



## Full wwPDB EM Validation Report ⓘ

Nov 10, 2024 – 10:01 am GMT

PDB ID : 7P3Z  
EMDB ID : EMD-13189  
Title : Homology model of the full-length AP-3 complex in a stretched open conformation  
Authors : Schubert, E.; Raunser, S.  
Deposited on : 2021-07-09  
Resolution : 10.50 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.dev113  
MolProbity : 4.02b-467  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
MapQ : 1.9.13  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.39

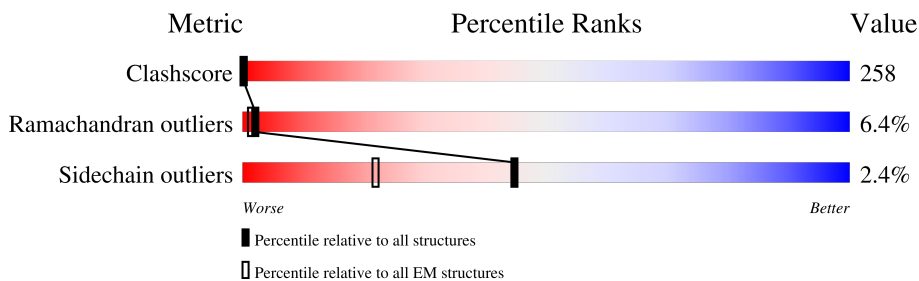
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*ELECTRON MICROSCOPY*

The reported resolution of this entry is 10.50 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

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	A	964	
2	B	809	
3	M	483	
4	S	194	

## 2 Entry composition

There are 4 unique types of molecules in this entry. The entry contains 14041 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called AP-3 complex subunit delta.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	576	Total	C	N	O	S	0	0
			4625	2978	738	881	28		

There are 32 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	933	ARG	-	expression tag	UNP A0A7I9C4X2
A	934	THR	-	expression tag	UNP A0A7I9C4X2
A	935	LEU	-	expression tag	UNP A0A7I9C4X2
A	936	GLN	-	expression tag	UNP A0A7I9C4X2
A	937	VAL	-	expression tag	UNP A0A7I9C4X2
A	938	ASP	-	expression tag	UNP A0A7I9C4X2
A	939	GLY	-	expression tag	UNP A0A7I9C4X2
A	940	SER	-	expression tag	UNP A0A7I9C4X2
A	941	ASP	-	expression tag	UNP A0A7I9C4X2
A	942	TYR	-	expression tag	UNP A0A7I9C4X2
A	943	LYS	-	expression tag	UNP A0A7I9C4X2
A	944	ASP	-	expression tag	UNP A0A7I9C4X2
A	945	ASP	-	expression tag	UNP A0A7I9C4X2
A	946	ASP	-	expression tag	UNP A0A7I9C4X2
A	947	ASP	-	expression tag	UNP A0A7I9C4X2
A	948	LYS	-	expression tag	UNP A0A7I9C4X2
A	949	ASP	-	expression tag	UNP A0A7I9C4X2
A	950	TYR	-	expression tag	UNP A0A7I9C4X2
A	951	LYS	-	expression tag	UNP A0A7I9C4X2
A	952	ASP	-	expression tag	UNP A0A7I9C4X2
A	953	ASP	-	expression tag	UNP A0A7I9C4X2
A	954	ASP	-	expression tag	UNP A0A7I9C4X2
A	955	ASP	-	expression tag	UNP A0A7I9C4X2
A	956	LYS	-	expression tag	UNP A0A7I9C4X2
A	957	ASP	-	expression tag	UNP A0A7I9C4X2
A	958	TYR	-	expression tag	UNP A0A7I9C4X2
A	959	LYS	-	expression tag	UNP A0A7I9C4X2
A	960	ASP	-	expression tag	UNP A0A7I9C4X2

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Chain	Residue	Modelled	Actual	Comment	Reference
A	961	ASP	-	expression tag	UNP A0A7I9C4X2
A	962	ASP	-	expression tag	UNP A0A7I9C4X2
A	963	ASP	-	expression tag	UNP A0A7I9C4X2
A	964	LYS	-	expression tag	UNP A0A7I9C4X2

- Molecule 2 is a protein called Y55\_G0035830.mRNA.1.CDS.1.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	B	621	Total	C	N	O	S	0	0
			4954	3160	830	936	28		

- Molecule 3 is a protein called AP-3 complex subunit mu.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	M	391	Total	C	N	O	S	0	0
			3106	1986	509	599	12		

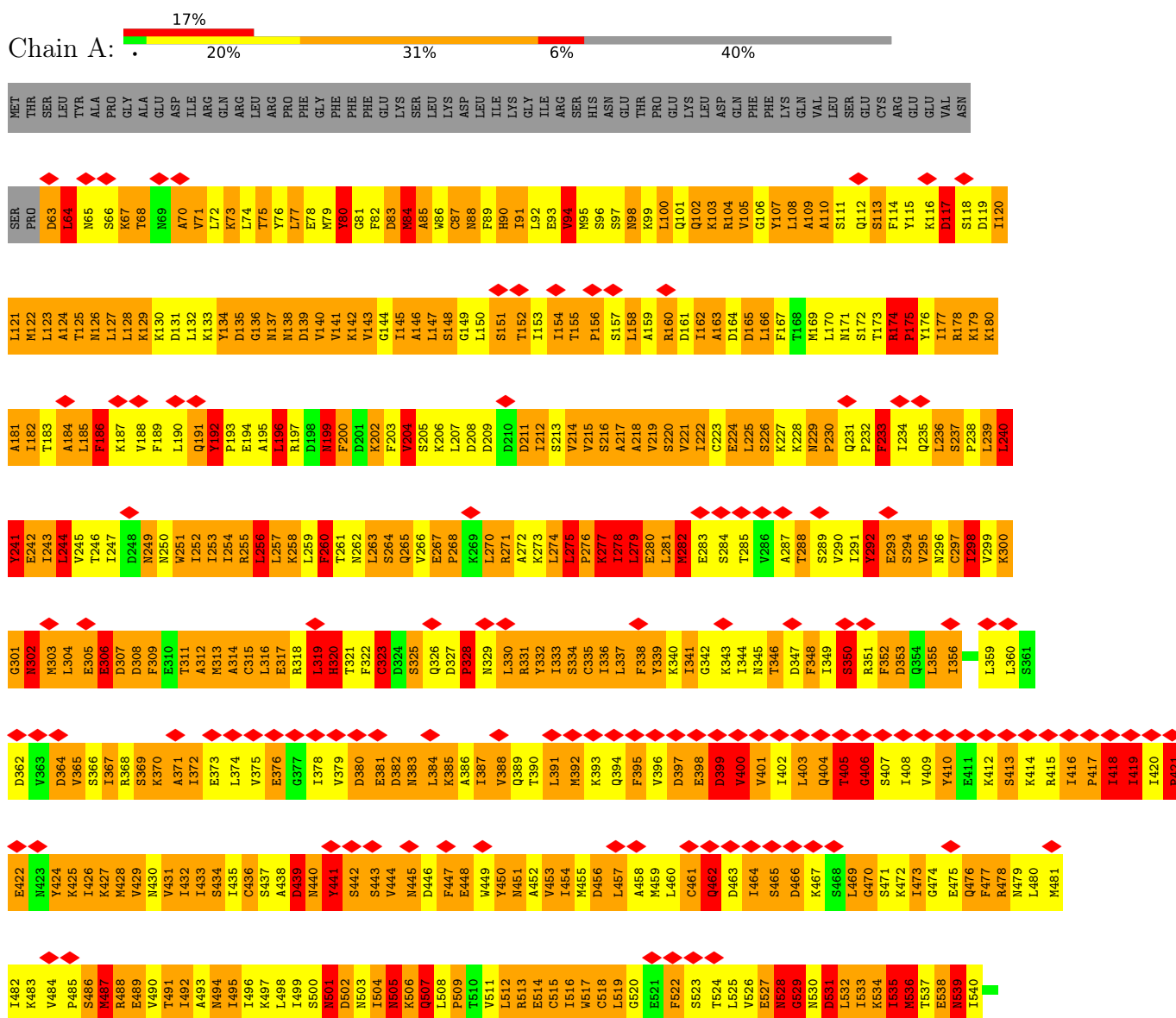
- Molecule 4 is a protein called AP complex subunit sigma.

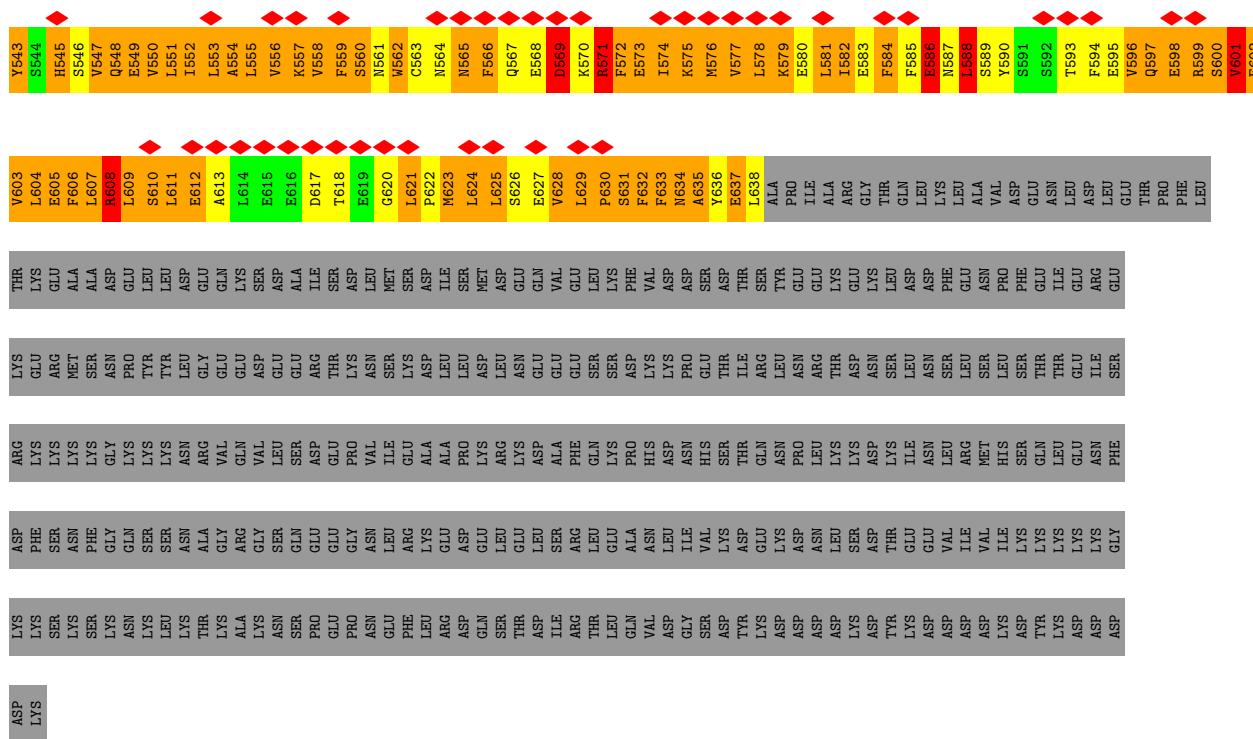
Mol	Chain	Residues	Atoms					AltConf	Trace
4	S	168	Total	C	N	O	S	0	0
			1356	867	215	270	4		

### 3 Residue-property plots

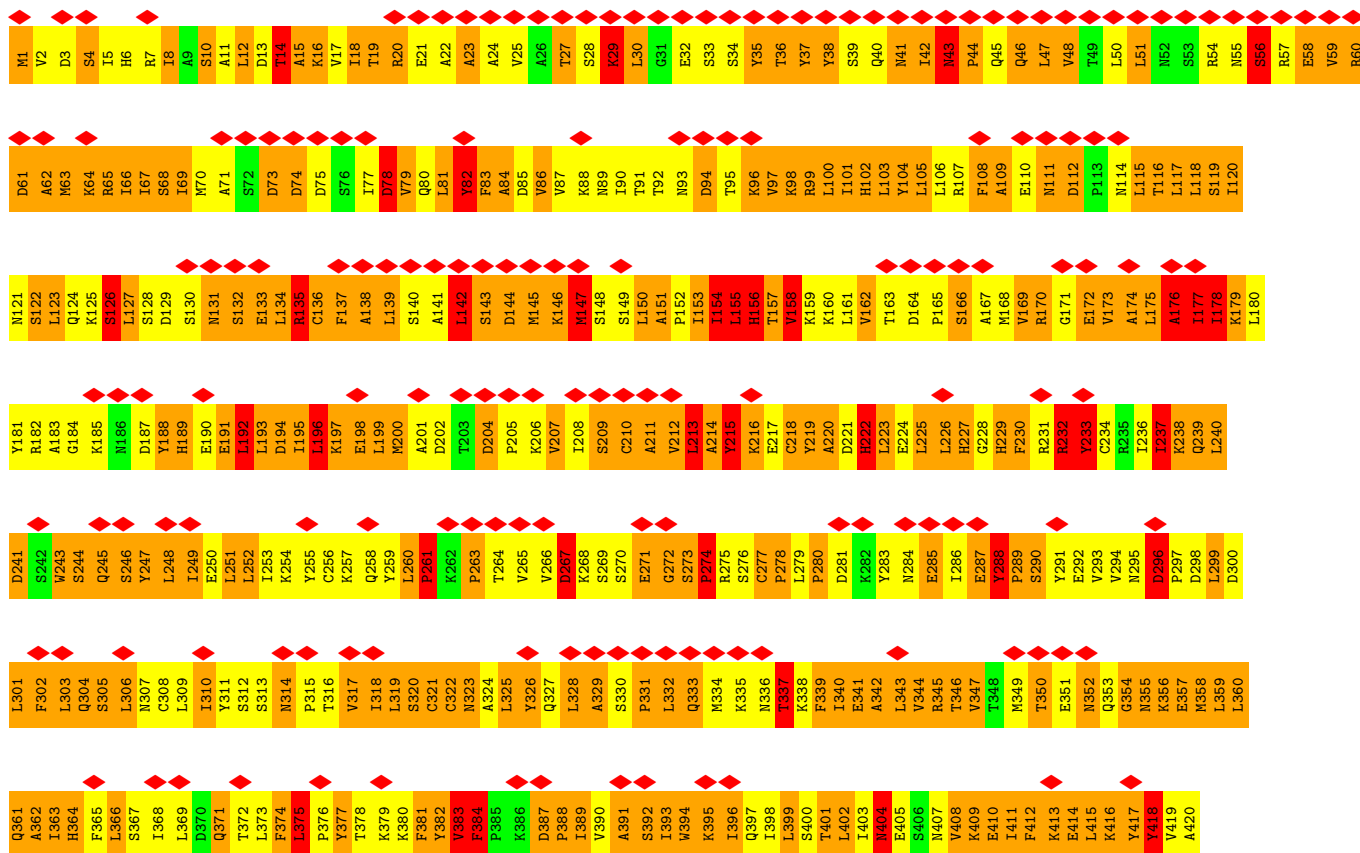
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion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

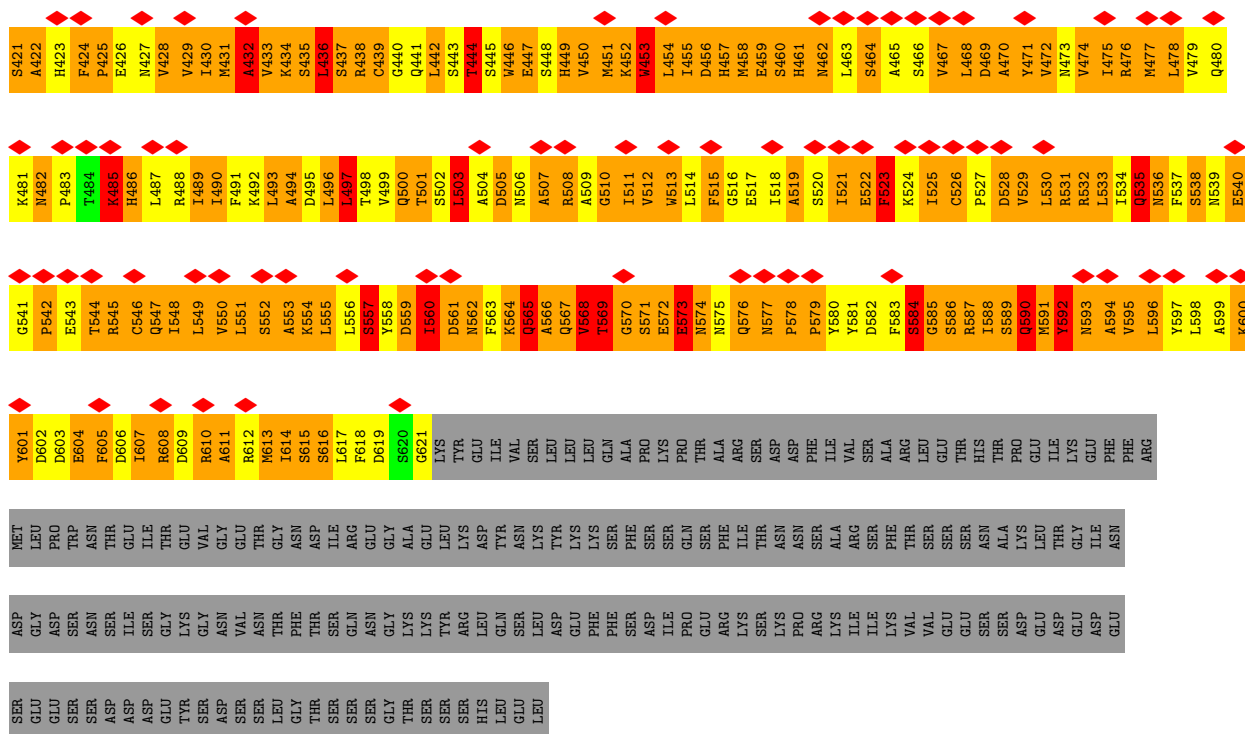
#### • Molecule 1: AP-3 complex subunit delta



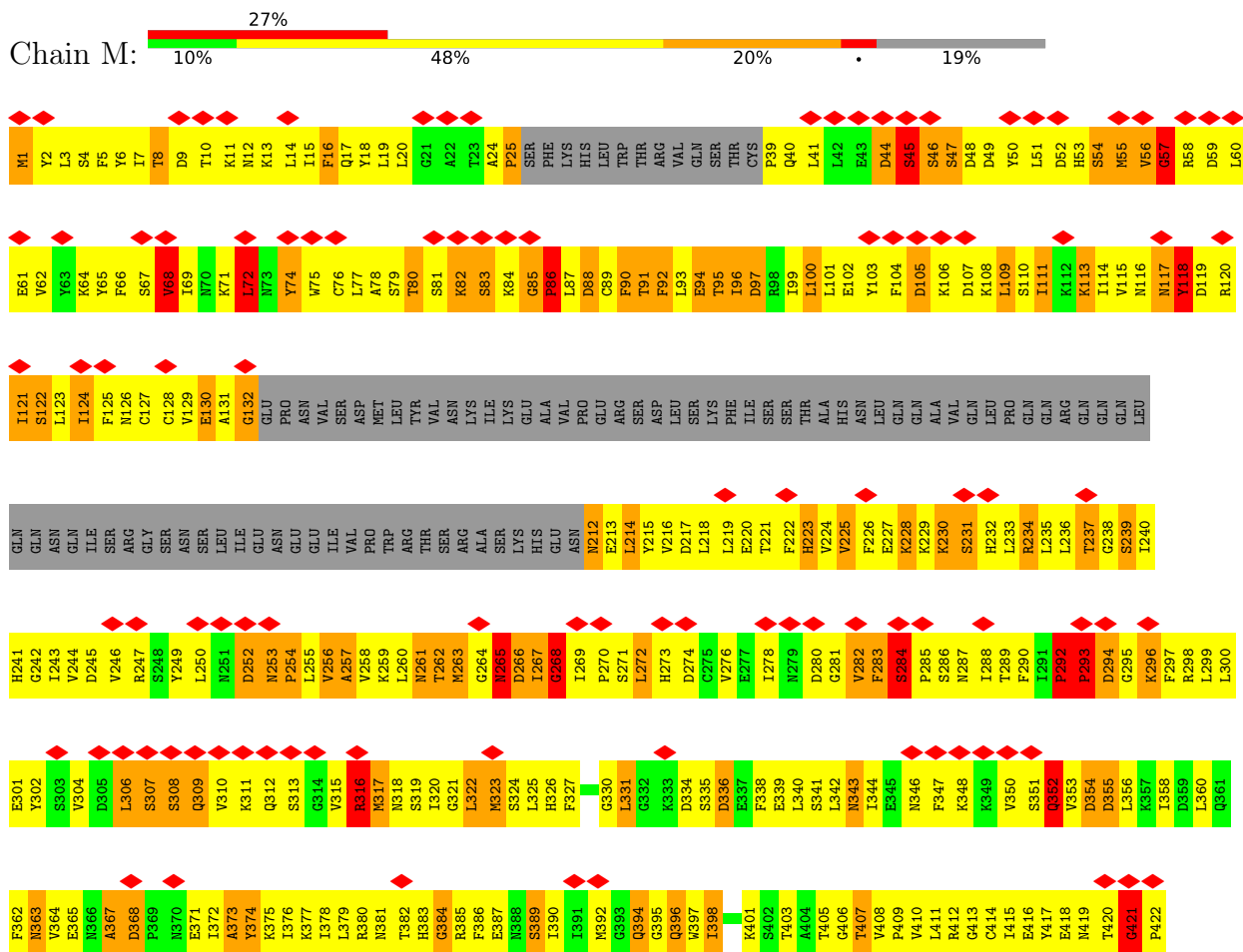


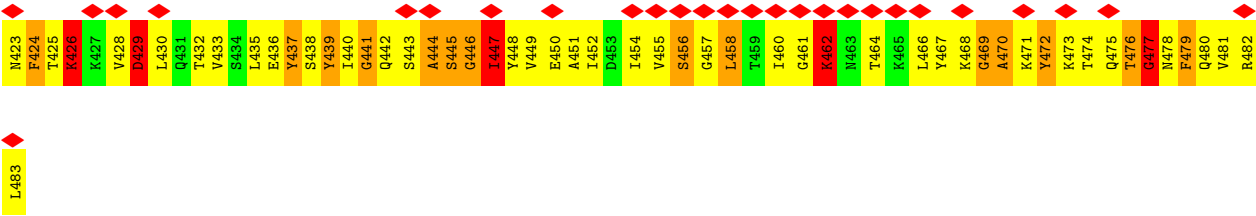
• Molecule 2: Y55\_G0035830.mRNA.1.CDS.1



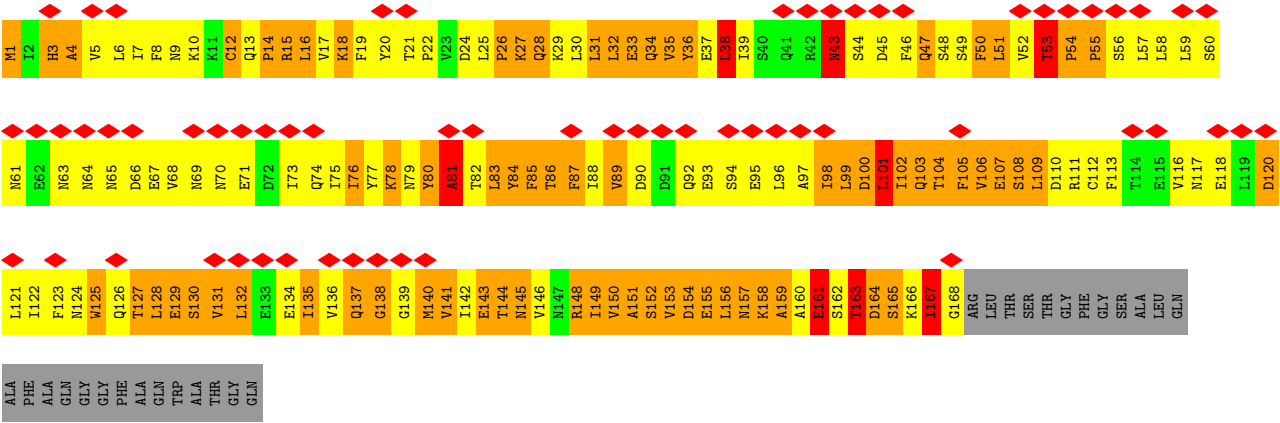


• Molecule 3: AP-3 complex subunit mu





• Molecule 4: AP complex subunit sigma



## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	20312	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING ONLY	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	81	Depositor
Minimum defocus (nm)	1500	Depositor
Maximum defocus (nm)	3600	Depositor
Magnification	130000	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor
Maximum map value	0.046	Depositor
Minimum map value	-0.015	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.002	Depositor
Recommended contour level	0.0155	Depositor
Map size ( $\text{\AA}$ )	282.48, 282.48, 282.48	wwPDB
Map dimensions	264, 264, 264	wwPDB
Map angles ( $^\circ$ )	90.0, 90.0, 90.0	wwPDB
Pixel spacing ( $\text{\AA}$ )	1.07, 1.07, 1.07	Depositor

## 5 Model quality ⓘ

### 5.1 Standard geometry ⓘ

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
1	A	1.68	70/4699 (1.5%)	2.61	607/6358 (9.5%)
2	B	1.52	52/5047 (1.0%)	2.32	540/6841 (7.9%)
3	M	1.59	61/3163 (1.9%)	1.85	139/4271 (3.3%)
4	S	1.84	33/1377 (2.4%)	2.17	116/1872 (6.2%)
All	All	1.62	216/14286 (1.5%)	2.31	1402/19342 (7.2%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	49
2	B	0	28
3	M	0	14
4	S	0	8
All	All	0	99

All (216) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	M	132	GLY	CA-C	-17.19	1.24	1.51
4	S	53	THR	N-CA	-14.56	1.17	1.46
3	M	132	GLY	N-CA	-14.40	1.24	1.46
1	A	406	GLY	CA-C	13.34	1.73	1.51
3	M	293	PRO	N-CD	12.51	1.65	1.47
4	S	163	THR	CA-C	11.13	1.81	1.52
4	S	46	PHE	CA-C	-11.05	1.24	1.52
4	S	55	PRO	N-CA	10.71	1.65	1.47
3	M	56	VAL	CA-C	-10.60	1.25	1.52
1	A	218	ALA	CA-C	-10.18	1.26	1.52
1	A	139	ASP	N-CA	-9.65	1.27	1.46
1	A	407	SER	N-CA	9.46	1.65	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
4	S	51	LEU	N-CA	-9.35	1.27	1.46
1	A	175	PRO	N-CD	9.26	1.60	1.47
3	M	41	LEU	CA-C	-9.12	1.29	1.52
2	B	337	THR	N-CA	-9.11	1.28	1.46
3	M	295	GLY	CA-C	9.02	1.66	1.51
4	S	54	PRO	N-CA	8.98	1.62	1.47
4	S	144	THR	N-CA	-8.82	1.28	1.46
3	M	231	SER	N-CA	8.77	1.63	1.46
4	S	4	ALA	CA-CB	8.56	1.70	1.52
1	A	467	LYS	N-CA	-8.49	1.29	1.46
2	B	289	PRO	CA-C	-8.40	1.36	1.52
1	A	394	GLN	CA-C	-8.30	1.31	1.52
1	A	405	THR	CA-C	-8.16	1.31	1.52
3	M	225	VAL	CA-C	-8.04	1.32	1.52
3	M	335	SER	CA-C	-8.04	1.32	1.52
2	B	212	VAL	C-O	8.02	1.38	1.23
4	S	83	LEU	CA-C	7.92	1.73	1.52
3	M	354	ASP	CA-C	7.83	1.73	1.52
4	S	140	MET	N-CA	-7.76	1.30	1.46
1	A	87	CYS	N-CA	-7.71	1.30	1.46
3	M	294	ASP	N-CA	-7.71	1.30	1.46
3	M	283	PHE	N-CA	-7.54	1.31	1.46
4	S	55	PRO	CA-C	-7.50	1.37	1.52
3	M	45	SER	CA-C	7.49	1.72	1.52
1	A	406	GLY	C-N	7.45	1.51	1.34
1	A	217	ALA	C-N	7.39	1.51	1.34
3	M	239	SER	N-CA	-7.35	1.31	1.46
3	M	109	LEU	N-CA	-7.32	1.31	1.46
3	M	44	ASP	C-N	7.23	1.50	1.34
4	S	167	ILE	C-N	7.19	1.46	1.33
3	M	293	PRO	CA-C	-7.19	1.38	1.52
3	M	441	GLY	CA-C	-7.16	1.40	1.51
3	M	95	THR	N-CA	7.10	1.60	1.46
1	A	529	GLY	N-CA	-7.03	1.35	1.46
2	B	330	SER	N-CA	-6.99	1.32	1.46
2	B	330	SER	C-N	-6.96	1.21	1.34
1	A	243	ILE	N-CA	-6.96	1.32	1.46
1	A	301	GLY	CA-C	-6.95	1.40	1.51
2	B	329	ALA	CA-C	-6.86	1.35	1.52
3	M	441	GLY	N-CA	-6.81	1.35	1.46
1	A	218	ALA	CA-CB	-6.77	1.38	1.52
1	A	230	PRO	N-CD	-6.74	1.38	1.47

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	M	55	MET	N-CA	-6.73	1.32	1.46
3	M	420	THR	N-CA	-6.69	1.32	1.46
3	M	55	MET	C-N	6.68	1.49	1.34
1	A	465	SER	N-CA	-6.67	1.33	1.46
3	M	230	LYS	CA-C	6.67	1.70	1.52
1	A	509	PRO	N-CD	6.65	1.57	1.47
3	M	322	LEU	C-O	6.61	1.35	1.23
3	M	392	MET	N-CA	-6.59	1.33	1.46
4	S	55	PRO	N-CD	6.53	1.56	1.47
4	S	108	SER	N-CA	-6.50	1.33	1.46
3	M	212	ASN	N-CA	-6.48	1.33	1.46
4	S	143	GLU	CA-C	-6.48	1.36	1.52
1	A	533	ILE	N-CA	-6.46	1.33	1.46
4	S	138	GLY	C-N	6.46	1.44	1.33
4	S	103	GLN	CA-C	-6.43	1.36	1.52
3	M	336	ASP	N-CA	-6.34	1.33	1.46
1	A	298	ILE	C-O	6.30	1.35	1.23
4	S	3	HIS	N-CA	-6.26	1.33	1.46
2	B	458	MET	C-O	6.25	1.35	1.23
2	B	329	ALA	N-CA	-6.23	1.33	1.46
1	A	410	TYR	N-CA	-6.21	1.33	1.46
4	S	81	ALA	CA-CB	-6.21	1.39	1.52
1	A	71	VAL	CA-C	-6.20	1.36	1.52
4	S	124	ASN	N-CA	-6.18	1.33	1.46
2	B	568	VAL	N-CA	-6.18	1.33	1.46
1	A	221	VAL	N-CA	6.16	1.58	1.46
1	A	566	PHE	CA-C	-6.15	1.36	1.52
1	A	394	GLN	N-CA	-6.14	1.34	1.46
2	B	585	GLY	N-CA	-6.11	1.36	1.46
1	A	278	ILE	N-CA	-6.10	1.34	1.46
1	A	464	ILE	CA-C	-6.10	1.37	1.52
2	B	172	GLU	CD-OE2	-6.02	1.19	1.25
3	M	56	VAL	C-N	-6.00	1.22	1.33
2	B	522	GLU	CA-C	-5.98	1.37	1.52
3	M	91	THR	CA-C	-5.95	1.37	1.52
1	A	418	ILE	C-N	5.93	1.47	1.34
1	A	281	LEU	N-CA	5.92	1.58	1.46
1	A	444	VAL	N-CA	-5.92	1.34	1.46
1	A	466	ASP	C-N	-5.91	1.20	1.34
1	A	534	LYS	CA-C	-5.91	1.37	1.52
1	A	277	LYS	C-N	-5.91	1.20	1.34
3	M	97	ASP	CA-C	-5.90	1.37	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	M	252	ASP	C-N	-5.90	1.20	1.34
3	M	257	ALA	N-CA	-5.90	1.34	1.46
3	M	262	THR	CA-C	-5.90	1.37	1.52
4	S	14	PRO	C-N	-5.87	1.20	1.34
3	M	446	GLY	CA-C	5.86	1.61	1.51
3	M	25	PRO	CA-C	-5.85	1.41	1.52
1	A	84	MET	N-CA	-5.84	1.34	1.46
3	M	263	MET	N-CA	-5.78	1.34	1.46
3	M	456	SER	C-N	5.78	1.43	1.33
1	A	302	ASN	CA-C	-5.77	1.38	1.52
3	M	83	SER	N-CA	-5.76	1.34	1.46
2	B	573	GLU	CD-OE2	-5.75	1.19	1.25
4	S	139	GLY	CA-C	-5.75	1.42	1.51
2	B	150	LEU	N-CA	-5.75	1.34	1.46
2	B	567	GLN	CA-C	-5.75	1.38	1.52
1	A	588	LEU	C-N	-5.73	1.20	1.34
2	B	576	GLN	C-N	-5.73	1.20	1.34
1	A	303	MET	N-CA	-5.72	1.34	1.46
1	A	241	TYR	N-CA	-5.71	1.34	1.46
1	A	508	LEU	N-CA	-5.70	1.34	1.46
4	S	95	GLU	CD-OE2	-5.70	1.19	1.25
2	B	290	SER	N-CA	-5.69	1.34	1.46
2	B	187	ASP	CA-C	-5.68	1.38	1.52
1	A	406	GLY	N-CA	-5.68	1.37	1.46
2	B	579	PRO	N-CD	-5.67	1.40	1.47
2	B	222	HIS	N-CA	5.65	1.57	1.46
3	M	96	ILE	C-N	5.61	1.47	1.34
3	M	113	LYS	CA-C	-5.61	1.38	1.52
1	A	387	ILE	CA-C	-5.61	1.38	1.52
2	B	501	THR	N-CA	-5.60	1.35	1.46
1	A	531	ASP	CA-C	-5.60	1.38	1.52
1	A	348	PHE	CA-C	-5.58	1.38	1.52
2	B	82	TYR	N-CA	-5.58	1.35	1.46
4	S	83	LEU	C-N	-5.58	1.21	1.34
1	A	405	THR	C-N	-5.57	1.23	1.33
2	B	526	CYS	N-CA	-5.57	1.35	1.46
4	S	158	LYS	CA-C	-5.55	1.38	1.52
3	M	296	LYS	CA-C	-5.54	1.38	1.52
2	B	219	TYR	C-N	-5.52	1.21	1.34
2	B	485	LYS	CA-C	-5.51	1.38	1.52
4	S	99	LEU	CA-C	-5.51	1.38	1.52
1	A	263	LEU	N-CA	-5.51	1.35	1.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	M	323	MET	N-CA	-5.50	1.35	1.46
4	S	80	TYR	N-CA	-5.50	1.35	1.46
4	S	83	LEU	N-CA	-5.50	1.35	1.46
2	B	584	SER	C-N	-5.49	1.23	1.33
4	S	84	TYR	C-N	-5.48	1.21	1.34
3	M	367	ALA	CA-CB	-5.47	1.41	1.52
3	M	25	PRO	N-CA	-5.44	1.38	1.47
4	S	144	THR	C-N	-5.43	1.21	1.34
1	A	373	GLU	CD-OE2	-5.42	1.19	1.25
3	M	284	SER	C-N	-5.42	1.24	1.34
3	M	457	GLY	CA-C	-5.40	1.43	1.51
2	B	574	ASN	N-CA	-5.39	1.35	1.46
2	B	601	TYR	N-CA	-5.39	1.35	1.46
3	M	354	ASP	C-N	-5.39	1.21	1.34
1	A	308	ASP	N-CA	-5.39	1.35	1.46
2	B	382	TYR	N-CA	-5.38	1.35	1.46
1	A	216	SER	C-N	5.38	1.46	1.34
1	A	83	ASP	CA-C	-5.37	1.39	1.52
4	S	143	GLU	C-N	-5.36	1.21	1.34
3	M	262	THR	N-CA	-5.34	1.35	1.46
1	A	224	GLU	CD-OE2	-5.33	1.19	1.25
1	A	392	MET	N-CA	-5.33	1.35	1.46
2	B	83	PHE	C-N	5.33	1.46	1.34
1	A	623	MET	CA-C	-5.33	1.39	1.52
3	M	223	HIS	C-N	-5.33	1.21	1.34
1	A	404	GLN	N-CA	-5.33	1.35	1.46
2	B	290	SER	CA-C	-5.32	1.39	1.52
2	B	383	VAL	CA-C	-5.32	1.39	1.52
1	A	192	TYR	CA-C	-5.32	1.39	1.52
1	A	535	ILE	N-CA	-5.30	1.35	1.46
2	B	190	GLU	CD-OE1	-5.30	1.19	1.25
2	B	500	GLN	C-N	-5.30	1.21	1.34
1	A	138	ASN	C-N	-5.29	1.21	1.34
2	B	331	PRO	N-CA	-5.28	1.38	1.47
2	B	425	PRO	CA-C	-5.28	1.42	1.52
1	A	464	ILE	C-N	-5.27	1.22	1.34
2	B	75	ASP	C-N	-5.27	1.22	1.34
2	B	219	TYR	CA-C	-5.26	1.39	1.52
2	B	407	ASN	N-CA	-5.25	1.35	1.46
2	B	443	SER	CA-C	-5.25	1.39	1.52
2	B	35	TYR	CA-C	-5.25	1.39	1.52
2	B	226	LEU	CA-C	-5.24	1.39	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	B	273	SER	N-CA	-5.24	1.35	1.46
1	A	388	VAL	CA-C	-5.23	1.39	1.52
2	B	133	GLU	CD-OE2	-5.22	1.20	1.25
2	B	384	PHE	N-CA	-5.21	1.35	1.46
3	M	444	ALA	CA-CB	5.21	1.63	1.52
2	B	424	PHE	N-CA	-5.21	1.35	1.46
2	B	109	ALA	CA-CB	-5.20	1.41	1.52
3	M	256	VAL	C-O	5.20	1.33	1.23
4	S	159	ALA	CA-CB	5.18	1.63	1.52
2	B	190	GLU	CD-OE2	-5.17	1.20	1.25
1	A	120	ILE	CA-C	-5.17	1.39	1.52
3	M	253	ASN	N-CA	-5.17	1.36	1.46
1	A	445	ASN	CA-C	-5.17	1.39	1.52
3	M	374	TYR	N-CA	-5.15	1.36	1.46
3	M	365	GLU	N-CA	-5.13	1.36	1.46
2	B	573	GLU	CD-OE1	-5.13	1.20	1.25
3	M	476	THR	C-N	-5.13	1.23	1.33
1	A	399	ASP	N-CA	-5.12	1.36	1.46
1	A	507	GLN	C-N	-5.11	1.22	1.34
4	S	29	LYS	CA-C	-5.11	1.39	1.52
3	M	41	LEU	C-N	5.10	1.45	1.34
1	A	224	GLU	CD-OE1	-5.10	1.20	1.25
3	M	228	LYS	C-N	-5.10	1.22	1.34
2	B	289	PRO	C-N	-5.10	1.22	1.34
3	M	238	GLY	CA-C	-5.10	1.43	1.51
2	B	500	GLN	CA-C	-5.09	1.39	1.52
3	M	373	ALA	C-N	-5.08	1.22	1.34
2	B	515	PHE	CA-C	5.06	1.66	1.52
3	M	227	GLU	CD-OE2	-5.06	1.20	1.25
1	A	85	ALA	CA-CB	-5.05	1.41	1.52
1	A	302	ASN	N-CA	-5.05	1.36	1.46
1	A	514	GLU	CD-OE1	-5.05	1.20	1.25
1	A	466	ASP	CA-C	-5.04	1.39	1.52
1	A	304	LEU	N-CA	-5.04	1.36	1.46
1	A	522	PHE	CA-C	-5.02	1.39	1.52
1	A	242	GLU	C-O	-5.02	1.13	1.23

All (1402) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	242	GLU	C-N-CA	22.76	178.60	121.70
1	A	265	GLN	N-CA-C	-19.23	59.09	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	80	TYR	N-CA-C	-18.37	61.39	111.00
3	M	45	SER	C-N-CA	-17.46	78.06	121.70
1	A	277	LYS	C-N-CA	17.23	164.78	121.70
3	M	354	ASP	N-CA-C	-16.54	66.34	111.00
3	M	354	ASP	C-N-CA	16.54	163.04	121.70
1	A	444	VAL	N-CA-C	-16.42	66.65	111.00
3	M	373	ALA	CB-CA-C	15.46	133.29	110.10
1	A	98	ASN	C-N-CA	15.35	160.08	121.70
2	B	584	SER	N-CA-C	15.03	151.59	111.00
1	A	304	LEU	C-N-CA	-14.99	84.22	121.70
2	B	263	PRO	C-N-CA	14.60	158.20	121.70
1	A	534	LYS	CA-C-N	14.58	149.28	117.20
2	B	290	SER	C-N-CA	-14.53	85.37	121.70
1	A	534	LYS	C-N-CA	14.19	157.16	121.70
3	M	294	ASP	N-CA-C	-13.94	73.35	111.00
2	B	600	LYS	C-N-CA	-13.51	87.92	121.70
1	A	532	LEU	C-N-CA	-13.43	88.14	121.70
1	A	80	TYR	O-C-N	-13.25	100.68	123.20
1	A	418	ILE	C-N-CA	-13.24	88.60	121.70
2	B	78	ASP	C-N-CA	-13.24	88.60	121.70
3	M	293	PRO	CA-N-CD	-12.75	93.65	111.50
1	A	98	ASN	N-CA-C	-12.73	76.62	111.00
1	A	464	ILE	C-N-CA	-12.72	89.90	121.70
4	S	168	GLY	N-CA-C	12.69	144.83	113.10
1	A	64	LEU	O-C-N	-12.60	102.54	122.70
1	A	302	ASN	N-CA-C	-12.60	76.99	111.00
1	A	260	PHE	O-C-N	-12.55	102.61	122.70
1	A	405	THR	C-N-CA	-12.50	96.04	122.30
1	A	536	MET	O-C-N	-12.30	103.02	122.70
1	A	151	SER	C-N-CA	-12.26	91.06	121.70
3	M	54	SER	N-CA-C	12.18	143.88	111.00
1	A	469	LEU	C-N-CA	-12.12	96.86	122.30
1	A	135	ASP	C-N-CA	-11.93	97.25	122.30
2	B	330	SER	N-CA-C	-11.78	79.19	111.00
1	A	233	PHE	C-N-CA	-11.76	92.31	121.70
1	A	323	CYS	C-N-CA	11.63	150.78	121.70
1	A	136	GLY	C-N-CA	-11.47	93.02	121.70
1	A	265	GLN	CA-C-N	11.46	142.42	117.20
1	A	536	MET	C-N-CA	-11.42	93.16	121.70
4	S	53	THR	C-N-CD	11.41	152.36	128.40
3	M	458	LEU	O-C-N	11.37	140.89	122.70
2	B	329	ALA	CB-CA-C	-11.32	93.12	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	404	ASN	C-N-CA	-11.29	93.48	121.70
3	M	462	LYS	N-CA-C	-11.24	80.64	111.00
1	A	465	SER	CA-C-N	-11.18	92.60	117.20
1	A	242	GLU	CA-C-N	11.16	141.76	117.20
1	A	534	LYS	N-CA-C	-11.16	80.85	111.00
3	M	46	SER	C-N-CA	-10.93	94.37	121.70
2	B	310	ILE	C-N-CA	-10.93	94.38	121.70
4	S	109	LEU	O-C-N	-10.91	105.24	122.70
1	A	416	ILE	C-N-CD	10.83	151.15	128.40
4	S	43	ASN	O-C-N	-10.74	105.51	122.70
2	B	497	LEU	O-C-N	-10.67	105.62	122.70
1	A	80	TYR	CA-C-N	10.67	137.54	116.20
4	S	163	THR	C-N-CA	10.67	148.37	121.70
4	S	54	PRO	CA-N-CD	-10.63	96.61	111.50
1	A	204	VAL	O-C-N	-10.58	105.78	122.70
1	A	586	GLU	C-N-CA	-10.53	95.37	121.70
2	B	82	TYR	C-N-CA	-10.53	95.37	121.70
1	A	244	LEU	O-C-N	-10.47	105.95	122.70
2	B	584	SER	C-N-CA	-10.44	100.38	122.30
1	A	629	LEU	C-N-CD	10.32	150.08	128.40
1	A	84	MET	C-N-CA	-10.31	95.92	121.70
4	S	164	ASP	C-N-CA	-10.30	95.94	121.70
1	A	519	LEU	C-N-CA	-10.28	100.72	122.30
1	A	621	LEU	C-N-CD	10.16	149.74	128.40
2	B	571	SER	N-CA-C	-10.09	83.77	111.00
1	A	569	ASP	C-N-CA	-10.04	96.61	121.70
3	M	83	SER	C-N-CA	10.02	146.75	121.70
2	B	289	PRO	C-N-CA	-9.94	96.84	121.70
2	B	162	VAL	C-N-CA	-9.91	96.93	121.70
1	A	84	MET	O-C-N	-9.90	106.86	122.70
1	A	320	HIS	O-C-N	-9.90	106.86	122.70
1	A	380	ASP	N-CA-C	-9.90	84.27	111.00
2	B	230	PHE	C-N-CA	-9.86	97.05	121.70
2	B	570	GLY	N-CA-C	-9.73	88.77	113.10
1	A	346	THR	C-N-CA	-9.72	97.39	121.70
1	A	586	GLU	O-C-N	-9.70	107.18	122.70
2	B	576	GLN	N-CA-C	-9.66	84.92	111.00
3	M	268	GLY	C-N-CA	-9.63	97.62	121.70
1	A	504	ILE	O-C-N	-9.63	107.30	122.70
3	M	57	GLY	C-N-CA	9.58	145.66	121.70
2	B	577	ASN	C-N-CD	9.56	148.48	128.40
1	A	462	GLN	O-C-N	-9.55	107.41	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	319	LEU	O-C-N	-9.53	107.45	122.70
2	B	212	VAL	CA-C-O	9.50	140.05	120.10
2	B	584	SER	N-CA-CB	-9.47	96.30	110.50
1	A	212	ILE	C-N-CA	-9.45	98.06	121.70
1	A	305	GLU	N-CA-C	9.45	136.51	111.00
1	A	242	GLU	N-CA-C	-9.43	85.53	111.00
2	B	102	HIS	O-C-N	-9.39	107.68	122.70
1	A	569	ASP	N-CA-C	-9.37	85.69	111.00
2	B	187	ASP	C-N-CA	-9.37	98.27	121.70
1	A	403	LEU	C-N-CA	-9.36	98.30	121.70
3	M	252	ASP	N-CA-C	-9.34	85.78	111.00
2	B	366	LEU	O-C-N	-9.33	107.77	122.70
2	B	577	ASN	N-CA-C	9.33	136.19	111.00
1	A	282	MET	O-C-N	-9.32	107.78	122.70
1	A	381	GLU	C-N-CA	-9.30	98.44	121.70
2	B	535	GLN	C-N-CA	-9.29	98.48	121.70
4	S	143	GLU	N-CA-C	-9.29	85.93	111.00
1	A	80	TYR	C-N-CA	9.28	141.79	122.30
2	B	560	ILE	C-N-CA	-9.24	98.59	121.70
3	M	306	LEU	O-C-N	-9.24	107.91	122.70
2	B	461	HIS	C-N-CA	-9.23	98.62	121.70
2	B	505	ASP	C-N-CA	-9.19	98.72	121.70
1	A	504	ILE	C-N-CA	-9.18	98.76	121.70
1	A	110	ALA	C-N-CA	-9.16	98.79	121.70
2	B	205	PRO	C-N-CA	-9.13	98.89	121.70
1	A	275	LEU	C-N-CD	9.12	147.55	128.40
1	A	365	VAL	C-N-CA	-9.11	98.92	121.70
1	A	163	ALA	C-N-CA	-9.08	99.00	121.70
1	A	534	LYS	CA-C-O	-9.07	101.04	120.10
3	M	41	LEU	O-C-N	9.07	137.21	122.70
2	B	211	ALA	O-C-N	9.05	137.18	122.70
1	A	465	SER	C-N-CA	9.01	144.21	121.70
1	A	103	LYS	O-C-N	-8.99	108.31	122.70
3	M	80	THR	N-CA-C	8.99	135.26	111.00
2	B	568	VAL	CA-C-N	-8.98	97.44	117.20
2	B	497	LEU	C-N-CA	-8.98	99.25	121.70
1	A	539	ASN	O-C-N	-8.88	108.49	122.70
1	A	242	GLU	O-C-N	-8.87	108.50	122.70
1	A	86	TRP	C-N-CA	-8.86	99.55	121.70
1	A	100	LEU	C-N-CA	-8.85	99.59	121.70
2	B	569	THR	N-CA-C	8.83	134.84	111.00
3	M	477	GLY	C-N-CA	-8.83	99.62	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	196	LEU	C-N-CA	-8.82	99.64	121.70
1	A	240	LEU	C-N-CA	-8.82	99.64	121.70
1	A	529	GLY	CA-C-O	-8.81	104.74	120.60
1	A	461	CYS	O-C-N	-8.81	108.61	122.70
2	B	559	ASP	CA-C-O	-8.80	101.62	120.10
1	A	350	SER	O-C-N	-8.79	108.64	122.70
2	B	212	VAL	O-C-N	-8.78	108.65	122.70
2	B	404	ASN	O-C-N	-8.78	108.66	122.70
1	A	545	HIS	C-N-CA	-8.77	99.79	121.70
4	S	104	THR	O-C-N	8.72	136.65	122.70
1	A	88	ASN	C-N-CA	-8.71	99.92	121.70
2	B	557	SER	O-C-N	-8.71	108.77	122.70
4	S	167	ILE	C-N-CA	-8.68	104.08	122.30
2	B	523	PHE	O-C-N	-8.62	108.91	122.70
1	A	154	ILE	N-CA-C	8.61	134.25	111.00
4	S	98	ILE	O-C-N	8.61	136.47	122.70
2	B	109	ALA	C-N-CA	-8.60	100.21	121.70
1	A	325	SER	N-CA-C	-8.56	87.87	111.00
2	B	223	LEU	C-N-CA	-8.53	100.39	121.70
3	M	92	PHE	O-C-N	8.52	136.34	122.70
1	A	413	SER	N-CA-C	-8.52	88.01	111.00
3	M	367	ALA	C-N-CA	8.51	142.97	121.70
4	S	81	ALA	CB-CA-C	8.51	122.86	110.10
4	S	100	ASP	O-C-N	8.51	136.31	122.70
1	A	233	PHE	O-C-N	-8.49	109.11	122.70
3	M	420	THR	C-N-CA	-8.48	104.49	122.30
1	A	367	ILE	O-C-N	8.47	136.25	122.70
2	B	339	PHE	O-C-N	8.44	136.21	122.70
1	A	461	CYS	CA-C-O	8.44	137.83	120.10
3	M	292	PRO	C-N-CA	8.44	157.45	122.00
1	A	432	ILE	O-C-N	8.43	136.19	122.70
1	A	328	PRO	C-N-CA	-8.43	100.64	121.70
1	A	281	LEU	CA-C-O	8.42	137.79	120.10
1	A	391	LEU	C-N-CA	-8.42	100.66	121.70
1	A	508	LEU	N-CA-C	8.42	133.72	111.00
2	B	147	MET	O-C-N	-8.41	109.24	122.70
2	B	146	LYS	C-N-CA	-8.41	100.67	121.70
1	A	450	TYR	O-C-N	8.39	136.13	122.70
4	S	103	GLN	O-C-N	8.38	136.10	122.70
2	B	574	ASN	C-N-CA	-8.36	100.79	121.70
1	A	215	VAL	CA-C-O	8.36	137.65	120.10
2	B	472	VAL	O-C-N	-8.33	109.38	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	120	ILE	O-C-N	8.32	136.02	122.70
4	S	46	PHE	CA-C-O	-8.31	102.64	120.10
1	A	424	TYR	O-C-N	8.31	135.99	122.70
1	A	94	VAL	O-C-N	-8.31	109.41	122.70
3	M	457	GLY	N-CA-C	-8.31	92.33	113.10
4	S	161	GLU	O-C-N	-8.30	109.42	122.70
2	B	108	PHE	O-C-N	-8.30	109.42	122.70
1	A	270	LEU	O-C-N	8.27	135.93	122.70
3	M	68	VAL	N-CA-C	8.27	133.34	111.00
3	M	335	SER	O-C-N	8.27	135.94	122.70
2	B	325	LEU	O-C-N	-8.27	109.47	122.70
3	M	97	ASP	O-C-N	8.27	135.92	122.70
2	B	444	THR	O-C-N	-8.26	109.48	122.70
2	B	366	LEU	C-N-CA	-8.25	101.08	121.70
1	A	275	LEU	C-N-CA	-8.24	87.37	122.00
1	A	534	LYS	O-C-N	-8.24	109.52	122.70
2	B	56	SER	O-C-N	-8.24	109.52	122.70
3	M	54	SER	O-C-N	8.22	135.86	122.70
1	A	573	GLU	O-C-N	-8.22	109.55	122.70
2	B	205	PRO	O-C-N	-8.19	109.59	122.70
3	M	330	GLY	C-N-CA	8.19	142.17	121.70
1	A	192	TYR	CA-C-O	-8.19	102.91	120.10
3	M	91	THR	CA-C-O	-8.17	102.94	120.10
1	A	631	SER	O-C-N	8.16	135.76	122.70
4	S	158	LYS	O-C-N	8.16	135.76	122.70
1	A	443	SER	N-CA-C	8.15	133.01	111.00
2	B	83	PHE	O-C-N	8.14	135.72	122.70
4	S	46	PHE	N-CA-C	-8.13	89.05	111.00
1	A	431	VAL	O-C-N	8.12	135.70	122.70
2	B	105	LEU	O-C-N	-8.12	109.71	122.70
2	B	505	ASP	O-C-N	-8.12	109.71	122.70
2	B	132	SER	C-N-CA	-8.11	101.42	121.70
1	A	415	ARG	N-CA-C	-8.11	89.10	111.00
1	A	487	MET	O-C-N	-8.11	109.72	122.70
1	A	100	LEU	O-C-N	-8.11	109.73	122.70
1	A	637	GLU	C-N-CA	-8.10	101.44	121.70
1	A	103	LYS	CA-C-O	8.09	137.10	120.10
1	A	434	SER	O-C-N	8.09	135.65	122.70
3	M	74	TYR	CA-C-O	8.08	137.07	120.10
2	B	132	SER	O-C-N	-8.08	109.77	122.70
4	S	74	GLN	O-C-N	-8.07	109.79	122.70
1	A	138	ASN	N-CA-C	-8.05	89.26	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	142	LEU	O-C-N	-8.04	109.84	122.70
2	B	115	LEU	O-C-N	8.02	135.53	122.70
2	B	375	LEU	O-C-N	-8.02	105.86	121.10
2	B	515	PHE	C-N-CA	-8.02	105.45	122.30
2	B	296	ASP	CA-C-O	-8.01	103.28	120.10
1	A	547	VAL	O-C-N	8.01	135.51	122.70
2	B	273	SER	N-CA-C	8.01	132.62	111.00
1	A	220	SER	O-C-N	7.99	135.49	122.70
2	B	240	LEU	N-CA-C	7.98	132.55	111.00
2	B	35	TYR	O-C-N	7.97	135.46	122.70
3	M	55	MET	CA-C-N	7.97	134.74	117.20
3	M	294	ASP	C-N-CA	7.97	139.03	122.30
1	A	264	SER	C-N-CA	-7.96	101.79	121.70
2	B	389	ILE	O-C-N	-7.94	109.99	122.70
1	A	545	HIS	O-C-N	-7.94	110.00	122.70
2	B	108	PHE	CA-C-O	7.94	136.77	120.10
2	B	566	ALA	C-N-CA	7.94	141.54	121.70
3	M	262	THR	C-N-CA	-7.92	101.90	121.70
2	B	299	LEU	O-C-N	7.91	135.36	122.70
1	A	281	LEU	C-N-CA	-7.91	101.93	121.70
2	B	126	SER	O-C-N	-7.91	110.05	122.70
3	M	292	PRO	N-CA-C	-7.90	91.57	112.10
1	A	218	ALA	O-C-N	7.89	135.33	122.70
2	B	508	ARG	O-C-N	-7.89	110.08	122.70
2	B	325	LEU	C-N-CA	-7.88	101.99	121.70
1	A	212	ILE	O-C-N	-7.86	110.13	122.70
1	A	281	LEU	O-C-N	-7.86	110.13	122.70
1	A	448	GLU	N-CA-C	-7.84	89.82	111.00
2	B	261	PRO	CA-N-CD	-7.84	100.52	111.50
1	A	365	VAL	O-C-N	-7.84	110.16	122.70
1	A	158	LEU	O-C-N	7.84	135.24	122.70
4	S	108	SER	O-C-N	7.84	135.24	122.70
2	B	153	ILE	O-C-N	7.83	135.23	122.70
3	M	72	LEU	C-N-CA	7.82	141.24	121.70
2	B	317	VAL	O-C-N	7.80	135.18	122.70
1	A	70	ALA	O-C-N	7.80	135.18	122.70
2	B	172	GLU	O-C-N	-7.80	110.22	122.70
3	M	263	MET	C-N-CA	7.80	138.68	122.30
1	A	298	ILE	O-C-N	-7.78	110.25	122.70
4	S	132	LEU	O-C-N	7.77	135.13	122.70
2	B	418	TYR	C-N-CA	-7.76	102.29	121.70
2	B	528	ASP	O-C-N	7.76	135.12	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	588	LEU	O-C-N	-7.75	110.29	122.70
1	A	218	ALA	CB-CA-C	7.75	121.72	110.10
2	B	147	MET	C-N-CA	-7.74	102.34	121.70
3	M	237	THR	C-N-CA	-7.74	106.05	122.30
1	A	328	PRO	O-C-N	-7.71	110.36	122.70
1	A	330	LEU	O-C-N	7.71	135.04	122.70
1	A	394	GLN	CA-C-O	-7.71	103.92	120.10
4	S	109	LEU	CA-C-O	7.70	136.28	120.10
2	B	478	LEU	O-C-N	-7.70	110.39	122.70
3	M	132	GLY	CA-C-O	7.69	134.45	120.60
1	A	461	CYS	C-N-CA	-7.69	102.47	121.70
2	B	584	SER	O-C-N	-7.68	110.14	123.20
1	A	304	LEU	O-C-N	-7.68	110.42	122.70
2	B	174	ALA	O-C-N	-7.68	110.42	122.70
1	A	346	THR	O-C-N	-7.67	110.43	122.70
2	B	560	ILE	O-C-N	-7.66	110.44	122.70
1	A	527	GLU	C-N-CA	7.66	140.84	121.70
2	B	237	ILE	O-C-N	-7.65	110.46	122.70
1	A	559	PHE	O-C-N	7.65	134.93	122.70
4	S	36	TYR	O-C-N	7.64	134.93	122.70
3	M	232	HIS	CA-C-O	7.64	136.15	120.10
1	A	156	PRO	O-C-N	-7.63	110.49	122.70
2	B	220	ALA	N-CA-C	-7.63	90.40	111.00
1	A	601	VAL	O-C-N	-7.62	110.50	122.70
4	S	139	GLY	C-N-CA	-7.62	102.64	121.70
1	A	469	LEU	CA-C-O	7.61	136.07	120.10
2	B	51	LEU	C-N-CA	-7.61	102.68	121.70
4	S	28	GLN	O-C-N	7.60	134.86	122.70
1	A	601	VAL	C-N-CA	-7.59	102.72	121.70
2	B	486	HIS	O-C-N	7.59	134.85	122.70
2	B	585	GLY	N-CA-C	-7.58	94.14	113.10
4	S	27	LYS	O-C-N	7.57	134.81	122.70
1	A	279	LEU	O-C-N	-7.57	110.60	122.70
1	A	504	ILE	CA-C-O	7.55	135.96	120.10
1	A	465	SER	N-CA-C	7.55	131.38	111.00
1	A	98	ASN	CA-C-N	7.54	133.80	117.20
2	B	237	ILE	C-N-CA	-7.54	102.86	121.70
1	A	528	ASN	CA-C-O	7.54	135.93	120.10
3	M	265	ASN	N-CA-C	-7.53	90.66	111.00
2	B	488	ARG	O-C-N	7.53	134.75	122.70
1	A	88	ASN	CA-C-O	7.53	135.91	120.10
2	B	277	CYS	C-N-CD	7.53	144.20	128.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	338	PHE	O-C-N	7.52	134.73	122.70
3	M	230	LYS	CA-C-N	7.51	133.73	117.20
1	A	596	VAL	O-C-N	7.50	134.70	122.70
2	B	46	GLN	O-C-N	7.50	134.70	122.70
4	S	55	PRO	CA-N-CD	-7.50	101.00	111.50
2	B	416	LYS	O-C-N	7.48	134.66	122.70
2	B	215	TYR	O-C-N	7.47	134.65	122.70
1	A	394	GLN	O-C-N	7.46	134.64	122.70
2	B	442	LEU	C-N-CA	-7.45	103.07	121.70
3	M	266	ASP	CA-C-N	-7.44	100.83	117.20
3	M	367	ALA	CB-CA-C	7.44	121.26	110.10
4	S	96	LEU	O-C-N	7.43	134.59	122.70
2	B	325	LEU	CA-C-O	7.43	135.70	120.10
1	A	465	SER	O-C-N	7.42	134.58	122.70
2	B	328	LEU	C-N-CA	-7.42	103.14	121.70
1	A	606	PHE	O-C-N	7.42	134.57	122.70
2	B	494	ALA	O-C-N	7.42	134.56	122.70
1	A	519	LEU	O-C-N	-7.41	110.61	123.20
2	B	428	VAL	O-C-N	7.41	134.55	122.70
2	B	362	ALA	O-C-N	7.37	134.49	122.70
1	A	311	THR	O-C-N	7.37	134.49	122.70
3	M	407	THR	N-CA-C	-7.36	91.12	111.00
1	A	391	LEU	O-C-N	-7.35	110.94	122.70
2	B	142	LEU	C-N-CA	-7.34	103.35	121.70
3	M	421	GLY	O-C-N	-7.34	107.16	121.10
4	S	54	PRO	C-N-CD	7.33	143.79	128.40
1	A	405	THR	N-CA-C	7.33	130.78	111.00
2	B	493	LEU	O-C-N	7.32	134.42	122.70
1	A	177	ILE	O-C-N	7.32	134.41	122.70
1	A	256	LEU	O-C-N	7.32	134.41	122.70
4	S	31	LEU	O-C-N	7.31	134.40	122.70
2	B	223	LEU	O-C-N	-7.30	111.01	122.70
4	S	149	ILE	O-C-N	7.30	134.39	122.70
2	B	192	LEU	O-C-N	7.30	134.38	122.70
2	B	288	TYR	N-CA-C	-7.30	91.29	111.00
1	A	260	PHE	C-N-CA	-7.30	103.46	121.70
1	A	162	ILE	O-C-N	7.29	134.37	122.70
2	B	515	PHE	O-C-N	-7.29	110.80	123.20
4	S	167	ILE	N-CA-C	7.29	130.68	111.00
1	A	511	VAL	O-C-N	7.29	134.36	122.70
4	S	80	TYR	CA-C-O	7.28	135.40	120.10
2	B	408	VAL	O-C-N	7.27	134.34	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	220	ALA	CB-CA-C	7.27	121.00	110.10
2	B	270	SER	N-CA-C	-7.27	91.37	111.00
2	B	531	ARG	O-C-N	7.27	134.33	122.70
1	A	399	ASP	O-C-N	7.26	134.32	122.70
4	S	127	THR	O-C-N	7.26	134.32	122.70
1	A	384	LEU	O-C-N	7.26	134.31	122.70
2	B	267	ASP	O-C-N	-7.26	111.09	122.70
2	B	554	LYS	O-C-N	7.25	134.30	122.70
3	M	426	LYS	C-N-CA	7.25	139.82	121.70
3	M	460	ILE	C-N-CA	7.23	137.49	122.30
1	A	573	GLU	C-N-CA	-7.23	103.64	121.70
1	A	529	GLY	N-CA-C	-7.21	95.08	113.10
1	A	265	GLN	CA-C-O	-7.17	105.04	120.10
4	S	83	LEU	CA-C-O	7.17	135.16	120.10
1	A	306	GLU	O-C-N	-7.17	111.23	122.70
4	S	125	TRP	CA-C-O	-7.16	105.06	120.10
3	M	54	SER	C-N-CA	7.16	139.60	121.70
1	A	298	ILE	CA-C-O	7.16	135.14	120.10
2	B	150	LEU	O-C-N	7.15	134.14	122.70
2	B	243	TRP	O-C-N	7.15	134.14	122.70
1	A	441	TYR	CA-C-N	-7.15	101.48	117.20
2	B	478	LEU	CA-C-O	7.14	135.10	120.10
1	A	300	LYS	N-CA-C	7.14	130.28	111.00
2	B	415	LEU	O-C-N	7.14	134.12	122.70
1	A	263	LEU	C-N-CA	-7.13	103.87	121.70
3	M	96	ILE	O-C-N	7.13	134.10	122.70
1	A	388	VAL	O-C-N	7.12	134.09	122.70
1	A	447	PHE	O-C-N	7.12	134.09	122.70
3	M	118	TYR	CA-C-O	-7.11	105.17	120.10
1	A	517	TRP	O-C-N	7.11	134.08	122.70
1	A	488	ARG	O-C-N	7.11	134.07	122.70
1	A	355	LEU	O-C-N	7.10	134.06	122.70
2	B	23	ALA	N-CA-CB	7.09	120.03	110.10
1	A	582	ILE	O-C-N	7.09	134.04	122.70
1	A	381	GLU	O-C-N	-7.08	111.36	122.70
4	S	84	TYR	O-C-N	-7.08	111.37	122.70
2	B	611	ALA	O-C-N	7.08	134.02	122.70
1	A	539	ASN	C-N-CA	-7.08	104.01	121.70
2	B	226	LEU	O-C-N	7.08	134.02	122.70
1	A	453	VAL	O-C-N	7.07	134.01	122.70
2	B	521	ILE	C-N-CA	-7.07	104.03	121.70
2	B	213	LEU	O-C-N	7.05	133.99	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
4	S	99	LEU	CA-C-O	-7.05	105.29	120.10
3	M	110	SER	O-C-N	7.05	133.98	122.70
2	B	555	LEU	O-C-N	7.05	133.97	122.70
1	A	566	PHE	N-CA-C	7.04	130.02	111.00
2	B	504	ALA	O-C-N	7.04	133.97	122.70
2	B	526	CYS	O-C-N	7.04	134.48	121.10
1	A	166	LEU	O-C-N	7.03	133.95	122.70
1	A	441	TYR	CA-C-O	7.03	134.86	120.10
1	A	302	ASN	CA-C-N	-7.03	101.74	117.20
4	S	30	LEU	O-C-N	7.03	133.94	122.70
1	A	442	SER	N-CA-C	-7.02	92.04	111.00
1	A	225	LEU	CA-C-O	-7.02	105.36	120.10
1	A	635	ALA	O-C-N	7.02	133.93	122.70
2	B	389	ILE	C-N-CA	-7.01	104.17	121.70
2	B	143	SER	C-N-CA	-7.01	104.17	121.70
3	M	458	LEU	C-N-CA	7.00	139.21	121.70
3	M	425	THR	C-N-CA	7.00	139.19	121.70
1	A	271	ARG	O-C-N	6.99	133.88	122.70
2	B	97	VAL	O-C-N	6.99	133.88	122.70
1	A	225	LEU	O-C-N	6.99	133.88	122.70
2	B	422	ALA	N-CA-CB	6.99	119.88	110.10
1	A	301	GLY	N-CA-C	-6.98	95.64	113.10
2	B	572	GLU	O-C-N	6.98	133.87	122.70
1	A	605	GLU	O-C-N	6.98	133.87	122.70
4	S	143	GLU	CA-C-N	-6.98	101.84	117.20
1	A	476	GLN	O-C-N	6.98	133.87	122.70
2	B	216	LYS	C-N-CA	-6.96	104.29	121.70
1	A	140	VAL	O-C-N	-6.96	111.56	122.70
1	A	257	LEU	O-C-N	6.96	133.83	122.70
2	B	56	SER	C-N-CA	-6.96	104.31	121.70
1	A	156	PRO	C-N-CA	-6.96	104.31	121.70
2	B	102	HIS	CA-C-O	6.95	134.70	120.10
2	B	355	ASN	O-C-N	6.95	133.82	122.70
1	A	138	ASN	O-C-N	6.95	133.82	122.70
2	B	324	ALA	O-C-N	6.95	133.82	122.70
2	B	425	PRO	O-C-N	6.95	133.82	122.70
2	B	104	TYR	O-C-N	6.94	133.80	122.70
2	B	559	ASP	O-C-N	6.93	133.79	122.70
2	B	252	LEU	O-C-N	6.92	133.78	122.70
1	A	204	VAL	C-N-CA	-6.92	104.40	121.70
1	A	607	LEU	O-C-N	6.91	133.75	122.70
2	B	99	ARG	O-C-N	6.90	133.74	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	240	LEU	O-C-N	-6.90	111.66	122.70
1	A	348	PHE	O-C-N	6.90	133.73	122.70
2	B	271	GLU	C-N-CA	-6.89	107.83	122.30
2	B	511	ILE	O-C-N	-6.89	111.68	122.70
1	A	383	ASN	O-C-N	6.89	133.72	122.70
1	A	621	LEU	C-N-CA	-6.88	93.09	122.00
2	B	169	VAL	O-C-N	6.88	133.72	122.70
1	A	279	LEU	C-N-CA	-6.88	104.50	121.70
1	A	457	LEU	O-C-N	6.88	133.71	122.70
1	A	608	ARG	O-C-N	6.88	133.70	122.70
4	S	18	LYS	O-C-N	-6.88	111.69	122.70
2	B	573	GLU	C-N-CA	-6.87	104.52	121.70
1	A	632	PHE	O-C-N	6.87	133.69	122.70
2	B	323	ASN	O-C-N	6.87	133.69	122.70
4	S	157	ASN	O-C-N	6.87	133.69	122.70
2	B	134	LEU	O-C-N	6.86	133.68	122.70
4	S	99	LEU	O-C-N	6.86	133.67	122.70
1	A	140	VAL	CA-C-O	6.86	134.50	120.10
2	B	105	LEU	C-N-CA	-6.85	104.58	121.70
4	S	33	GLU	O-C-N	6.85	133.65	122.70
3	M	89	CYS	O-C-N	6.84	133.65	122.70
2	B	544	THR	O-C-N	6.84	133.65	122.70
1	A	102	GLN	O-C-N	6.84	133.64	122.70
1	A	602	GLU	O-C-N	6.83	133.63	122.70
1	A	555	LEU	O-C-N	6.83	133.63	122.70
2	B	63	MET	O-C-N	6.83	133.63	122.70
1	A	610	SER	O-C-N	6.82	133.61	122.70
2	B	568	VAL	C-N-CA	6.82	138.74	121.70
2	B	594	ALA	O-C-N	6.81	133.60	122.70
3	M	123	LEU	O-C-N	6.81	133.60	122.70
1	A	623	MET	O-C-N	6.81	133.60	122.70
2	B	66	ILE	O-C-N	6.81	133.60	122.70
1	A	477	PHE	O-C-N	6.81	133.59	122.70
2	B	457	HIS	CA-C-O	6.81	134.39	120.10
1	A	236	LEU	O-C-N	6.81	133.59	122.70
1	A	188	VAL	O-C-N	6.80	133.59	122.70
4	S	148	ARG	O-C-N	6.80	133.58	122.70
1	A	496	ILE	O-C-N	6.79	133.56	122.70
1	A	624	LEU	O-C-N	6.79	133.56	122.70
2	B	454	LEU	O-C-N	6.79	133.56	122.70
3	M	54	SER	CB-CA-C	-6.78	97.21	110.10
2	B	174	ALA	C-N-CA	-6.78	104.76	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	M	54	SER	CA-C-N	-6.78	102.29	117.20
4	S	101	LEU	O-C-N	6.78	133.54	122.70
2	B	489	ILE	O-C-N	6.78	133.54	122.70
1	A	466	ASP	O-C-N	6.77	133.54	122.70
2	B	100	LEU	O-C-N	6.76	133.52	122.70
3	M	47	SER	N-CA-C	-6.76	92.74	111.00
2	B	411	ILE	O-C-N	6.75	133.51	122.70
3	M	385	ARG	O-C-N	-6.75	111.89	122.70
4	S	16	LEU	O-C-N	-6.75	111.89	122.70
1	A	581	LEU	O-C-N	6.75	133.50	122.70
1	A	601	VAL	CA-C-O	6.75	134.28	120.10
2	B	78	ASP	O-C-N	-6.75	111.90	122.70
1	A	67	LYS	O-C-N	6.75	133.49	122.70
1	A	418	ILE	N-CA-C	6.73	129.18	111.00
2	B	253	ILE	O-C-N	6.73	133.47	122.70
1	A	125	THR	O-C-N	6.72	133.46	122.70
3	M	424	PHE	N-CA-C	-6.72	92.85	111.00
2	B	452	LYS	O-C-N	6.72	133.45	122.70
3	M	425	THR	N-CA-C	-6.72	92.86	111.00
1	A	91	ILE	O-C-N	6.71	133.44	122.70
1	A	265	GLN	N-CA-CB	6.71	122.69	110.60
4	S	128	LEU	O-C-N	6.71	133.44	122.70
2	B	566	ALA	CB-CA-C	6.71	120.17	110.10
3	M	85	GLY	N-CA-C	-6.71	96.33	113.10
2	B	592	TYR	O-C-N	6.70	133.42	122.70
1	A	432	ILE	CA-C-O	-6.69	106.05	120.10
1	A	87	CYS	O-C-N	6.69	133.40	122.70
1	A	117	ASP	O-C-N	6.69	133.40	122.70
1	A	307	ASP	N-CA-C	-6.68	92.95	111.00
3	M	265	ASN	CA-C-N	6.68	131.91	117.20
3	M	284	SER	O-C-N	-6.68	108.41	121.10
4	S	137	GLN	CA-C-N	-6.68	102.84	116.20
1	A	128	LEU	O-C-N	6.68	133.38	122.70
1	A	472	LYS	O-C-N	6.68	133.38	122.70
1	A	196	LEU	CA-C-O	6.68	134.12	120.10
1	A	569	ASP	O-C-N	-6.68	112.02	122.70
3	M	113	LYS	O-C-N	6.68	133.38	122.70
2	B	356	LYS	O-C-N	6.67	133.38	122.70
2	B	154	ILE	CA-C-O	-6.67	106.09	120.10
3	M	16	PHE	CA-C-O	6.67	134.10	120.10
2	B	139	LEU	O-C-N	6.67	133.37	122.70
4	S	150	VAL	O-C-N	6.67	133.36	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	361	GLN	O-C-N	6.66	133.36	122.70
1	A	392	MET	O-C-N	6.66	133.35	122.70
2	B	460	SER	N-CA-C	6.66	128.97	111.00
2	B	607	ILE	O-C-N	6.66	133.35	122.70
4	S	54	PRO	N-CA-C	-6.66	94.79	112.10
2	B	244	SER	O-C-N	6.65	133.34	122.70
2	B	589	SER	O-C-N	6.65	133.34	122.70
2	B	337	THR	CA-C-O	-6.65	106.14	120.10
2	B	399	LEU	C-N-CA	-6.65	105.08	121.70
2	B	412	PHE	O-C-N	6.65	133.34	122.70
1	A	107	TYR	O-C-N	6.64	133.33	122.70
2	B	273	SER	N-CA-CB	-6.64	100.53	110.50
1	A	265	GLN	O-C-N	-6.64	112.07	122.70
2	B	65	ARG	O-C-N	6.64	133.32	122.70
1	A	522	PHE	O-C-N	6.63	133.31	122.70
1	A	600	SER	O-C-N	6.63	133.31	122.70
2	B	447	GLU	O-C-N	6.63	133.31	122.70
1	A	462	GLN	C-N-CA	-6.63	105.13	121.70
1	A	296	ASN	O-C-N	6.63	133.30	122.70
3	M	394	GLN	CA-C-O	6.63	134.02	120.10
2	B	552	SER	O-C-N	6.62	133.30	122.70
4	S	106	VAL	O-C-N	6.62	133.30	122.70
2	B	521	ILE	N-CA-C	-6.62	93.12	111.00
1	A	515	CYS	O-C-N	6.62	133.29	122.70
1	A	556	VAL	O-C-N	6.60	133.26	122.70
1	A	436	CYS	O-C-N	-6.59	112.15	122.70
4	S	35	VAL	O-C-N	6.59	133.25	122.70
1	A	80	TYR	N-CA-CB	6.59	122.46	110.60
2	B	123	LEU	CA-C-O	-6.59	106.27	120.10
2	B	267	ASP	C-N-CA	-6.58	105.24	121.70
1	A	139	ASP	O-C-N	6.58	133.23	122.70
1	A	558	VAL	O-C-N	6.58	133.23	122.70
2	B	267	ASP	CA-C-O	6.58	133.93	120.10
2	B	591	MET	O-C-N	6.58	133.23	122.70
1	A	554	ALA	O-C-N	6.58	133.23	122.70
1	A	611	LEU	O-C-N	6.58	133.23	122.70
4	S	32	LEU	O-C-N	6.58	133.23	122.70
2	B	123	LEU	O-C-N	6.58	133.22	122.70
2	B	251	LEU	O-C-N	6.58	133.22	122.70
1	A	495	ILE	O-C-N	6.57	133.21	122.70
1	A	332	TYR	O-C-N	6.57	133.21	122.70
2	B	475	ILE	O-C-N	6.57	133.21	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	406	GLY	N-CA-C	-6.57	96.69	113.10
2	B	62	ALA	O-C-N	6.57	133.21	122.70
2	B	474	VAL	O-C-N	6.57	133.20	122.70
1	A	454	ILE	O-C-N	6.56	133.19	122.70
4	S	29	LYS	O-C-N	6.56	133.19	122.70
2	B	359	LEU	O-C-N	6.55	133.19	122.70
1	A	507	GLN	C-N-CA	-6.55	105.33	121.70
2	B	513	TRP	O-C-N	6.54	133.17	122.70
2	B	508	ARG	CA-C-O	6.54	133.83	120.10
1	A	562	TRP	O-C-N	6.54	133.16	122.70
1	A	145	ILE	O-C-N	6.53	133.15	122.70
1	A	451	ASN	O-C-N	6.53	133.15	122.70
2	B	346	THR	O-C-N	6.53	133.15	122.70
1	A	386	ALA	O-C-N	6.53	133.15	122.70
2	B	604	GLU	O-C-N	6.53	133.15	122.70
2	B	301	LEU	O-C-N	6.53	133.14	122.70
1	A	492	ILE	O-C-N	6.52	133.14	122.70
2	B	588	ILE	O-C-N	6.52	133.13	122.70
2	B	453	TRP	O-C-N	6.51	133.12	122.70
3	M	458	LEU	N-CA-C	-6.51	93.42	111.00
3	M	394	GLN	O-C-N	-6.51	112.14	123.20
2	B	467	VAL	O-C-N	6.51	133.11	122.70
2	B	154	ILE	O-C-N	6.50	133.11	122.70
4	S	129	GLU	O-C-N	6.50	133.10	122.70
2	B	388	PRO	C-N-CA	6.50	137.94	121.70
2	B	177	ILE	O-C-N	6.50	133.09	122.70
1	A	612	GLU	O-C-N	6.50	133.09	122.70
2	B	344	VAL	O-C-N	6.49	133.09	122.70
4	S	83	LEU	O-C-N	-6.49	112.31	122.70
2	B	137	PHE	O-C-N	6.49	133.09	122.70
2	B	146	LYS	CA-C-N	6.49	131.48	117.20
4	S	4	ALA	CA-C-O	6.49	133.73	120.10
2	B	507	ALA	O-C-N	6.49	133.08	122.70
1	A	569	ASP	CA-C-O	-6.49	106.48	120.10
4	S	124	ASN	O-C-N	6.49	133.08	122.70
2	B	119	SER	O-C-N	6.48	133.07	122.70
3	M	91	THR	O-C-N	6.48	133.07	122.70
3	M	124	ILE	O-C-N	6.48	133.07	122.70
4	S	106	VAL	CA-C-O	-6.48	106.49	120.10
4	S	130	SER	O-C-N	6.48	133.07	122.70
2	B	548	ILE	O-C-N	6.48	133.06	122.70
3	M	122	SER	O-C-N	6.48	133.06	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	145	MET	C-N-CA	-6.46	105.56	121.70
2	B	230	PHE	O-C-N	-6.46	112.37	122.70
1	A	146	ALA	O-C-N	6.45	133.02	122.70
4	S	145	ASN	O-C-N	6.45	133.02	122.70
1	A	105	VAL	O-C-N	6.45	134.16	123.20
3	M	94	GLU	O-C-N	6.45	133.01	122.70
3	M	389	SER	O-C-N	6.44	133.01	122.70
1	A	388	VAL	CA-C-O	-6.44	106.57	120.10
2	B	229	HIS	C-N-CA	6.44	137.80	121.70
1	A	64	LEU	CA-C-O	6.44	133.62	120.10
1	A	185	LEU	O-C-N	6.43	132.99	122.70
2	B	81	LEU	C-N-CA	-6.43	105.62	121.70
2	B	363	ILE	O-C-N	6.43	132.99	122.70
2	B	587	ARG	O-C-N	6.42	132.96	122.70
3	M	457	GLY	CA-C-N	-6.41	103.10	117.20
2	B	490	ILE	O-C-N	6.41	132.95	122.70
2	B	306	LEU	CA-C-O	-6.40	106.65	120.10
1	A	252	ILE	O-C-N	6.40	132.94	122.70
1	A	313	MET	O-C-N	6.40	132.94	122.70
1	A	550	VAL	O-C-N	6.40	132.94	122.70
2	B	414	GLU	O-C-N	6.40	132.93	122.70
1	A	159	ALA	O-C-N	6.39	132.93	122.70
1	A	407	SER	N-CA-C	-6.39	93.74	111.00
1	A	467	LYS	O-C-N	6.39	132.92	122.70
2	B	135	ARG	O-C-N	6.39	132.92	122.70
1	A	331	ARG	O-C-N	6.39	132.92	122.70
1	A	364	ASP	O-C-N	6.38	132.92	122.70
2	B	193	LEU	O-C-N	6.38	132.92	122.70
1	A	492	ILE	CA-C-O	-6.38	106.70	120.10
1	A	516	ILE	O-C-N	6.38	132.91	122.70
4	S	155	GLU	O-C-N	6.38	132.91	122.70
1	A	598	GLU	O-C-N	6.38	132.90	122.70
1	A	629	LEU	C-N-CA	-6.38	95.22	122.00
2	B	102	HIS	C-N-CA	-6.38	105.76	121.70
1	A	415	ARG	C-N-CA	-6.37	105.78	121.70
1	A	552	ILE	O-C-N	6.37	132.89	122.70
1	A	215	VAL	O-C-N	-6.37	112.52	122.70
2	B	178	ILE	O-C-N	6.36	132.88	122.70
2	B	271	GLU	N-CA-C	6.36	128.18	111.00
4	S	165	SER	C-N-CA	-6.36	105.79	121.70
2	B	170	ARG	O-C-N	-6.36	112.38	123.20
1	A	474	GLY	O-C-N	6.36	132.88	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	186	PHE	O-C-N	6.36	132.87	122.70
1	A	604	LEU	O-C-N	6.36	132.87	122.70
4	S	153	VAL	O-C-N	6.36	132.87	122.70
3	M	111	ILE	O-C-N	6.35	132.87	122.70
2	B	395	LYS	O-C-N	6.35	132.86	122.70
4	S	156	LEU	O-C-N	6.35	132.87	122.70
2	B	151	ALA	O-C-N	6.35	133.17	121.10
1	A	471	SER	O-C-N	6.35	132.86	122.70
2	B	248	LEU	O-C-N	6.35	132.85	122.70
2	B	510	GLY	CA-C-O	-6.35	109.18	120.60
1	A	475	GLU	O-C-N	6.34	132.85	122.70
3	M	272	LEU	C-N-CA	6.34	137.55	121.70
2	B	79	VAL	CA-C-O	-6.34	106.79	120.10
1	A	603	VAL	O-C-N	6.33	132.84	122.70
1	A	297	CYS	O-C-N	-6.33	112.57	122.70
1	A	147	LEU	O-C-N	6.33	132.83	122.70
2	B	519	ALA	N-CA-CB	-6.33	101.24	110.10
2	B	67	ILE	O-C-N	6.33	132.82	122.70
2	B	417	TYR	O-C-N	6.33	132.82	122.70
2	B	432	ALA	O-C-N	6.33	132.82	122.70
1	A	628	VAL	O-C-N	6.32	132.82	122.70
1	A	387	ILE	O-C-N	6.32	132.81	122.70
4	S	37	GLU	O-C-N	6.32	132.81	122.70
2	B	150	LEU	CA-C-O	-6.32	106.83	120.10
2	B	204	ASP	O-C-N	6.32	133.10	121.10
1	A	202	LYS	O-C-N	6.31	132.80	122.70
2	B	59	VAL	O-C-N	6.31	132.80	122.70
2	B	455	ILE	O-C-N	6.31	132.80	122.70
2	B	158	VAL	CA-C-O	-6.31	106.85	120.10
2	B	305	SER	O-C-N	6.31	132.79	122.70
2	B	306	LEU	O-C-N	6.31	132.79	122.70
2	B	612	ARG	O-C-N	6.31	132.80	122.70
1	A	578	LEU	O-C-N	6.31	132.79	122.70
2	B	209	SER	O-C-N	6.31	132.79	122.70
2	B	341	GLU	O-C-N	6.30	132.79	122.70
2	B	302	PHE	O-C-N	6.30	132.78	122.70
3	M	86	PRO	O-C-N	6.30	132.78	122.70
1	A	182	ILE	O-C-N	6.30	132.78	122.70
2	B	429	VAL	O-C-N	6.30	132.78	122.70
2	B	610	ARG	O-C-N	6.30	132.78	122.70
4	S	38	LEU	O-C-N	6.30	132.78	122.70
4	S	80	TYR	O-C-N	-6.30	112.62	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	429	VAL	O-C-N	6.30	132.77	122.70
1	A	508	LEU	CA-C-O	-6.30	106.88	120.10
2	B	613	MET	O-C-N	6.30	132.77	122.70
4	S	105	PHE	O-C-N	6.30	132.77	122.70
1	A	574	ILE	O-C-N	6.29	132.77	122.70
2	B	111	ASN	N-CA-C	6.29	127.99	111.00
1	A	337	LEU	O-C-N	6.29	132.76	122.70
1	A	613	ALA	O-C-N	6.29	132.76	122.70
1	A	421	PRO	O-C-N	6.29	132.76	122.70
4	S	120	ASP	O-C-N	6.29	132.76	122.70
4	S	107	GLU	O-C-N	6.28	132.75	122.70
2	B	360	LEU	O-C-N	6.28	132.75	122.70
2	B	437	SER	O-C-N	6.28	132.75	122.70
4	S	1	MET	O-C-N	-6.28	112.65	122.70
1	A	522	PHE	CA-C-O	-6.28	106.92	120.10
2	B	116	THR	O-C-N	6.28	132.74	122.70
1	A	302	ASN	CA-C-O	6.27	133.27	120.10
1	A	508	LEU	O-C-N	6.27	133.02	121.10
3	M	55	MET	C-N-CA	-6.27	106.02	121.70
2	B	43	ASN	O-C-N	6.27	133.01	121.10
1	A	249	ASN	O-C-N	6.27	132.73	122.70
2	B	249	ILE	O-C-N	6.27	132.73	122.70
2	B	158	VAL	O-C-N	6.27	132.73	122.70
1	A	68	THR	O-C-N	6.26	132.72	122.70
2	B	244	SER	CA-C-O	-6.26	106.95	120.10
2	B	472	VAL	CA-C-O	6.26	133.25	120.10
2	B	485	LYS	O-C-N	6.26	132.71	122.70
2	B	595	VAL	O-C-N	6.26	132.71	122.70
1	A	395	PHE	C-N-CA	-6.25	106.06	121.70
1	A	385	LYS	O-C-N	6.25	132.70	122.70
2	B	374	PHE	O-C-N	6.25	132.70	122.70
2	B	605	PHE	O-C-N	6.25	132.70	122.70
1	A	278	ILE	C-N-CA	6.25	137.32	121.70
2	B	321	CYS	O-C-N	6.25	132.69	122.70
2	B	410	GLU	O-C-N	6.25	132.69	122.70
1	A	83	ASP	CA-C-N	-6.24	103.46	117.20
1	A	315	CYS	O-C-N	6.24	132.69	122.70
1	A	428	MET	O-C-N	6.24	132.69	122.70
2	B	191	GLU	O-C-N	6.24	132.69	122.70
1	A	625	LEU	O-C-N	-6.24	112.72	122.70
2	B	58	GLU	O-C-N	6.24	132.68	122.70
1	A	353	ASP	O-C-N	6.23	132.67	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	419	ILE	N-CA-C	6.23	127.83	111.00
2	B	214	ALA	O-C-N	6.23	132.67	122.70
2	B	436	LEU	O-C-N	6.23	132.67	122.70
1	A	124	ALA	O-C-N	6.23	132.66	122.70
1	A	404	GLN	C-N-CA	6.22	137.26	121.70
4	S	85	PHE	CA-C-O	6.22	133.17	120.10
1	A	473	ILE	O-C-N	6.22	133.78	123.20
2	B	151	ALA	CA-C-O	-6.22	107.03	120.10
2	B	105	LEU	CA-C-O	6.22	133.16	120.10
1	A	579	LYS	O-C-N	6.22	132.65	122.70
2	B	155	LEU	O-C-N	6.22	132.65	122.70
2	B	615	SER	O-C-N	6.22	132.65	122.70
1	A	200	PHE	O-C-N	6.21	132.64	122.70
2	B	409	LYS	O-C-N	6.21	132.64	122.70
2	B	157	THR	O-C-N	6.21	132.64	122.70
2	B	246	SER	O-C-N	6.21	132.63	122.70
3	M	223	HIS	O-C-N	-6.21	112.77	122.70
1	A	298	ILE	C-N-CA	-6.21	106.19	121.70
1	A	421	PRO	CA-C-O	-6.20	105.31	120.20
1	A	309	PHE	O-C-N	6.20	132.62	122.70
2	B	593	ASN	O-C-N	6.20	132.62	122.70
1	A	141	VAL	O-C-N	6.20	132.61	122.70
2	B	136	CYS	O-C-N	6.19	132.61	122.70
2	B	614	ILE	O-C-N	6.19	132.60	122.70
2	B	27	THR	C-N-CA	6.19	137.17	121.70
2	B	433	VAL	O-C-N	6.19	132.60	122.70
1	A	535	ILE	CA-C-N	-6.18	103.60	117.20
1	A	297	CYS	CA-C-O	6.18	133.08	120.10
1	A	316	LEU	O-C-N	6.18	132.59	122.70
1	A	551	LEU	O-C-N	6.18	132.59	122.70
2	B	233	TYR	O-C-N	6.18	132.59	122.70
2	B	337	THR	O-C-N	6.18	132.58	122.70
2	B	468	LEU	O-C-N	6.17	132.58	122.70
2	B	342	ALA	O-C-N	6.17	132.57	122.70
1	A	162	ILE	CA-C-O	-6.17	107.15	120.10
2	B	550	VAL	O-C-N	6.16	132.56	122.70
2	B	209	SER	CA-C-O	-6.16	107.17	120.10
1	A	533	ILE	CA-C-O	-6.16	107.17	120.10
1	A	81	GLY	C-N-CA	6.15	137.08	121.70
1	A	607	LEU	CA-C-O	-6.15	107.18	120.10
1	A	274	LEU	C-N-CA	-6.15	106.33	121.70
2	B	430	ILE	O-C-N	6.14	132.53	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	127	LEU	O-C-N	6.14	132.52	122.70
2	B	138	ALA	O-C-N	6.14	132.52	122.70
2	B	296	ASP	O-C-N	6.14	132.76	121.10
2	B	319	LEU	O-C-N	6.14	132.52	122.70
1	A	143	VAL	O-C-N	6.14	133.63	123.20
1	A	323	CYS	CA-C-N	6.14	130.70	117.20
2	B	272	GLY	N-CA-C	-6.14	97.76	113.10
2	B	343	LEU	O-C-N	6.13	132.52	122.70
1	A	295	VAL	CA-C-O	-6.13	107.22	120.10
1	A	335	CYS	O-C-N	6.13	132.51	122.70
2	B	512	VAL	O-C-N	6.13	132.51	122.70
1	A	295	VAL	O-C-N	6.13	132.50	122.70
2	B	195	ILE	O-C-N	6.13	132.50	122.70
2	B	139	LEU	CA-C-O	-6.12	107.24	120.10
2	B	324	ALA	CA-C-O	-6.12	107.24	120.10
1	A	155	THR	CA-C-O	-6.12	107.25	120.10
2	B	98	LYS	O-C-N	6.12	132.49	122.70
1	A	454	ILE	CA-C-O	-6.12	107.25	120.10
2	B	464	SER	O-C-N	6.12	132.49	122.70
2	B	549	LEU	O-C-N	6.11	132.47	122.70
2	B	196	LEU	O-C-N	6.11	132.47	122.70
1	A	633	PHE	CA-C-O	-6.10	107.30	120.10
1	A	390	THR	O-C-N	6.10	132.45	122.70
2	B	585	GLY	O-C-N	6.09	132.45	122.70
1	A	105	VAL	C-N-CA	6.09	135.09	122.30
4	S	81	ALA	N-CA-C	-6.09	94.56	111.00
2	B	82	TYR	O-C-N	-6.09	112.96	122.70
1	A	557	LYS	O-C-N	6.08	132.44	122.70
2	B	190	GLU	O-C-N	6.08	132.43	122.70
1	A	336	ILE	O-C-N	6.08	132.43	122.70
2	B	44	PRO	O-C-N	6.08	132.43	122.70
2	B	250	GLU	O-C-N	6.08	132.43	122.70
3	M	308	SER	O-C-N	6.08	132.43	122.70
1	A	271	ARG	CA-C-O	-6.08	107.34	120.10
1	A	584	PHE	O-C-N	6.07	132.42	122.70
2	B	166	SER	O-C-N	6.07	132.42	122.70
3	M	368	ASP	CA-C-O	-6.07	107.35	120.10
2	B	47	LEU	O-C-N	6.07	132.41	122.70
3	M	367	ALA	N-CA-C	-6.07	94.61	111.00
2	B	546	CYS	O-C-N	6.06	132.40	122.70
2	B	568	VAL	O-C-N	6.06	132.40	122.70
2	B	462	ASN	C-N-CA	-6.06	106.55	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	M	232	HIS	O-C-N	-6.06	113.00	122.70
2	B	451	MET	O-C-N	6.06	132.40	122.70
2	B	466	SER	O-C-N	6.06	132.40	122.70
2	B	216	LYS	O-C-N	-6.06	113.01	122.70
2	B	522	GLU	O-C-N	6.06	132.39	122.70
2	B	103	LEU	O-C-N	6.06	132.39	122.70
4	S	139	GLY	N-CA-C	-6.05	97.96	113.10
1	A	211	ASP	O-C-N	6.05	132.38	122.70
1	A	400	VAL	O-C-N	6.05	132.38	122.70
3	M	429	ASP	C-N-CA	-6.05	106.58	121.70
1	A	465	SER	CA-C-O	6.05	132.80	120.10
1	A	142	LYS	O-C-N	6.05	132.38	122.70
1	A	278	ILE	CA-C-O	-6.05	107.40	120.10
1	A	566	PHE	C-N-CA	6.05	136.82	121.70
2	B	596	LEU	O-C-N	6.05	132.38	122.70
2	B	579	PRO	N-CA-C	6.04	127.82	112.10
4	S	16	LEU	CA-C-O	6.04	132.79	120.10
1	A	369	SER	O-C-N	6.04	132.36	122.70
4	S	150	VAL	CA-C-O	-6.04	107.42	120.10
2	B	435	SER	O-C-N	6.04	132.36	122.70
1	A	107	TYR	CA-C-O	-6.04	107.43	120.10
2	B	101	ILE	O-C-N	6.04	132.35	122.70
1	A	339	TYR	O-C-N	6.03	132.35	122.70
1	A	389	GLN	O-C-N	6.03	132.35	122.70
3	M	90	PHE	O-C-N	6.03	132.35	122.70
1	A	158	LEU	CA-C-O	-6.03	107.43	120.10
1	A	372	ILE	O-C-N	6.03	132.35	122.70
1	A	163	ALA	O-C-N	-6.03	113.06	122.70
1	A	258	LYS	O-C-N	6.03	132.35	122.70
1	A	631	SER	CA-C-O	-6.03	107.44	120.10
2	B	525	ILE	O-C-N	6.03	132.34	122.70
3	M	283	PHE	N-CA-C	-6.03	94.73	111.00
1	A	518	CYS	CA-C-O	-6.02	107.45	120.10
2	B	511	ILE	CA-C-O	6.02	132.75	120.10
1	A	325	SER	CA-C-N	-6.02	103.96	117.20
1	A	620	GLY	CA-C-O	-6.02	109.77	120.60
2	B	64	LYS	O-C-N	6.02	132.33	122.70
2	B	543	GLU	O-C-N	6.02	132.33	122.70
2	B	590	GLN	O-C-N	6.02	132.33	122.70
4	S	151	ALA	O-C-N	6.01	132.32	122.70
1	A	436	CYS	C-N-CA	-6.01	106.67	121.70
2	B	450	VAL	O-C-N	6.01	132.32	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	312	ALA	O-C-N	6.01	132.31	122.70
3	M	120	ARG	O-C-N	6.01	132.31	122.70
2	B	320	SER	O-C-N	6.01	132.31	122.70
2	B	145	MET	CA-C-N	6.00	130.41	117.20
2	B	446	TRP	O-C-N	6.00	132.31	122.70
3	M	367	ALA	N-CA-CB	-6.00	101.70	110.10
1	A	140	VAL	C-N-CA	-6.00	106.70	121.70
2	B	74	ASP	C-N-CA	6.00	136.70	121.70
2	B	510	GLY	O-C-N	6.00	132.30	122.70
1	A	493	ALA	O-C-N	6.00	132.29	122.70
2	B	238	LYS	C-N-CA	5.99	136.68	121.70
2	B	232	ARG	O-C-N	5.99	132.29	122.70
2	B	529	VAL	O-C-N	5.99	132.29	122.70
1	A	129	LYS	O-C-N	5.98	132.27	122.70
3	M	130	GLU	N-CA-C	-5.98	94.84	111.00
4	S	43	ASN	C-N-CA	-5.98	106.74	121.70
1	A	576	MET	O-C-N	5.98	132.27	122.70
2	B	469	ASP	CA-C-O	-5.98	107.55	120.10
1	A	71	VAL	CA-C-O	-5.98	107.55	120.10
2	B	207	VAL	O-C-N	5.98	132.26	122.70
1	A	229	ASN	C-N-CD	5.97	140.94	128.40
1	A	577	VAL	O-C-N	5.97	132.25	122.70
1	A	121	LEU	O-C-N	5.97	132.25	122.70
2	B	569	THR	CA-C-N	-5.97	104.26	116.20
1	A	104	ARG	O-C-N	5.96	132.24	122.70
1	A	296	ASN	CA-C-O	-5.96	107.57	120.10
3	M	121	ILE	O-C-N	5.96	132.24	122.70
1	A	633	PHE	O-C-N	5.96	132.24	122.70
2	B	55	ASN	O-C-N	5.96	132.24	122.70
2	B	112	ASP	N-CA-C	5.96	127.09	111.00
2	B	225	LEU	CA-C-N	-5.96	104.08	117.20
1	A	130	LYS	O-C-N	5.96	132.23	122.70
1	A	553	LEU	O-C-N	5.96	132.23	122.70
1	A	108	LEU	O-C-N	5.96	132.23	122.70
2	B	471	TYR	O-C-N	5.96	132.23	122.70
1	A	486	SER	N-CA-C	-5.95	94.92	111.00
2	B	322	CYS	O-C-N	5.95	132.22	122.70
1	A	490	VAL	O-C-N	5.95	132.22	122.70
2	B	585	GLY	CA-C-O	-5.95	109.89	120.60
1	A	237	SER	CA-C-O	-5.95	107.61	120.10
1	A	425	LYS	O-C-N	5.95	132.22	122.70
1	A	491	THR	O-C-N	5.95	132.22	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	96	LYS	O-C-N	5.95	132.22	122.70
1	A	513	ARG	O-C-N	5.95	132.22	122.70
1	A	538	GLU	C-N-CA	-5.95	106.83	121.70
2	B	333	GLN	N-CA-C	-5.95	94.95	111.00
1	A	214	VAL	O-C-N	5.94	132.21	122.70
2	B	553	ALA	O-C-N	5.94	132.21	122.70
4	S	85	PHE	O-C-N	-5.94	113.19	122.70
2	B	358	MET	O-C-N	5.94	132.20	122.70
1	A	165	ASP	O-C-N	5.93	132.19	122.70
2	B	314	ASN	O-C-N	5.93	132.38	121.10
2	B	131	ASN	O-C-N	5.93	132.19	122.70
1	A	528	ASN	O-C-N	-5.93	113.12	123.20
1	A	625	LEU	C-N-CA	-5.93	106.88	121.70
1	A	154	ILE	C-N-CA	-5.92	106.90	121.70
2	B	530	LEU	O-C-N	5.92	132.17	122.70
2	B	567	GLN	C-N-CA	-5.92	106.91	121.70
2	B	469	ASP	O-C-N	5.91	132.16	122.70
1	A	98	ASN	O-C-N	-5.91	113.24	122.70
3	M	426	LYS	N-CA-C	-5.91	95.04	111.00
1	A	458	ALA	O-C-N	5.90	132.15	122.70
1	A	230	PRO	CA-C-O	-5.90	106.03	120.20
2	B	396	ILE	O-C-N	5.90	132.14	122.70
4	S	159	ALA	O-C-N	5.90	132.13	122.70
1	A	588	LEU	CA-C-O	5.89	132.47	120.10
2	B	371	GLN	O-C-N	5.89	132.12	122.70
2	B	476	ARG	O-C-N	5.89	132.12	122.70
2	B	604	GLU	CA-C-N	-5.89	104.25	117.20
1	A	499	ILE	C-N-CA	-5.88	106.99	121.70
2	B	418	TYR	O-C-N	-5.88	113.29	122.70
2	B	413	LYS	O-C-N	5.88	132.11	122.70
3	M	261	ASN	C-N-CA	-5.88	107.00	121.70
1	A	110	ALA	O-C-N	-5.88	113.30	122.70
1	A	257	LEU	CA-C-O	-5.88	107.75	120.10
1	A	371	ALA	O-C-N	5.88	132.10	122.70
1	A	609	LEU	O-C-N	5.88	132.10	122.70
2	B	247	TYR	O-C-N	5.88	132.10	122.70
1	A	575	LYS	O-C-N	5.87	132.10	122.70
2	B	526	CYS	CA-C-O	-5.87	107.77	120.10
3	M	265	ASN	CA-C-O	-5.87	107.77	120.10
1	A	217	ALA	O-C-N	5.86	132.08	122.70
2	B	33	SER	O-C-N	5.86	132.08	122.70
1	A	600	SER	CA-C-O	-5.86	107.79	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	449	HIS	O-C-N	5.86	132.07	122.70
1	A	276	PRO	O-C-N	-5.86	113.33	122.70
2	B	48	VAL	O-C-N	5.86	132.07	122.70
1	A	512	LEU	O-C-N	5.85	132.07	122.70
2	B	523	PHE	C-N-CA	-5.85	107.07	121.70
2	B	43	ASN	CA-C-O	-5.85	107.82	120.10
3	M	256	VAL	O-C-N	5.84	132.05	122.70
2	B	245	GLN	O-C-N	5.83	132.04	122.70
4	S	98	ILE	CA-C-O	-5.83	107.86	120.10
2	B	60	ARG	O-C-N	5.82	132.02	122.70
1	A	122	MET	O-C-N	5.82	132.01	122.70
2	B	212	VAL	C-N-CA	-5.82	107.15	121.70
2	B	366	LEU	CA-C-O	5.82	132.33	120.10
1	A	494	ASN	O-C-N	5.82	132.01	122.70
1	A	603	VAL	CA-C-O	-5.82	107.88	120.10
4	S	1	MET	CA-C-O	5.82	132.32	120.10
1	A	317	GLU	O-C-N	5.81	132.00	122.70
2	B	176	ALA	O-C-N	5.81	132.00	122.70
2	B	355	ASN	CA-C-O	-5.81	107.89	120.10
1	A	514	GLU	O-C-N	5.81	132.00	122.70
2	B	194	ASP	O-C-N	5.80	131.99	122.70
2	B	444	THR	CA-C-O	5.80	132.29	120.10
2	B	490	ILE	CA-C-O	-5.80	107.92	120.10
2	B	551	LEU	O-C-N	5.80	131.98	122.70
2	B	555	LEU	CA-C-O	-5.80	107.92	120.10
2	B	303	LEU	CA-C-O	-5.80	107.93	120.10
4	S	32	LEU	CA-C-O	-5.79	107.93	120.10
1	A	549	GLU	O-C-N	5.79	131.96	122.70
1	A	64	LEU	C-N-CA	-5.79	107.23	121.70
3	M	306	LEU	C-N-CA	-5.79	107.23	121.70
1	A	545	HIS	CA-C-O	5.79	132.25	120.10
1	A	417	PRO	N-CA-C	-5.78	97.06	112.10
2	B	133	GLU	O-C-N	5.78	131.95	122.70
2	B	189	HIS	O-C-N	5.78	131.95	122.70
1	A	294	SER	O-C-N	5.78	131.95	122.70
2	B	492	LYS	O-C-N	5.78	131.94	122.70
2	B	586	SER	O-C-N	5.78	131.95	122.70
2	B	402	LEU	N-CA-C	5.77	126.58	111.00
1	A	356	ILE	O-C-N	5.77	131.93	122.70
2	B	79	VAL	O-C-N	5.77	131.93	122.70
2	B	412	PHE	CA-C-O	-5.77	107.98	120.10
1	A	152	THR	C-N-CA	-5.77	107.28	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	495	ASP	O-C-N	5.77	131.93	122.70
1	A	268	PRO	O-C-N	5.76	131.92	122.70
3	M	421	GLY	CA-C-O	5.76	130.97	120.60
1	A	275	LEU	O-C-N	-5.75	110.17	121.10
2	B	457	HIS	O-C-N	-5.75	113.50	122.70
1	A	350	SER	C-N-CA	-5.75	107.32	121.70
2	B	86	VAL	O-C-N	5.75	131.90	122.70
3	M	225	VAL	O-C-N	5.75	131.90	122.70
1	A	582	ILE	CA-C-O	-5.75	108.03	120.10
2	B	394	TRP	O-C-N	5.75	131.89	122.70
3	M	292	PRO	C-N-CD	-5.74	107.97	120.60
2	B	99	ARG	CA-C-O	-5.74	108.04	120.10
2	B	120	ILE	CA-C-O	-5.74	108.05	120.10
2	B	532	ARG	O-C-N	5.74	131.88	122.70
1	A	430	ASN	O-C-N	5.74	131.88	122.70
1	A	109	ALA	O-C-N	5.74	131.88	122.70
1	A	217	ALA	CA-C-O	-5.74	108.06	120.10
4	S	96	LEU	CA-C-O	-5.73	108.06	120.10
1	A	173	THR	C-N-CA	5.73	136.02	121.70
2	B	423	HIS	N-CA-C	-5.73	95.54	111.00
1	A	543	TYR	N-CA-C	5.73	126.46	111.00
1	A	126	ASN	O-C-N	5.72	131.85	122.70
2	B	241	ASP	O-C-N	5.72	131.84	122.70
3	M	85	GLY	CA-C-O	-5.72	110.31	120.60
3	M	447	ILE	CA-C-O	5.72	132.10	120.10
1	A	413	SER	C-N-CA	-5.71	107.41	121.70
2	B	174	ALA	CA-C-O	5.71	132.10	120.10
2	B	204	ASP	CA-C-O	-5.71	108.10	120.10
2	B	122	SER	O-C-N	5.71	131.84	122.70
1	A	422	GLU	CA-C-O	-5.71	108.11	120.10
2	B	434	LYS	O-C-N	5.71	131.83	122.70
4	S	152	SER	O-C-N	5.70	131.83	122.70
2	B	393	ILE	O-C-N	5.70	131.82	122.70
4	S	74	GLN	CA-C-O	5.70	132.07	120.10
1	A	123	LEU	O-C-N	5.70	131.81	122.70
1	A	108	LEU	CA-C-O	-5.69	108.14	120.10
2	B	318	ILE	CA-C-O	-5.69	108.15	120.10
2	B	341	GLU	CA-C-O	-5.69	108.15	120.10
2	B	392	SER	O-C-N	5.69	131.81	122.70
2	B	265	VAL	CA-C-N	-5.69	104.68	117.20
4	S	164	ASP	O-C-N	-5.69	113.60	122.70
1	A	325	SER	C-N-CA	5.69	135.92	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	300	ASP	O-C-N	5.69	131.80	122.70
2	B	593	ASN	CA-C-O	-5.69	108.16	120.10
1	A	75	THR	O-C-N	5.69	131.80	122.70
2	B	211	ALA	CA-C-O	-5.68	108.17	120.10
4	S	158	LYS	CA-C-O	-5.68	108.17	120.10
2	B	173	VAL	O-C-N	5.68	131.79	122.70
1	A	106	GLY	CA-C-O	-5.68	110.38	120.60
1	A	491	THR	CA-C-O	-5.68	108.18	120.10
1	A	558	VAL	CA-C-O	-5.67	108.18	120.10
1	A	514	GLU	CA-C-O	-5.67	108.19	120.10
2	B	438	ARG	O-C-N	5.67	131.78	122.70
2	B	436	LEU	CA-C-O	-5.67	108.19	120.10
3	M	88	ASP	O-C-N	5.67	131.77	122.70
1	A	456	ASP	O-C-N	5.67	131.77	122.70
2	B	448	SER	O-C-N	5.67	131.77	122.70
2	B	322	CYS	CA-C-O	-5.67	108.20	120.10
3	M	306	LEU	CA-C-O	5.67	132.00	120.10
1	A	155	THR	O-C-N	5.67	131.86	121.10
2	B	22	ALA	N-CA-C	-5.67	95.70	111.00
4	S	84	TYR	CA-C-O	5.67	132.00	120.10
1	A	478	ARG	O-C-N	5.66	131.76	122.70
3	M	237	THR	O-C-N	-5.66	113.58	123.20
3	M	398	ILE	O-C-N	-5.66	113.64	122.70
1	A	293	GLU	O-C-N	5.66	131.75	122.70
2	B	545	ARG	O-C-N	5.66	131.75	122.70
2	B	318	ILE	O-C-N	5.66	131.75	122.70
3	M	396	GLN	O-C-N	-5.66	113.65	122.70
1	A	253	ILE	O-C-N	5.65	131.75	122.70
2	B	189	HIS	CA-C-O	-5.65	108.23	120.10
1	A	452	ALA	O-C-N	5.65	131.74	122.70
2	B	383	VAL	CA-C-N	-5.65	104.77	117.20
2	B	456	ASP	O-C-N	5.65	131.74	122.70
3	M	284	SER	C-N-CD	5.65	140.27	128.40
4	S	154	ASP	O-C-N	5.65	131.74	122.70
1	A	448	GLU	C-N-CA	5.65	135.82	121.70
1	A	479	ASN	O-C-N	5.65	131.74	122.70
1	A	572	PHE	C-N-CA	5.65	135.82	121.70
4	S	137	GLN	CA-C-O	5.65	131.96	120.10
3	M	87	LEU	O-C-N	5.64	131.73	122.70
2	B	274	PRO	N-CA-C	-5.64	97.43	112.10
2	B	391	ALA	O-C-N	5.64	131.73	122.70
1	A	233	PHE	CA-C-O	5.64	131.95	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	M	421	GLY	C-N-CD	5.64	140.25	128.40
1	A	230	PRO	CA-N-CD	5.63	119.59	111.70
2	B	239	GLN	C-N-CA	5.63	135.78	121.70
1	A	88	ASN	O-C-N	-5.63	113.69	122.70
3	M	385	ARG	CA-C-O	5.63	131.93	120.10
1	A	241	TYR	O-C-N	5.63	131.71	122.70
2	B	407	ASN	O-C-N	5.63	131.71	122.70
3	M	230	LYS	O-C-N	-5.63	113.70	122.70
2	B	472	VAL	C-N-CA	-5.62	107.64	121.70
2	B	172	GLU	C-N-CA	-5.62	107.64	121.70
1	A	224	GLU	C-N-CA	-5.62	107.65	121.70
1	A	145	ILE	CA-C-O	-5.62	108.30	120.10
1	A	265	GLN	C-N-CA	5.62	135.75	121.70
2	B	117	LEU	O-C-N	5.62	131.69	122.70
3	M	444	ALA	N-CA-CB	-5.62	102.24	110.10
1	A	575	LYS	CA-C-O	-5.60	108.33	120.10
2	B	303	LEU	O-C-N	5.60	131.66	122.70
1	A	90	HIS	O-C-N	5.59	131.65	122.70
1	A	551	LEU	CA-C-O	-5.59	108.35	120.10
1	A	387	ILE	CA-C-O	-5.59	108.36	120.10
4	S	34	GLN	O-C-N	5.59	131.65	122.70
2	B	426	GLU	O-C-N	5.59	131.64	122.70
2	B	215	TYR	CA-C-O	-5.58	108.37	120.10
2	B	496	LEU	O-C-N	5.58	131.64	122.70
1	A	139	ASP	CA-C-O	-5.58	108.38	120.10
1	A	630	PRO	O-C-N	5.58	131.63	122.70
1	A	184	ALA	O-C-N	5.58	131.62	122.70
2	B	138	ALA	CA-C-O	-5.58	108.38	120.10
2	B	302	PHE	CA-C-O	-5.58	108.39	120.10
2	B	387	ASP	N-CA-C	5.58	126.06	111.00
3	M	384	GLY	O-C-N	5.57	131.62	122.70
2	B	155	LEU	CA-C-O	-5.57	108.41	120.10
1	A	334	SER	O-C-N	5.57	131.61	122.70
2	B	605	PHE	CA-C-O	-5.57	108.41	120.10
1	A	380	ASP	O-C-N	5.56	131.60	122.70
1	A	635	ALA	CA-C-O	-5.56	108.42	120.10
3	M	368	ASP	O-C-N	5.56	131.67	121.10
1	A	489	GLU	O-C-N	5.56	131.59	122.70
1	A	135	ASP	CA-C-N	5.55	127.31	116.20
1	A	137	ASN	C-N-CA	-5.55	107.81	121.70
1	A	319	LEU	CA-C-O	5.55	131.76	120.10
1	A	177	ILE	CA-C-O	-5.55	108.44	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	352	PHE	O-C-N	5.55	131.58	122.70
2	B	493	LEU	CA-C-O	-5.55	108.45	120.10
2	B	540	GLU	N-CA-C	5.55	125.98	111.00
1	A	193	PRO	CA-C-O	-5.54	106.89	120.20
2	B	569	THR	C-N-CA	5.54	133.94	122.30
1	A	222	ILE	O-C-N	5.54	131.57	122.70
1	A	156	PRO	CA-C-O	5.54	133.49	120.20
3	M	387	GLU	O-C-N	-5.54	113.84	122.70
4	S	86	THR	O-C-N	-5.54	113.84	122.70
1	A	578	LEU	CA-C-O	-5.54	108.47	120.10
1	A	416	ILE	C-N-CA	-5.53	98.76	122.00
2	B	415	LEU	CA-C-O	-5.53	108.48	120.10
4	S	109	LEU	C-N-CA	-5.53	107.87	121.70
2	B	522	GLU	N-CA-C	5.53	125.93	111.00
1	A	216	SER	O-C-N	5.53	131.54	122.70
1	A	254	ILE	O-C-N	5.53	131.54	122.70
3	M	46	SER	O-C-N	-5.53	113.86	122.70
2	B	477	MET	O-C-N	5.53	131.54	122.70
4	S	87	PHE	O-C-N	-5.53	113.86	122.70
1	A	80	TYR	CB-CA-C	5.52	121.45	110.40
1	A	487	MET	C-N-CA	-5.52	107.89	121.70
1	A	513	ARG	CA-C-O	-5.52	108.50	120.10
4	S	87	PHE	CA-C-O	5.52	131.70	120.10
1	A	583	GLU	O-C-N	5.52	131.53	122.70
1	A	72	LEU	O-C-N	5.52	131.53	122.70
1	A	341	ILE	O-C-N	5.52	132.58	123.20
1	A	251	TRP	O-C-N	5.52	131.53	122.70
2	B	457	HIS	C-N-CA	-5.51	107.91	121.70
1	A	333	ILE	CA-C-O	-5.51	108.52	120.10
1	A	419	ILE	CA-C-N	-5.51	105.07	117.20
1	A	229	ASN	C-N-CA	-5.51	98.87	122.00
2	B	73	ASP	C-N-CA	-5.51	107.94	121.70
1	A	501	ASN	C-N-CA	5.50	135.46	121.70
2	B	156	HIS	O-C-N	5.50	131.51	122.70
2	B	68	SER	O-C-N	5.50	131.51	122.70
1	A	265	GLN	CB-CA-C	5.50	121.40	110.40
1	A	63	ASP	N-CA-C	5.50	125.85	111.00
1	A	426	ILE	O-C-N	5.50	131.50	122.70
2	B	340	ILE	O-C-N	5.49	131.49	122.70
3	M	117	ASN	O-C-N	5.49	131.49	122.70
4	S	131	VAL	O-C-N	5.49	131.49	122.70
1	A	292	TYR	O-C-N	5.49	131.49	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	290	SER	N-CA-C	5.49	125.81	111.00
2	B	345	ARG	O-C-N	5.48	131.47	122.70
1	A	191	GLN	C-N-CA	5.48	135.40	121.70
1	A	333	ILE	O-C-N	5.48	131.47	122.70
1	A	580	GLU	O-C-N	5.48	131.47	122.70
2	B	245	GLN	CA-C-O	-5.48	108.59	120.10
2	B	458	MET	C-N-CA	-5.48	108.00	121.70
4	S	78	LYS	CA-C-O	5.48	131.61	120.10
2	B	36	THR	CA-C-O	-5.48	108.59	120.10
1	A	188	VAL	CA-C-O	-5.48	108.60	120.10
2	B	547	GLN	O-C-N	5.48	131.46	122.70
3	M	74	TYR	CA-C-N	-5.48	105.15	117.20
1	A	441	TYR	N-CA-C	-5.47	96.22	111.00
2	B	109	ALA	CB-CA-C	-5.47	101.89	110.10
2	B	531	ARG	CA-C-O	-5.47	108.61	120.10
4	S	138	GLY	C-N-CA	5.47	133.78	122.30
3	M	296	LYS	C-N-CA	-5.47	108.03	121.70
2	B	475	ILE	CA-C-O	-5.47	108.62	120.10
2	B	431	MET	O-C-N	5.46	131.44	122.70
2	B	126	SER	CA-C-O	5.46	131.57	120.10
2	B	365	PHE	O-C-N	5.46	131.43	122.70
2	B	36	THR	O-C-N	5.45	131.42	122.70
3	M	267	ILE	C-N-CA	-5.45	110.86	122.30
1	A	175	PRO	O-C-N	5.45	131.41	122.70
2	B	491	PHE	O-C-N	5.45	131.41	122.70
2	B	127	LEU	C-N-CA	-5.44	108.09	121.70
1	A	106	GLY	O-C-N	5.44	131.41	122.70
1	A	392	MET	CA-C-O	-5.44	108.67	120.10
1	A	255	ARG	O-C-N	5.44	131.41	122.70
1	A	634	ASN	O-C-N	5.44	131.40	122.70
2	B	388	PRO	O-C-N	5.44	131.40	122.70
3	M	469	GLY	C-N-CA	-5.44	108.10	121.70
1	A	91	ILE	CA-C-O	-5.44	108.68	120.10
1	A	427	LYS	O-C-N	5.44	131.40	122.70
2	B	29	LYS	C-N-CA	-5.43	108.11	121.70
1	A	71	VAL	O-C-N	5.43	131.39	122.70
2	B	586	SER	CA-C-O	-5.43	108.70	120.10
1	A	316	LEU	CA-C-O	-5.42	108.71	120.10
2	B	67	ILE	CA-C-O	-5.42	108.71	120.10
1	A	305	GLU	CA-C-N	5.42	129.13	117.20
1	A	611	LEU	CA-C-O	-5.42	108.72	120.10
3	M	354	ASP	N-CA-CB	5.42	120.35	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	313	MET	CA-C-O	-5.42	108.72	120.10
2	B	248	LEU	CA-C-O	-5.42	108.72	120.10
1	A	160	ARG	O-C-N	5.42	131.36	122.70
1	A	433	ILE	CA-C-O	-5.41	108.74	120.10
2	B	482	ASN	O-C-N	5.41	131.38	121.10
1	A	308	ASP	N-CA-C	5.41	125.60	111.00
3	M	307	SER	O-C-N	5.41	131.35	122.70
1	A	397	ASP	O-C-N	-5.41	114.05	122.70
1	A	593	THR	O-C-N	5.41	131.35	122.70
2	B	94	ASP	O-C-N	5.41	131.35	122.70
2	B	430	ILE	CA-C-O	-5.41	108.75	120.10
1	A	467	LYS	C-N-CA	-5.40	108.20	121.70
3	M	16	PHE	O-C-N	-5.40	114.06	122.70
2	B	609	ASP	O-C-N	5.40	131.34	122.70
2	B	439	CYS	CA-C-O	-5.40	108.77	120.10
1	A	136	GLY	CA-C-N	5.39	129.06	117.20
2	B	213	LEU	CA-C-O	-5.39	108.78	120.10
2	B	360	LEU	CA-C-O	-5.39	108.78	120.10
2	B	120	ILE	O-C-N	5.39	131.32	122.70
1	A	455	MET	O-C-N	5.39	131.32	122.70
2	B	363	ILE	CA-C-O	-5.39	108.79	120.10
1	A	368	ARG	O-C-N	5.38	131.31	122.70
1	A	572	PHE	O-C-N	5.38	131.30	122.70
1	A	604	LEU	CA-C-O	-5.37	108.82	120.10
1	A	566	PHE	O-C-N	5.37	131.30	122.70
1	A	226	SER	C-N-CA	-5.37	108.28	121.70
1	A	390	THR	CA-C-O	-5.37	108.83	120.10
1	A	624	LEU	CA-C-O	-5.37	108.83	120.10
2	B	544	THR	CA-C-O	-5.37	108.83	120.10
4	S	76	ILE	O-C-N	-5.37	114.11	122.70
4	S	125	TRP	O-C-N	5.37	131.28	122.70
2	B	577	ASN	CA-C-N	5.36	132.11	117.10
1	A	105	VAL	CA-C-O	-5.36	108.85	120.10
1	A	216	SER	CA-C-O	-5.36	108.85	120.10
1	A	391	LEU	CA-C-O	5.35	131.34	120.10
1	A	466	ASP	N-CA-C	-5.35	96.55	111.00
1	A	180	LYS	O-C-N	5.35	131.26	122.70
1	A	181	ALA	O-C-N	5.35	131.25	122.70
2	B	219	TYR	N-CA-C	5.34	125.43	111.00
1	A	559	PHE	CA-C-O	-5.34	108.88	120.10
1	A	518	CYS	O-C-N	5.34	131.24	122.70
1	A	172	SER	C-N-CA	-5.33	108.37	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	277	LYS	CA-C-N	5.33	128.93	117.20
2	B	364	HIS	O-C-N	5.33	131.23	122.70
3	M	25	PRO	N-CA-C	-5.33	98.25	112.10
2	B	61	ASP	O-C-N	5.33	131.22	122.70
2	B	178	ILE	CA-C-O	-5.33	108.91	120.10
1	A	312	ALA	CA-C-O	-5.32	108.92	120.10
1	A	434	SER	CA-C-O	-5.32	108.94	120.10
1	A	493	ALA	CA-C-O	-5.32	108.94	120.10
2	B	132	SER	CA-C-O	5.32	131.26	120.10
2	B	595	VAL	CA-C-O	-5.32	108.94	120.10
1	A	345	ASN	O-C-N	5.31	131.20	122.70
1	A	314	ALA	O-C-N	5.31	131.20	122.70
2	B	354	GLY	N-CA-C	-5.31	99.83	113.10
2	B	169	VAL	CA-C-O	-5.30	108.97	120.10
2	B	196	LEU	CA-C-O	-5.30	108.97	120.10
4	S	18	LYS	CA-C-O	5.30	131.22	120.10
1	A	110	ALA	CA-C-O	5.29	131.22	120.10
1	A	115	TYR	C-N-CA	5.29	134.93	121.70
4	S	95	GLU	CA-C-O	-5.29	108.99	120.10
2	B	616	SER	O-C-N	5.29	131.16	122.70
3	M	55	MET	O-C-N	-5.29	114.24	122.70
1	A	237	SER	O-C-N	5.28	131.14	121.10
1	A	335	CYS	CA-C-O	-5.28	109.01	120.10
2	B	357	GLU	O-C-N	5.28	131.15	122.70
2	B	467	VAL	CA-C-O	-5.28	109.00	120.10
1	A	244	LEU	CA-C-O	5.28	131.19	120.10
1	A	470	GLY	O-C-N	5.28	131.15	122.70
3	M	309	GLN	O-C-N	5.28	131.15	122.70
1	A	505	ASN	O-C-N	-5.28	114.25	122.70
3	M	284	SER	CA-C-O	5.28	131.18	120.10
1	A	382	ASP	O-C-N	5.28	131.14	122.70
1	A	398	GLU	N-CA-C	5.28	125.24	111.00
2	B	451	MET	CA-C-O	-5.27	109.03	120.10
2	B	188	TYR	O-C-N	5.27	131.13	122.70
3	M	1	MET	C-N-CA	-5.27	108.53	121.70
2	B	216	LYS	CA-C-O	5.27	131.16	120.10
1	A	125	THR	CA-C-O	-5.26	109.04	120.10
2	B	54	ARG	N-CA-C	5.26	125.21	111.00
2	B	533	LEU	O-C-N	5.26	131.12	122.70
2	B	222	HIS	N-CA-C	5.26	125.21	111.00
2	B	443	SER	CA-C-N	-5.26	105.63	117.20
2	B	323	ASN	CA-C-O	-5.26	109.06	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	437	SER	N-CA-C	5.25	125.19	111.00
2	B	588	ILE	CA-C-O	-5.25	109.07	120.10
1	A	370	LYS	CA-C-O	-5.25	109.07	120.10
1	A	528	ASN	C-N-CA	-5.25	111.27	122.30
2	B	346	THR	CA-C-O	-5.25	109.07	120.10
1	A	242	GLU	CA-C-O	-5.24	109.09	120.10
1	A	496	ILE	CA-C-O	-5.24	109.09	120.10
1	A	163	ALA	CA-C-O	5.24	131.10	120.10
2	B	554	LYS	CA-C-O	-5.24	109.10	120.10
2	B	589	SER	CA-C-O	-5.24	109.09	120.10
2	B	319	LEU	CA-C-O	-5.24	109.10	120.10
2	B	439	CYS	O-C-N	5.24	132.11	123.20
2	B	192	LEU	CA-C-O	-5.24	109.11	120.10
4	S	78	LYS	O-C-N	-5.24	114.32	122.70
1	A	516	ILE	CA-C-O	-5.23	109.11	120.10
2	B	63	MET	CA-C-O	-5.22	109.13	120.10
3	M	331	LEU	CA-C-N	-5.22	105.75	116.20
1	A	526	VAL	C-N-CA	5.22	134.75	121.70
1	A	327	ASP	O-C-N	5.22	131.01	121.10
1	A	85	ALA	O-C-N	5.21	131.03	122.70
2	B	118	LEU	O-C-N	5.21	131.03	122.70
2	B	489	ILE	CA-C-O	-5.21	109.16	120.10
1	A	341	ILE	CA-C-O	-5.21	109.17	120.10
1	A	562	TRP	CA-C-O	-5.20	109.17	120.10
3	M	90	PHE	CA-C-O	-5.20	109.19	120.10
3	M	323	MET	C-N-CA	-5.19	108.71	121.70
1	A	548	GLN	O-C-N	5.19	131.01	122.70
1	A	634	ASN	CA-C-O	-5.19	109.19	120.10
2	B	494	ALA	CA-C-O	-5.19	109.20	120.10
2	B	411	ILE	CA-C-O	-5.19	109.20	120.10
1	A	330	LEU	CA-C-O	-5.19	109.21	120.10
2	B	503	LEU	CA-C-N	-5.19	105.79	117.20
2	B	223	LEU	CA-C-O	5.18	130.99	120.10
1	A	146	ALA	CA-C-O	-5.18	109.22	120.10
1	A	278	ILE	O-C-N	-5.18	114.42	122.70
3	M	122	SER	CA-C-O	-5.17	109.24	120.10
1	A	370	LYS	O-C-N	5.17	130.98	122.70
4	S	141	VAL	N-CA-C	-5.17	97.03	111.00
1	A	178	ARG	O-C-N	5.17	130.97	122.70
1	A	253	ILE	CA-C-O	-5.17	109.25	120.10
3	M	354	ASP	CB-CA-C	5.17	120.73	110.40
2	B	399	LEU	O-C-N	-5.16	114.44	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	B	551	LEU	CA-C-O	-5.16	109.26	120.10
2	B	112	ASP	CA-C-O	-5.16	109.26	120.10
2	B	175	LEU	O-C-N	5.16	130.96	122.70
3	M	117	ASN	CA-C-O	-5.16	109.26	120.10
1	A	134	TYR	C-N-CA	-5.16	108.80	121.70
1	A	221	VAL	CA-C-O	-5.16	109.27	120.10
4	S	103	GLN	CA-C-O	-5.15	109.28	120.10
1	A	474	GLY	CA-C-O	-5.15	111.32	120.60
1	A	533	ILE	O-C-N	5.15	130.94	122.70
2	B	166	SER	C-N-CA	5.15	134.58	121.70
2	B	304	GLN	O-C-N	5.15	130.94	122.70
2	B	350	THR	N-CA-C	5.15	124.91	111.00
1	A	94	VAL	CA-C-O	5.15	130.91	120.10
1	A	174	ARG	C-N-CD	-5.15	109.28	120.60
1	A	239	LEU	C-N-CA	-5.15	108.83	121.70
2	B	566	ALA	N-CA-CB	-5.15	102.89	110.10
1	A	141	VAL	CA-C-O	-5.14	109.30	120.10
2	B	577	ASN	N-CA-CB	-5.14	101.34	110.60
1	A	597	GLN	O-C-N	5.14	130.92	122.70
2	B	470	ALA	O-C-N	5.14	130.92	122.70
2	B	280	PRO	N-CA-C	-5.13	98.75	112.10
1	A	380	ASP	CA-C-O	-5.13	109.32	120.10
2	B	608	ARG	O-C-N	5.13	130.91	122.70
2	B	84	ALA	O-C-N	5.13	130.91	122.70
4	S	154	ASP	CA-C-O	-5.13	109.33	120.10
1	A	555	LEU	CA-C-O	-5.13	109.33	120.10
2	B	134	LEU	CA-C-O	-5.13	109.33	120.10
2	B	401	THR	C-N-CA	5.13	134.52	121.70
2	B	530	LEU	CA-C-O	-5.12	109.34	120.10
2	B	592	TYR	CA-C-O	-5.12	109.34	120.10
4	S	164	ASP	CA-C-N	5.12	128.47	117.20
2	B	391	ALA	CA-C-O	-5.12	109.36	120.10
2	B	500	GLN	N-CA-C	5.12	124.81	111.00
1	A	174	ARG	O-C-N	5.11	130.82	121.10
2	B	421	SER	N-CA-C	5.11	124.81	111.00
1	A	230	PRO	O-C-N	5.11	130.87	122.70
2	B	615	SER	CA-C-O	-5.11	109.38	120.10
4	S	89	VAL	C-N-CA	-5.11	108.93	121.70
4	S	54	PRO	CA-C-N	5.10	131.39	117.10
2	B	116	THR	CA-C-O	-5.10	109.38	120.10
2	B	230	PHE	CA-C-O	5.10	130.80	120.10
2	B	492	LYS	CA-C-O	-5.09	109.40	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	302	ASN	N-CA-CB	5.09	119.77	110.60
2	B	365	PHE	CA-C-O	-5.09	109.41	120.10
1	A	121	LEU	CA-C-O	-5.08	109.43	120.10
2	B	590	GLN	CA-C-O	-5.08	109.43	120.10
1	A	148	SER	O-C-N	5.08	131.83	123.20
1	A	103	LYS	C-N-CA	-5.07	109.02	121.70
1	A	280	GLU	O-C-N	5.07	130.82	122.70
1	A	445	ASN	N-CA-C	-5.07	97.30	111.00
1	A	599	ARG	O-C-N	5.07	130.82	122.70
3	M	39	PRO	O-C-N	5.07	130.82	122.70
1	A	305	GLU	CA-C-O	-5.07	109.45	120.10
1	A	628	VAL	CA-C-O	-5.07	109.45	120.10
1	A	193	PRO	O-C-N	5.07	130.81	122.70
1	A	254	ILE	CA-C-O	-5.07	109.46	120.10
1	A	512	LEU	CA-C-O	-5.07	109.46	120.10
2	B	340	ILE	CA-C-O	-5.07	109.46	120.10
2	B	562	ASN	O-C-N	5.07	130.80	122.70
1	A	179	LYS	O-C-N	5.06	130.80	122.70
2	B	465	ALA	O-C-N	5.06	130.79	122.70
1	A	218	ALA	CA-C-O	-5.06	109.48	120.10
1	A	338	PHE	CA-C-O	-5.06	109.48	120.10
1	A	199	ASN	N-CA-C	5.05	124.64	111.00
1	A	337	LEU	CA-C-O	-5.05	109.49	120.10
1	A	422	GLU	O-C-N	5.05	130.78	122.70
1	A	502	ASP	O-C-N	5.05	130.77	122.70
2	B	426	GLU	CA-C-O	-5.05	109.50	120.10
2	B	548	ILE	CA-C-O	-5.04	109.51	120.10
1	A	113	SER	N-CA-C	5.04	124.60	111.00
1	A	252	ILE	CA-C-O	-5.04	109.52	120.10
1	A	602	GLU	CA-C-O	-5.04	109.52	120.10
3	M	316	ARG	C-N-CA	-5.04	109.10	121.70
1	A	73	LYS	O-C-N	5.04	130.76	122.70
1	A	383	ASN	CA-C-O	-5.04	109.53	120.10
2	B	86	VAL	CA-C-O	-5.04	109.53	120.10
3	M	57	GLY	O-C-N	-5.03	114.65	122.70
2	B	119	SER	CA-C-O	-5.03	109.54	120.10
3	M	234	ARG	O-C-N	-5.03	114.66	122.70
1	A	620	GLY	O-C-N	5.02	130.74	122.70
1	A	77	LEU	O-C-N	5.02	130.73	122.70
2	B	210	CYS	CA-C-O	-5.02	109.56	120.10
2	B	69	ILE	C-N-CA	-5.02	109.16	121.70
4	S	36	TYR	CA-C-O	-5.02	109.57	120.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	90	HIS	CA-C-O	-5.01	109.57	120.10
2	B	34	SER	O-C-N	5.01	130.72	122.70
2	B	60	ARG	CA-C-O	-5.01	109.58	120.10
2	B	157	THR	CA-C-O	-5.01	109.58	120.10
4	S	12	CYS	C-N-CA	-5.01	109.18	121.70
1	A	323	CYS	O-C-N	-5.01	114.69	122.70
4	S	132	LEU	CA-C-O	-5.01	109.58	120.10
1	A	185	LEU	CA-C-O	-5.00	109.59	120.10
1	A	469	LEU	CA-C-N	-5.00	106.19	116.20
4	S	157	ASN	CA-C-O	-5.00	109.59	120.10
1	A	85	ALA	CA-C-N	-5.00	106.19	117.20
2	B	482	ASN	CA-C-O	-5.00	109.59	120.10
1	A	431	VAL	CA-C-O	-5.00	109.60	120.10
1	A	560	SER	O-C-N	5.00	130.70	122.70

