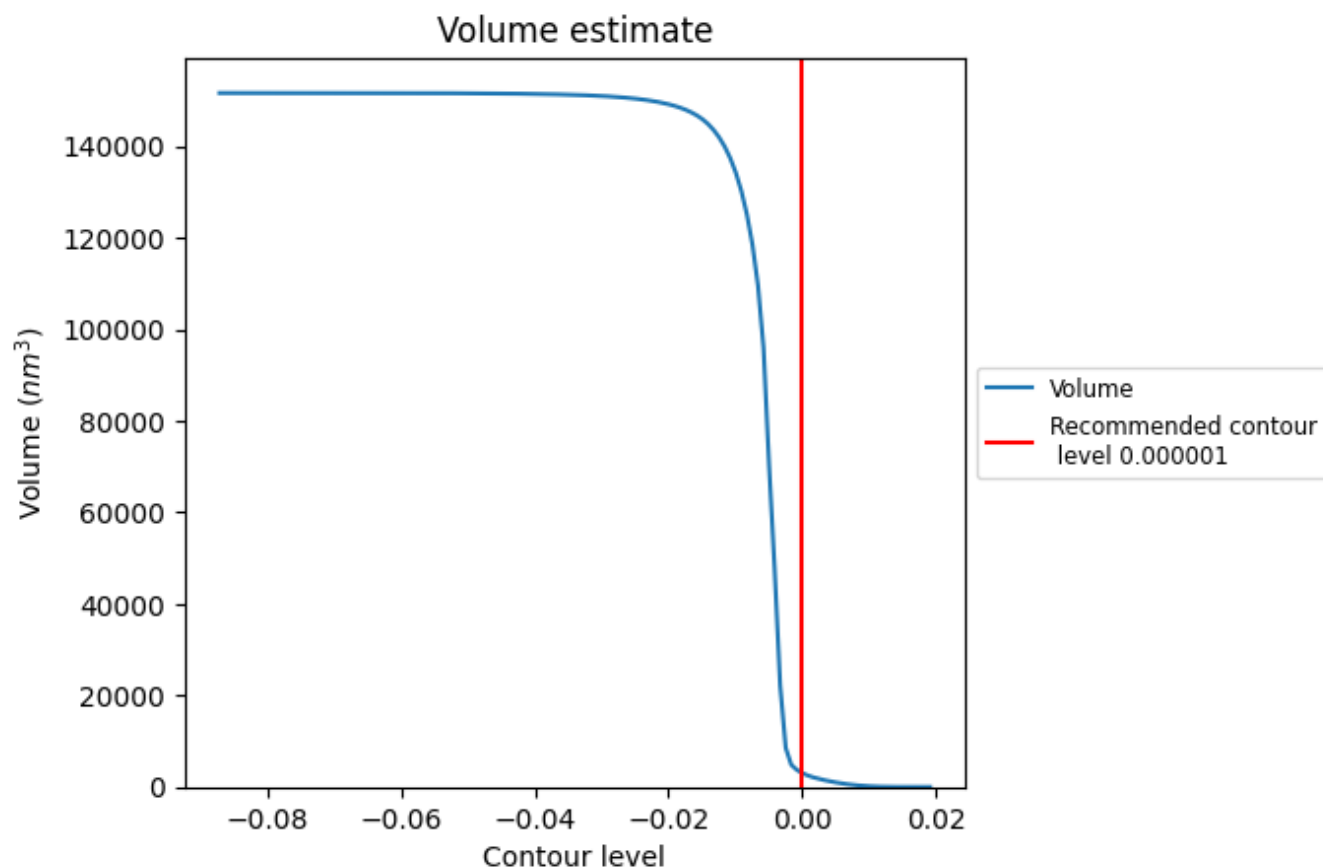
This section contains the results of statistical analysis of the map.