There are no chirality outliers.

All (99) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	117	ASP	Mainchain
1	A	174	ARG	Mainchain
1	A	192	TYR	Mainchain
1	A	199	ASN	Mainchain
1	A	204	VAL	Mainchain
1	A	219	VAL	Mainchain
1	A	233	PHE	Mainchain
1	A	240	LEU	Mainchain
1	A	244	LEU	Mainchain
1	A	260	PHE	Mainchain
1	A	275	LEU	Mainchain
1	A	277	LYS	Mainchain
1	A	278	ILE	Mainchain
1	A	282	MET	Mainchain
1	A	298	ILE	Mainchain
1	A	302	ASN	Mainchain
1	A	306	GLU	Mainchain
1	A	319	LEU	Mainchain
1	A	320	HIS	Mainchain
1	A	323	CYS	Mainchain
1	A	325	SER	Mainchain
1	A	328	PRO	Mainchain
1	A	350	SER	Mainchain

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Mol	Chain	Res	Type	Group
1	A	399	ASP	Peptide
1	A	400	VAL	Peptide
1	A	441	TYR	Mainchain
1	A	462	GLN	Mainchain
1	A	487	MET	Mainchain
1	A	500	SER	Mainchain
1	A	501	ASN	Mainchain
1	A	505	ASN	Mainchain
1	A	506	LYS	Mainchain
1	A	527	GLU	Mainchain
1	A	528	ASN	Mainchain
1	A	529	GLY	Mainchain
1	A	531	ASP	Mainchain
1	A	535	ILE	Mainchain
1	A	536	MET	Mainchain
1	A	538	GLU	Mainchain
1	A	539	ASN	Mainchain
1	A	565	ASN	Mainchain
1	A	569	ASP	Mainchain
1	A	571	ARG	Mainchain
1	A	586	GLU	Mainchain
1	A	588	LEU	Mainchain
1	A	64	LEU	Mainchain
1	A	80	TYR	Mainchain
1	A	84	MET	Mainchain
1	A	94	VAL	Mainchain
2	B	126	SER	Mainchain
2	B	142	LEU	Mainchain
2	B	147	MET	Mainchain
2	B	237	ILE	Mainchain
2	B	267	ASP	Mainchain
2	B	278	PRO	Peptide
2	B	288	TYR	Mainchain
2	B	326	TYR	Mainchain
2	B	375	LEU	Mainchain
2	B	377	TYR	Mainchain
2	B	381	PHE	Mainchain
2	B	384	PHE	Mainchain
2	B	404	ASN	Mainchain
2	B	41	ASN	Peptide
2	B	444	THR	Mainchain
2	B	459	GLU	Mainchain

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Mol	Chain	Res	Type	Group
2	B	485	LYS	Mainchain
2	B	497	LEU	Mainchain
2	B	523	PHE	Mainchain
2	B	536	ASN	Mainchain
2	B	557	SER	Mainchain
2	B	56	SER	Mainchain
2	B	565	GLN	Mainchain
2	B	569	THR	Mainchain
2	B	573	GLU	Mainchain
2	B	584	SER	Mainchain
2	B	78	ASP	Mainchain
2	B	82	TYR	Mainchain
3	M	265	ASN	Mainchain
3	M	284	SER	Mainchain
3	M	292	PRO	Peptide
3	M	40	GLN	Mainchain
3	M	421	GLY	Mainchain
3	M	426	LYS	Peptide
3	M	445	SER	Peptide
3	M	45	SER	Peptide
3	M	456	SER	Mainchain
3	M	462	LYS	Peptide
3	M	477	GLY	Peptide
3	M	57	GLY	Mainchain
3	M	8	THR	Mainchain
3	M	85	GLY	Mainchain
4	S	101	LEU	Mainchain
4	S	102	ILE	Mainchain
4	S	135	ILE	Mainchain
4	S	161	GLU	Mainchain
4	S	163	THR	Peptide
4	S	167	ILE	Peptide
4	S	43	ASN	Mainchain
4	S	53	THR	Mainchain

## 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	A	4625	0	4688	2296	0
2	B	4954	0	4969	3353	0
3	M	3106	0	3081	1519	0
4	S	1356	0	1329	910	0
All	All	14041	0	14067	7266	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 258.

All (7266) 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)
4:S:8:PHE:CE1	4:S:84:TYR:HB2	1.15	1.68
2:B:171:GLY:CA	2:B:207:VAL:HG13	1.22	1.64
1:A:140:VAL:HA	1:A:177:ILE:CG1	1.17	1.64
1:A:102:GLN:CG	4:S:166:LYS:H	1.11	1.63
1:A:633:PHE:CE1	2:B:513:TRP:CE3	1.74	1.63
2:B:127:LEU:HD13	2:B:157:THR:CG2	1.14	1.62
2:B:219:TYR:HB3	2:B:223:LEU:CD2	1.19	1.61
1:A:217:ALA:CA	4:S:142:ILE:HB	1.14	1.61
3:M:104:PHE:CE1	3:M:113:LYS:HE2	1.25	1.61
2:B:123:LEU:CD1	2:B:142:LEU:HG	1.29	1.61
2:B:252:LEU:HD13	2:B:302:PHE:CD1	1.15	1.61
4:S:5:VAL:HG21	4:S:132:LEU:CD2	1.26	1.61
1:A:288:THR:HG21	1:A:322:PHE:CZ	1.25	1.60
2:B:216:LYS:CB	2:B:251:LEU:HD13	1.19	1.60
4:S:8:PHE:CE1	4:S:84:TYR:CB	1.77	1.60
2:B:83:PHE:CZ	2:B:119:SER:HB3	1.14	1.60
2:B:260:LEU:HB3	2:B:291:TYR:CE1	1.14	1.60
3:M:449:VAL:CG1	3:M:452:ILE:HD11	1.26	1.60
2:B:556:LEU:HA	2:B:588:ILE:CD1	1.27	1.59
2:B:219:TYR:CE1	2:B:226:LEU:CB	1.80	1.59
2:B:260:LEU:CB	2:B:291:TYR:HE1	1.14	1.59
3:M:104:PHE:CE1	3:M:113:LYS:CE	1.74	1.59
2:B:83:PHE:CE2	2:B:119:SER:HB3	1.09	1.59
2:B:139:LEU:HD23	2:B:173:VAL:CA	1.23	1.59
1:A:557:LYS:CE	2:B:606:ASP:H	1.06	1.58
2:B:433:VAL:HG12	2:B:474:VAL:CG2	1.24	1.58
3:M:215:TYR:CD1	3:M:468:LYS:HA	1.35	1.58
1:A:105:VAL:HG23	4:S:167:ILE:CD1	1.11	1.58
1:A:105:VAL:H	4:S:167:ILE:CD1	1.13	1.58
3:M:69:ILE:CD1	3:M:90:PHE:HZ	1.10	1.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:244:VAL:HA	3:M:472:TYR:CD2	1.13	1.57
2:B:316:THR:HG21	3:M:90:PHE:CE2	1.38	1.57
2:B:546:CYS:HB2	2:B:607:ILE:CG1	1.20	1.57
3:M:104:PHE:CZ	3:M:113:LYS:CE	1.80	1.57
3:M:379:LEU:HD22	3:M:386:PHE:CD1	1.36	1.57
2:B:2:VAL:CG1	2:B:6:HIS:NE2	1.67	1.57
2:B:319:LEU:CD1	2:B:358:MET:CG	1.80	1.57
2:B:523:PHE:HZ	2:B:580:TYR:CD2	1.21	1.57
1:A:605:GLU:HG3	1:A:632:PHE:CD2	1.37	1.57
2:B:291:TYR:HD2	2:B:294:VAL:CG1	1.18	1.57
2:B:549:LEU:HD21	2:B:611:ALA:CA	1.29	1.56
2:B:275:ARG:HB2	2:B:294:VAL:CG1	1.32	1.56
1:A:630:PRO:CG	2:B:614:ILE:HG12	1.18	1.56
2:B:87:VAL:HG13	2:B:122:SER:CB	1.17	1.56
1:A:384:LEU:HD22	1:A:441:TYR:CE2	1.41	1.55
2:B:472:VAL:HG11	2:B:510:GLY:CA	1.35	1.55
2:B:523:PHE:CZ	2:B:580:TYR:HD2	1.21	1.55
2:B:275:ARG:CG	2:B:294:VAL:HG11	1.24	1.55
2:B:70:MET:CE	2:B:107:ARG:HB2	1.11	1.55
2:B:162:VAL:CG2	2:B:195:ILE:HG23	1.29	1.55
3:M:223:HIS:HA	3:M:479:PHE:CD2	1.38	1.55
4:S:73:ILE:CG2	4:S:88:ILE:HG23	1.37	1.55
2:B:24:ALA:HB3	2:B:35:TYR:CE2	1.41	1.54
1:A:633:PHE:CG	2:B:550:VAL:HG12	1.03	1.54
2:B:216:LYS:HA	2:B:251:LEU:CD1	1.32	1.54
2:B:513:TRP:HA	2:B:551:LEU:CD2	1.37	1.54
2:B:267:ASP:H	2:B:289:PRO:CB	1.15	1.54
3:M:69:ILE:CD1	3:M:90:PHE:CZ	1.87	1.54
1:A:217:ALA:HA	4:S:142:ILE:CB	1.06	1.54
1:A:595:GLU:HG3	2:B:469:ASP:CB	1.18	1.54
1:A:217:ALA:HB1	4:S:142:ILE:CD1	1.34	1.53
2:B:556:LEU:CA	2:B:588:ILE:HD11	1.34	1.53
2:B:127:LEU:CB	2:B:157:THR:HG23	1.35	1.53
2:B:162:VAL:CG2	2:B:195:ILE:CG2	1.85	1.53
2:B:219:TYR:CZ	2:B:226:LEU:CA	1.92	1.53
2:B:127:LEU:CD2	2:B:161:LEU:HD21	1.27	1.53
3:M:101:LEU:HG	3:M:106:LYS:CA	1.30	1.52
2:B:83:PHE:CE2	2:B:119:SER:CB	1.91	1.52
2:B:158:VAL:HG11	2:B:177:ILE:CG1	1.11	1.52
1:A:630:PRO:HG3	2:B:614:ILE:CG1	1.07	1.51
4:S:5:VAL:CG2	4:S:132:LEU:HD21	1.05	1.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:219:TYR:CE1	2:B:226:LEU:HB2	1.34	1.51
2:B:537:PHE:HB3	2:B:598:LEU:CD1	1.35	1.51
2:B:353:GLN:HG3	3:M:47:SER:C	1.20	1.51
2:B:549:LEU:CD2	2:B:611:ALA:N	1.72	1.50
2:B:139:LEU:CD2	2:B:173:VAL:HA	1.07	1.50
2:B:291:TYR:CD2	2:B:294:VAL:CG1	1.93	1.50
2:B:513:TRP:N	2:B:551:LEU:CD1	1.71	1.50
2:B:537:PHE:CD2	2:B:598:LEU:HB3	1.47	1.50
1:A:128:LEU:HD13	1:A:150:LEU:CD2	1.38	1.50
2:B:47:LEU:HD22	2:B:66:ILE:CG1	1.06	1.50
2:B:219:TYR:CB	2:B:223:LEU:HD23	1.42	1.50
2:B:47:LEU:CD2	2:B:66:ILE:HG13	1.36	1.49
2:B:208:ILE:CD1	2:B:236:ILE:HG21	1.42	1.49
2:B:219:TYR:CD1	2:B:226:LEU:HB2	1.46	1.49
3:M:283:PHE:CZ	3:M:289:THR:OG1	1.65	1.49
2:B:215:TYR:CD1	2:B:233:TYR:HE1	1.29	1.49
2:B:275:ARG:CB	2:B:294:VAL:HG11	1.39	1.49
2:B:278:PRO:HA	2:B:288:TYR:C	1.21	1.49
1:A:220:SER:HB3	4:S:142:ILE:CG2	1.36	1.49
2:B:127:LEU:HD22	2:B:161:LEU:CD2	1.38	1.49
4:S:163:THR:CA	4:S:163:THR:C	1.81	1.49
1:A:179:LYS:CE	4:S:143:GLU:HB2	1.40	1.48
2:B:223:LEU:CD1	2:B:259:TYR:CA	1.90	1.48
1:A:128:LEU:CD1	1:A:150:LEU:HD21	1.44	1.48
1:A:204:VAL:HG22	1:A:236:LEU:CD2	1.43	1.48
2:B:178:ILE:HG13	2:B:214:ALA:C	1.17	1.48
2:B:208:ILE:HD13	2:B:236:ILE:CG2	1.01	1.48
2:B:219:TYR:CZ	2:B:226:LEU:HB2	1.49	1.48
2:B:559:ASP:HB2	2:B:563:PHE:CD1	1.46	1.48
1:A:408:ILE:CG2	4:S:64:ASN:C	1.81	1.48
2:B:537:PHE:CB	2:B:598:LEU:HD13	1.43	1.48
1:A:408:ILE:HG22	4:S:65:ASN:N	1.20	1.47
2:B:275:ARG:CB	2:B:294:VAL:CG1	1.90	1.47
1:A:128:LEU:CD1	1:A:150:LEU:CD2	1.91	1.47
1:A:212:ILE:CD1	4:S:145:ASN:ND2	1.73	1.47
2:B:303:LEU:CD1	2:B:333:GLN:HB3	1.41	1.47
2:B:337:THR:HA	2:B:373:LEU:CD2	1.44	1.47
2:B:433:VAL:CG1	2:B:474:VAL:HG21	1.43	1.47
2:B:70:MET:HE1	2:B:107:ARG:CB	1.43	1.47
2:B:523:PHE:CZ	2:B:580:TYR:CD2	1.96	1.47
2:B:252:LEU:CD1	2:B:302:PHE:CD1	1.97	1.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:100:LEU:H	4:S:162:SER:CB	1.25	1.46
1:A:105:VAL:CG2	4:S:167:ILE:CD1	1.91	1.46
2:B:70:MET:CE	2:B:107:ARG:CB	1.93	1.46
2:B:230:PHE:CE1	2:B:252:LEU:HD23	1.50	1.46
2:B:20:ARG:NH1	2:B:21:GLU:HG3	1.23	1.46
2:B:260:LEU:CB	2:B:291:TYR:CE1	1.88	1.46
1:A:633:PHE:CG	2:B:550:VAL:CG1	1.97	1.46
2:B:83:PHE:CZ	2:B:119:SER:CB	1.91	1.46
2:B:219:TYR:CG	2:B:255:TYR:HE1	1.33	1.46
3:M:215:TYR:CG	3:M:468:LYS:HA	1.51	1.46
2:B:106:LEU:HD13	2:B:144:ASP:CB	1.45	1.45
1:A:211:ASP:OD1	4:S:148:ARG:CD	1.65	1.45
1:A:408:ILE:HG23	4:S:64:ASN:CB	1.01	1.45
4:S:53:THR:C	4:S:69:ASN:HB2	1.34	1.45
2:B:127:LEU:CD1	2:B:157:THR:HG21	0.98	1.45
3:M:101:LEU:CG	3:M:106:LYS:HA	1.47	1.45
2:B:230:PHE:CZ	2:B:252:LEU:CD2	1.97	1.45
1:A:147:LEU:HD22	1:A:166:LEU:CD2	1.46	1.44
1:A:408:ILE:CG2	4:S:64:ASN:CB	1.96	1.44
2:B:247:TYR:CE2	3:M:91:THR:HG21	1.50	1.44
2:B:311:TYR:HE2	2:B:342:ALA:CB	1.30	1.44
3:M:104:PHE:CE2	3:M:113:LYS:NZ	1.83	1.44
2:B:527:PRO:CB	2:B:587:ARG:HG3	1.47	1.44
2:B:216:LYS:CA	2:B:251:LEU:HD13	1.44	1.44
2:B:549:LEU:CD2	2:B:611:ALA:CA	1.96	1.44
2:B:219:TYR:CE1	2:B:226:LEU:CD1	1.99	1.43
4:S:109:LEU:HD12	4:S:113:PHE:CD1	1.48	1.43
1:A:96:SER:N	1:A:127:LEU:CD2	1.78	1.43
1:A:101:GLN:HG3	4:S:160:ALA:CB	1.44	1.43
2:B:178:ILE:HG13	2:B:215:TYR:N	1.30	1.43
2:B:566:ALA:O	2:B:574:ASN:CB	1.65	1.43
1:A:100:LEU:N	4:S:162:SER:HB2	1.24	1.43
1:A:107:TYR:CE2	1:A:128:LEU:HD21	1.49	1.43
4:S:109:LEU:CD1	4:S:113:PHE:CD1	2.00	1.43
1:A:185:LEU:CD1	1:A:203:PHE:HE1	1.30	1.42
2:B:219:TYR:CE1	2:B:226:LEU:HD13	1.53	1.42
2:B:260:LEU:CD2	2:B:291:TYR:OH	1.66	1.42
2:B:279:LEU:N	2:B:288:TYR:HB2	1.10	1.42
2:B:549:LEU:CD2	2:B:611:ALA:CB	1.96	1.42
2:B:87:VAL:CG1	2:B:122:SER:CB	1.79	1.42
2:B:106:LEU:CD1	2:B:144:ASP:CB	1.95	1.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:219:TYR:CD1	2:B:226:LEU:HD22	1.52	1.42
1:A:212:ILE:CD1	4:S:145:ASN:HD21	1.25	1.42
2:B:20:ARG:CD	2:B:21:GLU:HB2	1.49	1.42
1:A:629:LEU:CD1	2:B:610:ARG:NH1	1.80	1.42
2:B:311:TYR:CE2	2:B:342:ALA:HB2	1.54	1.42
2:B:546:CYS:CB	2:B:607:ILE:HG12	0.94	1.42
3:M:245:ASP:N	3:M:472:TYR:CD1	1.87	1.42
2:B:309:LEU:HB3	2:B:317:VAL:CG1	1.48	1.41
1:A:594:PHE:HB3	2:B:473:ASN:CB	1.49	1.41
1:A:631:SER:CB	2:B:557:SER:OG	1.66	1.41
2:B:20:ARG:NH1	2:B:21:GLU:CG	1.83	1.41
2:B:422:ALA:HB3	2:B:424:PHE:CE1	1.53	1.41
3:M:69:ILE:HD12	3:M:90:PHE:CZ	1.49	1.41
1:A:408:ILE:HG22	4:S:64:ASN:C	1.37	1.41
2:B:108:PHE:CZ	2:B:115:LEU:HG	1.52	1.41
2:B:546:CYS:HB2	2:B:607:ILE:CD1	1.48	1.41
2:B:546:CYS:CB	2:B:607:ILE:CG1	1.83	1.41
2:B:29:LYS:HE2	2:B:30:LEU:N	1.32	1.41
3:M:65:TYR:CZ	3:M:86:PRO:HB3	1.52	1.41
1:A:250:ASN:OD1	1:A:285:THR:CG2	1.67	1.40
2:B:319:LEU:CD1	2:B:358:MET:HG3	0.93	1.40
1:A:215:VAL:CG2	1:A:243:ILE:HD12	1.49	1.40
2:B:549:LEU:CD2	2:B:611:ALA:HB2	1.49	1.40
4:S:73:ILE:CG2	4:S:88:ILE:CG2	1.98	1.40
1:A:111:SER:CB	1:A:152:THR:OG1	1.67	1.40
4:S:17:VAL:HG21	4:S:19:PHE:CZ	1.54	1.40
1:A:102:GLN:HG2	4:S:166:LYS:N	1.13	1.40
2:B:352:ASN:CB	3:M:49:ASP:OD2	1.69	1.40
2:B:422:ALA:HB3	2:B:424:PHE:CD1	1.57	1.40
3:M:319:SER:CB	3:M:346:ASN:HB2	1.48	1.40
1:A:450:TYR:OH	1:A:476:GLN:CG	1.66	1.39
2:B:158:VAL:CG1	2:B:177:ILE:CG1	2.01	1.39
3:M:260:LEU:CD2	3:M:449:VAL:HG22	1.50	1.39
1:A:633:PHE:CB	2:B:550:VAL:HG12	1.52	1.39
3:M:222:PHE:O	3:M:479:PHE:CE2	1.75	1.39
2:B:215:TYR:CD1	2:B:233:TYR:CE1	2.11	1.39
4:S:109:LEU:CD1	4:S:113:PHE:CE1	2.04	1.39
1:A:217:ALA:C	4:S:142:ILE:HB	1.43	1.38
2:B:87:VAL:CG1	2:B:122:SER:HB3	0.92	1.38
2:B:170:ARG:HA	2:B:199:LEU:CD2	1.53	1.38
3:M:281:GLY:O	3:M:282:VAL:CG2	1.72	1.38

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:288:THR:CG2	1:A:322:PHE:CZ	2.06	1.38
1:A:633:PHE:CD2	2:B:550:VAL:C	1.97	1.38
2:B:62:ALA:O	2:B:66:ILE:CG1	1.70	1.38
2:B:171:GLY:CA	2:B:207:VAL:CG1	2.01	1.38
2:B:549:LEU:HD22	2:B:611:ALA:CB	1.50	1.38
2:B:223:LEU:HD13	2:B:259:TYR:CB	1.53	1.37
1:A:142:LYS:N	4:S:159:ALA:HB2	1.33	1.37
2:B:215:TYR:HD1	2:B:233:TYR:CE1	1.43	1.37
2:B:343:LEU:HD12	2:B:359:LEU:CD1	1.55	1.37
2:B:261:PRO:HG2	2:B:292:GLU:N	1.08	1.37
3:M:290:PHE:CE1	3:M:297:PHE:CD1	2.13	1.37
1:A:595:GLU:CG	2:B:469:ASP:HB3	1.53	1.37
1:A:629:LEU:HD11	2:B:610:ARG:NH1	1.09	1.36
2:B:123:LEU:HD12	2:B:142:LEU:CD2	1.52	1.36
2:B:230:PHE:CZ	2:B:252:LEU:HD22	1.57	1.36
3:M:443:SER:HB3	3:M:447:ILE:CG1	1.55	1.36
2:B:106:LEU:CD1	2:B:144:ASP:HB3	1.54	1.36
2:B:353:GLN:NE2	3:M:47:SER:CB	1.86	1.36
3:M:223:HIS:HA	3:M:479:PHE:CE2	1.58	1.36
4:S:8:PHE:CZ	4:S:84:TYR:HB3	1.58	1.36
2:B:261:PRO:CG	2:B:292:GLU:H	1.37	1.36
1:A:96:SER:CB	1:A:127:LEU:HD11	1.54	1.35
1:A:288:THR:HG21	1:A:322:PHE:CE2	1.57	1.35
4:S:17:VAL:CG2	4:S:19:PHE:CZ	2.09	1.35
3:M:339:GLU:CD	3:M:412:ARG:HE	1.24	1.35
1:A:182:ILE:HG22	1:A:221:VAL:CG2	1.54	1.35
1:A:102:GLN:CA	4:S:167:ILE:HG12	1.57	1.35
2:B:181:TYR:CD2	2:B:218:CYS:O	1.79	1.35
3:M:215:TYR:CD2	3:M:469:GLY:N	1.91	1.35
1:A:107:TYR:CE2	1:A:128:LEU:CD2	2.07	1.34
1:A:136:GLY:O	1:A:139:ASP:HB3	1.24	1.34
2:B:208:ILE:CD1	2:B:236:ILE:CG2	1.94	1.34
2:B:393:ILE:CG2	2:B:431:MET:HG2	1.54	1.34
1:A:219:VAL:O	1:A:259:LEU:HD11	1.26	1.34
1:A:224:GLU:CB	4:S:138:GLY:O	1.73	1.34
2:B:219:TYR:CD1	2:B:226:LEU:CG	2.09	1.34
4:S:53:THR:HB	4:S:69:ASN:N	1.02	1.34
2:B:223:LEU:HD13	2:B:259:TYR:CA	1.53	1.34
3:M:245:ASP:O	3:M:472:TYR:CE1	1.80	1.34
1:A:291:ILE:CG2	1:A:318:ARG:HB3	1.57	1.33
1:A:403:LEU:HD21	1:A:421:PRO:C	1.48	1.33

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:197:LYS:HA	2:B:229:HIS:CD2	1.61	1.33
4:S:8:PHE:CB	4:S:36:TYR:HE2	1.36	1.33
1:A:105:VAL:CG2	4:S:167:ILE:HD12	1.47	1.33
1:A:185:LEU:CD1	1:A:203:PHE:CE1	2.09	1.33
1:A:255:ARG:NH2	4:S:135:ILE:CG2	1.92	1.33
3:M:101:LEU:O	3:M:106:LYS:N	1.58	1.33
2:B:140:SER:HB2	2:B:172:GLU:CD	1.46	1.33
2:B:252:LEU:CD1	2:B:302:PHE:CE1	2.10	1.33
2:B:566:ALA:HA	2:B:574:ASN:CB	1.55	1.33
1:A:520:GLY:CA	1:A:558:VAL:HG22	1.56	1.33
2:B:353:GLN:NE2	3:M:47:SER:HB3	1.00	1.33
2:B:559:ASP:CB	2:B:563:PHE:CD1	2.08	1.33
1:A:463:ASP:O	2:B:1:MET:HG3	1.15	1.33
2:B:546:CYS:HA	2:B:607:ILE:CG2	1.58	1.33
4:S:53:THR:CB	4:S:69:ASN:N	1.90	1.33
1:A:141:VAL:HG12	4:S:159:ALA:CB	1.59	1.32
2:B:106:LEU:CD1	2:B:144:ASP:O	1.73	1.32
2:B:519:ALA:O	2:B:523:PHE:CB	1.74	1.32
2:B:2:VAL:HG12	2:B:6:HIS:NE2	1.21	1.32
2:B:41:ASN:HB3	2:B:43:ASN:OD1	1.23	1.32
2:B:219:TYR:CD1	2:B:226:LEU:CD2	2.11	1.32
2:B:393:ILE:HG23	2:B:431:MET:CB	1.56	1.32
2:B:230:PHE:CE2	2:B:298:ASP:O	1.80	1.32
2:B:556:LEU:CD2	2:B:588:ILE:HG12	1.57	1.32
2:B:193:LEU:O	2:B:195:ILE:N	1.60	1.32
2:B:219:TYR:CG	2:B:255:TYR:CE1	2.17	1.32
3:M:244:VAL:CA	3:M:472:TYR:CD2	2.09	1.32
1:A:103:LYS:HE3	1:A:131:ASP:OD1	1.19	1.32
2:B:352:ASN:CA	3:M:49:ASP:OD2	1.76	1.32
2:B:143:SER:C	2:B:179:LYS:CD	1.97	1.31
2:B:519:ALA:O	2:B:523:PHE:HB3	1.16	1.31
3:M:67:SER:OG	3:M:90:PHE:CD1	1.71	1.31
1:A:138:ASN:OD1	4:S:158:LYS:CE	1.77	1.31
1:A:255:ARG:NH2	4:S:135:ILE:HG23	1.41	1.31
3:M:104:PHE:CE1	3:M:113:LYS:NZ	1.84	1.31
3:M:125:PHE:O	3:M:129:VAL:CG2	1.77	1.31
2:B:62:ALA:O	2:B:66:ILE:CD1	1.79	1.31
2:B:274:PRO:HG2	2:B:295:ASN:CG	1.45	1.31
2:B:343:LEU:CD2	2:B:362:ALA:HB3	1.60	1.31
3:M:454:ILE:CG2	3:M:464:THR:HG21	1.61	1.31
4:S:53:THR:CA	4:S:69:ASN:HB2	1.60	1.31

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:105:VAL:CB	4:S:167:ILE:HD13	1.59	1.31
1:A:140:VAL:CA	1:A:177:ILE:CG1	2.07	1.31
1:A:170:LEU:O	1:A:206:LYS:HD2	1.26	1.31
1:A:251:TRP:CH2	4:S:103:GLN:HB2	1.66	1.31
3:M:283:PHE:CE2	3:M:289:THR:OG1	1.81	1.31
1:A:105:VAL:N	4:S:167:ILE:CD1	1.92	1.31
1:A:176:TYR:CB	4:S:155:GLU:HG3	1.61	1.31
1:A:332:TYR:CD1	1:A:366:SER:OG	1.77	1.31
1:A:631:SER:CB	2:B:557:SER:CB	2.07	1.31
2:B:117:LEU:HD21	2:B:149:SER:OG	1.25	1.31
2:B:230:PHE:HE2	2:B:298:ASP:O	1.06	1.31
1:A:429:VAL:CB	1:A:469:LEU:HD11	1.59	1.30
1:A:633:PHE:CD1	2:B:513:TRP:CZ3	2.19	1.30
2:B:123:LEU:CD1	2:B:142:LEU:CG	2.08	1.30
2:B:278:PRO:CA	2:B:288:TYR:C	1.99	1.30
3:M:65:TYR:CE1	3:M:86:PRO:HB3	1.63	1.30
1:A:252:ILE:HA	4:S:144:THR:OG1	1.13	1.30
2:B:158:VAL:HG11	2:B:177:ILE:CD1	1.61	1.30
2:B:181:TYR:HD2	2:B:218:CYS:C	1.32	1.30
2:B:216:LYS:CA	2:B:251:LEU:CD1	2.01	1.30
3:M:218:LEU:HA	3:M:472:TYR:CE2	1.67	1.30
4:S:8:PHE:CB	4:S:36:TYR:CE2	2.13	1.30
4:S:8:PHE:HB2	4:S:36:TYR:CE2	1.66	1.30
2:B:140:SER:CB	2:B:172:GLU:OE1	1.80	1.30
2:B:559:ASP:O	2:B:563:PHE:N	1.64	1.30
3:M:60:LEU:HD22	3:M:62:VAL:CG2	1.60	1.30
3:M:317:MET:HB3	3:M:320:ILE:O	1.13	1.30
1:A:384:LEU:CD2	1:A:441:TYR:CE2	2.14	1.29
2:B:274:PRO:C	2:B:295:ASN:HD21	1.35	1.29
3:M:217:ASP:CB	3:M:470:ALA:C	2.00	1.29
2:B:170:ARG:NH1	2:B:198:GLU:HG2	1.45	1.29
2:B:307:ASN:HD21	2:B:336:ASN:ND2	1.27	1.29
1:A:275:LEU:HD13	1:A:308:ASP:OD1	1.23	1.29
1:A:633:PHE:CD2	2:B:551:LEU:N	2.01	1.29
1:A:633:PHE:CD1	2:B:550:VAL:HG12	1.68	1.29
2:B:513:TRP:N	2:B:551:LEU:HD13	0.97	1.29
2:B:566:ALA:CA	2:B:574:ASN:HB3	1.62	1.29
1:A:128:LEU:HD13	1:A:150:LEU:CG	1.62	1.28
2:B:37:TYR:OH	2:B:46:GLN:CD	1.71	1.28
3:M:221:THR:CB	3:M:474:THR:O	1.81	1.28
4:S:39:ILE:HD11	4:S:77:TYR:CD2	1.68	1.28

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:147:LEU:CD1	1:A:181:ALA:HA	1.62	1.28
2:B:24:ALA:HB3	2:B:35:TYR:CZ	1.66	1.28
2:B:279:LEU:H	2:B:288:TYR:CB	1.44	1.28
2:B:418:TYR:OH	2:B:432:ALA:HB2	1.18	1.28
1:A:633:PHE:CD1	2:B:550:VAL:CG1	2.14	1.28
2:B:77:ILE:O	2:B:79:VAL:N	1.64	1.28
2:B:181:TYR:HD2	2:B:218:CYS:O	0.95	1.28
2:B:227:HIS:O	2:B:229:HIS:N	1.64	1.28
2:B:546:CYS:SG	2:B:607:ILE:HG12	1.71	1.28
3:M:241:HIS:O	3:M:474:THR:HB	1.29	1.28
3:M:319:SER:HB3	3:M:346:ASN:N	1.48	1.28
4:S:9:ASN:OD1	4:S:13:GLN:N	1.63	1.28
1:A:403:LEU:HD22	1:A:422:GLU:CG	1.64	1.28
2:B:116:THR:CG2	2:B:150:LEU:HD11	1.62	1.28
3:M:379:LEU:HD22	3:M:386:PHE:CG	1.65	1.28
1:A:633:PHE:CE2	2:B:551:LEU:N	1.98	1.28
2:B:226:LEU:HB3	2:B:255:TYR:OH	1.21	1.28
2:B:563:PHE:HD2	2:B:584:SER:CB	1.45	1.28
2:B:274:PRO:HG2	2:B:295:ASN:OD1	1.13	1.27
1:A:463:ASP:O	2:B:1:MET:CG	1.80	1.27
2:B:315:PRO:HB3	2:B:352:ASN:OD1	1.34	1.27
3:M:449:VAL:HG11	3:M:452:ILE:CD1	1.63	1.27
2:B:140:SER:HB2	2:B:172:GLU:OE1	1.10	1.27
2:B:225:LEU:HD12	2:B:283:TYR:OH	1.21	1.27
3:M:339:GLU:HG3	3:M:412:ARG:CG	1.63	1.27
1:A:516:ILE:HG22	1:A:554:ALA:CB	1.63	1.27
2:B:549:LEU:CD1	2:B:611:ALA:HA	1.63	1.27
2:B:127:LEU:CD1	2:B:157:THR:CG2	1.84	1.27
2:B:353:GLN:HG3	3:M:47:SER:O	1.26	1.27
2:B:512:VAL:C	2:B:551:LEU:HD13	1.55	1.27
2:B:513:TRP:CA	2:B:551:LEU:CD1	2.10	1.27
3:M:66:PHE:CA	3:M:77:LEU:HD11	1.63	1.27
3:M:443:SER:CB	3:M:447:ILE:HG13	1.65	1.27
2:B:162:VAL:HG22	2:B:199:LEU:CG	1.66	1.26
2:B:316:THR:CG2	3:M:90:PHE:CE2	2.15	1.26
2:B:252:LEU:HD12	2:B:302:PHE:CE1	1.65	1.26
2:B:278:PRO:CG	2:B:292:GLU:OE1	1.83	1.26
2:B:337:THR:CA	2:B:373:LEU:HD21	1.64	1.26
2:B:21:GLU:HA	2:B:24:ALA:CB	1.66	1.26
2:B:29:LYS:CE	2:B:30:LEU:H	1.48	1.26
2:B:123:LEU:HD13	2:B:142:LEU:CG	1.64	1.26

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:219:TYR:CB	2:B:255:TYR:CE1	2.16	1.26
3:M:268:GLY:N	3:M:302:TYR:OH	1.69	1.26
2:B:143:SER:C	2:B:179:LYS:HD2	1.50	1.25
2:B:566:ALA:C	2:B:574:ASN:CB	2.04	1.25
4:S:14:PRO:HA	4:S:36:TYR:OH	1.33	1.25
1:A:250:ASN:OD1	1:A:285:THR:HG22	1.20	1.25
1:A:633:PHE:CE1	2:B:513:TRP:CZ3	2.24	1.25
1:A:96:SER:CA	1:A:127:LEU:HD21	1.65	1.25
1:A:204:VAL:CG2	1:A:236:LEU:HD21	1.64	1.25
3:M:342:LEU:HD11	3:M:411:LEU:CD2	1.65	1.25
1:A:141:VAL:CG1	4:S:159:ALA:HB3	1.67	1.25
2:B:219:TYR:CG	2:B:226:LEU:HB2	1.72	1.25
2:B:245:GLN:CD	2:B:309:LEU:CD1	2.05	1.25
3:M:101:LEU:HA	3:M:109:LEU:CD1	1.65	1.25
3:M:224:VAL:H	3:M:479:PHE:CB	1.50	1.25
1:A:138:ASN:OD1	4:S:158:LYS:HE3	1.08	1.25
2:B:553:ALA:HB2	2:B:614:ILE:CD1	1.63	1.25
2:B:556:LEU:HD23	2:B:588:ILE:CG1	1.66	1.25
3:M:217:ASP:HB2	3:M:470:ALA:O	1.33	1.25
3:M:218:LEU:HA	3:M:472:TYR:CD2	1.72	1.25
3:M:241:HIS:O	3:M:474:THR:CB	1.83	1.25
3:M:260:LEU:HD23	3:M:449:VAL:CG2	1.67	1.25
1:A:102:GLN:HB2	4:S:163:THR:CB	1.65	1.25
1:A:399:ASP:O	1:A:420:ILE:O	1.55	1.25
1:A:633:PHE:CD2	2:B:551:LEU:CA	2.07	1.25
2:B:25:VAL:HG11	2:B:36:THR:OG1	1.34	1.25
2:B:225:LEU:HD13	2:B:283:TYR:CE1	1.70	1.25
2:B:291:TYR:CE2	2:B:294:VAL:HB	1.69	1.25
3:M:254:PRO:HB3	3:M:454:ILE:CD1	1.65	1.25
1:A:103:LYS:H	4:S:163:THR:CB	1.49	1.24
2:B:219:TYR:CZ	2:B:226:LEU:N	2.03	1.24
4:S:53:THR:CG2	4:S:67:GLU:O	1.85	1.24
1:A:96:SER:CA	1:A:127:LEU:HD11	1.66	1.24
1:A:220:SER:CB	4:S:142:ILE:HG22	1.65	1.24
2:B:158:VAL:CG1	2:B:177:ILE:HD11	1.66	1.24
2:B:159:LYS:HA	2:B:195:ILE:CD1	1.67	1.24
2:B:353:GLN:CG	3:M:47:SER:C	2.06	1.24
3:M:272:LEU:CD2	3:M:278:ILE:HB	1.67	1.24
2:B:143:SER:O	2:B:179:LYS:CD	1.85	1.24
2:B:178:ILE:CG1	2:B:214:ALA:C	2.05	1.24
4:S:56:SER:O	4:S:60:SER:CB	1.86	1.24

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:219:LEU:CB	3:M:472:TYR:O	1.85	1.24
3:M:244:VAL:HG13	3:M:472:TYR:CE2	1.72	1.24
3:M:449:VAL:CG1	3:M:452:ILE:CD1	2.14	1.24
1:A:101:GLN:CG	4:S:160:ALA:HB1	1.56	1.24
1:A:513:ARG:CD	1:A:550:VAL:HG21	1.65	1.24
1:A:637:GLU:CG	2:B:516:GLY:H	1.49	1.24
4:S:48:SER:CB	4:S:77:TYR:C	2.06	1.24
1:A:516:ILE:CG2	1:A:554:ALA:HB3	1.67	1.23
1:A:631:SER:HB2	2:B:557:SER:OG	1.23	1.23
2:B:24:ALA:CB	2:B:35:TYR:CZ	2.21	1.23
2:B:37:TYR:HD2	2:B:38:TYR:CD1	1.55	1.23
2:B:162:VAL:HG23	2:B:195:ILE:CG2	1.51	1.23
2:B:199:LEU:O	2:B:201:ALA:N	1.71	1.23
3:M:432:THR:OG1	3:M:480:GLN:CG	1.86	1.23
4:S:53:THR:HB	4:S:69:ASN:CA	1.66	1.23
2:B:86:VAL:CG1	2:B:101:ILE:HG23	1.66	1.23
2:B:223:LEU:CD1	2:B:259:TYR:N	2.01	1.23
2:B:584:SER:O	2:B:588:ILE:HG22	1.08	1.23
3:M:217:ASP:CB	3:M:470:ALA:O	1.85	1.23
4:S:8:PHE:CD1	4:S:84:TYR:HB2	1.71	1.23
1:A:147:LEU:HB3	1:A:184:ALA:CB	1.69	1.23
1:A:326:GLN:HA	1:A:331:ARG:NH2	1.54	1.23
1:A:557:LYS:HE2	2:B:606:ASP:N	0.92	1.23
2:B:143:SER:O	2:B:179:LYS:HD3	1.09	1.23
2:B:219:TYR:CZ	2:B:226:LEU:CB	2.01	1.23
2:B:278:PRO:CB	2:B:288:TYR:O	1.85	1.23
3:M:378:ILE:O	3:M:413:GLY:HA3	1.12	1.23
4:S:8:PHE:CZ	4:S:84:TYR:CB	2.16	1.23
1:A:217:ALA:CB	4:S:142:ILE:HD12	1.68	1.23
2:B:261:PRO:CG	2:B:292:GLU:N	1.98	1.23
2:B:277:CYS:O	2:B:288:TYR:HB3	1.06	1.23
3:M:66:PHE:CB	3:M:77:LEU:HD11	1.43	1.23
2:B:212:VAL:CG2	2:B:248:LEU:HD21	1.69	1.22
4:S:47:GLN:NE2	4:S:79:ASN:N	1.86	1.22
1:A:88:ASN:CB	1:A:120:ILE:HD12	1.68	1.22
1:A:212:ILE:HD12	4:S:145:ASN:ND2	0.91	1.22
1:A:291:ILE:HG23	1:A:318:ARG:CB	1.69	1.22
1:A:609:LEU:HG	1:A:628:VAL:CB	1.67	1.22
2:B:216:LYS:CB	2:B:251:LEU:CD1	2.16	1.22
2:B:472:VAL:CG1	2:B:510:GLY:HA3	1.68	1.22
3:M:214:LEU:C	3:M:467:TYR:HB3	1.57	1.22

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:563:CYS:CB	1:A:621:LEU:HD12	1.68	1.22
2:B:20:ARG:CZ	2:B:21:GLU:HG3	1.68	1.22
2:B:219:TYR:CD1	2:B:226:LEU:HD13	1.74	1.22
2:B:275:ARG:N	2:B:295:ASN:ND2	1.87	1.22
2:B:479:VAL:HG22	2:B:486:HIS:CD2	1.73	1.22
3:M:221:THR:OG1	3:M:474:THR:O	1.56	1.22
1:A:92:LEU:HD21	1:A:120:ILE:O	1.38	1.22
2:B:135:ARG:NH2	2:B:164:ASP:OD1	1.72	1.22
2:B:219:TYR:CB	2:B:255:TYR:HE1	1.52	1.22
2:B:542:PRO:HA	2:B:602:ASP:OD2	1.29	1.22
4:S:48:SER:CB	4:S:77:TYR:CB	2.18	1.22
4:S:73:ILE:HG21	4:S:88:ILE:CG2	1.65	1.22
1:A:96:SER:CB	1:A:127:LEU:CD1	2.16	1.21
2:B:171:GLY:N	2:B:207:VAL:HG13	1.53	1.21
2:B:223:LEU:CD2	2:B:255:TYR:CE1	2.23	1.21
4:S:131:VAL:CG2	4:S:153:VAL:HG22	1.69	1.21
1:A:638:LEU:HB2	2:B:516:GLY:O	1.36	1.21
2:B:20:ARG:NE	2:B:21:GLU:HB2	1.55	1.21
1:A:217:ALA:CB	4:S:142:ILE:CG1	2.19	1.21
1:A:219:VAL:CG1	1:A:259:LEU:HD13	1.70	1.21
2:B:231:ARG:HG3	2:B:298:ASP:OD1	1.39	1.21
2:B:277:CYS:O	2:B:288:TYR:CB	1.88	1.21
2:B:352:ASN:HA	3:M:49:ASP:CG	1.39	1.21
3:M:2:TYR:O	3:M:81:SER:HB2	1.38	1.21
1:A:105:VAL:CG2	4:S:167:ILE:HG23	1.69	1.21
1:A:213:SER:O	4:S:143:GLU:CG	1.87	1.21
1:A:219:VAL:O	1:A:259:LEU:CD1	1.87	1.21
1:A:557:LYS:NZ	2:B:604:GLU:CG	1.70	1.21
2:B:87:VAL:HG13	2:B:122:SER:OG	1.39	1.21
4:S:34:GLN:OE1	4:S:58:LEU:HD21	1.32	1.21
1:A:101:GLN:HE21	4:S:167:ILE:CG2	1.53	1.21
2:B:47:LEU:CD2	2:B:66:ILE:CG1	2.02	1.21
2:B:79:VAL:HB	2:B:108:PHE:CE1	1.74	1.21
2:B:105:LEU:HB3	2:B:145:MET:CE	1.71	1.21
2:B:219:TYR:CD1	2:B:226:LEU:CB	2.07	1.21
2:B:290:SER:O	2:B:292:GLU:N	1.67	1.21
2:B:219:TYR:HB3	2:B:255:TYR:CE1	1.76	1.20
4:S:15:ARG:NH1	4:S:122:ILE:HD11	1.54	1.20
1:A:101:GLN:C	4:S:167:ILE:HG13	1.59	1.20
1:A:140:VAL:CA	1:A:177:ILE:HG12	1.70	1.20
1:A:211:ASP:CG	4:S:148:ARG:HD3	1.60	1.20

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:557:LYS:HB3	2:B:605:PHE:CD2	1.76	1.20
2:B:124:GLN:HA	2:B:157:THR:OG1	1.38	1.20
2:B:344:VAL:HG13	2:B:381:PHE:CZ	1.76	1.20
2:B:531:ARG:HA	2:B:591:MET:SD	1.80	1.20
3:M:319:SER:CB	3:M:346:ASN:CB	2.18	1.20
1:A:125:THR:OG1	1:A:158:LEU:HD13	1.38	1.20
1:A:388:VAL:HG13	1:A:432:ILE:CD1	1.70	1.20
2:B:37:TYR:CD2	2:B:38:TYR:CD1	2.28	1.20
2:B:127:LEU:CB	2:B:157:THR:CG2	2.19	1.20
4:S:34:GLN:OE1	4:S:58:LEU:HD11	1.35	1.20
2:B:219:TYR:CE1	2:B:226:LEU:CG	2.22	1.20
2:B:278:PRO:HA	2:B:288:TYR:CA	1.69	1.20
3:M:222:PHE:CD1	3:M:240:ILE:HG23	1.75	1.20
1:A:96:SER:N	1:A:127:LEU:HD21	0.89	1.20
2:B:162:VAL:CB	2:B:195:ILE:HG23	1.72	1.20
2:B:245:GLN:CD	2:B:309:LEU:HD11	1.56	1.20
2:B:261:PRO:CA	2:B:290:SER:HB3	1.49	1.20
1:A:211:ASP:OD1	4:S:148:ARG:HD3	1.02	1.19
1:A:316:LEU:CD1	1:A:348:PHE:CD2	2.26	1.19
1:A:594:PHE:CB	2:B:473:ASN:HB2	1.72	1.19
2:B:62:ALA:O	2:B:66:ILE:HG13	1.21	1.19
2:B:219:TYR:CE2	2:B:226:LEU:HB2	1.75	1.19
2:B:513:TRP:CA	2:B:551:LEU:CD2	2.20	1.19
3:M:243:ILE:H	3:M:474:THR:CG2	1.53	1.19
4:S:47:GLN:NE2	4:S:79:ASN:H	1.38	1.19
1:A:559:PHE:CE1	1:A:581:LEU:HD22	1.76	1.19
1:A:631:SER:HA	2:B:557:SER:CB	1.71	1.19
2:B:25:VAL:HG23	2:B:35:TYR:CD2	1.77	1.19
2:B:267:ASP:N	2:B:289:PRO:CB	2.00	1.19
2:B:278:PRO:HG2	2:B:292:GLU:OE1	1.41	1.19
2:B:291:TYR:CD2	2:B:294:VAL:HG12	1.61	1.19
3:M:9:ASP:C	3:M:75:TRP:CD1	2.16	1.19
3:M:340:LEU:O	3:M:411:LEU:HB3	1.39	1.19
1:A:102:GLN:CB	4:S:166:LYS:HB2	1.71	1.19
2:B:197:LYS:O	2:B:199:LEU:N	1.74	1.19
2:B:279:LEU:N	2:B:288:TYR:CB	2.02	1.19
2:B:513:TRP:CA	2:B:551:LEU:HD13	1.66	1.19
1:A:217:ALA:HB2	4:S:143:GLU:HG3	1.23	1.19
2:B:291:TYR:CD2	2:B:294:VAL:HG11	1.68	1.19
2:B:311:TYR:CE2	2:B:342:ALA:CB	2.17	1.19
2:B:393:ILE:HG23	2:B:431:MET:CG	1.71	1.19

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:546:CYS:CA	2:B:607:ILE:HG23	1.72	1.19
3:M:246:VAL:HB	3:M:297:PHE:CZ	1.75	1.19
1:A:185:LEU:HD12	1:A:203:PHE:CE1	1.74	1.19
1:A:605:GLU:CG	1:A:632:PHE:CD2	2.26	1.19
2:B:108:PHE:CE2	2:B:115:LEU:HB2	1.78	1.19
2:B:151:ALA:O	2:B:188:TYR:CE2	1.85	1.19
2:B:223:LEU:HD22	2:B:255:TYR:CE1	1.78	1.19
2:B:582:ASP:O	2:B:584:SER:HB3	1.43	1.19
1:A:101:GLN:CG	4:S:167:ILE:HG21	1.72	1.18
1:A:179:LYS:HE3	4:S:143:GLU:CB	1.73	1.18
2:B:223:LEU:CD1	2:B:259:TYR:HA	1.57	1.18
1:A:502:ASP:OD2	1:A:506:LYS:NZ	1.72	1.18
1:A:275:LEU:HD21	1:A:311:THR:OG1	1.41	1.18
2:B:25:VAL:CG1	2:B:36:THR:OG1	1.92	1.18
2:B:225:LEU:CD1	2:B:283:TYR:OH	1.91	1.18
2:B:267:ASP:N	2:B:289:PRO:HB3	1.53	1.18
2:B:277:CYS:SG	2:B:292:GLU:HG3	1.83	1.18
2:B:303:LEU:CD1	2:B:333:GLN:CB	2.22	1.18
2:B:531:ARG:CA	2:B:591:MET:SD	2.31	1.18
3:M:66:PHE:CB	3:M:77:LEU:CD1	2.21	1.18
3:M:319:SER:OG	3:M:346:ASN:CB	1.92	1.18
4:S:135:ILE:O	4:S:141:VAL:HA	1.40	1.18
2:B:422:ALA:CB	2:B:424:PHE:CE1	2.26	1.18
3:M:338:PHE:CD2	3:M:415:ILE:HG13	1.78	1.18
1:A:103:LYS:O	1:A:107:TYR:CD1	1.97	1.18
2:B:98:LYS:NZ	2:B:134:LEU:HB3	1.58	1.18
4:S:130:SER:OG	4:S:156:LEU:CD1	1.89	1.18
2:B:12:LEU:O	2:B:16:LYS:HB2	1.42	1.17
2:B:56:SER:CB	2:B:92:THR:HG21	1.74	1.17
2:B:315:PRO:HB3	2:B:355:ASN:ND2	1.58	1.17
2:B:567:GLN:O	2:B:569:THR:OG1	1.54	1.17
2:B:47:LEU:HD22	2:B:66:ILE:HG12	1.20	1.17
2:B:170:ARG:HH12	2:B:198:GLU:CG	1.57	1.17
3:M:244:VAL:HA	3:M:472:TYR:CE2	1.77	1.17
1:A:102:GLN:HB3	4:S:166:LYS:CB	1.74	1.17
1:A:166:LEU:CD1	1:A:185:LEU:HD23	1.73	1.17
1:A:631:SER:CA	2:B:557:SER:CB	2.21	1.17
2:B:352:ASN:CA	3:M:49:ASP:CG	2.10	1.17
3:M:222:PHE:C	3:M:479:PHE:CZ	2.18	1.17
1:A:253:ILE:HD11	1:A:281:LEU:HD22	1.20	1.17
1:A:399:ASP:CA	1:A:420:ILE:HB	1.74	1.17

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:595:GLU:CG	2:B:469:ASP:CB	2.14	1.17
2:B:171:GLY:HA3	2:B:207:VAL:CG1	1.70	1.17
2:B:252:LEU:HB2	2:B:302:PHE:CZ	1.78	1.17
2:B:360:LEU:CD1	2:B:391:ALA:HA	1.73	1.17
1:A:71:VAL:CG1	1:A:105:VAL:HG12	1.71	1.17
2:B:278:PRO:CD	2:B:292:GLU:OE1	1.91	1.17
2:B:523:PHE:HZ	2:B:580:TYR:CE2	1.63	1.17
2:B:545:ARG:HD3	2:B:602:ASP:HB2	1.25	1.17
3:M:226:PHE:HB2	3:M:481:VAL:HG22	1.17	1.17
3:M:336:ASP:OD1	3:M:415:ILE:O	1.62	1.17
1:A:105:VAL:N	4:S:167:ILE:HD11	1.53	1.16
1:A:185:LEU:HD13	1:A:203:PHE:CE1	1.71	1.16
1:A:557:LYS:HD3	2:B:605:PHE:HB3	1.24	1.16
1:A:631:SER:HA	2:B:557:SER:HB3	1.24	1.16
3:M:260:LEU:CD2	3:M:449:VAL:CG2	2.20	1.16
1:A:99:LYS:HD3	4:S:164:ASP:H	1.06	1.16
1:A:107:TYR:CD2	1:A:128:LEU:HD21	1.80	1.16
2:B:479:VAL:CG2	2:B:486:HIS:CD2	2.29	1.16
3:M:339:GLU:CD	3:M:412:ARG:NE	1.98	1.16
2:B:83:PHE:HE2	2:B:119:SER:CA	1.58	1.16
2:B:216:LYS:CG	2:B:251:LEU:HD13	1.76	1.16
2:B:497:LEU:HD23	2:B:533:LEU:CD2	1.75	1.16
3:M:215:TYR:CD1	3:M:468:LYS:CA	2.27	1.16
3:M:339:GLU:HG3	3:M:412:ARG:CD	1.74	1.16
4:S:47:GLN:CD	4:S:79:ASN:N	1.97	1.16
1:A:147:LEU:CD2	1:A:166:LEU:CD2	2.23	1.16
1:A:219:VAL:HG21	1:A:256:LEU:HD21	1.22	1.16
2:B:105:LEU:HB3	2:B:145:MET:HE3	1.22	1.16
3:M:219:LEU:HD13	3:M:472:TYR:O	1.45	1.16
4:S:80:TYR:O	4:S:82:THR:N	1.78	1.16
1:A:103:LYS:CE	1:A:131:ASP:OD1	1.94	1.16
1:A:213:SER:O	4:S:143:GLU:HG2	0.99	1.16
1:A:216:SER:HB3	4:S:143:GLU:HA	1.18	1.16
2:B:175:LEU:CG	2:B:210:CYS:HB3	1.75	1.16
2:B:275:ARG:CG	2:B:294:VAL:CG1	2.16	1.16
2:B:275:ARG:N	2:B:295:ASN:HD21	1.41	1.16
3:M:219:LEU:CD1	3:M:472:TYR:O	1.94	1.16
1:A:217:ALA:O	4:S:142:ILE:HB	1.43	1.15
1:A:421:PRO:HG2	1:A:424:TYR:CE1	1.81	1.15
3:M:92:PHE:CZ	3:M:129:VAL:HG22	1.79	1.15
3:M:379:LEU:CD2	3:M:386:PHE:CD1	2.28	1.15

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:219:VAL:C	1:A:259:LEU:CD1	2.13	1.15
2:B:219:TYR:CZ	2:B:226:LEU:HA	1.78	1.15
2:B:556:LEU:CD2	2:B:588:ILE:CG1	2.24	1.15
4:S:53:THR:HG21	4:S:67:GLU:O	1.44	1.15
1:A:103:LYS:H	4:S:163:THR:CG2	1.57	1.15
1:A:104:ARG:HA	1:A:145:ILE:HG21	1.22	1.15
1:A:147:LEU:CD2	1:A:166:LEU:HD23	1.77	1.15
2:B:80:GLN:HG2	2:B:115:LEU:HD11	1.18	1.15
2:B:151:ALA:HA	2:B:180:LEU:CD1	1.74	1.15
2:B:275:ARG:HG3	2:B:291:TYR:CD2	1.79	1.15
2:B:393:ILE:CG2	2:B:431:MET:CG	2.23	1.15
3:M:319:SER:HB3	3:M:346:ASN:CB	1.74	1.15
3:M:338:PHE:CE2	3:M:415:ILE:HG13	1.81	1.15
4:S:34:GLN:OE1	4:S:58:LEU:CD2	1.93	1.15
1:A:163:ALA:CB	1:A:195:ALA:HB1	1.76	1.15
1:A:630:PRO:CB	2:B:614:ILE:HG12	1.75	1.15
2:B:162:VAL:CG2	2:B:195:ILE:HG22	1.76	1.15
2:B:223:LEU:HD12	2:B:259:TYR:CA	1.64	1.15
2:B:230:PHE:CD2	2:B:298:ASP:HB3	1.81	1.15
3:M:443:SER:OG	3:M:447:ILE:C	1.84	1.15
1:A:102:GLN:HB2	4:S:163:THR:CA	1.77	1.15
1:A:147:LEU:HD12	1:A:181:ALA:HA	1.29	1.15
1:A:219:VAL:HG13	1:A:259:LEU:HD13	1.24	1.15
1:A:450:TYR:OH	1:A:476:GLN:HG2	1.34	1.15
1:A:450:TYR:OH	1:A:476:GLN:HG3	1.42	1.15
2:B:132:SER:HA	2:B:169:VAL:HG21	1.20	1.15
1:A:630:PRO:HG2	2:B:614:ILE:HA	1.22	1.14
1:A:634:ASN:O	2:B:516:GLY:C	1.86	1.14
2:B:158:VAL:CG1	2:B:177:ILE:CD1	2.17	1.14
2:B:353:GLN:CG	3:M:49:ASP:H	1.59	1.14
2:B:513:TRP:HA	2:B:551:LEU:CG	1.76	1.14
1:A:92:LEU:HD11	1:A:120:ILE:HA	1.20	1.14
1:A:140:VAL:N	1:A:177:ILE:HD11	1.61	1.14
1:A:288:THR:OG1	1:A:291:ILE:HB	1.43	1.14
1:A:609:LEU:CG	1:A:628:VAL:HB	1.78	1.14
3:M:217:ASP:CG	3:M:471:LYS:HA	1.67	1.14
3:M:340:LEU:O	3:M:411:LEU:CB	1.92	1.14
1:A:101:GLN:HE21	4:S:167:ILE:HG22	1.10	1.14
1:A:629:LEU:HD11	2:B:610:ARG:CZ	1.77	1.14
2:B:2:VAL:HG12	2:B:6:HIS:CE1	1.82	1.14
2:B:154:ILE:HD12	2:B:180:LEU:CD1	1.78	1.14

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:523:PHE:CE1	2:B:580:TYR:HD2	1.66	1.14
1:A:251:TRP:HH2	4:S:103:GLN:OE1	1.27	1.14
1:A:275:LEU:O	1:A:276:PRO:C	1.73	1.14
1:A:555:LEU:HD13	1:A:581:LEU:HD11	1.21	1.14
1:A:609:LEU:CD2	1:A:628:VAL:HG21	1.78	1.14
2:B:127:LEU:CD2	2:B:161:LEU:CD2	2.06	1.14
2:B:127:LEU:HD22	2:B:161:LEU:HD22	1.24	1.14
1:A:67:LYS:H	4:S:165:SER:HB2	1.00	1.14
1:A:176:TYR:HB3	4:S:155:GLU:HG3	1.28	1.14
1:A:638:LEU:HD12	2:B:518:ILE:CG2	1.78	1.14
2:B:260:LEU:CA	2:B:291:TYR:CE1	2.21	1.14
2:B:556:LEU:HD22	2:B:588:ILE:HG12	1.16	1.14
3:M:343:ASN:HA	3:M:408:VAL:HG13	1.21	1.14
1:A:178:ARG:HD3	1:A:209:ASP:OD2	1.48	1.13
1:A:217:ALA:CB	4:S:142:ILE:CD1	2.24	1.13
2:B:79:VAL:HG23	2:B:108:PHE:CZ	1.82	1.13
2:B:196:LEU:HB2	2:B:229:HIS:CE1	1.83	1.13
4:S:7:ILE:HG21	4:S:121:LEU:HD21	1.26	1.13
1:A:262:ASN:O	1:A:265:GLN:O	1.65	1.13
1:A:399:ASP:HA	1:A:420:ILE:CB	1.77	1.13
1:A:631:SER:O	2:B:554:LYS:HG3	1.49	1.13
2:B:219:TYR:CD1	2:B:226:LEU:CD1	2.27	1.13
2:B:275:ARG:HB3	2:B:291:TYR:HB3	1.19	1.13
2:B:400:SER:HB2	2:B:439:CYS:SG	1.87	1.13
3:M:283:PHE:CE2	3:M:289:THR:CB	2.31	1.13
3:M:319:SER:OG	3:M:346:ASN:HB2	0.99	1.13
1:A:637:GLU:HG2	2:B:516:GLY:H	1.11	1.13
2:B:197:LYS:CA	2:B:229:HIS:NE2	2.11	1.13
2:B:260:LEU:HD22	2:B:291:TYR:CZ	1.83	1.13
2:B:563:PHE:CD2	2:B:584:SER:CB	2.32	1.13
3:M:125:PHE:O	3:M:129:VAL:HG23	0.98	1.13
3:M:224:VAL:N	3:M:479:PHE:CG	2.15	1.13
3:M:323:MET:SD	3:M:342:LEU:HG	1.88	1.13
4:S:5:VAL:CG2	4:S:132:LEU:CD2	2.00	1.13
4:S:34:GLN:OE1	4:S:58:LEU:CD1	1.95	1.13
1:A:253:ILE:HD13	1:A:281:LEU:HB3	1.25	1.13
1:A:595:GLU:HG3	2:B:469:ASP:CA	1.61	1.13
2:B:108:PHE:CE2	2:B:115:LEU:CB	2.30	1.13
2:B:513:TRP:CA	2:B:551:LEU:HD22	1.77	1.13
2:B:123:LEU:O	2:B:127:LEU:HG	1.48	1.12
2:B:566:ALA:CA	2:B:574:ASN:CB	2.21	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:584:SER:O	2:B:588:ILE:CG2	1.97	1.13
2:B:139:LEU:CD2	2:B:173:VAL:CA	1.95	1.12
2:B:200:MET:HG2	2:B:232:ARG:HB3	1.23	1.12
2:B:527:PRO:CB	2:B:587:ARG:CG	2.27	1.12
2:B:559:ASP:HB2	2:B:563:PHE:CE1	1.83	1.12
1:A:244:LEU:HD13	1:A:256:LEU:CD1	1.79	1.12
1:A:252:ILE:CA	4:S:144:THR:OG1	1.96	1.12
2:B:174:ALA:CB	2:B:211:ALA:HA	1.79	1.12
2:B:178:ILE:O	2:B:180:LEU:N	1.83	1.12
2:B:181:TYR:CD2	2:B:218:CYS:C	2.15	1.12
2:B:212:VAL:O	2:B:214:ALA:N	1.82	1.12
2:B:461:HIS:O	2:B:462:ASN:C	1.74	1.12
2:B:537:PHE:CZ	2:B:545:ARG:HG2	1.84	1.12
3:M:18:TYR:CD1	3:M:122:SER:CA	2.30	1.12
4:S:48:SER:CB	4:S:77:TYR:O	1.96	1.12
1:A:74:LEU:HD22	1:A:87:CYS:SG	1.90	1.12
1:A:219:VAL:HG11	1:A:256:LEU:HD23	1.31	1.12
1:A:225:LEU:HD13	1:A:233:PHE:CZ	1.83	1.12
1:A:255:ARG:HH22	4:S:135:ILE:CG2	1.58	1.12
2:B:418:TYR:OH	2:B:432:ALA:CB	1.96	1.12
2:B:483:PRO:HB3	2:B:521:ILE:CG2	1.79	1.12
3:M:342:LEU:CD1	3:M:411:LEU:HB2	1.78	1.12
1:A:141:VAL:C	4:S:159:ALA:HB2	1.70	1.12
1:A:215:VAL:HG22	1:A:243:ILE:HD12	1.23	1.12
1:A:225:LEU:CD1	1:A:233:PHE:CZ	2.31	1.12
2:B:230:PHE:CE1	2:B:252:LEU:CD2	2.24	1.12
2:B:393:ILE:HG22	2:B:431:MET:HG2	1.23	1.12
2:B:563:PHE:HD2	2:B:584:SER:HB2	1.05	1.12
1:A:182:ILE:CG2	1:A:221:VAL:HG21	1.79	1.12
1:A:637:GLU:CG	2:B:516:GLY:N	2.12	1.12
2:B:25:VAL:CG2	2:B:35:TYR:HD2	1.61	1.12
2:B:336:ASN:O	2:B:337:THR:HB	1.50	1.12
2:B:353:GLN:CD	3:M:47:SER:CB	2.18	1.12
2:B:549:LEU:HD11	2:B:611:ALA:CA	1.78	1.12
3:M:243:ILE:N	3:M:474:THR:HG22	1.64	1.12
3:M:272:LEU:HD22	3:M:278:ILE:HB	1.20	1.12
1:A:128:LEU:HD12	1:A:150:LEU:HD21	1.21	1.11
1:A:251:TRP:CZ2	4:S:103:GLN:CB	2.32	1.11
2:B:127:LEU:CG	2:B:157:THR:HG21	1.78	1.11
2:B:178:ILE:CG1	2:B:215:TYR:N	2.13	1.11
2:B:353:GLN:HG2	3:M:49:ASP:N	1.64	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:437:SER:HA	2:B:478:LEU:HD21	1.14	1.11
2:B:549:LEU:CD1	2:B:611:ALA:CB	2.27	1.11
3:M:223:HIS:CA	3:M:479:PHE:CD2	2.33	1.11
3:M:265:ASN:OD1	3:M:313:SER:OG	1.67	1.11
3:M:432:THR:OG1	3:M:480:GLN:HG3	0.94	1.11
4:S:53:THR:O	4:S:69:ASN:CB	1.97	1.11
1:A:99:LYS:NZ	4:S:164:ASP:HB2	1.65	1.11
2:B:144:ASP:HA	2:B:179:LYS:NZ	1.63	1.11
2:B:219:TYR:CD2	2:B:226:LEU:HB2	1.84	1.11
2:B:234:CYS:O	2:B:237:ILE:HG22	1.49	1.11
3:M:45:SER:HB2	3:M:51:LEU:HD11	1.18	1.11
1:A:102:GLN:HB2	4:S:163:THR:HB	1.15	1.11
1:A:217:ALA:HB1	4:S:142:ILE:CG1	1.81	1.11
1:A:403:LEU:HD22	1:A:422:GLU:HG3	1.16	1.11
2:B:352:ASN:HB3	3:M:49:ASP:OD2	1.42	1.11
2:B:472:VAL:CG1	2:B:510:GLY:CA	2.25	1.11
2:B:483:PRO:CB	2:B:521:ILE:HG21	1.81	1.11
3:M:18:TYR:CD1	3:M:122:SER:HA	1.66	1.11
4:S:73:ILE:HG23	4:S:88:ILE:CG2	1.77	1.11
2:B:219:TYR:CG	2:B:226:LEU:HD22	1.83	1.11
2:B:319:LEU:HD11	2:B:358:MET:CG	1.70	1.11
3:M:219:LEU:HD22	3:M:473:LYS:HA	1.13	1.11
1:A:103:LYS:CB	4:S:163:THR:HG21	1.80	1.11
1:A:204:VAL:HG22	1:A:236:LEU:HD21	1.18	1.11
2:B:24:ALA:CB	2:B:35:TYR:CE2	2.32	1.11
2:B:106:LEU:HD12	2:B:144:ASP:O	1.43	1.11
2:B:307:ASN:ND2	2:B:336:ASN:ND2	1.98	1.11
2:B:334:MET:HA	2:B:334:MET:HE2	1.30	1.11
1:A:408:ILE:CG2	4:S:64:ASN:CA	2.28	1.10
2:B:37:TYR:HH	2:B:46:GLN:CD	1.47	1.10
2:B:121:ASN:OD1	2:B:153:ILE:CD1	1.97	1.10
2:B:127:LEU:HB3	2:B:161:LEU:CD1	1.80	1.10
2:B:343:LEU:HD22	2:B:362:ALA:CB	1.80	1.10
3:M:319:SER:HB3	3:M:346:ASN:CA	1.79	1.10
4:S:53:THR:C	4:S:69:ASN:CB	2.18	1.10
4:S:53:THR:CB	4:S:68:VAL:C	2.18	1.10
1:A:163:ALA:HB2	1:A:195:ALA:CB	1.80	1.10
2:B:25:VAL:HG23	2:B:35:TYR:HD2	0.94	1.10
2:B:37:TYR:CD2	2:B:38:TYR:HD1	1.65	1.10
2:B:212:VAL:HG22	2:B:248:LEU:HD21	1.30	1.10
3:M:221:THR:HB	3:M:474:THR:O	1.50	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:101:GLN:HG2	4:S:167:ILE:HG21	1.32	1.10
1:A:102:GLN:CB	4:S:163:THR:HB	1.72	1.10
2:B:231:ARG:CG	2:B:298:ASP:OD1	1.98	1.10
2:B:267:ASP:HB3	2:B:289:PRO:HD3	1.31	1.10
2:B:497:LEU:HD11	2:B:508:ARG:NH1	1.67	1.10
2:B:549:LEU:HD21	2:B:611:ALA:N	0.78	1.10
1:A:631:SER:HB3	2:B:557:SER:OG	1.45	1.10
2:B:227:HIS:CD2	2:B:292:GLU:OE2	2.05	1.10
3:M:317:MET:CB	3:M:320:ILE:O	1.98	1.10
3:M:339:GLU:HG3	3:M:412:ARG:HG2	1.16	1.10
4:S:5:VAL:CB	4:S:132:LEU:CD2	2.29	1.10
4:S:9:ASN:HD21	4:S:13:GLN:CG	1.64	1.10
1:A:204:VAL:CG2	1:A:236:LEU:CD2	2.26	1.10
1:A:408:ILE:HG23	4:S:64:ASN:CA	1.82	1.10
2:B:2:VAL:HG12	2:B:6:HIS:CD2	1.86	1.10
2:B:25:VAL:HG21	2:B:36:THR:OG1	1.51	1.10
2:B:127:LEU:CG	2:B:157:THR:CG2	2.27	1.10
2:B:139:LEU:HD21	2:B:173:VAL:O	1.49	1.10
2:B:162:VAL:HG21	2:B:195:ILE:CG2	1.62	1.10
2:B:512:VAL:HG11	2:B:548:ILE:HA	1.17	1.10
3:M:217:ASP:HB3	3:M:470:ALA:C	1.67	1.10
1:A:253:ILE:HG12	1:A:281:LEU:CD1	1.80	1.09
1:A:520:GLY:HA3	1:A:558:VAL:CG2	1.81	1.09
2:B:212:VAL:O	2:B:213:LEU:C	1.83	1.09
3:M:101:LEU:HA	3:M:109:LEU:HD11	1.23	1.09
1:A:488:ARG:HG2	1:A:522:PHE:CE2	1.88	1.09
1:A:637:GLU:OE1	2:B:513:TRP:CD1	2.06	1.09
2:B:98:LYS:HZ1	2:B:134:LEU:CB	1.63	1.09
2:B:108:PHE:CZ	2:B:115:LEU:CG	2.34	1.09
2:B:170:ARG:CA	2:B:199:LEU:HD22	1.82	1.09
2:B:178:ILE:HA	2:B:218:CYS:HB2	1.28	1.09
2:B:208:ILE:HD13	2:B:236:ILE:HG23	1.17	1.09
2:B:343:LEU:CD1	2:B:359:LEU:HD13	1.80	1.09
2:B:353:GLN:HG2	3:M:49:ASP:H	1.06	1.09
3:M:92:PHE:HZ	3:M:129:VAL:HG22	0.98	1.09
1:A:142:LYS:N	4:S:159:ALA:CB	2.16	1.09
1:A:254:ILE:HG23	1:A:293:GLU:HG2	1.32	1.09
2:B:20:ARG:HD2	2:B:21:GLU:CB	1.83	1.09
2:B:79:VAL:CG2	2:B:108:PHE:CE1	2.36	1.09
2:B:151:ALA:HA	2:B:180:LEU:HD11	1.16	1.09
2:B:553:ALA:HB2	2:B:614:ILE:HD13	1.11	1.09

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:342:LEU:HD11	3:M:411:LEU:HD22	1.18	1.09
4:S:109:LEU:HD11	4:S:113:PHE:CD1	1.85	1.09
1:A:103:LYS:HG3	4:S:163:THR:CG2	1.82	1.09
1:A:586:GLU:O	1:A:587:ASN:C	1.83	1.09
2:B:219:TYR:OH	2:B:226:LEU:HA	1.50	1.09
2:B:515:PHE:HE2	2:B:529:VAL:HG21	1.15	1.09
2:B:530:LEU:HG	2:B:591:MET:HB3	1.32	1.09
2:B:566:ALA:C	2:B:574:ASN:HD22	1.56	1.09
2:B:568:VAL:O	2:B:571:SER:HB2	1.49	1.09
3:M:443:SER:HB3	3:M:447:ILE:HG13	1.17	1.09
3:M:451:ALA:C	3:M:452:ILE:HD12	1.72	1.09
4:S:5:VAL:CB	4:S:132:LEU:HD21	1.81	1.09
1:A:182:ILE:HD13	1:A:218:ALA:HB2	1.34	1.09
1:A:403:LEU:HD22	1:A:422:GLU:CD	1.70	1.09
2:B:225:LEU:HD13	2:B:283:TYR:HE1	1.05	1.09
2:B:292:GLU:CG	2:B:296:ASP:HB2	1.82	1.09
3:M:267:ILE:HD12	3:M:445:SER:OG	1.52	1.09
1:A:67:LYS:N	4:S:165:SER:HB2	1.68	1.08
1:A:105:VAL:HB	4:S:167:ILE:HD13	1.34	1.08
1:A:141:VAL:HG21	4:S:158:LYS:H	1.03	1.08
1:A:513:ARG:HD2	1:A:550:VAL:CG2	1.81	1.08
2:B:197:LYS:N	2:B:229:HIS:NE2	2.01	1.08
3:M:217:ASP:H	3:M:470:ALA:HB3	1.07	1.08
1:A:132:LEU:HD22	1:A:169:MET:CG	1.82	1.08
1:A:140:VAL:HA	1:A:177:ILE:HG13	1.22	1.08
1:A:215:VAL:HG22	1:A:243:ILE:CD1	1.82	1.08
1:A:298:ILE:HD11	1:A:311:THR:HG21	1.33	1.08
1:A:408:ILE:CG2	4:S:64:ASN:HB3	1.70	1.08
1:A:429:VAL:HB	1:A:469:LEU:CD1	1.82	1.08
1:A:448:GLU:HB2	1:A:487:MET:SD	1.93	1.08
2:B:343:LEU:HD12	2:B:359:LEU:HD13	1.23	1.08
3:M:65:TYR:CE2	3:M:86:PRO:HB3	1.88	1.08
3:M:323:MET:HB3	3:M:340:LEU:HD11	1.21	1.08
3:M:435:LEU:O	3:M:479:PHE:CD2	2.06	1.08
4:S:5:VAL:CG1	4:S:132:LEU:HD22	1.83	1.08
1:A:132:LEU:O	1:A:169:MET:CE	2.00	1.08
1:A:217:ALA:HA	4:S:142:ILE:CA	1.63	1.08
1:A:224:GLU:HB2	4:S:138:GLY:O	0.92	1.08
2:B:107:ARG:O	2:B:110:GLU:N	1.87	1.08
2:B:347:VAL:HG22	2:B:359:LEU:HB3	1.23	1.08
2:B:534:ILE:HD13	2:B:594:ALA:HB3	1.33	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:566:ALA:O	2:B:574:ASN:HB2	0.91	1.08
3:M:245:ASP:N	3:M:472:TYR:CE1	2.20	1.08
1:A:103:LYS:N	4:S:163:THR:CB	2.16	1.08
2:B:56:SER:HB3	2:B:92:THR:HG21	1.31	1.08
2:B:175:LEU:HG	2:B:210:CYS:HB3	1.11	1.08
2:B:215:TYR:HB3	2:B:226:LEU:HD11	1.29	1.08
2:B:400:SER:CB	2:B:439:CYS:SG	2.42	1.08
2:B:497:LEU:HD23	2:B:533:LEU:HD21	1.09	1.08
2:B:567:GLN:HG3	2:B:569:THR:OG1	1.53	1.08
3:M:336:ASP:O	3:M:414:CYS:SG	2.10	1.08
1:A:68:THR:HA	4:S:167:ILE:N	1.67	1.08
1:A:132:LEU:O	1:A:169:MET:HE1	1.52	1.08
1:A:170:LEU:O	1:A:206:LYS:CD	2.00	1.08
1:A:219:VAL:HG21	1:A:256:LEU:CD2	1.82	1.08
1:A:609:LEU:HD21	1:A:628:VAL:CG2	1.83	1.08
2:B:175:LEU:HD21	2:B:210:CYS:HA	1.32	1.08
2:B:216:LYS:HB2	2:B:251:LEU:HD13	1.31	1.08
2:B:307:ASN:ND2	2:B:336:ASN:HD21	1.48	1.08
2:B:319:LEU:HD11	2:B:358:MET:HG3	1.29	1.08
3:M:454:ILE:HG21	3:M:464:THR:CG2	1.84	1.08
4:S:15:ARG:CZ	4:S:122:ILE:HD11	1.84	1.08
1:A:132:LEU:HD22	1:A:169:MET:HG3	1.13	1.07
1:A:182:ILE:HG22	1:A:221:VAL:HG21	1.20	1.07
2:B:21:GLU:HA	2:B:24:ALA:HB2	1.35	1.07
2:B:159:LYS:CA	2:B:195:ILE:HD11	1.84	1.07
2:B:175:LEU:HD11	2:B:210:CYS:SG	1.94	1.07
2:B:279:LEU:HG	2:B:288:TYR:HD1	1.12	1.07
2:B:292:GLU:OE2	2:B:296:ASP:OD2	1.71	1.07
2:B:352:ASN:HA	3:M:49:ASP:OD2	1.37	1.07
2:B:396:ILE:HD13	2:B:432:ALA:HB2	1.36	1.07
2:B:433:VAL:HG12	2:B:474:VAL:CB	1.84	1.07
3:M:254:PRO:HB3	3:M:454:ILE:HD12	1.26	1.07
4:S:53:THR:HG21	4:S:68:VAL:CA	1.83	1.07
2:B:197:LYS:CA	2:B:229:HIS:CD2	2.36	1.07
2:B:208:ILE:HG12	2:B:236:ILE:HD13	1.31	1.07
2:B:247:TYR:CE2	3:M:91:THR:CG2	2.36	1.07
2:B:247:TYR:CZ	3:M:91:THR:HG21	1.87	1.07
2:B:316:THR:OG1	3:M:90:PHE:HZ	1.37	1.07
2:B:458:MET:SD	2:B:471:TYR:HB3	1.94	1.07
3:M:222:PHE:C	3:M:479:PHE:CE2	2.28	1.07
1:A:251:TRP:CZ2	4:S:103:GLN:HB2	1.90	1.07

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:322:PHE:CD2	1:A:330:LEU:HD21	1.88	1.07
1:A:509:PRO:HB3	1:A:547:VAL:CG2	1.84	1.07
1:A:555:LEU:HD13	1:A:581:LEU:CD1	1.84	1.07
1:A:633:PHE:CB	2:B:550:VAL:CG1	2.22	1.07
1:A:633:PHE:HB2	2:B:550:VAL:CG1	1.80	1.07
2:B:123:LEU:HD12	2:B:142:LEU:CG	1.79	1.07
2:B:278:PRO:HD3	2:B:289:PRO:O	1.52	1.07
2:B:353:GLN:CB	3:M:49:ASP:N	2.09	1.07
3:M:217:ASP:HB2	3:M:470:ALA:C	1.67	1.07
3:M:217:ASP:N	3:M:470:ALA:HB3	1.70	1.07
3:M:219:LEU:HB2	3:M:472:TYR:O	1.54	1.07
4:S:53:THR:O	4:S:69:ASN:HB2	1.50	1.07
4:S:73:ILE:HG21	4:S:88:ILE:HG21	1.33	1.07
2:B:278:PRO:CA	2:B:288:TYR:O	2.00	1.07
4:S:5:VAL:HG11	4:S:132:LEU:HD22	1.09	1.07
1:A:166:LEU:HD13	1:A:185:LEU:CD2	1.85	1.07
1:A:384:LEU:HD22	1:A:441:TYR:CZ	1.88	1.07
1:A:384:LEU:HD13	1:A:435:ILE:CG2	1.85	1.07
2:B:154:ILE:HD12	2:B:180:LEU:HD13	1.12	1.07
2:B:274:PRO:C	2:B:295:ASN:ND2	2.08	1.07
2:B:290:SER:O	2:B:291:TYR:C	1.75	1.07
2:B:367:SER:OG	2:B:401:THR:HG21	1.55	1.07
2:B:549:LEU:HD23	2:B:607:ILE:O	1.54	1.07
2:B:560:ILE:HG23	2:B:564:LYS:HB2	1.35	1.07
1:A:207:LEU:HD23	1:A:239:LEU:HB2	1.29	1.06
1:A:237:SER:HB2	1:A:270:LEU:HD13	1.37	1.06
1:A:594:PHE:HB3	2:B:473:ASN:HB3	1.32	1.06
2:B:77:ILE:HG22	2:B:82:TYR:HE1	1.18	1.06
2:B:120:ILE:HA	2:B:142:LEU:HD21	1.34	1.06
2:B:132:SER:HA	2:B:169:VAL:CG2	1.85	1.06
2:B:158:VAL:HG11	2:B:177:ILE:HG13	1.34	1.06
2:B:219:TYR:CE1	2:B:226:LEU:CA	2.27	1.06
2:B:515:PHE:CE2	2:B:529:VAL:HG21	1.89	1.06
2:B:537:PHE:CE1	2:B:545:ARG:HG2	1.91	1.06
3:M:45:SER:CB	3:M:51:LEU:HD11	1.85	1.06
4:S:53:THR:HG21	4:S:68:VAL:HA	1.36	1.06
2:B:175:LEU:HD23	2:B:210:CYS:O	1.53	1.06
2:B:291:TYR:HE2	2:B:294:VAL:HB	0.90	1.06
2:B:343:LEU:HD11	2:B:359:LEU:HA	1.06	1.06
2:B:353:GLN:CD	3:M:47:SER:HB2	1.76	1.06
3:M:215:TYR:CG	3:M:468:LYS:CA	2.38	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:67:LYS:HB3	1:A:94:VAL:HG22	1.32	1.06
1:A:80:TYR:HB2	1:A:82:PHE:CD2	1.90	1.06
1:A:204:VAL:HG22	1:A:236:LEU:HD22	1.21	1.06
1:A:211:ASP:OD2	4:S:148:ARG:NH1	1.89	1.06
2:B:225:LEU:CD1	2:B:283:TYR:CE1	2.36	1.06
2:B:303:LEU:HD11	2:B:333:GLN:HB3	1.31	1.06
2:B:527:PRO:CG	2:B:587:ARG:HG3	1.85	1.06
3:M:374:TYR:O	3:M:390:ILE:HD12	1.51	1.06
1:A:295:VAL:HG22	1:A:315:CYS:HB3	1.25	1.06
2:B:29:LYS:CE	2:B:30:LEU:N	2.11	1.06
2:B:127:LEU:HD23	2:B:161:LEU:HD21	1.07	1.06
2:B:178:ILE:CG2	2:B:217:GLU:HB2	1.85	1.06
2:B:274:PRO:CG	2:B:295:ASN:OD1	2.03	1.06
2:B:353:GLN:CG	3:M:47:SER:O	2.02	1.06
3:M:302:TYR:CD2	3:M:445:SER:HB3	1.90	1.06
3:M:338:PHE:CE2	3:M:415:ILE:CG1	2.39	1.06
1:A:207:LEU:CD2	1:A:239:LEU:HB2	1.85	1.06
1:A:215:VAL:HG21	1:A:243:ILE:HD12	1.31	1.06
1:A:224:GLU:HB2	4:S:138:GLY:C	1.75	1.06
1:A:258:LYS:HZ1	4:S:97:ALA:HB2	1.20	1.06
2:B:136:CYS:C	2:B:172:GLU:HG3	1.75	1.06
2:B:177:ILE:HB	2:B:196:LEU:HD21	1.37	1.06
2:B:256:CYS:SG	2:B:299:LEU:HD23	1.96	1.06
2:B:293:VAL:O	2:B:299:LEU:CB	2.04	1.06
2:B:319:LEU:HD13	2:B:358:MET:HG3	1.09	1.06
2:B:437:SER:CA	2:B:478:LEU:HD21	1.86	1.06
2:B:508:ARG:O	2:B:512:VAL:HG23	1.54	1.06
2:B:527:PRO:HB3	2:B:587:ARG:HG3	1.32	1.06
1:A:533:ILE:HG12	1:A:562:TRP:CH2	1.91	1.05
2:B:25:VAL:CG2	2:B:36:THR:OG1	2.04	1.05
2:B:219:TYR:OH	2:B:226:LEU:N	1.87	1.05
2:B:433:VAL:CG1	2:B:474:VAL:CG2	2.12	1.05
2:B:546:CYS:CA	2:B:607:ILE:HG12	1.85	1.05
3:M:226:PHE:N	3:M:480:GLN:O	1.89	1.05
3:M:360:LEU:HD23	3:M:362:PHE:CZ	1.90	1.05
1:A:225:LEU:HD13	1:A:233:PHE:HZ	0.99	1.05
1:A:630:PRO:O	2:B:554:LYS:CA	2.05	1.05
2:B:79:VAL:CB	2:B:108:PHE:CE1	2.38	1.05
2:B:116:THR:HG22	2:B:150:LEU:HD11	1.35	1.05
2:B:549:LEU:CD1	2:B:611:ALA:CA	2.31	1.05
3:M:60:LEU:CD2	3:M:62:VAL:HG23	1.86	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:66:PHE:HA	3:M:77:LEU:HD11	1.38	1.05
3:M:244:VAL:O	3:M:299:LEU:N	1.88	1.05
1:A:67:LYS:H	4:S:165:SER:CB	1.68	1.05
1:A:166:LEU:CD1	1:A:185:LEU:CD2	2.33	1.05
1:A:200:PHE:CE1	1:A:232:PRO:O	2.09	1.05
1:A:260:PHE:CD2	1:A:274:LEU:HG	1.91	1.05
1:A:629:LEU:CG	2:B:610:ARG:NH1	2.18	1.05
1:A:630:PRO:HG3	2:B:614:ILE:HG13	1.27	1.05
1:A:633:PHE:CD1	2:B:513:TRP:CE3	2.40	1.05
2:B:77:ILE:HG22	2:B:82:TYR:CE1	1.92	1.05
2:B:566:ALA:CA	2:B:574:ASN:CG	2.25	1.05
3:M:348:LYS:HG3	3:M:405:THR:HG22	1.38	1.05
1:A:557:LYS:HE3	2:B:606:ASP:HB2	1.38	1.05
1:A:631:SER:CA	2:B:557:SER:HB3	1.82	1.05
2:B:286:ILE:O	2:B:287:GLU:O	1.73	1.05
2:B:315:PRO:CB	2:B:352:ASN:OD1	2.04	1.05
3:M:341:SER:OG	3:M:343:ASN:ND2	1.88	1.05
3:M:342:LEU:HD13	3:M:411:LEU:HB2	1.38	1.05
1:A:101:GLN:HG2	4:S:167:ILE:CG2	1.86	1.05
1:A:176:TYR:HB2	4:S:155:GLU:HG3	1.33	1.05
2:B:20:ARG:HD2	2:B:21:GLU:HB2	1.15	1.05
2:B:245:GLN:CG	2:B:309:LEU:HD11	1.85	1.05
2:B:309:LEU:HB3	2:B:317:VAL:HG12	1.07	1.05
2:B:545:ARG:CD	2:B:602:ASP:HB2	1.86	1.05
2:B:566:ALA:C	2:B:574:ASN:ND2	2.11	1.05
3:M:64:LYS:NZ	3:M:79:SER:O	1.90	1.05
3:M:244:VAL:HB	3:M:300:LEU:HG	1.39	1.05
4:S:8:PHE:HB2	4:S:36:TYR:HE2	0.90	1.05
1:A:182:ILE:HG22	1:A:221:VAL:HG23	1.33	1.04
1:A:260:PHE:O	1:A:261:THR:C	1.78	1.04
1:A:440:ASN:CG	1:A:442:SER:HB3	1.77	1.04
2:B:2:VAL:CG1	2:B:6:HIS:CE1	2.36	1.04
2:B:20:ARG:HD2	2:B:35:TYR:OH	1.56	1.04
2:B:154:ILE:HD13	2:B:180:LEU:HB2	1.37	1.04
2:B:162:VAL:HG22	2:B:199:LEU:HG	1.11	1.04
2:B:219:TYR:HE1	2:B:226:LEU:CD1	1.51	1.04
2:B:223:LEU:HD11	2:B:258:GLN:C	1.77	1.04
2:B:563:PHE:O	2:B:567:GLN:N	1.88	1.04
3:M:49:ASP:HA	3:M:75:TRP:CH2	1.91	1.04
1:A:67:LYS:O	1:A:71:VAL:HG23	1.58	1.04
1:A:182:ILE:CG2	1:A:221:VAL:CG2	2.33	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:630:PRO:HG2	2:B:614:ILE:CA	1.87	1.04
1:A:633:PHE:CD2	2:B:551:LEU:HA	1.54	1.04
2:B:158:VAL:HG13	2:B:173:VAL:HG12	1.37	1.04
2:B:162:VAL:HG21	2:B:195:ILE:HG22	1.34	1.04
2:B:178:ILE:HG23	2:B:217:GLU:HB2	1.38	1.04
2:B:215:TYR:HB3	2:B:226:LEU:CD1	1.87	1.04
2:B:278:PRO:HA	2:B:288:TYR:CB	1.87	1.04
2:B:396:ILE:HG12	2:B:418:TYR:CE2	1.93	1.04
3:M:327:PHE:HE1	3:M:336:ASP:HB2	1.19	1.04
4:S:53:THR:HB	4:S:68:VAL:C	1.77	1.04
1:A:71:VAL:HG11	1:A:105:VAL:HG12	1.36	1.04
1:A:147:LEU:HD22	1:A:166:LEU:HD21	1.07	1.04
2:B:162:VAL:HG22	2:B:199:LEU:CD1	1.87	1.04
2:B:537:PHE:CE2	2:B:598:LEU:O	2.11	1.04
2:B:567:GLN:CG	2:B:569:THR:OG1	2.05	1.04
3:M:215:TYR:HD1	3:M:467:TYR:O	1.39	1.04
3:M:245:ASP:N	3:M:472:TYR:CG	2.09	1.04
1:A:150:LEU:HB3	1:A:162:ILE:HD13	1.37	1.04
1:A:557:LYS:HE2	2:B:606:ASP:CA	1.86	1.04
1:A:557:LYS:HE3	2:B:606:ASP:CB	1.86	1.04
1:A:594:PHE:CB	2:B:473:ASN:CB	2.34	1.04
2:B:248:LEU:O	2:B:252:LEU:HG	1.58	1.04
2:B:340:ILE:HD11	2:B:366:LEU:HB3	1.38	1.04
2:B:542:PRO:HA	2:B:602:ASP:CG	1.76	1.04
3:M:244:VAL:HG13	3:M:472:TYR:CZ	1.92	1.04
1:A:96:SER:HB3	1:A:127:LEU:CD1	1.86	1.04
1:A:251:TRP:CH2	4:S:103:GLN:CB	2.41	1.04
1:A:636:TYR:HB2	2:B:554:LYS:HZ2	1.19	1.04
2:B:41:ASN:CB	2:B:43:ASN:OD1	2.04	1.04
2:B:261:PRO:CA	2:B:290:SER:CB	2.35	1.04
2:B:303:LEU:HD11	2:B:333:GLN:CB	1.84	1.04
2:B:309:LEU:CB	2:B:317:VAL:CG1	2.35	1.04
2:B:519:ALA:O	2:B:523:PHE:CD2	2.11	1.04
3:M:222:PHE:HB2	3:M:479:PHE:CZ	1.92	1.04
3:M:223:HIS:CA	3:M:479:PHE:CE2	2.41	1.04
3:M:226:PHE:HZ	3:M:321:GLY:O	1.41	1.04
1:A:103:LYS:H	4:S:163:THR:HG21	1.23	1.03
1:A:104:ARG:CA	1:A:145:ILE:HG21	1.88	1.03
1:A:105:VAL:HG21	4:S:167:ILE:HG23	1.40	1.03
2:B:178:ILE:HD11	2:B:215:TYR:HA	1.07	1.03
2:B:267:ASP:C	2:B:276:SER:HB2	1.79	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:340:ILE:HD13	2:B:366:LEU:HD13	1.32	1.03
2:B:343:LEU:HD21	2:B:362:ALA:HB3	1.35	1.03
2:B:537:PHE:CG	2:B:598:LEU:HB3	1.92	1.03
3:M:214:LEU:O	3:M:467:TYR:N	1.89	1.03
1:A:251:TRP:CZ2	4:S:103:GLN:HB3	1.93	1.03
2:B:158:VAL:CG1	2:B:177:ILE:HG13	1.86	1.03
2:B:175:LEU:HD21	2:B:210:CYS:CA	1.88	1.03
2:B:343:LEU:HD12	2:B:359:LEU:HD12	1.33	1.03
2:B:493:LEU:HG	2:B:511:ILE:HG23	1.40	1.03
2:B:530:LEU:CG	2:B:591:MET:HB3	1.88	1.03
3:M:319:SER:HB3	3:M:346:ASN:H	0.98	1.03
3:M:449:VAL:HG12	3:M:452:ILE:HD11	1.36	1.03
1:A:323:CYS:SG	1:A:334:SER:HB3	1.97	1.03
1:A:397:ASP:O	1:A:418:ILE:HG13	1.59	1.03
2:B:154:ILE:CD1	2:B:180:LEU:HB2	1.87	1.03
2:B:175:LEU:HA	2:B:214:ALA:HB2	1.35	1.03
2:B:508:ARG:CB	2:B:544:THR:HG23	1.87	1.03
3:M:60:LEU:HD23	3:M:61:GLU:N	1.71	1.03
3:M:243:ILE:O	3:M:472:TYR:CD2	2.12	1.03
3:M:290:PHE:CZ	3:M:297:PHE:CG	2.47	1.03
3:M:319:SER:CB	3:M:346:ASN:H	1.70	1.03
1:A:68:THR:HA	4:S:166:LYS:C	1.76	1.03
1:A:140:VAL:CG2	1:A:177:ILE:HG13	1.87	1.03
2:B:424:PHE:CD2	2:B:428:VAL:HG11	1.93	1.03
2:B:542:PRO:CA	2:B:602:ASP:OD2	2.07	1.03
3:M:2:TYR:N	3:M:81:SER:CB	2.22	1.03
3:M:224:VAL:O	3:M:480:GLN:N	1.91	1.03
3:M:243:ILE:H	3:M:474:THR:HG22	0.87	1.03
1:A:132:LEU:HD13	1:A:165:ASP:HB3	1.40	1.03
1:A:176:TYR:CB	4:S:155:GLU:CG	2.35	1.03
1:A:183:THR:O	1:A:186:PHE:HB3	1.59	1.03
1:A:398:GLU:O	1:A:420:ILE:N	1.90	1.03
1:A:594:PHE:HB3	2:B:473:ASN:HB2	1.09	1.03
1:A:605:GLU:HG3	1:A:632:PHE:CE2	1.93	1.03
2:B:120:ILE:HA	2:B:142:LEU:CD2	1.88	1.03
2:B:227:HIS:C	2:B:229:HIS:H	1.59	1.03
2:B:371:GLN:NE2	2:B:401:THR:O	1.90	1.03
1:A:96:SER:HA	1:A:127:LEU:HD11	1.39	1.02
1:A:140:VAL:HG22	1:A:177:ILE:HG13	1.33	1.02
1:A:225:LEU:HB3	1:A:233:PHE:CE1	1.94	1.02
1:A:323:CYS:SG	1:A:334:SER:CB	2.47	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:135:ARG:NH2	2:B:164:ASP:CG	2.12	1.02
2:B:215:TYR:CZ	2:B:229:HIS:HB3	1.94	1.02
2:B:567:GLN:N	2:B:574:ASN:HD22	1.55	1.02
3:M:214:LEU:HD11	3:M:256:VAL:HG21	1.41	1.02
3:M:242:GLY:HA2	3:M:474:THR:HG21	1.40	1.02
1:A:96:SER:HB2	1:A:127:LEU:HD11	1.39	1.02
1:A:111:SER:HB2	1:A:152:THR:OG1	0.86	1.02
1:A:219:VAL:HG22	1:A:240:LEU:HD22	1.42	1.02
1:A:463:ASP:O	2:B:1:MET:SD	2.16	1.02
1:A:563:CYS:HA	1:A:566:PHE:HD2	1.23	1.02
1:A:630:PRO:CG	2:B:614:ILE:CG1	1.92	1.02
2:B:243:TRP:CZ3	3:M:91:THR:HA	1.94	1.02
2:B:553:ALA:HB2	2:B:614:ILE:CG1	1.88	1.02
2:B:564:LYS:HD2	2:B:621:GLY:O	1.58	1.02
2:B:588:ILE:HG23	2:B:618:PHE:CZ	1.93	1.02
3:M:243:ILE:HB	3:M:473:LYS:O	1.55	1.02
4:S:8:PHE:HB3	4:S:36:TYR:CE2	1.92	1.02
4:S:17:VAL:HG22	4:S:19:PHE:CZ	1.94	1.02
1:A:102:GLN:HA	4:S:167:ILE:CG1	1.88	1.02
1:A:102:GLN:OE1	4:S:166:LYS:HG3	1.58	1.02
1:A:217:ALA:O	4:S:142:ILE:CG2	2.06	1.02
1:A:251:TRP:CH2	4:S:103:GLN:OE1	2.11	1.02
2:B:106:LEU:CD1	2:B:144:ASP:HB2	1.74	1.02
2:B:418:TYR:O	2:B:419:VAL:C	1.88	1.02
3:M:7:ILE:HA	3:M:76:CYS:HA	1.40	1.02
3:M:65:TYR:CE1	3:M:86:PRO:CB	2.43	1.02
1:A:141:VAL:HB	4:S:159:ALA:N	1.73	1.02
2:B:127:LEU:HB2	2:B:157:THR:HG23	1.39	1.02
2:B:143:SER:C	2:B:179:LYS:HD3	1.70	1.02
2:B:219:TYR:CB	2:B:223:LEU:CD2	2.14	1.02
2:B:267:ASP:H	2:B:289:PRO:HB3	0.88	1.02
2:B:343:LEU:HD22	2:B:362:ALA:HB3	1.33	1.02
4:S:53:THR:CG2	4:S:67:GLU:C	2.27	1.02
1:A:67:LYS:HB2	4:S:165:SER:OG	1.59	1.02
1:A:96:SER:CA	1:A:127:LEU:CD1	2.38	1.02
1:A:200:PHE:HE1	1:A:232:PRO:O	1.39	1.02
1:A:223:CYS:HB2	1:A:259:LEU:HG	1.42	1.02
2:B:38:TYR:HA	2:B:42:ILE:H	1.23	1.02
2:B:70:MET:HE3	2:B:107:ARG:CB	1.82	1.02
2:B:79:VAL:CG2	2:B:108:PHE:CZ	2.43	1.02
2:B:278:PRO:C	2:B:288:TYR:HB2	1.78	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:537:PHE:CD2	2:B:598:LEU:CB	2.43	1.02
4:S:48:SER:CB	4:S:77:TYR:HB2	1.86	1.02
2:B:56:SER:HB3	2:B:92:THR:CG2	1.89	1.01
2:B:62:ALA:C	2:B:66:ILE:HD12	1.80	1.01
2:B:155:LEU:HB2	2:B:188:TYR:HD2	1.22	1.01
2:B:261:PRO:HA	2:B:290:SER:HB3	1.03	1.01
2:B:415:LEU:HD13	2:B:436:LEU:HD21	1.39	1.01
2:B:550:VAL:HG22	2:B:610:ARG:HD3	1.37	1.01
3:M:344:ILE:HG23	3:M:347:PHE:CB	1.89	1.01
1:A:101:GLN:HG2	4:S:167:ILE:CB	1.90	1.01
1:A:104:ARG:HG3	1:A:145:ILE:CG1	1.89	1.01
1:A:217:ALA:O	4:S:142:ILE:CB	2.00	1.01
1:A:253:ILE:CG1	1:A:281:LEU:HD13	1.90	1.01
2:B:70:MET:HE1	2:B:104:TYR:O	1.59	1.01
2:B:178:ILE:HG21	2:B:214:ALA:O	1.60	1.01
2:B:219:TYR:O	2:B:223:LEU:CD2	2.08	1.01
2:B:293:VAL:O	2:B:299:LEU:HB2	1.58	1.01
2:B:513:TRP:HA	2:B:551:LEU:HD22	1.02	1.01
3:M:101:LEU:CG	3:M:106:LYS:CA	2.17	1.01
3:M:212:ASN:CB	3:M:250:LEU:HD23	1.90	1.01
3:M:256:VAL:HG12	3:M:290:PHE:HB3	1.39	1.01
3:M:342:LEU:CD1	3:M:411:LEU:CB	2.37	1.01
4:S:56:SER:O	4:S:60:SER:HB2	1.61	1.01
1:A:253:ILE:CD1	1:A:281:LEU:HD22	1.88	1.01
2:B:171:GLY:HA3	2:B:207:VAL:HG13	1.04	1.01
2:B:182:ARG:HD2	2:B:217:GLU:OE1	1.61	1.01
2:B:549:LEU:HD13	2:B:611:ALA:CB	1.89	1.01
2:B:562:ASN:HB3	2:B:580:TYR:HB3	1.39	1.01
3:M:443:SER:OG	3:M:447:ILE:O	1.77	1.01
1:A:147:LEU:HD22	1:A:166:LEU:HD23	1.35	1.01
1:A:295:VAL:HG11	1:A:319:LEU:HD11	1.42	1.01
1:A:398:GLU:O	1:A:420:ILE:HG13	1.61	1.01
1:A:426:ILE:HG13	1:A:464:ILE:HD13	1.40	1.01
2:B:120:ILE:CG2	2:B:154:ILE:HG13	1.90	1.01
2:B:174:ALA:HB1	2:B:211:ALA:HA	1.40	1.01
2:B:291:TYR:CE2	2:B:294:VAL:CB	2.43	1.01
2:B:513:TRP:CG	2:B:551:LEU:HD21	1.96	1.01
2:B:559:ASP:HB3	2:B:563:PHE:CD1	1.93	1.01
3:M:257:ALA:O	3:M:452:ILE:HG23	1.58	1.01
3:M:378:ILE:O	3:M:413:GLY:CA	2.07	1.01
1:A:105:VAL:HG23	4:S:167:ILE:CG1	1.90	1.01