### 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.

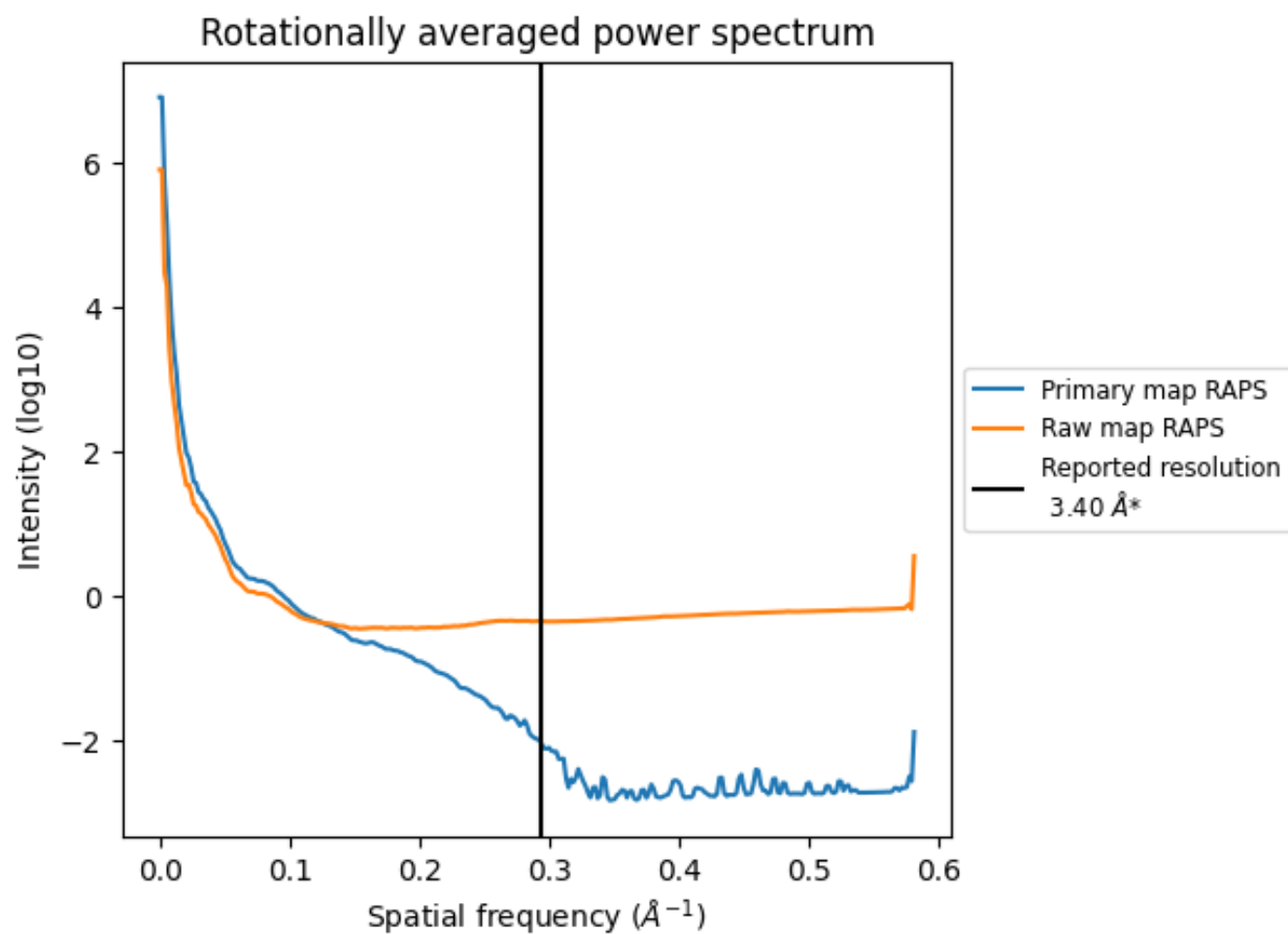
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 3055 nm<sup>3</sup>; this corresponds to an approximate mass of 2760 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 [i](#)