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:136:GLY:O	1:A:139:ASP:CB	2.09	1.01
1:A:516:ILE:HD13	1:A:551:LEU:HA	1.42	1.01
2:B:178:ILE:CD1	2:B:215:TYR:HA	1.89	1.01
3:M:16:PHE:CZ	3:M:18:TYR:HB2	1.96	1.01
3:M:339:GLU:CG	3:M:412:ARG:HG2	1.90	1.01
3:M:379:LEU:CD2	3:M:386:PHE:HB2	1.90	1.01
4:S:48:SER:CA	4:S:77:TYR:HB2	1.91	1.01
2:B:77:ILE:CG2	2:B:82:TYR:HE1	1.73	1.00
2:B:127:LEU:HB3	2:B:157:THR:HG23	1.04	1.00
2:B:178:ILE:HD11	2:B:215:TYR:CA	1.91	1.00
2:B:230:PHE:O	2:B:231:ARG:O	1.79	1.00
2:B:303:LEU:HD11	2:B:333:GLN:CG	1.91	1.00
2:B:393:ILE:CG2	2:B:431:MET:CB	2.38	1.00
2:B:497:LEU:CD2	2:B:533:LEU:CD2	2.38	1.00
2:B:508:ARG:HB2	2:B:544:THR:CG2	1.91	1.00
2:B:508:ARG:HB2	2:B:544:THR:HG23	1.37	1.00
4:S:53:THR:HG22	4:S:57:LEU:HB2	1.39	1.00
1:A:288:THR:CG2	1:A:322:PHE:HZ	1.71	1.00
1:A:429:VAL:CG1	1:A:469:LEU:HD11	1.90	1.00
2:B:77:ILE:CG2	2:B:82:TYR:CE1	2.44	1.00
2:B:127:LEU:HB3	2:B:161:LEU:HD11	1.39	1.00
2:B:223:LEU:HD23	2:B:255:TYR:CE1	1.93	1.00
2:B:226:LEU:HB3	2:B:255:TYR:CZ	1.85	1.00
3:M:344:ILE:CG2	3:M:347:PHE:HB3	1.90	1.00
4:S:73:ILE:HG22	4:S:88:ILE:HG23	1.41	1.00
4:S:80:TYR:O	4:S:81:ALA:C	1.98	1.00
1:A:178:ARG:CD	1:A:209:ASP:OD2	2.08	1.00
1:A:254:ILE:CG1	1:A:290:VAL:HG22	1.91	1.00
2:B:108:PHE:CE2	2:B:115:LEU:CG	2.43	1.00
2:B:185:LYS:HG3	2:B:222:HIS:CE1	1.96	1.00
2:B:189:HIS:NE2	2:B:193:LEU:HD11	1.75	1.00
2:B:393:ILE:HG23	2:B:431:MET:HB3	1.42	1.00
2:B:404:ASN:O	2:B:405:GLU:C	1.80	1.00
2:B:447:GLU:OE1	2:B:485:LYS:HG3	1.61	1.00
3:M:327:PHE:CE1	3:M:336:ASP:HB2	1.97	1.00
3:M:339:GLU:CG	3:M:412:ARG:NE	2.23	1.00
1:A:104:ARG:HG3	1:A:145:ILE:HG13	1.41	1.00
1:A:104:ARG:CG	1:A:145:ILE:HG13	1.91	1.00
1:A:141:VAL:HG21	4:S:158:LYS:N	1.77	1.00
1:A:244:LEU:CD2	1:A:277:LYS:O	2.10	1.00
1:A:408:ILE:HG21	4:S:64:ASN:C	1.82	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:546:CYS:SG	2:B:607:ILE:CG1	2.40	1.00
3:M:290:PHE:CZ	3:M:297:PHE:CD1	2.49	1.00
1:A:217:ALA:HB2	4:S:142:ILE:HG13	1.43	1.00
1:A:408:ILE:HG23	4:S:64:ASN:HB3	1.01	1.00
2:B:158:VAL:HG12	2:B:195:ILE:HG21	1.40	1.00
3:M:217:ASP:OD1	3:M:471:LYS:HA	1.58	1.00
3:M:353:VAL:HG23	3:M:438:SER:O	1.61	1.00
1:A:166:LEU:HD12	1:A:185:LEU:HD23	1.41	1.00
1:A:220:SER:CB	4:S:142:ILE:CG2	2.32	1.00
1:A:528:ASN:O	1:A:529:GLY:C	1.92	1.00
1:A:633:PHE:HE1	2:B:513:TRP:CE3	1.33	1.00
2:B:380:LYS:NZ	3:M:236:LEU:HG	1.76	1.00
2:B:487:LEU:HD21	2:B:522:GLU:HB3	1.43	1.00
3:M:244:VAL:HA	3:M:472:TYR:HD2	1.26	1.00
3:M:379:LEU:HD23	3:M:386:PHE:HB2	1.41	1.00
2:B:182:ARG:HD2	2:B:217:GLU:HB3	1.41	1.00
3:M:65:TYR:CZ	3:M:86:PRO:CB	2.44	1.00
3:M:69:ILE:HG13	3:M:90:PHE:CE1	1.97	1.00
1:A:429:VAL:HB	1:A:469:LEU:HD11	1.03	0.99
1:A:629:LEU:CG	2:B:610:ARG:HH11	1.74	0.99
2:B:120:ILE:HG23	2:B:154:ILE:HG13	1.43	0.99
3:M:326:HIS:O	3:M:338:PHE:HA	1.61	0.99
4:S:131:VAL:HG22	4:S:153:VAL:HG22	1.40	0.99
2:B:80:GLN:HG2	2:B:115:LEU:CD1	1.91	0.99
2:B:303:LEU:HD13	2:B:333:GLN:HB3	1.05	0.99
2:B:307:ASN:OD1	2:B:339:PHE:CE2	2.15	0.99
2:B:513:TRP:HA	2:B:551:LEU:CD1	1.83	0.99
1:A:275:LEU:CD1	1:A:308:ASP:CG	2.31	0.99
1:A:388:VAL:HG13	1:A:432:ILE:HD12	1.03	0.99
2:B:106:LEU:HD11	2:B:144:ASP:HB3	1.02	0.99
3:M:101:LEU:HD21	3:M:106:LYS:O	1.62	0.99
2:B:63:MET:HG3	2:B:100:LEU:HB3	1.44	0.99
2:B:219:TYR:OH	2:B:226:LEU:CA	2.05	0.99
3:M:339:GLU:CG	3:M:412:ARG:HE	1.75	0.99
1:A:101:GLN:NE2	4:S:167:ILE:CG2	2.25	0.99
2:B:37:TYR:CE2	2:B:38:TYR:CE1	2.50	0.99
2:B:162:VAL:CG2	2:B:199:LEU:CD1	2.40	0.99
2:B:523:PHE:CD1	2:B:559:ASP:OD1	2.16	0.99
2:B:549:LEU:CD2	2:B:607:ILE:O	2.10	0.99
3:M:245:ASP:CA	3:M:472:TYR:CD1	2.45	0.99
3:M:290:PHE:CE1	3:M:297:PHE:CE1	2.50	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:34:GLN:OE1	4:S:58:LEU:CG	2.09	0.99
1:A:88:ASN:O	1:A:89:PHE:C	1.91	0.99
1:A:128:LEU:CD1	1:A:150:LEU:HD23	1.87	0.99
1:A:638:LEU:HD12	2:B:518:ILE:HG23	1.42	0.99
2:B:121:ASN:OD1	2:B:153:ILE:HD11	1.59	0.99
2:B:159:LYS:HA	2:B:195:ILE:HD11	1.00	0.99
2:B:490:ILE:CG2	2:B:515:PHE:CE2	2.44	0.99
2:B:519:ALA:O	2:B:523:PHE:CG	2.15	0.99
4:S:6:LEU:HD22	4:S:32:LEU:HD22	1.42	0.99
1:A:189:PHE:CD2	1:A:225:LEU:HD21	1.96	0.99
2:B:225:LEU:CD1	2:B:283:TYR:CZ	2.45	0.99
2:B:277:CYS:SG	2:B:292:GLU:CG	2.50	0.99
2:B:592:TYR:OH	2:B:619:ASP:OD2	1.81	0.99
1:A:264:SER:HB2	1:A:271:ARG:CD	1.93	0.99
1:A:275:LEU:CD1	1:A:308:ASP:OD1	2.11	0.99
1:A:287:ALA:HB3	1:A:289:SER:N	1.76	0.99
1:A:563:CYS:HB3	1:A:621:LEU:HD12	1.41	0.99
2:B:279:LEU:HG	2:B:288:TYR:CD1	1.96	0.99
1:A:384:LEU:HD22	1:A:441:TYR:HE2	1.27	0.99
2:B:83:PHE:HZ	2:B:119:SER:CB	1.51	0.99
2:B:86:VAL:HG12	2:B:101:ILE:HG23	1.01	0.99
2:B:243:TRP:HZ3	3:M:91:THR:HA	1.24	0.99
2:B:530:LEU:HD11	2:B:595:VAL:CG2	1.92	0.99
3:M:69:ILE:HD11	3:M:90:PHE:CZ	1.66	0.99
1:A:179:LYS:CE	4:S:143:GLU:CB	2.36	0.99
1:A:217:ALA:CB	4:S:142:ILE:HG13	1.90	0.99
2:B:303:LEU:HD11	2:B:333:GLN:NE2	1.77	0.99
2:B:309:LEU:HB3	2:B:317:VAL:HG11	1.41	0.99
3:M:224:VAL:HG23	3:M:479:PHE:CD1	1.98	0.99
4:S:8:PHE:HB3	4:S:36:TYR:CZ	1.98	0.99
4:S:54:PRO:CG	4:S:57:LEU:HD11	1.89	0.99
1:A:128:LEU:HD13	1:A:150:LEU:HG	1.39	0.98
2:B:60:ARG:HD2	2:B:96:LYS:HG2	1.41	0.98
2:B:274:PRO:CG	2:B:295:ASN:CG	2.32	0.98
2:B:459:GLU:HA	2:B:496:LEU:HD22	1.44	0.98
3:M:66:PHE:HB3	3:M:77:LEU:CD1	1.90	0.98
1:A:183:THR:OG1	4:S:142:ILE:CD1	2.11	0.98
2:B:50:LEU:HB3	2:B:62:ALA:HB2	1.43	0.98
2:B:178:ILE:HG22	2:B:179:LYS:N	1.76	0.98
2:B:350:THR:HB	2:B:352:ASN:ND2	1.78	0.98
3:M:293:PRO:HD2	3:M:293:PRO:O	1.61	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:388:VAL:CG1	1:A:432:ILE:HD12	1.92	0.98
2:B:127:LEU:CA	2:B:161:LEU:HD11	1.93	0.98
2:B:307:ASN:OD1	2:B:339:PHE:CD2	2.15	0.98
2:B:490:ILE:HG22	2:B:515:PHE:CE2	1.97	0.98
1:A:84:MET:O	1:A:85:ALA:C	1.93	0.98
2:B:78:ASP:OD2	2:B:81:LEU:HG	1.62	0.98
2:B:86:VAL:HG13	2:B:101:ILE:HG12	1.45	0.98
2:B:230:PHE:CZ	2:B:252:LEU:HD23	1.80	0.98
2:B:261:PRO:HA	2:B:290:SER:CB	1.93	0.98
2:B:530:LEU:HD11	2:B:595:VAL:HG21	1.45	0.98
2:B:549:LEU:HD11	2:B:611:ALA:HA	0.99	0.98
3:M:67:SER:OG	3:M:90:PHE:CG	2.16	0.98
1:A:132:LEU:CD2	1:A:169:MET:HG3	1.93	0.98
1:A:215:VAL:CG2	1:A:243:ILE:CD1	2.36	0.98
1:A:408:ILE:CG2	4:S:65:ASN:N	2.14	0.98
2:B:105:LEU:HG	2:B:119:SER:HB2	1.44	0.98
2:B:296:ASP:OD1	2:B:297:PRO:HD2	1.63	0.98
1:A:88:ASN:HB3	1:A:120:ILE:HD12	0.99	0.98
1:A:589:SER:O	1:A:597:GLN:HG3	1.61	0.98
2:B:268:LYS:O	2:B:273:SER:CB	2.12	0.98
2:B:527:PRO:HB2	2:B:587:ARG:CG	1.93	0.98
4:S:130:SER:OG	4:S:156:LEU:HD12	1.59	0.98
1:A:320:HIS:HB2	1:A:352:PHE:CE2	1.99	0.98
1:A:332:TYR:HD1	1:A:366:SER:OG	1.21	0.98
1:A:408:ILE:HG23	4:S:64:ASN:HB2	0.99	0.98
2:B:2:VAL:HG13	2:B:6:HIS:NE2	1.79	0.98
2:B:243:TRP:CZ3	3:M:91:THR:O	2.16	0.98
2:B:275:ARG:CG	2:B:291:TYR:CD2	2.47	0.98
2:B:508:ARG:CB	2:B:544:THR:CG2	2.42	0.98
2:B:549:LEU:CG	2:B:611:ALA:HB2	1.93	0.98
3:M:66:PHE:HA	3:M:77:LEU:CD1	1.94	0.98
1:A:101:GLN:C	4:S:167:ILE:CG1	2.30	0.98
1:A:516:ILE:CG2	1:A:554:ALA:CB	2.34	0.98
2:B:25:VAL:HG11	2:B:36:THR:HG1	1.26	0.98
2:B:140:SER:CA	2:B:172:GLU:OE1	2.11	0.98
2:B:275:ARG:HG3	2:B:294:VAL:HG11	1.02	0.98
2:B:353:GLN:CG	3:M:49:ASP:N	2.19	0.98
1:A:142:LYS:HA	4:S:159:ALA:HB1	1.42	0.98
1:A:288:THR:CG2	1:A:322:PHE:CE2	2.39	0.98
2:B:223:LEU:CD1	2:B:259:TYR:CB	2.31	0.98
2:B:517:GLU:OE2	2:B:554:LYS:NZ	1.96	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:211:ASP:OD1	4:S:148:ARG:NE	1.96	0.98
1:A:537:THR:HB	1:A:584:PHE:CE1	1.99	0.98
2:B:37:TYR:CZ	2:B:46:GLN:NE2	1.78	0.98
2:B:212:VAL:CG2	2:B:248:LEU:CD2	2.42	0.98
2:B:275:ARG:CB	2:B:291:TYR:HD2	1.76	0.98
2:B:582:ASP:O	2:B:584:SER:N	1.96	0.98
1:A:105:VAL:CG2	4:S:167:ILE:HD13	1.75	0.97
1:A:219:VAL:C	1:A:259:LEU:HD12	1.81	0.97
1:A:275:LEU:HD13	1:A:308:ASP:CG	1.83	0.97
2:B:566:ALA:HA	2:B:574:ASN:CG	1.83	0.97
3:M:281:GLY:O	3:M:282:VAL:HG23	0.80	0.97
4:S:109:LEU:HD11	4:S:113:PHE:CE1	1.92	0.97
1:A:96:SER:N	4:S:166:LYS:NZ	2.10	0.97
1:A:96:SER:HA	1:A:127:LEU:CD1	1.94	0.97
1:A:217:ALA:CA	4:S:142:ILE:CB	1.92	0.97
2:B:86:VAL:HG12	2:B:101:ILE:CG2	1.93	0.97
2:B:90:ILE:HD13	2:B:123:LEU:HD23	1.46	0.97
2:B:279:LEU:CG	2:B:288:TYR:HD1	1.76	0.97
2:B:377:TYR:O	2:B:380:LYS:HB2	1.62	0.97
2:B:418:TYR:CD1	2:B:418:TYR:C	2.33	0.97
2:B:553:ALA:CB	2:B:614:ILE:HG12	1.93	0.97
3:M:241:HIS:HB2	3:M:476:THR:CG2	1.93	0.97
1:A:107:TYR:CE2	1:A:128:LEU:HD23	1.96	0.97
3:M:217:ASP:HB3	3:M:471:LYS:N	1.78	0.97
2:B:37:TYR:CD2	2:B:38:TYR:CE1	2.52	0.97
2:B:191:GLU:O	2:B:193:LEU:N	1.97	0.97
2:B:196:LEU:O	2:B:215:TYR:OH	1.82	0.97
2:B:212:VAL:HG21	2:B:248:LEU:HD21	1.46	0.97
3:M:65:TYR:CD1	3:M:86:PRO:HG3	2.00	0.97
4:S:5:VAL:HG11	4:S:132:LEU:CD2	1.94	0.97
4:S:8:PHE:HE1	4:S:84:TYR:CB	1.70	0.97
1:A:323:CYS:SG	1:A:338:PHE:HE1	1.86	0.97
1:A:627:GLU:CB	2:B:617:LEU:HB3	1.91	0.97
1:A:633:PHE:CD1	2:B:550:VAL:HG11	1.94	0.97
2:B:70:MET:CE	2:B:107:ARG:CG	2.41	0.97
2:B:208:ILE:CG1	2:B:236:ILE:HG21	1.92	0.97
2:B:223:LEU:HD13	2:B:259:TYR:N	1.68	0.97
3:M:347:PHE:HE1	3:M:439:TYR:CD2	1.83	0.97
4:S:53:THR:HG21	4:S:67:GLU:C	1.83	0.97
2:B:102:HIS:HD1	2:B:137:PHE:HB3	1.26	0.97
2:B:219:TYR:CB	2:B:255:TYR:CD1	2.46	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:343:LEU:CD1	2:B:359:LEU:CD1	2.38	0.97
2:B:346:THR:CG2	2:B:350:THR:HG23	1.94	0.97
3:M:273:HIS:HB2	3:M:298:ARG:O	1.62	0.97
3:M:360:LEU:CD2	3:M:362:PHE:CE2	2.47	0.97
1:A:92:LEU:CD2	1:A:120:ILE:O	2.12	0.97
1:A:212:ILE:HB	1:A:247:ILE:CD1	1.95	0.97
1:A:399:ASP:O	1:A:420:ILE:C	2.02	0.97
2:B:116:THR:HG22	2:B:150:LEU:CD1	1.94	0.97
2:B:127:LEU:CB	2:B:161:LEU:HD11	1.94	0.97
2:B:181:TYR:CE2	2:B:222:HIS:N	2.32	0.97
2:B:344:VAL:HG22	2:B:363:ILE:HD11	1.46	0.97
3:M:319:SER:CB	3:M:346:ASN:N	2.25	0.97
1:A:630:PRO:O	2:B:554:LYS:HA	1.62	0.97
2:B:38:TYR:CD2	2:B:65:ARG:HD3	1.96	0.97
3:M:95:THR:O	3:M:99:ILE:CD1	2.13	0.97
3:M:246:VAL:HA	3:M:470:ALA:CB	1.93	0.97
2:B:87:VAL:HG12	2:B:122:SER:HB3	0.97	0.97
2:B:219:TYR:O	2:B:223:LEU:HG	1.65	0.97
3:M:8:THR:N	3:M:75:TRP:O	1.96	0.97
1:A:88:ASN:HB3	1:A:120:ILE:CD1	1.94	0.97
1:A:557:LYS:NZ	2:B:604:GLU:HG3	1.78	0.97
1:A:584:PHE:O	1:A:587:ASN:N	1.97	0.97
1:A:636:TYR:HB2	2:B:554:LYS:NZ	1.78	0.97
3:M:261:ASN:N	3:M:448:TYR:O	1.97	0.97
1:A:166:LEU:HD13	1:A:185:LEU:HD23	1.45	0.96
1:A:185:LEU:HD13	1:A:203:PHE:HE1	0.82	0.96
1:A:255:ARG:HH22	4:S:135:ILE:HG22	1.29	0.96
2:B:20:ARG:CD	2:B:21:GLU:CB	2.42	0.96
2:B:79:VAL:CB	2:B:108:PHE:HE1	1.74	0.96
2:B:219:TYR:HD1	2:B:226:LEU:HD22	1.21	0.96
2:B:437:SER:HA	2:B:478:LEU:CD2	1.94	0.96
2:B:549:LEU:CD2	2:B:611:ALA:H	1.51	0.96
1:A:77:LEU:O	1:A:80:TYR:O	1.81	0.96
2:B:38:TYR:HA	2:B:42:ILE:N	1.81	0.96
2:B:291:TYR:HD2	2:B:294:VAL:HG11	0.82	0.96
2:B:340:ILE:HG13	2:B:373:LEU:HD23	1.45	0.96
2:B:394:TRP:HZ3	2:B:397:GLN:OE1	1.47	0.96
3:M:224:VAL:N	3:M:479:PHE:HA	1.80	0.96
3:M:302:TYR:CE2	3:M:445:SER:HB3	2.00	0.96
1:A:151:SER:HB2	1:A:187:LYS:CB	1.94	0.96
1:A:260:PHE:CZ	1:A:274:LEU:HD11	2.00	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:630:PRO:CG	2:B:614:ILE:CB	2.43	0.96
2:B:274:PRO:HG2	2:B:295:ASN:ND2	1.80	0.96
2:B:343:LEU:HD11	2:B:359:LEU:CA	1.94	0.96
2:B:559:ASP:HB2	2:B:563:PHE:HD1	1.20	0.96
3:M:244:VAL:CG1	3:M:472:TYR:CE2	2.47	0.96
4:S:83:LEU:HD11	4:S:116:VAL:HG21	1.44	0.96
1:A:176:TYR:HB3	4:S:155:GLU:CG	1.95	0.96
2:B:278:PRO:CD	2:B:289:PRO:O	2.13	0.96
3:M:4:SER:O	3:M:78:ALA:CB	1.90	0.96
3:M:8:THR:C	3:M:75:TRP:HB2	1.86	0.96
1:A:244:LEU:HD11	1:A:281:LEU:CD1	1.94	0.96
1:A:637:GLU:HB3	2:B:516:GLY:CA	1.95	0.96
2:B:44:PRO:O	2:B:47:LEU:HB2	1.65	0.96
3:M:101:LEU:HA	3:M:109:LEU:HD13	1.46	0.96
3:M:319:SER:CB	3:M:346:ASN:CA	2.42	0.96
1:A:103:LYS:HG3	4:S:163:THR:HG22	1.44	0.96
1:A:540:ILE:HG13	1:A:551:LEU:HD23	1.46	0.96
2:B:197:LYS:HB2	2:B:229:HIS:NE2	1.80	0.96
2:B:216:LYS:HA	2:B:251:LEU:HD11	0.97	0.96
1:A:93:GLU:O	4:S:166:LYS:NZ	1.89	0.96
2:B:139:LEU:HD22	2:B:173:VAL:HA	1.48	0.96
2:B:159:LYS:HE3	2:B:191:GLU:OE1	1.64	0.96
3:M:222:PHE:O	3:M:479:PHE:HE2	1.23	0.96
3:M:226:PHE:CZ	3:M:321:GLY:O	2.18	0.96
2:B:212:VAL:HG22	2:B:248:LEU:CD2	1.95	0.96
2:B:343:LEU:CD2	2:B:362:ALA:CB	2.38	0.96
2:B:343:LEU:CD1	2:B:359:LEU:HA	1.94	0.96
1:A:65:ASN:C	4:S:165:SER:CB	2.34	0.96
1:A:506:LYS:HE2	3:M:58:ARG:HA	1.46	0.96
2:B:151:ALA:CA	2:B:180:LEU:HD11	1.70	0.96
2:B:286:ILE:HG23	2:B:288:TYR:CE2	2.00	0.96
2:B:389:ILE:CG2	2:B:427:ASN:HB2	1.96	0.96
3:M:283:PHE:CE2	3:M:289:THR:HB	1.98	0.96
2:B:366:LEU:O	2:B:367:SER:C	1.85	0.95
3:M:101:LEU:HD11	3:M:106:LYS:C	1.85	0.95
4:S:109:LEU:CD1	4:S:113:PHE:HE1	1.78	0.95
2:B:48:VAL:HG23	2:B:82:TYR:HE2	1.30	0.95
2:B:117:LEU:HD21	2:B:149:SER:HG	1.18	0.95
1:A:101:GLN:HG3	4:S:160:ALA:HB3	1.46	0.95
1:A:316:LEU:HD13	1:A:348:PHE:CG	2.01	0.95
2:B:80:GLN:CG	2:B:115:LEU:HD11	1.94	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:363:ILE:HG21	2:B:398:ILE:CD1	1.96	0.95
3:M:6:TYR:CD2	3:M:17:GLN:HA	2.00	0.95
3:M:339:GLU:CG	3:M:412:ARG:CD	2.44	0.95
3:M:433:VAL:HG12	3:M:481:VAL:HB	1.47	0.95
4:S:135:ILE:O	4:S:141:VAL:CA	2.13	0.95
1:A:186:PHE:CE1	1:A:224:GLU:CG	2.50	0.95
2:B:230:PHE:HE1	2:B:252:LEU:HD23	1.15	0.95
3:M:243:ILE:O	3:M:472:TYR:CB	2.14	0.95
1:A:64:LEU:CG	1:A:102:GLN:HE22	1.79	0.95
1:A:253:ILE:HG12	1:A:281:LEU:HD13	0.96	0.95
2:B:79:VAL:HB	2:B:108:PHE:HE1	1.17	0.95
2:B:559:ASP:CB	2:B:563:PHE:CE1	2.45	0.95
3:M:101:LEU:CD1	3:M:106:LYS:C	2.34	0.95
4:S:25:LEU:HB2	4:S:26:PRO:HD3	1.49	0.95
2:B:37:TYR:CE2	2:B:38:TYR:HE1	1.84	0.95
2:B:211:ALA:O	2:B:214:ALA:HB3	1.65	0.95
3:M:71:LYS:O	3:M:72:LEU:HB2	1.65	0.95
3:M:244:VAL:HA	3:M:472:TYR:CG	2.02	0.95
4:S:89:VAL:HG11	4:S:98:ILE:HG21	1.45	0.95
1:A:64:LEU:CB	1:A:102:GLN:HE22	1.78	0.95
1:A:140:VAL:CA	1:A:177:ILE:HG13	1.86	0.95
1:A:258:LYS:NZ	4:S:97:ALA:HB2	1.80	0.95
2:B:127:LEU:HD13	2:B:157:THR:CB	1.97	0.95
2:B:185:LYS:HG3	2:B:222:HIS:NE2	1.82	0.95
3:M:243:ILE:O	3:M:472:TYR:HD2	1.49	0.95
3:M:323:MET:HE3	3:M:342:LEU:HD23	1.49	0.95
1:A:186:PHE:CE1	1:A:224:GLU:HB2	2.02	0.95
1:A:403:LEU:CD2	1:A:422:GLU:HG3	1.97	0.95
2:B:196:LEU:HB3	2:B:215:TYR:CE2	2.01	0.95
2:B:268:LYS:O	2:B:273:SER:HB3	1.65	0.95
3:M:51:LEU:HD13	3:M:75:TRP:CZ3	2.01	0.95
4:S:89:VAL:HG11	4:S:98:ILE:CG2	1.95	0.95
1:A:163:ALA:HB2	1:A:195:ALA:HB1	0.96	0.95
1:A:251:TRP:HZ2	4:S:103:GLN:HB3	1.26	0.95
2:B:42:ILE:CD1	2:B:65:ARG:HD3	1.95	0.95
2:B:174:ALA:CB	2:B:211:ALA:CA	2.44	0.95
2:B:174:ALA:CB	2:B:211:ALA:CB	2.44	0.95
2:B:178:ILE:HG13	2:B:214:ALA:CA	1.97	0.95
2:B:437:SER:O	2:B:478:LEU:CD2	2.14	0.95
3:M:18:TYR:CD1	3:M:122:SER:OG	2.20	0.95
3:M:218:LEU:O	3:M:441:GLY:N	2.00	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:222:PHE:CB	3:M:479:PHE:CZ	2.49	0.95
1:A:145:ILE:CD1	4:S:156:LEU:HD22	1.97	0.95
1:A:186:PHE:CE1	1:A:224:GLU:HG2	2.02	0.95
1:A:610:SER:OG	1:A:625:LEU:HD13	1.67	0.95
1:A:147:LEU:HD13	1:A:166:LEU:HD22	1.47	0.94
1:A:217:ALA:HA	4:S:142:ILE:CG1	1.95	0.94
2:B:117:LEU:CD2	2:B:149:SER:OG	2.14	0.94
2:B:158:VAL:HG11	2:B:177:ILE:HG12	0.96	0.94
2:B:178:ILE:CG1	2:B:214:ALA:HB1	1.97	0.94
3:M:7:ILE:HG12	3:M:76:CYS:SG	2.07	0.94
3:M:347:PHE:CE1	3:M:439:TYR:HD2	1.84	0.94
1:A:64:LEU:CB	1:A:102:GLN:NE2	2.30	0.94
1:A:101:GLN:HG3	4:S:160:ALA:HB1	0.95	0.94
1:A:581:LEU:HD23	1:A:607:LEU:CD2	1.96	0.94
2:B:12:LEU:O	2:B:16:LYS:CB	2.13	0.94
2:B:175:LEU:CD2	2:B:210:CYS:O	2.14	0.94
2:B:267:ASP:H	2:B:289:PRO:HB2	1.32	0.94
3:M:67:SER:CB	3:M:90:PHE:HD1	1.78	0.94
1:A:102:GLN:CD	4:S:166:LYS:H	1.70	0.94
1:A:366:SER:O	1:A:370:LYS:HG2	1.66	0.94
1:A:520:GLY:CA	1:A:558:VAL:CG2	2.42	0.94
2:B:123:LEU:CD1	2:B:142:LEU:CD2	2.39	0.94
2:B:394:TRP:CZ3	2:B:397:GLN:OE1	2.20	0.94
1:A:264:SER:HB2	1:A:271:ARG:HD3	1.49	0.94
1:A:408:ILE:HD12	1:A:410:TYR:CE1	2.03	0.94
2:B:143:SER:O	2:B:145:MET:N	1.99	0.94
2:B:226:LEU:CB	2:B:255:TYR:OH	2.13	0.94
2:B:566:ALA:HA	2:B:574:ASN:HB3	0.96	0.94
3:M:60:LEU:HD22	3:M:62:VAL:HG23	0.95	0.94
3:M:222:PHE:HB2	3:M:479:PHE:HZ	1.28	0.94
3:M:343:ASN:CA	3:M:408:VAL:HG13	1.97	0.94
4:S:131:VAL:HG21	4:S:153:VAL:HG22	1.48	0.94
1:A:244:LEU:HD13	1:A:256:LEU:HD13	1.45	0.94
1:A:485:PRO:O	1:A:488:ARG:HG3	1.67	0.94
1:A:555:LEU:CD1	1:A:581:LEU:HD11	1.98	0.94
2:B:219:TYR:O	2:B:223:LEU:CG	2.15	0.94
4:S:17:VAL:CG2	4:S:19:PHE:CE2	2.51	0.94
4:S:83:LEU:HD11	4:S:116:VAL:HG11	1.47	0.94
1:A:107:TYR:CZ	1:A:128:LEU:HD23	2.02	0.94
2:B:24:ALA:CB	2:B:35:TYR:CE1	2.50	0.94
2:B:62:ALA:C	2:B:66:ILE:CD1	2.35	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:171:GLY:HA2	2:B:207:VAL:CG1	1.94	0.94
2:B:451:MET:SD	2:B:489:ILE:HG12	2.08	0.94
2:B:476:ARG:HA	2:B:514:LEU:CD1	1.96	0.94
4:S:53:THR:OG1	4:S:68:VAL:C	2.06	0.94
1:A:399:ASP:O	1:A:420:ILE:CA	2.14	0.94
2:B:48:VAL:HG23	2:B:82:TYR:CE2	2.02	0.94
2:B:123:LEU:HD12	2:B:142:LEU:HD21	1.48	0.94
2:B:123:LEU:O	2:B:127:LEU:CG	2.15	0.94
2:B:267:ASP:HB3	2:B:289:PRO:CD	1.98	0.94
2:B:523:PHE:CZ	2:B:580:TYR:CE2	2.44	0.94
3:M:215:TYR:CB	3:M:467:TYR:HD2	1.80	0.94
3:M:443:SER:HB3	3:M:447:ILE:HG12	1.47	0.94
1:A:101:GLN:NE2	4:S:167:ILE:HG22	1.83	0.94
1:A:150:LEU:HB3	1:A:162:ILE:CD1	1.97	0.94
1:A:212:ILE:CG1	4:S:145:ASN:ND2	2.31	0.94
1:A:316:LEU:HD13	1:A:348:PHE:CD2	1.99	0.94
1:A:557:LYS:CE	2:B:606:ASP:N	1.85	0.94
2:B:117:LEU:HD23	2:B:150:LEU:HD23	1.45	0.94
2:B:135:ARG:HB3	2:B:161:LEU:HG	1.48	0.94
2:B:162:VAL:HG21	2:B:195:ILE:C	1.88	0.94
2:B:239:GLN:OE1	3:M:278:ILE:HG23	1.66	0.94
2:B:303:LEU:HD11	2:B:333:GLN:CD	1.88	0.94
2:B:374:PHE:HE1	2:B:381:PHE:CE1	1.86	0.94
3:M:218:LEU:CA	3:M:472:TYR:CE2	2.50	0.94
3:M:432:THR:HG1	3:M:480:GLN:HG3	1.14	0.94
4:S:8:PHE:CE1	4:S:84:TYR:CG	2.56	0.94
1:A:509:PRO:HB3	1:A:547:VAL:HG23	1.50	0.94
2:B:215:TYR:HD2	2:B:219:TYR:HE1	1.04	0.94
2:B:363:ILE:HG22	2:B:398:ILE:HG12	1.48	0.94
3:M:302:TYR:CE2	3:M:445:SER:CB	2.51	0.94
1:A:557:LYS:CE	2:B:606:ASP:CB	2.45	0.94
2:B:103:LEU:HD13	3:M:132:GLY:HA3	1.50	0.94
2:B:556:LEU:HD23	2:B:588:ILE:HG13	1.49	0.94
3:M:272:LEU:HD22	3:M:278:ILE:CB	1.98	0.94
3:M:364:VAL:O	3:M:367:ALA:O	1.85	0.94
1:A:332:TYR:CE1	1:A:366:SER:OG	2.12	0.93
2:B:162:VAL:CG2	2:B:199:LEU:HD11	1.97	0.93
2:B:352:ASN:CB	3:M:49:ASP:CG	2.35	0.93
2:B:353:GLN:CD	3:M:47:SER:HB3	1.83	0.93
2:B:490:ILE:HG13	2:B:518:ILE:HG21	1.51	0.93
2:B:546:CYS:CA	2:B:607:ILE:CG2	2.39	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:582:ASP:O	2:B:584:SER:CB	2.16	0.93
3:M:319:SER:HG	3:M:346:ASN:HB2	1.19	0.93
1:A:244:LEU:HD23	1:A:277:LYS:O	1.66	0.93
3:M:222:PHE:O	3:M:479:PHE:CZ	2.20	0.93
3:M:443:SER:HG	3:M:447:ILE:C	1.67	0.93
2:B:117:LEU:HD21	2:B:149:SER:CB	1.99	0.93
2:B:223:LEU:HD22	2:B:255:TYR:CD1	2.01	0.93
2:B:256:CYS:HA	2:B:293:VAL:HG21	1.47	0.93
2:B:396:ILE:HD11	2:B:418:TYR:CZ	2.04	0.93
2:B:553:ALA:HB1	2:B:614:ILE:HG12	1.48	0.93
2:B:563:PHE:CD2	2:B:584:SER:HB2	1.95	0.93
3:M:100:LEU:HD22	3:M:101:LEU:N	1.81	0.93
1:A:178:ARG:NH1	1:A:209:ASP:HB3	1.83	0.93
1:A:182:ILE:C	1:A:221:VAL:HG21	1.89	0.93
1:A:207:LEU:HD23	1:A:239:LEU:CB	1.97	0.93
2:B:29:LYS:HE2	2:B:30:LEU:H	0.87	0.93
2:B:106:LEU:HD13	2:B:144:ASP:HB2	0.95	0.93
2:B:216:LYS:HG3	2:B:251:LEU:CD1	1.99	0.93
3:M:215:TYR:HB2	3:M:467:TYR:CD2	2.02	0.93
3:M:219:LEU:HB3	3:M:472:TYR:C	1.87	0.93
3:M:290:PHE:CZ	3:M:297:PHE:CD2	2.57	0.93
1:A:71:VAL:HG12	1:A:105:VAL:HG12	1.47	0.93
1:A:282:MET:O	1:A:283:GLU:C	2.01	0.93
2:B:51:LEU:CD2	2:B:59:VAL:HG13	1.98	0.93
2:B:159:LYS:HD2	2:B:191:GLU:HG3	1.49	0.93
2:B:552:SER:O	2:B:556:LEU:HG	1.68	0.93
3:M:244:VAL:CA	3:M:472:TYR:CE2	2.43	0.93
1:A:180:LYS:HE3	4:S:156:LEU:HD21	1.51	0.93
2:B:158:VAL:HG13	2:B:173:VAL:CG1	1.98	0.93
2:B:267:ASP:C	2:B:276:SER:CB	2.36	0.93
3:M:2:TYR:O	3:M:81:SER:CB	2.16	0.93
3:M:224:VAL:N	3:M:479:PHE:CA	2.31	0.93
3:M:317:MET:HB3	3:M:320:ILE:C	1.87	0.93
3:M:327:PHE:HE1	3:M:336:ASP:CB	1.81	0.93
4:S:63:ASN:O	4:S:66:ASP:OD1	1.85	0.93
1:A:213:SER:OG	4:S:143:GLU:OE1	1.86	0.93
2:B:24:ALA:HB3	2:B:35:TYR:CD2	2.04	0.93
2:B:123:LEU:HD22	2:B:138:ALA:O	1.67	0.93
2:B:311:TYR:HE2	2:B:342:ALA:HB2	0.80	0.93
2:B:549:LEU:CG	2:B:611:ALA:HA	1.98	0.93
3:M:104:PHE:CZ	3:M:113:LYS:NZ	0.73	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:102:GLN:HB3	4:S:166:LYS:HB2	0.96	0.93
1:A:103:LYS:CG	4:S:163:THR:HG21	1.98	0.93
1:A:401:VAL:HB	1:A:419:ILE:HD12	1.49	0.93
2:B:24:ALA:HB1	2:B:35:TYR:CE1	2.03	0.93
2:B:275:ARG:HG3	2:B:294:VAL:CG1	1.85	0.93
3:M:69:ILE:HD11	3:M:90:PHE:HZ	0.77	0.93
4:S:4:ALA:HA	4:S:18:LYS:O	1.69	0.93
1:A:606:PHE:HZ	2:B:550:VAL:HG11	1.34	0.93
1:A:629:LEU:HD11	2:B:610:ARG:HH11	1.28	0.93
2:B:132:SER:CA	2:B:169:VAL:CG2	2.47	0.93
2:B:252:LEU:HD12	2:B:302:PHE:HE1	1.34	0.93
2:B:549:LEU:HG	2:B:614:ILE:HD12	1.50	0.93
1:A:132:LEU:CD1	1:A:165:ASP:HB3	1.99	0.93
1:A:316:LEU:CD1	1:A:348:PHE:CE2	2.52	0.93
1:A:332:TYR:CZ	1:A:336:ILE:HD11	2.03	0.93
1:A:523:SER:CB	1:A:562:TRP:HE1	1.82	0.93
2:B:67:ILE:CD1	2:B:103:LEU:HB2	1.99	0.93
2:B:219:TYR:O	2:B:223:LEU:HD21	1.66	0.93
2:B:553:ALA:HA	2:B:614:ILE:CG2	1.99	0.93
2:B:559:ASP:O	2:B:562:ASN:C	2.08	0.93
1:A:189:PHE:HD2	1:A:225:LEU:CD2	1.81	0.92
1:A:605:GLU:OE2	1:A:632:PHE:CZ	2.22	0.92
1:A:633:PHE:HD1	2:B:513:TRP:CZ3	1.76	0.92
2:B:106:LEU:HD11	2:B:144:ASP:O	1.56	0.92
2:B:197:LYS:CB	2:B:229:HIS:NE2	2.32	0.92
4:S:87:PHE:CD1	4:S:102:ILE:HG12	2.04	0.92
1:A:99:LYS:NZ	4:S:164:ASP:CB	2.32	0.92
1:A:563:CYS:HA	1:A:566:PHE:CD2	2.03	0.92
2:B:197:LYS:N	2:B:229:HIS:CE1	2.37	0.92
2:B:277:CYS:CA	2:B:292:GLU:HG3	1.99	0.92
2:B:396:ILE:CD1	2:B:418:TYR:CE2	2.52	0.92
2:B:497:LEU:CD2	2:B:533:LEU:HD21	1.97	0.92
3:M:4:SER:O	3:M:78:ALA:HB1	1.17	0.92
3:M:69:ILE:HG13	3:M:90:PHE:HE1	1.32	0.92
4:S:54:PRO:C	4:S:57:LEU:HD13	1.90	0.92
1:A:104:ARG:HA	1:A:145:ILE:CG2	1.97	0.92
1:A:138:ASN:OD1	4:S:158:LYS:CD	2.13	0.92
1:A:141:VAL:CG1	4:S:159:ALA:CB	2.35	0.92
1:A:264:SER:HB3	1:A:271:ARG:CG	1.99	0.92
1:A:319:LEU:O	1:A:320:HIS:C	1.98	0.92
1:A:557:LYS:CB	2:B:605:PHE:HD2	1.80	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:609:LEU:HD21	1:A:628:VAL:HG21	0.92	0.92
2:B:20:ARG:HG3	2:B:21:GLU:H	1.31	0.92
2:B:182:ARG:CD	2:B:217:GLU:HB3	2.00	0.92
2:B:286:ILE:CG2	2:B:288:TYR:CZ	2.52	0.92
2:B:566:ALA:C	2:B:574:ASN:HB2	1.78	0.92
3:M:52:ASP:HA	3:M:67:SER:C	1.90	0.92
3:M:435:LEU:O	3:M:479:PHE:CG	2.22	0.92
2:B:195:ILE:O	2:B:197:LYS:N	2.02	0.92
2:B:478:LEU:O	2:B:479:VAL:C	2.00	0.92
3:M:81:SER:O	3:M:82:LYS:HB3	1.69	0.92
4:S:15:ARG:HD2	4:S:122:ILE:CG1	2.00	0.92
1:A:216:SER:CB	4:S:143:GLU:HA	1.99	0.92
1:A:609:LEU:HD23	1:A:628:VAL:HG11	1.51	0.92
2:B:176:ALA:O	2:B:178:ILE:N	2.02	0.92
3:M:215:TYR:CE1	3:M:468:LYS:HG2	2.04	0.92
3:M:224:VAL:N	3:M:479:PHE:CB	2.31	0.92
1:A:253:ILE:CD1	1:A:281:LEU:HB3	1.99	0.92
1:A:516:ILE:HG21	1:A:554:ALA:HB3	1.51	0.92
2:B:174:ALA:HB1	2:B:211:ALA:CA	2.00	0.92
2:B:351:GLU:N	2:B:351:GLU:OE2	2.01	0.92
3:M:66:PHE:CA	3:M:77:LEU:CD1	2.41	0.92
3:M:245:ASP:O	3:M:472:TYR:HE1	1.39	0.92
4:S:48:SER:CB	4:S:77:TYR:HB3	1.99	0.92
1:A:244:LEU:HD13	1:A:256:LEU:HD12	1.51	0.92
2:B:62:ALA:O	2:B:66:ILE:HD12	1.66	0.92
2:B:132:SER:CA	2:B:169:VAL:HG21	1.99	0.92
3:M:281:GLY:C	3:M:282:VAL:HG23	1.90	0.92
4:S:53:THR:HG23	4:S:67:GLU:O	1.67	0.92
2:B:102:HIS:O	2:B:103:LEU:C	1.98	0.92
2:B:144:ASP:CA	2:B:179:LYS:NZ	2.27	0.92
2:B:174:ALA:O	2:B:175:LEU:C	1.99	0.92
2:B:219:TYR:CE2	2:B:226:LEU:CB	2.44	0.92
2:B:278:PRO:HA	2:B:288:TYR:O	1.63	0.92
3:M:4:SER:OG	3:M:6:TYR:CE1	2.23	0.92
3:M:256:VAL:CG1	3:M:290:PHE:HB3	1.99	0.92
3:M:347:PHE:CE1	3:M:439:TYR:CD2	2.58	0.92
4:S:5:VAL:O	4:S:17:VAL:HA	1.68	0.92
4:S:7:ILE:HG21	4:S:121:LEU:CD2	2.00	0.92
1:A:103:LYS:HB3	1:A:107:TYR:CE1	2.04	0.92
1:A:105:VAL:CG2	4:S:167:ILE:CG2	2.46	0.92
1:A:170:LEU:HD12	1:A:206:LYS:HG3	1.48	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:292:GLU:HG2	2:B:296:ASP:HB2	1.52	0.92
2:B:578:PRO:CB	2:B:579:PRO:CD	2.48	0.92
3:M:245:ASP:CB	3:M:472:TYR:CD1	2.52	0.92
1:A:64:LEU:HG	1:A:102:GLN:HE22	1.35	0.92
1:A:144:GLY:HA2	1:A:180:LYS:CB	2.00	0.92
2:B:158:VAL:C	2:B:195:ILE:HD13	1.90	0.92
2:B:483:PRO:HB3	2:B:521:ILE:HG21	0.95	0.92
2:B:578:PRO:HB2	2:B:579:PRO:HD2	1.49	0.92
3:M:254:PRO:HB3	3:M:454:ILE:HD11	1.51	0.92
4:S:8:PHE:HB3	4:S:36:TYR:OH	1.69	0.92
4:S:130:SER:CB	4:S:156:LEU:HD13	1.99	0.92
1:A:189:PHE:HD2	1:A:225:LEU:HD21	1.29	0.91
2:B:151:ALA:HA	2:B:180:LEU:CG	2.00	0.91
2:B:181:TYR:HE2	2:B:222:HIS:N	1.64	0.91
2:B:193:LEU:C	2:B:195:ILE:N	2.20	0.91
2:B:215:TYR:HD2	2:B:219:TYR:CE1	1.87	0.91
2:B:230:PHE:HZ	2:B:252:LEU:HD22	0.87	0.91
4:S:21:THR:HB	4:S:22:PRO:HD2	1.50	0.91
2:B:412:PHE:HE2	2:B:446:TRP:HB3	1.32	0.91
3:M:433:VAL:CG1	3:M:481:VAL:HB	2.00	0.91
1:A:186:PHE:CE1	1:A:224:GLU:CB	2.53	0.91
1:A:291:ILE:HG23	1:A:318:ARG:HB3	0.93	0.91
2:B:93:ASN:HA	2:B:134:LEU:HD11	1.51	0.91
2:B:162:VAL:HG23	2:B:195:ILE:HG23	1.09	0.91
2:B:170:ARG:HA	2:B:199:LEU:HD22	0.92	0.91
2:B:340:ILE:CG1	2:B:373:LEU:HD23	2.01	0.91
4:S:109:LEU:HD13	4:S:113:PHE:CE1	2.03	0.91
1:A:147:LEU:CD2	1:A:166:LEU:HD21	1.92	0.91
1:A:633:PHE:CD2	2:B:550:VAL:HG12	2.02	0.91
2:B:116:THR:CG2	2:B:150:LEU:CD1	2.48	0.91
2:B:154:ILE:CD1	2:B:180:LEU:HD13	2.00	0.91
2:B:252:LEU:HB3	2:B:302:PHE:CD2	2.05	0.91
2:B:274:PRO:CG	2:B:295:ASN:ND2	2.33	0.91
2:B:319:LEU:HD11	2:B:358:MET:SD	2.10	0.91
3:M:256:VAL:O	3:M:289:THR:HA	1.69	0.91
1:A:179:LYS:HE3	4:S:143:GLU:HB2	0.94	0.91
1:A:629:LEU:CD2	2:B:610:ARG:HH11	1.82	0.91
1:A:637:GLU:HG3	2:B:516:GLY:N	1.84	0.91
2:B:286:ILE:HG23	2:B:288:TYR:CZ	2.06	0.91
3:M:19:LEU:HD13	3:M:24:ALA:HB3	1.51	0.91
4:S:16:LEU:HD23	4:S:128:LEU:HD23	1.50	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:103:LYS:HD3	1:A:131:ASP:CG	1.91	0.91
2:B:87:VAL:HG11	2:B:122:SER:HB3	1.48	0.91
2:B:237:ILE:HG21	2:B:305:SER:HB3	1.50	0.91
2:B:319:LEU:HD13	2:B:354:GLY:O	1.69	0.91
2:B:518:ILE:O	2:B:518:ILE:HD12	1.70	0.91
3:M:2:TYR:N	3:M:81:SER:OG	2.04	0.91
1:A:99:LYS:HZ3	4:S:164:ASP:HB2	1.28	0.91
1:A:143:VAL:O	1:A:147:LEU:HG	1.71	0.91
1:A:147:LEU:HD13	1:A:181:ALA:HA	1.49	0.91
2:B:307:ASN:HD21	2:B:336:ASN:HD22	1.13	0.91
2:B:549:LEU:CG	2:B:611:ALA:CA	2.49	0.91
3:M:101:LEU:HG	3:M:106:LYS:C	1.89	0.91
3:M:362:PHE:O	3:M:363:ASN:C	2.07	0.91
1:A:102:GLN:CB	4:S:163:THR:CB	2.27	0.91
2:B:143:SER:CB	2:B:179:LYS:HD2	2.01	0.91
2:B:219:TYR:CE2	2:B:226:LEU:N	2.28	0.91
3:M:45:SER:CA	3:M:47:SER:H	1.83	0.91
1:A:220:SER:HB3	4:S:142:ILE:HG23	1.51	0.91
1:A:250:ASN:OD1	1:A:285:THR:HG21	1.69	0.91
1:A:403:LEU:CD2	1:A:422:GLU:CD	2.38	0.91
1:A:573:GLU:O	1:A:574:ILE:C	1.97	0.91
1:A:631:SER:CA	2:B:557:SER:HB2	1.98	0.91
2:B:351:GLU:O	3:M:48:ASP:HB2	1.70	0.91
2:B:486:HIS:CD2	2:B:518:ILE:HG12	2.06	0.91
2:B:596:LEU:HD22	2:B:615:SER:CB	2.00	0.91
3:M:245:ASP:C	3:M:472:TYR:CE1	2.44	0.91
3:M:262:THR:HG22	3:M:264:GLY:N	1.86	0.91
4:S:53:THR:O	4:S:69:ASN:ND2	2.04	0.91
4:S:53:THR:N	4:S:69:ASN:HB2	1.85	0.91
1:A:100:LEU:O	4:S:160:ALA:HA	1.30	0.91
1:A:101:GLN:CG	4:S:160:ALA:CB	2.17	0.91
1:A:102:GLN:HA	4:S:167:ILE:HG12	0.92	0.91
1:A:148:SER:O	1:A:151:SER:OG	1.87	0.91
2:B:70:MET:HE3	2:B:107:ARG:CG	2.01	0.91
3:M:344:ILE:HB	3:M:407:THR:O	1.70	0.91
4:S:16:LEU:HD12	4:S:17:VAL:H	1.35	0.91
1:A:65:ASN:C	4:S:165:SER:HB2	1.91	0.90
1:A:80:TYR:CB	1:A:82:PHE:CE2	2.54	0.90
1:A:623:MET:O	2:B:617:LEU:HD21	1.71	0.90
2:B:588:ILE:HG23	2:B:618:PHE:HZ	1.31	0.90
3:M:45:SER:HA	3:M:47:SER:H	1.36	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:65:TYR:CD1	3:M:86:PRO:HB3	2.05	0.90
1:A:92:LEU:HD13	1:A:123:LEU:HD12	1.50	0.90
1:A:179:LYS:CD	4:S:143:GLU:HB2	1.99	0.90
1:A:381:GLU:O	1:A:382:ASP:C	1.97	0.90
2:B:346:THR:HG22	2:B:350:THR:CG2	2.00	0.90
3:M:92:PHE:HZ	3:M:129:VAL:CG2	1.84	0.90
1:A:105:VAL:HG21	4:S:167:ILE:CA	2.01	0.90
1:A:609:LEU:HG	1:A:628:VAL:HB	0.90	0.90
1:A:638:LEU:CD1	2:B:518:ILE:HG23	2.00	0.90
2:B:20:ARG:NH1	2:B:35:TYR:OH	2.03	0.90
2:B:29:LYS:CD	2:B:30:LEU:H	1.83	0.90
2:B:441:GLN:HG2	2:B:481:LYS:HG3	1.53	0.90
4:S:8:PHE:HE1	4:S:84:TYR:CG	1.89	0.90
1:A:634:ASN:O	2:B:517:GLU:N	2.02	0.90
2:B:353:GLN:HB2	3:M:49:ASP:N	1.84	0.90
3:M:317:MET:HB2	3:M:322:LEU:N	1.84	0.90
1:A:67:LYS:C	4:S:166:LYS:HA	1.91	0.90
1:A:107:TYR:CZ	1:A:128:LEU:CD2	2.54	0.90
1:A:326:GLN:HA	1:A:331:ARG:HH21	1.33	0.90
2:B:123:LEU:HD12	2:B:142:LEU:HD23	1.50	0.90
2:B:354:GLY:O	2:B:358:MET:HG2	1.71	0.90
2:B:513:TRP:N	2:B:551:LEU:HD11	1.85	0.90
2:B:537:PHE:CZ	2:B:545:ARG:CG	2.54	0.90
3:M:375:LYS:N	3:M:416:GLU:O	2.03	0.90
3:M:379:LEU:HD22	3:M:386:PHE:HD1	1.34	0.90
4:S:56:SER:O	4:S:60:SER:OG	1.88	0.90
1:A:102:GLN:HE21	4:S:165:SER:H	1.18	0.90
1:A:224:GLU:OE1	4:S:138:GLY:O	1.90	0.90
2:B:20:ARG:HH12	2:B:21:GLU:HG3	1.34	0.90
2:B:83:PHE:CZ	2:B:119:SER:HB2	2.05	0.90
2:B:108:PHE:CE2	2:B:115:LEU:HG	2.05	0.90
3:M:223:HIS:HB3	3:M:478:ASN:HA	1.54	0.90
4:S:135:ILE:HG22	4:S:141:VAL:HG22	1.51	0.90
1:A:403:LEU:HD21	1:A:422:GLU:N	1.86	0.90
1:A:454:ILE:HG23	1:A:473:ILE:HG23	1.51	0.90
1:A:498:LEU:O	1:A:501:ASN:O	1.88	0.90
2:B:63:MET:CE	2:B:104:TYR:HB2	2.02	0.90
2:B:70:MET:SD	2:B:107:ARG:HB2	2.11	0.90
2:B:527:PRO:HB2	2:B:587:ARG:HG3	1.52	0.90
2:B:553:ALA:CB	2:B:614:ILE:CG1	2.48	0.90
4:S:15:ARG:NH1	4:S:122:ILE:CD1	2.35	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:17:VAL:HG21	4:S:19:PHE:HZ	1.22	0.90
1:A:96:SER:HA	1:A:127:LEU:CG	2.00	0.90
2:B:135:ARG:HH22	2:B:164:ASP:CG	1.73	0.90
2:B:278:PRO:HB3	2:B:288:TYR:O	1.69	0.90
2:B:319:LEU:HD12	2:B:358:MET:HG3	0.91	0.90
2:B:418:TYR:CD1	2:B:424:PHE:CE1	2.59	0.90
2:B:525:ILE:C	2:B:527:PRO:HD2	1.92	0.90
1:A:634:ASN:O	2:B:517:GLU:HA	1.71	0.90
2:B:67:ILE:HD11	2:B:103:LEU:HB2	1.51	0.90
2:B:219:TYR:CD1	2:B:255:TYR:CE1	2.49	0.90
2:B:360:LEU:HD13	2:B:391:ALA:HA	1.51	0.90
2:B:549:LEU:HD22	2:B:611:ALA:HB2	0.91	0.90
2:B:602:ASP:O	2:B:608:ARG:NH2	2.04	0.90
3:M:104:PHE:HE1	3:M:113:LYS:CE	1.42	0.90
3:M:219:LEU:CB	3:M:472:TYR:C	2.39	0.90
1:A:186:PHE:CD1	1:A:224:GLU:CB	2.54	0.90
1:A:631:SER:HA	2:B:557:SER:HB2	1.53	0.90
1:A:637:GLU:OE1	2:B:513:TRP:HD1	1.50	0.90
2:B:155:LEU:HB2	2:B:188:TYR:CD2	2.07	0.90
2:B:311:TYR:CE2	2:B:342:ALA:HB1	2.07	0.90
2:B:319:LEU:HD13	2:B:358:MET:CG	1.72	0.90
3:M:222:PHE:CD1	3:M:240:ILE:CG2	2.55	0.90
3:M:379:LEU:CD2	3:M:386:PHE:CG	2.51	0.90
4:S:39:ILE:CD1	4:S:77:TYR:CD2	2.54	0.90
1:A:105:VAL:HG21	4:S:167:ILE:CG2	2.01	0.89
2:B:20:ARG:CZ	2:B:21:GLU:CG	2.37	0.89
2:B:563:PHE:CE2	2:B:584:SER:CA	2.55	0.89
3:M:101:LEU:HG	3:M:106:LYS:CB	2.02	0.89
3:M:265:ASN:OD1	3:M:313:SER:CB	2.18	0.89
1:A:102:GLN:CB	4:S:163:THR:CA	2.48	0.89
1:A:533:ILE:CG1	1:A:562:TRP:CH2	2.56	0.89
1:A:531:ASP:C	1:A:534:LYS:O	2.10	0.89
2:B:237:ILE:O	2:B:238:LYS:C	2.04	0.89
2:B:286:ILE:HG21	2:B:288:TYR:OH	1.73	0.89
3:M:101:LEU:CG	3:M:106:LYS:C	2.41	0.89
3:M:215:TYR:CB	3:M:467:TYR:CD2	2.54	0.89
1:A:219:VAL:CG1	1:A:259:LEU:CD1	2.51	0.89
2:B:346:THR:HG22	2:B:350:THR:HG23	1.53	0.89
3:M:219:LEU:HD22	3:M:473:LYS:CA	2.00	0.89
1:A:99:LYS:HD3	4:S:164:ASP:N	1.86	0.89
1:A:557:LYS:CE	2:B:606:ASP:HB2	2.03	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:605:GLU:HG3	1:A:632:PHE:CG	2.07	0.89
2:B:24:ALA:HB1	2:B:35:TYR:CD1	2.07	0.89
2:B:120:ILE:HD13	2:B:142:LEU:HD23	1.54	0.89
2:B:275:ARG:HB2	2:B:294:VAL:HG13	0.90	0.89
2:B:563:PHE:CD2	2:B:584:SER:CA	2.55	0.89
3:M:224:VAL:H	3:M:479:PHE:CA	1.85	0.89
3:M:246:VAL:HA	3:M:470:ALA:HB2	1.52	0.89
1:A:531:ASP:O	1:A:534:LYS:O	1.90	0.89
2:B:378:THR:HG23	2:B:379:LYS:N	1.86	0.89
2:B:562:ASN:HB3	2:B:580:TYR:CB	2.02	0.89
3:M:243:ILE:O	3:M:472:TYR:HB2	1.72	0.89
3:M:338:PHE:CD2	3:M:415:ILE:CG1	2.56	0.89
1:A:214:VAL:CG2	4:S:148:ARG:NH2	2.35	0.89
1:A:225:LEU:CD1	1:A:233:PHE:HZ	1.72	0.89
1:A:372:ILE:HG21	1:A:427:LYS:HE3	1.53	0.89
1:A:586:GLU:HB2	1:A:604:LEU:HD11	1.54	0.89
1:A:631:SER:HB3	2:B:557:SER:CB	1.93	0.89
2:B:140:SER:HB2	2:B:172:GLU:OE2	1.71	0.89
2:B:177:ILE:HD13	2:B:196:LEU:HG	1.54	0.89
2:B:216:LYS:CA	2:B:251:LEU:HD11	1.86	0.89
2:B:243:TRP:CZ2	3:M:95:THR:N	2.40	0.89
2:B:353:GLN:HG3	3:M:47:SER:CA	2.02	0.89
2:B:474:VAL:O	2:B:477:MET:N	2.06	0.89
3:M:436:GLU:HA	3:M:479:PHE:CE2	2.08	0.89
4:S:109:LEU:HD13	4:S:113:PHE:HE1	1.36	0.89
1:A:254:ILE:HG12	1:A:290:VAL:HG22	1.52	0.89
1:A:392:MET:SD	1:A:457:LEU:CD2	2.61	0.89
2:B:20:ARG:CZ	2:B:21:GLU:CB	2.51	0.89
2:B:226:LEU:CB	2:B:255:TYR:CZ	2.42	0.89
2:B:291:TYR:CE2	2:B:294:VAL:CG1	2.55	0.89
2:B:475:ILE:HG22	2:B:514:LEU:HD21	1.54	0.89
3:M:479:PHE:H	3:M:479:PHE:HD2	1.17	0.89
1:A:147:LEU:HB3	1:A:184:ALA:HB2	1.55	0.89
1:A:288:THR:HG1	1:A:291:ILE:HB	1.38	0.89
2:B:346:THR:CG2	2:B:350:THR:CG2	2.50	0.89
2:B:415:LEU:HD13	2:B:436:LEU:CD2	2.02	0.89
2:B:553:ALA:CB	2:B:614:ILE:CD1	2.51	0.89
3:M:218:LEU:HG	3:M:244:VAL:HG22	1.54	0.89
3:M:255:LEU:O	3:M:454:ILE:HA	1.71	0.89
4:S:53:THR:HB	4:S:69:ASN:H	1.33	0.89
1:A:128:LEU:HD12	1:A:150:LEU:CD2	1.77	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:147:LEU:HD13	1:A:166:LEU:CD2	2.02	0.89
1:A:150:LEU:HD13	1:A:162:ILE:HG12	1.52	0.89
1:A:605:GLU:CG	1:A:632:PHE:CG	2.55	0.89
2:B:161:LEU:HB3	2:B:173:VAL:HG22	1.55	0.89
2:B:219:TYR:C	2:B:223:LEU:HD21	1.93	0.89
2:B:219:TYR:HB2	2:B:255:TYR:CD1	2.07	0.89
4:S:54:PRO:HG2	4:S:57:LEU:HD11	1.55	0.89
1:A:95:MET:SD	1:A:107:TYR:CD2	2.65	0.88
2:B:278:PRO:CA	2:B:288:TYR:CB	2.50	0.88
3:M:219:LEU:CG	3:M:472:TYR:O	2.21	0.88
1:A:105:VAL:CB	4:S:167:ILE:CD1	2.33	0.88
1:A:516:ILE:HG22	1:A:554:ALA:HB1	1.54	0.88
1:A:630:PRO:CG	2:B:614:ILE:HA	2.02	0.88
2:B:56:SER:HB2	2:B:92:THR:HG21	1.55	0.88
2:B:79:VAL:HG23	2:B:108:PHE:CE1	2.06	0.88
2:B:208:ILE:HD13	2:B:236:ILE:HG22	1.54	0.88
2:B:275:ARG:CG	2:B:291:TYR:HD2	1.83	0.88
3:M:242:GLY:CA	3:M:444:ALA:HB2	2.03	0.88
1:A:101:GLN:O	4:S:167:ILE:HG13	1.73	0.88
1:A:213:SER:C	4:S:143:GLU:OE1	2.12	0.88
1:A:217:ALA:CB	4:S:143:GLU:HG3	2.02	0.88
1:A:316:LEU:HD12	1:A:348:PHE:CE2	2.07	0.88
2:B:98:LYS:HD2	2:B:138:ALA:HB2	1.55	0.88
2:B:219:TYR:HB3	2:B:223:LEU:HD21	1.50	0.88
2:B:274:PRO:CD	2:B:295:ASN:ND2	2.36	0.88
2:B:337:THR:HG23	2:B:373:LEU:HD11	1.55	0.88
2:B:550:VAL:HG22	2:B:610:ARG:CD	2.02	0.88
1:A:96:SER:CA	1:A:127:LEU:CD2	2.37	0.88
1:A:102:GLN:CA	4:S:167:ILE:CG1	2.49	0.88
1:A:216:SER:O	1:A:219:VAL:HB	1.71	0.88
1:A:320:HIS:HB2	1:A:352:PHE:HE2	1.33	0.88
1:A:326:GLN:HA	1:A:331:ARG:CZ	2.02	0.88
2:B:20:ARG:CZ	2:B:21:GLU:HB2	2.03	0.88
2:B:227:HIS:NE2	2:B:292:GLU:OE2	2.06	0.88
2:B:268:LYS:O	2:B:273:SER:OG	1.91	0.88
2:B:303:LEU:HD13	2:B:333:GLN:CB	1.97	0.88
3:M:283:PHE:HE2	3:M:289:THR:HB	1.37	0.88
1:A:225:LEU:CB	1:A:233:PHE:CE1	2.55	0.88
2:B:309:LEU:CB	2:B:317:VAL:HG12	1.99	0.88
3:M:10:THR:HA	3:M:75:TRP:NE1	1.88	0.88
3:M:374:TYR:HB3	3:M:417:TYR:HD2	1.38	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:64:LEU:HG	1:A:102:GLN:NE2	1.87	0.88
1:A:103:LYS:N	4:S:163:THR:HG21	1.89	0.88
1:A:141:VAL:C	4:S:159:ALA:CB	2.41	0.88
1:A:513:ARG:HD2	1:A:550:VAL:HG21	0.88	0.88
2:B:105:LEU:O	2:B:106:LEU:C	2.00	0.88
2:B:592:TYR:OH	2:B:619:ASP:CG	2.11	0.88
1:A:151:SER:HB2	1:A:187:LYS:HB2	1.55	0.88
2:B:78:ASP:OD1	2:B:80:GLN:HB2	1.73	0.88
2:B:116:THR:HG21	2:B:150:LEU:HD11	1.55	0.88
3:M:220:GLU:HG3	3:M:439:TYR:HD1	1.36	0.88
3:M:272:LEU:HD21	3:M:278:ILE:HB	1.54	0.88
4:S:53:THR:CG2	4:S:57:LEU:HB2	2.04	0.88
1:A:96:SER:CA	1:A:127:LEU:CG	2.52	0.88
1:A:154:ILE:HG22	1:A:191:GLN:HG3	1.54	0.88
1:A:264:SER:CB	1:A:271:ARG:HD3	2.03	0.88
1:A:595:GLU:OE2	2:B:472:VAL:HB	1.73	0.88
2:B:120:ILE:HG13	2:B:150:LEU:HB3	1.54	0.88
2:B:158:VAL:CG1	2:B:177:ILE:HG12	1.84	0.88
2:B:260:LEU:HD22	2:B:291:TYR:OH	0.70	0.88
3:M:214:LEU:CD1	3:M:256:VAL:HG21	2.04	0.88
3:M:220:GLU:HG3	3:M:439:TYR:CD1	2.08	0.88
3:M:334:ASP:O	3:M:417:TYR:N	2.06	0.88
1:A:67:LYS:CB	4:S:165:SER:OG	2.22	0.88
1:A:100:LEU:HG	4:S:162:SER:H	1.38	0.88
1:A:537:THR:HB	1:A:584:PHE:CD1	2.08	0.88
2:B:162:VAL:CG2	2:B:199:LEU:HG	2.03	0.88
2:B:353:GLN:CG	3:M:48:ASP:N	2.36	0.88
2:B:513:TRP:H	2:B:551:LEU:CD1	1.83	0.88
1:A:536:MET:O	1:A:537:THR:C	1.98	0.88
2:B:178:ILE:HD13	2:B:218:CYS:HB3	1.55	0.88
2:B:527:PRO:HG2	2:B:587:ARG:CG	2.02	0.88
2:B:531:ARG:N	2:B:591:MET:SD	2.46	0.88
3:M:226:PHE:HB2	3:M:481:VAL:CG2	2.04	0.88
3:M:317:MET:HB2	3:M:321:GLY:C	1.94	0.88
4:S:34:GLN:CD	4:S:58:LEU:HD11	1.95	0.88
4:S:73:ILE:HG23	4:S:88:ILE:HG23	1.37	0.88
2:B:360:LEU:CD1	2:B:391:ALA:CA	2.52	0.87
2:B:577:ASN:O	2:B:578:PRO:O	1.93	0.87
3:M:4:SER:O	3:M:78:ALA:CA	2.21	0.87
4:S:17:VAL:HG21	4:S:19:PHE:CE2	2.09	0.87
1:A:178:ARG:NE	1:A:209:ASP:OD2	2.07	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:464:ILE:O	1:A:465:SER:C	1.96	0.87
2:B:215:TYR:HE2	2:B:229:HIS:HD1	1.22	0.87
2:B:317:VAL:O	2:B:321:CYS:SG	2.32	0.87
3:M:243:ILE:CB	3:M:473:LYS:O	2.21	0.87
3:M:245:ASP:HB3	3:M:472:TYR:HD1	1.39	0.87
3:M:336:ASP:OD2	3:M:415:ILE:HB	1.74	0.87
4:S:130:SER:OG	4:S:156:LEU:HD13	1.70	0.87
1:A:64:LEU:CA	1:A:102:GLN:HE22	1.88	0.87
1:A:212:ILE:HD12	4:S:145:ASN:CG	1.95	0.87
1:A:629:LEU:HD21	2:B:610:ARG:HH11	1.38	0.87
2:B:127:LEU:HB3	2:B:157:THR:CG2	1.95	0.87
2:B:154:ILE:HD12	2:B:180:LEU:CG	2.04	0.87
2:B:247:TYR:HE2	3:M:91:THR:HG21	1.09	0.87
2:B:278:PRO:C	2:B:288:TYR:CB	2.38	0.87
2:B:393:ILE:HG23	2:B:431:MET:HG2	1.35	0.87
2:B:451:MET:HG3	2:B:489:ILE:CD1	2.04	0.87
3:M:215:TYR:N	3:M:467:TYR:HB3	1.88	0.87
4:S:54:PRO:O	4:S:57:LEU:HD13	1.72	0.87
1:A:140:VAL:CB	1:A:177:ILE:HG13	2.02	0.87
1:A:633:PHE:HD2	2:B:550:VAL:C	1.74	0.87
2:B:215:TYR:CD2	2:B:219:TYR:HE1	1.92	0.87
2:B:344:VAL:CG1	2:B:381:PHE:CZ	2.58	0.87
2:B:374:PHE:CE1	2:B:381:PHE:CE1	2.63	0.87
3:M:215:TYR:HB3	3:M:467:TYR:HD2	1.39	0.87
3:M:454:ILE:HG21	3:M:464:THR:HG21	0.88	0.87
1:A:557:LYS:HB3	2:B:605:PHE:HD2	1.24	0.87
2:B:121:ASN:OD1	2:B:153:ILE:HD13	1.73	0.87
2:B:174:ALA:HB3	2:B:211:ALA:HA	1.56	0.87
2:B:321:CYS:O	2:B:325:LEU:HG	1.73	0.87
2:B:592:TYR:OH	2:B:619:ASP:OD1	1.93	0.87
3:M:340:LEU:HB3	3:M:411:LEU:HG	1.54	0.87
3:M:374:TYR:OH	3:M:394:GLN:C	2.13	0.87
1:A:105:VAL:H	4:S:167:ILE:HD11	0.72	0.87
1:A:461:CYS:SG	1:A:469:LEU:HD23	2.14	0.87
2:B:20:ARG:NH1	2:B:21:GLU:HG2	1.85	0.87
2:B:143:SER:HA	2:B:179:LYS:HB2	1.56	0.87
2:B:178:ILE:HG21	2:B:214:ALA:HA	1.55	0.87
2:B:297:PRO:O	2:B:301:LEU:HG	1.73	0.87
2:B:513:TRP:CB	2:B:551:LEU:HD21	2.02	0.87
3:M:3:LEU:HB2	3:M:20:LEU:HD12	1.55	0.87
3:M:18:TYR:HD1	3:M:122:SER:HA	1.04	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:217:ASP:CG	3:M:470:ALA:O	2.13	0.87
3:M:259:LYS:O	3:M:449:VAL:HA	1.73	0.87
1:A:557:LYS:CB	2:B:605:PHE:CD2	2.55	0.87
2:B:83:PHE:HE2	2:B:119:SER:CB	1.51	0.87
2:B:214:ALA:O	2:B:216:LYS:N	2.07	0.87
3:M:214:LEU:C	3:M:467:TYR:CB	2.43	0.87
1:A:294:SER:O	1:A:298:ILE:HG12	1.75	0.87
2:B:153:ILE:O	2:B:155:LEU:N	2.08	0.87
2:B:223:LEU:CD2	2:B:255:TYR:CD1	2.56	0.87
2:B:236:ILE:HG22	2:B:240:LEU:HD11	1.54	0.87
2:B:408:VAL:CG1	2:B:412:PHE:CE2	2.58	0.87
2:B:497:LEU:O	2:B:498:THR:C	2.03	0.87
2:B:559:ASP:HA	2:B:562:ASN:HB2	1.53	0.87
3:M:44:ASP:O	3:M:46:SER:N	2.07	0.87
3:M:212:ASN:HB3	3:M:250:LEU:HD23	1.56	0.87
3:M:219:LEU:O	3:M:474:THR:CG2	2.23	0.87
3:M:306:LEU:O	3:M:307:SER:C	2.04	0.87
4:S:105:PHE:CZ	4:S:128:LEU:HD11	2.10	0.87
1:A:213:SER:CB	4:S:143:GLU:OE1	2.23	0.87
1:A:503:ASN:OD1	3:M:59:ASP:OD2	1.93	0.87
2:B:106:LEU:CD1	2:B:144:ASP:C	2.43	0.87
2:B:215:TYR:CD2	2:B:226:LEU:CD1	2.58	0.87
2:B:219:TYR:CB	2:B:226:LEU:HD22	2.02	0.87
2:B:452:LYS:NZ	2:B:456:ASP:OD1	2.08	0.87
3:M:219:LEU:CD2	3:M:473:LYS:HA	2.04	0.87
4:S:83:LEU:HD11	4:S:116:VAL:CG2	2.05	0.87
1:A:67:LYS:HG3	4:S:165:SER:HG	1.40	0.86
1:A:125:THR:HG1	1:A:158:LEU:HD13	1.40	0.86
1:A:633:PHE:CE1	2:B:513:TRP:HE3	1.39	0.86
2:B:196:LEU:HB2	2:B:229:HIS:ND1	1.90	0.86
2:B:208:ILE:CD1	2:B:236:ILE:HG23	1.80	0.86
2:B:267:ASP:O	2:B:276:SER:OG	1.93	0.86
2:B:378:THR:HG23	2:B:379:LYS:H	1.38	0.86
3:M:223:HIS:HD2	3:M:478:ASN:HB2	1.40	0.86
3:M:379:LEU:CD2	3:M:386:PHE:CB	2.53	0.86
4:S:50:PHE:HB2	4:S:76:ILE:HD13	1.56	0.86
1:A:76:TYR:HH	4:S:125:TRP:HZ3	1.20	0.86
1:A:264:SER:CB	1:A:271:ARG:CG	2.54	0.86
1:A:633:PHE:HD2	2:B:551:LEU:HA	1.09	0.86
2:B:165:PRO:HA	2:B:170:ARG:HH21	1.38	0.86
2:B:245:GLN:OE1	2:B:309:LEU:HD12	1.75	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:252:LEU:HD13	2:B:302:PHE:HD1	1.04	0.86
2:B:267:ASP:H	2:B:289:PRO:CG	1.87	0.86
1:A:64:LEU:HA	1:A:102:GLN:NE2	1.90	0.86
2:B:232:ARG:O	2:B:236:ILE:HG13	1.74	0.86
4:S:111:ARG:HB2	4:S:150:VAL:HG22	1.58	0.86
2:B:219:TYR:CD2	2:B:223:LEU:HA	2.11	0.86
2:B:243:TRP:CH2	3:M:91:THR:O	2.27	0.86
2:B:353:GLN:HB2	3:M:50:TYR:N	1.87	0.86
3:M:223:HIS:CG	3:M:478:ASN:HA	2.11	0.86
1:A:426:ILE:HG13	1:A:464:ILE:CD1	2.04	0.86
1:A:637:GLU:O	2:B:518:ILE:N	1.86	0.86
2:B:161:LEU:HB3	2:B:173:VAL:CG2	2.05	0.86
2:B:232:ARG:HG3	2:B:236:ILE:CD1	2.06	0.86
2:B:252:LEU:CB	2:B:302:PHE:CZ	2.59	0.86
2:B:275:ARG:HB3	2:B:291:TYR:CB	2.02	0.86
2:B:143:SER:CA	2:B:179:LYS:HD2	2.05	0.86
3:M:45:SER:HB2	3:M:51:LEU:CD1	2.05	0.86
3:M:242:GLY:HA3	3:M:444:ALA:HB2	1.58	0.86
1:A:76:TYR:OH	4:S:125:TRP:HZ3	1.57	0.86
1:A:316:LEU:HD11	1:A:348:PHE:CD2	2.07	0.86
1:A:421:PRO:HG2	1:A:424:TYR:CD1	2.11	0.86
1:A:630:PRO:O	2:B:554:LYS:N	2.07	0.86
2:B:232:ARG:HG3	2:B:236:ILE:HD11	1.58	0.86
2:B:389:ILE:HG21	2:B:427:ASN:HB2	1.56	0.86
4:S:8:PHE:CZ	4:S:84:TYR:HB2	1.96	0.86
4:S:135:ILE:HG23	4:S:141:VAL:CG1	2.04	0.86
1:A:141:VAL:CB	4:S:159:ALA:N	2.38	0.86
2:B:178:ILE:CD1	2:B:215:TYR:CA	2.51	0.86
2:B:215:TYR:CD2	2:B:226:LEU:HD12	2.10	0.86
2:B:534:ILE:O	2:B:598:LEU:HD11	1.76	0.86
2:B:546:CYS:HB2	2:B:607:ILE:HD11	1.55	0.86
2:B:566:ALA:CA	2:B:574:ASN:ND2	2.38	0.86
3:M:6:TYR:CD2	3:M:17:GLN:CA	2.47	0.86
3:M:323:MET:HE1	3:M:342:LEU:HB3	1.56	0.86
4:S:48:SER:CA	4:S:77:TYR:CB	2.51	0.86
1:A:220:SER:HB2	4:S:141:VAL:O	1.76	0.86
2:B:79:VAL:HG23	2:B:108:PHE:HZ	1.40	0.86
2:B:513:TRP:CD2	2:B:551:LEU:HD21	2.11	0.86
4:S:39:ILE:HD11	4:S:77:TYR:CG	2.09	0.86
1:A:132:LEU:HD13	1:A:165:ASP:CB	2.05	0.86
2:B:267:ASP:C	2:B:289:PRO:HG3	1.95	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:553:ALA:CB	2:B:614:ILE:HD13	2.02	0.86
2:B:553:ALA:HA	2:B:614:ILE:HG21	1.56	0.86
3:M:258:VAL:HG22	3:M:452:ILE:CG2	2.06	0.86
4:S:54:PRO:CG	4:S:57:LEU:CD1	2.46	0.86
1:A:103:LYS:N	4:S:163:THR:HB	1.91	0.85
2:B:223:LEU:CD1	2:B:258:GLN:C	2.38	0.85
3:M:18:TYR:CE1	3:M:122:SER:OG	2.29	0.85
3:M:253:ASN:OD1	3:M:292:PRO:HG2	1.75	0.85
3:M:290:PHE:CZ	3:M:297:PHE:CE1	2.64	0.85
3:M:405:THR:HG22	3:M:406:GLY:H	1.40	0.85
4:S:14:PRO:CA	4:S:36:TYR:OH	2.22	0.85
2:B:2:VAL:HG11	2:B:6:HIS:NE2	1.90	0.85
2:B:127:LEU:CG	2:B:157:THR:HG23	2.00	0.85
2:B:151:ALA:O	2:B:188:TYR:CD2	2.29	0.85
2:B:196:LEU:CB	2:B:229:HIS:ND1	2.39	0.85
2:B:280:PRO:O	2:B:283:TYR:HB2	1.77	0.85
3:M:2:TYR:C	3:M:81:SER:HB2	1.95	0.85
3:M:323:MET:SD	3:M:342:LEU:CG	2.64	0.85
4:S:135:ILE:HG23	4:S:141:VAL:HG13	1.56	0.85
3:M:2:TYR:N	3:M:81:SER:HB2	1.88	0.85
3:M:222:PHE:CG	3:M:439:TYR:HE1	1.94	0.85
3:M:360:LEU:HD23	3:M:362:PHE:CE2	2.08	0.85
1:A:147:LEU:CB	1:A:184:ALA:CB	2.54	0.85
1:A:332:TYR:CE1	1:A:366:SER:CB	2.58	0.85
2:B:56:SER:CB	2:B:92:THR:CG2	2.52	0.85
2:B:191:GLU:C	2:B:193:LEU:H	1.78	0.85
2:B:355:ASN:C	2:B:359:LEU:HD23	1.96	0.85
3:M:215:TYR:HD2	3:M:470:ALA:N	1.74	0.85
1:A:64:LEU:CA	1:A:102:GLN:NE2	2.39	0.85
1:A:240:LEU:O	1:A:241:TYR:C	2.00	0.85
1:A:252:ILE:CG1	4:S:145:ASN:N	2.36	0.85
1:A:516:ILE:HD13	1:A:551:LEU:CA	2.05	0.85
2:B:70:MET:CE	2:B:104:TYR:O	2.24	0.85
2:B:196:LEU:CB	2:B:229:HIS:CE1	2.58	0.85
2:B:316:THR:CG2	3:M:90:PHE:CZ	2.59	0.85
4:S:135:ILE:O	4:S:141:VAL:HG22	1.77	0.85
1:A:217:ALA:CB	4:S:142:ILE:HB	2.05	0.85
2:B:374:PHE:CZ	2:B:381:PHE:CD1	2.65	0.85
2:B:396:ILE:HG12	2:B:418:TYR:HE2	1.37	0.85
2:B:566:ALA:N	2:B:574:ASN:ND2	2.24	0.85
1:A:95:MET:C	1:A:127:LEU:HD21	1.94	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:381:GLU:HA	1:A:384:LEU:HD23	1.59	0.85
1:A:384:LEU:CD2	1:A:441:TYR:HE2	1.81	0.85
2:B:162:VAL:HB	2:B:195:ILE:HG23	1.57	0.85
2:B:177:ILE:HG21	2:B:196:LEU:HG	1.56	0.85
2:B:404:ASN:O	2:B:408:VAL:HG23	1.76	0.85
3:M:6:TYR:O	3:M:77:LEU:N	2.10	0.85
3:M:7:ILE:CA	3:M:75:TRP:O	2.24	0.85
4:S:28:GLN:O	4:S:32:LEU:HG	1.75	0.85
1:A:178:ARG:HH11	1:A:209:ASP:HB3	1.40	0.85
1:A:196:LEU:O	1:A:196:LEU:HD22	1.75	0.85
2:B:21:GLU:CA	2:B:24:ALA:HB3	2.07	0.85
2:B:352:ASN:HB2	3:M:49:ASP:CB	2.06	0.85
2:B:396:ILE:CG1	2:B:418:TYR:CE2	2.59	0.85
2:B:519:ALA:C	2:B:523:PHE:HB3	1.96	0.85
2:B:592:TYR:CD2	2:B:618:PHE:CE2	2.64	0.85
3:M:258:VAL:HA	3:M:452:ILE:HG13	1.56	0.85
1:A:225:LEU:CB	1:A:233:PHE:CZ	2.60	0.85
1:A:288:THR:CB	1:A:322:PHE:CZ	2.59	0.85
1:A:298:ILE:HD11	1:A:311:THR:CG2	2.07	0.85
2:B:20:ARG:NE	2:B:21:GLU:CB	2.39	0.85
3:M:121:ILE:HG22	3:M:125:PHE:CE1	2.12	0.85
3:M:215:TYR:CD1	3:M:467:TYR:O	2.27	0.85
3:M:226:PHE:CB	3:M:481:VAL:HG22	2.04	0.85
1:A:95:MET:C	1:A:127:LEU:CD2	2.45	0.85
1:A:608:ARG:NH2	1:A:632:PHE:HZ	1.74	0.85
1:A:633:PHE:HB2	2:B:550:VAL:HG13	1.56	0.85
2:B:143:SER:OG	2:B:175:LEU:HB3	1.75	0.85
2:B:230:PHE:HZ	2:B:252:LEU:CD2	1.58	0.85
2:B:562:ASN:CG	2:B:580:TYR:HB2	1.97	0.85
2:B:562:ASN:CG	2:B:580:TYR:CB	2.45	0.85
3:M:342:LEU:CD1	3:M:411:LEU:HD22	2.05	0.85
4:S:57:LEU:HB3	4:S:67:GLU:O	1.76	0.85
1:A:101:GLN:HG2	4:S:167:ILE:CG1	2.07	0.84
1:A:147:LEU:CD1	1:A:181:ALA:CA	2.53	0.84
1:A:482:ILE:HG12	1:A:517:TRP:CH2	2.12	0.84
1:A:557:LYS:CG	2:B:605:PHE:HD2	1.90	0.84
1:A:563:CYS:HB2	1:A:621:LEU:HD12	1.58	0.84
1:A:606:PHE:HZ	2:B:550:VAL:CG1	1.90	0.84
1:A:636:TYR:CB	2:B:554:LYS:HZ2	1.89	0.84
2:B:268:LYS:HA	2:B:276:SER:HB2	1.58	0.84
1:A:251:TRP:CD2	4:S:104:THR:OG1	2.30	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:636:TYR:CB	2:B:554:LYS:NZ	2.39	0.84
2:B:82:TYR:O	2:B:83:PHE:C	2.06	0.84
2:B:260:LEU:HA	2:B:291:TYR:CE1	2.11	0.84
1:A:88:ASN:CB	1:A:120:ILE:CD1	2.51	0.84
1:A:132:LEU:O	1:A:169:MET:HE3	1.77	0.84
1:A:638:LEU:CB	2:B:516:GLY:O	2.23	0.84
2:B:380:LYS:HZ3	3:M:236:LEU:HG	1.36	0.84
3:M:6:TYR:HA	3:M:16:PHE:O	1.77	0.84
1:A:260:PHE:CZ	1:A:274:LEU:CD1	2.59	0.84
1:A:266:VAL:O	1:A:267:GLU:CB	2.25	0.84
1:A:557:LYS:O	2:B:605:PHE:HE2	1.57	0.84
2:B:223:LEU:HD12	2:B:259:TYR:HA	0.87	0.84
2:B:549:LEU:CG	2:B:611:ALA:CB	2.54	0.84
3:M:224:VAL:HG23	3:M:479:PHE:CE1	2.13	0.84
3:M:290:PHE:CE2	3:M:297:PHE:CZ	2.64	0.84
3:M:347:PHE:CD2	3:M:350:VAL:HB	2.13	0.84
4:S:48:SER:CB	4:S:77:TYR:CA	2.56	0.84
1:A:103:LYS:HG3	4:S:163:THR:HG21	1.55	0.84
1:A:401:VAL:HG23	1:A:418:ILE:O	1.76	0.84
1:A:408:ILE:CG2	4:S:64:ASN:HB2	1.82	0.84
2:B:162:VAL:O	2:B:164:ASP:N	2.10	0.84
2:B:245:GLN:OE1	2:B:309:LEU:CD1	2.24	0.84
2:B:418:TYR:C	2:B:418:TYR:HD1	1.77	0.84
2:B:569:THR:C	2:B:571:SER:H	1.76	0.84
1:A:96:SER:HB3	1:A:127:LEU:CD2	2.07	0.84
1:A:504:ILE:O	1:A:505:ASN:C	2.06	0.84
1:A:589:SER:HB2	1:A:601:VAL:CG2	2.08	0.84
2:B:42:ILE:CG1	2:B:65:ARG:HD3	2.06	0.84
3:M:323:MET:HB3	3:M:340:LEU:CD1	2.06	0.84
4:S:83:LEU:CD1	4:S:116:VAL:HG21	2.07	0.84
1:A:103:LYS:HB2	4:S:163:THR:HG21	1.57	0.84
1:A:125:THR:OG1	1:A:158:LEU:CD1	2.23	0.84
1:A:259:LEU:O	1:A:262:ASN:N	2.10	0.84
1:A:533:ILE:HG12	1:A:562:TRP:HH2	1.41	0.84
1:A:595:GLU:CB	2:B:469:ASP:HB3	2.07	0.84
1:A:605:GLU:OE2	1:A:632:PHE:CE2	2.31	0.84
2:B:418:TYR:O	2:B:419:VAL:O	1.93	0.84
3:M:19:LEU:CD1	3:M:24:ALA:HB3	2.07	0.84
3:M:82:LYS:O	3:M:84:LYS:HG2	1.78	0.84
3:M:212:ASN:CG	3:M:250:LEU:HD23	1.97	0.84
1:A:103:LYS:CD	1:A:131:ASP:CG	2.46	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:629:LEU:O	1:A:630:PRO:C	2.00	0.84
2:B:490:ILE:O	2:B:515:PHE:CZ	2.30	0.84
3:M:254:PRO:CB	3:M:454:ILE:HD12	2.06	0.84
4:S:83:LEU:CD1	4:S:116:VAL:HG11	2.07	0.84
1:A:64:LEU:HA	1:A:102:GLN:HE22	1.40	0.84
1:A:179:LYS:HG3	1:A:217:ALA:HB2	1.58	0.84
2:B:48:VAL:CG2	2:B:82:TYR:CE2	2.60	0.84
3:M:233:LEU:HD22	3:M:324:SER:HA	1.59	0.84
1:A:225:LEU:HD12	1:A:233:PHE:CZ	2.13	0.84
2:B:182:ARG:CD	2:B:217:GLU:OE1	2.25	0.84
2:B:353:GLN:HE21	3:M:47:SER:HB3	1.04	0.84
2:B:396:ILE:HD13	2:B:432:ALA:CB	2.08	0.84
3:M:293:PRO:O	3:M:293:PRO:CD	2.26	0.84
4:S:15:ARG:HD2	4:S:122:ILE:HG12	1.58	0.84
1:A:67:LYS:HB2	4:S:165:SER:C	1.97	0.83
1:A:101:GLN:HG2	4:S:167:ILE:HG13	1.59	0.83
1:A:253:ILE:HD13	1:A:281:LEU:CB	2.07	0.83
2:B:264:THR:O	2:B:266:VAL:HG23	1.78	0.83
2:B:534:ILE:CD1	2:B:594:ALA:HB3	1.99	0.83
3:M:215:TYR:CG	3:M:469:GLY:N	2.25	0.83
3:M:223:HIS:CD2	3:M:478:ASN:CB	2.61	0.83
4:S:5:VAL:HG21	4:S:132:LEU:CG	2.08	0.83
1:A:66:SER:N	4:S:165:SER:HB2	1.93	0.83
2:B:70:MET:HE1	2:B:107:ARG:CG	2.07	0.83
2:B:274:PRO:CA	2:B:295:ASN:HD21	1.90	0.83
2:B:316:THR:OG1	3:M:90:PHE:CZ	2.19	0.83
2:B:336:ASN:O	2:B:337:THR:CB	2.20	0.83
2:B:344:VAL:HG22	2:B:363:ILE:CD1	2.06	0.83
3:M:101:LEU:O	3:M:106:LYS:CA	2.26	0.83
1:A:101:GLN:HE21	4:S:167:ILE:HG21	1.40	0.83
1:A:103:LYS:N	4:S:163:THR:CG2	2.40	0.83
1:A:211:ASP:CG	4:S:148:ARG:CD	2.31	0.83
1:A:320:HIS:O	1:A:321:THR:C	2.14	0.83
1:A:581:LEU:HD23	1:A:607:LEU:CD1	2.06	0.83
1:A:586:GLU:HB2	1:A:604:LEU:CD1	2.07	0.83
1:A:630:PRO:HG2	2:B:614:ILE:CB	2.05	0.83
1:A:634:ASN:O	2:B:517:GLU:CA	2.27	0.83
2:B:25:VAL:CG2	2:B:35:TYR:CD2	2.49	0.83
2:B:102:HIS:ND1	2:B:137:PHE:HB3	1.90	0.83
2:B:275:ARG:CB	2:B:291:TYR:CD2	2.61	0.83
2:B:512:VAL:HG21	2:B:548:ILE:HG12	1.58	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:531:ARG:CB	2:B:591:MET:SD	2.66	0.83
2:B:549:LEU:HD13	2:B:611:ALA:HB1	1.56	0.83
3:M:403:THR:CG2	3:M:407:THR:OG1	2.26	0.83
1:A:253:ILE:O	1:A:257:LEU:HG	1.78	0.83
1:A:384:LEU:HD13	1:A:441:TYR:HE2	1.44	0.83
1:A:585:PHE:CE2	1:A:607:LEU:HD11	2.14	0.83
2:B:177:ILE:HD12	2:B:196:LEU:HD23	1.59	0.83
2:B:232:ARG:HG3	2:B:236:ILE:CG1	2.06	0.83
2:B:513:TRP:CA	2:B:551:LEU:HD11	2.07	0.83
4:S:61:ASN:OD1	4:S:66:ASP:OD2	1.95	0.83
1:A:255:ARG:HH21	4:S:135:ILE:HG23	1.43	0.83
1:A:331:ARG:NH1	1:A:362:ASP:OD2	2.12	0.83
2:B:159:LYS:CA	2:B:195:ILE:CD1	2.50	0.83
4:S:53:THR:CA	4:S:69:ASN:CB	2.53	0.83
1:A:140:VAL:HA	1:A:177:ILE:CD1	2.08	0.83
1:A:306:GLU:O	1:A:307:ASP:C	2.12	0.83
1:A:332:TYR:HE1	1:A:366:SER:CB	1.92	0.83
2:B:173:VAL:HB	2:B:199:LEU:CD1	2.09	0.83
2:B:177:ILE:O	2:B:181:TYR:HB2	1.78	0.83
2:B:292:GLU:HG2	2:B:296:ASP:CB	2.09	0.83
3:M:121:ILE:O	3:M:125:PHE:CD1	2.31	0.83
3:M:261:ASN:O	3:M:446:GLY:O	1.97	0.83
4:S:5:VAL:CG1	4:S:132:LEU:CD2	2.49	0.83
1:A:403:LEU:HD11	1:A:421:PRO:O	1.79	0.83
1:A:429:VAL:HG21	1:A:469:LEU:HD21	1.60	0.83
2:B:178:ILE:CG2	2:B:214:ALA:O	2.26	0.83
3:M:220:GLU:CG	3:M:439:TYR:HB2	2.07	0.83
3:M:262:THR:HG22	3:M:264:GLY:CA	2.09	0.83
1:A:95:MET:SD	1:A:107:TYR:HD2	2.00	0.83
1:A:163:ALA:O	1:A:164:ASP:C	2.08	0.83
3:M:218:LEU:HG	3:M:472:TYR:HE2	1.43	0.83
3:M:224:VAL:O	3:M:479:PHE:HB3	1.79	0.83
2:B:20:ARG:HH11	2:B:21:GLU:CG	1.91	0.83
2:B:155:LEU:CB	2:B:188:TYR:HD2	1.91	0.83
2:B:243:TRP:HZ2	3:M:94:GLU:C	1.82	0.83
1:A:182:ILE:CD1	1:A:218:ALA:HB2	2.07	0.83
1:A:275:LEU:HD21	1:A:311:THR:HG1	1.44	0.83
1:A:403:LEU:CD2	1:A:422:GLU:CG	2.52	0.83
1:A:637:GLU:HG3	2:B:515:PHE:N	1.68	0.83
4:S:9:ASN:ND2	4:S:13:GLN:HB2	1.93	0.83
4:S:53:THR:O	4:S:69:ASN:CG	2.17	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:559:PHE:HE1	1:A:581:LEU:HD22	1.43	0.82
1:A:633:PHE:CD2	2:B:550:VAL:O	2.31	0.82
2:B:50:LEU:HG	2:B:58:GLU:O	1.79	0.82
2:B:231:ARG:HH21	2:B:297:PRO:HD2	1.44	0.82
2:B:281:ASP:OD1	2:B:287:GLU:OE2	1.97	0.82
2:B:566:ALA:C	2:B:574:ASN:CG	2.38	0.82
3:M:219:LEU:HB3	3:M:472:TYR:O	1.79	0.82
1:A:65:ASN:O	4:S:165:SER:HA	1.79	0.82
1:A:140:VAL:N	1:A:177:ILE:CD1	2.41	0.82
1:A:151:SER:HB2	1:A:187:LYS:HB3	1.59	0.82
1:A:348:PHE:O	1:A:352:PHE:CD1	2.32	0.82
1:A:566:PHE:CE1	1:A:618:THR:HB	2.14	0.82
2:B:393:ILE:HG12	2:B:428:VAL:HA	1.62	0.82
1:A:103:LYS:CD	1:A:131:ASP:OD1	2.28	0.82
1:A:163:ALA:HB1	1:A:199:ASN:ND2	1.92	0.82
1:A:621:LEU:O	1:A:622:PRO:C	1.97	0.82
1:A:634:ASN:O	2:B:516:GLY:O	1.95	0.82
2:B:42:ILE:HG12	2:B:65:ARG:HD3	1.60	0.82
2:B:171:GLY:HA2	2:B:207:VAL:HG12	1.61	0.82
2:B:505:ASP:HA	2:B:544:THR:OG1	1.79	0.82
2:B:522:GLU:O	2:B:522:GLU:HG2	1.79	0.82
3:M:262:THR:HG22	3:M:264:GLY:H	1.43	0.82
4:S:47:GLN:HE22	4:S:78:LYS:HA	1.45	0.82
1:A:260:PHE:CE2	1:A:274:LEU:HG	2.14	0.82
1:A:564:ASN:OD1	1:A:622:PRO:HG3	1.78	0.82
2:B:127:LEU:HB3	2:B:161:LEU:HD13	1.61	0.82
2:B:158:VAL:CG1	2:B:195:ILE:HG21	2.08	0.82
2:B:197:LYS:HD2	2:B:283:TYR:CE2	2.14	0.82
2:B:310:ILE:HA	2:B:318:ILE:HG12	1.61	0.82
2:B:527:PRO:HB2	2:B:587:ARG:CD	2.09	0.82
2:B:562:ASN:CB	2:B:580:TYR:HB3	2.08	0.82
4:S:83:LEU:HD11	4:S:116:VAL:CG1	2.07	0.82
1:A:114:PHE:CG	1:A:153:ILE:HG23	2.13	0.82
1:A:140:VAL:CG1	1:A:176:TYR:HB2	2.10	0.82
1:A:144:GLY:HA2	1:A:180:LYS:HB2	1.61	0.82
2:B:70:MET:HE3	2:B:107:ARG:HG3	1.59	0.82
2:B:178:ILE:C	2:B:180:LEU:H	1.82	0.82
2:B:311:TYR:HE2	2:B:342:ALA:HB1	1.40	0.82
2:B:418:TYR:CD1	2:B:419:VAL:N	2.47	0.82
2:B:493:LEU:CG	2:B:511:ILE:HG23	2.09	0.82
2:B:542:PRO:O	2:B:602:ASP:OD2	1.95	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:592:TYR:CD2	2:B:618:PHE:CD2	2.67	0.82
3:M:231:SER:O	3:M:326:HIS:HE1	1.62	0.82
1:A:166:LEU:HD13	1:A:185:LEU:CG	2.09	0.82
1:A:432:ILE:HG21	1:A:457:LEU:HD11	1.59	0.82
2:B:42:ILE:O	2:B:46:GLN:OE1	1.96	0.82
2:B:309:LEU:HD12	2:B:317:VAL:HG11	1.62	0.82
2:B:344:VAL:HG13	2:B:381:PHE:HZ	1.42	0.82
2:B:396:ILE:CD1	2:B:418:TYR:CZ	2.62	0.82
2:B:512:VAL:HB	2:B:551:LEU:HD12	1.62	0.82
1:A:217:ALA:O	4:S:142:ILE:HG22	1.77	0.82
1:A:563:CYS:CB	1:A:621:LEU:CD1	2.55	0.82
1:A:563:CYS:SG	1:A:621:LEU:CD1	2.68	0.82
2:B:182:ARG:CZ	2:B:217:GLU:OE1	2.27	0.82
2:B:431:MET:O	2:B:434:LYS:N	2.12	0.82
3:M:3:LEU:HG	3:M:80:THR:O	1.79	0.82
3:M:101:LEU:CA	3:M:109:LEU:HD13	2.09	0.82
3:M:243:ILE:CG1	3:M:473:LYS:O	2.27	0.82
1:A:140:VAL:CG1	1:A:176:TYR:CB	2.52	0.82
1:A:255:ARG:HD3	4:S:141:VAL:O	1.80	0.82
1:A:258:LYS:HZ1	4:S:97:ALA:CB	1.92	0.82
1:A:258:LYS:NZ	4:S:97:ALA:CB	2.42	0.82
2:B:230:PHE:HB3	2:B:298:ASP:OD2	1.80	0.82
2:B:177:ILE:HD11	2:B:195:ILE:CG2	2.10	0.82
2:B:197:LYS:HD2	2:B:283:TYR:CD2	2.15	0.82
2:B:252:LEU:HB2	2:B:302:PHE:CE1	2.15	0.82
2:B:568:VAL:O	2:B:571:SER:CB	2.28	0.82
2:B:569:THR:C	2:B:571:SER:N	2.28	0.82
1:A:288:THR:CB	1:A:322:PHE:HZ	1.93	0.82
1:A:520:GLY:HA3	1:A:558:VAL:HG22	0.86	0.82
2:B:353:GLN:CB	3:M:50:TYR:N	2.39	0.82
2:B:360:LEU:HD12	2:B:391:ALA:HA	1.60	0.82
2:B:437:SER:HB2	2:B:474:VAL:HG13	1.60	0.82
3:M:290:PHE:CE1	3:M:297:PHE:CG	2.66	0.82
3:M:353:VAL:HG22	3:M:354:ASP:O	1.80	0.82
1:A:71:VAL:CG1	1:A:105:VAL:CG1	2.56	0.81
1:A:219:VAL:HG12	1:A:259:LEU:CD1	2.08	0.81
2:B:120:ILE:HG23	2:B:154:ILE:CG1	2.10	0.81
2:B:199:LEU:C	2:B:201:ALA:N	2.30	0.81
2:B:578:PRO:HB2	2:B:579:PRO:CD	2.07	0.81
3:M:241:HIS:O	3:M:474:THR:OG1	1.97	0.81
4:S:54:PRO:CB	4:S:57:LEU:HD11	2.10	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:129:LYS:HD2	1:A:161:ASP:CG	2.00	0.81
1:A:163:ALA:HB1	1:A:199:ASN:HD21	1.43	0.81
1:A:189:PHE:CD2	1:A:225:LEU:CD2	2.60	0.81
1:A:219:VAL:CG1	1:A:256:LEU:HD23	2.09	0.81
1:A:263:LEU:O	1:A:266:VAL:N	2.12	0.81
1:A:298:ILE:O	1:A:299:VAL:C	2.16	0.81
2:B:170:ARG:HG2	2:B:199:LEU:HD23	1.61	0.81
2:B:340:ILE:CD1	2:B:366:LEU:HB3	2.09	0.81
3:M:7:ILE:HA	3:M:75:TRP:O	1.80	0.81
3:M:101:LEU:CD2	3:M:106:LYS:O	2.28	0.81
3:M:344:ILE:HG23	3:M:347:PHE:HB3	0.93	0.81
1:A:147:LEU:CB	1:A:184:ALA:HB2	2.10	0.81
1:A:178:ARG:NH1	1:A:209:ASP:CG	2.33	0.81
1:A:566:PHE:HE1	1:A:618:THR:HB	1.43	0.81
2:B:90:ILE:CD1	2:B:123:LEU:CD2	2.58	0.81
2:B:116:THR:HG22	2:B:150:LEU:HD21	1.60	0.81
2:B:171:GLY:N	2:B:207:VAL:CG1	2.34	0.81
2:B:239:GLN:OE1	3:M:278:ILE:O	1.98	0.81
2:B:374:PHE:CD2	2:B:402:LEU:HD11	2.15	0.81
2:B:490:ILE:HG23	2:B:515:PHE:CE2	2.16	0.81
2:B:527:PRO:HG2	2:B:587:ARG:HG2	1.61	0.81
2:B:556:LEU:CB	2:B:588:ILE:HD11	2.10	0.81
3:M:245:ASP:CA	3:M:472:TYR:CE1	2.63	0.81
3:M:257:ALA:HB2	3:M:455:VAL:HG21	1.60	0.81
3:M:374:TYR:HA	3:M:417:TYR:HA	1.63	0.81
1:A:298:ILE:CD1	1:A:311:THR:HG21	2.10	0.81
3:M:339:GLU:OE2	3:M:412:ARG:NE	2.07	0.81
1:A:96:SER:HB3	1:A:127:LEU:HD13	1.60	0.81
1:A:103:LYS:CE	1:A:131:ASP:CG	2.49	0.81
1:A:104:ARG:HE	4:S:160:ALA:HB2	1.45	0.81
1:A:408:ILE:HG21	4:S:64:ASN:O	1.79	0.81
2:B:120:ILE:HG23	2:B:142:LEU:HD22	1.63	0.81
2:B:182:ARG:HD2	2:B:217:GLU:CB	2.10	0.81
2:B:182:ARG:CG	2:B:217:GLU:HB3	2.10	0.81
2:B:553:ALA:CA	2:B:614:ILE:HG21	2.11	0.81
3:M:458:LEU:HD22	3:M:462:LYS:HG2	1.62	0.81
1:A:211:ASP:OD1	1:A:213:SER:N	2.14	0.81
1:A:634:ASN:HD22	2:B:554:LYS:HB3	1.46	0.81
2:B:109:ALA:O	2:B:110:GLU:C	2.06	0.81
2:B:123:LEU:CB	2:B:142:LEU:HD11	2.09	0.81
2:B:158:VAL:HG13	2:B:177:ILE:HD11	1.61	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:223:LEU:O	2:B:224:GLU:C	2.11	0.81
2:B:277:CYS:HA	2:B:292:GLU:HG3	1.61	0.81
2:B:303:LEU:CD1	2:B:333:GLN:CG	2.55	0.81
3:M:245:ASP:HB3	3:M:472:TYR:CD1	2.16	0.81
4:S:48:SER:C	4:S:77:TYR:HB2	2.00	0.81
1:A:71:VAL:HG11	1:A:105:VAL:CG1	2.11	0.81
1:A:260:PHE:CE1	1:A:274:LEU:HD12	2.15	0.81
1:A:567:GLN:O	1:A:568:GLU:C	2.17	0.81
1:A:586:GLU:N	1:A:604:LEU:HD12	1.94	0.81
2:B:90:ILE:HD13	2:B:123:LEU:CD2	2.10	0.81
3:M:49:ASP:HA	3:M:75:TRP:HH2	1.40	0.81
3:M:101:LEU:HG	3:M:106:LYS:HA	0.81	0.81
1:A:96:SER:H	1:A:127:LEU:HD21	0.98	0.81
1:A:102:GLN:NE2	4:S:165:SER:H	1.79	0.81
1:A:225:LEU:HD12	1:A:233:PHE:CE2	2.16	0.81
1:A:346:THR:O	1:A:347:ASP:C	2.10	0.81
1:A:532:LEU:O	1:A:533:ILE:C	2.00	0.81
2:B:151:ALA:CA	2:B:180:LEU:CD1	2.44	0.81
2:B:396:ILE:CD1	2:B:418:TYR:OH	2.29	0.81
2:B:563:PHE:C	2:B:566:ALA:HB3	2.02	0.81
4:S:9:ASN:ND2	4:S:13:GLN:CG	2.44	0.81
1:A:99:LYS:CE	4:S:164:ASP:HB2	2.11	0.81
1:A:140:VAL:HA	1:A:177:ILE:HG12	0.81	0.81
1:A:140:VAL:CA	1:A:177:ILE:CD1	2.59	0.81
1:A:163:ALA:CB	1:A:199:ASN:HD21	1.94	0.81
1:A:237:SER:HB2	1:A:270:LEU:CD1	2.11	0.81
2:B:47:LEU:HD22	2:B:62:ALA:O	1.80	0.81
2:B:214:ALA:O	2:B:217:GLU:N	2.13	0.81
2:B:219:TYR:O	2:B:220:ALA:C	2.16	0.81
2:B:353:GLN:CB	3:M:49:ASP:H	1.78	0.81
2:B:433:VAL:CG1	2:B:474:VAL:CB	2.51	0.81
2:B:477:MET:O	2:B:480:GLN:HB2	1.80	0.81
2:B:527:PRO:CG	2:B:587:ARG:CG	2.51	0.81
3:M:69:ILE:CD1	3:M:90:PHE:CE1	2.64	0.81
3:M:101:LEU:HD23	3:M:106:LYS:HG3	1.62	0.81
3:M:212:ASN:HB3	3:M:250:LEU:HA	1.63	0.81
1:A:103:LYS:HD3	1:A:131:ASP:OD2	1.81	0.81
1:A:140:VAL:CG1	1:A:176:TYR:HB3	2.09	0.81
1:A:182:ILE:O	1:A:221:VAL:HG21	1.81	0.81
1:A:202:LYS:O	1:A:205:SER:N	2.13	0.81
1:A:631:SER:HB2	2:B:557:SER:CB	1.92	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:77:ILE:O	2:B:78:ASP:C	2.17	0.81
2:B:87:VAL:HG12	2:B:122:SER:CB	1.77	0.81
2:B:343:LEU:HD22	2:B:362:ALA:HB1	1.62	0.81
2:B:545:ARG:HH11	2:B:602:ASP:HA	1.45	0.81
3:M:224:VAL:HG11	3:M:226:PHE:CZ	2.16	0.81
3:M:310:VAL:HG13	3:M:315:VAL:O	1.81	0.81
3:M:342:LEU:CD1	3:M:411:LEU:CD2	2.56	0.81
1:A:68:THR:CA	4:S:167:ILE:N	2.44	0.80
1:A:105:VAL:CA	4:S:167:ILE:CD1	2.59	0.80
1:A:215:VAL:HG22	1:A:243:ILE:CG1	2.10	0.80
1:A:463:ASP:C	2:B:1:MET:SD	2.59	0.80
1:A:637:GLU:HB3	2:B:516:GLY:N	1.96	0.80
2:B:136:CYS:O	2:B:172:GLU:HB3	1.80	0.80
2:B:260:LEU:CD2	2:B:291:TYR:CZ	2.53	0.80
2:B:380:LYS:HZ3	3:M:236:LEU:CG	1.94	0.80
2:B:472:VAL:HG11	2:B:510:GLY:C	2.01	0.80
2:B:560:ILE:HA	2:B:564:LYS:H	1.46	0.80
3:M:7:ILE:CG2	3:M:75:TRP:O	2.29	0.80
1:A:488:ARG:O	1:A:491:THR:OG1	1.99	0.80
2:B:296:ASP:OD1	2:B:297:PRO:CD	2.29	0.80
2:B:337:THR:CA	2:B:373:LEU:CD2	2.40	0.80
2:B:425:PRO:HD2	2:B:428:VAL:HG21	1.61	0.80
2:B:508:ARG:HB3	2:B:544:THR:CG2	2.12	0.80
3:M:223:HIS:HA	3:M:479:PHE:CG	2.13	0.80
3:M:340:LEU:HG	3:M:411:LEU:CB	2.09	0.80
4:S:108:SER:CB	4:S:149:ILE:HG21	2.10	0.80
1:A:244:LEU:CD1	1:A:256:LEU:HD13	2.11	0.80
1:A:264:SER:CB	1:A:271:ARG:CD	2.59	0.80
1:A:408:ILE:HD12	1:A:410:TYR:CD1	2.15	0.80
1:A:609:LEU:CG	1:A:628:VAL:CB	2.48	0.80
1:A:637:GLU:CB	2:B:516:GLY:N	2.44	0.80
2:B:90:ILE:CD1	2:B:123:LEU:HD23	2.11	0.80
2:B:135:ARG:NH1	2:B:164:ASP:HB2	1.97	0.80
2:B:177:ILE:HG21	2:B:196:LEU:CG	2.10	0.80
2:B:178:ILE:HG22	2:B:179:LYS:H	1.44	0.80
2:B:252:LEU:CB	2:B:302:PHE:CE2	2.65	0.80
2:B:256:CYS:HA	2:B:293:VAL:CG2	2.05	0.80
2:B:563:PHE:O	2:B:566:ALA:HB3	1.80	0.80
3:M:215:TYR:CD2	3:M:469:GLY:CA	2.65	0.80
4:S:7:ILE:CG2	4:S:121:LEU:HD21	2.09	0.80
1:A:64:LEU:O	1:A:65:ASN:C	2.19	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:433:VAL:HG11	2:B:471:TYR:HA	1.61	0.80
1:A:96:SER:CB	1:A:127:LEU:CD2	2.60	0.80
1:A:222:ILE:HG23	1:A:233:PHE:HB3	1.64	0.80
2:B:315:PRO:CG	2:B:352:ASN:OD1	2.28	0.80
2:B:546:CYS:HA	2:B:607:ILE:HG23	0.82	0.80
3:M:71:LYS:O	3:M:74:TYR:CE2	2.34	0.80
1:A:101:GLN:NE2	4:S:167:ILE:HG21	1.93	0.80
1:A:207:LEU:HD11	1:A:240:LEU:HD23	1.62	0.80
1:A:253:ILE:HD11	1:A:281:LEU:CD2	2.09	0.80
1:A:279:LEU:O	1:A:280:GLU:C	2.11	0.80
1:A:465:SER:H	2:B:1:MET:HG2	1.47	0.80
1:A:557:LYS:HB3	2:B:605:PHE:CE2	2.15	0.80
1:A:608:ARG:NH2	1:A:632:PHE:CZ	2.49	0.80
2:B:230:PHE:CD2	2:B:298:ASP:CB	2.63	0.80
2:B:486:HIS:NE2	2:B:518:ILE:HB	1.95	0.80
2:B:560:ILE:CG2	2:B:564:LYS:HB2	2.11	0.80
3:M:18:TYR:CD1	3:M:122:SER:CB	2.64	0.80
1:A:170:LEU:O	1:A:206:LYS:CE	2.28	0.80
1:A:213:SER:C	4:S:143:GLU:HG2	2.01	0.80
1:A:384:LEU:CD1	1:A:441:TYR:HE2	1.94	0.80
1:A:399:ASP:HA	1:A:420:ILE:HB	0.87	0.80
1:A:545:HIS:O	1:A:546:SER:C	2.07	0.80
1:A:582:ILE:HD11	1:A:608:ARG:HA	1.64	0.80
2:B:230:PHE:O	2:B:231:ARG:C	1.96	0.80
2:B:266:VAL:HG13	2:B:291:TYR:H	1.46	0.80
2:B:344:VAL:CG2	2:B:363:ILE:HD11	2.12	0.80
2:B:371:GLN:CD	2:B:401:THR:O	2.20	0.80
2:B:525:ILE:C	2:B:527:PRO:CD	2.50	0.80
4:S:14:PRO:HB3	4:S:36:TYR:HE1	1.47	0.80
4:S:108:SER:HB3	4:S:149:ILE:CG2	2.12	0.80
1:A:224:GLU:CG	4:S:138:GLY:O	2.30	0.80
1:A:392:MET:SD	1:A:457:LEU:HD23	2.22	0.80
1:A:537:THR:CB	1:A:584:PHE:CE1	2.64	0.80
2:B:200:MET:HE3	2:B:232:ARG:H	1.47	0.80
3:M:66:PHE:HB3	3:M:77:LEU:CD2	2.10	0.80
4:S:87:PHE:CG	4:S:102:ILE:HG12	2.15	0.80
4:S:109:LEU:O	4:S:110:ASP:C	2.13	0.80
1:A:426:ILE:CG1	1:A:464:ILE:HD13	2.12	0.80
1:A:436:CYS:SG	1:A:450:TYR:CZ	2.75	0.80
2:B:527:PRO:HB2	2:B:587:ARG:HD2	1.62	0.80
2:B:550:VAL:CG2	2:B:610:ARG:CD	2.60	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:71:LYS:O	3:M:72:LEU:CB	2.30	0.80
1:A:151:SER:CB	1:A:187:LYS:HB2	2.12	0.80
2:B:62:ALA:HB1	2:B:66:ILE:HD11	1.64	0.80
2:B:337:THR:HG23	2:B:373:LEU:CD1	2.12	0.80
3:M:245:ASP:O	3:M:472:TYR:CZ	2.34	0.80
1:A:178:ARG:NH1	1:A:209:ASP:CB	2.44	0.79
1:A:291:ILE:HG21	1:A:322:PHE:HE1	1.44	0.79
2:B:47:LEU:CD2	2:B:62:ALA:O	2.30	0.79
2:B:219:TYR:HE1	2:B:226:LEU:HD12	1.47	0.79
2:B:352:ASN:O	2:B:355:ASN:HB2	1.81	0.79
2:B:519:ALA:O	2:B:523:PHE:HD2	1.65	0.79
3:M:223:HIS:CB	3:M:478:ASN:HA	2.12	0.79
1:A:125:THR:CB	1:A:158:LEU:HD13	2.12	0.79
1:A:145:ILE:HG12	4:S:156:LEU:HD22	1.62	0.79
2:B:170:ARG:NH1	2:B:198:GLU:O	2.15	0.79
2:B:189:HIS:CD2	2:B:222:HIS:HB3	2.17	0.79
2:B:341:GLU:HG2	2:B:345:ARG:HE	1.46	0.79
2:B:389:ILE:O	2:B:390:VAL:C	2.17	0.79
2:B:458:MET:O	2:B:459:GLU:C	2.19	0.79
1:A:185:LEU:HD22	1:A:189:PHE:CZ	2.16	0.79
2:B:50:LEU:CB	2:B:62:ALA:HB2	2.13	0.79
2:B:63:MET:CG	2:B:100:LEU:HB3	2.12	0.79
2:B:178:ILE:HG21	2:B:214:ALA:CA	2.12	0.79
2:B:217:GLU:O	2:B:219:TYR:N	2.15	0.79
2:B:245:GLN:NE2	2:B:309:LEU:CD1	2.45	0.79
2:B:374:PHE:CE2	2:B:398:ILE:CG2	2.66	0.79
3:M:379:LEU:HD23	3:M:397:TRP:CD1	2.17	0.79
1:A:96:SER:CB	1:A:127:LEU:HD21	2.11	0.79
1:A:137:ASN:C	1:A:139:ASP:N	2.28	0.79
1:A:182:ILE:HD13	1:A:218:ALA:CB	2.12	0.79
1:A:291:ILE:HG21	1:A:322:PHE:CE1	2.17	0.79
2:B:63:MET:SD	2:B:101:ILE:HA	2.22	0.79
2:B:139:LEU:CD2	2:B:173:VAL:C	2.50	0.79
2:B:162:VAL:HG23	2:B:199:LEU:HD11	1.61	0.79
2:B:174:ALA:HB3	2:B:211:ALA:CA	2.09	0.79
2:B:178:ILE:O	2:B:181:TYR:N	2.16	0.79
3:M:65:TYR:CD1	3:M:86:PRO:CB	2.65	0.79
3:M:69:ILE:CG1	3:M:90:PHE:CE1	2.66	0.79
1:A:102:GLN:N	4:S:167:ILE:HG12	1.97	0.79
1:A:170:LEU:C	1:A:206:LYS:HD2	2.02	0.79
1:A:461:CYS:O	1:A:462:GLN:C	2.08	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:116:THR:HG22	2:B:150:LEU:CD2	2.12	0.79
2:B:353:GLN:HG2	3:M:48:ASP:N	1.97	0.79
2:B:490:ILE:CG2	2:B:515:PHE:CD2	2.66	0.79
3:M:65:TYR:CD1	3:M:86:PRO:CG	2.65	0.79
1:A:217:ALA:CB	4:S:142:ILE:CB	2.52	0.79
1:A:260:PHE:CE1	1:A:274:LEU:CD1	2.65	0.79
1:A:605:GLU:HG2	1:A:632:PHE:CG	2.17	0.79
1:A:636:TYR:H	2:B:554:LYS:HD3	1.47	0.79
2:B:42:ILE:HD13	2:B:65:ARG:HD3	1.61	0.79
1:A:137:ASN:C	1:A:139:ASP:H	1.80	0.79
1:A:220:SER:HB2	4:S:141:VAL:C	2.02	0.79
1:A:326:GLN:CA	1:A:331:ARG:NH2	2.42	0.79
1:A:393:LYS:NZ	1:A:397:ASP:OD2	2.16	0.79
2:B:42:ILE:O	2:B:43:ASN:HB2	1.83	0.79
2:B:216:LYS:HG3	2:B:251:LEU:HD12	1.63	0.79
2:B:236:ILE:HG22	2:B:240:LEU:CD1	2.12	0.79
2:B:367:SER:OG	2:B:401:THR:CG2	2.30	0.79
2:B:556:LEU:HB3	2:B:618:PHE:CD1	2.18	0.79
1:A:142:LYS:CA	4:S:159:ALA:CB	2.61	0.79
1:A:143:VAL:HG11	1:A:169:MET:HB3	1.65	0.79
1:A:189:PHE:CD2	1:A:225:LEU:HD11	2.17	0.79
1:A:244:LEU:CD1	1:A:256:LEU:CD1	2.59	0.79
1:A:254:ILE:CG2	1:A:293:GLU:HG2	2.10	0.79
2:B:136:CYS:C	2:B:172:GLU:CG	2.50	0.79
2:B:170:ARG:CA	2:B:199:LEU:CD2	2.48	0.79
3:M:262:THR:C	3:M:264:GLY:H	1.86	0.79
3:M:374:TYR:CB	3:M:417:TYR:HD2	1.95	0.79
1:A:557:LYS:HD3	2:B:605:PHE:CB	2.10	0.79
1:A:628:VAL:O	1:A:631:SER:OG	2.00	0.79
2:B:137:PHE:N	2:B:172:GLU:HG3	1.98	0.79
4:S:53:THR:HB	4:S:69:ASN:CB	2.13	0.79
1:A:371:ALA:O	1:A:374:LEU:N	2.16	0.79
1:A:461:CYS:SG	1:A:469:LEU:HB3	2.23	0.79
2:B:123:LEU:O	2:B:127:LEU:CD1	2.30	0.79
2:B:151:ALA:HB3	2:B:184:GLY:HA3	1.64	0.79
2:B:155:LEU:O	2:B:157:THR:N	2.16	0.79
2:B:309:LEU:CB	2:B:317:VAL:HG11	2.06	0.79
2:B:567:GLN:HG3	2:B:585:GLY:HA3	1.62	0.79
3:M:356:LEU:CD2	3:M:358:ILE:HG13	2.13	0.79
3:M:374:TYR:O	3:M:390:ILE:CD1	2.31	0.79
4:S:3:HIS:O	4:S:19:PHE:HA	1.82	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:6:LEU:HD13	4:S:32:LEU:HD13	1.65	0.79
1:A:145:ILE:CG1	4:S:156:LEU:HD22	2.13	0.78
2:B:25:VAL:CB	2:B:36:THR:OG1	2.31	0.78
2:B:37:TYR:O	2:B:40:GLN:N	2.16	0.78
2:B:165:PRO:HA	2:B:170:ARG:NH2	1.96	0.78
2:B:178:ILE:HB	2:B:214:ALA:HB1	1.65	0.78
4:S:9:ASN:HD21	4:S:13:GLN:HG3	1.46	0.78
1:A:74:LEU:CD2	1:A:87:CYS:SG	2.70	0.78
1:A:102:GLN:CG	4:S:166:LYS:N	1.94	0.78
1:A:440:ASN:OD1	1:A:442:SER:HB3	1.83	0.78
2:B:123:LEU:HB2	2:B:142:LEU:HD11	1.66	0.78
2:B:151:ALA:CB	2:B:184:GLY:HA3	2.13	0.78
2:B:219:TYR:CE1	2:B:226:LEU:HA	2.06	0.78
2:B:219:TYR:HB2	2:B:255:TYR:HD1	1.45	0.78
2:B:267:ASP:O	2:B:276:SER:CB	2.31	0.78
2:B:311:TYR:CZ	2:B:342:ALA:HB2	2.19	0.78
2:B:339:PHE:O	2:B:343:LEU:HB2	1.81	0.78
2:B:531:ARG:HB2	2:B:591:MET:SD	2.23	0.78
3:M:9:ASP:C	3:M:75:TRP:HD1	1.85	0.78
3:M:16:PHE:HA	3:M:118:TYR:CD1	2.18	0.78
3:M:320:ILE:HG21	3:M:439:TYR:CE2	2.18	0.78
3:M:342:LEU:HD12	3:M:411:LEU:CB	2.10	0.78
1:A:80:TYR:CB	1:A:82:PHE:CD2	2.65	0.78
1:A:519:LEU:O	1:A:520:GLY:C	2.13	0.78
2:B:120:ILE:CD1	2:B:142:LEU:HD23	2.13	0.78
2:B:267:ASP:CA	2:B:289:PRO:HG3	2.14	0.78
2:B:537:PHE:CE2	2:B:598:LEU:HB3	2.13	0.78
3:M:101:LEU:CB	3:M:106:LYS:HA	2.14	0.78
3:M:104:PHE:HZ	3:M:113:LYS:CE	1.54	0.78
1:A:67:LYS:CB	1:A:94:VAL:HG22	2.12	0.78
1:A:114:PHE:CD1	1:A:153:ILE:HG23	2.18	0.78
2:B:83:PHE:CE2	2:B:119:SER:CA	2.47	0.78
2:B:148:SER:O	2:B:183:ALA:HB3	1.70	0.78
2:B:501:THR:HA	2:B:508:ARG:HH22	1.47	0.78
3:M:95:THR:O	3:M:99:ILE:HD13	1.82	0.78
3:M:233:LEU:HD22	3:M:323:MET:O	1.82	0.78
3:M:338:PHE:HD2	3:M:415:ILE:HG13	1.48	0.78
3:M:362:PHE:O	3:M:363:ASN:O	2.00	0.78
3:M:449:VAL:HG12	3:M:452:ILE:CD1	1.96	0.78
4:S:8:PHE:HZ	4:S:84:TYR:HB3	1.45	0.78
1:A:241:TYR:C	1:A:242:GLU:O	2.02	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:629:LEU:HD21	2:B:610:ARG:HD3	1.65	0.78
1:A:637:GLU:CG	2:B:515:PHE:N	2.36	0.78
3:M:67:SER:OG	3:M:90:PHE:HB2	1.84	0.78
1:A:147:LEU:HD12	1:A:181:ALA:CA	2.10	0.78
1:A:384:LEU:HD13	1:A:441:TYR:CE2	2.19	0.78
2:B:291:TYR:CD2	2:B:294:VAL:CB	2.66	0.78
2:B:350:THR:HB	2:B:352:ASN:HD21	1.45	0.78
2:B:515:PHE:CE2	2:B:529:VAL:HG11	2.19	0.78
3:M:244:VAL:CB	3:M:472:TYR:CE2	2.66	0.78
3:M:354:ASP:HB2	3:M:440:ILE:HD12	1.63	0.78
1:A:141:VAL:CG1	4:S:159:ALA:N	2.47	0.78
2:B:20:ARG:NH1	2:B:35:TYR:CE1	2.51	0.78
2:B:396:ILE:CD1	2:B:432:ALA:HB2	2.14	0.78
2:B:409:LYS:O	2:B:413:LYS:HG3	1.83	0.78
2:B:566:ALA:C	2:B:574:ASN:HB3	1.86	0.78
3:M:9:ASP:O	3:M:75:TRP:CD1	2.37	0.78
3:M:228:LYS:NZ	3:M:326:HIS:HA	1.98	0.78
3:M:403:THR:HG23	3:M:407:THR:OG1	1.82	0.78
1:A:429:VAL:HG11	1:A:473:ILE:HD11	1.64	0.78
1:A:634:ASN:OD1	2:B:556:LEU:N	2.10	0.78
2:B:104:TYR:O	2:B:107:ARG:N	2.16	0.78
2:B:144:ASP:HA	2:B:179:LYS:CE	2.13	0.78
2:B:146:LYS:O	2:B:147:MET:SD	2.42	0.78
2:B:174:ALA:O	2:B:214:ALA:CB	2.32	0.78
2:B:219:TYR:CG	2:B:226:LEU:CB	2.48	0.78
2:B:336:ASN:ND2	2:B:338:LYS:HB2	1.98	0.78
2:B:374:PHE:CE2	2:B:398:ILE:HG21	2.18	0.78
2:B:437:SER:O	2:B:478:LEU:HD23	1.83	0.78
2:B:509:ALA:HB1	2:B:547:GLN:HG3	1.47	0.78
3:M:45:SER:CA	3:M:47:SER:N	2.47	0.78
3:M:360:LEU:CD2	3:M:362:PHE:CZ	2.66	0.78
4:S:8:PHE:CG	4:S:36:TYR:HE2	2.01	0.78
1:A:182:ILE:HG21	1:A:218:ALA:HA	1.66	0.78
1:A:244:LEU:HD21	1:A:277:LYS:O	1.83	0.78
1:A:333:ILE:O	1:A:337:LEU:HG	1.84	0.78
2:B:37:TYR:HE2	2:B:38:TYR:CE1	1.97	0.78
2:B:178:ILE:HG21	2:B:214:ALA:C	2.04	0.78
2:B:275:ARG:HG3	2:B:291:TYR:HD2	1.37	0.78
3:M:379:LEU:HD22	3:M:386:PHE:CB	2.13	0.78
4:S:34:GLN:CB	4:S:58:LEU:HD11	2.14	0.78
1:A:90:HIS:O	1:A:94:VAL:HG23	1.83	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:408:ILE:HG22	4:S:65:ASN:CA	2.13	0.78
1:A:631:SER:CB	2:B:557:SER:HB2	2.10	0.78
2:B:472:VAL:CG1	2:B:510:GLY:C	2.51	0.78
2:B:500:GLN:OE1	2:B:503:LEU:HD21	1.83	0.78
3:M:101:LEU:CA	3:M:109:LEU:CD1	2.55	0.78
2:B:178:ILE:CG1	2:B:214:ALA:CB	2.61	0.77
2:B:223:LEU:HD23	2:B:255:TYR:HE1	1.47	0.77
2:B:230:PHE:HD2	2:B:298:ASP:HB3	1.48	0.77
2:B:475:ILE:CG2	2:B:514:LEU:HD21	2.13	0.77
3:M:283:PHE:CZ	3:M:289:THR:CB	2.62	0.77
2:B:20:ARG:HG3	2:B:21:GLU:N	1.99	0.77
2:B:170:ARG:HA	2:B:199:LEU:HD23	1.64	0.77
2:B:252:LEU:HB2	2:B:302:PHE:CE2	2.19	0.77
2:B:278:PRO:CB	2:B:288:TYR:C	2.44	0.77
2:B:331:PRO:HB3	2:B:369:LEU:HD11	1.64	0.77
2:B:513:TRP:O	2:B:551:LEU:HD22	1.83	0.77
3:M:224:VAL:N	3:M:479:PHE:CD1	2.53	0.77
3:M:380:ARG:HH11	3:M:410:VAL:HG11	1.47	0.77
4:S:16:LEU:CD2	4:S:128:LEU:HD23	2.14	0.77
1:A:139:ASP:C	1:A:177:ILE:HD11	2.04	0.77
1:A:140:VAL:HG22	1:A:177:ILE:CG1	2.14	0.77
1:A:323:CYS:SG	1:A:334:SER:HB2	2.25	0.77
1:A:401:VAL:CG2	1:A:418:ILE:N	2.47	0.77
2:B:353:GLN:HE22	3:M:47:SER:HB3	1.42	0.77
2:B:380:LYS:NZ	3:M:236:LEU:CG	2.47	0.77
2:B:537:PHE:HE2	2:B:598:LEU:O	1.68	0.77
3:M:290:PHE:CZ	3:M:297:PHE:CE2	2.71	0.77
4:S:109:LEU:HD12	4:S:113:PHE:HD1	0.92	0.77
1:A:256:LEU:O	1:A:260:PHE:CD1	2.37	0.77
1:A:498:LEU:HB3	1:A:504:ILE:HG13	1.66	0.77
2:B:170:ARG:HH12	2:B:198:GLU:HG2	0.69	0.77
2:B:364:HIS:ND1	2:B:397:GLN:HB3	2.00	0.77
3:M:228:LYS:HZ2	3:M:326:HIS:HA	1.50	0.77
3:M:360:LEU:HD13	3:M:433:VAL:HB	1.66	0.77
4:S:53:THR:CB	4:S:69:ASN:CA	2.54	0.77
4:S:108:SER:OG	4:S:149:ILE:HG21	1.83	0.77
1:A:147:LEU:CD1	1:A:166:LEU:CD2	2.62	0.77
1:A:176:TYR:HB2	4:S:155:GLU:CG	2.09	0.77
1:A:275:LEU:CD2	1:A:311:THR:OG1	2.28	0.77
1:A:304:LEU:HD22	1:A:344:ILE:HG21	1.66	0.77
2:B:123:LEU:HD22	2:B:138:ALA:C	2.04	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:219:TYR:HB3	2:B:255:TYR:CD1	2.17	0.77
2:B:340:ILE:HG13	2:B:373:LEU:CD2	2.13	0.77
2:B:479:VAL:HG21	2:B:486:HIS:CD2	2.20	0.77
2:B:530:LEU:CD2	2:B:591:MET:HB3	2.14	0.77
2:B:562:ASN:O	2:B:580:TYR:O	2.03	0.77
3:M:101:LEU:CD1	3:M:106:LYS:HA	2.14	0.77
3:M:304:VAL:HG11	3:M:445:SER:HA	1.65	0.77
3:M:428:VAL:O	3:M:429:ASP:C	2.19	0.77
1:A:142:LYS:HA	4:S:159:ALA:CB	2.13	0.77
1:A:401:VAL:CB	1:A:419:ILE:HD12	2.14	0.77
2:B:63:MET:HE1	2:B:104:TYR:HB2	1.66	0.77
2:B:127:LEU:CD1	2:B:157:THR:CB	2.58	0.77
2:B:284:ASN:CG	2:B:285:GLU:HG3	2.03	0.77
2:B:437:SER:O	2:B:478:LEU:HD21	1.81	0.77
2:B:545:ARG:HB2	2:B:602:ASP:OD2	1.84	0.77
2:B:562:ASN:CB	2:B:580:TYR:CB	2.62	0.77
3:M:69:ILE:CG1	3:M:90:PHE:CZ	2.67	0.77
3:M:99:ILE:H	3:M:99:ILE:HD12	1.48	0.77
3:M:284:SER:O	3:M:285:PRO:C	2.09	0.77
3:M:302:TYR:CE2	3:M:445:SER:HB2	2.19	0.77
1:A:401:VAL:HB	1:A:419:ILE:CD1	2.14	0.77
1:A:460:LEU:O	1:A:463:ASP:N	2.16	0.77
2:B:2:VAL:HG11	2:B:6:HIS:CE1	2.17	0.77
2:B:106:LEU:HD13	2:B:144:ASP:CA	2.14	0.77
3:M:243:ILE:N	3:M:474:THR:CG2	2.35	0.77
3:M:256:VAL:HG22	3:M:452:ILE:CG2	2.15	0.77
3:M:257:ALA:HB1	3:M:287:ASN:HD21	1.50	0.77
4:S:75:ILE:HG22	4:S:86:THR:HG23	1.67	0.77
1:A:244:LEU:HG	1:A:277:LYS:HG3	1.65	0.77
1:A:384:LEU:HD13	1:A:435:ILE:HG22	1.67	0.77
2:B:139:LEU:HD23	2:B:173:VAL:N	2.00	0.77
2:B:513:TRP:HB2	2:B:551:LEU:HD11	1.65	0.77
2:B:563:PHE:CE1	2:B:588:ILE:HD12	2.19	0.77
3:M:235:LEU:CD1	3:M:306:LEU:HB3	2.14	0.77
3:M:348:LYS:HG3	3:M:405:THR:CG2	2.13	0.77
1:A:178:ARG:HB3	1:A:214:VAL:HG13	1.67	0.77
1:A:353:ASP:OD1	1:A:378:ILE:HD13	1.84	0.77
1:A:636:TYR:H	2:B:554:LYS:CD	1.97	0.77
2:B:191:GLU:C	2:B:193:LEU:N	2.37	0.77
2:B:292:GLU:HG3	2:B:296:ASP:HB2	1.46	0.77
3:M:223:HIS:CE1	3:M:476:THR:H	2.03	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:235:LEU:HD11	3:M:306:LEU:CB	2.14	0.77
3:M:340:LEU:HG	3:M:411:LEU:HB3	1.66	0.77
3:M:379:LEU:CD2	3:M:397:TRP:HE1	1.98	0.77
2:B:86:VAL:CG1	2:B:101:ILE:CG2	2.57	0.77
2:B:105:LEU:HB3	2:B:145:MET:HE1	1.65	0.77
2:B:124:GLN:HB2	2:B:153:ILE:HG23	1.67	0.77
2:B:424:PHE:HD2	2:B:428:VAL:HG11	1.48	0.77
3:M:4:SER:O	3:M:78:ALA:C	2.23	0.77
4:S:57:LEU:HA	4:S:60:SER:HB2	1.67	0.77
2:B:120:ILE:CD1	2:B:142:LEU:CD2	2.63	0.76
3:M:3:LEU:HB2	3:M:20:LEU:CD1	2.13	0.76
3:M:246:VAL:O	3:M:297:PHE:CE2	2.37	0.76
4:S:47:GLN:OE1	4:S:79:ASN:N	2.18	0.76
4:S:87:PHE:CE1	4:S:102:ILE:HG23	2.20	0.76
4:S:89:VAL:CG1	4:S:98:ILE:HG21	2.15	0.76
4:S:112:CYS:SG	4:S:153:VAL:HG21	2.25	0.76
1:A:79:MET:HG2	1:A:112:GLN:OE1	1.84	0.76
1:A:255:ARG:O	1:A:258:LYS:N	2.18	0.76
1:A:288:THR:OG1	1:A:291:ILE:CB	2.28	0.76
1:A:316:LEU:HD11	1:A:341:ILE:HG21	1.67	0.76
2:B:144:ASP:HA	2:B:179:LYS:CD	2.16	0.76
2:B:334:MET:HE2	2:B:339:PHE:CE1	2.20	0.76
3:M:215:TYR:CE1	3:M:468:LYS:HA	2.18	0.76
3:M:253:ASN:OD1	3:M:292:PRO:CG	2.33	0.76
1:A:252:ILE:HA	4:S:144:THR:HG1	1.49	0.76
2:B:90:ILE:HD11	2:B:102:HIS:CD2	2.21	0.76
2:B:184:GLY:O	2:B:188:TYR:HB2	1.84	0.76
2:B:197:LYS:C	2:B:199:LEU:N	2.36	0.76
2:B:227:HIS:C	2:B:229:HIS:N	2.21	0.76
2:B:243:TRP:CZ2	3:M:94:GLU:C	2.55	0.76
2:B:279:LEU:CG	2:B:288:TYR:CD1	2.64	0.76
2:B:526:CYS:N	2:B:527:PRO:CD	2.48	0.76
2:B:549:LEU:HD21	2:B:610:ARG:C	2.01	0.76
4:S:75:ILE:HG22	4:S:77:TYR:CE1	2.21	0.76
1:A:102:GLN:HE21	4:S:165:SER:N	1.83	0.76
1:A:594:PHE:CE2	2:B:474:VAL:HG22	2.20	0.76
2:B:29:LYS:HE2	2:B:30:LEU:CA	2.16	0.76
2:B:196:LEU:HB3	2:B:215:TYR:CZ	2.20	0.76
2:B:458:MET:SD	2:B:471:TYR:CB	2.72	0.76
4:S:53:THR:HG23	4:S:67:GLU:C	1.98	0.76
1:A:63:ASP:O	4:S:165:SER:OG	2.01	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:67:LYS:HB3	1:A:94:VAL:CG2	2.14	0.76
1:A:103:LYS:HB3	1:A:107:TYR:HE1	1.46	0.76
1:A:105:VAL:CA	4:S:167:ILE:HD13	2.16	0.76
1:A:183:THR:OG1	4:S:142:ILE:HD11	1.83	0.76
1:A:215:VAL:O	1:A:216:SER:C	2.18	0.76
2:B:5:ILE:HA	2:B:8:ILE:HD12	1.66	0.76
2:B:6:HIS:ND1	3:M:25:PRO:HB3	2.00	0.76
2:B:93:ASN:HA	2:B:134:LEU:CD1	2.15	0.76
2:B:136:CYS:HA	2:B:172:GLU:HB2	1.67	0.76
2:B:182:ARG:NH1	2:B:217:GLU:OE1	2.18	0.76
2:B:232:ARG:HG3	2:B:236:ILE:HG13	1.65	0.76
2:B:261:PRO:HG2	2:B:292:GLU:CA	2.15	0.76
2:B:334:MET:HA	2:B:334:MET:CE	2.14	0.76
4:S:17:VAL:HG13	4:S:19:PHE:CE1	2.21	0.76
4:S:53:THR:HG21	4:S:68:VAL:N	2.00	0.76
1:A:140:VAL:C	4:S:155:GLU:HB3	2.05	0.76
1:A:516:ILE:CD1	1:A:551:LEU:HD13	2.15	0.76
2:B:105:LEU:CB	2:B:145:MET:HE3	2.11	0.76
2:B:215:TYR:CE2	2:B:229:HIS:CB	2.69	0.76
3:M:348:LYS:O	3:M:405:THR:HG23	1.85	0.76
4:S:131:VAL:HG21	4:S:153:VAL:CG2	2.14	0.76
1:A:101:GLN:O	4:S:167:ILE:CG1	2.31	0.76
1:A:370:LYS:O	1:A:374:LEU:HD13	1.86	0.76
2:B:47:LEU:HB3	2:B:66:ILE:HG12	1.67	0.76
2:B:172:GLU:O	2:B:173:VAL:C	2.15	0.76
2:B:212:VAL:C	2:B:214:ALA:N	2.35	0.76
2:B:260:LEU:HB3	2:B:291:TYR:CZ	2.12	0.76
2:B:479:VAL:CG2	2:B:486:HIS:NE2	2.48	0.76
3:M:65:TYR:CD2	3:M:86:PRO:HA	2.20	0.76
3:M:373:ALA:HB3	3:M:418:GLU:O	1.86	0.76
3:M:443:SER:HB3	3:M:447:ILE:N	2.00	0.76
4:S:53:THR:CB	4:S:69:ASN:HB2	2.16	0.76
4:S:69:ASN:O	4:S:73:ILE:O	2.04	0.76
4:S:75:ILE:CG2	4:S:86:THR:CG2	2.64	0.76
1:A:99:LYS:HZ3	4:S:164:ASP:CB	1.97	0.76
1:A:606:PHE:CG	1:A:629:LEU:HG	2.20	0.76
2:B:83:PHE:O	2:B:87:VAL:HG23	1.85	0.76
2:B:178:ILE:HG23	2:B:217:GLU:CB	2.15	0.76
2:B:334:MET:HB2	2:B:369:LEU:HD23	1.67	0.76
3:M:214:LEU:O	3:M:467:TYR:HB3	1.85	0.76
3:M:273:HIS:CB	3:M:298:ARG:O	2.32	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:101:GLN:HG2	4:S:160:ALA:HB1	1.66	0.76
1:A:207:LEU:O	1:A:243:ILE:HD11	1.86	0.76
2:B:36:THR:HA	2:B:39:SER:OG	1.84	0.76
2:B:396:ILE:CG1	2:B:418:TYR:HE2	1.98	0.76
3:M:101:LEU:C	3:M:106:LYS:HA	2.06	0.76
3:M:223:HIS:CD2	3:M:478:ASN:HB2	2.20	0.76
3:M:224:VAL:O	3:M:479:PHE:CA	2.34	0.76
4:S:85:PHE:CE2	4:S:109:LEU:HD23	2.21	0.76
1:A:207:LEU:CD2	1:A:239:LEU:CB	2.58	0.76
1:A:244:LEU:HD11	1:A:281:LEU:HD11	1.68	0.76
1:A:623:MET:O	2:B:617:LEU:CD2	2.34	0.76
1:A:627:GLU:HG2	2:B:617:LEU:HA	1.67	0.76
2:B:549:LEU:HD21	2:B:611:ALA:H	0.94	0.76
2:B:559:ASP:O	2:B:562:ASN:CA	2.33	0.76
2:B:563:PHE:O	2:B:564:LYS:O	2.03	0.76
3:M:217:ASP:OD1	3:M:440:ILE:HG23	1.85	0.76
3:M:342:LEU:HD12	3:M:411:LEU:HB3	1.68	0.76
1:A:74:LEU:HD22	1:A:87:CYS:CB	2.14	0.75
1:A:101:GLN:N	4:S:162:SER:N	2.34	0.75
1:A:179:LYS:NZ	4:S:143:GLU:HB2	2.00	0.75
1:A:254:ILE:HG13	1:A:290:VAL:HG22	1.68	0.75
1:A:264:SER:HB3	1:A:271:ARG:HG3	1.68	0.75
1:A:421:PRO:CG	1:A:424:TYR:CE1	2.66	0.75
1:A:481:MET:HE2	1:A:518:CYS:HB3	1.68	0.75
1:A:595:GLU:HG3	2:B:469:ASP:HB3	0.75	0.75
2:B:67:ILE:HD13	2:B:103:LEU:CB	2.15	0.75
2:B:178:ILE:HD13	2:B:218:CYS:CB	2.15	0.75
2:B:433:VAL:CB	2:B:474:VAL:HG21	2.15	0.75
2:B:500:GLN:HB3	2:B:503:LEU:HG	1.68	0.75
2:B:589:SER:OG	2:B:618:PHE:CE2	2.39	0.75
3:M:60:LEU:HD22	3:M:62:VAL:HG22	1.67	0.75
3:M:99:ILE:HG12	3:M:128:CYS:SG	2.27	0.75
3:M:222:PHE:CB	3:M:479:PHE:HZ	1.93	0.75
3:M:224:VAL:C	3:M:479:PHE:HA	2.06	0.75
3:M:245:ASP:CB	3:M:472:TYR:HD1	1.95	0.75
3:M:306:LEU:HD22	3:M:317:MET:HE3	1.68	0.75
3:M:383:HIS:HB2	3:M:403:THR:OG1	1.85	0.75
1:A:103:LYS:CG	4:S:163:THR:CG2	2.57	0.75
1:A:461:CYS:CB	1:A:469:LEU:HD23	2.17	0.75
2:B:29:LYS:O	2:B:32:GLU:HG3	1.78	0.75
2:B:154:ILE:CD1	2:B:180:LEU:CB	2.64	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:178:ILE:CB	2:B:214:ALA:HB1	2.16	0.75
2:B:231:ARG:HG2	2:B:298:ASP:OD1	1.84	0.75
2:B:556:LEU:HD23	2:B:588:ILE:CD1	2.17	0.75
2:B:596:LEU:HD22	2:B:615:SER:OG	1.84	0.75
3:M:101:LEU:CG	3:M:106:LYS:O	2.35	0.75
3:M:104:PHE:HZ	3:M:113:LYS:HZ2	0.77	0.75
3:M:320:ILE:HG13	3:M:347:PHE:CD1	2.20	0.75
3:M:379:LEU:HD23	3:M:397:TRP:NE1	2.01	0.75
4:S:54:PRO:CB	4:S:57:LEU:CD1	2.64	0.75
1:A:166:LEU:HD12	1:A:185:LEU:CD2	2.04	0.75
1:A:178:ARG:CZ	1:A:209:ASP:CG	2.55	0.75
1:A:320:HIS:O	1:A:322:PHE:N	2.19	0.75
2:B:346:THR:O	2:B:350:THR:N	2.18	0.75
2:B:390:VAL:O	2:B:393:ILE:HB	1.86	0.75
2:B:563:PHE:O	2:B:566:ALA:CA	2.35	0.75
2:B:592:TYR:CE2	2:B:618:PHE:CD2	2.74	0.75
3:M:437:TYR:HB2	3:M:439:TYR:CZ	2.21	0.75
4:S:9:ASN:HD21	4:S:13:GLN:HG2	1.51	0.75
4:S:47:GLN:HE22	4:S:79:ASN:N	1.84	0.75
4:S:73:ILE:HG23	4:S:88:ILE:HG22	1.65	0.75
1:A:178:ARG:CZ	1:A:209:ASP:OD2	2.35	0.75
1:A:264:SER:CB	1:A:271:ARG:HG3	2.15	0.75
2:B:108:PHE:CD2	2:B:115:LEU:HB2	2.22	0.75
2:B:307:ASN:O	2:B:310:ILE:N	2.19	0.75
2:B:343:LEU:HD21	2:B:359:LEU:O	1.86	0.75
2:B:363:ILE:HG21	2:B:398:ILE:HD13	1.66	0.75
2:B:469:ASP:OD1	2:B:506:ASN:HB3	1.86	0.75
3:M:6:TYR:OH	3:M:17:GLN:NE2	2.18	0.75
3:M:16:PHE:HA	3:M:118:TYR:CE1	2.22	0.75
3:M:18:TYR:HE2	3:M:20:LEU:CD2	1.98	0.75
4:S:131:VAL:CG2	4:S:153:VAL:CG2	2.58	0.75
1:A:117:ASP:CG	1:A:120:ILE:HG12	2.07	0.75
1:A:260:PHE:CG	1:A:274:LEU:HG	2.22	0.75
2:B:106:LEU:HD11	2:B:144:ASP:CB	1.80	0.75
2:B:267:ASP:N	2:B:289:PRO:CG	2.49	0.75
3:M:374:TYR:CB	3:M:417:TYR:CD2	2.69	0.75
1:A:104:ARG:HG3	1:A:145:ILE:CD1	2.15	0.75
1:A:403:LEU:CD2	1:A:421:PRO:C	2.44	0.75
1:A:638:LEU:O	2:B:486:HIS:HE1	1.70	0.75
2:B:177:ILE:CG2	2:B:196:LEU:HD11	2.16	0.75
2:B:277:CYS:C	2:B:288:TYR:HB3	2.04	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:334:MET:CE	2:B:339:PHE:CE1	2.70	0.75
3:M:221:THR:HB	3:M:474:THR:C	2.06	0.75
3:M:290:PHE:CD1	3:M:297:PHE:CE1	2.75	0.75
3:M:350:VAL:HG22	3:M:442:GLN:CD	2.06	0.75
3:M:421:GLY:O	3:M:422:PRO:C	2.22	0.75
4:S:8:PHE:CB	4:S:36:TYR:CZ	2.62	0.75
1:A:322:PHE:CG	1:A:330:LEU:HD21	2.22	0.75
1:A:557:LYS:CD	2:B:605:PHE:HB3	2.12	0.75
2:B:118:LEU:O	2:B:121:ASN:N	2.20	0.75
2:B:286:ILE:CG2	2:B:288:TYR:OH	2.34	0.75
2:B:473:ASN:O	2:B:476:ARG:HB3	1.86	0.75
2:B:550:VAL:CG2	2:B:610:ARG:HD3	2.13	0.75
3:M:121:ILE:CG2	3:M:125:PHE:CE1	2.70	0.75
3:M:219:LEU:O	3:M:474:THR:HG23	1.86	0.75
3:M:222:PHE:CD1	3:M:439:TYR:HE1	2.04	0.75
3:M:241:HIS:HB2	3:M:476:THR:HG23	1.67	0.75
3:M:454:ILE:CG2	3:M:464:THR:CG2	2.53	0.75
4:S:16:LEU:HD23	4:S:128:LEU:CD2	2.16	0.75
1:A:304:LEU:HD12	1:A:312:ALA:HB2	1.68	0.75
2:B:602:ASP:OD1	2:B:603:ASP:N	2.19	0.75
3:M:258:VAL:HG13	3:M:452:ILE:CG1	2.16	0.75
3:M:290:PHE:CZ	3:M:297:PHE:CZ	2.74	0.75
3:M:437:TYR:N	3:M:437:TYR:CD1	2.53	0.75
1:A:581:LEU:HD23	1:A:607:LEU:HD21	1.69	0.75
1:A:633:PHE:HD2	2:B:551:LEU:CA	1.71	0.75
1:A:636:TYR:CG	2:B:554:LYS:NZ	2.55	0.75
2:B:60:ARG:HD2	2:B:96:LYS:CG	2.16	0.75
2:B:171:GLY:HA3	2:B:207:VAL:CA	2.17	0.75
2:B:189:HIS:NE2	2:B:225:LEU:HD23	2.00	0.75
2:B:200:MET:CG	2:B:232:ARG:HB3	2.13	0.75
2:B:215:TYR:CB	2:B:226:LEU:CD1	2.63	0.75
2:B:310:ILE:HG12	2:B:318:ILE:HA	1.69	0.75
2:B:319:LEU:HD12	2:B:358:MET:CG	1.79	0.75
2:B:328:LEU:HB2	2:B:333:GLN:HE22	1.52	0.75
2:B:346:THR:HG23	2:B:350:THR:HG23	1.66	0.75
2:B:444:THR:O	2:B:445:SER:C	2.15	0.75
2:B:597:TYR:O	2:B:601:TYR:CE1	2.39	0.75
3:M:52:ASP:HA	3:M:67:SER:CA	2.17	0.75
1:A:129:LYS:HA	1:A:165:ASP:OD2	1.86	0.74
2:B:219:TYR:CE1	2:B:226:LEU:HD12	2.19	0.74
2:B:225:LEU:HD13	2:B:283:TYR:CZ	2.15	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:596:LEU:HD13	2:B:611:ALA:O	1.87	0.74
3:M:67:SER:CB	3:M:90:PHE:CD1	2.61	0.74
3:M:101:LEU:CD1	3:M:109:LEU:CD1	2.65	0.74
3:M:342:LEU:HD11	3:M:411:LEU:HD23	1.65	0.74
1:A:291:ILE:HG21	1:A:318:ARG:HB3	1.63	0.74
1:A:372:ILE:CG2	1:A:427:LYS:HE3	2.17	0.74
1:A:556:VAL:HG11	2:B:610:ARG:NH2	2.02	0.74
2:B:219:TYR:CG	2:B:223:LEU:HD23	2.20	0.74
2:B:299:LEU:O	2:B:302:PHE:HB3	1.87	0.74
2:B:337:THR:HA	2:B:373:LEU:HD22	1.63	0.74
2:B:500:GLN:CB	2:B:503:LEU:HG	2.17	0.74
4:S:54:PRO:HB2	4:S:57:LEU:HD11	1.70	0.74
1:A:99:LYS:CD	4:S:164:ASP:H	1.94	0.74
1:A:186:PHE:CD1	1:A:224:GLU:HB3	2.22	0.74
1:A:609:LEU:CD2	1:A:628:VAL:HG11	2.17	0.74
2:B:37:TYR:CE2	2:B:42:ILE:HA	2.22	0.74
2:B:37:TYR:O	2:B:39:SER:N	2.20	0.74
2:B:63:MET:HE2	2:B:104:TYR:HB2	1.69	0.74
2:B:278:PRO:N	2:B:292:GLU:CD	2.37	0.74
2:B:546:CYS:CB	2:B:607:ILE:CD1	2.42	0.74
3:M:18:TYR:HD1	3:M:122:SER:CA	1.80	0.74
3:M:449:VAL:HG11	3:M:452:ILE:HD11	0.75	0.74
1:A:163:ALA:CA	1:A:199:ASN:HD21	2.00	0.74
1:A:287:ALA:HB3	1:A:288:THR:C	2.06	0.74
2:B:38:TYR:CD2	2:B:65:ARG:CD	2.33	0.74
2:B:70:MET:HB2	2:B:104:TYR:CE1	2.22	0.74
2:B:143:SER:CA	2:B:179:LYS:HB2	2.16	0.74
2:B:599:ALA:C	2:B:601:TYR:N	2.39	0.74
3:M:443:SER:CA	3:M:447:ILE:HG13	2.18	0.74
1:A:397:ASP:O	1:A:418:ILE:CG1	2.35	0.74
2:B:310:ILE:O	2:B:311:TYR:C	2.13	0.74
4:S:17:VAL:HG22	4:S:19:PHE:CE2	2.21	0.74
4:S:85:PHE:CD1	4:S:106:VAL:HG22	2.21	0.74
1:A:183:THR:OG1	4:S:142:ILE:HD13	1.86	0.74
1:A:220:SER:HB3	4:S:142:ILE:HG22	0.75	0.74
2:B:378:THR:CG2	2:B:379:LYS:H	1.99	0.74
2:B:390:VAL:HA	2:B:393:ILE:HD12	1.69	0.74
2:B:602:ASP:O	2:B:608:ARG:CZ	2.36	0.74
3:M:67:SER:OG	3:M:90:PHE:CB	2.35	0.74
3:M:95:THR:O	3:M:99:ILE:HD12	1.87	0.74
1:A:80:TYR:HB2	1:A:82:PHE:CE2	2.22	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:110:ALA:O	1:A:111:SER:C	2.14	0.74
1:A:142:LYS:CA	4:S:159:ALA:HB1	2.17	0.74
1:A:179:LYS:NZ	4:S:149:ILE:HG12	2.01	0.74
1:A:212:ILE:CG2	1:A:247:ILE:HD13	2.17	0.74
1:A:398:GLU:O	1:A:420:ILE:CG1	2.34	0.74
2:B:178:ILE:HG13	2:B:214:ALA:CB	2.18	0.74
2:B:197:LYS:C	2:B:199:LEU:H	1.91	0.74
3:M:222:PHE:CE1	3:M:240:ILE:CG2	2.70	0.74
3:M:360:LEU:HD21	3:M:362:PHE:CE2	2.23	0.74
1:A:304:LEU:HA	1:A:308:ASP:HB2	1.67	0.74
1:A:540:ILE:CG1	1:A:551:LEU:HD23	2.16	0.74
2:B:20:ARG:CG	2:B:21:GLU:N	2.49	0.74
2:B:225:LEU:CD1	2:B:283:TYR:HE1	1.83	0.74
1:A:105:VAL:CG2	4:S:167:ILE:CB	2.66	0.74
1:A:175:PRO:HB3	1:A:211:ASP:CG	2.07	0.74
1:A:506:LYS:HE2	3:M:58:ARG:CA	2.18	0.74
1:A:634:ASN:ND2	2:B:554:LYS:HB3	1.79	0.74
2:B:175:LEU:CD2	2:B:210:CYS:CA	2.65	0.74
2:B:277:CYS:CB	2:B:292:GLU:HG3	2.16	0.74
3:M:17:GLN:O	3:M:118:TYR:OH	2.02	0.74
3:M:241:HIS:C	3:M:474:THR:HB	2.06	0.74
1:A:67:LYS:HB3	4:S:166:LYS:HA	1.70	0.74
1:A:67:LYS:CG	4:S:165:SER:HG	2.01	0.74
1:A:391:LEU:O	1:A:392:MET:C	2.17	0.74
1:A:436:CYS:HB2	1:A:450:TYR:CE1	2.23	0.74
1:A:606:PHE:CZ	2:B:550:VAL:HG11	2.21	0.74
1:A:624:LEU:O	2:B:617:LEU:HD11	1.88	0.74
2:B:490:ILE:HG23	2:B:515:PHE:CD2	2.23	0.74
3:M:5:PHE:CD2	3:M:78:ALA:HB3	2.16	0.74
3:M:64:LYS:HG2	3:M:79:SER:C	2.07	0.74
3:M:101:LEU:CD1	3:M:106:LYS:CA	2.65	0.74
1:A:213:SER:H	4:S:148:ARG:HD3	1.52	0.73
1:A:383:ASN:O	1:A:387:ILE:HG12	1.88	0.73
2:B:90:ILE:HG12	2:B:98:LYS:HE2	1.69	0.73
2:B:139:LEU:HD21	2:B:173:VAL:C	2.08	0.73
2:B:162:VAL:HG11	2:B:195:ILE:HA	1.67	0.73
2:B:171:GLY:HA3	2:B:207:VAL:CB	2.18	0.73
2:B:199:LEU:C	2:B:201:ALA:H	1.89	0.73
2:B:497:LEU:CD2	2:B:533:LEU:HD22	2.18	0.73
3:M:10:THR:HG23	3:M:11:LYS:HG3	1.70	0.73
3:M:94:GLU:O	3:M:97:ASP:HB2	1.88	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:132:LEU:CD1	1:A:165:ASP:CB	2.65	0.73
1:A:384:LEU:CD1	1:A:441:TYR:CE2	2.71	0.73
1:A:450:TYR:O	1:A:454:ILE:HG12	1.88	0.73
1:A:513:ARG:CD	1:A:550:VAL:CG2	2.53	0.73
1:A:633:PHE:CZ	2:B:513:TRP:CE3	2.70	0.73
2:B:114:ASN:O	2:B:117:LEU:HB2	1.87	0.73
2:B:132:SER:HB2	2:B:166:SER:HB3	1.70	0.73
2:B:430:ILE:HG23	2:B:470:ALA:CB	2.18	0.73
2:B:478:LEU:O	2:B:480:GLN:N	2.21	0.73
2:B:553:ALA:CB	2:B:614:ILE:CG2	2.66	0.73
2:B:83:PHE:HE2	2:B:119:SER:HA	1.53	0.73
2:B:141:ALA:O	2:B:143:SER:N	2.21	0.73
2:B:175:LEU:CG	2:B:210:CYS:CB	2.63	0.73
3:M:214:LEU:C	3:M:214:LEU:HD23	2.08	0.73
3:M:217:ASP:H	3:M:470:ALA:CB	1.93	0.73
3:M:222:PHE:CG	3:M:439:TYR:CE1	2.76	0.73
4:S:109:LEU:CD1	4:S:113:PHE:HD1	1.70	0.73
1:A:378:ILE:O	1:A:378:ILE:HG13	1.87	0.73
2:B:479:VAL:HG22	2:B:486:HIS:CG	2.22	0.73
2:B:513:TRP:CB	2:B:551:LEU:HD11	2.18	0.73
3:M:45:SER:CB	3:M:51:LEU:CD1	2.62	0.73
4:S:4:ALA:CA	4:S:18:LYS:O	2.35	0.73
1:A:186:PHE:HE2	1:A:187:LYS:HD3	1.53	0.73
1:A:254:ILE:HG23	1:A:293:GLU:CG	2.16	0.73
2:B:108:PHE:HE2	2:B:115:LEU:HB2	1.50	0.73
3:M:247:ARG:H	3:M:470:ALA:HB2	1.53	0.73
3:M:443:SER:CB	3:M:447:ILE:N	2.51	0.73
1:A:102:GLN:NE2	4:S:165:SER:N	2.37	0.73
1:A:233:PHE:O	1:A:234:ILE:C	2.05	0.73
1:A:319:LEU:O	1:A:320:HIS:O	2.07	0.73
1:A:322:PHE:CD2	1:A:330:LEU:CD2	2.71	0.73
1:A:516:ILE:HG12	1:A:551:LEU:HD13	1.68	0.73
1:A:633:PHE:HE1	2:B:513:TRP:CZ3	1.88	0.73
2:B:212:VAL:HG21	2:B:248:LEU:CD2	2.14	0.73
2:B:343:LEU:O	2:B:347:VAL:HG23	1.88	0.73
2:B:438:ARG:HA	2:B:441:GLN:HE21	1.52	0.73
2:B:476:ARG:HA	2:B:514:LEU:HD13	1.69	0.73
2:B:567:GLN:N	2:B:574:ASN:ND2	2.33	0.73
3:M:101:LEU:HD12	3:M:109:LEU:CD1	2.18	0.73
3:M:437:TYR:CD1	3:M:479:PHE:CE1	2.76	0.73
4:S:43:ASN:O	4:S:44:SER:C	2.20	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:211:ASP:CG	4:S:148:ARG:CZ	2.57	0.73
2:B:219:TYR:C	2:B:223:LEU:CD2	2.52	0.73
2:B:567:GLN:C	2:B:569:THR:OG1	2.26	0.73
3:M:96:ILE:HG21	3:M:125:PHE:CZ	2.24	0.73
3:M:262:THR:O	3:M:264:GLY:N	2.20	0.73
3:M:338:PHE:HE2	3:M:415:ILE:CG1	2.02	0.73
4:S:17:VAL:CG1	4:S:19:PHE:CE1	2.71	0.73
1:A:630:PRO:CB	2:B:614:ILE:HG23	2.19	0.73
2:B:208:ILE:HD13	2:B:236:ILE:HG21	0.73	0.73
2:B:374:PHE:CZ	2:B:398:ILE:HG21	2.24	0.73
2:B:508:ARG:HD2	2:B:544:THR:HG21	1.70	0.73
4:S:146:VAL:O	4:S:150:VAL:HG23	1.88	0.73
1:A:185:LEU:HD12	1:A:203:PHE:CZ	2.23	0.73
1:A:495:ILE:HD12	1:A:515:CYS:SG	2.28	0.73
2:B:237:ILE:HG12	2:B:309:LEU:HD21	1.70	0.73
2:B:374:PHE:CE1	2:B:381:PHE:CD1	2.77	0.73
2:B:403:ILE:HD11	2:B:442:LEU:HD12	1.70	0.73
2:B:418:TYR:CD1	2:B:424:PHE:CD1	2.75	0.73
2:B:508:ARG:CD	2:B:544:THR:HG21	2.18	0.73
2:B:515:PHE:CD2	2:B:529:VAL:HG11	2.23	0.73
4:S:109:LEU:HD12	4:S:113:PHE:CE1	1.91	0.73
1:A:609:LEU:HG	1:A:628:VAL:CG1	2.19	0.73
2:B:154:ILE:HG23	2:B:176:ALA:HB1	1.71	0.73
2:B:174:ALA:O	2:B:214:ALA:HB1	1.88	0.73
2:B:177:ILE:HG21	2:B:196:LEU:CD1	2.18	0.73
2:B:275:ARG:CB	2:B:294:VAL:HG13	1.82	0.73
2:B:393:ILE:CG2	2:B:431:MET:HB2	2.18	0.73
3:M:65:TYR:CE2	3:M:86:PRO:CB	2.68	0.73
3:M:290:PHE:HZ	3:M:297:PHE:CD2	2.07	0.73
3:M:343:ASN:HD22	3:M:343:ASN:N	1.87	0.73
4:S:10:LYS:HB3	4:S:82:THR:O	1.89	0.73
1:A:100:LEU:N	4:S:162:SER:CA	2.48	0.72
1:A:252:ILE:HG12	4:S:145:ASN:N	2.04	0.72
2:B:154:ILE:HD12	2:B:180:LEU:CB	2.18	0.72
2:B:271:GLU:O	2:B:272:GLY:C	2.20	0.72
3:M:126:ASN:O	3:M:130:GLU:HG2	1.88	0.72
3:M:257:ALA:CB	3:M:455:VAL:HG21	2.19	0.72
4:S:50:PHE:CB	4:S:76:ILE:HD13	2.19	0.72
4:S:53:THR:CG2	4:S:68:VAL:CA	2.63	0.72
1:A:223:CYS:O	1:A:226:SER:OG	2.03	0.72
1:A:263:LEU:O	1:A:266:VAL:O	2.07	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:408:ILE:HD12	1:A:410:TYR:HE1	1.51	0.72
1:A:539:ASN:O	1:A:540:ILE:C	2.23	0.72
1:A:557:LYS:CE	2:B:606:ASP:CA	2.57	0.72
1:A:633:PHE:HB3	2:B:550:VAL:O	1.88	0.72
3:M:7:ILE:HG23	3:M:75:TRP:O	1.89	0.72
3:M:316:ARG:O	3:M:318:ASN:N	2.21	0.72
1:A:147:LEU:HD21	1:A:166:LEU:HD23	1.66	0.72
1:A:402:ILE:O	1:A:402:ILE:CG2	2.36	0.72
1:A:636:TYR:N	2:B:554:LYS:HD3	1.95	0.72
2:B:63:MET:CG	2:B:100:LEU:CB	2.67	0.72
2:B:267:ASP:C	2:B:276:SER:OG	2.27	0.72
2:B:352:ASN:H	2:B:352:ASN:HD22	1.37	0.72
2:B:553:ALA:CA	2:B:614:ILE:CG2	2.66	0.72
2:B:159:LYS:HD2	2:B:191:GLU:CG	2.18	0.72
2:B:166:SER:O	2:B:170:ARG:HG3	1.89	0.72
2:B:212:VAL:C	2:B:214:ALA:H	1.93	0.72
2:B:490:ILE:O	2:B:515:PHE:HZ	1.71	0.72
4:S:135:ILE:CG2	4:S:141:VAL:HG22	2.20	0.72
1:A:461:CYS:HB2	1:A:469:LEU:HD23	1.71	0.72
1:A:599:ARG:HD2	2:B:547:GLN:NE2	2.04	0.72
2:B:64:LYS:HA	2:B:100:LEU:HD22	1.71	0.72
2:B:106:LEU:CD1	2:B:144:ASP:CA	2.68	0.72
2:B:162:VAL:HG21	2:B:195:ILE:CA	2.20	0.72
2:B:252:LEU:CB	2:B:302:PHE:CE1	2.73	0.72
2:B:278:PRO:N	2:B:292:GLU:OE1	2.22	0.72
2:B:368:ILE:CD1	2:B:401:THR:HG22	2.19	0.72
2:B:416:LYS:HD2	2:B:453:TRP:CE2	2.25	0.72
3:M:353:VAL:CG2	3:M:438:SER:O	2.37	0.72
3:M:379:LEU:HD23	3:M:386:PHE:CB	2.15	0.72
1:A:564:ASN:OD1	1:A:622:PRO:CG	2.38	0.72
1:A:606:PHE:CZ	2:B:550:VAL:CG1	2.72	0.72
1:A:637:GLU:HB2	2:B:551:LEU:HD22	1.69	0.72
2:B:277:CYS:O	2:B:288:TYR:CA	2.37	0.72
2:B:398:ILE:O	2:B:401:THR:N	2.18	0.72
2:B:472:VAL:HG11	2:B:510:GLY:HA3	0.73	0.72
4:S:44:SER:O	4:S:47:GLN:HB2	1.89	0.72
4:S:108:SER:HB3	4:S:149:ILE:HG22	1.70	0.72
1:A:63:ASP:O	1:A:67:LYS:HG3	1.90	0.72
1:A:94:VAL:HG13	1:A:102:GLN:HB3	1.64	0.72
1:A:223:CYS:CB	1:A:259:LEU:HG	2.17	0.72
1:A:291:ILE:HG13	1:A:318:ARG:HG2	1.70	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:133:GLU:HA	2:B:168:MET:SD	2.28	0.72
2:B:170:ARG:CZ	2:B:198:GLU:HG2	2.20	0.72
2:B:554:LYS:O	2:B:557:SER:N	2.21	0.72
3:M:6:TYR:CD2	3:M:14:LEU:HD11	2.24	0.72
3:M:225:VAL:HA	3:M:480:GLN:O	1.90	0.72
3:M:243:ILE:HG22	3:M:472:TYR:HB3	1.72	0.72
1:A:100:LEU:O	4:S:160:ALA:CA	1.85	0.72
1:A:350:SER:O	1:A:351:ARG:C	2.26	0.72
1:A:523:SER:OG	1:A:562:TRP:NE1	1.85	0.72
1:A:557:LYS:HG3	2:B:606:ASP:HB2	1.71	0.72
2:B:20:ARG:HD2	2:B:21:GLU:CA	2.18	0.72
2:B:98:LYS:HZ1	2:B:134:LEU:HB3	0.70	0.72
2:B:284:ASN:CB	2:B:285:GLU:HG3	2.20	0.72
2:B:505:ASP:O	2:B:506:ASN:C	2.17	0.72
2:B:564:LYS:CD	2:B:621:GLY:O	2.37	0.72
3:M:250:LEU:HD13	3:M:254:PRO:HG2	1.71	0.72
3:M:381:ASN:OD1	3:M:382:THR:N	2.23	0.72
4:S:75:ILE:CG2	4:S:77:TYR:CE1	2.72	0.72
1:A:105:VAL:CG2	4:S:167:ILE:CA	2.67	0.72
1:A:166:LEU:HD13	1:A:185:LEU:HG	1.70	0.72
1:A:213:SER:H	4:S:148:ARG:CD	1.90	0.72
1:A:226:SER:O	1:A:230:PRO:HG3	1.89	0.72
1:A:295:VAL:CG2	1:A:315:CYS:HB3	2.14	0.72
2:B:51:LEU:HD23	2:B:59:VAL:HG13	1.69	0.72
2:B:158:VAL:HG12	2:B:177:ILE:HD11	1.67	0.72
1:A:121:LEU:CD1	1:A:155:THR:HG23	2.20	0.72
1:A:214:VAL:HG22	4:S:148:ARG:NH2	2.03	0.72
1:A:252:ILE:HG12	4:S:144:THR:OG1	1.89	0.72
2:B:177:ILE:HD11	2:B:195:ILE:HG21	1.72	0.72
2:B:227:HIS:CG	2:B:292:GLU:OE2	2.43	0.72
2:B:493:LEU:HG	2:B:511:ILE:CG2	2.18	0.72
3:M:290:PHE:HE1	3:M:297:PHE:CD1	2.05	0.72
1:A:441:TYR:HB3	1:A:444:VAL:CG2	2.20	0.71
2:B:63:MET:HG3	2:B:100:LEU:CB	2.19	0.71
2:B:363:ILE:CG2	2:B:398:ILE:HG12	2.19	0.71
2:B:374:PHE:CE2	2:B:402:LEU:HD11	2.25	0.71
3:M:96:ILE:O	3:M:100:LEU:HD12	1.90	0.71
3:M:219:LEU:HG	3:M:440:ILE:HG12	1.72	0.71
4:S:55:PRO:C	4:S:57:LEU:H	1.91	0.71
1:A:95:MET:C	4:S:166:LYS:NZ	2.43	0.71
1:A:149:GLY:O	1:A:152:THR:N	2.15	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:204:VAL:O	1:A:205:SER:C	2.25	0.71
1:A:268:PRO:HG3	1:A:271:ARG:HH21	1.55	0.71
1:A:554:ALA:O	1:A:557:LYS:N	2.23	0.71
1:A:603:VAL:HG22	2:B:610:ARG:HH21	1.55	0.71
2:B:578:PRO:CB	2:B:579:PRO:HD2	2.18	0.71
3:M:243:ILE:O	3:M:472:TYR:CG	2.42	0.71
3:M:258:VAL:HG13	3:M:449:VAL:HG11	1.71	0.71
4:S:14:PRO:O	4:S:15:ARG:HD3	1.90	0.71
4:S:57:LEU:CB	4:S:67:GLU:O	2.38	0.71
1:A:80:TYR:CG	1:A:82:PHE:CE2	2.78	0.71
1:A:170:LEU:CD1	1:A:206:LYS:HG3	2.21	0.71
1:A:212:ILE:CB	1:A:247:ILE:CD1	2.68	0.71
1:A:353:ASP:OD1	1:A:378:ILE:CD1	2.39	0.71
1:A:603:VAL:O	1:A:606:PHE:HB2	1.89	0.71
2:B:403:ILE:HB	2:B:408:VAL:HG22	1.72	0.71
2:B:545:ARG:CD	2:B:602:ASP:CB	2.65	0.71
2:B:546:CYS:HA	2:B:607:ILE:CB	2.20	0.71
2:B:550:VAL:O	2:B:553:ALA:HB3	1.89	0.71
3:M:225:VAL:HG22	3:M:480:GLN:HB3	1.71	0.71
1:A:100:LEU:HD22	1:A:138:ASN:HB3	1.72	0.71
1:A:101:GLN:N	4:S:163:THR:H	1.88	0.71
1:A:103:LYS:CA	4:S:163:THR:HG21	2.20	0.71
1:A:121:LEU:HD11	1:A:155:THR:HG23	1.73	0.71
2:B:154:ILE:HD12	2:B:180:LEU:HB2	1.70	0.71
2:B:215:TYR:CE2	2:B:229:HIS:HB3	2.25	0.71
1:A:141:VAL:CB	4:S:159:ALA:CB	2.67	0.71
1:A:189:PHE:HB2	1:A:225:LEU:HD21	1.72	0.71
1:A:275:LEU:O	1:A:277:LYS:N	2.23	0.71
1:A:585:PHE:CE2	1:A:603:VAL:HG12	2.25	0.71
2:B:2:VAL:O	2:B:6:HIS:CD2	2.43	0.71
2:B:93:ASN:CA	2:B:134:LEU:HD11	2.21	0.71
2:B:431:MET:O	2:B:433:VAL:N	2.23	0.71
3:M:18:TYR:HE2	3:M:20:LEU:HD23	1.54	0.71
3:M:99:ILE:HD12	3:M:99:ILE:N	2.04	0.71
3:M:220:GLU:N	3:M:439:TYR:O	2.23	0.71
3:M:287:ASN:OD1	3:M:288:ILE:N	2.22	0.71
3:M:341:SER:HG	3:M:343:ASN:HD21	1.38	0.71
3:M:347:PHE:CZ	3:M:350:VAL:HG11	2.26	0.71
3:M:374:TYR:HB3	3:M:417:TYR:CD2	2.23	0.71
3:M:376:ILE:HD12	3:M:415:ILE:HG12	1.71	0.71
4:S:53:THR:H	4:S:69:ASN:CB	2.04	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:135:ILE:O	4:S:141:VAL:CG2	2.38	0.71
1:A:179:LYS:CD	4:S:143:GLU:CB	2.68	0.71
1:A:212:ILE:HB	1:A:247:ILE:HD12	1.73	0.71
1:A:329:ASN:OD1	4:S:50:PHE:HZ	1.72	0.71
1:A:626:SER:C	2:B:617:LEU:HG	2.10	0.71
2:B:108:PHE:CD2	2:B:115:LEU:CB	2.73	0.71
2:B:120:ILE:CG1	2:B:150:LEU:HD22	2.21	0.71
2:B:231:ARG:NH2	2:B:297:PRO:HD2	2.04	0.71
2:B:247:TYR:HE2	3:M:91:THR:CG2	1.88	0.71
2:B:337:THR:HA	2:B:373:LEU:HD21	0.75	0.71
2:B:403:ILE:HG22	2:B:411:ILE:HD12	1.71	0.71
3:M:52:ASP:HA	3:M:68:VAL:N	2.04	0.71
3:M:101:LEU:CB	3:M:109:LEU:HD13	2.20	0.71
3:M:258:VAL:HG22	3:M:452:ILE:CG1	2.20	0.71
3:M:290:PHE:CE2	3:M:297:PHE:CE2	2.77	0.71
3:M:343:ASN:HA	3:M:408:VAL:CG1	2.11	0.71
3:M:405:THR:HG22	3:M:406:GLY:N	2.04	0.71
3:M:448:TYR:HE2	3:M:450:GLU:OE1	1.74	0.71
1:A:266:VAL:O	1:A:267:GLU:HB3	1.90	0.71
1:A:332:TYR:HE1	1:A:366:SER:HB2	1.56	0.71
1:A:436:CYS:CB	1:A:450:TYR:CE1	2.73	0.71
2:B:20:ARG:NH1	2:B:35:TYR:CZ	2.48	0.71
2:B:120:ILE:HG13	2:B:150:LEU:HD22	1.73	0.71
2:B:169:VAL:O	2:B:173:VAL:HG23	1.91	0.71
2:B:278:PRO:HB2	2:B:288:TYR:O	1.87	0.71
2:B:396:ILE:HD13	2:B:418:TYR:OH	1.89	0.71
3:M:81:SER:O	3:M:82:LYS:CB	2.38	0.71
3:M:256:VAL:CG2	3:M:452:ILE:HG22	2.20	0.71
4:S:98:ILE:O	4:S:102:ILE:HG13	1.91	0.71
1:A:638:LEU:HD11	2:B:526:CYS:SG	2.31	0.71
2:B:129:ASP:O	2:B:135:ARG:HD3	1.91	0.71
2:B:137:PHE:O	2:B:140:SER:HB3	1.91	0.71
2:B:267:ASP:HB3	2:B:289:PRO:CG	2.21	0.71
2:B:563:PHE:CA	2:B:566:ALA:HB3	2.19	0.71
3:M:65:TYR:CE1	3:M:86:PRO:CG	2.74	0.71
3:M:293:PRO:HB2	3:M:294:ASP:O	1.91	0.71
4:S:105:PHE:CZ	4:S:128:LEU:CD1	2.74	0.71
1:A:147:LEU:HB3	1:A:184:ALA:HB1	1.70	0.71
1:A:400:VAL:O	1:A:403:LEU:HB2	1.90	0.71
1:A:520:GLY:HA2	1:A:558:VAL:HG22	1.67	0.71
1:A:594:PHE:CD2	2:B:474:VAL:HG22	2.25	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:169:VAL:HG12	2:B:173:VAL:CG2	2.20	0.71
2:B:303:LEU:HD23	2:B:325:LEU:HD23	1.73	0.71
2:B:549:LEU:HG	2:B:614:ILE:CD1	2.20	0.71
3:M:340:LEU:C	3:M:411:LEU:HB3	2.12	0.71
4:S:7:ILE:HD12	4:S:16:LEU:HD23	1.71	0.71
1:A:179:LYS:O	4:S:142:ILE:HD11	1.91	0.71
1:A:244:LEU:O	1:A:245:VAL:C	2.28	0.71
2:B:6:HIS:CG	3:M:25:PRO:HB3	2.26	0.71
2:B:47:LEU:CD2	2:B:66:ILE:HG12	1.93	0.71
2:B:50:LEU:HD23	2:B:62:ALA:HB2	1.73	0.71
2:B:537:PHE:CE2	2:B:598:LEU:C	2.63	0.71
3:M:341:SER:HA	3:M:411:LEU:H	1.56	0.71
3:M:360:LEU:HD23	3:M:362:PHE:HZ	1.55	0.71
3:M:372:ILE:CD1	3:M:428:VAL:HG22	2.21	0.71
2:B:153:ILE:C	2:B:155:LEU:N	2.44	0.70
2:B:252:LEU:HB3	2:B:302:PHE:CE2	2.25	0.70
2:B:275:ARG:HB3	2:B:291:TYR:CD2	2.25	0.70
2:B:368:ILE:HD13	2:B:401:THR:HG22	1.72	0.70
2:B:378:THR:CG2	2:B:379:LYS:N	2.54	0.70
2:B:418:TYR:CD1	2:B:419:VAL:HA	2.26	0.70
3:M:219:LEU:O	3:M:474:THR:HG21	1.91	0.70
3:M:323:MET:CE	3:M:342:LEU:HB3	2.21	0.70
3:M:373:ALA:O	3:M:418:GLU:N	2.24	0.70
4:S:50:PHE:CB	4:S:76:ILE:HA	2.20	0.70
1:A:528:ASN:O	1:A:530:ASN:N	2.24	0.70
2:B:375:LEU:O	2:B:377:TYR:N	2.23	0.70
2:B:458:MET:O	2:B:460:SER:N	2.25	0.70
2:B:530:LEU:HD23	2:B:591:MET:CB	2.21	0.70
3:M:224:VAL:CG2	3:M:479:PHE:CD1	2.74	0.70
3:M:343:ASN:HD22	3:M:343:ASN:H	1.39	0.70
3:M:443:SER:CB	3:M:447:ILE:C	2.59	0.70
4:S:135:ILE:HG22	4:S:141:VAL:CG2	2.20	0.70
1:A:103:LYS:N	4:S:163:THR:OG1	2.14	0.70
1:A:179:LYS:O	4:S:142:ILE:CD1	2.40	0.70
1:A:462:GLN:O	1:A:463:ASP:C	2.26	0.70
2:B:127:LEU:HA	2:B:161:LEU:HD11	1.72	0.70
2:B:245:GLN:NE2	2:B:309:LEU:HD11	2.05	0.70
2:B:285:GLU:O	2:B:286:ILE:HG22	1.91	0.70
2:B:549:LEU:CD1	2:B:611:ALA:HB2	2.07	0.70
3:M:215:TYR:HB3	3:M:467:TYR:CD2	2.21	0.70
3:M:240:ILE:HD11	3:M:306:LEU:HD11	1.72	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:129:LYS:HD2	1:A:161:ASP:HB3	1.73	0.70
1:A:222:ILE:HD12	1:A:240:LEU:HD21	1.71	0.70
2:B:212:VAL:HG13	2:B:248:LEU:HD23	1.71	0.70
2:B:267:ASP:O	2:B:269:SER:N	2.25	0.70
2:B:515:PHE:CE2	2:B:529:VAL:CG2	2.73	0.70
2:B:546:CYS:CA	2:B:607:ILE:CG1	2.54	0.70
3:M:306:LEU:O	3:M:307:SER:O	2.10	0.70
3:M:331:LEU:HD12	3:M:331:LEU:O	1.91	0.70
1:A:105:VAL:HG21	4:S:167:ILE:HA	1.73	0.70
1:A:150:LEU:HD22	1:A:162:ILE:HG12	1.73	0.70
1:A:163:ALA:CB	1:A:195:ALA:CB	2.55	0.70
1:A:403:LEU:HD21	1:A:421:PRO:O	1.91	0.70
2:B:215:TYR:CE1	2:B:233:TYR:CE1	2.77	0.70
2:B:292:GLU:CG	2:B:296:ASP:CB	2.65	0.70
2:B:563:PHE:CD2	2:B:584:SER:HB3	2.24	0.70
3:M:65:TYR:CD2	3:M:86:PRO:HB3	2.26	0.70
3:M:244:VAL:CB	3:M:300:LEU:HG	2.20	0.70
3:M:258:VAL:HG22	3:M:452:ILE:HG21	1.72	0.70
3:M:283:PHE:CE1	3:M:289:THR:OG1	2.39	0.70
3:M:378:ILE:C	3:M:379:LEU:HD12	2.11	0.70
3:M:452:ILE:HD12	3:M:452:ILE:N	2.06	0.70
4:S:69:ASN:OD1	4:S:73:ILE:O	2.09	0.70
1:A:67:LYS:CB	4:S:165:SER:C	2.59	0.70
1:A:399:ASP:C	1:A:420:ILE:H	1.95	0.70
2:B:120:ILE:HD12	2:B:142:LEU:HD22	1.72	0.70
2:B:306:LEU:HD22	2:B:321:CYS:HA	1.72	0.70
3:M:215:TYR:O	3:M:246:VAL:HG13	1.90	0.70
3:M:256:VAL:CG2	3:M:452:ILE:CG2	2.70	0.70
2:B:159:LYS:CE	2:B:191:GLU:OE1	2.39	0.70
2:B:252:LEU:HB3	2:B:302:PHE:CG	2.27	0.70
2:B:584:SER:O	2:B:585:GLY:C	2.20	0.70
3:M:217:ASP:OD1	3:M:217:ASP:O	2.10	0.70
3:M:374:TYR:H	3:M:390:ILE:HG23	1.56	0.70
1:A:129:LYS:HD2	1:A:161:ASP:CB	2.22	0.70
1:A:182:ILE:HG21	1:A:218:ALA:CA	2.22	0.70
1:A:421:PRO:HB2	1:A:424:TYR:CD1	2.27	0.70
2:B:139:LEU:HD23	2:B:172:GLU:O	1.91	0.70
2:B:217:GLU:O	2:B:218:CYS:C	2.28	0.70
2:B:278:PRO:CA	2:B:292:GLU:OE1	2.40	0.70
2:B:475:ILE:HG22	2:B:514:LEU:CD2	2.22	0.70
2:B:515:PHE:HE2	2:B:529:VAL:CG2	2.01	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:589:SER:HG	2:B:618:PHE:HZ	1.35	0.70
3:M:246:VAL:HB	3:M:297:PHE:CE1	2.26	0.70
4:S:16:LEU:HD12	4:S:17:VAL:N	2.07	0.70
1:A:67:LYS:CG	4:S:165:SER:OG	2.40	0.70
1:A:609:LEU:CD2	1:A:628:VAL:CG2	2.56	0.70
2:B:230:PHE:HD2	2:B:298:ASP:CB	2.03	0.70
3:M:276:VAL:CG2	3:M:299:LEU:HD12	2.21	0.70
1:A:237:SER:N	1:A:238:PRO:HD2	2.07	0.70
1:A:582:ILE:CD1	1:A:608:ARG:HA	2.22	0.70
1:A:603:VAL:HG22	2:B:610:ARG:NH2	2.07	0.70
2:B:20:ARG:CZ	2:B:35:TYR:OH	2.38	0.70
2:B:216:LYS:HB2	2:B:251:LEU:CD1	2.04	0.70
2:B:422:ALA:HB2	2:B:424:PHE:CE1	2.27	0.70
3:M:217:ASP:OD2	3:M:470:ALA:O	2.09	0.70
3:M:226:PHE:O	3:M:481:VAL:HA	1.91	0.70
3:M:350:VAL:HG13	3:M:442:GLN:HB2	1.72	0.70
4:S:5:VAL:HB	4:S:132:LEU:CD2	2.21	0.70
4:S:75:ILE:CG2	4:S:77:TYR:CZ	2.75	0.70
4:S:75:ILE:HG21	4:S:77:TYR:CZ	2.26	0.70
1:A:450:TYR:CE1	1:A:454:ILE:HD11	2.25	0.69
1:A:488:ARG:CG	1:A:522:PHE:CE2	2.71	0.69
2:B:42:ILE:O	2:B:43:ASN:CB	2.40	0.69
2:B:408:VAL:HG11	2:B:412:PHE:CE2	2.27	0.69
2:B:546:CYS:SG	2:B:607:ILE:HG13	2.32	0.69
2:B:574:ASN:O	2:B:576:GLN:O	2.10	0.69
2:B:592:TYR:CZ	2:B:619:ASP:OD1	2.44	0.69
2:B:599:ALA:O	2:B:601:TYR:N	2.25	0.69
3:M:9:ASP:CA	3:M:75:TRP:HD1	2.04	0.69
1:A:107:TYR:HB3	1:A:149:GLY:HA3	1.74	0.69
1:A:211:ASP:CG	4:S:148:ARG:NE	2.46	0.69
1:A:217:ALA:HB1	4:S:142:ILE:HD12	0.73	0.69
2:B:36:THR:O	2:B:40:GLN:HB2	1.93	0.69
2:B:120:ILE:HG22	2:B:153:ILE:HB	1.74	0.69
3:M:53:HIS:HB3	3:M:65:TYR:CE1	2.28	0.69
3:M:323:MET:SD	3:M:342:LEU:HA	2.32	0.69
1:A:252:ILE:CG1	4:S:144:THR:OG1	2.41	0.69
1:A:282:MET:HE3	1:A:294:SER:HB2	1.75	0.69
1:A:398:GLU:O	1:A:419:ILE:C	2.30	0.69
1:A:421:PRO:HG2	1:A:424:TYR:CZ	2.27	0.69
2:B:68:SER:O	2:B:71:ALA:N	2.24	0.69
2:B:70:MET:HE1	2:B:107:ARG:HB2	0.70	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:319:LEU:HD13	2:B:358:MET:HG2	1.69	0.69
2:B:563:PHE:CE2	2:B:584:SER:HA	2.25	0.69
2:B:565:GLN:C	2:B:574:ASN:ND2	2.38	0.69
1:A:215:VAL:HG21	1:A:243:ILE:CD1	2.15	0.69
2:B:155:LEU:HD13	2:B:156:HIS:N	2.07	0.69
2:B:435:SER:O	2:B:437:SER:N	2.26	0.69
2:B:508:ARG:HB3	2:B:544:THR:HG23	1.68	0.69
2:B:550:VAL:CG2	2:B:610:ARG:HD2	2.22	0.69
3:M:101:LEU:HD11	3:M:106:LYS:O	1.91	0.69
3:M:231:SER:O	3:M:326:HIS:CE1	2.46	0.69
1:A:140:VAL:HG13	1:A:176:TYR:CB	2.21	0.69
1:A:147:LEU:HB3	1:A:184:ALA:HB3	1.67	0.69
1:A:219:VAL:C	4:S:140:MET:SD	2.49	0.69
1:A:225:LEU:HB3	1:A:233:PHE:CZ	2.26	0.69
2:B:67:ILE:CD1	2:B:103:LEU:CB	2.71	0.69
2:B:132:SER:O	2:B:133:GLU:C	2.24	0.69
2:B:162:VAL:HG23	2:B:195:ILE:HG22	1.54	0.69
2:B:174:ALA:HB1	2:B:211:ALA:CB	2.22	0.69
2:B:178:ILE:HA	2:B:218:CYS:CB	2.14	0.69
2:B:215:TYR:HD2	2:B:226:LEU:CD1	2.04	0.69
2:B:247:TYR:CZ	3:M:91:THR:CG2	2.68	0.69
2:B:306:LEU:HD12	2:B:325:LEU:HD23	1.73	0.69
2:B:353:GLN:CG	3:M:47:SER:CB	2.70	0.69
2:B:493:LEU:CD2	2:B:511:ILE:HG23	2.22	0.69
3:M:220:GLU:CD	3:M:442:GLN:HB3	2.12	0.69
4:S:14:PRO:HA	4:S:36:TYR:HH	1.56	0.69
1:A:250:ASN:OD1	1:A:285:THR:CB	2.38	0.69
1:A:408:ILE:N	4:S:64:ASN:HB2	2.07	0.69
2:B:219:TYR:CD2	2:B:226:LEU:CB	2.64	0.69
2:B:322:CYS:HA	2:B:325:LEU:HD12	1.73	0.69
2:B:500:GLN:CD	2:B:503:LEU:HD21	2.12	0.69
2:B:613:MET:HE3	2:B:617:LEU:HD11	1.74	0.69
1:A:97:SER:C	1:A:98:ASN:O	2.05	0.69
1:A:323:CYS:HG	1:A:338:PHE:HE1	0.74	0.69
2:B:64:LYS:HG3	2:B:100:LEU:HD11	1.73	0.69
2:B:206:LYS:O	2:B:210:CYS:SG	2.42	0.69
2:B:433:VAL:CG1	2:B:474:VAL:HB	2.22	0.69
2:B:569:THR:HG22	2:B:569:THR:O	1.92	0.69
4:S:9:ASN:ND2	4:S:13:GLN:CB	2.55	0.69
1:A:104:ARG:HE	4:S:160:ALA:CB	2.06	0.69
1:A:140:VAL:HG11	1:A:176:TYR:HB2	1.64	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:186:PHE:CE2	1:A:187:LYS:HD3	2.28	0.69
1:A:206:LYS:HE2	1:A:206:LYS:HA	1.74	0.69
1:A:207:LEU:HD21	1:A:239:LEU:HB2	1.75	0.69
1:A:288:THR:HA	1:A:291:ILE:HD13	1.73	0.69
1:A:477:PHE:CE2	1:A:481:MET:SD	2.86	0.69
1:A:594:PHE:CD1	2:B:434:LYS:HE2	2.27	0.69
2:B:37:TYR:CE2	2:B:38:TYR:CD1	2.77	0.69
2:B:121:ASN:O	2:B:124:GLN:HB3	1.93	0.69
2:B:177:ILE:CB	2:B:196:LEU:HD21	2.18	0.69
2:B:178:ILE:HG12	2:B:214:ALA:HB1	1.74	0.69
2:B:182:ARG:NE	2:B:217:GLU:OE1	2.26	0.69
2:B:230:PHE:CE2	2:B:234:CYS:SG	2.86	0.69
2:B:236:ILE:CG2	2:B:240:LEU:HD11	2.21	0.69
2:B:508:ARG:CB	2:B:544:THR:HG21	2.22	0.69
2:B:513:TRP:CB	2:B:551:LEU:CD2	2.67	0.69
2:B:519:ALA:O	2:B:523:PHE:CA	2.41	0.69
2:B:525:ILE:O	2:B:527:PRO:HD2	1.93	0.69
3:M:16:PHE:HZ	3:M:18:TYR:HB2	1.52	0.69
3:M:214:LEU:O	3:M:467:TYR:CA	2.41	0.69
3:M:215:TYR:O	3:M:470:ALA:HB2	1.92	0.69
3:M:224:VAL:O	3:M:479:PHE:CB	2.40	0.69
3:M:233:LEU:HD21	3:M:325:LEU:H	1.56	0.69
3:M:235:LEU:HD11	3:M:306:LEU:HB3	1.71	0.69
3:M:246:VAL:HA	3:M:470:ALA:HB1	1.74	0.69
1:A:189:PHE:CB	1:A:225:LEU:CD2	2.71	0.69
1:A:211:ASP:CG	4:S:148:ARG:NH1	2.46	0.69
1:A:252:ILE:HG13	4:S:145:ASN:N	2.07	0.69
1:A:272:ALA:O	1:A:276:PRO:HD3	1.93	0.69
1:A:298:ILE:CD1	1:A:311:THR:CG2	2.69	0.69
1:A:629:LEU:CD1	2:B:610:ARG:HH11	1.74	0.69
2:B:116:THR:O	2:B:120:ILE:HG12	1.93	0.69
2:B:215:TYR:CZ	2:B:229:HIS:CB	2.76	0.69
2:B:267:ASP:O	2:B:276:SER:HB2	1.93	0.69
2:B:275:ARG:HG3	2:B:291:TYR:CE2	2.28	0.69
2:B:357:GLU:HG2	2:B:361:GLN:NE2	2.06	0.69
2:B:374:PHE:HZ	2:B:381:PHE:CD1	2.09	0.69
3:M:55:MET:O	3:M:56:VAL:C	2.17	0.69
3:M:260:LEU:HD23	3:M:449:VAL:HG22	0.73	0.69
3:M:374:TYR:HB2	3:M:417:TYR:CD2	2.28	0.69
3:M:378:ILE:HB	3:M:413:GLY:HA2	1.73	0.69
4:S:7:ILE:CG2	4:S:121:LEU:CD2	2.70	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:231:GLN:HB2	1:A:232:PRO:HD3	1.74	0.69
1:A:322:PHE:HD2	1:A:330:LEU:HD21	1.53	0.69
1:A:332:TYR:CZ	1:A:336:ILE:CD1	2.75	0.69
1:A:482:ILE:HG12	1:A:517:TRP:CZ3	2.27	0.69
1:A:582:ILE:HG12	1:A:607:LEU:HB2	1.74	0.69
2:B:87:VAL:O	2:B:90:ILE:HG22	1.93	0.69
2:B:171:GLY:HA3	2:B:207:VAL:HA	1.75	0.69
2:B:332:LEU:HG	2:B:332:LEU:O	1.93	0.69
2:B:523:PHE:HB2	2:B:559:ASP:OD2	1.93	0.69
3:M:472:TYR:CD1	3:M:472:TYR:N	2.58	0.69
4:S:15:ARG:HH21	4:S:118:GLU:CD	1.96	0.69
4:S:53:THR:OG1	4:S:68:VAL:O	2.10	0.69
4:S:75:ILE:HG21	4:S:77:TYR:OH	1.92	0.69
1:A:64:LEU:CG	1:A:102:GLN:NE2	2.44	0.68
1:A:101:GLN:CD	4:S:167:ILE:HG21	2.12	0.68
1:A:186:PHE:HD2	1:A:187:LYS:N	1.91	0.68
1:A:211:ASP:OD2	4:S:148:ARG:HD3	1.93	0.68
1:A:223:CYS:HB2	1:A:259:LEU:CG	2.19	0.68
1:A:488:ARG:HG2	1:A:522:PHE:CD2	2.28	0.68
1:A:552:ILE:HG22	1:A:603:VAL:HG21	1.74	0.68
2:B:8:ILE:O	2:B:12:LEU:HD12	1.92	0.68
2:B:81:LEU:C	2:B:83:PHE:N	2.33	0.68
2:B:135:ARG:CZ	2:B:164:ASP:HB2	2.24	0.68
2:B:267:ASP:O	2:B:268:LYS:C	2.29	0.68
2:B:306:LEU:HB2	2:B:325:LEU:HD21	1.74	0.68
2:B:360:LEU:HD11	2:B:391:ALA:CB	2.22	0.68