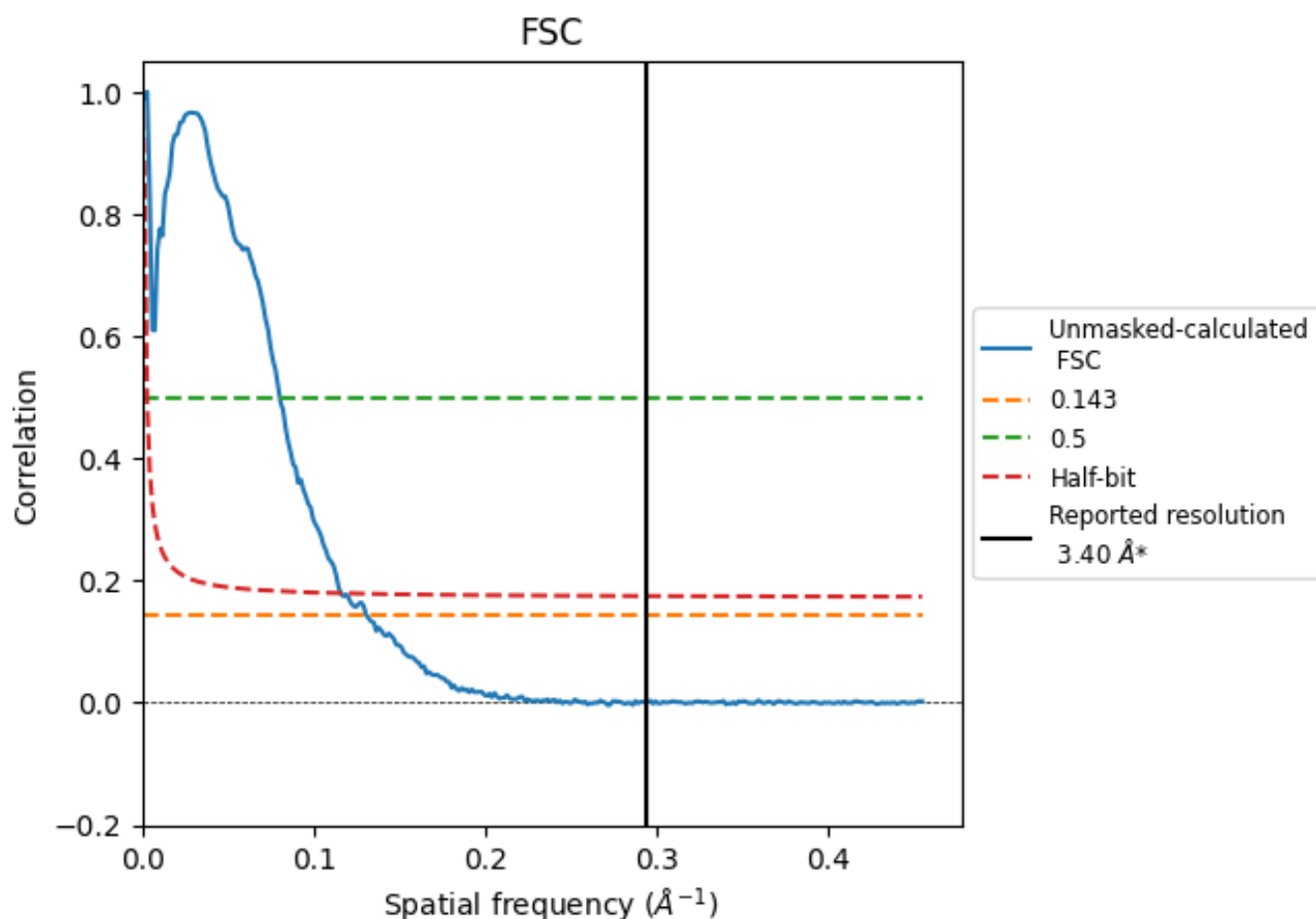


\*Reported resolution corresponds to spatial frequency of 0.294 Å<sup>-1</sup>

## 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 0.294 Å<sup>-1</sup>

## 8.2 Resolution estimates [i](#)

Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.40	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	7.66	12.47	8.64