2:B:542:PRO:O	2:B:545:ARG:N	2.26	0.68
2:B:592:TYR:HD2	2:B:618:PHE:CE2	2.09	0.68
3:M:339:GLU:HG3	3:M:412:ARG:HD3	1.75	0.68
3:M:410:VAL:HG13	3:M:412:ARG:NH1	2.08	0.68
4:S:87:PHE:CE1	4:S:102:ILE:HG12	2.27	0.68
1:A:244:LEU:HB2	1:A:256:LEU:HD13	1.74	0.68
1:A:566:PHE:C	1:A:568:GLU:H	1.96	0.68
1:A:630:PRO:CG	2:B:614:ILE:CA	2.60	0.68
2:B:48:VAL:O	2:B:51:LEU:N	2.22	0.68
2:B:307:ASN:OD1	2:B:339:PHE:HD2	1.76	0.68
2:B:340:ILE:CG1	2:B:373:LEU:CD2	2.70	0.68
2:B:352:ASN:CB	3:M:49:ASP:CB	2.48	0.68
3:M:245:ASP:OD1	3:M:297:PHE:C	2.32	0.68
1:A:636:TYR:H	2:B:554:LYS:NZ	1.90	0.68
2:B:542:PRO:C	2:B:602:ASP:OD2	2.31	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:66:PHE:HE2	3:M:77:LEU:O	1.77	0.68
3:M:67:SER:HG	3:M:90:PHE:HD1	0.70	0.68
3:M:101:LEU:CD1	3:M:109:LEU:HD13	2.22	0.68
3:M:222:PHE:CD1	3:M:439:TYR:CE1	2.81	0.68
3:M:244:VAL:N	3:M:300:LEU:O	2.27	0.68
3:M:260:LEU:HA	3:M:448:TYR:O	1.93	0.68
4:S:85:PHE:CZ	4:S:109:LEU:HD23	2.28	0.68
4:S:111:ARG:CB	4:S:150:VAL:HG22	2.22	0.68
1:A:96:SER:HA	1:A:127:LEU:HG	1.76	0.68
1:A:170:LEU:HB3	1:A:206:LYS:CG	2.23	0.68
1:A:566:PHE:CD1	1:A:570:LYS:HA	2.28	0.68
2:B:47:LEU:HD22	2:B:66:ILE:CB	2.15	0.68
2:B:176:ALA:C	2:B:178:ILE:N	2.44	0.68
2:B:177:ILE:HD12	2:B:196:LEU:CD2	2.24	0.68
2:B:219:TYR:HB2	2:B:226:LEU:HD22	1.73	0.68
2:B:352:ASN:HB2	3:M:49:ASP:CG	2.12	0.68
3:M:223:HIS:CD2	3:M:478:ASN:HA	2.27	0.68
3:M:242:GLY:HA2	3:M:474:THR:CG2	2.22	0.68
3:M:319:SER:O	3:M:343:ASN:O	2.12	0.68
3:M:443:SER:HB3	3:M:447:ILE:CA	2.23	0.68
4:S:50:PHE:HA	4:S:77:TYR:HD1	1.58	0.68
1:A:79:MET:CG	1:A:112:GLN:OE1	2.40	0.68
1:A:277:LYS:O	1:A:277:LYS:HG3	1.94	0.68
1:A:303:MET:HG2	1:A:308:ASP:OD2	1.92	0.68
1:A:441:TYR:HB3	1:A:444:VAL:HG23	1.75	0.68
1:A:509:PRO:CB	1:A:547:VAL:CG2	2.67	0.68
1:A:589:SER:CB	1:A:601:VAL:CG2	2.70	0.68
2:B:2:VAL:HG21	3:M:58:ARG:HD2	1.75	0.68
2:B:102:HIS:ND1	2:B:137:PHE:CB	2.52	0.68
2:B:563:PHE:O	2:B:566:ALA:CB	2.42	0.68
3:M:65:TYR:CG	3:M:86:PRO:HA	2.29	0.68
3:M:224:VAL:HG22	3:M:306:LEU:HD12	1.73	0.68
3:M:225:VAL:HB	3:M:237:THR:OG1	1.93	0.68
1:A:105:VAL:HG21	4:S:167:ILE:CB	2.23	0.68
1:A:178:ARG:HB3	1:A:214:VAL:CG1	2.23	0.68
1:A:189:PHE:CD2	1:A:225:LEU:CD1	2.76	0.68
1:A:192:TYR:CE2	1:A:194:GLU:HB2	2.28	0.68
1:A:271:ARG:NH1	1:A:302:ASN:O	2.26	0.68
1:A:461:CYS:HB2	1:A:469:LEU:CD2	2.23	0.68
2:B:24:ALA:CB	2:B:35:TYR:CD2	2.68	0.68
2:B:146:LYS:O	2:B:147:MET:CG	2.42	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:412:PHE:CE2	2:B:446:TRP:HB3	2.22	0.68
2:B:422:ALA:CB	2:B:424:PHE:CD1	2.54	0.68
2:B:512:VAL:HG21	2:B:548:ILE:CG1	2.22	0.68
3:M:222:PHE:HB2	3:M:479:PHE:CE1	2.29	0.68
4:S:105:PHE:HZ	4:S:128:LEU:HD11	1.58	0.68
1:A:214:VAL:HG23	4:S:148:ARG:NH2	2.08	0.68
1:A:240:LEU:O	1:A:241:TYR:O	2.11	0.68
1:A:244:LEU:CD1	1:A:281:LEU:HD11	2.24	0.68
1:A:482:ILE:HG12	1:A:517:TRP:HH2	1.58	0.68
2:B:37:TYR:O	2:B:38:TYR:C	2.31	0.68
2:B:90:ILE:HG13	2:B:98:LYS:HD3	1.74	0.68
2:B:247:TYR:OH	3:M:91:THR:HG21	1.94	0.68
2:B:334:MET:HE2	2:B:334:MET:CA	2.17	0.68
2:B:361:GLN:HG2	2:B:394:TRP:CZ2	2.29	0.68
2:B:433:VAL:O	2:B:474:VAL:HG11	1.93	0.68
2:B:589:SER:HA	2:B:592:TYR:HD2	1.58	0.68
3:M:306:LEU:HD22	3:M:317:MET:CE	2.23	0.68
3:M:317:MET:CB	3:M:321:GLY:C	2.57	0.68
1:A:605:GLU:CG	1:A:632:PHE:CE2	2.69	0.68
2:B:37:TYR:CE2	2:B:46:GLN:NE2	2.48	0.68
2:B:47:LEU:CD1	2:B:66:ILE:HA	2.23	0.68
2:B:216:LYS:CG	2:B:251:LEU:CD1	2.47	0.68
3:M:65:TYR:CD2	3:M:86:PRO:CA	2.76	0.68
3:M:244:VAL:CG1	3:M:472:TYR:CZ	2.75	0.68
3:M:246:VAL:HB	3:M:297:PHE:HZ	1.51	0.68
3:M:339:GLU:CD	3:M:412:ARG:CD	2.60	0.68
3:M:360:LEU:CD1	3:M:433:VAL:HB	2.23	0.68
4:S:9:ASN:ND2	4:S:118:GLU:OE1	2.27	0.68
1:A:102:GLN:N	4:S:167:ILE:CG1	2.55	0.68
1:A:558:VAL:O	1:A:561:ASN:N	2.15	0.68
2:B:20:ARG:CG	2:B:21:GLU:H	2.06	0.68
2:B:42:ILE:O	2:B:46:GLN:CD	2.31	0.68
2:B:140:SER:O	2:B:143:SER:N	2.21	0.68
2:B:523:PHE:CD1	2:B:559:ASP:CG	2.67	0.68
2:B:549:LEU:HD13	2:B:611:ALA:HB2	1.71	0.68
3:M:10:THR:OG1	3:M:48:ASP:OD1	2.09	0.68
3:M:101:LEU:CD2	3:M:106:LYS:HG3	2.24	0.68
3:M:300:LEU:HD11	3:M:447:ILE:HD12	1.76	0.68
4:S:8:PHE:CB	4:S:36:TYR:OH	2.41	0.68
1:A:100:LEU:CD2	1:A:138:ASN:HB3	2.23	0.68
1:A:189:PHE:CE2	1:A:225:LEU:HD11	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:401:VAL:CG1	1:A:419:ILE:HD12	2.23	0.68
1:A:594:PHE:CD2	2:B:474:VAL:CG2	2.77	0.68
2:B:144:ASP:HA	2:B:179:LYS:HD3	1.72	0.68
2:B:380:LYS:HZ1	3:M:236:LEU:CD1	2.07	0.68
1:A:102:GLN:HB3	4:S:166:LYS:CG	2.23	0.67
1:A:212:ILE:HG22	1:A:247:ILE:HD13	1.75	0.67
1:A:635:ALA:C	2:B:517:GLU:HA	2.15	0.67
2:B:124:GLN:CA	2:B:157:THR:OG1	2.31	0.67
2:B:148:SER:O	2:B:183:ALA:CB	2.36	0.67
3:M:223:HIS:HB3	3:M:478:ASN:CA	2.24	0.67
3:M:354:ASP:O	3:M:438:SER:O	2.12	0.67
3:M:371:GLU:HB3	3:M:424:PHE:HD1	1.60	0.67
4:S:14:PRO:O	4:S:15:ARG:CG	2.43	0.67
4:S:130:SER:CB	4:S:156:LEU:CD1	2.65	0.67
1:A:186:PHE:CD1	1:A:224:GLU:HB2	2.27	0.67
1:A:186:PHE:CZ	4:S:138:GLY:N	2.29	0.67
1:A:190:LEU:CD1	1:A:228:LYS:HE3	2.23	0.67
1:A:318:ARG:O	1:A:322:PHE:HD1	1.76	0.67
1:A:384:LEU:CG	1:A:441:TYR:HE2	2.06	0.67
1:A:446:ASP:O	1:A:448:GLU:O	2.11	0.67
2:B:67:ILE:HD13	2:B:103:LEU:HB2	1.76	0.67
2:B:204:ASP:O	2:B:207:VAL:HB	1.94	0.67
2:B:293:VAL:O	2:B:299:LEU:CG	2.42	0.67
2:B:493:LEU:HD21	2:B:511:ILE:HG12	1.74	0.67
3:M:269:ILE:O	3:M:302:TYR:CD1	2.47	0.67
3:M:281:GLY:O	3:M:282:VAL:CB	2.42	0.67
3:M:304:VAL:HG21	3:M:309:GLN:NE2	2.09	0.67
4:S:94:SER:O	4:S:98:ILE:HG12	1.94	0.67
1:A:70:ALA:O	1:A:74:LEU:HG	1.94	0.67
1:A:137:ASN:O	1:A:139:ASP:N	2.26	0.67
1:A:292:TYR:CD1	1:A:292:TYR:O	2.48	0.67
1:A:409:VAL:O	1:A:409:VAL:HG13	1.93	0.67
1:A:572:PHE:O	1:A:575:LYS:HB3	1.93	0.67
1:A:585:PHE:CE2	1:A:603:VAL:CG1	2.77	0.67
2:B:94:ASP:O	2:B:134:LEU:HD21	1.94	0.67
3:M:66:PHE:CB	3:M:77:LEU:CD2	2.64	0.67
3:M:215:TYR:HB2	3:M:468:LYS:N	2.09	0.67
3:M:222:PHE:CA	3:M:479:PHE:CZ	2.76	0.67
4:S:9:ASN:CG	4:S:13:GLN:HB2	2.14	0.67
4:S:47:GLN:OE1	4:S:78:LYS:C	2.32	0.67
1:A:189:PHE:HD2	1:A:225:LEU:CD1	2.06	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:249:ASN:HD21	4:S:146:VAL:HG23	1.58	0.67
1:A:282:MET:O	1:A:284:SER:N	2.28	0.67
1:A:429:VAL:HG11	1:A:469:LEU:HD11	1.77	0.67
1:A:465:SER:N	2:B:1:MET:HG2	2.09	0.67
2:B:92:THR:O	2:B:134:LEU:HD13	1.93	0.67
2:B:173:VAL:HB	2:B:199:LEU:HD13	1.75	0.67
2:B:215:TYR:HE2	2:B:229:HIS:ND1	1.90	0.67
2:B:216:LYS:HD3	2:B:251:LEU:HB2	1.77	0.67
3:M:96:ILE:HG23	3:M:125:PHE:CE1	2.30	0.67
3:M:245:ASP:HA	3:M:297:PHE:O	1.94	0.67
1:A:448:GLU:CB	1:A:487:MET:SD	2.78	0.67
2:B:226:LEU:O	2:B:229:HIS:HB2	1.95	0.67
2:B:260:LEU:CB	2:B:291:TYR:CZ	2.73	0.67
2:B:309:LEU:CD1	2:B:317:VAL:HG11	2.24	0.67
2:B:380:LYS:NZ	3:M:236:LEU:CD1	2.58	0.67
3:M:253:ASN:HA	3:M:292:PRO:HG2	1.75	0.67
1:A:99:LYS:HZ2	4:S:164:ASP:CB	2.05	0.67
1:A:103:LYS:NZ	1:A:131:ASP:HB2	2.08	0.67
1:A:105:VAL:HG23	4:S:167:ILE:HD12	0.68	0.67
1:A:189:PHE:CB	1:A:225:LEU:HD21	2.24	0.67
1:A:212:ILE:O	1:A:213:SER:C	2.21	0.67
1:A:392:MET:SD	1:A:457:LEU:HD21	2.34	0.67
2:B:4:SER:O	2:B:8:ILE:CD1	2.43	0.67
2:B:123:LEU:HB3	2:B:142:LEU:HD11	1.76	0.67
3:M:375:LYS:HG3	3:M:418:GLU:OE1	1.95	0.67
1:A:100:LEU:H	4:S:162:SER:CA	2.00	0.67
1:A:104:ARG:CD	1:A:145:ILE:HG13	2.25	0.67
1:A:140:VAL:CA	1:A:177:ILE:HD11	2.22	0.67
1:A:145:ILE:HD11	4:S:156:LEU:HD13	1.77	0.67
1:A:594:PHE:HB2	2:B:473:ASN:HB2	1.71	0.67
2:B:47:LEU:CG	2:B:66:ILE:HG12	2.25	0.67
3:M:327:PHE:CE1	3:M:336:ASP:CB	2.69	0.67
1:A:125:THR:O	1:A:127:LEU:N	2.27	0.67
1:A:320:HIS:CB	1:A:352:PHE:HE2	2.05	0.67
1:A:401:VAL:HB	1:A:419:ILE:CG1	2.24	0.67
1:A:552:ILE:HD13	1:A:600:SER:CB	2.25	0.67
1:A:625:LEU:CG	2:B:613:MET:SD	2.79	0.67
2:B:175:LEU:CA	2:B:214:ALA:HB2	2.19	0.67
2:B:208:ILE:HG21	2:B:236:ILE:HG21	1.77	0.67
2:B:315:PRO:HA	2:B:318:ILE:HD12	1.76	0.67
2:B:613:MET:CE	2:B:617:LEU:HD11	2.25	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:2:TYR:CE1	3:M:64:LYS:NZ	2.57	0.67
3:M:2:TYR:CA	3:M:81:SER:HB2	2.25	0.67
1:A:100:LEU:O	1:A:101:GLN:O	2.13	0.67
1:A:264:SER:HB3	1:A:271:ARG:HG2	1.75	0.67
2:B:37:TYR:HD2	2:B:38:TYR:CE1	2.01	0.67
2:B:120:ILE:HD12	2:B:142:LEU:CD2	2.24	0.67
2:B:139:LEU:HD23	2:B:172:GLU:C	2.15	0.67
2:B:560:ILE:HG22	2:B:561:ASP:N	2.10	0.67
3:M:7:ILE:C	3:M:75:TRP:O	2.32	0.67
4:S:53:THR:CB	4:S:69:ASN:CB	2.73	0.67
1:A:109:ALA:O	1:A:112:GLN:N	2.27	0.67
1:A:140:VAL:CG2	1:A:177:ILE:CG1	2.68	0.67
1:A:182:ILE:CG2	1:A:221:VAL:HG23	2.17	0.67
1:A:370:LYS:O	1:A:374:LEU:CD1	2.42	0.67
1:A:421:PRO:CG	1:A:424:TYR:CD1	2.76	0.67
1:A:478:ARG:HD3	1:A:514:GLU:OE1	1.95	0.67
1:A:556:VAL:CG2	1:A:603:VAL:CG1	2.73	0.67
1:A:637:GLU:CB	2:B:551:LEU:HD22	2.25	0.67
2:B:219:TYR:CA	2:B:223:LEU:HD23	2.24	0.67
2:B:340:ILE:CD1	2:B:366:LEU:CB	2.73	0.67
3:M:54:SER:HB2	3:M:66:PHE:CD1	2.30	0.67
3:M:222:PHE:CD2	3:M:439:TYR:CE1	2.82	0.67
3:M:233:LEU:CD2	3:M:323:MET:O	2.42	0.67
3:M:338:PHE:CD2	3:M:415:ILE:CD1	2.78	0.67
1:A:186:PHE:CD2	1:A:187:LYS:N	2.63	0.66
2:B:20:ARG:HD2	2:B:21:GLU:N	2.09	0.66
2:B:80:GLN:CG	2:B:115:LEU:HD21	2.25	0.66
2:B:195:ILE:C	2:B:197:LYS:N	2.43	0.66
2:B:277:CYS:HA	2:B:292:GLU:CG	2.25	0.66
2:B:303:LEU:HD11	2:B:333:GLN:HE21	1.58	0.66
2:B:347:VAL:HG22	2:B:359:LEU:CB	2.13	0.66
2:B:381:PHE:O	2:B:395:LYS:HD2	1.95	0.66
2:B:537:PHE:CE1	2:B:545:ARG:CG	2.75	0.66
3:M:3:LEU:HD12	3:M:3:LEU:N	2.10	0.66
3:M:215:TYR:HB3	3:M:470:ALA:H	1.60	0.66
4:S:9:ASN:HD21	4:S:13:GLN:CB	2.09	0.66
4:S:99:LEU:O	4:S:102:ILE:HB	1.95	0.66
1:A:104:ARG:HG3	1:A:145:ILE:CG2	2.25	0.66
1:A:214:VAL:HG23	4:S:148:ARG:CZ	2.26	0.66
1:A:557:LYS:HE3	2:B:606:ASP:HB3	1.78	0.66
1:A:566:PHE:HZ	1:A:618:THR:C	1.99	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:605:GLU:OE2	1:A:608:ARG:NH2	2.25	0.66
2:B:13:ASP:OD2	3:M:17:GLN:OE1	2.14	0.66
2:B:120:ILE:HD13	2:B:142:LEU:CD2	2.22	0.66
2:B:175:LEU:CD1	2:B:210:CYS:HB3	2.25	0.66
2:B:249:ILE:HG21	2:B:320:SER:HB3	1.76	0.66
2:B:363:ILE:CG2	2:B:398:ILE:CD1	2.73	0.66
2:B:364:HIS:O	2:B:367:SER:N	2.28	0.66
2:B:375:LEU:HD23	2:B:402:LEU:HD22	1.78	0.66
3:M:269:ILE:C	3:M:302:TYR:CE1	2.68	0.66
3:M:324:SER:O	3:M:339:GLU:O	2.13	0.66
3:M:340:LEU:N	3:M:411:LEU:O	2.28	0.66
3:M:443:SER:CB	3:M:447:ILE:CG1	2.37	0.66
4:S:14:PRO:O	4:S:15:ARG:HG2	1.95	0.66
1:A:429:VAL:CG1	1:A:469:LEU:CD1	2.69	0.66
2:B:343:LEU:CD1	2:B:359:LEU:HD12	2.18	0.66
2:B:363:ILE:HG21	2:B:398:ILE:HD11	1.77	0.66
2:B:534:ILE:HD13	2:B:594:ALA:CB	2.20	0.66
3:M:410:VAL:CG1	3:M:412:ARG:HH11	2.08	0.66
3:M:437:TYR:CD1	3:M:479:PHE:CZ	2.83	0.66
1:A:103:LYS:O	1:A:107:TYR:HD1	1.74	0.66
1:A:219:VAL:O	4:S:140:MET:SD	2.52	0.66
1:A:244:LEU:CD1	1:A:256:LEU:HD12	2.25	0.66
1:A:625:LEU:HG	2:B:613:MET:SD	2.35	0.66
2:B:77:ILE:O	2:B:79:VAL:CA	2.43	0.66
2:B:124:GLN:HB2	2:B:153:ILE:CG2	2.24	0.66
2:B:127:LEU:HD12	2:B:157:THR:OG1	1.96	0.66
2:B:472:VAL:O	2:B:473:ASN:C	2.33	0.66
2:B:512:VAL:CG1	2:B:548:ILE:HA	2.10	0.66
3:M:396:GLN:HG2	3:M:398:ILE:HD11	1.77	0.66
4:S:85:PHE:HE2	4:S:109:LEU:CD2	2.08	0.66
1:A:64:LEU:HB3	1:A:102:GLN:NE2	2.10	0.66
1:A:370:LYS:HE2	1:A:370:LYS:HA	1.76	0.66
1:A:581:LEU:HB3	1:A:607:LEU:HD13	1.77	0.66
1:A:631:SER:O	2:B:554:LYS:CG	2.35	0.66
2:B:85:ASP:O	2:B:89:ASN:ND2	2.28	0.66
2:B:173:VAL:HB	2:B:199:LEU:HD11	1.76	0.66
2:B:174:ALA:HB3	2:B:211:ALA:CB	2.22	0.66
2:B:204:ASP:HB3	2:B:207:VAL:CG2	2.25	0.66
2:B:208:ILE:HG21	2:B:236:ILE:CG2	2.25	0.66
2:B:344:VAL:HG22	2:B:381:PHE:HZ	1.61	0.66
2:B:553:ALA:HA	2:B:614:ILE:HG23	1.78	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:8:THR:HA	3:M:13:LYS:O	1.95	0.66
3:M:220:GLU:HB2	3:M:222:PHE:HE1	1.60	0.66
4:S:130:SER:HB2	4:S:156:LEU:HD13	1.76	0.66
1:A:516:ILE:CG1	1:A:551:LEU:HD13	2.26	0.66
1:A:557:LYS:CD	2:B:606:ASP:N	2.56	0.66
1:A:629:LEU:HG	2:B:610:ARG:NH1	2.05	0.66
2:B:90:ILE:CD1	2:B:123:LEU:HD21	2.24	0.66
2:B:94:ASP:C	2:B:134:LEU:HD21	2.16	0.66
2:B:139:LEU:CD2	2:B:173:VAL:O	2.32	0.66
2:B:315:PRO:HG3	2:B:352:ASN:OD1	1.94	0.66
3:M:437:TYR:N	3:M:437:TYR:HD1	1.91	0.66
4:S:110:ASP:O	4:S:113:PHE:O	2.14	0.66
1:A:221:VAL:HG23	4:S:142:ILE:HG21	1.77	0.66
1:A:241:TYR:CD2	1:A:242:GLU:N	2.63	0.66
1:A:399:ASP:O	1:A:420:ILE:CB	2.43	0.66
1:A:454:ILE:HG23	1:A:473:ILE:CG2	2.24	0.66
1:A:626:SER:C	2:B:617:LEU:CG	2.35	0.66
2:B:123:LEU:CD2	2:B:138:ALA:O	2.42	0.66
2:B:230:PHE:CD2	2:B:298:ASP:O	2.48	0.66
2:B:232:ARG:CG	2:B:236:ILE:HD11	2.26	0.66
2:B:234:CYS:O	2:B:237:ILE:CG2	2.38	0.66
2:B:437:SER:C	2:B:478:LEU:HD21	2.15	0.66
3:M:51:LEU:N	3:M:51:LEU:HD12	2.10	0.66
3:M:379:LEU:HD23	3:M:397:TRP:HE1	1.58	0.66
1:A:92:LEU:HD11	1:A:120:ILE:CA	2.13	0.66
1:A:401:VAL:CG2	1:A:419:ILE:HB	2.26	0.66
1:A:509:PRO:HB3	1:A:547:VAL:HG21	1.75	0.66
1:A:581:LEU:CD2	1:A:607:LEU:HD11	2.26	0.66
2:B:353:GLN:HG2	3:M:48:ASP:CA	2.26	0.66
2:B:375:LEU:O	2:B:376:PRO:C	2.33	0.66
2:B:403:ILE:HA	2:B:411:ILE:HD12	1.77	0.66
2:B:537:PHE:CG	2:B:598:LEU:CB	2.74	0.66
2:B:556:LEU:CA	2:B:588:ILE:CD1	2.21	0.66
2:B:563:PHE:HE2	2:B:584:SER:CA	2.06	0.66
3:M:259:LYS:O	3:M:449:VAL:HG13	1.96	0.66
4:S:48:SER:HA	4:S:77:TYR:CB	2.25	0.66
4:S:54:PRO:HB2	4:S:57:LEU:CD1	2.26	0.66
4:S:137:GLN:O	4:S:140:MET:HB3	1.96	0.66
1:A:92:LEU:O	1:A:95:MET:N	2.27	0.66
1:A:100:LEU:O	1:A:101:GLN:C	2.16	0.66
1:A:102:GLN:CA	4:S:166:LYS:HB2	2.26	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:150:LEU:HD22	1:A:162:ILE:CG1	2.26	0.66
1:A:288:THR:HB	1:A:322:PHE:HZ	1.59	0.66
1:A:318:ARG:O	1:A:322:PHE:CD1	2.49	0.66
1:A:625:LEU:O	1:A:627:GLU:N	2.28	0.66
2:B:215:TYR:HD2	2:B:226:LEU:HD13	1.60	0.66
2:B:400:SER:HB3	2:B:439:CYS:SG	2.35	0.66
3:M:215:TYR:HD2	3:M:469:GLY:H	1.29	0.66
3:M:265:ASN:HB3	3:M:309:GLN:HG3	1.77	0.66
1:A:92:LEU:HD22	1:A:123:LEU:HB2	1.78	0.66
1:A:215:VAL:O	1:A:219:VAL:HG23	1.95	0.66
2:B:141:ALA:O	2:B:142:LEU:C	2.31	0.66
2:B:390:VAL:HG12	2:B:394:TRP:CD1	2.31	0.66
2:B:418:TYR:CD1	2:B:419:VAL:CA	2.79	0.66
2:B:497:LEU:HD21	2:B:533:LEU:HD22	1.77	0.66
2:B:592:TYR:CE2	2:B:618:PHE:HD2	2.13	0.66
3:M:2:TYR:H	3:M:81:SER:CB	2.03	0.66
3:M:100:LEU:CD1	3:M:100:LEU:H	2.09	0.66
3:M:258:VAL:HG13	3:M:452:ILE:HG13	1.77	0.66
3:M:272:LEU:HD21	3:M:288:ILE:HD13	1.76	0.66
3:M:435:LEU:O	3:M:479:PHE:CE2	2.49	0.66
4:S:57:LEU:N	4:S:57:LEU:HD12	2.11	0.66
1:A:323:CYS:CB	1:A:355:LEU:HD21	2.26	0.65
2:B:123:LEU:HD13	2:B:142:LEU:HG	0.67	0.65
2:B:530:LEU:HD23	2:B:591:MET:HB2	1.78	0.65
3:M:213:GLU:C	3:M:467:TYR:HB2	2.16	0.65
3:M:360:LEU:HD13	3:M:433:VAL:CG2	2.26	0.65
1:A:520:GLY:HA2	1:A:558:VAL:CG2	2.23	0.65
2:B:67:ILE:HD11	2:B:100:LEU:O	1.95	0.65
2:B:103:LEU:CD1	3:M:132:GLY:HA3	2.24	0.65
2:B:120:ILE:HD11	2:B:150:LEU:HD13	1.77	0.65
2:B:175:LEU:CD2	2:B:210:CYS:C	2.64	0.65
2:B:336:ASN:HB3	2:B:339:PHE:CD1	2.32	0.65
3:M:2:TYR:CE2	3:M:62:VAL:HG13	2.31	0.65
3:M:302:TYR:HE2	3:M:304:VAL:HB	1.61	0.65
3:M:403:THR:HG23	3:M:407:THR:HG1	1.62	0.65
3:M:437:TYR:CE1	3:M:479:PHE:CE1	2.85	0.65
1:A:225:LEU:HB2	1:A:233:PHE:CZ	2.31	0.65
1:A:253:ILE:HG21	1:A:281:LEU:HB3	1.76	0.65
1:A:264:SER:HB2	1:A:271:ARG:CG	2.24	0.65
1:A:557:LYS:O	2:B:605:PHE:CE2	2.45	0.65
1:A:557:LYS:HG2	2:B:606:ASP:HA	1.76	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:609:LEU:O	1:A:612:GLU:N	2.29	0.65
2:B:123:LEU:HD22	2:B:138:ALA:CA	2.27	0.65
2:B:208:ILE:CG1	2:B:236:ILE:HD13	2.19	0.65
2:B:212:VAL:CG1	2:B:248:LEU:HD23	2.26	0.65
2:B:252:LEU:HD13	2:B:302:PHE:CG	2.14	0.65
2:B:331:PRO:HB3	2:B:369:LEU:CD1	2.27	0.65
2:B:334:MET:HE2	2:B:339:PHE:CD1	2.31	0.65
2:B:346:THR:O	2:B:349:MET:N	2.29	0.65
2:B:353:GLN:HG2	3:M:48:ASP:C	2.17	0.65
2:B:447:GLU:OE1	2:B:485:LYS:CG	2.41	0.65
2:B:545:ARG:HD2	2:B:602:ASP:CG	2.17	0.65
2:B:549:LEU:HD22	2:B:607:ILE:O	1.96	0.65
3:M:55:MET:O	3:M:57:GLY:N	2.30	0.65
3:M:229:LYS:HG2	3:M:230:LYS:HG3	1.79	0.65
3:M:379:LEU:HD12	3:M:379:LEU:N	2.10	0.65
4:S:108:SER:CB	4:S:149:ILE:CG2	2.73	0.65
1:A:132:LEU:CD2	1:A:169:MET:CG	2.62	0.65
1:A:170:LEU:HB3	1:A:206:LYS:HG3	1.77	0.65
1:A:557:LYS:CG	2:B:606:ASP:HB2	2.25	0.65
2:B:211:ALA:C	2:B:233:TYR:CZ	2.52	0.65
2:B:223:LEU:HD11	2:B:259:TYR:N	1.90	0.65
2:B:340:ILE:HD13	2:B:366:LEU:CD1	2.18	0.65
2:B:438:ARG:HA	2:B:441:GLN:NE2	2.11	0.65
3:M:214:LEU:CD1	3:M:256:VAL:CG2	2.74	0.65
3:M:241:HIS:HD2	3:M:475:GLN:H	1.42	0.65
3:M:340:LEU:CB	3:M:411:LEU:HG	2.14	0.65
4:S:24:ASP:OD2	4:S:27:LYS:HD2	1.96	0.65
1:A:128:LEU:HD13	1:A:150:LEU:HD21	1.22	0.65
1:A:189:PHE:HB2	1:A:225:LEU:CD2	2.26	0.65
1:A:485:PRO:O	1:A:488:ARG:CG	2.43	0.65
1:A:638:LEU:O	2:B:486:HIS:CE1	2.50	0.65
2:B:90:ILE:CG1	2:B:98:LYS:HD3	2.27	0.65
2:B:178:ILE:CG2	2:B:214:ALA:C	2.64	0.65
2:B:274:PRO:N	2:B:295:ASN:ND2	2.44	0.65
3:M:66:PHE:CB	3:M:77:LEU:HD21	2.03	0.65
3:M:242:GLY:HA3	3:M:444:ALA:CB	2.25	0.65
1:A:67:LYS:CB	4:S:166:LYS:HA	2.27	0.65
1:A:78:GLU:O	1:A:80:TYR:O	2.13	0.65
1:A:88:ASN:HB2	1:A:120:ILE:HD12	1.73	0.65
1:A:105:VAL:HG23	4:S:167:ILE:CB	2.26	0.65
1:A:211:ASP:OD2	4:S:148:ARG:CD	2.43	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:416:ILE:HG22	1:A:417:PRO:O	1.96	0.65
1:A:420:ILE:HG23	1:A:421:PRO:HD2	1.77	0.65
1:A:638:LEU:O	2:B:518:ILE:CD1	2.44	0.65
2:B:5:ILE:CA	2:B:8:ILE:HD12	2.26	0.65
2:B:67:ILE:HD13	2:B:103:LEU:HB3	1.77	0.65
2:B:159:LYS:HE3	2:B:191:GLU:CD	2.16	0.65
2:B:178:ILE:CG2	2:B:214:ALA:HA	2.24	0.65
2:B:479:VAL:HG21	2:B:486:HIS:NE2	2.11	0.65
2:B:490:ILE:CG1	2:B:518:ILE:HG21	2.23	0.65
4:S:105:PHE:HZ	4:S:128:LEU:CD1	2.08	0.65
1:A:103:LYS:HE3	1:A:131:ASP:CG	2.05	0.65
1:A:412:LYS:O	1:A:413:SER:C	2.31	0.65
1:A:530:ASN:OD1	1:A:577:VAL:HG21	1.96	0.65
2:B:177:ILE:HG22	2:B:196:LEU:HD11	1.79	0.65
2:B:277:CYS:HA	2:B:292:GLU:HA	1.77	0.65
2:B:351:GLU:O	3:M:48:ASP:CB	2.42	0.65
2:B:430:ILE:HG23	2:B:470:ALA:HB2	1.78	0.65
3:M:5:PHE:HB2	3:M:125:PHE:CE2	2.31	0.65
3:M:216:VAL:O	3:M:216:VAL:HG23	1.96	0.65
3:M:222:PHE:CE1	3:M:240:ILE:HG21	2.32	0.65
3:M:327:PHE:CE1	3:M:336:ASP:OD2	2.49	0.65
3:M:433:VAL:HG13	3:M:433:VAL:O	1.97	0.65
2:B:105:LEU:CB	2:B:145:MET:CE	2.65	0.65
2:B:178:ILE:CG2	2:B:179:LYS:N	2.51	0.65
2:B:435:SER:O	2:B:438:ARG:N	2.19	0.65
2:B:575:ASN:C	2:B:576:GLN:O	2.26	0.65
1:A:196:LEU:HD22	1:A:196:LEU:C	2.15	0.65
1:A:220:SER:HB2	4:S:142:ILE:HA	1.79	0.65
1:A:291:ILE:N	1:A:291:ILE:HD12	2.12	0.65
1:A:364:ASP:HB3	1:A:367:ILE:HD12	1.79	0.65
1:A:384:LEU:HD21	1:A:441:TYR:CE2	2.28	0.65
1:A:480:LEU:C	1:A:480:LEU:HD13	2.17	0.65
1:A:570:LYS:O	1:A:571:ARG:HB2	1.97	0.65
2:B:37:TYR:HH	2:B:46:GLN:HE21	0.83	0.65
2:B:158:VAL:HG13	2:B:177:ILE:CD1	2.19	0.65
2:B:196:LEU:HB3	2:B:229:HIS:ND1	2.10	0.65
2:B:567:GLN:HG2	2:B:569:THR:OG1	1.97	0.65
3:M:54:SER:HB2	3:M:66:PHE:HD1	1.62	0.65
3:M:101:LEU:HG	3:M:106:LYS:CG	2.26	0.65
3:M:271:SER:HB3	3:M:301:GLU:HG3	1.79	0.65
3:M:316:ARG:O	3:M:317:MET:C	2.34	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:319:SER:HB2	3:M:347:PHE:N	2.11	0.65
1:A:97:SER:H	4:S:166:LYS:HZ1	1.45	0.65
1:A:104:ARG:NE	4:S:160:ALA:HB2	2.11	0.65
1:A:136:GLY:O	1:A:137:ASN:C	2.27	0.65
2:B:50:LEU:HB3	2:B:62:ALA:CB	2.23	0.65
2:B:50:LEU:CG	2:B:62:ALA:HB2	2.27	0.65
2:B:141:ALA:C	2:B:143:SER:N	2.46	0.65
2:B:350:THR:CB	2:B:352:ASN:HD21	2.10	0.65
3:M:224:VAL:HB	3:M:226:PHE:CE1	2.32	0.65
3:M:244:VAL:HG13	3:M:472:TYR:HE2	1.50	0.65
1:A:185:LEU:HB3	1:A:189:PHE:CE2	2.32	0.64
1:A:638:LEU:HD13	2:B:515:PHE:O	1.98	0.64
2:B:73:ASP:OD1	2:B:111:ASN:ND2	2.28	0.64
2:B:157:THR:O	2:B:159:LYS:N	2.30	0.64
2:B:227:HIS:CE1	2:B:292:GLU:HG2	2.32	0.64
2:B:306:LEU:HB3	2:B:321:CYS:HB3	1.78	0.64
2:B:578:PRO:HB3	2:B:579:PRO:HD3	1.80	0.64
3:M:371:GLU:O	3:M:419:ASN:OD1	2.14	0.64
4:S:3:HIS:NE2	4:S:90:ASP:OD2	2.30	0.64
1:A:145:ILE:HD13	4:S:156:LEU:HD22	1.78	0.64
1:A:150:LEU:HD22	1:A:158:LEU:HD11	1.79	0.64
2:B:196:LEU:C	2:B:229:HIS:CE1	2.69	0.64
2:B:212:VAL:HG23	2:B:233:TYR:CE2	2.33	0.64
2:B:344:VAL:HG22	2:B:381:PHE:CZ	2.32	0.64
2:B:519:ALA:HB1	2:B:555:LEU:HD12	1.80	0.64
2:B:549:LEU:CG	2:B:614:ILE:HD12	2.27	0.64
3:M:3:LEU:HA	3:M:80:THR:HA	1.79	0.64
3:M:101:LEU:C	3:M:106:LYS:CA	2.64	0.64
1:A:323:CYS:SG	1:A:338:PHE:CE1	2.74	0.64
2:B:90:ILE:HD11	2:B:123:LEU:CD2	2.27	0.64
2:B:344:VAL:CG2	2:B:363:ILE:CD1	2.74	0.64
2:B:396:ILE:HG21	2:B:432:ALA:HA	1.78	0.64
2:B:513:TRP:CG	2:B:551:LEU:CD2	2.76	0.64
3:M:255:LEU:O	3:M:454:ILE:HG13	1.97	0.64
3:M:256:VAL:HG12	3:M:290:PHE:CB	2.23	0.64
3:M:352:GLN:HB3	3:M:403:THR:O	1.97	0.64
4:S:69:ASN:ND2	4:S:71:GLU:O	2.31	0.64
4:S:135:ILE:C	4:S:141:VAL:HG22	2.18	0.64
1:A:552:ILE:O	1:A:556:VAL:HG23	1.97	0.64
1:A:585:PHE:CD2	1:A:603:VAL:HG12	2.32	0.64
1:A:634:ASN:ND2	2:B:554:LYS:N	2.37	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:158:VAL:CG2	2:B:177:ILE:HG13	2.27	0.64
2:B:275:ARG:HB3	2:B:294:VAL:CG1	2.19	0.64
3:M:49:ASP:HA	3:M:75:TRP:CZ3	2.32	0.64
3:M:250:LEU:HD13	3:M:254:PRO:CG	2.27	0.64
3:M:350:VAL:CG1	3:M:442:GLN:HB2	2.26	0.64
1:A:67:LYS:HB2	4:S:166:LYS:N	2.11	0.64
1:A:96:SER:HB2	1:A:127:LEU:CD1	2.08	0.64
1:A:239:LEU:O	1:A:242:GLU:O	2.14	0.64
2:B:20:ARG:HH11	2:B:21:GLU:HG2	1.57	0.64
2:B:79:VAL:HG21	2:B:108:PHE:CZ	2.33	0.64
2:B:123:LEU:CD2	2:B:138:ALA:HB1	2.27	0.64
2:B:214:ALA:C	2:B:216:LYS:N	2.48	0.64
2:B:316:THR:CG2	3:M:90:PHE:HE2	1.75	0.64
2:B:346:THR:HG22	2:B:350:THR:HG21	1.78	0.64
3:M:74:TYR:OH	3:M:97:ASP:HB3	1.98	0.64
3:M:214:LEU:O	3:M:214:LEU:HD23	1.97	0.64
3:M:257:ALA:HB2	3:M:455:VAL:CG2	2.28	0.64
4:S:8:PHE:HE2	4:S:86:THR:OG1	1.81	0.64
1:A:91:ILE:HG21	1:A:110:ALA:HB2	1.80	0.64
1:A:103:LYS:C	1:A:107:TYR:CD1	2.70	0.64
2:B:87:VAL:HG13	2:B:122:SER:HG	1.57	0.64
2:B:266:VAL:HA	2:B:289:PRO:HB2	1.79	0.64
2:B:513:TRP:C	2:B:551:LEU:HD22	2.17	0.64
2:B:559:ASP:HB3	2:B:563:PHE:CG	2.33	0.64
3:M:96:ILE:CG2	3:M:125:PHE:CE1	2.81	0.64
3:M:260:LEU:HD22	3:M:449:VAL:CG2	2.24	0.64
1:A:119:ASP:O	1:A:123:LEU:HG	1.97	0.64
1:A:513:ARG:HB3	1:A:550:VAL:HG11	1.78	0.64
1:A:575:LYS:HG3	1:A:611:LEU:HG	1.77	0.64
2:B:29:LYS:HD3	2:B:30:LEU:H	1.62	0.64
2:B:30:LEU:C	2:B:30:LEU:HD12	2.18	0.64
2:B:219:TYR:HD1	2:B:226:LEU:CD2	1.83	0.64
2:B:306:LEU:HD12	2:B:325:LEU:CD2	2.28	0.64
3:M:217:ASP:CB	3:M:471:LYS:HA	2.27	0.64
1:A:255:ARG:O	1:A:257:LEU:N	2.31	0.64
1:A:405:THR:OG1	1:A:406:GLY:N	2.27	0.64
1:A:557:LYS:HG2	2:B:606:ASP:CA	2.28	0.64
1:A:581:LEU:HD23	1:A:607:LEU:HD22	1.80	0.64
1:A:631:SER:HB3	2:B:557:SER:HB2	1.74	0.64
2:B:132:SER:HB2	2:B:166:SER:CB	2.27	0.64
2:B:396:ILE:HD11	2:B:418:TYR:OH	1.92	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:214:LEU:O	3:M:467:TYR:CB	2.44	0.64
3:M:356:LEU:HD21	3:M:358:ILE:CG1	2.27	0.64
1:A:125:THR:C	1:A:127:LEU:N	2.50	0.64
1:A:436:CYS:SG	1:A:450:TYR:CE2	2.87	0.64
1:A:609:LEU:CD2	1:A:628:VAL:CB	2.76	0.64
2:B:79:VAL:HB	2:B:108:PHE:CD1	2.33	0.64
2:B:277:CYS:HA	2:B:292:GLU:CB	2.26	0.64
2:B:461:HIS:O	2:B:462:ASN:O	2.16	0.64
2:B:508:ARG:O	2:B:509:ALA:O	2.16	0.64
2:B:534:ILE:HD11	2:B:595:VAL:HG23	1.80	0.64
2:B:566:ALA:O	2:B:574:ASN:HB3	1.78	0.64
3:M:16:PHE:HE2	3:M:125:PHE:CE2	2.16	0.64
3:M:316:ARG:O	3:M:318:ASN:OD1	2.15	0.64
4:S:53:THR:CG2	4:S:69:ASN:N	2.61	0.64
4:S:157:ASN:O	4:S:161:GLU:HG3	1.97	0.64
1:A:67:LYS:HB3	4:S:166:LYS:CA	2.28	0.64
1:A:219:VAL:CG2	1:A:256:LEU:CD2	2.69	0.64
2:B:2:VAL:CG1	2:B:6:HIS:CD2	2.61	0.64
2:B:8:ILE:O	2:B:12:LEU:CD1	2.45	0.64
2:B:47:LEU:CB	2:B:66:ILE:HG12	2.28	0.64
2:B:50:LEU:HD23	2:B:62:ALA:N	2.13	0.64
2:B:136:CYS:O	2:B:172:GLU:CB	2.45	0.64
2:B:309:LEU:CG	2:B:317:VAL:HG11	2.27	0.64
3:M:316:ARG:HG3	3:M:322:LEU:HD13	1.80	0.64
3:M:375:LYS:HE3	3:M:418:GLU:OE1	1.97	0.64
4:S:8:PHE:CE2	4:S:86:THR:OG1	2.50	0.64
1:A:105:VAL:CG2	4:S:167:ILE:HA	2.28	0.63
1:A:484:VAL:O	1:A:486:SER:O	2.16	0.63
1:A:598:GLU:O	1:A:602:GLU:HG3	1.99	0.63
2:B:45:GLN:C	2:B:47:LEU:H	2.01	0.63
2:B:93:ASN:C	2:B:134:LEU:HD11	2.18	0.63
2:B:275:ARG:NE	2:B:275:ARG:HA	2.13	0.63
2:B:449:HIS:O	2:B:453:TRP:CD1	2.51	0.63
2:B:520:SER:O	2:B:523:PHE:HE2	1.81	0.63
3:M:45:SER:HA	3:M:47:SER:N	2.07	0.63
3:M:100:LEU:HD13	3:M:100:LEU:N	2.13	0.63
1:A:104:ARG:NH1	4:S:126:GLN:C	2.52	0.63
1:A:121:LEU:CD1	1:A:155:THR:CG2	2.77	0.63
1:A:196:LEU:O	1:A:197:ARG:C	2.31	0.63
1:A:297:CYS:O	1:A:298:ILE:C	2.32	0.63
1:A:451:ASN:OD1	1:A:480:LEU:HD12	1.98	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:63:MET:HA	2:B:66:ILE:HD12	1.78	0.63
2:B:83:PHE:HZ	2:B:119:SER:HB2	1.53	0.63
2:B:98:LYS:HB2	2:B:134:LEU:HD22	1.78	0.63
2:B:215:TYR:CD2	2:B:226:LEU:HD13	2.31	0.63
2:B:487:LEU:HD21	2:B:522:GLU:CB	2.25	0.63
2:B:545:ARG:HD2	2:B:602:ASP:HB2	1.79	0.63
3:M:100:LEU:H	3:M:100:LEU:HD13	1.61	0.63
3:M:215:TYR:HB2	3:M:467:TYR:C	2.18	0.63
3:M:241:HIS:CB	3:M:476:THR:HG23	2.29	0.63
1:A:67:LYS:C	4:S:166:LYS:CA	2.66	0.63
1:A:103:LYS:HZ2	1:A:131:ASP:HB2	1.63	0.63
1:A:322:PHE:HB3	1:A:330:LEU:HD21	1.79	0.63
2:B:157:THR:C	2:B:159:LYS:N	2.51	0.63
2:B:167:ALA:O	2:B:207:VAL:HG11	1.99	0.63
2:B:245:GLN:NE2	2:B:309:LEU:HD13	2.14	0.63
2:B:287:GLU:O	2:B:288:TYR:CD2	2.50	0.63
2:B:364:HIS:CE1	2:B:397:GLN:HB3	2.33	0.63
2:B:537:PHE:CD2	2:B:537:PHE:C	2.71	0.63
1:A:100:LEU:HD11	4:S:157:ASN:C	2.17	0.63
2:B:36:THR:O	2:B:40:GLN:CB	2.46	0.63
3:M:10:THR:HA	3:M:75:TRP:HE1	1.62	0.63
3:M:96:ILE:CG2	3:M:125:PHE:CZ	2.81	0.63
3:M:258:VAL:CA	3:M:452:ILE:HG13	2.28	0.63
3:M:265:ASN:HB3	3:M:309:GLN:CG	2.29	0.63
3:M:320:ILE:HG13	3:M:347:PHE:HD1	1.59	0.63
4:S:164:ASP:O	4:S:165:SER:C	2.20	0.63
1:A:104:ARG:HG2	4:S:126:GLN:OE1	1.97	0.63
1:A:139:ASP:OD1	1:A:177:ILE:HD11	1.98	0.63
1:A:156:PRO:O	1:A:160:ARG:HG3	1.98	0.63
1:A:175:PRO:HA	1:A:214:VAL:CG2	2.29	0.63
2:B:178:ILE:HG22	2:B:217:GLU:HB2	1.74	0.63
2:B:243:TRP:CZ3	3:M:91:THR:CA	2.75	0.63
2:B:274:PRO:CG	2:B:295:ASN:HD21	2.05	0.63
2:B:319:LEU:O	2:B:322:CYS:N	2.31	0.63
2:B:360:LEU:HD11	2:B:391:ALA:HB1	1.81	0.63
2:B:475:ILE:CG2	2:B:514:LEU:CD2	2.76	0.63
3:M:125:PHE:O	3:M:129:VAL:HG21	1.93	0.63
3:M:347:PHE:CE2	3:M:350:VAL:HB	2.34	0.63
1:A:68:THR:HA	4:S:167:ILE:CA	2.28	0.63
1:A:185:LEU:O	1:A:189:PHE:CG	2.51	0.63
1:A:571:ARG:NH2	1:A:573:GLU:OE1	2.30	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:90:ILE:HD11	2:B:123:LEU:HD21	1.80	0.63
2:B:295:ASN:O	2:B:296:ASP:HB2	1.99	0.63
2:B:334:MET:HB2	2:B:369:LEU:CD2	2.27	0.63
2:B:396:ILE:HD11	2:B:418:TYR:CE2	2.25	0.63
2:B:452:LYS:HZ3	2:B:456:ASP:CG	2.01	0.63
2:B:515:PHE:O	2:B:516:GLY:C	2.33	0.63
2:B:526:CYS:N	2:B:527:PRO:HD3	2.14	0.63
3:M:354:ASP:CB	3:M:440:ILE:HD12	2.28	0.63
3:M:360:LEU:HD13	3:M:433:VAL:CB	2.27	0.63
4:S:6:LEU:CD2	4:S:32:LEU:HD22	2.24	0.63
1:A:141:VAL:HG12	4:S:159:ALA:HB3	0.73	0.63
1:A:179:LYS:CD	4:S:143:GLU:CG	2.74	0.63
1:A:313:MET:HA	1:A:348:PHE:CZ	2.33	0.63
1:A:563:CYS:SG	1:A:621:LEU:HD11	2.39	0.63
1:A:575:LYS:HE3	1:A:611:LEU:HD21	1.80	0.63
1:A:581:LEU:CD2	1:A:607:LEU:CD1	2.77	0.63
2:B:362:ALA:O	2:B:366:LEU:HG	1.99	0.63
2:B:592:TYR:CE2	2:B:619:ASP:OD1	2.52	0.63
3:M:10:THR:HG23	3:M:11:LYS:N	2.13	0.63
3:M:215:TYR:CD2	3:M:470:ALA:N	2.64	0.63
1:A:96:SER:H	4:S:166:LYS:NZ	1.96	0.63
1:A:141:VAL:HG11	4:S:159:ALA:N	2.12	0.63
1:A:178:ARG:HB2	1:A:214:VAL:HG22	1.78	0.63
1:A:253:ILE:CD1	1:A:281:LEU:CD2	2.70	0.63
1:A:369:SER:HB2	1:A:424:TYR:CE2	2.32	0.63
1:A:566:PHE:HZ	1:A:618:THR:O	1.82	0.63
1:A:607:LEU:O	1:A:609:LEU:N	2.32	0.63
2:B:268:LYS:CA	2:B:276:SER:HB2	2.29	0.63
2:B:293:VAL:O	2:B:299:LEU:HG	1.99	0.63
2:B:481:LYS:C	2:B:483:PRO:HD3	2.18	0.63
3:M:306:LEU:CD2	3:M:317:MET:CE	2.77	0.63
4:S:7:ILE:CD1	4:S:16:LEU:HD23	2.28	0.63
4:S:20:TYR:HD2	4:S:21:THR:HG23	1.62	0.63
4:S:80:TYR:HB2	4:S:106:VAL:HG11	1.80	0.63
1:A:100:LEU:N	4:S:162:SER:CB	2.04	0.63
1:A:102:GLN:CD	4:S:166:LYS:N	2.42	0.63
1:A:225:LEU:HB2	1:A:233:PHE:CE1	2.33	0.63
1:A:563:CYS:SG	1:A:621:LEU:HD12	2.37	0.63
1:A:581:LEU:HD23	1:A:607:LEU:HD11	1.81	0.63
2:B:4:SER:O	2:B:8:ILE:HD12	1.99	0.63
2:B:136:CYS:C	2:B:172:GLU:CB	2.67	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:431:MET:C	2:B:433:VAL:H	2.03	0.63
2:B:518:ILE:O	2:B:518:ILE:CD1	2.47	0.63
2:B:545:ARG:HD2	2:B:602:ASP:CB	2.28	0.63
3:M:99:ILE:CD1	3:M:99:ILE:H	2.12	0.63
3:M:245:ASP:N	3:M:472:TYR:CZ	2.66	0.63
3:M:377:LYS:HE3	3:M:416:GLU:CB	2.28	0.63
3:M:437:TYR:HD1	3:M:479:PHE:CZ	2.17	0.63
1:A:144:GLY:HA2	1:A:180:LYS:HB3	1.78	0.62
2:B:90:ILE:HD11	2:B:102:HIS:NE2	2.14	0.62
2:B:124:GLN:HA	2:B:157:THR:HG1	1.63	0.62
2:B:154:ILE:CD1	2:B:180:LEU:CG	2.77	0.62
2:B:154:ILE:O	2:B:158:VAL:HG23	1.98	0.62
2:B:353:GLN:HG3	3:M:47:SER:CB	2.29	0.62
2:B:353:GLN:CG	3:M:47:SER:HB2	2.29	0.62
3:M:320:ILE:HG22	3:M:321:GLY:N	2.14	0.62
4:S:8:PHE:CG	4:S:36:TYR:CE2	2.83	0.62
4:S:47:GLN:HE22	4:S:78:LYS:CA	2.11	0.62
4:S:137:GLN:C	4:S:140:MET:H	2.02	0.62
1:A:100:LEU:CA	4:S:162:SER:HB2	2.22	0.62
1:A:117:ASP:OD2	1:A:120:ILE:HG12	2.00	0.62
1:A:142:LYS:H	4:S:159:ALA:HB2	1.53	0.62
1:A:237:SER:CB	1:A:270:LEU:HD13	2.23	0.62
1:A:342:GLY:O	1:A:343:LYS:C	2.35	0.62
2:B:63:MET:HE1	2:B:104:TYR:CB	2.29	0.62
2:B:435:SER:C	2:B:437:SER:N	2.53	0.62
2:B:512:VAL:HG12	2:B:551:LEU:HB2	1.82	0.62
2:B:567:GLN:CA	2:B:569:THR:OG1	2.47	0.62
3:M:66:PHE:CE2	3:M:77:LEU:O	2.53	0.62
3:M:104:PHE:HE1	3:M:113:LYS:HE2	0.77	0.62
3:M:272:LEU:CD2	3:M:288:ILE:HD13	2.30	0.62
1:A:80:TYR:CD1	1:A:82:PHE:CE2	2.88	0.62
1:A:180:LYS:HE3	4:S:156:LEU:HD11	1.81	0.62
1:A:240:LEU:O	1:A:242:GLU:O	2.18	0.62
2:B:157:THR:O	2:B:160:LYS:N	2.32	0.62
2:B:182:ARG:HD2	2:B:217:GLU:CG	2.29	0.62
2:B:307:ASN:OD1	2:B:339:PHE:HE2	1.77	0.62
2:B:416:LYS:HD2	2:B:453:TRP:CD2	2.34	0.62
2:B:500:GLN:O	2:B:508:ARG:NH2	2.32	0.62
3:M:10:THR:CA	3:M:75:TRP:NE1	2.62	0.62
1:A:76:TYR:OH	4:S:125:TRP:CZ3	2.39	0.62
1:A:96:SER:N	4:S:166:LYS:HZ1	1.97	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:101:GLN:N	4:S:162:SER:H	1.96	0.62
1:A:176:TYR:CG	4:S:155:GLU:HG2	2.34	0.62
1:A:212:ILE:HB	4:S:145:ASN:HD22	1.63	0.62
1:A:255:ARG:HH21	4:S:135:ILE:CG2	1.99	0.62
1:A:403:LEU:HD21	1:A:421:PRO:CA	2.28	0.62
1:A:513:ARG:CG	1:A:550:VAL:HG21	2.29	0.62
2:B:136:CYS:CA	2:B:172:GLU:HB2	2.28	0.62
2:B:153:ILE:C	2:B:155:LEU:H	2.03	0.62
3:M:242:GLY:CA	3:M:444:ALA:CB	2.77	0.62
4:S:48:SER:HA	4:S:77:TYR:HB3	1.81	0.62
4:S:164:ASP:HA	4:S:167:ILE:HB	1.79	0.62
1:A:141:VAL:CB	4:S:159:ALA:HB2	2.29	0.62
2:B:135:ARG:NH2	2:B:164:ASP:CB	2.62	0.62
2:B:196:LEU:O	2:B:199:LEU:HB2	1.99	0.62
2:B:196:LEU:CB	2:B:229:HIS:HD1	2.13	0.62
2:B:316:THR:HG23	3:M:90:PHE:CE2	2.31	0.62
2:B:319:LEU:CD1	2:B:358:MET:SD	2.78	0.62
2:B:472:VAL:HG11	2:B:510:GLY:HA2	1.68	0.62
3:M:407:THR:O	3:M:409:PRO:HD3	1.98	0.62
1:A:175:PRO:O	1:A:214:VAL:HG22	2.00	0.62
3:M:62:VAL:O	3:M:62:VAL:HG12	2.00	0.62
3:M:224:VAL:CA	3:M:479:PHE:HA	2.29	0.62
3:M:356:LEU:C	3:M:356:LEU:HD23	2.19	0.62
4:S:85:PHE:CE1	4:S:106:VAL:HG22	2.34	0.62
1:A:80:TYR:HB3	1:A:82:PHE:CZ	2.34	0.62
1:A:99:LYS:HZ2	4:S:164:ASP:HB3	1.65	0.62
1:A:212:ILE:HB	1:A:247:ILE:HD13	1.78	0.62
1:A:480:LEU:O	1:A:483:LYS:O	2.18	0.62
1:A:623:MET:C	2:B:617:LEU:HD21	2.20	0.62
2:B:106:LEU:CD2	2:B:144:ASP:CB	2.76	0.62
2:B:177:ILE:HG21	2:B:196:LEU:HD11	1.79	0.62
2:B:309:LEU:CG	2:B:317:VAL:CG1	2.77	0.62
2:B:511:ILE:O	2:B:515:PHE:HD1	1.83	0.62
2:B:588:ILE:HG23	2:B:618:PHE:CE1	2.34	0.62
3:M:216:VAL:HG11	3:M:452:ILE:HG12	1.81	0.62
3:M:225:VAL:HA	3:M:480:GLN:H	1.65	0.62
3:M:362:PHE:O	3:M:364:VAL:HG13	2.00	0.62
4:S:75:ILE:CG2	4:S:86:THR:HG23	2.26	0.62
1:A:101:GLN:O	1:A:104:ARG:N	2.29	0.62
1:A:589:SER:O	1:A:597:GLN:NE2	2.32	0.62
2:B:47:LEU:HD13	2:B:66:ILE:HA	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:143:SER:HB3	2:B:175:LEU:C	2.18	0.62
2:B:486:HIS:HA	2:B:489:ILE:HD12	1.82	0.62
1:A:67:LYS:HB3	4:S:166:LYS:CG	2.29	0.62
1:A:91:ILE:CG2	1:A:110:ALA:HB2	2.30	0.62
1:A:182:ILE:CG2	1:A:218:ALA:HA	2.29	0.62
1:A:341:ILE:O	1:A:344:ILE:HB	1.99	0.62
1:A:408:ILE:CG2	4:S:64:ASN:O	2.34	0.62
2:B:360:LEU:HD13	2:B:391:ALA:O	2.00	0.62
2:B:476:ARG:CA	2:B:514:LEU:HD13	2.30	0.62
2:B:501:THR:C	2:B:508:ARG:NH2	2.53	0.62
3:M:4:SER:OG	3:M:6:TYR:CZ	2.51	0.62
3:M:24:ALA:O	3:M:25:PRO:C	2.34	0.62
3:M:222:PHE:CB	3:M:479:PHE:CE1	2.82	0.62
3:M:320:ILE:HG21	3:M:439:TYR:CZ	2.35	0.62
4:S:32:LEU:O	4:S:36:TYR:CD1	2.53	0.62
4:S:47:GLN:OE1	4:S:79:ASN:CA	2.42	0.62
1:A:80:TYR:HB3	1:A:82:PHE:CE2	2.33	0.62
1:A:326:GLN:CA	1:A:331:ARG:HH21	2.11	0.62
1:A:556:VAL:CG2	1:A:603:VAL:HG11	2.29	0.62
2:B:70:MET:HE2	2:B:104:TYR:HA	1.80	0.62
2:B:545:ARG:HD3	2:B:602:ASP:CB	2.17	0.62
2:B:545:ARG:NH1	2:B:602:ASP:HA	2.14	0.62
3:M:1:MET:CA	3:M:81:SER:OG	2.48	0.62
3:M:215:TYR:CD1	3:M:468:LYS:HG2	2.35	0.62
3:M:223:HIS:CD2	3:M:478:ASN:CA	2.82	0.62
3:M:449:VAL:HG11	3:M:452:ILE:CG1	2.30	0.62
4:S:9:ASN:OD1	4:S:13:GLN:CA	2.46	0.62
1:A:212:ILE:CB	1:A:247:ILE:HD13	2.30	0.61
1:A:399:ASP:O	1:A:420:ILE:N	2.33	0.61
2:B:216:LYS:HB2	2:B:251:LEU:CG	2.30	0.61
2:B:302:PHE:CD2	2:B:328:LEU:HD11	2.35	0.61
2:B:336:ASN:CG	2:B:338:LYS:HB2	2.21	0.61
2:B:512:VAL:HG11	2:B:548:ILE:CA	2.11	0.61
2:B:362:ALA:O	2:B:366:LEU:N	2.28	0.61
3:M:214:LEU:N	3:M:467:TYR:HB2	2.15	0.61
3:M:215:TYR:CE1	3:M:468:LYS:CG	2.82	0.61
3:M:386:PHE:CD1	3:M:386:PHE:O	2.54	0.61
3:M:469:GLY:C	3:M:470:ALA:O	2.38	0.61
1:A:182:ILE:O	1:A:221:VAL:CG2	2.47	0.61
1:A:183:THR:C	4:S:137:GLN:OE1	2.39	0.61
1:A:630:PRO:CG	2:B:614:ILE:HG23	2.30	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:21:GLU:O	2:B:24:ALA:HB3	2.01	0.61
2:B:162:VAL:C	2:B:164:ASP:N	2.53	0.61
2:B:463:LEU:O	2:B:468:LEU:HD11	2.00	0.61
2:B:508:ARG:HB2	2:B:544:THR:HG21	1.78	0.61
2:B:520:SER:O	2:B:523:PHE:CE2	2.53	0.61
3:M:233:LEU:CD2	3:M:325:LEU:H	2.13	0.61
3:M:258:VAL:HG22	3:M:452:ILE:HG12	1.81	0.61
3:M:267:ILE:HG23	3:M:445:SER:OG	1.99	0.61
3:M:338:PHE:HE2	3:M:415:ILE:HG13	1.55	0.61
4:S:3:HIS:O	4:S:20:TYR:N	2.31	0.61
4:S:17:VAL:HG22	4:S:19:PHE:CE1	2.34	0.61
1:A:259:LEU:CD1	4:S:140:MET:HE1	2.31	0.61
1:A:312:ALA:O	1:A:315:CYS:HB2	2.00	0.61
1:A:556:VAL:HG22	1:A:603:VAL:CG1	2.30	0.61
1:A:589:SER:CB	1:A:601:VAL:HG22	2.30	0.61
2:B:276:SER:C	2:B:289:PRO:HG2	2.14	0.61
2:B:311:TYR:O	2:B:312:SER:C	2.38	0.61
3:M:44:ASP:C	3:M:46:SER:N	2.53	0.61
3:M:69:ILE:HD12	3:M:90:PHE:CE2	2.25	0.61
3:M:224:VAL:CG1	3:M:226:PHE:CE1	2.84	0.61
3:M:242:GLY:N	3:M:444:ALA:CB	2.64	0.61
3:M:323:MET:CE	3:M:342:LEU:HD23	2.27	0.61
3:M:356:LEU:CD2	3:M:358:ILE:CG1	2.77	0.61
3:M:478:ASN:O	3:M:479:PHE:C	2.37	0.61
1:A:273:LYS:O	1:A:276:PRO:HD2	2.01	0.61
1:A:630:PRO:CG	2:B:614:ILE:CG2	2.78	0.61
2:B:133:GLU:O	2:B:168:MET:HE1	2.00	0.61
2:B:534:ILE:C	2:B:536:ASN:N	2.53	0.61
3:M:121:ILE:O	3:M:125:PHE:HD1	1.81	0.61
3:M:265:ASN:HA	3:M:313:SER:OG	2.00	0.61
4:S:53:THR:CG2	4:S:57:LEU:CB	2.77	0.61
1:A:328:PRO:O	1:A:329:ASN:C	2.25	0.61
1:A:439:ASP:O	1:A:440:ASN:HB2	2.00	0.61
2:B:44:PRO:HB3	2:B:82:TYR:OH	2.00	0.61
2:B:178:ILE:HB	2:B:214:ALA:CB	2.29	0.61
2:B:340:ILE:HG12	2:B:373:LEU:HD23	1.82	0.61
2:B:355:ASN:O	2:B:359:LEU:CD2	2.48	0.61
2:B:498:THR:HG21	2:B:532:ARG:HG3	1.82	0.61
2:B:556:LEU:HD11	2:B:592:TYR:HB2	1.82	0.61
3:M:244:VAL:HG13	3:M:472:TYR:OH	1.99	0.61
3:M:443:SER:CB	3:M:447:ILE:CA	2.79	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:104:ARG:HG3	1:A:145:ILE:HD12	1.81	0.61
1:A:224:GLU:CD	4:S:138:GLY:O	2.39	0.61
1:A:266:VAL:O	1:A:267:GLU:HB2	2.01	0.61
1:A:384:LEU:HG	1:A:385:LYS:N	2.15	0.61
1:A:573:GLU:O	1:A:577:VAL:HG23	2.00	0.61
2:B:101:ILE:O	2:B:104:TYR:HB3	2.00	0.61
2:B:158:VAL:O	2:B:195:ILE:HD13	1.99	0.61
2:B:162:VAL:C	2:B:164:ASP:H	2.04	0.61
2:B:237:ILE:CG1	2:B:309:LEU:HD21	2.31	0.61
2:B:267:ASP:N	2:B:289:PRO:HG3	2.14	0.61
2:B:353:GLN:CB	3:M:47:SER:O	2.48	0.61
3:M:451:ALA:O	3:M:452:ILE:HD12	2.00	0.61
4:S:108:SER:O	4:S:112:CYS:SG	2.58	0.61
1:A:369:SER:HB2	1:A:424:TYR:HE2	1.66	0.61
1:A:376:GLU:H	1:A:376:GLU:CD	2.03	0.61
1:A:634:ASN:HA	2:B:516:GLY:HA3	1.83	0.61
2:B:127:LEU:HD23	2:B:135:ARG:O	2.01	0.61
2:B:274:PRO:HD2	2:B:295:ASN:ND2	2.13	0.61
2:B:498:THR:HG22	2:B:532:ARG:CB	2.31	0.61
3:M:118:TYR:C	3:M:118:TYR:CD2	2.73	0.61
3:M:215:TYR:HE1	3:M:468:LYS:HG2	1.60	0.61
3:M:461:GLY:C	3:M:462:LYS:O	2.23	0.61
1:A:100:LEU:CA	4:S:163:THR:HG23	2.27	0.61
1:A:166:LEU:O	1:A:170:LEU:CD2	2.48	0.61
1:A:180:LYS:HE3	4:S:156:LEU:CD2	2.29	0.61
1:A:186:PHE:CD1	1:A:224:GLU:HG2	2.35	0.61
1:A:244:LEU:O	1:A:246:THR:N	2.34	0.61
2:B:204:ASP:HB3	2:B:207:VAL:HG23	1.82	0.61
2:B:271:GLU:C	2:B:273:SER:N	2.44	0.61
2:B:339:PHE:O	2:B:343:LEU:CB	2.49	0.61
2:B:356:LYS:O	2:B:359:LEU:HB2	2.00	0.61
3:M:6:TYR:CA	3:M:16:PHE:O	2.49	0.61
3:M:217:ASP:HB3	3:M:471:LYS:CA	2.31	0.61
3:M:244:VAL:HG22	3:M:472:TYR:CE2	2.36	0.61
3:M:435:LEU:H	3:M:479:PHE:HB2	1.66	0.61
2:B:38:TYR:CD1	2:B:42:ILE:HA	2.35	0.61
2:B:127:LEU:HB2	2:B:157:THR:CG2	2.11	0.61
2:B:216:LYS:HB2	2:B:251:LEU:HD22	1.82	0.61
2:B:216:LYS:HG3	2:B:251:LEU:HA	1.83	0.61
2:B:362:ALA:HB1	2:B:366:LEU:HD11	1.81	0.61
2:B:512:VAL:C	2:B:551:LEU:CD1	2.46	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:9:ASP:CA	3:M:75:TRP:CD1	2.83	0.61
3:M:9:ASP:OD2	3:M:13:LYS:HB3	2.01	0.61
3:M:220:GLU:HG3	3:M:439:TYR:HB2	1.82	0.61
3:M:220:GLU:HG2	3:M:439:TYR:O	2.01	0.61
3:M:228:LYS:NZ	3:M:327:PHE:H	1.98	0.61
3:M:479:PHE:CD2	3:M:479:PHE:N	2.59	0.61
4:S:53:THR:CG2	4:S:68:VAL:N	2.62	0.61
1:A:582:ILE:HD11	1:A:608:ARG:CA	2.30	0.60
1:A:627:GLU:HG2	2:B:617:LEU:CA	1.96	0.60
2:B:486:HIS:NE2	2:B:518:ILE:CB	2.48	0.60
2:B:497:LEU:O	2:B:499:VAL:N	2.34	0.60
2:B:563:PHE:O	2:B:566:ALA:C	2.38	0.60
3:M:84:LYS:O	3:M:88:ASP:HB3	2.01	0.60
4:S:14:PRO:O	4:S:15:ARG:CD	2.49	0.60
4:S:25:LEU:CB	4:S:26:PRO:HD3	2.24	0.60
2:B:77:ILE:C	2:B:79:VAL:N	2.48	0.60
2:B:124:GLN:HA	2:B:127:LEU:HD12	1.82	0.60
2:B:178:ILE:CG1	2:B:214:ALA:CA	2.70	0.60
2:B:278:PRO:C	2:B:288:TYR:HB3	2.19	0.60
2:B:343:LEU:HD21	2:B:362:ALA:CB	2.18	0.60
2:B:354:GLY:O	2:B:358:MET:CG	2.47	0.60
2:B:435:SER:C	2:B:437:SER:H	2.04	0.60
2:B:602:ASP:O	2:B:608:ARG:NE	2.33	0.60
3:M:380:ARG:CZ	3:M:412:ARG:HD2	2.31	0.60
4:S:47:GLN:OE1	4:S:84:TYR:HD2	1.84	0.60
1:A:111:SER:HB3	1:A:152:THR:OG1	1.91	0.60
1:A:288:THR:CA	1:A:291:ILE:HD13	2.30	0.60
1:A:309:PHE:CE1	1:A:348:PHE:CZ	2.89	0.60
2:B:132:SER:HB2	2:B:169:VAL:HG23	1.82	0.60
2:B:279:LEU:HD12	2:B:288:TYR:CE1	2.35	0.60
2:B:393:ILE:HG23	2:B:431:MET:HB2	1.66	0.60
3:M:60:LEU:CD2	3:M:62:VAL:CG2	2.56	0.60
3:M:220:GLU:HG3	3:M:439:TYR:CB	2.31	0.60
4:S:6:LEU:CD1	4:S:32:LEU:HD13	2.32	0.60
1:A:186:PHE:CD2	1:A:186:PHE:C	2.75	0.60
1:A:270:LEU:HD12	1:A:274:LEU:HD23	1.84	0.60
1:A:436:CYS:SG	1:A:450:TYR:CE1	2.94	0.60
1:A:631:SER:CB	2:B:557:SER:HB3	2.12	0.60
2:B:47:LEU:HD22	2:B:66:ILE:HG13	0.63	0.60
2:B:57:ARG:O	2:B:60:ARG:HB3	2.01	0.60
2:B:90:ILE:HG12	2:B:98:LYS:CD	2.31	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:103:LEU:HD13	3:M:132:GLY:CA	2.28	0.60
2:B:178:ILE:HG23	2:B:218:CYS:H	1.63	0.60
2:B:231:ARG:O	2:B:233:TYR:N	2.34	0.60
2:B:303:LEU:CD2	2:B:325:LEU:HD23	2.31	0.60
3:M:65:TYR:CG	3:M:86:PRO:HB3	2.35	0.60
3:M:68:VAL:O	3:M:68:VAL:HG12	2.01	0.60
3:M:364:VAL:C	3:M:367:ALA:O	2.39	0.60
4:S:10:LYS:HA	4:S:84:TYR:CE1	2.36	0.60
1:A:156:PRO:O	1:A:157:SER:C	2.30	0.60
1:A:399:ASP:O	1:A:420:ILE:HB	2.01	0.60
1:A:624:LEU:HA	2:B:617:LEU:HD21	1.84	0.60
1:A:633:PHE:HE1	2:B:513:TRP:HE3	0.80	0.60
2:B:136:CYS:SG	2:B:168:MET:HG2	2.41	0.60
2:B:154:ILE:CG2	2:B:176:ALA:HB1	2.31	0.60
2:B:197:LYS:HA	2:B:229:HIS:HD2	1.53	0.60
2:B:219:TYR:HB3	2:B:223:LEU:HD23	0.61	0.60
2:B:319:LEU:CD1	2:B:358:MET:HG2	2.12	0.60
2:B:341:GLU:CG	2:B:345:ARG:HE	2.13	0.60
2:B:546:CYS:SG	2:B:607:ILE:HA	2.42	0.60
2:B:553:ALA:CB	2:B:614:ILE:HG21	2.28	0.60
2:B:589:SER:HG	2:B:618:PHE:HE2	1.43	0.60
2:B:592:TYR:CE2	2:B:618:PHE:CE2	2.89	0.60
1:A:121:LEU:HD11	1:A:155:THR:CG2	2.30	0.60
1:A:360:LEU:HD21	1:A:375:VAL:HG21	1.83	0.60
1:A:403:LEU:HD22	1:A:422:GLU:OE2	2.01	0.60
1:A:586:GLU:O	1:A:587:ASN:O	2.17	0.60
2:B:25:VAL:N	2:B:35:TYR:CD2	2.70	0.60
2:B:178:ILE:C	2:B:180:LEU:N	2.45	0.60
2:B:195:ILE:C	2:B:197:LYS:H	2.03	0.60
2:B:351:GLU:H	2:B:351:GLU:CD	2.04	0.60
2:B:374:PHE:HD2	2:B:402:LEU:HD11	1.63	0.60
2:B:417:TYR:O	2:B:421:SER:HB2	2.01	0.60
3:M:104:PHE:CE1	3:M:113:LYS:HE3	2.18	0.60
3:M:223:HIS:N	3:M:479:PHE:CE2	2.69	0.60
3:M:250:LEU:CD1	3:M:254:PRO:HG2	2.30	0.60
4:S:135:ILE:CG2	4:S:141:VAL:CG2	2.78	0.60
1:A:78:GLU:CD	1:A:113:SER:HB3	2.20	0.60
1:A:416:ILE:O	1:A:418:ILE:HG22	2.02	0.60
2:B:196:LEU:HB3	2:B:215:TYR:OH	2.02	0.60
2:B:208:ILE:CG2	2:B:236:ILE:HG21	2.31	0.60
2:B:431:MET:C	2:B:433:VAL:N	2.55	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:451:MET:HG3	2:B:489:ILE:HD11	1.83	0.60
2:B:582:ASP:O	2:B:584:SER:CA	2.49	0.60
3:M:235:LEU:HD23	3:M:235:LEU:C	2.22	0.60
1:A:88:ASN:O	1:A:90:HIS:N	2.33	0.60
1:A:96:SER:HB3	1:A:127:LEU:HD22	1.83	0.60
1:A:104:ARG:HG3	1:A:145:ILE:HG21	1.83	0.60
2:B:98:LYS:HD2	2:B:102:HIS:NE2	2.15	0.60
2:B:196:LEU:HA	2:B:199:LEU:HD12	1.83	0.60
2:B:493:LEU:HD21	2:B:511:ILE:HA	1.83	0.60
2:B:530:LEU:CD2	2:B:591:MET:CB	2.79	0.60
3:M:121:ILE:CG2	3:M:125:PHE:HE1	2.15	0.60
3:M:124:ILE:CG2	3:M:128:CYS:SG	2.90	0.60
3:M:271:SER:O	3:M:300:LEU:HA	2.01	0.60
3:M:454:ILE:HG22	3:M:464:THR:HG21	1.77	0.60
1:A:67:LYS:HD2	1:A:102:GLN:OE1	2.02	0.60
1:A:150:LEU:CD1	1:A:162:ILE:HG12	2.27	0.60
1:A:166:LEU:CD1	1:A:185:LEU:HD21	2.30	0.60
1:A:212:ILE:CD1	4:S:145:ASN:CG	2.62	0.60
1:A:292:TYR:O	1:A:295:VAL:HB	2.01	0.60
1:A:633:PHE:CG	2:B:550:VAL:CB	2.83	0.60
2:B:174:ALA:CB	2:B:211:ALA:HB2	2.30	0.60
2:B:274:PRO:N	2:B:295:ASN:HD21	1.99	0.60
2:B:286:ILE:O	2:B:286:ILE:HG23	2.00	0.60
2:B:562:ASN:HB3	2:B:580:TYR:O	2.02	0.60
3:M:99:ILE:HG22	3:M:103:TYR:CD1	2.37	0.60
3:M:247:ARG:N	3:M:470:ALA:HB2	2.17	0.60
1:A:139:ASP:C	1:A:177:ILE:CD1	2.69	0.60
1:A:251:TRP:HH2	4:S:103:GLN:CD	2.03	0.60
1:A:392:MET:HE3	1:A:428:MET:HE2	1.84	0.60
2:B:237:ILE:O	2:B:239:GLN:N	2.35	0.60
2:B:357:GLU:O	2:B:361:GLN:HG3	2.01	0.60
2:B:482:ASN:N	2:B:483:PRO:HD3	2.16	0.60
2:B:502:SER:O	2:B:503:LEU:C	2.35	0.60
2:B:549:LEU:CD2	2:B:611:ALA:HA	2.10	0.60
2:B:596:LEU:HD22	2:B:615:SER:HB2	1.83	0.60
3:M:225:VAL:HG13	3:M:480:GLN:HB3	1.83	0.60
3:M:226:PHE:O	3:M:482:ARG:N	2.29	0.60
1:A:170:LEU:CA	1:A:206:LYS:HD2	2.32	0.59
2:B:98:LYS:HD2	2:B:138:ALA:CB	2.29	0.59
2:B:146:LYS:C	2:B:147:MET:HG2	2.23	0.59
2:B:182:ARG:HG3	2:B:217:GLU:HB3	1.83	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:254:LYS:O	2:B:258:GLN:HG2	2.02	0.59
2:B:353:GLN:HE22	2:B:356:LYS:HD2	1.67	0.59
2:B:360:LEU:HB3	2:B:394:TRP:CB	2.31	0.59
2:B:394:TRP:CZ3	2:B:397:GLN:CD	2.75	0.59
2:B:544:THR:O	2:B:548:ILE:HG13	2.01	0.59
3:M:53:HIS:HB3	3:M:65:TYR:CZ	2.37	0.59
3:M:386:PHE:HB2	3:M:397:TRP:CD1	2.38	0.59
4:S:48:SER:CA	4:S:77:TYR:HB3	2.27	0.59
1:A:88:ASN:HB2	1:A:120:ILE:CD1	2.29	0.59
1:A:147:LEU:CG	1:A:166:LEU:CD2	2.80	0.59
1:A:630:PRO:C	2:B:554:LYS:HA	2.22	0.59
2:B:37:TYR:OH	2:B:46:GLN:NE2	0.65	0.59
2:B:80:GLN:CB	2:B:115:LEU:HD11	2.32	0.59
2:B:144:ASP:CA	2:B:179:LYS:HD3	2.32	0.59
2:B:215:TYR:O	2:B:219:TYR:HD1	1.85	0.59
2:B:412:PHE:HE2	2:B:446:TRP:CB	2.13	0.59
3:M:217:ASP:CG	3:M:471:LYS:CA	2.58	0.59
1:A:331:ARG:O	1:A:334:SER:N	2.35	0.59
1:A:533:ILE:HG13	1:A:562:TRP:CH2	2.36	0.59
2:B:143:SER:HB2	2:B:179:LYS:HD2	1.81	0.59
2:B:237:ILE:CG1	2:B:245:GLN:HG2	2.33	0.59
2:B:316:THR:HG21	3:M:90:PHE:HE2	0.87	0.59
2:B:563:PHE:O	2:B:566:ALA:N	2.35	0.59
3:M:212:ASN:CB	3:M:250:LEU:CD2	2.73	0.59
3:M:220:GLU:OE2	3:M:442:GLN:HB3	2.02	0.59
3:M:222:PHE:HD1	3:M:240:ILE:HG23	1.57	0.59
3:M:257:ALA:O	3:M:452:ILE:CG2	2.41	0.59
3:M:344:ILE:O	3:M:344:ILE:HG22	2.02	0.59
3:M:437:TYR:CB	3:M:439:TYR:CZ	2.84	0.59
1:A:134:TYR:O	1:A:135:ASP:C	2.36	0.59
1:A:145:ILE:O	1:A:148:SER:N	2.33	0.59
1:A:213:SER:C	4:S:143:GLU:CG	2.67	0.59
1:A:224:GLU:O	1:A:225:LEU:C	2.38	0.59
1:A:253:ILE:CG2	1:A:281:LEU:HB3	2.32	0.59
1:A:314:ALA:O	1:A:318:ARG:HD3	2.01	0.59
1:A:447:PHE:O	1:A:450:TYR:HB3	2.03	0.59
1:A:606:PHE:CD1	1:A:629:LEU:HG	2.37	0.59
1:A:607:LEU:O	1:A:610:SER:N	2.35	0.59
1:A:633:PHE:HD2	2:B:550:VAL:O	1.79	0.59
2:B:176:ALA:C	2:B:178:ILE:H	2.04	0.59
2:B:178:ILE:CB	2:B:214:ALA:C	2.71	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:297:PRO:O	2:B:301:LEU:CG	2.49	0.59
2:B:352:ASN:HB2	3:M:49:ASP:HB3	1.81	0.59
2:B:563:PHE:HE2	2:B:584:SER:HA	1.64	0.59
3:M:354:ASP:HA	3:M:401:LYS:HB3	1.83	0.59
3:M:374:TYR:OH	3:M:395:GLY:N	2.35	0.59
3:M:378:ILE:C	3:M:413:GLY:HA3	2.13	0.59
3:M:443:SER:HB3	3:M:447:ILE:CB	2.29	0.59
1:A:68:THR:OG1	4:S:165:SER:C	2.38	0.59
1:A:153:ILE:HB	1:A:158:LEU:HD21	1.85	0.59
1:A:219:VAL:HG12	1:A:259:LEU:HD12	1.84	0.59
1:A:584:PHE:C	1:A:587:ASN:H	2.02	0.59
1:A:630:PRO:HG2	2:B:614:ILE:CG2	2.33	0.59
2:B:64:LYS:HG3	2:B:100:LEU:CD1	2.33	0.59
2:B:161:LEU:HB3	2:B:173:VAL:HG21	1.84	0.59
2:B:215:TYR:CD1	2:B:233:TYR:CZ	2.87	0.59
2:B:215:TYR:HD1	2:B:233:TYR:CZ	2.12	0.59
3:M:99:ILE:CG2	3:M:103:TYR:HE1	2.15	0.59
3:M:225:VAL:CA	3:M:480:GLN:O	2.51	0.59
3:M:278:ILE:HG23	3:M:278:ILE:O	2.02	0.59
3:M:323:MET:SD	3:M:342:LEU:CB	2.91	0.59
3:M:338:PHE:CE2	3:M:415:ILE:HG12	2.33	0.59
1:A:68:THR:OG1	4:S:167:ILE:N	2.36	0.59
1:A:125:THR:C	1:A:127:LEU:H	2.04	0.59
1:A:179:LYS:HD2	4:S:143:GLU:CB	2.30	0.59
1:A:244:LEU:CB	1:A:256:LEU:HD13	2.32	0.59
1:A:561:ASN:O	1:A:564:ASN:N	2.35	0.59
2:B:38:TYR:CZ	2:B:42:ILE:HG12	2.21	0.59
2:B:537:PHE:CZ	2:B:545:ARG:CD	2.85	0.59
3:M:220:GLU:CG	3:M:439:TYR:CD1	2.85	0.59
3:M:290:PHE:CD1	3:M:299:LEU:HD13	2.37	0.59
1:A:170:LEU:O	1:A:206:LYS:NZ	2.35	0.59
1:A:322:PHE:HD2	1:A:330:LEU:CD2	2.14	0.59
1:A:375:VAL:O	1:A:378:ILE:O	2.21	0.59
1:A:634:ASN:ND2	2:B:554:LYS:CB	2.49	0.59
2:B:117:LEU:HA	2:B:150:LEU:CD2	2.33	0.59
2:B:135:ARG:HH12	2:B:164:ASP:HB2	1.66	0.59
2:B:597:TYR:O	2:B:601:TYR:CD1	2.55	0.59
1:A:107:TYR:HE2	1:A:128:LEU:CD2	2.03	0.59
1:A:153:ILE:HB	1:A:158:LEU:CD2	2.33	0.59
1:A:291:ILE:HD12	1:A:291:ILE:H	1.67	0.59
1:A:404:GLN:OE1	2:B:7:ARG:NH2	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:556:VAL:CG2	1:A:603:VAL:HG13	2.33	0.59
1:A:621:LEU:HD13	1:A:621:LEU:C	2.23	0.59
2:B:24:ALA:HB1	2:B:35:TYR:CG	2.37	0.59
2:B:91:THR:HA	2:B:126:SER:HB2	1.85	0.59
2:B:197:LYS:CD	2:B:283:TYR:CE2	2.86	0.59
2:B:334:MET:C	2:B:336:ASN:H	2.06	0.59
3:M:3:LEU:CG	3:M:80:THR:O	2.49	0.59
3:M:104:PHE:CD1	3:M:117:ASN:ND2	2.71	0.59
3:M:240:ILE:HG22	3:M:444:ALA:HB1	1.85	0.59
3:M:467:TYR:CG	3:M:468:LYS:N	2.71	0.59
1:A:67:LYS:HB2	4:S:165:SER:CB	2.33	0.59
1:A:140:VAL:O	1:A:141:VAL:C	2.34	0.59
2:B:37:TYR:C	2:B:39:SER:N	2.55	0.59
2:B:50:LEU:HD23	2:B:62:ALA:CA	2.33	0.59
2:B:142:LEU:HD13	2:B:154:ILE:HG12	1.85	0.59
2:B:155:LEU:HD21	2:B:192:LEU:N	2.17	0.59
2:B:178:ILE:CB	2:B:214:ALA:CA	2.81	0.59
2:B:215:TYR:CE2	2:B:229:HIS:ND1	2.70	0.59
2:B:275:ARG:HG2	2:B:294:VAL:HG21	1.83	0.59
2:B:451:MET:CG	2:B:489:ILE:HG12	2.33	0.59
2:B:461:HIS:HB3	2:B:463:LEU:HD23	1.84	0.59
2:B:556:LEU:CD2	2:B:588:ILE:CD1	2.79	0.59
3:M:243:ILE:HG21	3:M:298:ARG:HG2	1.83	0.59
1:A:182:ILE:C	1:A:221:VAL:CG2	2.67	0.59
1:A:402:ILE:O	1:A:402:ILE:HG22	2.02	0.59
1:A:483:LYS:C	1:A:484:VAL:HG23	2.23	0.59
2:B:387:ASP:HB3	2:B:391:ALA:CB	2.32	0.59
2:B:389:ILE:HG23	2:B:427:ASN:HB2	1.82	0.59
2:B:592:TYR:C	2:B:592:TYR:CD1	2.76	0.59
3:M:244:VAL:HG22	3:M:472:TYR:HE2	1.66	0.59
4:S:38:LEU:HB3	4:S:51:LEU:HD13	1.83	0.59
1:A:260:PHE:CE2	1:A:274:LEU:CG	2.84	0.58
1:A:516:ILE:HD13	1:A:551:LEU:CB	2.32	0.58
2:B:90:ILE:HG12	2:B:98:LYS:CE	2.32	0.58
2:B:182:ARG:HD2	2:B:217:GLU:CD	2.22	0.58
2:B:257:LYS:HA	2:B:260:LEU:CG	2.33	0.58
2:B:360:LEU:O	2:B:363:ILE:HB	2.03	0.58
3:M:99:ILE:CG2	3:M:103:TYR:CE1	2.86	0.58
3:M:218:LEU:HG	3:M:244:VAL:CG2	2.29	0.58
3:M:225:VAL:HA	3:M:480:GLN:N	2.18	0.58
3:M:347:PHE:CZ	3:M:350:VAL:CG1	2.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:97:SER:H	4:S:166:LYS:NZ	2.02	0.58
1:A:176:TYR:CB	4:S:155:GLU:HG2	2.31	0.58
1:A:270:LEU:CD1	1:A:274:LEU:HD23	2.32	0.58
1:A:278:ILE:O	1:A:280:GLU:N	2.36	0.58
1:A:607:LEU:C	1:A:609:LEU:N	2.55	0.58
2:B:175:LEU:O	2:B:178:ILE:HB	2.03	0.58
2:B:263:PRO:HA	2:B:266:VAL:CG2	2.33	0.58
2:B:355:ASN:O	2:B:359:LEU:HD23	2.03	0.58
2:B:549:LEU:HD22	2:B:611:ALA:HB3	1.72	0.58
2:B:562:ASN:CB	2:B:580:TYR:HB2	2.30	0.58
3:M:6:TYR:CD2	3:M:16:PHE:O	2.56	0.58
3:M:245:ASP:O	3:M:246:VAL:HG22	2.03	0.58
3:M:290:PHE:HZ	3:M:297:PHE:CG	2.11	0.58
3:M:410:VAL:HG13	3:M:412:ARG:HH11	1.66	0.58
4:S:8:PHE:CE1	4:S:84:TYR:HB3	1.80	0.58
1:A:180:LYS:CE	4:S:156:LEU:HD11	2.33	0.58
1:A:223:CYS:SG	1:A:262:ASN:ND2	2.66	0.58
1:A:638:LEU:HD12	2:B:518:ILE:HG22	1.77	0.58
2:B:87:VAL:CG2	2:B:119:SER:HA	2.33	0.58
2:B:120:ILE:C	2:B:153:ILE:HG21	2.23	0.58
2:B:177:ILE:CD1	2:B:196:LEU:CD2	2.82	0.58
2:B:278:PRO:HB3	2:B:288:TYR:C	2.16	0.58
2:B:534:ILE:O	2:B:535:GLN:C	2.38	0.58
2:B:537:PHE:CB	2:B:598:LEU:CD1	2.30	0.58
2:B:563:PHE:HA	2:B:566:ALA:HB3	1.84	0.58
2:B:569:THR:CA	2:B:571:SER:H	2.16	0.58
2:B:589:SER:OG	2:B:618:PHE:CZ	2.46	0.58
3:M:56:VAL:O	3:M:56:VAL:CG1	2.50	0.58
3:M:217:ASP:HB2	3:M:470:ALA:CA	2.33	0.58
3:M:235:LEU:HD13	3:M:306:LEU:HB3	1.85	0.58
3:M:245:ASP:N	3:M:472:TYR:CD2	2.57	0.58
3:M:443:SER:OG	3:M:448:TYR:N	2.35	0.58
1:A:229:ASN:O	1:A:230:PRO:C	2.36	0.58
1:A:533:ILE:HG12	1:A:562:TRP:CZ2	2.39	0.58
1:A:566:PHE:HD1	1:A:570:LYS:HA	1.68	0.58
2:B:120:ILE:HG22	2:B:153:ILE:CG2	2.33	0.58
2:B:181:TYR:CD2	2:B:218:CYS:CA	2.85	0.58
2:B:237:ILE:CD1	2:B:309:LEU:HD21	2.34	0.58
2:B:308:CYS:O	2:B:311:TYR:N	2.37	0.58
2:B:588:ILE:CG2	2:B:618:PHE:CE1	2.86	0.58
2:B:588:ILE:CG2	2:B:618:PHE:CZ	2.78	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:9:ASP:HA	3:M:75:TRP:HD1	1.67	0.58
3:M:19:LEU:CD1	3:M:24:ALA:CB	2.80	0.58
3:M:442:GLN:OE1	3:M:442:GLN:HA	2.02	0.58
4:S:53:THR:N	4:S:69:ASN:CB	2.58	0.58
2:B:51:LEU:HD22	2:B:59:VAL:HG13	1.84	0.58
2:B:497:LEU:HD11	2:B:508:ARG:CZ	2.31	0.58
2:B:501:THR:C	2:B:508:ARG:HH21	2.06	0.58
3:M:215:TYR:CB	3:M:469:GLY:H	2.13	0.58
3:M:219:LEU:HA	3:M:440:ILE:HA	1.85	0.58
1:A:190:LEU:HD11	1:A:228:LYS:HE3	1.85	0.58
1:A:338:PHE:HE2	1:A:352:PHE:CZ	2.21	0.58
1:A:634:ASN:OD1	2:B:553:ALA:C	2.41	0.58
2:B:120:ILE:HG21	2:B:154:ILE:HG13	1.82	0.58
2:B:177:ILE:HD11	2:B:195:ILE:HG22	1.84	0.58
2:B:299:LEU:C	2:B:299:LEU:HD13	2.24	0.58
2:B:452:LYS:NZ	2:B:456:ASP:CG	2.55	0.58
3:M:317:MET:CG	3:M:320:ILE:O	2.51	0.58
3:M:354:ASP:HB2	3:M:440:ILE:CD1	2.33	0.58
3:M:360:LEU:CD2	3:M:362:PHE:HE2	2.14	0.58
1:A:581:LEU:HG	1:A:607:LEU:HD11	1.86	0.58
2:B:29:LYS:O	2:B:32:GLU:CG	2.50	0.58
2:B:50:LEU:CD2	2:B:62:ALA:HB2	2.34	0.58
2:B:133:GLU:O	2:B:168:MET:CE	2.51	0.58
2:B:174:ALA:CB	2:B:211:ALA:HB1	2.31	0.58
2:B:188:TYR:O	2:B:192:LEU:HD13	2.02	0.58
2:B:316:THR:HG23	3:M:90:PHE:CZ	2.39	0.58
2:B:346:THR:CG2	2:B:350:THR:HG21	2.29	0.58
3:M:380:ARG:NH1	3:M:412:ARG:HD2	2.19	0.58
1:A:488:ARG:CG	1:A:522:PHE:CD2	2.87	0.58
2:B:29:LYS:O	2:B:30:LEU:C	2.40	0.58
2:B:37:TYR:HH	2:B:46:GLN:NE2	0.27	0.58
3:M:1:MET:HA	3:M:81:SER:OG	2.04	0.58
3:M:437:TYR:HB2	3:M:439:TYR:CE1	2.38	0.58
4:S:55:PRO:HB3	4:S:71:GLU:HA	1.86	0.58
4:S:117:ASN:HB2	4:S:120:ASP:OD2	2.04	0.58
1:A:402:ILE:C	1:A:404:GLN:N	2.56	0.58
2:B:30:LEU:HD12	2:B:30:LEU:O	2.04	0.58
2:B:37:TYR:CE2	2:B:46:GLN:OE1	2.56	0.58
2:B:83:PHE:CZ	2:B:105:LEU:HG	2.39	0.58
2:B:90:ILE:O	2:B:98:LYS:HE2	2.04	0.58
2:B:159:LYS:HG3	2:B:191:GLU:OE1	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:267:ASP:N	2:B:289:PRO:HB2	2.03	0.58
3:M:101:LEU:CD2	3:M:106:LYS:CG	2.81	0.58
3:M:218:LEU:N	3:M:218:LEU:HD12	2.18	0.58
3:M:242:GLY:O	3:M:301:GLU:HA	2.04	0.58
3:M:347:PHE:HE2	3:M:352:GLN:N	2.02	0.58
1:A:68:THR:CB	4:S:167:ILE:N	2.66	0.58
1:A:101:GLN:CG	4:S:167:ILE:HG13	2.31	0.58
1:A:102:GLN:HG2	4:S:166:LYS:CA	2.22	0.58
1:A:104:ARG:CG	1:A:145:ILE:HG21	2.34	0.58
1:A:322:PHE:CB	1:A:330:LEU:HD21	2.32	0.58
1:A:555:LEU:HD12	1:A:585:PHE:CE1	2.39	0.58
1:A:581:LEU:CD2	1:A:607:LEU:HD21	2.33	0.58
1:A:596:VAL:O	1:A:599:ARG:HB2	2.03	0.58
2:B:87:VAL:O	2:B:88:LYS:C	2.39	0.58
2:B:334:MET:CB	2:B:369:LEU:HD23	2.34	0.58
2:B:367:SER:OG	2:B:401:THR:CB	2.51	0.58
2:B:400:SER:CA	2:B:439:CYS:SG	2.91	0.58
2:B:403:ILE:HG22	2:B:411:ILE:CD1	2.33	0.58
1:A:189:PHE:HD2	1:A:225:LEU:HD11	1.62	0.57
1:A:204:VAL:HG13	1:A:239:LEU:HD11	1.86	0.57
1:A:384:LEU:HD13	1:A:435:ILE:HG23	1.80	0.57
1:A:438:ALA:O	1:A:441:TYR:CD1	2.56	0.57
1:A:464:ILE:HG23	1:A:465:SER:HA	1.86	0.57
1:A:516:ILE:HG12	1:A:551:LEU:CD1	2.34	0.57
1:A:625:LEU:O	1:A:626:SER:C	2.36	0.57
2:B:42:ILE:C	2:B:46:GLN:OE1	2.43	0.57
2:B:178:ILE:CD1	2:B:215:TYR:N	2.67	0.57
2:B:189:HIS:CE1	2:B:193:LEU:HD11	2.38	0.57
2:B:223:LEU:HD22	2:B:255:TYR:CZ	2.34	0.57
2:B:278:PRO:CG	2:B:289:PRO:O	2.51	0.57
2:B:350:THR:HB	2:B:352:ASN:HD22	1.69	0.57
2:B:470:ALA:O	2:B:473:ASN:N	2.35	0.57
2:B:508:ARG:O	2:B:509:ALA:C	2.30	0.57
2:B:519:ALA:HB2	2:B:555:LEU:HD13	1.86	0.57
2:B:534:ILE:O	2:B:536:ASN:N	2.37	0.57
3:M:435:LEU:HB2	3:M:437:TYR:CE1	2.39	0.57
1:A:241:TYR:HD2	1:A:242:GLU:N	2.01	0.57
1:A:355:LEU:O	1:A:359:LEU:HG	2.04	0.57
1:A:603:VAL:O	1:A:606:PHE:N	2.36	0.57
1:A:637:GLU:HG3	2:B:515:PHE:C	2.24	0.57
2:B:123:LEU:HD22	2:B:138:ALA:HB1	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:175:LEU:CD2	2:B:210:CYS:HB3	2.34	0.57
2:B:200:MET:CE	2:B:232:ARG:H	2.15	0.57
2:B:219:TYR:CA	2:B:223:LEU:CD2	2.82	0.57
2:B:247:TYR:OH	3:M:91:THR:CB	2.51	0.57
2:B:278:PRO:HD3	2:B:290:SER:HA	1.85	0.57