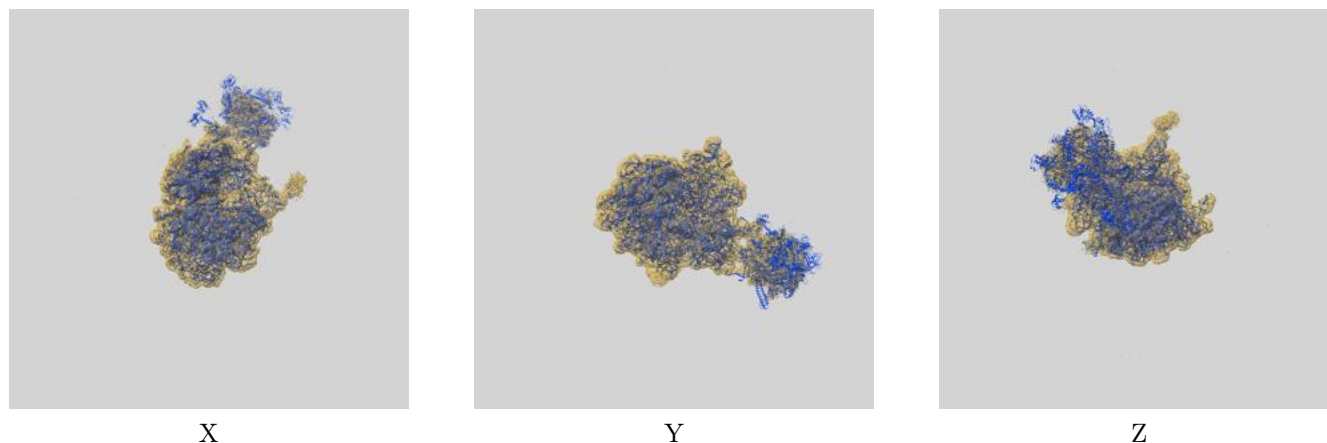
\*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 7.66 differs from the reported value 3.4 by more than 10 %



## 9 Map-model fit [i](#)

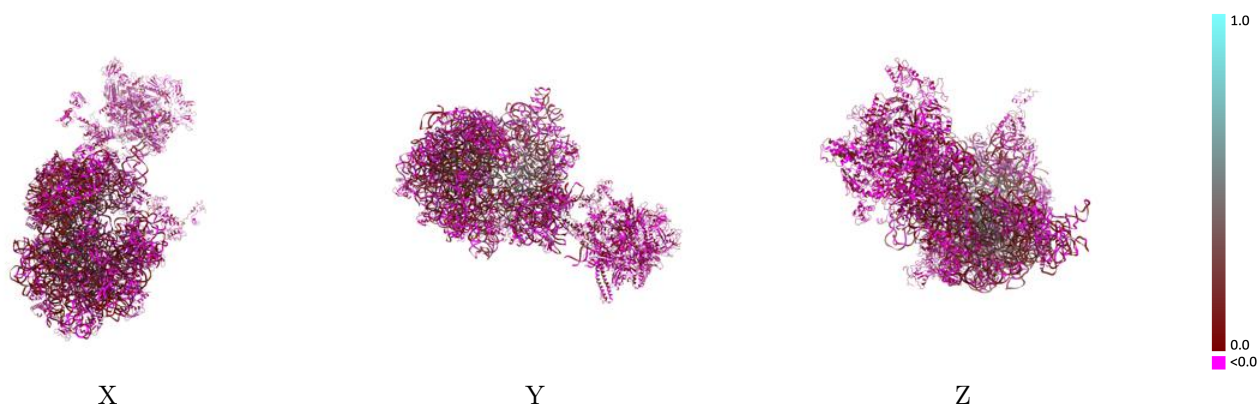
This section contains information regarding the fit between EMDB map EMD-43387 and PDB model 8VOO. Per-residue inclusion information can be found in section [3](#) on page [17](#).