2:B:310:ILE:O	2:B:311:TYR:O	2.21	0.57
2:B:433:VAL:HG12	2:B:474:VAL:HG21	0.60	0.57
2:B:537:PHE:CZ	2:B:599:ALA:HA	2.39	0.57
3:M:52:ASP:CA	3:M:67:SER:CA	2.82	0.57
3:M:218:LEU:HG	3:M:472:TYR:CE2	2.32	0.57
3:M:380:ARG:N	3:M:412:ARG:O	2.37	0.57
3:M:440:ILE:O	3:M:440:ILE:HG22	2.03	0.57
4:S:34:GLN:CD	4:S:58:LEU:HD21	2.17	0.57
4:S:107:GLU:HG2	4:S:146:VAL:HG21	1.85	0.57
1:A:92:LEU:HD13	1:A:123:LEU:CD1	2.27	0.57
1:A:101:GLN:O	4:S:167:ILE:CD1	2.53	0.57
1:A:189:PHE:CG	1:A:225:LEU:HD21	2.37	0.57
1:A:637:GLU:HB3	2:B:516:GLY:HA3	1.80	0.57
2:B:44:PRO:O	2:B:47:LEU:CB	2.48	0.57
2:B:178:ILE:CG2	2:B:214:ALA:CA	2.83	0.57
2:B:178:ILE:CA	2:B:218:CYS:HB2	2.18	0.57
2:B:189:HIS:NE2	2:B:222:HIS:HB3	2.20	0.57
2:B:306:LEU:HD22	2:B:321:CYS:SG	2.45	0.57
2:B:472:VAL:HG13	2:B:510:GLY:C	2.24	0.57
2:B:508:ARG:O	2:B:512:VAL:CG2	2.43	0.57
2:B:542:PRO:CA	2:B:602:ASP:CG	2.60	0.57
3:M:2:TYR:C	3:M:3:LEU:HD12	2.24	0.57
4:S:80:TYR:CE2	4:S:110:ASP:HB2	2.39	0.57
1:A:132:LEU:HD23	1:A:143:VAL:HG22	1.86	0.57
1:A:485:PRO:C	1:A:486:SER:O	2.39	0.57
1:A:492:ILE:HD11	1:A:522:PHE:HB3	1.85	0.57
1:A:627:GLU:O	1:A:630:PRO:HD2	2.05	0.57
2:B:64:LYS:CG	2:B:100:LEU:HD11	2.34	0.57
2:B:559:ASP:OD2	2:B:563:PHE:CE1	2.57	0.57
3:M:17:GLN:H	3:M:118:TYR:HE1	1.51	0.57
3:M:74:TYR:HD2	3:M:109:LEU:O	1.88	0.57
3:M:223:HIS:HB3	3:M:478:ASN:C	2.24	0.57
4:S:15:ARG:HD2	4:S:122:ILE:HG13	1.84	0.57
1:A:234:ILE:HG23	1:A:267:GLU:HG2	1.86	0.57
1:A:253:ILE:HG12	1:A:281:LEU:CG	2.35	0.57
1:A:594:PHE:HD2	2:B:474:VAL:HG23	1.70	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:134:LEU:O	2:B:136:CYS:N	2.37	0.57
2:B:158:VAL:HG21	2:B:177:ILE:HG13	1.86	0.57
2:B:162:VAL:HG21	2:B:195:ILE:CB	2.31	0.57
2:B:313:SER:O	2:B:315:PRO:HD3	2.04	0.57
2:B:396:ILE:CD1	2:B:418:TYR:HE2	2.11	0.57
3:M:374:TYR:CZ	3:M:390:ILE:HD13	2.40	0.57
4:S:8:PHE:HA	4:S:13:GLN:O	2.04	0.57
4:S:35:VAL:HB	4:S:77:TYR:OH	2.05	0.57
1:A:183:THR:HG21	4:S:134:GLU:HA	1.87	0.57
1:A:253:ILE:CG1	1:A:281:LEU:HD22	2.35	0.57
1:A:637:GLU:HB2	2:B:551:LEU:CD2	2.35	0.57
2:B:17:VAL:O	2:B:18:ILE:O	2.21	0.57
2:B:108:PHE:CZ	2:B:115:LEU:CD2	2.88	0.57
2:B:251:LEU:O	2:B:254:LYS:N	2.36	0.57
3:M:114:ILE:O	3:M:118:TYR:N	2.38	0.57
3:M:121:ILE:HG23	3:M:125:PHE:HE1	1.69	0.57
3:M:317:MET:HB2	3:M:322:LEU:CB	2.35	0.57
4:S:14:PRO:HB3	4:S:36:TYR:CE1	2.33	0.57
4:S:111:ARG:HB2	4:S:150:VAL:CG2	2.32	0.57
1:A:105:VAL:HG23	4:S:167:ILE:CG2	2.27	0.57
1:A:600:SER:C	1:A:602:GLU:N	2.56	0.57
1:A:630:PRO:HG2	2:B:614:ILE:HG23	1.86	0.57
1:A:637:GLU:HG2	2:B:516:GLY:N	1.95	0.57
2:B:21:GLU:CA	2:B:24:ALA:CB	2.58	0.57
2:B:64:LYS:N	2:B:100:LEU:HD13	2.19	0.57
2:B:70:MET:HE1	2:B:107:ARG:CA	2.31	0.57
2:B:155:LEU:C	2:B:157:THR:N	2.55	0.57
2:B:231:ARG:O	2:B:234:CYS:N	2.38	0.57
2:B:277:CYS:SG	2:B:292:GLU:OE2	2.62	0.57
2:B:360:LEU:HD13	2:B:391:ALA:CA	2.25	0.57
1:A:63:ASP:O	4:S:165:SER:CB	2.51	0.57
1:A:114:PHE:CD2	1:A:153:ILE:HG23	2.40	0.57
1:A:163:ALA:O	1:A:165:ASP:N	2.37	0.57
1:A:225:LEU:CD1	1:A:233:PHE:CE2	2.81	0.57
1:A:316:LEU:O	1:A:319:LEU:N	2.35	0.57
1:A:535:ILE:O	1:A:535:ILE:HG22	2.04	0.57
2:B:257:LYS:HA	2:B:260:LEU:HD21	1.86	0.57
2:B:275:ARG:HE	2:B:275:ARG:CA	2.17	0.57
2:B:340:ILE:HG21	2:B:373:LEU:HG	1.87	0.57
2:B:451:MET:CE	2:B:489:ILE:HG12	2.34	0.57
2:B:574:ASN:C	2:B:576:GLN:H	2.08	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:96:ILE:HD12	3:M:125:PHE:HA	1.86	0.57
3:M:104:PHE:HD1	3:M:117:ASN:ND2	2.02	0.57
3:M:235:LEU:HD22	3:M:307:SER:HA	1.86	0.57
4:S:15:ARG:HH11	4:S:122:ILE:HD11	1.57	0.57
4:S:55:PRO:HD3	4:S:71:GLU:HA	1.87	0.57
1:A:255:ARG:C	1:A:257:LEU:N	2.56	0.57
1:A:292:TYR:CD1	1:A:292:TYR:C	2.78	0.57
2:B:50:LEU:HD21	2:B:61:ASP:HB2	1.86	0.57
2:B:151:ALA:HA	2:B:180:LEU:HG	1.82	0.57
2:B:208:ILE:HG21	2:B:240:LEU:HD11	1.87	0.57
2:B:336:ASN:HB2	2:B:339:PHE:CE1	2.39	0.57
2:B:453:TRP:HE3	2:B:453:TRP:HA	1.70	0.57
2:B:519:ALA:HB1	2:B:555:LEU:CD1	2.35	0.57
3:M:308:SER:O	3:M:312:GLN:N	2.33	0.57
3:M:327:PHE:HE1	3:M:336:ASP:CG	2.07	0.57
1:A:94:VAL:O	1:A:95:MET:C	2.39	0.57
1:A:170:LEU:HB2	1:A:202:LYS:HG2	1.86	0.57
1:A:421:PRO:CB	1:A:424:TYR:CD1	2.88	0.57
1:A:465:SER:H	2:B:1:MET:CG	2.15	0.57
1:A:563:CYS:HB3	1:A:621:LEU:CD1	2.23	0.57
2:B:20:ARG:CD	2:B:21:GLU:N	2.68	0.57
2:B:60:ARG:CD	2:B:96:LYS:HG2	2.25	0.57
2:B:245:GLN:CB	2:B:309:LEU:HD11	2.34	0.57
2:B:310:ILE:HG23	2:B:318:ILE:HG23	1.87	0.57
2:B:399:LEU:O	2:B:400:SER:C	2.38	0.57
3:M:60:LEU:HD23	3:M:60:LEU:C	2.22	0.57
3:M:249:TYR:CE1	3:M:467:TYR:CZ	2.93	0.57
4:S:111:ARG:CB	4:S:150:VAL:CG2	2.83	0.57
1:A:67:LYS:N	4:S:165:SER:CB	2.44	0.56
1:A:253:ILE:CG1	1:A:281:LEU:HB3	2.34	0.56
2:B:80:GLN:HG3	2:B:115:LEU:HD21	1.87	0.56
2:B:143:SER:CB	2:B:179:LYS:HB2	2.35	0.56
2:B:197:LYS:CD	2:B:283:TYR:CD2	2.88	0.56
2:B:387:ASP:HB3	2:B:391:ALA:HB3	1.86	0.56
3:M:241:HIS:HB2	3:M:476:THR:HG22	1.84	0.56
3:M:302:TYR:CE2	3:M:304:VAL:HB	2.40	0.56
3:M:327:PHE:HE1	3:M:336:ASP:OD2	1.88	0.56
4:S:50:PHE:HB3	4:S:76:ILE:HA	1.86	0.56
4:S:93:GLU:HA	4:S:93:GLU:OE1	2.04	0.56
2:B:24:ALA:CB	2:B:35:TYR:CD1	2.84	0.56
2:B:275:ARG:N	2:B:295:ASN:CG	2.54	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:278:PRO:CB	2:B:292:GLU:OE1	2.52	0.56
2:B:355:ASN:C	2:B:359:LEU:CD2	2.73	0.56
2:B:498:THR:HG22	2:B:532:ARG:HB2	1.86	0.56
3:M:432:THR:HG23	3:M:432:THR:O	2.03	0.56
1:A:158:LEU:HG	1:A:162:ILE:HD11	1.86	0.56
1:A:166:LEU:O	1:A:170:LEU:HD22	2.05	0.56
1:A:186:PHE:CD1	1:A:224:GLU:CG	2.88	0.56
1:A:384:LEU:HG	1:A:385:LYS:H	1.70	0.56
1:A:404:GLN:OE1	2:B:7:ARG:CZ	2.53	0.56
2:B:41:ASN:CG	2:B:43:ASN:OD1	2.44	0.56
2:B:86:VAL:HG13	2:B:101:ILE:CG1	2.29	0.56
2:B:123:LEU:O	2:B:127:LEU:HD12	2.04	0.56
2:B:140:SER:N	2:B:172:GLU:OE1	2.37	0.56
2:B:181:TYR:CE2	2:B:218:CYS:O	2.52	0.56
2:B:199:LEU:O	2:B:200:MET:C	2.40	0.56
2:B:220:ALA:HA	2:B:258:GLN:HB3	1.88	0.56
2:B:436:LEU:HD12	2:B:454:LEU:HD21	1.87	0.56
2:B:461:HIS:O	2:B:463:LEU:N	2.35	0.56
2:B:468:LEU:O	2:B:472:VAL:HG23	2.06	0.56
3:M:66:PHE:HB3	3:M:77:LEU:HD13	1.83	0.56
3:M:96:ILE:CD1	3:M:125:PHE:HA	2.35	0.56
3:M:222:PHE:CE1	3:M:439:TYR:CE1	2.93	0.56
3:M:373:ALA:O	3:M:418:GLU:O	2.24	0.56
3:M:403:THR:CG2	3:M:407:THR:HG1	2.16	0.56
4:S:58:LEU:O	4:S:59:LEU:HB3	2.04	0.56
1:A:97:SER:O	1:A:98:ASN:C	2.37	0.56
1:A:99:LYS:HB3	4:S:163:THR:C	2.25	0.56
1:A:170:LEU:CB	1:A:206:LYS:HG3	2.34	0.56
1:A:214:VAL:N	4:S:143:GLU:OE1	2.39	0.56
1:A:477:PHE:HE2	1:A:481:MET:SD	2.29	0.56
2:B:21:GLU:C	2:B:24:ALA:HB3	2.25	0.56
2:B:37:TYR:HE2	2:B:38:TYR:HE1	1.35	0.56
2:B:278:PRO:CA	2:B:288:TYR:HB3	2.25	0.56
2:B:352:ASN:ND2	2:B:352:ASN:H	2.02	0.56
2:B:523:PHE:HD1	2:B:559:ASP:CG	2.09	0.56
2:B:556:LEU:HD23	2:B:588:ILE:HD11	1.88	0.56
2:B:559:ASP:C	2:B:563:PHE:HB2	2.26	0.56
2:B:599:ALA:C	2:B:601:TYR:H	2.08	0.56
3:M:222:PHE:CE1	3:M:240:ILE:HG23	2.28	0.56
3:M:428:VAL:O	3:M:430:LEU:N	2.39	0.56
4:S:17:VAL:CG2	4:S:19:PHE:CE1	2.81	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:141:VAL:CA	4:S:159:ALA:HB2	2.35	0.56
1:A:274:LEU:O	1:A:275:LEU:C	2.37	0.56
2:B:50:LEU:HD23	2:B:62:ALA:CB	2.35	0.56
2:B:98:LYS:CD	2:B:138:ALA:HB2	2.31	0.56
2:B:139:LEU:HD23	2:B:173:VAL:C	2.11	0.56
2:B:247:TYR:OH	3:M:91:THR:CG2	2.53	0.56
3:M:360:LEU:HD21	3:M:362:PHE:HE2	1.66	0.56
3:M:372:ILE:HD13	3:M:424:PHE:CE1	2.40	0.56
1:A:67:LYS:CB	4:S:166:LYS:N	2.69	0.56
1:A:101:GLN:CG	4:S:167:ILE:CG2	2.53	0.56
1:A:179:LYS:HE3	4:S:143:GLU:CA	2.33	0.56
1:A:397:ASP:O	1:A:418:ILE:CD1	2.53	0.56
3:M:5:PHE:HB2	3:M:125:PHE:CD2	2.40	0.56
1:A:208:ASP:OD1	1:A:239:LEU:HD22	2.05	0.56
1:A:275:LEU:HD12	1:A:308:ASP:CG	2.22	0.56
1:A:481:MET:SD	1:A:518:CYS:SG	3.04	0.56
1:A:556:VAL:HG21	1:A:603:VAL:CG1	2.36	0.56
2:B:102:HIS:O	2:B:103:LEU:O	2.24	0.56
2:B:108:PHE:CE2	2:B:112:ASP:HB3	2.41	0.56
2:B:134:LEU:C	2:B:136:CYS:N	2.57	0.56
2:B:175:LEU:HD21	2:B:210:CYS:CB	2.35	0.56
2:B:215:TYR:CG	2:B:226:LEU:CD1	2.88	0.56
2:B:277:CYS:HA	2:B:292:GLU:CA	2.34	0.56
2:B:560:ILE:CA	2:B:564:LYS:H	2.17	0.56
2:B:585:GLY:O	2:B:589:SER:N	2.29	0.56
3:M:67:SER:CB	3:M:90:PHE:HB2	2.34	0.56
3:M:219:LEU:HB2	3:M:472:TYR:C	2.14	0.56
3:M:348:LYS:CG	3:M:405:THR:HG22	2.23	0.56
1:A:121:LEU:HD23	1:A:121:LEU:C	2.24	0.56
1:A:186:PHE:CZ	1:A:224:GLU:HG2	2.40	0.56
1:A:215:VAL:HG22	1:A:243:ILE:HG13	1.87	0.56
1:A:401:VAL:HG23	1:A:418:ILE:CA	2.36	0.56
2:B:173:VAL:CG1	2:B:199:LEU:HD11	2.36	0.56
2:B:275:ARG:HA	2:B:275:ARG:HE	1.70	0.56
2:B:388:PRO:O	2:B:391:ALA:HB3	2.05	0.56
2:B:560:ILE:HA	2:B:563:PHE:HB2	1.88	0.56
3:M:49:ASP:CA	3:M:75:TRP:HH2	1.96	0.56
3:M:338:PHE:CE2	3:M:415:ILE:CD1	2.89	0.56
4:S:53:THR:CB	4:S:68:VAL:CA	2.84	0.56
1:A:179:LYS:HZ1	4:S:149:ILE:HG12	1.70	0.56
1:A:225:LEU:HB3	1:A:233:PHE:HE1	1.66	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:384:LEU:CG	1:A:385:LYS:N	2.68	0.56
2:B:81:LEU:O	2:B:82:TYR:C	2.35	0.56
2:B:178:ILE:CG1	2:B:215:TYR:CA	2.78	0.56
2:B:189:HIS:NE2	2:B:193:LEU:CD1	2.61	0.56
2:B:189:HIS:CD2	2:B:222:HIS:CB	2.87	0.56
2:B:226:LEU:HG	2:B:255:TYR:CE2	2.40	0.56
2:B:245:GLN:HG2	2:B:309:LEU:HD11	1.84	0.56
2:B:350:THR:CG2	2:B:352:ASN:HD21	2.18	0.56
2:B:374:PHE:HD2	2:B:402:LEU:HD21	1.71	0.56
2:B:578:PRO:HB3	2:B:579:PRO:CD	2.33	0.56
3:M:4:SER:O	3:M:79:SER:N	2.39	0.56
3:M:18:TYR:CE2	3:M:20:LEU:CD2	2.86	0.56
3:M:105:ASP:O	3:M:106:LYS:HB3	2.04	0.56
3:M:212:ASN:HB3	3:M:250:LEU:CD2	2.31	0.56
4:S:50:PHE:HA	4:S:75:ILE:O	2.06	0.56
4:S:68:VAL:O	4:S:75:ILE:HD12	2.06	0.56
1:A:88:ASN:HB3	1:A:120:ILE:HG23	1.87	0.56
1:A:107:TYR:CD1	1:A:146:ALA:HA	2.41	0.56
1:A:162:ILE:O	1:A:165:ASP:HB2	2.06	0.56
1:A:167:PHE:CD1	1:A:202:LYS:HB2	2.41	0.56
1:A:178:ARG:HB2	1:A:214:VAL:CG2	2.35	0.56
2:B:214:ALA:C	2:B:216:LYS:H	2.09	0.56
2:B:483:PRO:HA	2:B:486:HIS:HB2	1.88	0.56
2:B:519:ALA:O	2:B:523:PHE:N	2.39	0.56
3:M:223:HIS:HB3	3:M:479:PHE:N	2.21	0.56
1:A:155:THR:O	1:A:158:LEU:HB3	2.06	0.55
1:A:255:ARG:HD3	4:S:141:VAL:C	2.25	0.55
1:A:295:VAL:HG11	1:A:337:LEU:HD13	1.88	0.55
2:B:172:GLU:O	2:B:174:ALA:N	2.38	0.55
2:B:174:ALA:HB3	2:B:211:ALA:HB2	1.88	0.55
2:B:215:TYR:CD2	2:B:219:TYR:CE1	2.79	0.55
2:B:307:ASN:HD22	2:B:336:ASN:HD21	1.43	0.55
2:B:340:ILE:HG13	2:B:373:LEU:CG	2.36	0.55
3:M:101:LEU:HD13	3:M:109:LEU:HD13	1.88	0.55
3:M:262:THR:HG23	3:M:267:ILE:HG12	1.87	0.55
3:M:437:TYR:HD1	3:M:437:TYR:H	1.51	0.55
2:B:69:ILE:O	2:B:70:MET:C	2.36	0.55
2:B:512:VAL:HB	2:B:551:LEU:CD1	2.34	0.55
2:B:596:LEU:HD13	2:B:611:ALA:C	2.27	0.55
3:M:16:PHE:CE2	3:M:125:PHE:CE2	2.94	0.55
3:M:18:TYR:OH	3:M:126:ASN:HB2	2.05	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:216:VAL:HB	3:M:472:TYR:OH	2.06	0.55
3:M:224:VAL:O	3:M:479:PHE:HA	2.04	0.55
3:M:244:VAL:O	3:M:299:LEU:HB3	2.06	0.55
3:M:272:LEU:HD22	3:M:278:ILE:CG2	2.35	0.55
1:A:99:LYS:HG2	1:A:101:GLN:N	2.21	0.55
1:A:283:GLU:OE1	1:A:318:ARG:NH2	2.38	0.55
1:A:631:SER:C	2:B:554:LYS:HG3	2.24	0.55
2:B:10:SER:O	2:B:13:ASP:N	2.38	0.55
2:B:50:LEU:CG	2:B:58:GLU:O	2.51	0.55
2:B:139:LEU:HD22	2:B:173:VAL:HG13	1.89	0.55
2:B:267:ASP:CA	2:B:289:PRO:CG	2.84	0.55
2:B:537:PHE:C	2:B:539:ASN:H	2.09	0.55
3:M:300:LEU:O	3:M:300:LEU:HD12	2.07	0.55
4:S:15:ARG:NH2	4:S:118:GLU:CD	2.59	0.55
1:A:114:PHE:CD2	1:A:153:ILE:HG12	2.42	0.55
1:A:215:VAL:CG2	1:A:243:ILE:CG1	2.79	0.55
1:A:516:ILE:HG21	1:A:551:LEU:HA	1.88	0.55
2:B:108:PHE:CE2	2:B:115:LEU:CD2	2.90	0.55
2:B:116:THR:HG22	2:B:150:LEU:CG	2.36	0.55
2:B:120:ILE:HG13	2:B:150:LEU:CB	2.32	0.55
2:B:430:ILE:O	2:B:433:VAL:HB	2.07	0.55
3:M:253:ASN:N	3:M:254:PRO:CD	2.70	0.55
4:S:89:VAL:CG1	4:S:98:ILE:CG2	2.76	0.55
1:A:185:LEU:O	1:A:189:PHE:CD2	2.59	0.55
1:A:279:LEU:HA	1:A:282:MET:HG2	1.89	0.55
1:A:301:GLY:O	1:A:302:ASN:CB	2.52	0.55
1:A:441:TYR:O	1:A:442:SER:C	2.44	0.55
1:A:441:TYR:C	1:A:443:SER:N	2.58	0.55
1:A:606:PHE:CD2	1:A:629:LEU:HG	2.41	0.55
2:B:159:LYS:CD	2:B:191:GLU:OE1	2.55	0.55
2:B:225:LEU:HD11	2:B:283:TYR:CE1	2.39	0.55
3:M:65:TYR:CG	3:M:86:PRO:CB	2.89	0.55
3:M:215:TYR:O	3:M:246:VAL:CG1	2.55	0.55
3:M:443:SER:OG	3:M:447:ILE:HG13	2.05	0.55
1:A:96:SER:HB3	1:A:127:LEU:CG	2.37	0.55
1:A:136:GLY:O	1:A:139:ASP:CA	2.55	0.55
1:A:606:PHE:HB3	1:A:629:LEU:HD12	1.88	0.55
2:B:177:ILE:HD13	2:B:196:LEU:CG	2.34	0.55
2:B:453:TRP:HA	2:B:453:TRP:CE3	2.40	0.55
2:B:559:ASP:C	2:B:563:PHE:H	2.05	0.55
3:M:101:LEU:HD12	3:M:109:LEU:HD12	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:217:ASP:CB	3:M:471:LYS:CA	2.85	0.55
3:M:262:THR:HG22	3:M:264:GLY:HA2	1.88	0.55
3:M:262:THR:C	3:M:264:GLY:N	2.55	0.55
3:M:327:PHE:CD2	3:M:430:LEU:HD22	2.41	0.55
3:M:358:ILE:HB	3:M:397:TRP:HE3	1.70	0.55
4:S:75:ILE:HG23	4:S:86:THR:CG2	2.36	0.55
1:A:528:ASN:C	1:A:530:ASN:N	2.52	0.55
1:A:575:LYS:HE3	1:A:611:LEU:CD2	2.35	0.55
1:A:586:GLU:CB	1:A:604:LEU:CD1	2.84	0.55
1:A:609:LEU:HD13	1:A:609:LEU:C	2.26	0.55
2:B:175:LEU:HD11	2:B:210:CYS:CB	2.37	0.55
3:M:124:ILE:HG23	3:M:128:CYS:SG	2.47	0.55
3:M:131:ALA:O	3:M:132:GLY:C	2.42	0.55
3:M:376:ILE:CD1	3:M:415:ILE:HG12	2.37	0.55
4:S:21:THR:HB	4:S:22:PRO:CD	2.33	0.55
1:A:118:SER:O	1:A:122:MET:HG2	2.07	0.55
1:A:182:ILE:HD13	1:A:218:ALA:CA	2.36	0.55
1:A:398:GLU:O	1:A:420:ILE:CB	2.55	0.55
1:A:429:VAL:HG11	1:A:469:LEU:CD1	2.36	0.55
1:A:487:MET:O	1:A:488:ARG:C	2.43	0.55
1:A:624:LEU:O	2:B:617:LEU:CD1	2.54	0.55
2:B:60:ARG:O	2:B:63:MET:N	2.34	0.55
2:B:83:PHE:HE2	2:B:119:SER:N	2.03	0.55
2:B:158:VAL:HG12	2:B:195:ILE:HD13	1.87	0.55
2:B:159:LYS:N	2:B:195:ILE:HD13	2.22	0.55
2:B:286:ILE:O	2:B:287:GLU:C	2.45	0.55
2:B:323:ASN:O	2:B:327:GLN:HG2	2.07	0.55
3:M:2:TYR:OH	3:M:62:VAL:CG1	2.55	0.55
3:M:16:PHE:CE2	3:M:18:TYR:HB2	2.41	0.55
3:M:293:PRO:C	3:M:294:ASP:O	2.30	0.55
3:M:331:LEU:HD22	3:M:426:LYS:NZ	2.22	0.55
4:S:9:ASN:ND2	4:S:13:GLN:HG2	2.17	0.55
1:A:396:VAL:O	1:A:396:VAL:HG12	2.05	0.55
1:A:451:ASN:OD1	1:A:480:LEU:CD1	2.55	0.55
2:B:108:PHE:O	2:B:109:ALA:C	2.31	0.55
2:B:132:SER:C	2:B:169:VAL:CG2	2.76	0.55
2:B:450:VAL:O	2:B:453:TRP:HB2	2.07	0.55
2:B:546:CYS:N	2:B:607:ILE:CG2	2.70	0.55
3:M:100:LEU:HD22	3:M:101:LEU:H	1.71	0.55
3:M:222:PHE:CE2	3:M:439:TYR:CE1	2.94	0.55
3:M:222:PHE:HA	3:M:240:ILE:HA	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:223:HIS:N	3:M:479:PHE:CZ	2.74	0.55
3:M:242:GLY:CA	3:M:474:THR:HG21	2.27	0.55
3:M:269:ILE:C	3:M:302:TYR:CD1	2.80	0.55
4:S:50:PHE:CA	4:S:75:ILE:O	2.55	0.55
1:A:105:VAL:HG21	4:S:167:ILE:O	2.07	0.55
1:A:236:LEU:C	1:A:238:PRO:HD2	2.27	0.55
1:A:401:VAL:HG21	1:A:418:ILE:N	2.21	0.55
1:A:495:ILE:CG2	1:A:515:CYS:HB3	2.37	0.55
1:A:537:THR:HB	1:A:584:PHE:CZ	2.41	0.55
1:A:582:ILE:CG1	1:A:607:LEU:HB2	2.37	0.55
1:A:606:PHE:CZ	2:B:550:VAL:HG13	2.40	0.55
2:B:45:GLN:C	2:B:47:LEU:N	2.59	0.55
2:B:196:LEU:C	2:B:215:TYR:OH	2.46	0.55
2:B:226:LEU:HD12	2:B:226:LEU:O	2.07	0.55
2:B:277:CYS:CA	2:B:292:GLU:CG	2.80	0.55
2:B:392:SER:HB3	2:B:424:PHE:HE2	1.71	0.55
3:M:101:LEU:HD13	3:M:109:LEU:CD1	2.36	0.55
3:M:443:SER:HA	3:M:447:ILE:HG13	1.89	0.55
4:S:54:PRO:CA	4:S:57:LEU:HD13	2.37	0.55
1:A:461:CYS:C	1:A:463:ASP:N	2.51	0.54
2:B:70:MET:HB2	2:B:104:TYR:HE1	1.69	0.54
2:B:120:ILE:CD1	2:B:142:LEU:HD22	2.34	0.54
2:B:132:SER:CB	2:B:169:VAL:CG2	2.85	0.54
2:B:177:ILE:CD1	2:B:196:LEU:HG	2.33	0.54
2:B:193:LEU:O	2:B:195:ILE:CA	2.49	0.54
2:B:249:ILE:HD13	2:B:320:SER:HB2	1.87	0.54
2:B:306:LEU:CD1	2:B:325:LEU:CD2	2.85	0.54
2:B:418:TYR:HD1	2:B:424:PHE:CD1	2.22	0.54
2:B:534:ILE:CG1	2:B:595:VAL:HG23	2.37	0.54
4:S:134:GLU:O	4:S:137:GLN:HG2	2.07	0.54
1:A:105:VAL:CB	4:S:167:ILE:HA	2.37	0.54
1:A:151:SER:CB	1:A:187:LYS:CB	2.74	0.54
1:A:332:TYR:CE1	1:A:336:ILE:HD11	2.40	0.54
1:A:506:LYS:O	1:A:507:GLN:HB2	2.07	0.54
1:A:594:PHE:HD2	2:B:474:VAL:CG2	2.20	0.54
1:A:637:GLU:HG3	2:B:515:PHE:CA	2.37	0.54
2:B:90:ILE:CG1	2:B:98:LYS:CD	2.85	0.54
2:B:316:THR:CB	3:M:90:PHE:CZ	2.90	0.54
2:B:340:ILE:HG21	2:B:373:LEU:O	2.07	0.54
2:B:564:LYS:O	2:B:567:GLN:N	2.40	0.54
2:B:589:SER:HA	2:B:618:PHE:CE2	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:52:ASP:CA	3:M:67:SER:C	2.69	0.54
4:S:89:VAL:HG11	4:S:98:ILE:HG23	1.87	0.54
4:S:131:VAL:HG22	4:S:153:VAL:CG2	2.26	0.54
1:A:92:LEU:HD22	1:A:124:ALA:N	2.23	0.54
1:A:163:ALA:HA	1:A:199:ASN:HD21	1.72	0.54
1:A:185:LEU:HD22	1:A:189:PHE:CE1	2.43	0.54
2:B:248:LEU:O	2:B:252:LEU:CG	2.45	0.54
2:B:449:HIS:O	2:B:453:TRP:CG	2.60	0.54
3:M:101:LEU:CD1	3:M:109:LEU:HD12	2.36	0.54
1:A:101:GLN:O	4:S:167:ILE:HD11	2.08	0.54
1:A:320:HIS:ND1	1:A:352:PHE:CD2	2.76	0.54
1:A:606:PHE:O	1:A:609:LEU:HB3	2.07	0.54
2:B:18:ILE:H	2:B:18:ILE:HD13	1.70	0.54
2:B:28:SER:N	2:B:32:GLU:OE1	2.41	0.54
2:B:155:LEU:CD1	2:B:156:HIS:N	2.70	0.54
2:B:174:ALA:O	2:B:175:LEU:O	2.26	0.54
2:B:247:TYR:OH	3:M:91:THR:HB	2.08	0.54
2:B:415:LEU:O	2:B:418:TYR:N	2.33	0.54
2:B:563:PHE:CD2	2:B:584:SER:C	2.81	0.54
2:B:592:TYR:CG	2:B:593:ASN:N	2.75	0.54
3:M:250:LEU:CD1	3:M:254:PRO:HB2	2.38	0.54
3:M:380:ARG:NH1	3:M:412:ARG:HH11	2.05	0.54
4:S:25:LEU:O	4:S:28:GLN:HG3	2.08	0.54
1:A:170:LEU:HD12	1:A:206:LYS:CG	2.29	0.54
1:A:178:ARG:HH11	1:A:209:ASP:CB	2.10	0.54
1:A:182:ILE:HG21	1:A:218:ALA:N	2.22	0.54
1:A:418:ILE:O	1:A:418:ILE:HG12	2.05	0.54
1:A:505:ASN:O	1:A:506:LYS:C	2.43	0.54
2:B:28:SER:N	2:B:32:GLU:CD	2.55	0.54
2:B:127:LEU:CD2	2:B:157:THR:CG2	2.86	0.54
2:B:362:ALA:HB1	2:B:366:LEU:CD1	2.38	0.54
2:B:418:TYR:CE1	2:B:419:VAL:HA	2.41	0.54
2:B:505:ASP:OD1	2:B:541:GLY:HA3	2.08	0.54
3:M:218:LEU:CG	3:M:472:TYR:HE2	2.16	0.54
4:S:54:PRO:CB	4:S:57:LEU:HD13	2.37	0.54
1:A:129:LYS:CD	1:A:161:ASP:HB3	2.38	0.54
1:A:513:ARG:CB	1:A:550:VAL:HG11	2.37	0.54
2:B:41:ASN:HB3	2:B:43:ASN:CG	2.19	0.54
2:B:105:LEU:O	2:B:106:LEU:O	2.26	0.54
2:B:134:LEU:C	2:B:136:CYS:H	2.11	0.54
2:B:143:SER:HA	2:B:179:LYS:CB	2.32	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:144:ASP:CA	2:B:179:LYS:CD	2.84	0.54
2:B:451:MET:HG3	2:B:489:ILE:HG12	1.90	0.54
2:B:483:PRO:HA	2:B:486:HIS:CB	2.38	0.54
2:B:534:ILE:CD1	2:B:595:VAL:HG23	2.38	0.54
2:B:562:ASN:OD1	2:B:580:TYR:HB2	2.06	0.54
3:M:16:PHE:HE1	3:M:122:SER:HB2	1.71	0.54
3:M:65:TYR:CD2	3:M:86:PRO:CB	2.88	0.54
3:M:476:THR:OG1	3:M:477:GLY:N	2.34	0.54
1:A:100:LEU:HG	4:S:162:SER:N	2.18	0.54
1:A:150:LEU:CD2	1:A:158:LEU:HD11	2.37	0.54
1:A:428:MET:O	1:A:431:VAL:HB	2.07	0.54
1:A:495:ILE:HG23	1:A:515:CYS:HB3	1.89	0.54
1:A:532:LEU:O	1:A:533:ILE:O	2.26	0.54
2:B:159:LYS:CE	2:B:191:GLU:CD	2.75	0.54
2:B:170:ARG:NH1	2:B:198:GLU:C	2.61	0.54
2:B:178:ILE:CB	2:B:214:ALA:CB	2.84	0.54
2:B:197:LYS:O	2:B:198:GLU:C	2.42	0.54
2:B:451:MET:HG3	2:B:489:ILE:CG1	2.37	0.54
3:M:225:VAL:HG22	3:M:480:GLN:CB	2.38	0.54
3:M:265:ASN:O	3:M:266:ASP:C	2.44	0.54
1:A:132:LEU:HD23	1:A:169:MET:SD	2.47	0.54
1:A:255:ARG:C	1:A:257:LEU:H	2.11	0.54
1:A:524:THR:HG21	1:A:565:ASN:ND2	2.22	0.54
1:A:586:GLU:CA	1:A:604:LEU:HD12	2.38	0.54
1:A:602:GLU:CD	1:A:633:PHE:HE1	1.97	0.54
1:A:636:TYR:N	2:B:554:LYS:CD	2.63	0.54
1:A:638:LEU:O	2:B:518:ILE:HD12	2.08	0.54
2:B:117:LEU:CD2	2:B:149:SER:HG	2.07	0.54
2:B:121:ASN:O	2:B:124:GLN:CB	2.56	0.54
2:B:237:ILE:HG13	2:B:245:GLN:HG2	1.90	0.54
2:B:366:LEU:O	2:B:367:SER:O	2.25	0.54
2:B:397:GLN:NE2	2:B:431:MET:SD	2.81	0.54
2:B:418:TYR:CZ	2:B:432:ALA:HB2	2.30	0.54
2:B:455:ILE:O	2:B:459:GLU:N	2.31	0.54
3:M:10:THR:CA	3:M:75:TRP:HE1	2.20	0.54
3:M:101:LEU:HD11	3:M:108:LYS:N	2.23	0.54
4:S:56:SER:O	4:S:60:SER:HB3	1.98	0.54
1:A:102:GLN:O	4:S:167:ILE:CD1	2.56	0.54
1:A:105:VAL:HG22	4:S:167:ILE:HG23	1.81	0.54
1:A:143:VAL:CG1	1:A:169:MET:HB3	2.36	0.54
1:A:440:ASN:CB	1:A:442:SER:HB3	2.37	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:609:LEU:CG	1:A:628:VAL:CG1	2.85	0.54
2:B:79:VAL:C	2:B:108:PHE:HE1	2.12	0.54
2:B:120:ILE:HG22	2:B:153:ILE:CB	2.38	0.54
2:B:335:LYS:O	2:B:335:LYS:HG2	2.08	0.54
2:B:553:ALA:HB2	2:B:614:ILE:CG2	2.38	0.54
3:M:252:ASP:C	3:M:254:PRO:HD2	2.28	0.54
3:M:257:ALA:CB	3:M:455:VAL:CG2	2.85	0.54
3:M:271:SER:HB3	3:M:301:GLU:CG	2.37	0.54
3:M:293:PRO:CA	3:M:294:ASP:O	2.55	0.54
3:M:316:ARG:CG	3:M:322:LEU:HD13	2.37	0.54
3:M:317:MET:HB2	3:M:322:LEU:HB2	1.89	0.54
4:S:12:CYS:HB3	4:S:36:TYR:HB3	1.89	0.54
4:S:14:PRO:HA	4:S:36:TYR:CZ	2.36	0.54
4:S:136:VAL:O	4:S:140:MET:N	2.41	0.54
1:A:64:LEU:HB3	1:A:102:GLN:HE21	1.71	0.54
1:A:103:LYS:O	1:A:104:ARG:C	2.38	0.54
1:A:104:ARG:CG	4:S:126:GLN:OE1	2.47	0.54
1:A:288:THR:CB	1:A:291:ILE:HD13	2.37	0.54
1:A:322:PHE:HB3	1:A:330:LEU:CD2	2.38	0.54
2:B:18:ILE:O	2:B:23:ALA:HB3	2.07	0.54
2:B:155:LEU:CB	2:B:188:TYR:CD2	2.79	0.54
2:B:275:ARG:CB	2:B:291:TYR:HB3	2.13	0.54
2:B:498:THR:CG2	2:B:532:ARG:CB	2.86	0.54
3:M:243:ILE:HG13	3:M:473:LYS:O	2.07	0.54
3:M:372:ILE:O	3:M:372:ILE:HG22	2.07	0.54
1:A:254:ILE:O	1:A:257:LEU:HB2	2.09	0.53
1:A:384:LEU:CD1	1:A:435:ILE:CG2	2.74	0.53
1:A:384:LEU:CG	1:A:441:TYR:CE2	2.85	0.53
1:A:488:ARG:HD2	1:A:522:PHE:CG	2.43	0.53
2:B:124:GLN:OE1	2:B:153:ILE:HG23	2.07	0.53
2:B:158:VAL:HG12	2:B:195:ILE:CG2	2.27	0.53
2:B:275:ARG:HG3	2:B:294:VAL:CB	2.37	0.53
2:B:302:PHE:CE2	2:B:328:LEU:HD11	2.44	0.53
2:B:559:ASP:O	2:B:562:ASN:N	2.41	0.53
2:B:562:ASN:ND2	2:B:580:TYR:CG	2.77	0.53
3:M:215:TYR:CG	3:M:468:LYS:C	2.81	0.53
3:M:272:LEU:N	3:M:272:LEU:HD12	2.23	0.53
3:M:353:VAL:HG13	3:M:353:VAL:O	2.08	0.53
4:S:53:THR:OG1	4:S:68:VAL:N	2.40	0.53
1:A:220:SER:CB	4:S:141:VAL:C	2.75	0.53
1:A:304:LEU:HD22	1:A:344:ILE:CG2	2.36	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:356:ILE:HD13	1:A:374:LEU:HB3	1.89	0.53
1:A:356:ILE:O	1:A:359:LEU:N	2.37	0.53
2:B:185:LYS:HA	2:B:222:HIS:NE2	2.23	0.53
2:B:243:TRP:CH2	3:M:95:THR:N	2.76	0.53
2:B:302:PHE:O	2:B:306:LEU:HG	2.08	0.53
2:B:433:VAL:HG22	2:B:471:TYR:CE2	2.42	0.53
2:B:505:ASP:CA	2:B:544:THR:OG1	2.55	0.53
1:A:401:VAL:HG23	1:A:418:ILE:C	2.28	0.53
1:A:494:ASN:O	1:A:498:LEU:HG	2.08	0.53
1:A:522:PHE:C	1:A:524:THR:H	2.11	0.53
1:A:601:VAL:O	1:A:602:GLU:C	2.31	0.53
2:B:95:THR:OG1	2:B:133:GLU:OE1	2.24	0.53
2:B:189:HIS:HB2	2:B:222:HIS:CE1	2.43	0.53
2:B:237:ILE:CG2	2:B:305:SER:HB3	2.31	0.53
2:B:408:VAL:CG1	2:B:412:PHE:CD2	2.91	0.53
2:B:512:VAL:CB	2:B:551:LEU:HD12	2.36	0.53
2:B:596:LEU:HB2	2:B:615:SER:OG	2.08	0.53
3:M:215:TYR:CE2	3:M:469:GLY:N	2.67	0.53
1:A:561:ASN:O	1:A:562:TRP:C	2.44	0.53
2:B:209:SER:HB2	2:B:244:SER:OG	2.08	0.53
2:B:352:ASN:HD22	2:B:352:ASN:N	1.99	0.53
2:B:363:ILE:CG2	2:B:398:ILE:CG1	2.85	0.53
3:M:67:SER:OG	3:M:90:PHE:HD1	1.31	0.53
3:M:74:TYR:CD1	3:M:76:CYS:SG	3.02	0.53
3:M:217:ASP:O	3:M:472:TYR:CD1	2.62	0.53
3:M:220:GLU:HG2	3:M:439:TYR:HB2	1.89	0.53
3:M:374:TYR:HH	3:M:394:GLN:C	2.09	0.53
4:S:33:GLU:HA	4:S:36:TYR:HD1	1.73	0.53
1:A:80:TYR:CD1	1:A:82:PHE:HE2	2.26	0.53
1:A:182:ILE:HG23	1:A:203:PHE:HE2	1.72	0.53
1:A:498:LEU:HB3	1:A:504:ILE:CG1	2.37	0.53
1:A:610:SER:HG	1:A:625:LEU:HD13	1.72	0.53
2:B:2:VAL:HA	2:B:5:ILE:HD12	1.90	0.53
2:B:296:ASP:OD1	2:B:297:PRO:N	2.41	0.53
2:B:578:PRO:CB	2:B:579:PRO:HD3	2.29	0.53
2:B:586:SER:O	2:B:590:GLN:HG3	2.09	0.53
3:M:5:PHE:CB	3:M:125:PHE:HE2	2.21	0.53
3:M:276:VAL:HG22	3:M:299:LEU:HD12	1.90	0.53
3:M:354:ASP:CB	3:M:440:ILE:CD1	2.85	0.53
1:A:263:LEU:O	1:A:266:VAL:C	2.46	0.53
1:A:338:PHE:HE2	1:A:352:PHE:CE2	2.27	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:589:SER:HB3	1:A:601:VAL:HG22	1.91	0.53
2:B:182:ARG:N	2:B:218:CYS:HA	2.23	0.53
2:B:350:THR:CB	2:B:352:ASN:ND2	2.61	0.53
1:A:163:ALA:CB	1:A:199:ASN:ND2	2.62	0.53
1:A:171:ASN:OD1	1:A:202:LYS:NZ	2.38	0.53
1:A:252:ILE:CB	4:S:144:THR:OG1	2.54	0.53
1:A:408:ILE:CD1	1:A:410:TYR:CD1	2.90	0.53
1:A:585:PHE:CE2	1:A:603:VAL:HG11	2.44	0.53
2:B:83:PHE:CE1	2:B:105:LEU:HA	2.43	0.53
2:B:120:ILE:HG12	2:B:150:LEU:HD22	1.89	0.53
2:B:133:GLU:CA	2:B:168:MET:SD	2.96	0.53
2:B:136:CYS:CA	2:B:172:GLU:CB	2.86	0.53
2:B:184:GLY:O	2:B:188:TYR:CD1	2.62	0.53
2:B:523:PHE:CE1	2:B:580:TYR:CD2	2.58	0.53
3:M:99:ILE:HG22	3:M:103:TYR:CE1	2.44	0.53
3:M:276:VAL:HG21	3:M:299:LEU:HA	1.91	0.53
3:M:306:LEU:CD2	3:M:317:MET:HE3	2.38	0.53
1:A:104:ARG:CB	1:A:145:ILE:HG13	2.39	0.53
1:A:104:ARG:HD3	1:A:145:ILE:HG13	1.91	0.53
1:A:105:VAL:HG21	4:S:167:ILE:C	2.30	0.53
1:A:245:VAL:HG13	1:A:246:THR:HG23	1.90	0.53
1:A:609:LEU:CD2	1:A:628:VAL:CG1	2.85	0.53
1:A:637:GLU:HB2	2:B:513:TRP:CD1	2.43	0.53
2:B:108:PHE:CE2	2:B:115:LEU:HD23	2.44	0.53
2:B:136:CYS:O	2:B:139:LEU:N	2.40	0.53
2:B:151:ALA:N	2:B:152:PRO:CD	2.71	0.53
2:B:167:ALA:O	2:B:207:VAL:HG21	2.09	0.53
2:B:523:PHE:CE1	2:B:582:ASP:OD1	2.62	0.53
2:B:594:ALA:O	2:B:598:LEU:HG	2.07	0.53
3:M:74:TYR:CB	3:M:114:ILE:HD11	2.37	0.53
3:M:258:VAL:HG13	3:M:449:VAL:CG1	2.39	0.53
3:M:293:PRO:CB	3:M:294:ASP:O	2.55	0.53
4:S:45:ASP:C	4:S:47:GLN:H	2.12	0.53
1:A:88:ASN:CG	1:A:120:ILE:HG21	2.29	0.53
1:A:204:VAL:HG13	1:A:239:LEU:CD1	2.39	0.53
1:A:435:ILE:HG23	1:A:441:TYR:CE2	2.44	0.53
1:A:481:MET:CE	1:A:518:CYS:HB3	2.36	0.53
2:B:38:TYR:CD1	2:B:38:TYR:N	2.76	0.53
2:B:189:HIS:HB2	2:B:222:HIS:CD2	2.43	0.53
2:B:189:HIS:CG	2:B:222:HIS:CG	2.96	0.53
2:B:277:CYS:HB2	2:B:295:ASN:C	2.28	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:292:GLU:OE2	2:B:296:ASP:CG	2.46	0.53
2:B:464:SER:OG	2:B:467:VAL:HG23	2.08	0.53
3:M:7:ILE:CG2	3:M:74:TYR:HB2	2.39	0.53
3:M:220:GLU:HA	3:M:474:THR:HG21	1.91	0.53
3:M:263:MET:HG2	3:M:263:MET:O	2.09	0.53
1:A:101:GLN:HG3	4:S:167:ILE:HG21	1.80	0.53
1:A:125:THR:HG21	1:A:158:LEU:HA	1.90	0.53
1:A:463:ASP:CA	2:B:1:MET:SD	2.96	0.53
1:A:581:LEU:CG	1:A:607:LEU:HD11	2.38	0.53
2:B:80:GLN:HG2	2:B:115:LEU:HD21	1.90	0.53
2:B:306:LEU:CB	2:B:325:LEU:HD21	2.38	0.53
2:B:560:ILE:N	2:B:563:PHE:HB2	2.24	0.53
3:M:215:TYR:HB2	3:M:467:TYR:CG	2.42	0.53
3:M:242:GLY:N	3:M:444:ALA:HB2	2.23	0.53
4:S:1:MET:N	4:S:93:GLU:OE1	2.41	0.53
4:S:16:LEU:HD11	4:S:129:GLU:HG2	1.91	0.53
4:S:39:ILE:HD12	4:S:48:SER:HA	1.90	0.53
1:A:366:SER:O	1:A:370:LYS:CG	2.49	0.52
1:A:512:LEU:HD13	1:A:543:TYR:CE1	2.44	0.52
2:B:65:ARG:O	2:B:68:SER:N	2.34	0.52
2:B:87:VAL:HG22	2:B:119:SER:HA	1.89	0.52
2:B:100:LEU:O	2:B:103:LEU:HB2	2.10	0.52
2:B:132:SER:CA	2:B:169:VAL:HG23	2.38	0.52
2:B:155:LEU:HD21	2:B:192:LEU:H	1.75	0.52
2:B:196:LEU:HB3	2:B:215:TYR:HE2	1.62	0.52
2:B:400:SER:HA	2:B:439:CYS:SG	2.48	0.52
2:B:493:LEU:HD11	2:B:511:ILE:HG12	1.90	0.52
3:M:372:ILE:HD12	3:M:428:VAL:HG22	1.89	0.52
3:M:437:TYR:CE1	3:M:479:PHE:CD1	2.97	0.52
4:S:6:LEU:HD22	4:S:32:LEU:CD2	2.29	0.52
4:S:34:GLN:HB3	4:S:58:LEU:HD11	1.89	0.52
1:A:103:LYS:CB	1:A:107:TYR:CE1	2.87	0.52
1:A:557:LYS:HD3	2:B:605:PHE:CD2	2.44	0.52
1:A:588:LEU:O	1:A:589:SER:C	2.42	0.52
2:B:42:ILE:HG22	2:B:43:ASN:N	2.24	0.52
2:B:157:THR:HG22	2:B:158:VAL:N	2.24	0.52
4:S:76:ILE:O	4:S:86:THR:HA	2.09	0.52
1:A:140:VAL:HG13	1:A:176:TYR:HB3	1.86	0.52
1:A:204:VAL:HG23	1:A:236:LEU:HD21	1.80	0.52
1:A:309:PHE:CE1	1:A:348:PHE:CE2	2.97	0.52
1:A:400:VAL:HA	1:A:403:LEU:HD12	1.90	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:420:ILE:C	1:A:421:PRO:O	2.48	0.52
1:A:536:MET:O	1:A:537:THR:O	2.24	0.52
1:A:600:SER:C	1:A:602:GLU:H	2.11	0.52
2:B:36:THR:O	2:B:40:GLN:HG3	2.08	0.52
2:B:36:THR:CA	2:B:39:SER:OG	2.57	0.52
2:B:136:CYS:HB3	2:B:172:GLU:CG	2.39	0.52
2:B:181:TYR:C	2:B:218:CYS:HA	2.30	0.52
2:B:352:ASN:ND2	2:B:352:ASN:O	2.43	0.52
2:B:461:HIS:CB	2:B:463:LEU:HD23	2.39	0.52
2:B:519:ALA:CB	2:B:555:LEU:CD1	2.88	0.52
2:B:546:CYS:O	2:B:549:LEU:HB3	2.09	0.52
3:M:267:ILE:CD1	3:M:445:SER:O	2.57	0.52
4:S:9:ASN:OD1	4:S:13:GLN:CB	2.57	0.52
1:A:104:ARG:CB	1:A:145:ILE:HG21	2.37	0.52
1:A:244:LEU:HD11	1:A:281:LEU:HD13	1.88	0.52
1:A:401:VAL:HG23	1:A:418:ILE:N	2.24	0.52
2:B:123:LEU:HD23	2:B:138:ALA:HB1	1.91	0.52
2:B:132:SER:HB2	2:B:169:VAL:CG2	2.38	0.52
2:B:143:SER:HB2	2:B:179:LYS:HB2	1.91	0.52
2:B:170:ARG:CG	2:B:199:LEU:HD23	2.34	0.52
2:B:184:GLY:O	2:B:188:TYR:HD1	1.92	0.52
2:B:197:LYS:O	2:B:199:LEU:CA	2.56	0.52
2:B:256:CYS:O	2:B:260:LEU:HG	2.10	0.52
2:B:513:TRP:HA	2:B:551:LEU:HD13	1.60	0.52
2:B:542:PRO:HA	2:B:602:ASP:OD1	2.08	0.52
3:M:104:PHE:HD2	3:M:105:ASP:OD2	1.93	0.52
1:A:117:ASP:O	1:A:120:ILE:HB	2.09	0.52
1:A:251:TRP:CG	4:S:104:THR:OG1	2.53	0.52
1:A:581:LEU:HD23	1:A:607:LEU:HD13	1.92	0.52
1:A:589:SER:O	1:A:597:GLN:CG	2.45	0.52
2:B:94:ASP:O	2:B:134:LEU:CD2	2.57	0.52
2:B:98:LYS:HG3	2:B:102:HIS:CE1	2.44	0.52
2:B:175:LEU:CD2	2:B:210:CYS:CB	2.87	0.52
2:B:219:TYR:HD1	2:B:226:LEU:HD13	1.54	0.52
2:B:275:ARG:NE	2:B:275:ARG:CA	2.73	0.52
2:B:404:ASN:O	2:B:405:GLU:O	2.24	0.52
2:B:542:PRO:CB	2:B:602:ASP:OD1	2.58	0.52
2:B:566:ALA:CB	2:B:581:TYR:HA	2.39	0.52
3:M:7:ILE:O	3:M:15:ILE:N	2.41	0.52
3:M:53:HIS:CB	3:M:65:TYR:CE1	2.93	0.52
1:A:100:LEU:H	4:S:162:SER:HB2	0.43	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:212:ILE:CG1	4:S:145:ASN:HD22	2.21	0.52
1:A:431:VAL:O	1:A:434:SER:HB2	2.09	0.52
1:A:463:ASP:HA	2:B:1:MET:SD	2.49	0.52
1:A:528:ASN:O	1:A:529:GLY:O	2.27	0.52
1:A:554:ALA:O	1:A:557:LYS:HB2	2.10	0.52
2:B:21:GLU:HA	2:B:24:ALA:HB3	1.39	0.52
2:B:62:ALA:CB	2:B:66:ILE:HD11	2.37	0.52
2:B:92:THR:O	2:B:134:LEU:CD1	2.58	0.52
2:B:476:ARG:O	2:B:480:GLN:HG3	2.10	0.52
2:B:584:SER:O	2:B:585:GLY:O	2.27	0.52
3:M:7:ILE:CG1	3:M:76:CYS:SG	2.91	0.52
3:M:80:THR:O	3:M:81:SER:HB3	2.10	0.52
3:M:455:VAL:HG12	3:M:455:VAL:O	2.09	0.52
4:S:10:LYS:HD3	4:S:82:THR:HB	1.92	0.52
4:S:126:GLN:NE2	4:S:127:THR:OG1	2.42	0.52
1:A:348:PHE:HB3	1:A:352:PHE:HE1	1.74	0.52
1:A:426:ILE:HD13	1:A:466:ASP:OD2	2.09	0.52
1:A:630:PRO:O	2:B:554:LYS:CB	2.56	0.52
2:B:63:MET:CG	2:B:100:LEU:HB2	2.39	0.52
2:B:189:HIS:CD2	2:B:189:HIS:O	2.63	0.52
2:B:234:CYS:CB	2:B:301:LEU:HB3	2.40	0.52
2:B:243:TRP:HZ2	3:M:95:THR:N	1.90	0.52
2:B:295:ASN:O	2:B:296:ASP:CB	2.57	0.52
2:B:368:ILE:HD11	2:B:401:THR:HG22	1.91	0.52
2:B:596:LEU:O	2:B:599:ALA:N	2.42	0.52
3:M:5:PHE:CB	3:M:125:PHE:CE2	2.93	0.52
3:M:74:TYR:CD2	3:M:109:LEU:O	2.63	0.52
3:M:104:PHE:HZ	3:M:113:LYS:NZ	0.39	0.52
3:M:347:PHE:HE2	3:M:352:GLN:CA	2.22	0.52
4:S:4:ALA:CB	4:S:18:LYS:O	2.58	0.52
1:A:249:ASN:ND2	4:S:146:VAL:H	2.08	0.52
1:A:301:GLY:O	1:A:302:ASN:HB3	2.09	0.52
1:A:585:PHE:CD2	1:A:607:LEU:HD11	2.44	0.52
2:B:4:SER:O	2:B:8:ILE:HG13	2.09	0.52
2:B:123:LEU:CB	2:B:142:LEU:CD1	2.84	0.52
2:B:174:ALA:C	2:B:214:ALA:CB	2.77	0.52
2:B:189:HIS:CG	2:B:189:HIS:O	2.62	0.52
2:B:196:LEU:HB2	2:B:229:HIS:HE1	1.61	0.52
2:B:208:ILE:HD11	2:B:236:ILE:HG23	1.83	0.52
2:B:223:LEU:CD1	2:B:258:GLN:O	2.58	0.52
2:B:501:THR:HA	2:B:508:ARG:NH2	2.20	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:537:PHE:CZ	2:B:598:LEU:O	2.62	0.52
3:M:82:LYS:O	3:M:83:SER:C	2.48	0.52
3:M:270:PRO:HA	3:M:302:TYR:HD1	1.74	0.52
4:S:50:PHE:HA	4:S:76:ILE:HA	1.91	0.52
1:A:92:LEU:CD1	1:A:123:LEU:HD12	2.32	0.52
1:A:260:PHE:O	1:A:262:ASN:N	2.40	0.52
1:A:566:PHE:CZ	1:A:618:THR:O	2.63	0.52
1:A:637:GLU:CB	2:B:516:GLY:H	2.05	0.52
2:B:98:LYS:HG3	2:B:137:PHE:HB2	1.91	0.52
2:B:140:SER:HA	2:B:172:GLU:OE1	2.05	0.52
2:B:227:HIS:CE1	2:B:292:GLU:OE2	2.45	0.52
2:B:451:MET:HG3	2:B:489:ILE:HD13	1.88	0.52
2:B:500:GLN:HB2	2:B:503:LEU:HG	1.89	0.52
2:B:559:ASP:OD2	2:B:582:ASP:OD2	2.28	0.52
3:M:10:THR:CG2	3:M:11:LYS:N	2.73	0.52
3:M:44:ASP:C	3:M:46:SER:H	2.13	0.52
3:M:102:GLU:HA	3:M:106:LYS:HB2	1.91	0.52
3:M:443:SER:HB3	3:M:447:ILE:H	1.74	0.52
1:A:100:LEU:HB3	1:A:142:LYS:HG3	1.91	0.52
1:A:384:LEU:CD2	1:A:441:TYR:CD2	2.86	0.52
1:A:441:TYR:HD1	1:A:441:TYR:H	1.58	0.52
1:A:516:ILE:CD1	1:A:551:LEU:CD1	2.86	0.52
2:B:36:THR:O	2:B:40:GLN:N	2.43	0.52
2:B:139:LEU:CD2	2:B:172:GLU:O	2.55	0.52
2:B:216:LYS:HB2	2:B:251:LEU:CD2	2.40	0.52
2:B:307:ASN:CG	2:B:339:PHE:CE2	2.82	0.52
2:B:550:VAL:HA	2:B:614:ILE:HD11	1.91	0.52
3:M:222:PHE:CA	3:M:479:PHE:HZ	2.17	0.52
3:M:258:VAL:HG22	3:M:452:ILE:HG23	1.90	0.52
3:M:280:ASP:OD1	3:M:281:GLY:N	2.43	0.52
3:M:317:MET:O	3:M:322:LEU:HB3	2.10	0.52
3:M:368:ASP:O	3:M:371:GLU:HB2	2.10	0.52
4:S:83:LEU:HD11	4:S:116:VAL:CB	2.39	0.52
1:A:67:LYS:CB	4:S:166:LYS:CA	2.87	0.51
1:A:91:ILE:O	1:A:94:VAL:HB	2.10	0.51
1:A:102:GLN:CB	4:S:166:LYS:CB	2.56	0.51
1:A:320:HIS:CA	1:A:338:PHE:HZ	2.22	0.51
1:A:566:PHE:HZ	1:A:618:THR:CA	2.22	0.51
1:A:624:LEU:CA	2:B:617:LEU:HD21	2.40	0.51
2:B:70:MET:CE	2:B:107:ARG:HG3	2.23	0.51
2:B:196:LEU:HB2	2:B:229:HIS:HD1	1.73	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:344:VAL:CG1	2:B:381:PHE:CE1	2.93	0.51
2:B:403:ILE:CG2	2:B:411:ILE:HD12	2.39	0.51
2:B:537:PHE:C	2:B:539:ASN:N	2.62	0.51
3:M:100:LEU:C	3:M:109:LEU:HD21	2.30	0.51
3:M:379:LEU:N	3:M:379:LEU:CD1	2.73	0.51
4:S:57:LEU:CD1	4:S:57:LEU:N	2.73	0.51
1:A:140:VAL:CG2	1:A:177:ILE:CD1	2.88	0.51
1:A:187:LYS:O	1:A:190:LEU:HB3	2.09	0.51
1:A:190:LEU:HD12	1:A:228:LYS:HE3	1.91	0.51
1:A:224:GLU:HG2	4:S:138:GLY:HA2	1.88	0.51
1:A:298:ILE:HG13	1:A:311:THR:HG22	1.91	0.51
1:A:403:LEU:CD1	1:A:421:PRO:O	2.55	0.51
1:A:425:LYS:HB3	1:A:464:ILE:HD12	1.93	0.51
1:A:626:SER:O	2:B:614:ILE:HA	2.10	0.51
2:B:60:ARG:CD	2:B:96:LYS:CG	2.87	0.51
2:B:123:LEU:C	2:B:127:LEU:HD12	2.30	0.51
2:B:165:PRO:C	2:B:170:ARG:HE	2.12	0.51
2:B:173:VAL:CB	2:B:199:LEU:HD11	2.40	0.51
3:M:343:ASN:CB	3:M:408:VAL:HG13	2.40	0.51
4:S:47:GLN:OE1	4:S:84:TYR:CD2	2.63	0.51
1:A:71:VAL:HG12	1:A:105:VAL:CG1	2.31	0.51
1:A:185:LEU:HB3	1:A:189:PHE:CZ	2.46	0.51
2:B:47:LEU:HD23	2:B:62:ALA:O	2.11	0.51
2:B:81:LEU:C	2:B:83:PHE:H	2.13	0.51
2:B:108:PHE:CE1	2:B:115:LEU:HG	2.31	0.51
2:B:112:ASP:C	2:B:112:ASP:OD1	2.49	0.51
2:B:123:LEU:HB3	2:B:142:LEU:CD1	2.39	0.51
2:B:208:ILE:O	2:B:209:SER:C	2.48	0.51
2:B:246:SER:O	2:B:249:ILE:HB	2.09	0.51
2:B:393:ILE:HG21	2:B:431:MET:CE	2.41	0.51
3:M:6:TYR:HB3	3:M:14:LEU:HG	1.92	0.51
3:M:18:TYR:HE2	3:M:20:LEU:HD21	1.73	0.51
3:M:379:LEU:CD2	3:M:386:PHE:HD1	1.99	0.51
1:A:261:THR:O	1:A:264:SER:OG	2.17	0.51
1:A:624:LEU:O	1:A:627:GLU:HB2	2.11	0.51
2:B:24:ALA:CB	2:B:35:TYR:CG	2.92	0.51
2:B:132:SER:CB	2:B:166:SER:HB3	2.39	0.51
2:B:135:ARG:NH2	2:B:164:ASP:HB2	2.24	0.51
2:B:162:VAL:CG1	2:B:195:ILE:HA	2.37	0.51
2:B:313:SER:OG	3:M:301:GLU:OE2	2.20	0.51
2:B:344:VAL:CG1	2:B:381:PHE:HZ	2.12	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:553:ALA:HB2	2:B:614:ILE:HG21	1.91	0.51
4:S:87:PHE:HB2	4:S:102:ILE:HD11	1.92	0.51
1:A:144:GLY:CA	1:A:180:LYS:CB	2.82	0.51
1:A:418:ILE:O	1:A:419:ILE:HB	2.09	0.51
1:A:557:LYS:CG	2:B:605:PHE:CD2	2.81	0.51
1:A:579:LYS:O	1:A:582:ILE:HB	2.09	0.51
1:A:633:PHE:CD1	2:B:513:TRP:HZ3	2.16	0.51
1:A:638:LEU:HD11	2:B:518:ILE:HG23	1.88	0.51
2:B:15:ALA:O	2:B:17:VAL:HA	2.10	0.51
2:B:189:HIS:HB2	2:B:222:HIS:NE2	2.26	0.51
2:B:341:GLU:HG2	2:B:345:ARG:NE	2.22	0.51
2:B:360:LEU:CD1	2:B:391:ALA:CB	2.87	0.51
2:B:501:THR:CA	2:B:508:ARG:HH22	2.19	0.51
3:M:72:LEU:CD1	3:M:101:LEU:HD22	2.41	0.51
3:M:101:LEU:CD1	3:M:106:LYS:O	2.50	0.51
3:M:380:ARG:NH1	3:M:410:VAL:HG11	2.07	0.51
3:M:443:SER:HG	3:M:447:ILE:HG13	1.75	0.51
4:S:6:LEU:HD12	4:S:16:LEU:O	2.10	0.51
1:A:129:LYS:HD2	1:A:161:ASP:OD2	2.09	0.51
1:A:141:VAL:N	4:S:155:GLU:HB3	2.24	0.51
1:A:158:LEU:HG	1:A:162:ILE:CD1	2.40	0.51
1:A:399:ASP:C	1:A:420:ILE:HB	2.26	0.51
1:A:412:LYS:O	1:A:414:LYS:N	2.44	0.51
1:A:425:LYS:HD2	1:A:428:MET:HE3	1.92	0.51
1:A:563:CYS:CA	1:A:566:PHE:CD2	2.86	0.51
1:A:566:PHE:C	1:A:568:GLU:N	2.59	0.51
2:B:144:ASP:N	2:B:179:LYS:CD	2.71	0.51
2:B:216:LYS:CD	2:B:251:LEU:HA	2.41	0.51
2:B:227:HIS:CG	2:B:292:GLU:CD	2.84	0.51
2:B:408:VAL:HG12	2:B:412:PHE:CD2	2.45	0.51
2:B:522:GLU:O	2:B:522:GLU:CG	2.40	0.51
3:M:2:TYR:CZ	3:M:62:VAL:HG13	2.46	0.51
3:M:60:LEU:HD23	3:M:61:GLU:H	1.68	0.51
3:M:64:LYS:HG2	3:M:79:SER:O	2.10	0.51
3:M:235:LEU:CD2	3:M:307:SER:HA	2.41	0.51
3:M:478:ASN:O	3:M:479:PHE:O	2.29	0.51
4:S:87:PHE:CD2	4:S:102:ILE:HG12	2.45	0.51
1:A:105:VAL:HB	4:S:167:ILE:HA	1.93	0.51
1:A:195:ALA:O	1:A:199:ASN:ND2	2.44	0.51
1:A:295:VAL:CG1	1:A:319:LEU:HD11	2.28	0.51
1:A:450:TYR:CZ	1:A:476:GLN:HG3	2.39	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:522:PHE:C	1:A:524:THR:N	2.60	0.51
1:A:530:ASN:HD21	1:A:573:GLU:HB3	1.75	0.51
1:A:545:HIS:O	1:A:546:SER:O	2.28	0.51
2:B:189:HIS:CG	2:B:222:HIS:ND1	2.79	0.51
2:B:252:LEU:CG	2:B:302:PHE:CD1	2.88	0.51
2:B:530:LEU:HD23	2:B:591:MET:HB3	1.86	0.51
3:M:10:THR:N	3:M:75:TRP:CD1	2.76	0.51
3:M:101:LEU:CA	3:M:106:LYS:HA	2.40	0.51
3:M:218:LEU:CG	3:M:472:TYR:CE2	2.94	0.51
1:A:176:TYR:CG	4:S:155:GLU:CG	2.91	0.51
1:A:253:ILE:CD1	1:A:281:LEU:CB	2.79	0.51
1:A:537:THR:OG1	1:A:584:PHE:CE1	2.63	0.51
2:B:278:PRO:C	2:B:292:GLU:OE1	2.29	0.51
2:B:374:PHE:HA	2:B:377:TYR:HD1	1.75	0.51
2:B:454:LEU:O	2:B:457:HIS:HB2	2.09	0.51
2:B:463:LEU:O	2:B:468:LEU:CD1	2.58	0.51
3:M:3:LEU:N	3:M:3:LEU:CD1	2.73	0.51
3:M:220:GLU:CD	3:M:222:PHE:CE1	2.85	0.51
3:M:342:LEU:HD12	3:M:342:LEU:N	2.25	0.51
1:A:129:LYS:O	1:A:132:LEU:N	2.44	0.51
1:A:638:LEU:HD21	2:B:523:PHE:HB3	1.93	0.51
2:B:59:VAL:HG11	2:B:89:ASN:HB3	1.91	0.51
2:B:275:ARG:HB3	2:B:291:TYR:CG	2.45	0.51
2:B:560:ILE:CA	2:B:563:PHE:HB2	2.41	0.51
2:B:574:ASN:C	2:B:576:GLN:N	2.57	0.51
3:M:220:GLU:CG	3:M:439:TYR:HD1	2.15	0.51
3:M:224:VAL:C	3:M:480:GLN:H	2.07	0.51
3:M:340:LEU:HG	3:M:342:LEU:HD11	1.91	0.51
1:A:65:ASN:O	4:S:165:SER:CA	2.57	0.51
1:A:101:GLN:CD	4:S:167:ILE:CG2	2.77	0.51
1:A:582:ILE:CD1	1:A:608:ARG:CA	2.88	0.51
2:B:38:TYR:HA	2:B:42:ILE:CA	2.41	0.51
2:B:82:TYR:HB2	2:B:104:TYR:OH	2.10	0.51
2:B:196:LEU:HB3	2:B:229:HIS:HD1	1.75	0.51
2:B:294:VAL:HG13	2:B:295:ASN:N	2.26	0.51
2:B:336:ASN:CB	2:B:339:PHE:CE1	2.94	0.51
3:M:101:LEU:HD23	3:M:106:LYS:CG	2.35	0.51
3:M:245:ASP:O	3:M:246:VAL:CG2	2.59	0.51
3:M:350:VAL:HG22	3:M:442:GLN:OE1	2.11	0.51
4:S:10:LYS:HA	4:S:84:TYR:HE1	1.76	0.51
4:S:55:PRO:C	4:S:57:LEU:N	2.59	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:166:LEU:CB	1:A:185:LEU:HD21	2.42	0.50
1:A:219:VAL:C	1:A:259:LEU:HD13	2.23	0.50
1:A:300:LYS:C	1:A:302:ASN:N	2.57	0.50
1:A:438:ALA:O	1:A:441:TYR:CE1	2.64	0.50
1:A:606:PHE:CD2	1:A:629:LEU:CD1	2.78	0.50
2:B:63:MET:HG2	2:B:100:LEU:HB2	1.93	0.50
2:B:63:MET:CA	2:B:66:ILE:HD12	2.41	0.50
2:B:185:LYS:CG	2:B:222:HIS:NE2	2.66	0.50
2:B:219:TYR:HE1	2:B:226:LEU:HD13	1.12	0.50
2:B:389:ILE:O	2:B:393:ILE:HG13	2.10	0.50
3:M:258:VAL:CG2	3:M:452:ILE:CG2	2.86	0.50
3:M:348:LYS:CG	3:M:405:THR:CG2	2.85	0.50
3:M:376:ILE:HD13	3:M:415:ILE:HG23	1.93	0.50
1:A:83:ASP:CG	1:A:85:ALA:H	2.14	0.50
1:A:107:TYR:OH	1:A:128:LEU:HD23	2.11	0.50
1:A:137:ASN:HA	1:A:139:ASP:HB3	1.93	0.50
1:A:237:SER:N	1:A:238:PRO:CD	2.73	0.50
1:A:309:PHE:CZ	1:A:348:PHE:CZ	2.99	0.50
1:A:461:CYS:SG	1:A:469:LEU:CD2	2.95	0.50
2:B:29:LYS:CD	2:B:30:LEU:N	2.58	0.50
2:B:108:PHE:CD2	2:B:115:LEU:HB3	2.46	0.50
2:B:155:LEU:HD13	2:B:155:LEU:C	2.31	0.50
2:B:159:LYS:HA	2:B:195:ILE:HD13	1.82	0.50
2:B:159:LYS:N	2:B:195:ILE:CD1	2.73	0.50
2:B:281:ASP:C	2:B:283:TYR:N	2.64	0.50
2:B:343:LEU:HD11	2:B:359:LEU:HD13	1.86	0.50
2:B:562:ASN:O	2:B:581:TYR:HA	2.10	0.50
3:M:56:VAL:O	3:M:56:VAL:HG12	2.10	0.50
3:M:101:LEU:CG	3:M:106:LYS:HG3	2.42	0.50
3:M:222:PHE:CZ	3:M:439:TYR:CE1	3.00	0.50
3:M:300:LEU:HD12	3:M:300:LEU:C	2.31	0.50
3:M:383:HIS:CB	3:M:403:THR:OG1	2.56	0.50
4:S:16:LEU:HD22	4:S:125:TRP:HD1	1.76	0.50
1:A:68:THR:CA	4:S:166:LYS:C	2.65	0.50
1:A:103:LYS:C	1:A:107:TYR:CE1	2.85	0.50
1:A:147:LEU:HB2	1:A:184:ALA:HB2	1.92	0.50
1:A:260:PHE:O	1:A:261:THR:O	2.26	0.50
1:A:566:PHE:C	1:A:566:PHE:CD1	2.83	0.50
1:A:595:GLU:O	1:A:599:ARG:HG2	2.11	0.50
2:B:62:ALA:O	2:B:66:ILE:HD11	1.95	0.50
2:B:137:PHE:O	2:B:140:SER:CB	2.59	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:215:TYR:CB	2:B:226:LEU:HD13	2.39	0.50
2:B:219:TYR:OH	2:B:225:LEU:C	2.47	0.50
2:B:344:VAL:CG2	2:B:381:PHE:HZ	2.24	0.50
2:B:501:THR:CA	2:B:508:ARG:NH2	2.75	0.50
2:B:599:ALA:O	2:B:602:ASP:N	2.27	0.50
3:M:1:MET:C	3:M:81:SER:OG	2.50	0.50
3:M:254:PRO:CB	3:M:454:ILE:CD1	2.60	0.50
3:M:265:ASN:HB3	3:M:309:GLN:NE2	2.26	0.50
3:M:348:LYS:HA	3:M:405:THR:C	2.32	0.50
3:M:353:VAL:CG2	3:M:354:ASP:O	2.55	0.50
1:A:125:THR:HG23	1:A:158:LEU:HD12	1.94	0.50
1:A:178:ARG:CB	1:A:214:VAL:HG13	2.39	0.50
1:A:557:LYS:CD	2:B:605:PHE:HD2	2.24	0.50
1:A:559:PHE:CD2	1:A:578:LEU:CD2	2.94	0.50
1:A:581:LEU:CG	1:A:607:LEU:CD1	2.89	0.50
1:A:633:PHE:CD1	2:B:550:VAL:CB	2.93	0.50
2:B:127:LEU:C	2:B:161:LEU:HD11	2.31	0.50
2:B:136:CYS:CB	2:B:168:MET:HG2	2.42	0.50
2:B:175:LEU:CD1	2:B:210:CYS:SG	2.84	0.50
2:B:281:ASP:O	2:B:283:TYR:O	2.29	0.50
2:B:306:LEU:CD2	2:B:321:CYS:SG	3.00	0.50
2:B:337:THR:O	2:B:373:LEU:HD21	2.10	0.50
2:B:352:ASN:HB2	3:M:49:ASP:OD2	1.92	0.50
2:B:371:GLN:C	2:B:373:LEU:H	2.15	0.50
3:M:45:SER:HB3	3:M:51:LEU:CD1	2.40	0.50
3:M:51:LEU:HD13	3:M:75:TRP:CE3	2.43	0.50
3:M:100:LEU:CD1	3:M:100:LEU:N	2.73	0.50
3:M:246:VAL:CA	3:M:470:ALA:HB2	2.34	0.50
3:M:259:LYS:O	3:M:449:VAL:CA	2.53	0.50
3:M:339:GLU:CG	3:M:412:ARG:CG	2.53	0.50
4:S:34:GLN:HB2	4:S:58:LEU:HD11	1.94	0.50
4:S:97:ALA:O	4:S:101:LEU:HB2	2.12	0.50
1:A:371:ALA:O	1:A:374:LEU:HB2	2.12	0.50
1:A:488:ARG:CD	1:A:522:PHE:CD2	2.95	0.50
1:A:547:VAL:O	1:A:550:VAL:HB	2.12	0.50
1:A:581:LEU:CB	1:A:607:LEU:HD13	2.42	0.50
1:A:636:TYR:HB2	2:B:513:TRP:CZ2	2.46	0.50
2:B:25:VAL:HG22	2:B:35:TYR:CD2	2.44	0.50
2:B:68:SER:O	2:B:71:ALA:HB3	2.12	0.50
2:B:132:SER:CB	2:B:166:SER:CB	2.89	0.50
2:B:223:LEU:CB	2:B:259:TYR:CB	2.89	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:316:THR:CB	3:M:90:PHE:CE2	2.91	0.50
2:B:331:PRO:C	2:B:333:GLN:H	2.14	0.50
2:B:553:ALA:CB	2:B:614:ILE:HG23	2.42	0.50
2:B:560:ILE:HD11	2:B:618:PHE:HA	1.92	0.50
2:B:610:ARG:O	2:B:614:ILE:HG13	2.11	0.50
3:M:52:ASP:CB	3:M:67:SER:HA	2.42	0.50
3:M:52:ASP:HB3	3:M:67:SER:HA	1.93	0.50
1:A:96:SER:CB	1:A:127:LEU:HD13	2.18	0.50
1:A:254:ILE:HG21	4:S:100:ASP:OD2	2.12	0.50
1:A:488:ARG:HD2	1:A:522:PHE:CD2	2.47	0.50
1:A:492:ILE:HD11	1:A:522:PHE:CB	2.42	0.50
2:B:21:GLU:OE2	2:B:35:TYR:O	2.30	0.50
2:B:47:LEU:HD23	2:B:66:ILE:HG13	1.69	0.50
2:B:120:ILE:HG21	2:B:150:LEU:O	2.11	0.50
2:B:127:LEU:HA	2:B:135:ARG:HG2	1.93	0.50
2:B:245:GLN:HB3	2:B:309:LEU:CD1	2.41	0.50
2:B:355:ASN:O	2:B:359:LEU:HD22	2.12	0.50
2:B:411:ILE:O	2:B:414:GLU:N	2.23	0.50
2:B:475:ILE:HG23	2:B:489:ILE:HG21	1.93	0.50
3:M:18:TYR:CE2	3:M:20:LEU:HD21	2.47	0.50
3:M:51:LEU:CD1	3:M:51:LEU:N	2.73	0.50
3:M:338:PHE:HE2	3:M:415:ILE:HG12	1.75	0.50
3:M:374:TYR:H	3:M:390:ILE:CG2	2.25	0.50
4:S:70:ASN:CG	4:S:73:ILE:HD12	2.31	0.50
4:S:75:ILE:HG22	4:S:77:TYR:CZ	2.45	0.50
1:A:95:MET:HB2	1:A:127:LEU:HD23	1.92	0.50
1:A:96:SER:N	1:A:127:LEU:CG	2.66	0.50
1:A:101:GLN:HG2	4:S:167:ILE:HB	1.84	0.50
2:B:44:PRO:O	2:B:47:LEU:N	2.45	0.50
2:B:60:ARG:O	2:B:100:LEU:CD1	2.60	0.50
2:B:162:VAL:O	2:B:163:THR:C	2.35	0.50
2:B:267:ASP:CB	2:B:289:PRO:CG	2.90	0.50
2:B:461:HIS:HB2	2:B:463:LEU:CD2	2.42	0.50
2:B:577:ASN:C	2:B:578:PRO:O	2.49	0.50
3:M:215:TYR:CE1	3:M:468:LYS:CB	2.95	0.50
3:M:220:GLU:CD	3:M:222:PHE:CZ	2.85	0.50
3:M:224:VAL:O	3:M:479:PHE:C	2.49	0.50
4:S:16:LEU:HB2	4:S:125:TRP:CD1	2.47	0.50
4:S:17:VAL:CG1	4:S:19:PHE:CZ	2.95	0.50
4:S:75:ILE:HB	4:S:77:TYR:HE1	1.76	0.50
1:A:91:ILE:HG22	1:A:95:MET:HE2	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:92:LEU:HD13	1:A:123:LEU:HB2	1.93	0.50
1:A:141:VAL:CG1	4:S:159:ALA:CA	2.89	0.50
1:A:150:LEU:CD2	1:A:162:ILE:HG12	2.41	0.50
1:A:251:TRP:CE3	1:A:254:ILE:HD12	2.47	0.50
1:A:432:ILE:O	1:A:435:ILE:HB	2.11	0.50
2:B:162:VAL:HG23	2:B:173:VAL:HG11	1.94	0.50
2:B:252:LEU:CB	2:B:302:PHE:CD1	2.93	0.50
2:B:325:LEU:HB3	2:B:334:MET:HE1	1.93	0.50
2:B:512:VAL:O	2:B:515:PHE:HB2	2.12	0.50
3:M:380:ARG:NH1	3:M:412:ARG:NH1	2.60	0.50
1:A:78:GLU:C	1:A:80:TYR:O	2.50	0.50
1:A:145:ILE:HG12	4:S:156:LEU:CD2	2.37	0.50
1:A:200:PHE:CZ	1:A:232:PRO:O	2.62	0.50
1:A:497:LYS:O	1:A:501:ASN:N	2.44	0.50
1:A:556:VAL:HG22	1:A:603:VAL:HG13	1.93	0.50
1:A:573:GLU:O	1:A:575:LYS:N	2.45	0.50
2:B:143:SER:HB2	2:B:179:LYS:CG	2.42	0.50
2:B:144:ASP:N	2:B:179:LYS:HD2	2.22	0.50
2:B:197:LYS:CB	2:B:229:HIS:CD2	2.89	0.50
2:B:452:LYS:NZ	2:B:456:ASP:OD2	2.44	0.50
2:B:553:ALA:CA	2:B:614:ILE:HG23	2.39	0.50
3:M:115:VAL:O	3:M:116:ASN:C	2.46	0.50
3:M:375:LYS:CE	3:M:418:GLU:OE1	2.60	0.50
4:S:78:LYS:O	4:S:84:TYR:HA	2.12	0.50
1:A:166:LEU:O	1:A:170:LEU:HD23	2.12	0.49
1:A:212:ILE:CB	4:S:145:ASN:HD22	2.25	0.49
2:B:256:CYS:CA	2:B:293:VAL:HG21	2.31	0.49
2:B:560:ILE:HA	2:B:564:LYS:N	2.22	0.49
3:M:6:TYR:CG	3:M:14:LEU:HD11	2.46	0.49
3:M:235:LEU:HD23	3:M:235:LEU:O	2.11	0.49
3:M:256:VAL:HG22	3:M:452:ILE:HG23	1.92	0.49
3:M:373:ALA:CB	3:M:418:GLU:O	2.57	0.49
3:M:396:GLN:HG2	3:M:398:ILE:CD1	2.40	0.49
3:M:424:PHE:HZ	3:M:428:VAL:CG2	2.25	0.49
3:M:452:ILE:CD1	3:M:452:ILE:N	2.73	0.49
4:S:4:ALA:HB2	4:S:19:PHE:HD2	1.77	0.49
1:A:104:ARG:HB3	4:S:160:ALA:HB2	1.95	0.49
1:A:133:LYS:HE2	1:A:165:ASP:OD1	2.11	0.49
1:A:139:ASP:OD1	1:A:177:ILE:CD1	2.60	0.49
1:A:186:PHE:HD1	1:A:224:GLU:HB3	1.76	0.49
1:A:349:ILE:O	1:A:352:PHE:N	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:395:PHE:CD2	1:A:420:ILE:HD13	2.47	0.49
1:A:420:ILE:O	1:A:421:PRO:O	2.30	0.49
1:A:433:ILE:HD11	1:A:473:ILE:HA	1.94	0.49
1:A:461:CYS:O	1:A:462:GLN:O	2.30	0.49
1:A:549:GLU:O	1:A:553:LEU:HG	2.12	0.49
2:B:81:LEU:O	2:B:84:ALA:N	2.45	0.49
2:B:109:ALA:HB2	2:B:145:MET:SD	2.53	0.49
2:B:332:LEU:O	2:B:332:LEU:CG	2.60	0.49
2:B:349:MET:O	2:B:351:GLU:OE2	2.30	0.49
2:B:418:TYR:CG	2:B:424:PHE:CE1	3.00	0.49
2:B:494:ALA:HB2	2:B:529:VAL:HG22	1.94	0.49
2:B:534:ILE:HG12	2:B:595:VAL:HG23	1.93	0.49
3:M:260:LEU:CA	3:M:448:TYR:O	2.59	0.49
3:M:288:ILE:HG22	3:M:289:THR:N	2.27	0.49
3:M:323:MET:CE	3:M:342:LEU:CB	2.89	0.49
3:M:379:LEU:CD2	3:M:397:TRP:NE1	2.66	0.49
1:A:100:LEU:HD11	4:S:157:ASN:O	2.12	0.49
1:A:217:ALA:CA	4:S:142:ILE:CA	2.53	0.49
1:A:456:ASP:O	1:A:459:MET:N	2.37	0.49
1:A:602:GLU:OE1	1:A:633:PHE:HE1	1.95	0.49
1:A:623:MET:O	2:B:617:LEU:CG	2.60	0.49
2:B:343:LEU:HD23	2:B:363:ILE:HD13	1.93	0.49
2:B:421:SER:C	2:B:422:ALA:O	2.42	0.49
3:M:244:VAL:CG2	3:M:472:TYR:CE2	2.94	0.49
3:M:296:LYS:HG3	3:M:296:LYS:O	2.12	0.49
3:M:320:ILE:CG2	3:M:321:GLY:N	2.75	0.49
3:M:371:GLU:HB3	3:M:424:PHE:CD1	2.45	0.49
1:A:179:LYS:HG3	1:A:217:ALA:CB	2.35	0.49
1:A:244:LEU:CG	1:A:277:LYS:HG3	2.39	0.49
1:A:441:TYR:HB3	1:A:444:VAL:HG22	1.91	0.49
1:A:630:PRO:CB	2:B:614:ILE:CG2	2.88	0.49
2:B:125:LYS:O	2:B:128:SER:N	2.45	0.49
2:B:132:SER:CB	2:B:169:VAL:HG21	2.41	0.49
2:B:267:ASP:CA	2:B:289:PRO:HB3	2.39	0.49
2:B:284:ASN:HB3	2:B:285:GLU:HG3	1.91	0.49
2:B:334:MET:CE	2:B:339:PHE:CD1	2.94	0.49
3:M:74:TYR:HB3	3:M:114:ILE:HD11	1.95	0.49
3:M:222:PHE:CD1	3:M:222:PHE:N	2.80	0.49
3:M:317:MET:CB	3:M:322:LEU:N	2.65	0.49
4:S:14:PRO:CA	4:S:36:TYR:CZ	2.95	0.49
4:S:17:VAL:HG13	4:S:17:VAL:O	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:51:LEU:HB2	4:S:77:TYR:CE1	2.47	0.49
1:A:153:ILE:CG2	1:A:158:LEU:HD23	2.42	0.49
1:A:403:LEU:HG	1:A:420:ILE:O	2.12	0.49
1:A:440:ASN:OD1	1:A:442:SER:CB	2.58	0.49
1:A:552:ILE:HD13	1:A:600:SER:OG	2.12	0.49
2:B:143:SER:C	2:B:145:MET:H	2.14	0.49
2:B:223:LEU:HD11	2:B:258:GLN:O	2.08	0.49
2:B:267:ASP:OD1	2:B:269:SER:HB2	2.12	0.49
2:B:393:ILE:HG21	2:B:431:MET:HB2	1.95	0.49
3:M:244:VAL:C	3:M:472:TYR:CD2	2.81	0.49
3:M:350:VAL:HG22	3:M:442:GLN:NE2	2.28	0.49
1:A:317:GLU:CG	1:A:351:ARG:HH12	2.26	0.49
1:A:421:PRO:HB2	1:A:424:TYR:HD1	1.78	0.49
2:B:132:SER:O	2:B:136:CYS:SG	2.70	0.49
2:B:221:ASP:O	2:B:223:LEU:HG	2.11	0.49
2:B:257:LYS:HA	2:B:260:LEU:CD2	2.42	0.49
2:B:299:LEU:O	2:B:299:LEU:HD13	2.13	0.49
2:B:315:PRO:CB	2:B:355:ASN:ND2	2.51	0.49
2:B:340:ILE:O	2:B:343:LEU:HB3	2.12	0.49
2:B:424:PHE:CE2	2:B:428:VAL:HG11	2.45	0.49
2:B:427:ASN:HA	2:B:430:ILE:HD12	1.94	0.49
2:B:436:LEU:HB3	2:B:450:VAL:HG13	1.93	0.49
4:S:69:ASN:CG	4:S:71:GLU:O	2.51	0.49
1:A:65:ASN:C	4:S:165:SER:CA	2.81	0.49
1:A:123:LEU:O	1:A:127:LEU:HB2	2.13	0.49
1:A:170:LEU:CG	1:A:206:LYS:HG3	2.42	0.49
1:A:224:GLU:H	4:S:140:MET:HE3	1.78	0.49
1:A:557:LYS:CG	2:B:606:ASP:CA	2.89	0.49
2:B:120:ILE:CG2	2:B:153:ILE:HB	2.41	0.49
2:B:132:SER:O	2:B:169:VAL:CG2	2.60	0.49
2:B:227:HIS:O	2:B:230:PHE:N	2.44	0.49
2:B:292:GLU:CG	2:B:295:ASN:O	2.60	0.49
2:B:362:ALA:O	2:B:366:LEU:CG	2.58	0.49
2:B:455:ILE:O	2:B:458:MET:N	2.46	0.49
2:B:494:ALA:CB	2:B:529:VAL:HG13	2.43	0.49
2:B:513:TRP:HE1	2:B:517:GLU:HG3	1.77	0.49
3:M:103:TYR:CG	3:M:124:ILE:HD11	2.48	0.49
3:M:106:LYS:O	3:M:106:LYS:HG2	2.13	0.49
3:M:218:LEU:HB3	3:M:442:GLN:O	2.11	0.49
3:M:315:VAL:O	3:M:315:VAL:HG12	2.11	0.49
3:M:443:SER:HB2	3:M:447:ILE:N	2.27	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:9:ASN:CG	4:S:13:GLN:CB	2.79	0.49
1:A:65:ASN:C	4:S:165:SER:HA	2.33	0.49
1:A:67:LYS:CD	1:A:102:GLN:OE1	2.60	0.49
1:A:589:SER:HB2	1:A:601:VAL:HG23	1.92	0.49
1:A:624:LEU:O	2:B:613:MET:CE	2.50	0.49
2:B:106:LEU:HD22	2:B:144:ASP:OD2	2.13	0.49
2:B:123:LEU:HD22	2:B:138:ALA:CB	2.43	0.49
2:B:237:ILE:HG21	2:B:305:SER:CB	2.35	0.49
2:B:275:ARG:CG	2:B:294:VAL:HG21	2.42	0.49
2:B:343:LEU:HD21	2:B:363:ILE:N	2.27	0.49
2:B:346:THR:HG23	2:B:350:THR:CG2	2.30	0.49
2:B:389:ILE:CG2	2:B:427:ASN:CB	2.81	0.49
3:M:225:VAL:C	3:M:480:GLN:O	2.50	0.49
3:M:265:ASN:O	3:M:267:ILE:N	2.45	0.49
1:A:84:MET:SD	1:A:113:SER:HA	2.52	0.49
2:B:150:LEU:O	2:B:154:ILE:HD12	2.08	0.49
2:B:178:ILE:HG23	2:B:217:GLU:CA	2.43	0.49
2:B:200:MET:O	2:B:202:ASP:N	2.45	0.49
2:B:394:TRP:CE3	2:B:397:GLN:OE1	2.64	0.49
2:B:513:TRP:CE3	2:B:551:LEU:HD21	2.40	0.49
2:B:572:GLU:O	2:B:575:ASN:N	2.28	0.49
3:M:101:LEU:HB2	3:M:109:LEU:HD13	1.94	0.49
3:M:276:VAL:HG21	3:M:299:LEU:HD12	1.93	0.49
3:M:331:LEU:HD22	3:M:426:LYS:HZ2	1.77	0.49
4:S:33:GLU:O	4:S:36:TYR:HB2	2.12	0.49
1:A:275:LEU:HA	1:A:278:ILE:CG1	2.43	0.49
1:A:398:GLU:O	1:A:419:ILE:CA	2.61	0.49
1:A:627:GLU:HB3	2:B:617:LEU:HB3	1.90	0.49
1:A:633:PHE:CB	2:B:550:VAL:O	2.58	0.49
2:B:80:GLN:HA	2:B:115:LEU:HD11	1.95	0.49
2:B:151:ALA:HB2	2:B:180:LEU:O	2.13	0.49
2:B:559:ASP:O	2:B:562:ASN:CB	2.61	0.49
3:M:1:MET:HB3	3:M:81:SER:HB3	1.95	0.49
3:M:235:LEU:HD11	3:M:306:LEU:CD1	2.42	0.49
1:A:105:VAL:O	1:A:108:LEU:HB2	2.13	0.48
1:A:252:ILE:HA	4:S:144:THR:CB	2.30	0.48
1:A:319:LEU:O	1:A:321:THR:N	2.46	0.48
1:A:582:ILE:HD11	1:A:608:ARG:N	2.28	0.48
1:A:589:SER:CB	1:A:601:VAL:HG23	2.42	0.48
2:B:38:TYR:HD1	2:B:38:TYR:N	2.10	0.48
2:B:77:ILE:HG23	2:B:82:TYR:CE1	2.42	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:92:THR:HG22	2:B:93:ASN:N	2.27	0.48
2:B:143:SER:C	2:B:145:MET:N	2.66	0.48
2:B:231:ARG:C	2:B:233:TYR:N	2.65	0.48
2:B:353:GLN:HB2	3:M:50:TYR:H	1.71	0.48
2:B:520:SER:C	2:B:523:PHE:CD2	2.86	0.48
3:M:3:LEU:CD1	3:M:80:THR:O	2.61	0.48
3:M:101:LEU:CG	3:M:106:LYS:CG	2.91	0.48
3:M:224:VAL:CG1	3:M:226:PHE:CZ	2.89	0.48
3:M:226:PHE:CD2	3:M:235:LEU:HA	2.48	0.48
3:M:245:ASP:C	3:M:246:VAL:CG2	2.81	0.48
3:M:389:SER:N	3:M:394:GLN:O	2.33	0.48
1:A:179:LYS:HD3	4:S:152:SER:OG	2.13	0.48
1:A:408:ILE:HG21	4:S:64:ASN:HB3	1.82	0.48
1:A:512:LEU:HD13	1:A:543:TYR:CZ	2.48	0.48
2:B:146:LYS:C	2:B:147:MET:CG	2.80	0.48
2:B:174:ALA:O	2:B:214:ALA:HB2	2.07	0.48
2:B:219:TYR:HD1	2:B:226:LEU:CD1	2.06	0.48
2:B:237:ILE:CG2	2:B:238:LYS:N	2.76	0.48
3:M:93:LEU:O	3:M:96:ILE:HB	2.14	0.48
3:M:215:TYR:CD1	3:M:468:LYS:CB	2.94	0.48
3:M:222:PHE:CG	3:M:240:ILE:HG12	2.48	0.48
3:M:235:LEU:HD13	3:M:310:VAL:HG21	1.95	0.48
3:M:386:PHE:CB	3:M:397:TRP:HD1	2.26	0.48
3:M:435:LEU:O	3:M:437:TYR:CE1	2.66	0.48
4:S:53:THR:CG2	4:S:68:VAL:C	2.80	0.48
1:A:102:GLN:O	4:S:167:ILE:HD11	2.13	0.48
2:B:5:ILE:HA	2:B:8:ILE:CD1	2.38	0.48
2:B:56:SER:O	2:B:97:VAL:CG2	2.61	0.48
2:B:293:VAL:O	2:B:299:LEU:HB3	2.04	0.48
2:B:418:TYR:HD1	2:B:424:PHE:CE1	2.26	0.48
2:B:447:GLU:O	2:B:450:VAL:HB	2.13	0.48
2:B:580:TYR:O	2:B:582:ASP:N	2.43	0.48
3:M:7:ILE:HA	3:M:76:CYS:CA	2.28	0.48
3:M:226:PHE:CD1	3:M:481:VAL:HG22	2.49	0.48
3:M:235:LEU:HD11	3:M:306:LEU:HD13	1.94	0.48
4:S:80:TYR:CG	4:S:106:VAL:CG1	2.96	0.48
1:A:316:LEU:HD12	1:A:348:PHE:CZ	2.45	0.48
2:B:98:LYS:HD2	2:B:102:HIS:CE1	2.48	0.48
2:B:230:PHE:CE2	2:B:298:ASP:C	2.77	0.48
2:B:302:PHE:CE1	2:B:306:LEU:HD21	2.48	0.48
2:B:389:ILE:HG21	2:B:427:ASN:CB	2.35	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:513:TRP:NE1	2:B:517:GLU:CG	2.76	0.48
2:B:538:SER:CA	2:B:598:LEU:HD22	2.43	0.48
2:B:542:PRO:O	2:B:545:ARG:HB2	2.14	0.48
2:B:589:SER:OG	2:B:618:PHE:HE2	1.90	0.48
3:M:224:VAL:HG22	3:M:306:LEU:CD1	2.41	0.48
3:M:228:LYS:HZ2	3:M:327:PHE:H	1.58	0.48
3:M:249:TYR:CE1	3:M:467:TYR:CE1	3.02	0.48
3:M:323:MET:SD	3:M:342:LEU:CA	3.01	0.48
3:M:338:PHE:CE2	3:M:415:ILE:HD11	2.48	0.48
3:M:347:PHE:HA	3:M:350:VAL:HG23	1.96	0.48
1:A:207:LEU:CD1	1:A:240:LEU:HD23	2.38	0.48
1:A:304:LEU:C	1:A:304:LEU:HD23	2.34	0.48
1:A:384:LEU:C	1:A:384:LEU:HD12	2.34	0.48
1:A:478:ARG:O	1:A:482:ILE:HG13	2.13	0.48
2:B:80:GLN:HG2	2:B:115:LEU:CD2	2.43	0.48
2:B:139:LEU:CD2	2:B:173:VAL:CB	2.87	0.48
2:B:277:CYS:HA	2:B:295:ASN:O	2.13	0.48
2:B:360:LEU:HB3	2:B:394:TRP:HB2	1.95	0.48
2:B:393:ILE:CG1	2:B:428:VAL:HA	2.40	0.48
2:B:541:GLY:O	2:B:544:THR:HB	2.14	0.48
3:M:65:TYR:CG	3:M:86:PRO:CA	2.93	0.48
3:M:240:ILE:HG22	3:M:444:ALA:CB	2.42	0.48
3:M:319:SER:CB	3:M:346:ASN:C	2.81	0.48
3:M:380:ARG:HH11	3:M:412:ARG:HH11	1.59	0.48
1:A:67:LYS:HB3	4:S:166:LYS:HG2	1.86	0.48
1:A:78:GLU:O	1:A:80:TYR:C	2.52	0.48
1:A:121:LEU:HD13	1:A:155:THR:CG2	2.44	0.48
1:A:153:ILE:HG22	1:A:158:LEU:HD23	1.94	0.48
1:A:350:SER:O	1:A:352:PHE:N	2.46	0.48
1:A:384:LEU:HD13	1:A:435:ILE:HG21	1.86	0.48