### 9.1 Map-model overlay [i](#)



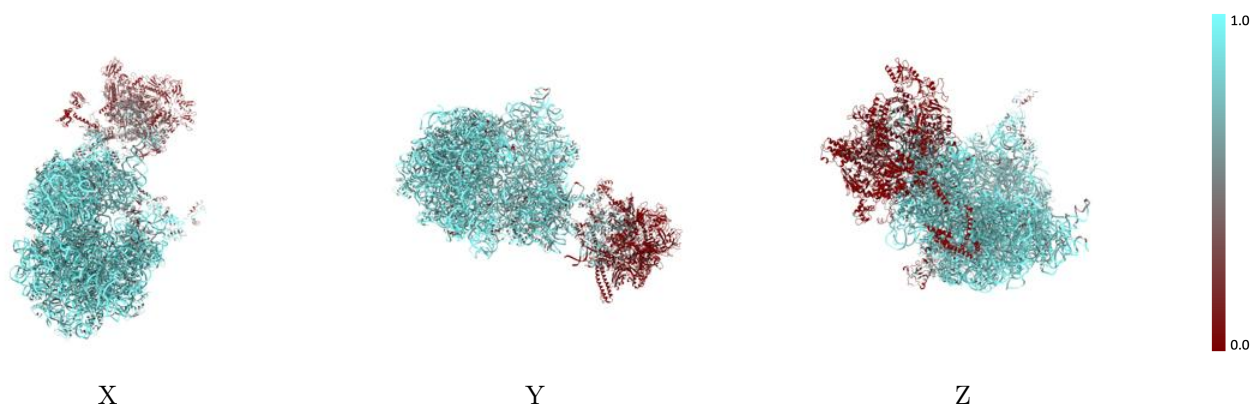
The images above show the 3D surface view of the map at the recommended contour level 1e-06 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

## 9.2 Q-score mapped to coordinate model [i](#)



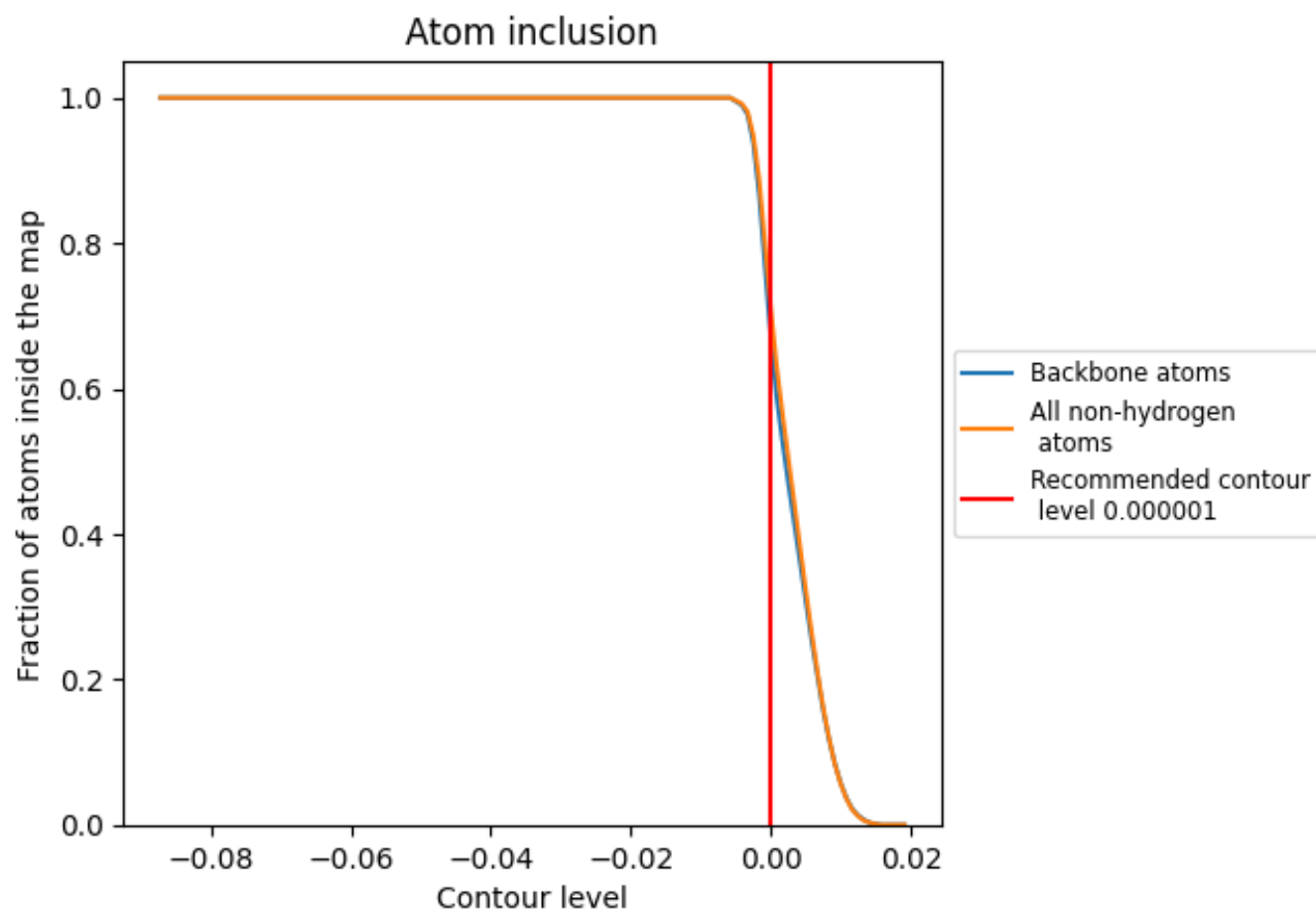
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.000001).







































































## 9.4 Atom inclusion [i](#)



At the recommended contour level, 67% of all backbone atoms, 72% 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 (0.000001) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.7150	 0.0820
0	 0.8140	 0.0300
1	 0.8300	 0.1110
2	 0.8040	 0.0360
3	 0.7890	 0.0030
4	 0.8460	 0.0400
5	 0.4390	 0.0320
6	 0.3940	 0.0500
7	 0.6240	 0.0340
9	 0.7630	 0.0470
A	 0.9100	 0.1480
AA	 0.1830	 0.0160
AB	 0.5300	 0.0830
AC	 0.0480	 0.0050
AD	 0.0110	 -0.0050
AE	 0.2190	 0.0260
AF	 0.0820	 0.0210
AG	 0.2940	 0.0160
B	 0.7960	 0.0470
C	 0.8180	 0.0220
D	 0.9290	 0.1250
E	 0.8100	 0.0310
F	 0.8180	 0.1220
G	 0.7390	 0.0360
H	 0.3640	 0.0350
I	 0.7120	 0.0100
J	 0.7680	 0.0140
K	 0.8580	 0.0970
L	 0.8280	 0.0290
M	 0.7650	 0.0540
N	 0.8560	 0.0890
O	 0.7650	 0.0250
P	 0.7430	 0.0550
Q	 0.8440	 0.0920
R	 0.8880	 0.1990



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Chain	Atom inclusion	Q-score
S	 0.7890	 0.0130
T	 0.8870	 0.1380
U	 0.7800	 0.0030
V	 0.8730	 0.1410
W	 0.6860	 0.0220
X	 0.7120	 0.0150
Y	 0.7340	 0.0180
Z	 0.5150	 0.0100
a	 0.9330	 0.1160
b	 0.8290	 0.0400
c	 0.8150	 0.0890
d	 0.8920	 0.0560
e	 0.8040	 0.0220
f	 0.8780	 0.0990
g	 0.7690	 0.0300
h	 0.8340	 0.1030
i	 0.8390	 0.1070
j	 0.8140	 0.0290
k	 0.8250	 0.0300
l	 0.7980	 0.0560
m	 0.9240	 0.2220
n	 0.7950	 0.0350
o	 0.8000	 0.0480
p	 0.7680	 -0.0060
q	 0.7940	 0.0120
r	 0.5770	 0.0270
s	 0.8040	 0.0090
t	 0.8240	 0.1020
u	 0.8160	 0.0440
v	 0.8590	 0.0840
w	 0.8600	 0.0660
x	 0.7870	 0.0210
y	 0.7620	 0.0360
z	 0.8290	 0.0470