1:A:509:PRO:CB	1:A:547:VAL:HG21	2.38	0.48
1:A:594:PHE:CG	2:B:434:LYS:HE2	2.48	0.48
2:B:27:THR:HG22	2:B:27:THR:O	2.14	0.48
2:B:56:SER:HB2	2:B:92:THR:CG2	2.30	0.48
2:B:126:SER:O	2:B:127:LEU:C	2.49	0.48
2:B:429:VAL:O	2:B:433:VAL:HG23	2.12	0.48
3:M:241:HIS:CD2	3:M:475:GLN:H	2.29	0.48
4:S:57:LEU:CA	4:S:60:SER:HB2	2.41	0.48
1:A:132:LEU:CB	1:A:165:ASP:HB3	2.44	0.48
1:A:196:LEU:O	1:A:197:ARG:O	2.31	0.48
1:A:320:HIS:ND1	1:A:352:PHE:HD2	2.12	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:402:ILE:O	1:A:402:ILE:HG23	2.12	0.48
1:A:516:ILE:HD13	1:A:551:LEU:CD1	2.43	0.48
1:A:556:VAL:HG22	1:A:603:VAL:HG11	1.94	0.48
1:A:556:VAL:HG21	1:A:603:VAL:HG13	1.96	0.48
2:B:83:PHE:CE2	2:B:119:SER:HA	2.38	0.48
2:B:279:LEU:CB	2:B:288:TYR:HD1	2.23	0.48
2:B:377:TYR:O	2:B:380:LYS:CB	2.49	0.48
2:B:394:TRP:CZ3	2:B:397:GLN:NE2	2.82	0.48
2:B:396:ILE:HD13	2:B:418:TYR:CE2	2.43	0.48
2:B:515:PHE:CD2	2:B:529:VAL:HG21	2.44	0.48
2:B:542:PRO:CA	2:B:602:ASP:OD1	2.61	0.48
1:A:154:ILE:O	1:A:155:THR:C	2.49	0.48
1:A:316:LEU:O	1:A:319:LEU:HB2	2.13	0.48
1:A:416:ILE:O	1:A:417:PRO:C	2.40	0.48
2:B:106:LEU:CG	2:B:144:ASP:CB	2.86	0.48
2:B:227:HIS:CE1	2:B:292:GLU:CG	2.96	0.48
2:B:237:ILE:HG12	2:B:245:GLN:HG2	1.95	0.48
2:B:433:VAL:C	2:B:474:VAL:HG21	2.33	0.48
2:B:440:GLY:C	2:B:442:LEU:N	2.61	0.48
2:B:515:PHE:CE2	2:B:529:VAL:CG1	2.95	0.48
2:B:519:ALA:CB	2:B:555:LEU:HD13	2.44	0.48
2:B:538:SER:HB3	2:B:598:LEU:CD2	2.44	0.48
2:B:589:SER:C	2:B:591:MET:N	2.65	0.48
3:M:6:TYR:CB	3:M:16:PHE:O	2.62	0.48
3:M:217:ASP:C	3:M:472:TYR:CZ	2.87	0.48
3:M:221:THR:HA	3:M:437:TYR:O	2.13	0.48
3:M:234:ARG:O	3:M:235:LEU:C	2.51	0.48
3:M:243:ILE:CG2	3:M:298:ARG:HG2	2.43	0.48
3:M:319:SER:HG	3:M:346:ASN:CB	2.05	0.48
4:S:5:VAL:O	4:S:18:LYS:N	2.44	0.48
4:S:17:VAL:HG13	4:S:19:PHE:HE1	1.74	0.48
1:A:144:GLY:HA3	1:A:180:LYS:HG3	1.96	0.48
1:A:215:VAL:O	1:A:216:SER:O	2.31	0.48
1:A:381:GLU:CA	1:A:384:LEU:HD23	2.39	0.48
1:A:454:ILE:CG2	1:A:473:ILE:CG2	2.92	0.48
1:A:488:ARG:CD	1:A:522:PHE:CE2	2.96	0.48
2:B:80:GLN:CA	2:B:115:LEU:HD11	2.43	0.48
2:B:417:TYR:O	2:B:421:SER:CB	2.62	0.48
3:M:7:ILE:HG21	3:M:74:TYR:HB2	1.96	0.48
3:M:215:TYR:CD1	3:M:468:LYS:CG	2.97	0.48
3:M:234:ARG:O	3:M:236:LEU:N	2.47	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:125:THR:CG2	1:A:158:LEU:HA	2.43	0.48
1:A:306:GLU:O	1:A:308:ASP:N	2.47	0.48
1:A:446:ASP:C	1:A:448:GLU:O	2.52	0.48
2:B:44:PRO:HA	2:B:47:LEU:HD12	1.96	0.48
2:B:155:LEU:O	2:B:158:VAL:N	2.47	0.48
2:B:340:ILE:CB	2:B:373:LEU:HG	2.44	0.48
2:B:380:LYS:NZ	3:M:236:LEU:HD11	2.28	0.48
2:B:522:GLU:C	2:B:524:LYS:N	2.67	0.48
3:M:10:THR:HA	3:M:75:TRP:CE2	2.47	0.48
3:M:10:THR:N	3:M:75:TRP:NE1	2.62	0.48
3:M:260:LEU:O	3:M:286:SER:HB3	2.14	0.48
3:M:300:LEU:HD11	3:M:447:ILE:CD1	2.41	0.48
1:A:100:LEU:CB	1:A:142:LYS:HG3	2.44	0.47
1:A:408:ILE:CD1	1:A:410:TYR:HD1	2.27	0.47
1:A:610:SER:OG	1:A:625:LEU:CD1	2.52	0.47
2:B:136:CYS:HA	2:B:172:GLU:CB	2.41	0.47
2:B:540:GLU:OE1	2:B:548:ILE:CD1	2.62	0.47
2:B:542:PRO:HB3	2:B:602:ASP:OD1	2.14	0.47
3:M:9:ASP:C	3:M:75:TRP:NE1	2.64	0.47
3:M:16:PHE:HZ	3:M:18:TYR:CD1	2.32	0.47
3:M:244:VAL:CA	3:M:472:TYR:CG	2.76	0.47
3:M:324:SER:O	3:M:340:LEU:HA	2.14	0.47
4:S:47:GLN:NE2	4:S:78:LYS:C	2.63	0.47
1:A:79:MET:HG3	1:A:112:GLN:OE1	2.14	0.47
1:A:216:SER:HB3	4:S:143:GLU:CA	2.13	0.47
1:A:384:LEU:HD12	1:A:385:LYS:N	2.28	0.47
1:A:429:VAL:CG1	1:A:473:ILE:HD11	2.40	0.47
1:A:563:CYS:SG	1:A:621:LEU:HG	2.54	0.47
1:A:625:LEU:C	1:A:627:GLU:N	2.68	0.47
1:A:633:PHE:CA	2:B:554:LYS:HB2	2.15	0.47
2:B:117:LEU:HD21	2:B:149:SER:HB2	1.92	0.47
2:B:207:VAL:O	2:B:210:CYS:HB2	2.14	0.47
2:B:278:PRO:CA	2:B:288:TYR:HB2	2.29	0.47
2:B:304:GLN:O	2:B:307:ASN:HB2	2.14	0.47
2:B:414:GLU:O	2:B:417:TYR:HB3	2.14	0.47
2:B:425:PRO:HD2	2:B:428:VAL:CG2	2.39	0.47
3:M:124:ILE:O	3:M:127:CYS:N	2.47	0.47
3:M:245:ASP:N	3:M:472:TYR:CE2	2.82	0.47
3:M:364:VAL:HG12	3:M:372:ILE:HG13	1.95	0.47
4:S:35:VAL:HG12	4:S:68:VAL:CG1	2.44	0.47
4:S:50:PHE:HB2	4:S:75:ILE:O	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:50:PHE:HD1	4:S:52:VAL:HG23	1.79	0.47
1:A:180:LYS:NZ	4:S:156:LEU:HD11	2.29	0.47
1:A:213:SER:C	4:S:143:GLU:CD	2.72	0.47
1:A:220:SER:CA	4:S:142:ILE:HG22	2.37	0.47
1:A:291:ILE:H	1:A:291:ILE:CD1	2.28	0.47
1:A:605:GLU:CD	1:A:632:PHE:CE2	2.88	0.47
2:B:42:ILE:HD13	2:B:65:ARG:CD	2.39	0.47
2:B:260:LEU:HA	2:B:291:TYR:CZ	2.49	0.47
2:B:588:ILE:HG21	2:B:618:PHE:HE1	1.79	0.47
3:M:8:THR:CA	3:M:13:LYS:O	2.62	0.47
3:M:223:HIS:N	3:M:239:SER:O	2.46	0.47
3:M:240:ILE:HG21	3:M:444:ALA:O	2.14	0.47
3:M:252:ASP:O	3:M:254:PRO:HD2	2.14	0.47
1:A:182:ILE:HG23	1:A:221:VAL:HG21	1.83	0.47
1:A:588:LEU:O	1:A:590:TYR:N	2.47	0.47
2:B:155:LEU:C	2:B:157:THR:H	2.17	0.47
2:B:277:CYS:SG	2:B:292:GLU:CD	2.91	0.47
2:B:279:LEU:HD12	2:B:288:TYR:HE1	1.79	0.47
2:B:494:ALA:CB	2:B:529:VAL:HG22	2.45	0.47
3:M:8:THR:O	3:M:75:TRP:HB2	2.14	0.47
3:M:226:PHE:CE2	3:M:235:LEU:HB2	2.50	0.47
3:M:235:LEU:CD1	3:M:306:LEU:HD13	2.44	0.47
3:M:284:SER:O	3:M:286:SER:N	2.48	0.47
4:S:4:ALA:CB	4:S:19:PHE:CD2	2.98	0.47
4:S:69:ASN:O	4:S:70:ASN:HB3	2.14	0.47
4:S:101:LEU:HD11	4:S:136:VAL:HG22	1.96	0.47
1:A:270:LEU:CD1	1:A:274:LEU:CD2	2.92	0.47
1:A:633:PHE:CG	2:B:550:VAL:O	2.68	0.47
2:B:127:LEU:CB	2:B:161:LEU:CD1	2.63	0.47
2:B:352:ASN:ND2	2:B:355:ASN:HB2	2.30	0.47
2:B:534:ILE:O	2:B:598:LEU:CD1	2.56	0.47
2:B:566:ALA:HB1	2:B:581:TYR:HA	1.95	0.47
3:M:6:TYR:HD2	3:M:16:PHE:O	1.94	0.47
3:M:218:LEU:HD12	3:M:218:LEU:H	1.80	0.47
4:S:101:LEU:HG	4:S:135:ILE:HG21	1.96	0.47
4:S:127:THR:HG21	4:S:157:ASN:OD1	2.15	0.47
1:A:489:GLU:HB2	1:A:525:LEU:HD21	1.96	0.47
1:A:586:GLU:O	1:A:588:LEU:N	2.44	0.47
2:B:25:VAL:HG22	2:B:36:THR:N	2.30	0.47
2:B:42:ILE:CD1	2:B:65:ARG:CD	2.81	0.47
2:B:42:ILE:HG23	2:B:46:GLN:OE1	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:141:ALA:C	2:B:143:SER:H	2.18	0.47
2:B:232:ARG:O	2:B:236:ILE:N	2.43	0.47
2:B:243:TRP:HZ3	3:M:91:THR:CA	2.12	0.47
2:B:274:PRO:CB	2:B:295:ASN:HD21	2.28	0.47
2:B:531:ARG:HG3	2:B:591:MET:CE	2.45	0.47
3:M:62:VAL:O	3:M:64:LYS:HG3	2.15	0.47
3:M:273:HIS:HB3	3:M:276:VAL:HG23	1.97	0.47
1:A:141:VAL:CG2	4:S:156:LEU:C	2.75	0.47
1:A:156:PRO:O	1:A:160:ARG:CG	2.62	0.47
1:A:178:ARG:O	1:A:182:ILE:HG13	2.15	0.47
1:A:220:SER:CB	4:S:142:ILE:HG23	2.27	0.47
1:A:316:LEU:HD21	1:A:341:ILE:HG13	1.95	0.47
1:A:388:VAL:HG22	1:A:435:ILE:HD12	1.96	0.47
1:A:463:ASP:C	2:B:1:MET:CG	2.73	0.47
1:A:563:CYS:O	1:A:566:PHE:CE2	2.68	0.47
1:A:609:LEU:HD12	1:A:625:LEU:HD12	1.96	0.47
2:B:25:VAL:HG22	2:B:35:TYR:C	2.35	0.47
2:B:37:TYR:HE2	2:B:46:GLN:OE1	1.97	0.47
2:B:51:LEU:HD11	2:B:66:ILE:HD13	1.95	0.47
2:B:85:ASP:C	2:B:89:ASN:HD22	2.18	0.47
2:B:117:LEU:O	2:B:120:ILE:HB	2.14	0.47
2:B:140:SER:CB	2:B:172:GLU:OE2	2.55	0.47
2:B:216:LYS:CG	2:B:251:LEU:HA	2.45	0.47
2:B:336:ASN:C	2:B:338:LYS:H	2.17	0.47
2:B:421:SER:O	2:B:422:ALA:C	2.49	0.47
2:B:474:VAL:C	2:B:476:ARG:N	2.66	0.47
2:B:523:PHE:HD1	2:B:559:ASP:OD2	1.97	0.47
2:B:534:ILE:HG12	2:B:595:VAL:CG2	2.44	0.47
2:B:585:GLY:O	2:B:589:SER:CB	2.63	0.47
3:M:215:TYR:CD1	3:M:467:TYR:C	2.88	0.47
3:M:222:PHE:CD2	3:M:439:TYR:CZ	3.02	0.47
3:M:243:ILE:H	3:M:474:THR:HG21	1.64	0.47
3:M:244:VAL:O	3:M:299:LEU:CB	2.62	0.47
3:M:276:VAL:CG1	3:M:299:LEU:HD12	2.45	0.47
3:M:348:LYS:HA	3:M:406:GLY:N	2.30	0.47
3:M:353:VAL:HG11	3:M:356:LEU:HB2	1.97	0.47
3:M:362:PHE:CE1	3:M:374:TYR:CZ	3.03	0.47
4:S:1:MET:H2	4:S:93:GLU:CD	2.18	0.47
4:S:85:PHE:CE2	4:S:109:LEU:CD2	2.85	0.47
4:S:112:CYS:SG	4:S:153:VAL:CG2	2.99	0.47
1:A:67:LYS:O	4:S:166:LYS:HA	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:77:LEU:O	1:A:80:TYR:HB2	2.15	0.47
1:A:114:PHE:CE2	1:A:153:ILE:HA	2.50	0.47
1:A:220:SER:HB2	4:S:142:ILE:CA	2.43	0.47
2:B:94:ASP:OD1	2:B:95:THR:N	2.47	0.47
2:B:213:LEU:HD21	2:B:247:TYR:CG	2.50	0.47
2:B:389:ILE:HG21	2:B:427:ASN:HD22	1.80	0.47
2:B:515:PHE:HA	2:B:518:ILE:HG22	1.97	0.47
3:M:72:LEU:HD11	3:M:101:LEU:HD22	1.97	0.47
3:M:101:LEU:HD11	3:M:107:ASP:N	2.30	0.47
1:A:114:PHE:CE1	1:A:153:ILE:HG23	2.50	0.47
1:A:147:LEU:CD1	1:A:166:LEU:HD23	2.44	0.47
1:A:189:PHE:HD2	1:A:225:LEU:CG	2.27	0.47
1:A:530:ASN:OD1	1:A:577:VAL:CG2	2.63	0.47
2:B:139:LEU:O	2:B:176:ALA:HB2	2.14	0.47
2:B:143:SER:O	2:B:144:ASP:C	2.46	0.47
2:B:143:SER:CA	2:B:179:LYS:CB	2.90	0.47
2:B:175:LEU:CD1	2:B:210:CYS:CB	2.89	0.47
2:B:219:TYR:CB	2:B:223:LEU:HD21	2.20	0.47
2:B:245:GLN:CB	2:B:309:LEU:CD1	2.93	0.47
2:B:267:ASP:CB	2:B:289:PRO:HB3	2.45	0.47
2:B:399:LEU:O	2:B:401:THR:N	2.48	0.47
3:M:224:VAL:CB	3:M:479:PHE:CD1	2.98	0.47
1:A:95:MET:C	1:A:127:LEU:HD23	2.33	0.47
1:A:408:ILE:HD12	1:A:410:TYR:HD1	1.77	0.47
1:A:476:GLN:OE1	1:A:476:GLN:HA	2.15	0.47
1:A:594:PHE:CB	2:B:473:ASN:HB3	2.22	0.47
2:B:38:TYR:HB3	2:B:42:ILE:HD12	1.45	0.47
2:B:174:ALA:HB1	2:B:211:ALA:HB1	1.94	0.47
2:B:215:TYR:CE2	2:B:229:HIS:CG	3.02	0.47
2:B:299:LEU:HD21	2:B:328:LEU:HD13	1.98	0.47
2:B:553:ALA:O	2:B:556:LEU:HB2	2.15	0.47
2:B:559:ASP:CG	2:B:582:ASP:OD2	2.53	0.47
3:M:5:PHE:HB2	3:M:125:PHE:HE2	1.77	0.47
3:M:215:TYR:CE1	3:M:468:LYS:CA	2.90	0.47
3:M:270:PRO:CA	3:M:302:TYR:HD1	2.28	0.47
3:M:336:ASP:CG	3:M:415:ILE:O	2.44	0.47
3:M:421:GLY:O	3:M:423:ASN:N	2.48	0.47
3:M:473:LYS:O	3:M:474:THR:HG22	2.15	0.47
1:A:100:LEU:CD1	4:S:161:GLU:H	2.28	0.46
1:A:141:VAL:HB	4:S:159:ALA:CB	2.38	0.46
1:A:219:VAL:HG21	1:A:256:LEU:HD23	1.86	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:241:TYR:C	1:A:241:TYR:CD2	2.88	0.46
1:A:335:CYS:O	1:A:338:PHE:HB2	2.15	0.46
1:A:491:THR:O	1:A:495:ILE:HG12	2.15	0.46
1:A:557:LYS:CG	2:B:606:ASP:CB	2.93	0.46
1:A:636:TYR:HD2	2:B:513:TRP:HZ2	1.63	0.46
2:B:4:SER:O	2:B:8:ILE:CG1	2.63	0.46
2:B:10:SER:O	2:B:11:ALA:C	2.52	0.46
2:B:196:LEU:HD13	2:B:215:TYR:CE2	2.50	0.46
2:B:212:VAL:N	2:B:233:TYR:CZ	2.83	0.46
2:B:336:ASN:C	2:B:338:LYS:N	2.67	0.46
2:B:570:GLY:O	2:B:571:SER:C	2.51	0.46
3:M:9:ASP:C	3:M:9:ASP:OD1	2.54	0.46
3:M:217:ASP:CB	3:M:471:LYS:N	2.54	0.46
3:M:223:HIS:CA	3:M:479:PHE:CG	2.86	0.46
3:M:375:LYS:CG	3:M:418:GLU:OE1	2.63	0.46
4:S:35:VAL:HG12	4:S:68:VAL:HG11	1.96	0.46
4:S:53:THR:HG22	4:S:57:LEU:CB	2.26	0.46
1:A:170:LEU:HB3	1:A:202:LYS:HG3	1.98	0.46
1:A:226:SER:HB3	1:A:263:LEU:HD21	1.97	0.46
1:A:316:LEU:CD1	1:A:341:ILE:HG21	2.42	0.46
1:A:403:LEU:CD2	1:A:422:GLU:OE2	2.59	0.46
1:A:636:TYR:H	2:B:554:LYS:CE	2.27	0.46
2:B:97:VAL:HG12	2:B:101:ILE:CD1	2.45	0.46
2:B:115:LEU:HD13	2:B:115:LEU:HA	1.83	0.46
2:B:216:LYS:HB2	2:B:251:LEU:HB2	1.96	0.46
2:B:261:PRO:HD2	2:B:293:VAL:HG23	1.97	0.46
2:B:592:TYR:CE1	2:B:615:SER:CB	2.98	0.46
3:M:6:TYR:OH	3:M:17:GLN:CD	2.41	0.46
3:M:290:PHE:CE2	3:M:297:PHE:CE1	3.00	0.46
3:M:325:LEU:HG	3:M:338:PHE:HD1	1.80	0.46
3:M:339:GLU:CG	3:M:412:ARG:HD3	2.39	0.46
4:S:17:VAL:HB	4:S:32:LEU:HD11	1.98	0.46
1:A:105:VAL:N	4:S:167:ILE:HD12	1.98	0.46
1:A:435:ILE:CG2	1:A:441:TYR:CE2	2.98	0.46
1:A:515:CYS:O	1:A:519:LEU:HG	2.15	0.46
2:B:96:LYS:O	2:B:99:ARG:HB3	2.16	0.46
2:B:403:ILE:CG2	2:B:411:ILE:CD1	2.92	0.46
2:B:522:GLU:C	2:B:524:LYS:H	2.18	0.46
2:B:546:CYS:N	2:B:607:ILE:HG21	2.31	0.46
2:B:592:TYR:O	2:B:595:VAL:HB	2.16	0.46
2:B:604:GLU:HB3	2:B:607:ILE:HD12	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:253:ASN:O	3:M:254:PRO:C	2.53	0.46
3:M:360:LEU:HD13	3:M:433:VAL:HG23	1.95	0.46
1:A:251:TRP:NE1	4:S:104:THR:HA	2.29	0.46
1:A:287:ALA:HB3	1:A:288:THR:CA	2.45	0.46
1:A:287:ALA:CB	1:A:289:SER:HB3	2.44	0.46
1:A:288:THR:HG1	1:A:291:ILE:CB	2.16	0.46
1:A:469:LEU:O	1:A:470:GLY:C	2.43	0.46
2:B:7:ARG:O	2:B:8:ILE:C	2.53	0.46
2:B:102:HIS:HA	2:B:141:ALA:HB2	1.97	0.46
2:B:307:ASN:O	2:B:309:LEU:N	2.49	0.46
2:B:437:SER:HA	2:B:478:LEU:CG	2.45	0.46
2:B:537:PHE:CG	2:B:598:LEU:HD13	2.36	0.46
2:B:559:ASP:O	2:B:562:ASN:HB2	2.15	0.46
3:M:17:GLN:N	3:M:118:TYR:CE1	2.74	0.46
3:M:220:GLU:HB2	3:M:222:PHE:CE1	2.46	0.46
3:M:220:GLU:OE2	3:M:439:TYR:CG	2.69	0.46
3:M:449:VAL:HG12	3:M:452:ILE:HD12	1.91	0.46
1:A:96:SER:H	1:A:127:LEU:CD2	1.79	0.46
1:A:129:LYS:HG2	1:A:165:ASP:OD2	2.16	0.46
1:A:537:THR:CB	1:A:584:PHE:CZ	2.97	0.46
1:A:607:LEU:C	1:A:609:LEU:H	2.18	0.46
2:B:59:VAL:O	2:B:63:MET:HB2	2.15	0.46
2:B:132:SER:O	2:B:169:VAL:HG22	2.16	0.46
2:B:143:SER:HB2	2:B:179:LYS:CD	2.45	0.46
2:B:169:VAL:HG12	2:B:173:VAL:HG23	1.94	0.46
2:B:223:LEU:HB3	2:B:259:TYR:CB	2.46	0.46
2:B:355:ASN:O	2:B:358:MET:N	2.49	0.46
2:B:513:TRP:H	2:B:551:LEU:HD11	1.61	0.46
2:B:546:CYS:N	2:B:607:ILE:HD13	2.30	0.46
3:M:217:ASP:CB	3:M:470:ALA:CA	2.89	0.46
3:M:224:VAL:HG11	3:M:235:LEU:HD12	1.98	0.46
3:M:240:ILE:CG2	3:M:444:ALA:HA	2.45	0.46
3:M:338:PHE:CD2	3:M:415:ILE:HD11	2.50	0.46
1:A:103:LYS:O	1:A:107:TYR:CG	2.61	0.46
1:A:323:CYS:HB2	1:A:355:LEU:HD21	1.96	0.46
1:A:374:LEU:C	1:A:376:GLU:OE1	2.54	0.46
1:A:409:VAL:O	1:A:409:VAL:CG1	2.63	0.46
1:A:604:LEU:HD23	1:A:604:LEU:C	2.35	0.46
2:B:20:ARG:CD	2:B:35:TYR:OH	2.31	0.46
2:B:59:VAL:O	2:B:59:VAL:HG12	2.16	0.46
2:B:136:CYS:HB3	2:B:172:GLU:HG3	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:169:VAL:C	2:B:173:VAL:HG23	2.35	0.46
2:B:199:LEU:O	2:B:201:ALA:CA	2.59	0.46
2:B:219:TYR:HE2	2:B:226:LEU:H	1.48	0.46
2:B:419:VAL:O	2:B:420:ALA:C	2.53	0.46
2:B:461:HIS:CB	2:B:463:LEU:CD2	2.94	0.46
2:B:477:MET:O	2:B:480:GLN:CB	2.58	0.46
2:B:497:LEU:HD13	2:B:503:LEU:HD12	1.97	0.46
2:B:564:LYS:O	2:B:574:ASN:ND2	2.47	0.46
2:B:569:THR:HA	2:B:571:SER:H	1.80	0.46
3:M:130:GLU:OE1	3:M:130:GLU:HA	2.16	0.46
3:M:217:ASP:C	3:M:472:TYR:CE1	2.89	0.46
1:A:95:MET:CB	1:A:127:LEU:HD23	2.46	0.46
1:A:128:LEU:CD1	1:A:150:LEU:HG	2.29	0.46
1:A:219:VAL:CA	1:A:259:LEU:CD1	2.93	0.46
1:A:275:LEU:C	1:A:275:LEU:HD23	2.35	0.46
1:A:287:ALA:N	1:A:288:THR:HA	2.31	0.46
1:A:503:ASN:OD1	3:M:59:ASP:CG	2.52	0.46
2:B:143:SER:CB	2:B:175:LEU:HB3	2.45	0.46
2:B:177:ILE:HB	2:B:196:LEU:CD2	2.26	0.46
2:B:490:ILE:CD1	2:B:518:ILE:HG21	2.45	0.46
2:B:572:GLU:C	2:B:574:ASN:N	2.69	0.46
3:M:235:LEU:HD11	3:M:306:LEU:HB2	1.94	0.46
3:M:290:PHE:HD1	3:M:299:LEU:CD1	2.28	0.46
1:A:166:LEU:HD12	1:A:185:LEU:HD21	1.95	0.46
1:A:166:LEU:HB2	1:A:185:LEU:HD21	1.98	0.46
1:A:320:HIS:N	1:A:338:PHE:HZ	2.14	0.46
1:A:384:LEU:HD21	1:A:441:TYR:CD2	2.49	0.46
2:B:56:SER:O	2:B:97:VAL:HG21	2.15	0.46
2:B:106:LEU:CG	2:B:144:ASP:HB2	2.43	0.46
2:B:120:ILE:HG13	2:B:150:LEU:CD2	2.44	0.46
2:B:208:ILE:C	2:B:210:CYS:N	2.66	0.46
2:B:215:TYR:HB3	2:B:226:LEU:HD13	1.87	0.46
2:B:257:LYS:HA	2:B:260:LEU:HG	1.97	0.46
2:B:389:ILE:HG22	2:B:393:ILE:HD11	1.98	0.46
2:B:487:LEU:CD2	2:B:522:GLU:HB3	2.29	0.46
2:B:556:LEU:HD22	2:B:588:ILE:CG1	2.07	0.46
3:M:2:TYR:O	3:M:80:THR:C	2.54	0.46
3:M:246:VAL:CA	3:M:470:ALA:HB1	2.45	0.46
3:M:258:VAL:CG1	3:M:452:ILE:HG13	2.45	0.46
3:M:435:LEU:HD12	3:M:435:LEU:N	2.31	0.46
4:S:135:ILE:O	4:S:141:VAL:CB	2.62	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:175:PRO:CB	1:A:211:ASP:CG	2.82	0.46
1:A:252:ILE:HG13	4:S:145:ASN:HA	1.67	0.46
1:A:562:TRP:CE3	1:A:574:ILE:HD12	2.51	0.46
1:A:633:PHE:CB	2:B:554:LYS:HB2	2.45	0.46
2:B:36:THR:O	2:B:40:GLN:CG	2.64	0.46
2:B:77:ILE:HG23	2:B:82:TYR:HE1	1.71	0.46
2:B:267:ASP:CB	2:B:289:PRO:HG3	2.46	0.46
3:M:339:GLU:CD	3:M:412:ARG:HD3	2.35	0.46
3:M:356:LEU:HD23	3:M:358:ILE:HG13	1.93	0.46
4:S:53:THR:HG1	4:S:68:VAL:C	1.98	0.46
4:S:127:THR:CG2	4:S:157:ASN:OD1	2.64	0.46
4:S:160:ALA:HB1	4:S:167:ILE:HG21	1.96	0.46
1:A:67:LYS:CG	1:A:94:VAL:HG22	2.45	0.46
1:A:132:LEU:HD12	1:A:165:ASP:CB	2.45	0.46
2:B:14:THR:O	2:B:16:LYS:N	2.48	0.46
2:B:87:VAL:HG21	2:B:119:SER:HA	1.98	0.46
2:B:157:THR:C	2:B:159:LYS:H	2.18	0.46
2:B:175:LEU:HD21	2:B:210:CYS:O	2.02	0.46
2:B:215:TYR:O	2:B:219:TYR:CD1	2.67	0.46
2:B:252:LEU:CG	2:B:302:PHE:CE1	2.94	0.46
2:B:433:VAL:HG12	2:B:474:VAL:CG1	2.43	0.46
2:B:493:LEU:HD21	2:B:511:ILE:CG1	2.46	0.46
3:M:18:TYR:CE2	3:M:20:LEU:HD23	2.43	0.46
4:S:34:GLN:CG	4:S:58:LEU:HD11	2.45	0.46
4:S:51:LEU:H	4:S:77:TYR:HE1	1.64	0.46
1:A:63:ASP:C	4:S:165:SER:HG	2.11	0.45
1:A:121:LEU:HD13	1:A:155:THR:HG21	1.97	0.45
1:A:128:LEU:CB	1:A:150:LEU:HD21	2.46	0.45
1:A:132:LEU:CD2	1:A:169:MET:SD	3.04	0.45
1:A:200:PHE:CE1	1:A:236:LEU:HG	2.51	0.45
1:A:353:ASP:OD1	1:A:378:ILE:HD12	2.16	0.45
1:A:401:VAL:HG23	1:A:419:ILE:HB	1.98	0.45
1:A:401:VAL:HG12	1:A:402:ILE:N	2.30	0.45
1:A:404:GLN:OE1	2:B:7:ARG:NH1	2.50	0.45
2:B:143:SER:CA	2:B:179:LYS:CD	2.80	0.45
2:B:178:ILE:HG23	2:B:218:CYS:N	2.30	0.45
2:B:523:PHE:CD1	2:B:559:ASP:OD2	2.69	0.45
3:M:67:SER:HB3	3:M:90:PHE:HD1	1.70	0.45
3:M:216:VAL:O	3:M:216:VAL:CG2	2.64	0.45
3:M:228:LYS:NZ	3:M:327:PHE:N	2.63	0.45
3:M:319:SER:HB2	3:M:346:ASN:N	2.26	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:469:GLY:O	3:M:470:ALA:C	2.53	0.45
4:S:50:PHE:CA	4:S:76:ILE:HA	2.46	0.45
1:A:74:LEU:HD22	1:A:87:CYS:HB3	1.96	0.45
1:A:95:MET:SD	1:A:107:TYR:CE2	3.09	0.45
1:A:105:VAL:CG2	4:S:167:ILE:CG1	2.71	0.45
1:A:171:ASN:CG	1:A:202:LYS:NZ	2.69	0.45
1:A:264:SER:HB2	1:A:271:ARG:HG3	1.92	0.45
1:A:533:ILE:CG1	1:A:562:TRP:HH2	2.12	0.45
1:A:633:PHE:CZ	2:B:513:TRP:HE3	2.13	0.45
2:B:29:LYS:HE2	2:B:30:LEU:CB	2.46	0.45
2:B:106:LEU:HD22	2:B:144:ASP:CB	2.45	0.45
2:B:121:ASN:N	2:B:153:ILE:HD13	2.31	0.45
2:B:215:TYR:CE2	2:B:229:HIS:HB2	2.50	0.45
2:B:381:PHE:O	2:B:383:VAL:N	2.49	0.45
2:B:468:LEU:HD13	2:B:507:ALA:HB2	1.96	0.45
2:B:478:LEU:C	2:B:480:GLN:N	2.63	0.45
2:B:512:VAL:CG2	2:B:548:ILE:HG12	2.39	0.45
2:B:538:SER:N	2:B:598:LEU:HD22	2.32	0.45
3:M:323:MET:CG	3:M:342:LEU:HG	2.46	0.45
1:A:224:GLU:O	1:A:227:LYS:N	2.46	0.45
1:A:320:HIS:CB	1:A:352:PHE:CE2	2.84	0.45
1:A:609:LEU:CG	1:A:628:VAL:CG2	2.92	0.45
1:A:631:SER:HA	2:B:554:LYS:HA	1.98	0.45
2:B:37:TYR:CZ	2:B:46:GLN:CD	2.63	0.45
2:B:127:LEU:HD12	2:B:157:THR:CB	2.43	0.45
2:B:133:GLU:O	2:B:168:MET:SD	2.74	0.45
2:B:193:LEU:HB3	2:B:225:LEU:HD11	0.97	0.45
2:B:381:PHE:HD2	2:B:395:LYS:NZ	2.13	0.45
2:B:418:TYR:CG	2:B:419:VAL:N	2.83	0.45
3:M:1:MET:C	3:M:81:SER:CB	2.84	0.45
3:M:111:ILE:HA	3:M:114:ILE:HD12	1.98	0.45
3:M:222:PHE:HB3	3:M:479:PHE:CE1	2.52	0.45
3:M:347:PHE:CZ	3:M:439:TYR:HD2	2.33	0.45
4:S:135:ILE:HG23	4:S:141:VAL:HG11	1.93	0.45
4:S:151:ALA:O	4:S:154:ASP:HB2	2.16	0.45
1:A:102:GLN:HA	4:S:167:ILE:N	2.32	0.45
1:A:102:GLN:C	4:S:163:THR:HB	2.13	0.45
1:A:287:ALA:H	1:A:288:THR:HA	1.81	0.45
1:A:586:GLU:CA	1:A:604:LEU:CD1	2.94	0.45
2:B:274:PRO:CD	2:B:295:ASN:CG	2.78	0.45
2:B:278:PRO:CB	2:B:289:PRO:O	2.64	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:325:LEU:HD22	2:B:339:PHE:CE2	2.52	0.45
2:B:588:ILE:O	2:B:591:MET:HB2	2.15	0.45
3:M:128:CYS:O	3:M:131:ALA:HB3	2.16	0.45
3:M:378:ILE:HB	3:M:413:GLY:CA	2.44	0.45
4:S:27:LYS:O	4:S:31:LEU:HG	2.17	0.45
4:S:135:ILE:O	4:S:140:MET:O	2.34	0.45
1:A:88:ASN:OD1	1:A:120:ILE:HG21	2.16	0.45
1:A:129:LYS:CG	1:A:161:ASP:HB3	2.47	0.45
1:A:215:VAL:O	1:A:219:VAL:CG2	2.61	0.45
1:A:225:LEU:CG	1:A:233:PHE:CZ	2.99	0.45
1:A:482:ILE:CG1	1:A:517:TRP:CZ3	2.97	0.45
1:A:555:LEU:HD12	1:A:585:PHE:CZ	2.52	0.45
2:B:63:MET:HG2	2:B:100:LEU:CB	2.46	0.45
2:B:334:MET:HE1	2:B:339:PHE:CE1	2.50	0.45
3:M:215:TYR:CB	3:M:468:LYS:CA	2.92	0.45
3:M:224:VAL:CG1	3:M:235:LEU:HG	2.46	0.45
3:M:290:PHE:CD1	3:M:297:PHE:CD1	2.90	0.45
3:M:319:SER:O	3:M:320:ILE:HD13	2.16	0.45
4:S:53:THR:HG22	4:S:54:PRO:O	2.17	0.45
4:S:70:ASN:HB3	4:S:73:ILE:HB	1.96	0.45
4:S:129:GLU:O	4:S:132:LEU:HB3	2.17	0.45
1:A:71:VAL:HG11	1:A:105:VAL:C	2.37	0.45
1:A:141:VAL:HB	4:S:159:ALA:HB2	1.97	0.45
1:A:150:LEU:HD22	1:A:162:ILE:CD1	2.46	0.45
1:A:179:LYS:HD2	4:S:143:GLU:HB2	1.89	0.45
1:A:232:PRO:O	1:A:235:GLN:HB2	2.16	0.45
1:A:433:ILE:HD11	1:A:473:ILE:CG1	2.47	0.45
1:A:617:ASP:OD1	1:A:618:THR:N	2.45	0.45
2:B:47:LEU:HD21	2:B:66:ILE:N	2.31	0.45
2:B:97:VAL:HG12	2:B:101:ILE:HD12	1.98	0.45
2:B:196:LEU:O	2:B:197:LYS:O	2.35	0.45
2:B:241:ASP:HB2	3:M:274:ASP:OD1	2.16	0.45
2:B:279:LEU:CD1	2:B:288:TYR:CE1	2.99	0.45
2:B:394:TRP:CE3	2:B:397:GLN:CD	2.89	0.45
2:B:416:LYS:HE3	2:B:453:TRP:CZ3	2.52	0.45
2:B:463:LEU:HD13	2:B:467:VAL:HG11	1.99	0.45
2:B:511:ILE:O	2:B:515:PHE:CD1	2.66	0.45
2:B:577:ASN:O	2:B:578:PRO:C	2.39	0.45
3:M:8:THR:H	3:M:75:TRP:C	2.17	0.45
3:M:222:PHE:CE2	3:M:240:ILE:HD13	2.52	0.45
3:M:242:GLY:O	3:M:302:TYR:N	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:246:VAL:CA	3:M:470:ALA:CB	2.80	0.45
4:S:14:PRO:CB	4:S:36:TYR:HE1	2.25	0.45
4:S:47:GLN:HG2	4:S:84:TYR:HE2	1.81	0.45
1:A:135:ASP:O	1:A:139:ASP:HB3	2.16	0.45
1:A:170:LEU:HD13	1:A:170:LEU:HA	1.82	0.45
1:A:395:PHE:CD1	1:A:428:MET:HG3	2.51	0.45
1:A:625:LEU:C	2:B:617:LEU:HD11	2.33	0.45
1:A:631:SER:HB2	2:B:557:SER:HB3	1.88	0.45
2:B:79:VAL:HG21	2:B:108:PHE:CE1	2.39	0.45
2:B:117:LEU:CD2	2:B:149:SER:CB	2.85	0.45
2:B:136:CYS:SG	2:B:168:MET:CB	3.05	0.45
2:B:155:LEU:HD21	2:B:191:GLU:HB2	1.97	0.45
2:B:279:LEU:O	2:B:280:PRO:C	2.52	0.45
2:B:467:VAL:HG12	2:B:471:TYR:CD1	2.51	0.45
2:B:498:THR:HG22	2:B:532:ARG:HB3	1.98	0.45
3:M:222:PHE:CD2	3:M:439:TYR:OH	2.70	0.45
1:A:464:ILE:O	1:A:464:ILE:HG12	2.17	0.45
1:A:481:MET:O	1:A:482:ILE:C	2.53	0.45
1:A:600:SER:O	1:A:602:GLU:N	2.50	0.45
2:B:74:ASP:O	2:B:77:ILE:HB	2.17	0.45
2:B:106:LEU:CD2	2:B:144:ASP:HB3	2.42	0.45
2:B:196:LEU:CA	2:B:229:HIS:CE1	2.99	0.45
2:B:257:LYS:HA	2:B:260:LEU:HD11	1.99	0.45
2:B:259:TYR:O	2:B:260:LEU:HB2	2.16	0.45
2:B:266:VAL:HG13	2:B:291:TYR:N	2.24	0.45
2:B:292:GLU:HG3	2:B:295:ASN:O	2.17	0.45
2:B:387:ASP:CB	2:B:388:PRO:HD2	2.45	0.45
2:B:390:VAL:CG1	2:B:394:TRP:NE1	2.80	0.45
2:B:461:HIS:C	2:B:463:LEU:N	2.63	0.45
2:B:544:THR:HG22	2:B:548:ILE:HD11	1.99	0.45
3:M:245:ASP:OD1	3:M:297:PHE:O	2.35	0.45
3:M:331:LEU:HD12	3:M:331:LEU:C	2.36	0.45
3:M:376:ILE:HG22	3:M:379:LEU:HD11	1.97	0.45
4:S:118:GLU:O	4:S:122:ILE:HG13	2.16	0.45
1:A:63:ASP:OD1	1:A:64:LEU:N	2.50	0.45
1:A:282:MET:HE3	1:A:294:SER:CB	2.46	0.45
2:B:139:LEU:CD2	2:B:173:VAL:CG1	2.95	0.45
2:B:425:PRO:HB2	2:B:428:VAL:HG23	1.97	0.45
2:B:513:TRP:HE1	2:B:517:GLU:CG	2.30	0.45
2:B:534:ILE:HD12	2:B:591:MET:HA	1.71	0.45
3:M:215:TYR:HB2	3:M:467:TYR:CB	2.47	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:344:ILE:O	3:M:406:GLY:HA2	2.17	0.45
4:S:55:PRO:HB3	4:S:71:GLU:CA	2.46	0.45
4:S:85:PHE:HE2	4:S:109:LEU:HD22	1.81	0.45
4:S:93:GLU:HB3	4:S:98:ILE:HD11	1.98	0.45
1:A:67:LYS:CA	4:S:166:LYS:HA	2.45	0.45
1:A:145:ILE:HD11	4:S:156:LEU:HD22	1.93	0.45
1:A:211:ASP:OD2	4:S:148:ARG:CZ	2.57	0.45
1:A:298:ILE:HG21	1:A:312:ALA:HB2	1.99	0.45
1:A:300:LYS:O	1:A:301:GLY:C	2.55	0.45
1:A:566:PHE:CE1	1:A:570:LYS:HB3	2.52	0.45
2:B:132:SER:CB	2:B:169:VAL:HG23	2.45	0.45
2:B:141:ALA:HB1	2:B:145:MET:HE1	1.98	0.45
2:B:281:ASP:OD1	2:B:287:GLU:CD	2.55	0.45
2:B:306:LEU:HD22	2:B:321:CYS:CA	2.45	0.45
2:B:314:ASN:O	2:B:317:VAL:HB	2.17	0.45
3:M:52:ASP:CA	3:M:67:SER:HA	2.46	0.45
3:M:347:PHE:CZ	3:M:352:GLN:O	2.70	0.45
3:M:430:LEU:HD23	3:M:483:LEU:HD13	1.98	0.45
1:A:150:LEU:HD22	1:A:162:ILE:HD11	1.99	0.44
1:A:249:ASN:OD1	1:A:251:TRP:N	2.45	0.44
1:A:292:TYR:O	1:A:292:TYR:HD1	1.96	0.44
1:A:388:VAL:HG22	1:A:435:ILE:CD1	2.47	0.44
1:A:396:VAL:O	1:A:396:VAL:CG1	2.65	0.44
1:A:453:VAL:O	1:A:457:LEU:HG	2.16	0.44
1:A:464:ILE:O	1:A:464:ILE:HG23	2.17	0.44
1:A:594:PHE:O	1:A:598:GLU:HG2	2.17	0.44
2:B:208:ILE:CB	2:B:236:ILE:HG21	2.46	0.44
2:B:279:LEU:CD1	2:B:288:TYR:CD1	3.00	0.44
2:B:419:VAL:HG11	2:B:457:HIS:NE2	2.31	0.44
3:M:218:LEU:HD12	3:M:472:TYR:OH	2.16	0.44
3:M:220:GLU:HG3	3:M:439:TYR:CG	2.51	0.44
3:M:224:VAL:CB	3:M:226:PHE:CE1	2.99	0.44
3:M:253:ASN:OD1	3:M:292:PRO:HD2	2.16	0.44
4:S:15:ARG:NE	4:S:122:ILE:HD11	2.29	0.44
1:A:100:LEU:HB3	1:A:142:LYS:CG	2.47	0.44
1:A:103:LYS:O	1:A:104:ARG:O	2.35	0.44
1:A:141:VAL:HG13	4:S:156:LEU:HA	0.85	0.44
1:A:213:SER:O	4:S:143:GLU:CB	2.63	0.44
1:A:240:LEU:C	1:A:242:GLU:O	2.56	0.44
1:A:281:LEU:O	1:A:282:MET:C	2.36	0.44
1:A:326:GLN:HG2	1:A:331:ARG:NH2	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:609:LEU:CG	1:A:628:VAL:HG11	2.48	0.44
1:A:638:LEU:HD13	2:B:519:ALA:HB2	1.37	0.44
2:B:162:VAL:HG13	2:B:198:GLU:HB3	1.98	0.44
2:B:201:ALA:HA	2:B:232:ARG:HG2	1.99	0.44
2:B:374:PHE:CZ	2:B:381:PHE:HD1	2.26	0.44
2:B:436:LEU:HB3	2:B:450:VAL:CG1	2.47	0.44
2:B:526:CYS:HB2	2:B:555:LEU:HD11	1.98	0.44
2:B:592:TYR:CE1	2:B:615:SER:HB3	2.53	0.44
2:B:592:TYR:HE1	2:B:615:SER:HB3	1.82	0.44
3:M:18:TYR:HD1	3:M:122:SER:CB	2.16	0.44
3:M:56:VAL:O	3:M:57:GLY:O	2.34	0.44
3:M:220:GLU:OE1	3:M:222:PHE:CZ	2.71	0.44
3:M:347:PHE:CE1	3:M:350:VAL:HG11	2.52	0.44
3:M:380:ARG:HG2	3:M:410:VAL:HB	1.40	0.44
4:S:17:VAL:HG11	4:S:19:PHE:CE1	2.51	0.44
4:S:51:LEU:HD12	4:S:77:TYR:CZ	2.52	0.44
4:S:107:GLU:HG3	4:S:146:VAL:HG11	2.00	0.44
1:A:114:PHE:HZ	1:A:154:ILE:HG12	1.82	0.44
1:A:481:MET:O	1:A:522:PHE:HZ	2.00	0.44
1:A:581:LEU:HG	1:A:585:PHE:CE2	2.52	0.44
1:A:636:TYR:CD2	2:B:513:TRP:HZ2	2.35	0.44
2:B:17:VAL:O	2:B:18:ILE:C	2.54	0.44
2:B:42:ILE:HG12	2:B:65:ARG:CD	2.35	0.44
2:B:135:ARG:O	2:B:161:LEU:CD2	2.66	0.44
2:B:344:VAL:CG2	2:B:381:PHE:CZ	2.98	0.44
2:B:418:TYR:O	2:B:418:TYR:HD1	1.99	0.44
2:B:556:LEU:CD2	2:B:588:ILE:HD11	2.46	0.44
2:B:559:ASP:CA	2:B:562:ASN:HB2	2.36	0.44
3:M:212:ASN:CB	3:M:250:LEU:HA	2.40	0.44
3:M:217:ASP:N	3:M:472:TYR:CE1	2.86	0.44
4:S:102:ILE:O	4:S:105:PHE:HB3	2.17	0.44
1:A:171:ASN:CG	1:A:202:LYS:HZ1	2.18	0.44
2:B:81:LEU:O	2:B:85:ASP:N	2.37	0.44
2:B:109:ALA:O	2:B:110:GLU:O	2.35	0.44
2:B:120:ILE:O	2:B:153:ILE:HG21	2.18	0.44
2:B:197:LYS:HG2	2:B:198:GLU:N	2.32	0.44
2:B:430:ILE:HG23	2:B:470:ALA:HB1	1.97	0.44
2:B:502:SER:O	2:B:503:LEU:O	2.35	0.44
2:B:520:SER:HA	2:B:523:PHE:CD2	2.53	0.44
3:M:74:TYR:HD1	3:M:76:CYS:SG	2.40	0.44
3:M:222:PHE:CE1	3:M:439:TYR:CD1	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:228:LYS:HZ1	3:M:326:HIS:HA	1.81	0.44
3:M:269:ILE:N	3:M:302:TYR:CE1	2.86	0.44
3:M:340:LEU:HG	3:M:342:LEU:CD1	2.48	0.44
3:M:424:PHE:HZ	3:M:428:VAL:HG23	1.82	0.44
4:S:51:LEU:N	4:S:75:ILE:O	2.50	0.44
4:S:80:TYR:CB	4:S:106:VAL:HG11	2.47	0.44
1:A:114:PHE:CD1	1:A:114:PHE:C	2.91	0.44
1:A:219:VAL:O	4:S:140:MET:HE1	2.18	0.44
1:A:365:VAL:O	1:A:366:SER:C	2.46	0.44
2:B:63:MET:N	2:B:66:ILE:HD12	2.32	0.44
2:B:116:THR:O	2:B:150:LEU:HD22	2.17	0.44
2:B:195:ILE:HG22	2:B:196:LEU:N	2.31	0.44
2:B:382:TYR:HE1	2:B:399:LEU:HD21	1.83	0.44
2:B:556:LEU:O	2:B:563:PHE:HD1	2.01	0.44
2:B:589:SER:C	2:B:591:MET:H	2.20	0.44
3:M:258:VAL:CB	3:M:452:ILE:HG13	2.48	0.44
1:A:237:SER:HB2	1:A:270:LEU:HD22	2.00	0.44
1:A:392:MET:HE3	1:A:428:MET:CE	2.46	0.44
1:A:396:VAL:C	1:A:397:ASP:OD1	2.56	0.44
1:A:438:ALA:O	1:A:441:TYR:HD1	1.97	0.44
1:A:603:VAL:O	1:A:606:PHE:CB	2.62	0.44
2:B:42:ILE:HG22	2:B:43:ASN:O	2.17	0.44
2:B:47:LEU:O	2:B:50:LEU:HB3	2.18	0.44
2:B:234:CYS:HB2	2:B:301:LEU:HB3	1.99	0.44
2:B:281:ASP:C	2:B:283:TYR:H	2.20	0.44
2:B:493:LEU:CD2	2:B:511:ILE:HA	2.48	0.44
3:M:223:HIS:CD2	3:M:478:ASN:HB3	2.50	0.44
3:M:437:TYR:CG	3:M:439:TYR:OH	2.70	0.44
4:S:34:GLN:HB3	4:S:58:LEU:CD1	2.48	0.44
1:A:132:LEU:HD13	1:A:165:ASP:HB2	1.95	0.44
1:A:179:LYS:NZ	4:S:149:ILE:CG1	2.77	0.44
2:B:140:SER:OG	2:B:175:LEU:CD1	2.66	0.44
2:B:212:VAL:CG2	2:B:248:LEU:HD23	2.42	0.44
2:B:212:VAL:HG22	2:B:248:LEU:HD23	1.93	0.44
2:B:315:PRO:O	2:B:318:ILE:HB	2.17	0.44
2:B:483:PRO:O	2:B:486:HIS:HB3	2.17	0.44
2:B:556:LEU:CB	2:B:588:ILE:CD1	2.85	0.44
3:M:258:VAL:CG2	3:M:452:ILE:HG23	2.47	0.44
3:M:290:PHE:CD2	3:M:297:PHE:CZ	3.06	0.44
3:M:372:ILE:CD1	3:M:428:VAL:HG13	2.48	0.44
3:M:433:VAL:CG1	3:M:433:VAL:O	2.66	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:3:HIS:O	4:S:19:PHE:CA	2.62	0.44
4:S:80:TYR:CB	4:S:106:VAL:CG1	2.96	0.44
4:S:148:ARG:O	4:S:151:ALA:N	2.51	0.44
1:A:200:PHE:O	1:A:203:PHE:HB2	2.18	0.44
1:A:392:MET:CE	1:A:428:MET:CE	2.96	0.44
2:B:158:VAL:HG21	2:B:177:ILE:CG1	2.47	0.44
2:B:230:PHE:CD2	2:B:298:ASP:CA	3.01	0.44
2:B:305:SER:O	2:B:309:LEU:HD23	2.17	0.44
2:B:498:THR:CG2	2:B:532:ARG:HB2	2.47	0.44
2:B:537:PHE:O	2:B:539:ASN:N	2.50	0.44
3:M:219:LEU:H	3:M:472:TYR:CB	2.31	0.44
3:M:220:GLU:OE2	3:M:439:TYR:HB2	2.18	0.44
3:M:220:GLU:OE1	3:M:222:PHE:CE1	2.70	0.44
3:M:253:ASN:OD1	3:M:292:PRO:CD	2.65	0.44
3:M:347:PHE:CE2	3:M:352:GLN:O	2.70	0.44
4:S:39:ILE:CG1	4:S:77:TYR:CD2	3.01	0.44
4:S:87:PHE:CD1	4:S:102:ILE:CG1	2.89	0.44
1:A:128:LEU:HD22	1:A:146:ALA:HB1	1.99	0.44
1:A:139:ASP:CG	1:A:177:ILE:CD1	2.86	0.44
1:A:213:SER:CA	4:S:143:GLU:OE1	2.66	0.44
1:A:275:LEU:CD1	1:A:308:ASP:CB	2.95	0.44
1:A:323:CYS:HB3	1:A:355:LEU:HD21	1.97	0.44
1:A:399:ASP:OD1	1:A:420:ILE:CG2	2.66	0.44
1:A:581:LEU:HG	1:A:607:LEU:CD1	2.48	0.44
1:A:594:PHE:CD2	2:B:474:VAL:HG23	2.47	0.44
2:B:162:VAL:HG11	2:B:195:ILE:O	2.18	0.44
2:B:390:VAL:HG12	2:B:394:TRP:NE1	2.32	0.44
2:B:440:GLY:C	2:B:442:LEU:H	2.22	0.44
2:B:490:ILE:HG23	2:B:515:PHE:CZ	2.51	0.44
3:M:3:LEU:O	3:M:20:LEU:HD12	2.18	0.44
3:M:7:ILE:HG21	3:M:114:ILE:HD13	2.00	0.44
3:M:219:LEU:CG	3:M:440:ILE:HG12	2.45	0.44
3:M:223:HIS:CA	3:M:479:PHE:CZ	2.98	0.44
3:M:319:SER:HB3	3:M:346:ASN:CG	2.37	0.44
3:M:323:MET:HE3	3:M:342:LEU:CD2	2.35	0.44
4:S:48:SER:CB	4:S:78:LYS:N	2.76	0.44
4:S:50:PHE:CB	4:S:76:ILE:CD1	2.93	0.44
4:S:75:ILE:HB	4:S:77:TYR:CE1	2.52	0.44
1:A:114:PHE:CZ	1:A:154:ILE:HG12	2.53	0.43
1:A:178:ARG:CB	1:A:214:VAL:CG1	2.95	0.43
1:A:317:GLU:HG2	1:A:351:ARG:HH12	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:420:ILE:HG22	1:A:421:PRO:O	2.18	0.43
1:A:429:VAL:HG13	1:A:457:LEU:HD13	1.99	0.43
1:A:509:PRO:HA	1:A:547:VAL:HG21	1.98	0.43
2:B:127:LEU:HD22	2:B:157:THR:CG2	2.48	0.43
2:B:197:LYS:H	2:B:229:HIS:CE1	2.33	0.43
2:B:227:HIS:O	2:B:229:HIS:CA	2.58	0.43
2:B:467:VAL:HG12	2:B:471:TYR:HD1	1.81	0.43
2:B:537:PHE:CZ	2:B:545:ARG:HD3	2.52	0.43
2:B:589:SER:HA	2:B:592:TYR:CD2	2.45	0.43
3:M:380:ARG:HH11	3:M:412:ARG:NH1	2.16	0.43
4:S:6:LEU:HB3	4:S:86:THR:HB	1.99	0.43
4:S:127:THR:HG22	4:S:153:VAL:HG13	2.00	0.43
1:A:150:LEU:HD13	1:A:162:ILE:HG23	2.00	0.43
1:A:179:LYS:HG3	4:S:143:GLU:HG3	1.50	0.43
1:A:249:ASN:HB3	1:A:252:ILE:HD12	2.00	0.43
1:A:486:SER:O	1:A:487:MET:HB2	2.18	0.43
1:A:563:CYS:O	1:A:566:PHE:CD2	2.71	0.43
2:B:171:GLY:H	2:B:207:VAL:HG13	1.66	0.43
2:B:208:ILE:O	2:B:210:CYS:N	2.51	0.43
2:B:340:ILE:CG2	2:B:373:LEU:HG	2.48	0.43
2:B:343:LEU:O	2:B:359:LEU:HD12	2.17	0.43
2:B:398:ILE:C	2:B:400:SER:N	2.69	0.43
2:B:483:PRO:CB	2:B:521:ILE:CG2	2.65	0.43
3:M:100:LEU:HD22	3:M:100:LEU:C	2.36	0.43
3:M:214:LEU:N	3:M:467:TYR:CB	2.81	0.43
3:M:319:SER:O	3:M:320:ILE:HG12	2.18	0.43
3:M:372:ILE:HD12	3:M:428:VAL:CG2	2.48	0.43
3:M:376:ILE:CD1	3:M:415:ILE:HG23	2.48	0.43
3:M:450:GLU:OE1	3:M:450:GLU:HA	2.18	0.43
4:S:15:ARG:HE	4:S:118:GLU:HB2	1.84	0.43
4:S:55:PRO:O	4:S:57:LEU:N	2.51	0.43
1:A:182:ILE:HG23	1:A:221:VAL:CG2	2.39	0.43
1:A:264:SER:HB2	1:A:271:ARG:NE	2.30	0.43
1:A:320:HIS:HA	1:A:338:PHE:HZ	1.82	0.43
1:A:548:GLN:NE2	1:A:588:LEU:HD11	2.32	0.43
1:A:582:ILE:HG23	1:A:604:LEU:HG	2.00	0.43
1:A:621:LEU:O	1:A:621:LEU:HD13	2.18	0.43
1:A:634:ASN:HA	2:B:516:GLY:CA	2.48	0.43
2:B:14:THR:C	2:B:16:LYS:N	2.71	0.43
2:B:237:ILE:HB	2:B:248:LEU:HD13	2.00	0.43
2:B:359:LEU:HD13	2:B:359:LEU:HA	1.88	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:580:TYR:C	2:B:582:ASP:N	2.72	0.43
2:B:589:SER:O	2:B:592:TYR:N	2.51	0.43
2:B:591:MET:O	2:B:595:VAL:HG23	2.18	0.43
3:M:61:GLU:OE2	3:M:82:LYS:HB2	2.19	0.43
4:S:16:LEU:CB	4:S:125:TRP:CD1	3.01	0.43
1:A:295:VAL:HG22	1:A:315:CYS:CB	2.19	0.43
1:A:309:PHE:CZ	1:A:348:PHE:CE1	3.07	0.43
1:A:532:LEU:O	1:A:535:ILE:N	2.51	0.43
1:A:569:ASP:C	1:A:571:ARG:N	2.65	0.43
1:A:576:MET:O	1:A:579:LYS:HB3	2.18	0.43
1:A:637:GLU:HG2	2:B:512:VAL:O	2.19	0.43
3:M:9:ASP:CG	3:M:13:LYS:HB3	2.38	0.43
3:M:67:SER:CB	3:M:90:PHE:CB	2.96	0.43
3:M:326:HIS:O	3:M:338:PHE:CA	2.50	0.43
3:M:473:LYS:C	3:M:474:THR:CG2	2.86	0.43
1:A:128:LEU:HD12	1:A:150:LEU:HD23	1.69	0.43
1:A:200:PHE:HZ	1:A:235:GLN:HB2	1.82	0.43
1:A:224:GLU:N	4:S:140:MET:HE3	2.31	0.43
1:A:253:ILE:CG2	1:A:281:LEU:CB	2.97	0.43
1:A:552:ILE:CD1	1:A:600:SER:CB	2.96	0.43
1:A:563:CYS:CA	1:A:566:PHE:HD2	2.11	0.43
2:B:24:ALA:HB2	2:B:35:TYR:CZ	2.36	0.43
2:B:63:MET:HG2	2:B:101:ILE:N	2.33	0.43
2:B:158:VAL:HG13	2:B:177:ILE:HG13	1.91	0.43
2:B:178:ILE:HG23	2:B:217:GLU:N	2.33	0.43
2:B:455:ILE:HD11	2:B:489:ILE:HG23	1.99	0.43
2:B:537:PHE:CZ	2:B:598:LEU:C	2.91	0.43
2:B:563:PHE:CE2	2:B:582:ASP:HB2	2.52	0.43
3:M:220:GLU:O	3:M:439:TYR:HD1	2.02	0.43
3:M:355:ASP:O	3:M:356:LEU:C	2.55	0.43
3:M:373:ALA:O	3:M:417:TYR:HA	2.18	0.43
4:S:34:GLN:CD	4:S:58:LEU:CG	2.83	0.43
4:S:34:GLN:C	4:S:36:TYR:H	2.21	0.43
4:S:107:GLU:HG3	4:S:146:VAL:CG1	2.48	0.43
4:S:111:ARG:HB3	4:S:150:VAL:CG2	2.47	0.43
4:S:117:ASN:CG	4:S:120:ASP:OD2	2.57	0.43
1:A:402:ILE:O	1:A:403:LEU:C	2.54	0.43
1:A:631:SER:CA	2:B:554:LYS:HA	2.49	0.43
2:B:37:TYR:CE2	2:B:46:GLN:CD	2.87	0.43
2:B:123:LEU:C	2:B:127:LEU:CD1	2.86	0.43
2:B:151:ALA:CB	2:B:188:TYR:CE1	3.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:159:LYS:CG	2:B:191:GLU:OE1	2.66	0.43
2:B:326:TYR:O	2:B:327:GLN:C	2.56	0.43
2:B:334:MET:CE	2:B:334:MET:CA	2.86	0.43
2:B:404:ASN:C	2:B:408:VAL:HG23	2.36	0.43
2:B:433:VAL:HG11	2:B:474:VAL:CG2	2.35	0.43
3:M:24:ALA:C	3:M:25:PRO:O	2.50	0.43
3:M:118:TYR:HA	3:M:121:ILE:HD12	2.01	0.43
3:M:304:VAL:CG1	3:M:445:SER:HA	2.43	0.43
3:M:374:TYR:CE1	3:M:390:ILE:CD1	3.02	0.43
1:A:104:ARG:CG	1:A:145:ILE:CG1	2.66	0.43
1:A:144:GLY:CA	1:A:180:LYS:HG3	2.49	0.43
1:A:429:VAL:CG2	1:A:469:LEU:HD21	2.40	0.43
1:A:626:SER:O	1:A:630:PRO:CD	2.66	0.43
2:B:116:THR:CG2	2:B:150:LEU:HD21	2.41	0.43
2:B:212:VAL:HG23	2:B:233:TYR:CD2	2.53	0.43
2:B:287:GLU:OE2	2:B:287:GLU:HA	2.18	0.43
2:B:367:SER:CB	2:B:401:THR:OG1	2.67	0.43
2:B:410:GLU:HA	2:B:413:LYS:HE2	2.01	0.43
3:M:103:TYR:CE1	3:M:124:ILE:HG12	2.54	0.43
3:M:271:SER:C	3:M:272:LEU:HD12	2.39	0.43
3:M:380:ARG:HG2	3:M:410:VAL:O	2.19	0.43
4:S:4:ALA:HB1	4:S:19:PHE:CD2	2.53	0.43
4:S:14:PRO:CB	4:S:36:TYR:CE1	3.00	0.43
4:S:75:ILE:CG2	4:S:86:THR:HG21	2.44	0.43
4:S:87:PHE:CZ	4:S:102:ILE:HA	2.53	0.43
1:A:125:THR:CA	1:A:158:LEU:HD13	2.49	0.43
1:A:128:LEU:HB3	1:A:150:LEU:HD11	1.99	0.43
1:A:219:VAL:CG2	1:A:240:LEU:HD22	2.28	0.43
1:A:432:ILE:CG2	1:A:457:LEU:HD11	2.39	0.43
1:A:524:THR:CG2	1:A:565:ASN:ND2	2.81	0.43
2:B:13:ASP:O	2:B:14:THR:O	2.36	0.43
2:B:108:PHE:HE2	2:B:115:LEU:HD23	1.82	0.43
2:B:230:PHE:CG	2:B:298:ASP:HB3	2.42	0.43
2:B:279:LEU:HB3	2:B:280:PRO:HD2	2.01	0.43
2:B:329:ALA:HA	2:B:333:GLN:NE2	2.34	0.43
2:B:367:SER:O	2:B:368:ILE:C	2.55	0.43
2:B:537:PHE:CD2	2:B:537:PHE:O	2.71	0.43
2:B:563:PHE:C	2:B:564:LYS:O	2.56	0.43
2:B:567:GLN:OE1	2:B:618:PHE:CE1	2.71	0.43
3:M:94:GLU:O	3:M:97:ASP:CB	2.64	0.43
3:M:306:LEU:HD13	3:M:317:MET:HE3	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:377:LYS:HE3	3:M:416:GLU:HB2	1.97	0.43
4:S:56:SER:C	4:S:60:SER:HB2	2.34	0.43
1:A:107:TYR:HD1	1:A:145:ILE:HG22	1.84	0.43
1:A:478:ARG:HG3	1:A:517:TRP:CE3	2.54	0.43
1:A:589:SER:HB3	1:A:601:VAL:CG2	2.48	0.43
2:B:139:LEU:HD21	2:B:176:ALA:HB3	1.97	0.43
2:B:237:ILE:HD13	2:B:309:LEU:HD21	2.01	0.43
2:B:546:CYS:SG	2:B:607:ILE:CA	3.06	0.43
2:B:556:LEU:HA	2:B:588:ILE:HD11	0.49	0.43
2:B:577:ASN:OD1	2:B:577:ASN:N	2.52	0.43
2:B:588:ILE:HG23	2:B:589:SER:N	2.34	0.43
3:M:343:ASN:ND2	3:M:343:ASN:N	2.56	0.43
3:M:351:SER:O	3:M:352:GLN:HB3	2.19	0.43
3:M:435:LEU:O	3:M:437:TYR:HE1	2.02	0.43
4:S:38:LEU:HB3	4:S:51:LEU:CD1	2.47	0.43
1:A:92:LEU:CD1	1:A:120:ILE:HA	2.15	0.43
1:A:121:LEU:O	1:A:124:ALA:HB3	2.18	0.43
1:A:395:PHE:CZ	1:A:428:MET:HB2	2.54	0.43
2:B:83:PHE:CE2	2:B:119:SER:N	2.84	0.43
2:B:98:LYS:CG	2:B:137:PHE:HB2	2.48	0.43
2:B:123:LEU:CD2	2:B:138:ALA:CA	2.97	0.43
2:B:130:SER:O	2:B:135:ARG:NH2	2.52	0.43
2:B:200:MET:HE2	2:B:229:HIS:HA	1.69	0.43
2:B:374:PHE:HB3	2:B:402:LEU:HD21	2.01	0.43
3:M:96:ILE:HG21	3:M:125:PHE:CE2	2.54	0.43
3:M:226:PHE:CG	3:M:481:VAL:HG22	2.53	0.43
3:M:242:GLY:CA	3:M:474:THR:CG2	2.94	0.43
3:M:374:TYR:CE1	3:M:376:ILE:HD11	2.54	0.43
4:S:16:LEU:HD11	4:S:129:GLU:CG	2.49	0.43
4:S:17:VAL:CG1	4:S:19:PHE:HE1	2.24	0.43
1:A:63:ASP:C	1:A:63:ASP:OD1	2.57	0.42
1:A:226:SER:CB	1:A:263:LEU:HD23	2.49	0.42
1:A:379:VAL:HG22	1:A:380:ASP:N	2.34	0.42
1:A:555:LEU:O	1:A:558:VAL:HB	2.19	0.42
2:B:59:VAL:O	2:B:59:VAL:CG1	2.66	0.42
2:B:60:ARG:O	2:B:100:LEU:HD13	2.18	0.42
2:B:80:GLN:HA	2:B:115:LEU:CD1	2.48	0.42
2:B:83:PHE:HZ	2:B:119:SER:OG	1.94	0.42
2:B:127:LEU:CD2	2:B:139:LEU:HB2	2.49	0.42
2:B:159:LYS:HD2	2:B:191:GLU:CD	2.39	0.42
2:B:261:PRO:C	2:B:290:SER:CB	2.86	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:285:GLU:N	2:B:285:GLU:OE2	2.52	0.42
2:B:366:LEU:O	2:B:368:ILE:N	2.48	0.42
2:B:397:GLN:O	2:B:400:SER:OG	2.16	0.42
2:B:520:SER:C	2:B:523:PHE:HD2	2.22	0.42
2:B:588:ILE:HG21	2:B:618:PHE:CE1	2.54	0.42
3:M:60:LEU:CD2	3:M:60:LEU:C	2.87	0.42
3:M:215:TYR:HD2	3:M:469:GLY:C	2.23	0.42
3:M:240:ILE:HG21	3:M:444:ALA:HA	2.01	0.42
3:M:466:LEU:HD23	3:M:467:TYR:O	2.18	0.42
1:A:250:ASN:CG	1:A:285:THR:HB	2.40	0.42
1:A:375:VAL:O	1:A:376:GLU:C	2.57	0.42
1:A:419:ILE:O	1:A:419:ILE:HG12	2.20	0.42
1:A:507:GLN:O	1:A:509:PRO:HD2	2.18	0.42
1:A:509:PRO:CA	1:A:547:VAL:HG21	2.49	0.42
1:A:532:LEU:O	1:A:535:ILE:C	2.56	0.42
1:A:552:ILE:CD1	1:A:600:SER:HB2	2.48	0.42
1:A:575:LYS:CE	1:A:611:LEU:HD21	2.46	0.42
2:B:127:LEU:CD2	2:B:157:THR:HG21	2.41	0.42
2:B:167:ALA:O	2:B:202:ASP:CG	2.56	0.42
2:B:174:ALA:C	2:B:214:ALA:HB2	2.40	0.42
2:B:230:PHE:HB3	2:B:298:ASP:CG	2.38	0.42
2:B:230:PHE:HD2	2:B:298:ASP:CA	2.32	0.42
2:B:277:CYS:O	2:B:288:TYR:HA	2.18	0.42
2:B:393:ILE:HG21	2:B:431:MET:HE2	2.01	0.42
2:B:560:ILE:CG2	2:B:561:ASP:N	2.79	0.42
2:B:560:ILE:O	2:B:563:PHE:N	2.51	0.42
3:M:7:ILE:HG23	3:M:75:TRP:C	2.38	0.42
4:S:5:VAL:O	4:S:17:VAL:CA	2.55	0.42
4:S:56:SER:C	4:S:57:LEU:HD12	2.39	0.42
1:A:147:LEU:CG	1:A:166:LEU:HD21	2.45	0.42
1:A:222:ILE:CG2	1:A:233:PHE:HB3	2.42	0.42
1:A:359:LEU:HD22	1:A:367:ILE:CG2	2.48	0.42
1:A:450:TYR:OH	1:A:476:GLN:NE2	2.52	0.42
2:B:139:LEU:CD2	2:B:173:VAL:HG13	2.49	0.42
2:B:549:LEU:CD1	2:B:611:ALA:HB1	2.21	0.42
3:M:226:PHE:CE2	3:M:321:GLY:O	2.69	0.42
3:M:336:ASP:CG	3:M:415:ILE:HB	2.37	0.42
1:A:75:THR:O	1:A:79:MET:HG3	2.19	0.42
1:A:102:GLN:N	4:S:163:THR:C	2.71	0.42
1:A:213:SER:N	4:S:148:ARG:CD	2.57	0.42
1:A:514:GLU:OE1	1:A:514:GLU:HA	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:18:ILE:HD13	2:B:18:ILE:N	2.32	0.42
2:B:47:LEU:HD11	2:B:66:ILE:HA	1.98	0.42
2:B:189:HIS:CD2	2:B:222:HIS:CG	3.08	0.42
2:B:245:GLN:HB3	2:B:309:LEU:HD11	2.02	0.42
2:B:292:GLU:CD	2:B:296:ASP:OD2	2.50	0.42
2:B:303:LEU:CD1	2:B:333:GLN:HG2	2.45	0.42
2:B:310:ILE:HG12	2:B:318:ILE:HG23	2.01	0.42
2:B:350:THR:HA	2:B:351:GLU:OE2	2.20	0.42
2:B:403:ILE:CA	2:B:411:ILE:HD12	2.48	0.42
2:B:437:SER:CA	2:B:478:LEU:CD2	2.71	0.42
2:B:589:SER:O	2:B:591:MET:N	2.52	0.42
3:M:220:GLU:O	3:M:439:TYR:CD1	2.73	0.42
3:M:243:ILE:C	3:M:472:TYR:CD2	2.89	0.42
3:M:276:VAL:HG22	3:M:290:PHE:HD1	1.84	0.42
3:M:410:VAL:HG11	3:M:412:ARG:HH11	1.80	0.42
1:A:143:VAL:HG13	1:A:169:MET:HG3	2.00	0.42
1:A:213:SER:HB2	4:S:143:GLU:HB3	2.01	0.42
1:A:288:THR:OG1	1:A:291:ILE:CG1	2.67	0.42
1:A:339:TYR:HB2	1:A:374:LEU:HD11	2.01	0.42
1:A:349:ILE:O	1:A:350:SER:C	2.56	0.42
1:A:483:LYS:O	1:A:484:VAL:CB	2.65	0.42
2:B:37:TYR:C	2:B:39:SER:H	2.20	0.42
2:B:106:LEU:CD2	2:B:144:ASP:HB2	2.48	0.42
2:B:121:ASN:HA	2:B:153:ILE:HG21	2.01	0.42
2:B:256:CYS:O	2:B:260:LEU:N	2.53	0.42
2:B:340:ILE:CD1	2:B:366:LEU:HD13	2.23	0.42
2:B:399:LEU:HD12	2:B:415:LEU:HD23	2.01	0.42
3:M:243:ILE:HD13	3:M:301:GLU:HB3	2.00	0.42
1:A:102:GLN:C	4:S:167:ILE:HG12	2.32	0.42
1:A:200:PHE:CZ	1:A:236:LEU:HG	2.54	0.42
1:A:348:PHE:HB3	1:A:352:PHE:CE1	2.53	0.42
1:A:552:ILE:CG2	1:A:603:VAL:HG21	2.45	0.42
1:A:555:LEU:HD12	1:A:585:PHE:HE1	1.83	0.42
1:A:558:VAL:C	1:A:560:SER:N	2.73	0.42
2:B:50:LEU:CD1	2:B:58:GLU:HB3	2.49	0.42
2:B:178:ILE:CG2	2:B:217:GLU:CB	2.75	0.42
2:B:195:ILE:O	2:B:196:LEU:C	2.55	0.42
2:B:307:ASN:O	2:B:308:CYS:C	2.57	0.42
2:B:418:TYR:CE1	2:B:419:VAL:HG22	2.53	0.42
2:B:475:ILE:HG21	2:B:514:LEU:HD21	1.98	0.42
1:A:171:ASN:HB2	1:A:202:LYS:HE2	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:182:ILE:HG21	1:A:217:ALA:C	2.39	0.42
1:A:564:ASN:OD1	1:A:622:PRO:HG2	2.17	0.42
1:A:623:MET:O	2:B:617:LEU:HG	2.19	0.42
2:B:37:TYR:CD2	2:B:42:ILE:HA	2.53	0.42
2:B:62:ALA:C	2:B:66:ILE:HD11	2.34	0.42
2:B:215:TYR:OH	2:B:229:HIS:HB3	2.17	0.42
2:B:396:ILE:HD13	2:B:418:TYR:CZ	2.47	0.42
2:B:534:ILE:HA	2:B:598:LEU:HD12	2.02	0.42
2:B:556:LEU:HD12	2:B:614:ILE:CG2	2.49	0.42
2:B:592:TYR:CD1	2:B:593:ASN:N	2.88	0.42
3:M:2:TYR:C	3:M:81:SER:CB	2.77	0.42
3:M:428:VAL:O	3:M:430:LEU:HA	2.19	0.42
4:S:47:GLN:O	4:S:49:SER:N	2.49	0.42
4:S:50:PHE:CA	4:S:77:TYR:HD1	2.28	0.42
4:S:122:ILE:O	4:S:123:PHE:C	2.56	0.42
1:A:91:ILE:HD12	1:A:109:ALA:HB3	2.01	0.42
1:A:100:LEU:HD22	1:A:138:ASN:CB	2.48	0.42
1:A:183:THR:N	4:S:142:ILE:HD13	2.34	0.42
1:A:274:LEU:O	1:A:277:LYS:N	2.52	0.42
1:A:634:ASN:ND2	2:B:554:LYS:H	2.15	0.42
2:B:47:LEU:HD22	2:B:66:ILE:CA	2.50	0.42
2:B:154:ILE:HG21	2:B:180:LEU:CB	2.50	0.42
2:B:336:ASN:HD21	2:B:338:LYS:HB2	1.80	0.42
2:B:383:VAL:CG2	2:B:384:PHE:N	2.83	0.42
2:B:560:ILE:HG22	2:B:561:ASP:H	1.83	0.42
3:M:219:LEU:H	3:M:472:TYR:HB2	1.84	0.42
3:M:245:ASP:C	3:M:246:VAL:HG23	2.40	0.42
4:S:1:MET:H3	4:S:93:GLU:HB2	1.85	0.42
1:A:66:SER:N	4:S:165:SER:CB	2.66	0.42
1:A:97:SER:CA	1:A:98:ASN:O	2.66	0.42
1:A:219:VAL:CG2	1:A:256:LEU:HD23	2.48	0.42
1:A:271:ARG:HH11	1:A:303:MET:HA	1.85	0.42
1:A:336:ILE:HG23	1:A:340:LYS:HE3	2.02	0.42
1:A:401:VAL:HB	1:A:419:ILE:HB	2.02	0.42
2:B:105:LEU:CB	2:B:145:MET:HE1	2.39	0.42
2:B:114:ASN:O	2:B:117:LEU:CB	2.63	0.42
2:B:216:LYS:HB2	2:B:251:LEU:CB	2.50	0.42
2:B:380:LYS:HZ2	3:M:237:THR:CG2	2.33	0.42
2:B:562:ASN:CG	2:B:580:TYR:CG	2.92	0.42
3:M:74:TYR:CE1	3:M:76:CYS:SG	3.13	0.42
3:M:265:ASN:OD1	3:M:313:SER:HB3	2.15	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:16:LEU:CD1	4:S:17:VAL:N	2.80	0.42
4:S:47:GLN:NE2	4:S:47:GLN:C	2.73	0.42
1:A:121:LEU:HD11	1:A:155:THR:OG1	2.20	0.42
1:A:213:SER:HB2	4:S:143:GLU:OE1	2.12	0.42
1:A:263:LEU:O	1:A:264:SER:C	2.53	0.42
1:A:288:THR:OG1	1:A:291:ILE:HD13	2.19	0.42
1:A:421:PRO:CB	1:A:424:TYR:CE1	3.03	0.42
1:A:524:THR:CG2	1:A:565:ASN:HD22	2.32	0.42
1:A:552:ILE:HD13	1:A:600:SER:HB2	2.01	0.42
1:A:627:GLU:HA	2:B:617:LEU:HB2	1.18	0.42
2:B:13:ASP:O	2:B:14:THR:C	2.59	0.42
2:B:90:ILE:O	2:B:98:LYS:CE	2.68	0.42
2:B:151:ALA:C	2:B:153:ILE:H	2.24	0.42
2:B:158:VAL:CB	2:B:177:ILE:CG1	2.89	0.42
2:B:172:GLU:OE1	2:B:175:LEU:HD12	2.19	0.42
2:B:176:ALA:O	2:B:178:ILE:C	2.58	0.42
2:B:177:ILE:CD1	2:B:196:LEU:CG	2.96	0.42
2:B:240:LEU:HB3	2:B:241:ASP:H	1.77	0.42
2:B:408:VAL:HG13	2:B:412:PHE:CE2	2.48	0.42
2:B:527:PRO:HB3	2:B:587:ARG:O	2.20	0.42
3:M:218:LEU:N	3:M:218:LEU:CD1	2.82	0.42
3:M:222:PHE:HD1	3:M:222:PHE:N	2.18	0.42
4:S:7:ILE:CG2	4:S:121:LEU:HD23	2.50	0.42
4:S:14:PRO:C	4:S:15:ARG:HG2	2.40	0.42
1:A:134:TYR:CD2	1:A:136:GLY:N	2.84	0.41
1:A:180:LYS:O	1:A:183:THR:N	2.53	0.41
1:A:183:THR:OG1	4:S:142:ILE:CG1	2.65	0.41
1:A:226:SER:HB3	1:A:263:LEU:CD2	2.50	0.41
1:A:253:ILE:HG21	1:A:281:LEU:CB	2.47	0.41
1:A:374:LEU:CA	1:A:376:GLU:OE1	2.64	0.41
1:A:429:VAL:O	1:A:432:ILE:HB	2.19	0.41
1:A:509:PRO:CA	1:A:547:VAL:CG2	2.98	0.41
2:B:106:LEU:HD22	2:B:144:ASP:HB2	2.02	0.41
2:B:131:ASN:O	2:B:135:ARG:HG3	2.20	0.41
2:B:135:ARG:CZ	2:B:164:ASP:CB	2.95	0.41
2:B:135:ARG:O	2:B:161:LEU:HD21	2.20	0.41
2:B:154:ILE:O	2:B:157:THR:HB	2.20	0.41
2:B:227:HIS:ND1	2:B:292:GLU:HG2	2.35	0.41
2:B:306:LEU:CD1	2:B:325:LEU:HD23	2.44	0.41
2:B:424:PHE:HA	2:B:425:PRO:HD3	1.58	0.41
2:B:447:GLU:HG3	2:B:482:ASN:ND2	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:95:THR:HG22	3:M:99:ILE:HD11	2.01	0.41
3:M:214:LEU:C	3:M:214:LEU:CD2	2.80	0.41
3:M:283:PHE:CD2	3:M:289:THR:OG1	2.60	0.41
3:M:380:ARG:HB3	3:M:412:ARG:N	2.35	0.41
3:M:381:ASN:HD21	3:M:384:GLY:HA3	1.83	0.41
3:M:386:PHE:HB2	3:M:397:TRP:HD1	1.81	0.41
3:M:445:SER:OG	3:M:447:ILE:HG23	2.19	0.41
4:S:47:GLN:HE22	4:S:78:LYS:C	2.22	0.41
4:S:53:THR:OG1	4:S:68:VAL:CA	2.68	0.41
4:S:101:LEU:HG	4:S:135:ILE:CG2	2.49	0.41
1:A:100:LEU:CB	1:A:142:LYS:CG	2.98	0.41
1:A:140:VAL:CG2	1:A:174:ARG:CB	2.97	0.41
1:A:185:LEU:HD22	1:A:189:PHE:HZ	1.80	0.41
1:A:245:VAL:O	1:A:245:VAL:HG22	2.20	0.41
2:B:153:ILE:O	2:B:154:ILE:C	2.58	0.41
3:M:8:THR:OG1	3:M:75:TRP:CB	2.67	0.41
3:M:16:PHE:CE2	3:M:125:PHE:CZ	3.08	0.41
3:M:218:LEU:H	3:M:218:LEU:CD1	2.33	0.41
3:M:223:HIS:HE1	3:M:476:THR:H	1.61	0.41
3:M:256:VAL:HG11	3:M:290:PHE:HB3	1.95	0.41
3:M:276:VAL:CG1	3:M:299:LEU:CD1	2.98	0.41
3:M:343:ASN:ND2	3:M:343:ASN:H	2.13	0.41
3:M:360:LEU:CG	3:M:362:PHE:CE2	3.03	0.41
1:A:145:ILE:O	1:A:147:LEU:N	2.54	0.41
1:A:170:LEU:HB2	1:A:202:LYS:CG	2.50	0.41
1:A:175:PRO:HG3	1:A:211:ASP:HB2	2.01	0.41
1:A:220:SER:OG	4:S:141:VAL:N	2.26	0.41
1:A:270:LEU:HD12	1:A:274:LEU:CD2	2.50	0.41
1:A:385:LYS:NZ	1:A:445:ASN:OD1	2.52	0.41
1:A:403:LEU:CD2	1:A:421:PRO:O	2.64	0.41
1:A:557:LYS:CD	2:B:606:ASP:HB2	2.50	0.41
1:A:567:GLN:C	1:A:569:ASP:N	2.72	0.41
1:A:589:SER:HB2	1:A:601:VAL:HG22	1.89	0.41
2:B:127:LEU:HD11	2:B:142:LEU:CD1	2.50	0.41
2:B:169:VAL:O	2:B:173:VAL:N	2.47	0.41
2:B:212:VAL:CG1	2:B:248:LEU:CD2	2.97	0.41
2:B:260:LEU:CG	2:B:291:TYR:CE1	2.90	0.41
2:B:285:GLU:N	2:B:285:GLU:CD	2.73	0.41
3:M:244:VAL:CG2	3:M:472:TYR:HE2	2.32	0.41
3:M:246:VAL:CB	3:M:297:PHE:CE1	3.02	0.41
3:M:290:PHE:CD1	3:M:299:LEU:CD1	3.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:306:LEU:HD22	3:M:310:VAL:HG23	2.02	0.41
3:M:311:LYS:O	3:M:312:GLN:C	2.55	0.41
3:M:411:LEU:O	3:M:412:ARG:HG3	2.20	0.41
4:S:58:LEU:HD13	4:S:68:VAL:CG1	2.50	0.41
1:A:73:LYS:O	1:A:76:TYR:HB2	2.21	0.41
1:A:92:LEU:O	1:A:94:VAL:N	2.53	0.41
1:A:104:ARG:N	4:S:167:ILE:HD11	2.35	0.41
1:A:332:TYR:CE2	1:A:336:ILE:HD12	2.55	0.41
1:A:338:PHE:O	1:A:341:ILE:HB	2.21	0.41
1:A:356:ILE:O	1:A:359:LEU:HB2	2.20	0.41
1:A:480:LEU:C	1:A:480:LEU:CD1	2.88	0.41
2:B:252:LEU:CB	2:B:302:PHE:CG	2.99	0.41
2:B:296:ASP:C	2:B:298:ASP:N	2.72	0.41
2:B:389:ILE:O	2:B:393:ILE:CG1	2.68	0.41
2:B:519:ALA:CB	2:B:555:LEU:HD12	2.49	0.41
2:B:528:ASP:HA	2:B:531:ARG:NH1	2.35	0.41
3:M:319:SER:HB2	3:M:346:ASN:C	2.40	0.41
4:S:58:LEU:HA	4:S:68:VAL:HG22	2.02	0.41
1:A:125:THR:OG1	1:A:158:LEU:HB2	2.21	0.41
1:A:175:PRO:HA	1:A:214:VAL:HG21	2.01	0.41
1:A:215:VAL:HG21	1:A:243:ILE:HG23	1.18	0.41
1:A:420:ILE:HA	1:A:421:PRO:HD3	1.83	0.41
1:A:462:GLN:NE2	2:B:1:MET:CE	2.83	0.41
1:A:523:SER:OG	1:A:562:TRP:CD1	2.57	0.41
1:A:626:SER:OG	2:B:616:SER:HB2	2.21	0.41
2:B:63:MET:HB3	2:B:100:LEU:CD1	2.51	0.41
2:B:105:LEU:C	2:B:145:MET:HE1	2.40	0.41
2:B:266:VAL:CA	2:B:289:PRO:HB2	2.48	0.41
2:B:311:TYR:CD2	2:B:342:ALA:HB1	2.53	0.41
2:B:357:GLU:CG	2:B:361:GLN:NE2	2.80	0.41
2:B:458:MET:C	2:B:460:SER:N	2.73	0.41
2:B:533:LEU:O	2:B:536:ASN:N	2.42	0.41
2:B:559:ASP:CG	2:B:563:PHE:CE1	2.92	0.41
3:M:8:THR:OG1	3:M:75:TRP:HB2	2.20	0.41
3:M:223:HIS:C	3:M:479:PHE:CG	2.83	0.41
3:M:374:TYR:O	3:M:390:ILE:CG2	2.68	0.41
1:A:100:LEU:CB	4:S:162:SER:HB2	2.51	0.41
1:A:140:VAL:O	1:A:141:VAL:O	2.38	0.41
1:A:396:VAL:O	1:A:397:ASP:OD1	2.37	0.41
2:B:25:VAL:HG22	2:B:35:TYR:HB3	2.03	0.41
2:B:123:LEU:HB2	2:B:142:LEU:CD1	2.45	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:135:ARG:HH22	2:B:164:ASP:CB	2.27	0.41
2:B:167:ALA:O	2:B:207:VAL:CG1	2.68	0.41
2:B:537:PHE:CE1	2:B:545:ARG:CB	3.03	0.41
2:B:617:LEU:O	2:B:618:PHE:C	2.58	0.41
3:M:80:THR:O	3:M:81:SER:CB	2.69	0.41
3:M:118:TYR:CD2	3:M:118:TYR:O	2.74	0.41
3:M:221:THR:HG22	3:M:223:HIS:CE1	2.55	0.41
3:M:352:GLN:CB	3:M:401:LYS:O	2.69	0.41
3:M:379:LEU:HD13	3:M:386:PHE:CE1	2.56	0.41
3:M:429:ASP:O	3:M:430:LEU:C	2.57	0.41
1:A:100:LEU:CD1	1:A:141:VAL:HG11	2.49	0.41
1:A:140:VAL:HG13	1:A:176:TYR:HB2	1.87	0.41
1:A:163:ALA:C	1:A:165:ASP:N	2.71	0.41
1:A:222:ILE:HG23	1:A:233:PHE:CB	2.43	0.41
1:A:436:CYS:CB	1:A:450:TYR:CZ	3.04	0.41
1:A:536:MET:HB3	1:A:555:LEU:HD21	2.01	0.41
1:A:559:PHE:CD2	1:A:578:LEU:HD23	2.55	0.41
1:A:609:LEU:C	1:A:611:LEU:N	2.73	0.41
2:B:137:PHE:O	2:B:140:SER:N	2.50	0.41
2:B:159:LYS:CD	2:B:191:GLU:CD	2.89	0.41
2:B:181:TYR:CD2	2:B:218:CYS:CB	2.90	0.41
2:B:212:VAL:CG2	2:B:233:TYR:CD2	3.03	0.41
2:B:257:LYS:O	2:B:260:LEU:HG	2.20	0.41
2:B:319:LEU:C	2:B:321:CYS:N	2.73	0.41
2:B:467:VAL:CG1	2:B:471:TYR:CE1	3.04	0.41
2:B:537:PHE:CA	2:B:598:LEU:HD13	2.37	0.41
3:M:347:PHE:CE2	3:M:350:VAL:CG1	3.04	0.41
1:A:147:LEU:O	1:A:184:ALA:HB1	2.21	0.41
1:A:179:LYS:O	4:S:142:ILE:HD12	2.16	0.41
1:A:212:ILE:CB	4:S:145:ASN:ND2	2.81	0.41
1:A:401:VAL:HG11	1:A:419:ILE:HD12	2.00	0.41
1:A:516:ILE:CD1	1:A:551:LEU:HB2	2.51	0.41
1:A:630:PRO:O	2:B:554:LYS:HB2	2.21	0.41
2:B:78:ASP:OD1	2:B:80:GLN:CB	2.56	0.41
2:B:158:VAL:CB	2:B:177:ILE:HG13	2.48	0.41
2:B:328:LEU:O	2:B:329:ALA:C	2.50	0.41
2:B:344:VAL:HG22	2:B:363:ILE:HD12	1.94	0.41
2:B:520:SER:CA	2:B:523:PHE:CD2	3.03	0.41
2:B:537:PHE:HA	2:B:540:GLU:CD	2.40	0.41
3:M:217:ASP:OD2	3:M:440:ILE:CG2	2.69	0.41
3:M:342:LEU:CD1	3:M:411:LEU:HB3	2.27	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:410:VAL:CG1	3:M:412:ARG:NH1	2.73	0.41
1:A:99:LYS:HE2	1:A:101:GLN:HB3	2.02	0.41
1:A:375:VAL:O	1:A:378:ILE:N	2.53	0.41
1:A:420:ILE:HG23	1:A:424:TYR:HB2	2.03	0.41
1:A:443:SER:O	1:A:445:ASN:N	2.47	0.41
1:A:461:CYS:SG	1:A:469:LEU:CB	3.02	0.41
1:A:567:GLN:O	1:A:569:ASP:N	2.53	0.41
1:A:588:LEU:C	1:A:590:TYR:N	2.72	0.41
1:A:626:SER:OG	2:B:617:LEU:HG	2.21	0.41
2:B:48:VAL:HG22	2:B:82:TYR:CE2	2.53	0.41
2:B:56:SER:HB3	2:B:92:THR:HG23	1.90	0.41
2:B:106:LEU:HD11	2:B:144:ASP:C	2.29	0.41
2:B:169:VAL:HG12	2:B:173:VAL:HG21	2.01	0.41
2:B:216:LYS:HD3	2:B:251:LEU:CB	2.49	0.41
2:B:231:ARG:C	2:B:233:TYR:H	2.24	0.41
2:B:343:LEU:CD2	2:B:363:ILE:N	2.84	0.41
2:B:444:THR:O	2:B:446:TRP:N	2.53	0.41
2:B:567:GLN:CB	2:B:569:THR:OG1	2.68	0.41
2:B:587:ARG:O	2:B:591:MET:HG2	2.21	0.41
3:M:96:ILE:HD13	3:M:96:ILE:HA	1.94	0.41
3:M:218:LEU:CB	3:M:442:GLN:O	2.69	0.41
3:M:220:GLU:CD	3:M:439:TYR:HB2	2.41	0.41
3:M:222:PHE:CB	3:M:240:ILE:HG12	2.51	0.41
3:M:224:VAL:HB	3:M:479:PHE:HD1	1.84	0.41
3:M:261:ASN:CG	3:M:262:THR:N	2.74	0.41
3:M:356:LEU:HD21	3:M:358:ILE:HG12	1.99	0.41
4:S:4:ALA:HB2	4:S:19:PHE:CD2	2.56	0.41
4:S:117:ASN:CB	4:S:120:ASP:OD2	2.68	0.41
4:S:127:THR:CG2	4:S:153:VAL:HG13	2.51	0.41
1:A:104:ARG:HG3	1:A:145:ILE:CB	2.49	0.41
1:A:154:ILE:HG22	1:A:191:GLN:CG	2.39	0.41
1:A:180:LYS:HE3	4:S:156:LEU:CD1	2.50	0.41
1:A:509:PRO:O	1:A:547:VAL:HG22	2.22	0.41
1:A:563:CYS:SG	1:A:621:LEU:CG	3.08	0.41
2:B:13:ASP:O	2:B:16:LYS:O	2.38	0.41
2:B:95:THR:HA	2:B:134:LEU:HD23	2.02	0.41
2:B:143:SER:O	2:B:179:LYS:CB	2.69	0.41
2:B:151:ALA:HB3	2:B:188:TYR:CE1	2.56	0.41
2:B:178:ILE:CD1	2:B:214:ALA:C	2.82	0.41
2:B:309:LEU:HD13	2:B:309:LEU:HA	1.87	0.41
2:B:336:ASN:CB	2:B:339:PHE:CD1	3.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:323:MET:CE	3:M:342:LEU:CG	2.99	0.41
3:M:436:GLU:HG3	3:M:436:GLU:O	2.21	0.41
1:A:100:LEU:HD23	4:S:162:SER:OG	2.20	0.40
1:A:200:PHE:CZ	1:A:235:GLN:HB2	2.56	0.40
1:A:316:LEU:HD11	1:A:348:PHE:CE2	2.43	0.40
1:A:338:PHE:CE2	1:A:352:PHE:CE2	3.08	0.40
1:A:376:GLU:CD	1:A:376:GLU:N	2.73	0.40
1:A:594:PHE:CE1	2:B:434:LYS:CD	2.98	0.40
2:B:102:HIS:CD2	2:B:123:LEU:HD21	2.56	0.40
2:B:123:LEU:HB2	2:B:142:LEU:HD21	2.03	0.40
2:B:159:LYS:CA	2:B:195:ILE:HD13	2.39	0.40
2:B:170:ARG:O	2:B:171:GLY:C	2.58	0.40
2:B:178:ILE:HA	2:B:178:ILE:HD13	1.89	0.40
2:B:334:MET:C	2:B:336:ASN:N	2.73	0.40
2:B:374:PHE:CD2	2:B:402:LEU:HD21	2.53	0.40
2:B:518:ILE:O	2:B:518:ILE:CG1	2.69	0.40
3:M:12:ASN:OD1	3:M:45:SER:OG	2.35	0.40
3:M:66:PHE:HB3	3:M:77:LEU:HD21	1.78	0.40
3:M:240:ILE:HG22	3:M:444:ALA:CA	2.51	0.40
3:M:446:GLY:C	3:M:448:TYR:H	2.21	0.40
3:M:478:ASN:C	3:M:479:PHE:O	2.59	0.40
1:A:71:VAL:HG11	1:A:105:VAL:CB	2.51	0.40
1:A:78:GLU:OE1	1:A:113:SER:HB3	2.20	0.40
1:A:92:LEU:CD2	1:A:124:ALA:N	2.85	0.40
1:A:92:LEU:HD22	1:A:123:LEU:CB	2.50	0.40
1:A:141:VAL:HB	4:S:159:ALA:CA	2.46	0.40
1:A:145:ILE:C	1:A:147:LEU:N	2.72	0.40
1:A:182:ILE:HD13	1:A:218:ALA:N	2.37	0.40
1:A:251:TRP:CD1	4:S:104:THR:HG23	2.55	0.40
1:A:420:ILE:HG23	1:A:421:PRO:CD	2.48	0.40
1:A:429:VAL:CG2	1:A:469:LEU:HD11	2.42	0.40
1:A:461:CYS:O	1:A:463:ASP:N	2.49	0.40
1:A:629:LEU:O	1:A:631:SER:N	2.52	0.40
2:B:20:ARG:HH11	2:B:35:TYR:HE1	1.67	0.40
2:B:37:TYR:O	2:B:40:GLN:C	2.59	0.40
2:B:139:LEU:HD23	2:B:173:VAL:HA	0.42	0.40
2:B:258:GLN:HA	2:B:258:GLN:OE1	2.21	0.40
2:B:565:GLN:C	2:B:574:ASN:HD21	2.20	0.40
3:M:292:PRO:HA	3:M:293:PRO:HD3	1.53	0.40
3:M:360:LEU:CD1	3:M:433:VAL:CB	2.93	0.40
3:M:433:VAL:O	3:M:435:LEU:HD12	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S:90:ASP:C	4:S:92:GLN:H	2.25	0.40
4:S:108:SER:OG	4:S:149:ILE:CG2	2.60	0.40
1:A:102:GLN:CG	4:S:164:ASP:C	2.89	0.40
1:A:104:ARG:CG	1:A:145:ILE:CD1	2.95	0.40
1:A:231:GLN:CB	1:A:232:PRO:HD3	2.44	0.40
1:A:247:ILE:HG21	1:A:252:ILE:CG2	2.52	0.40
1:A:247:ILE:HG21	1:A:252:ILE:HB	2.03	0.40
1:A:392:MET:CE	1:A:460:LEU:HD12	2.51	0.40
1:A:535:ILE:O	1:A:536:MET:HG3	2.20	0.40
2:B:208:ILE:HD13	2:B:240:LEU:HD11	2.04	0.40
2:B:286:ILE:CG2	2:B:288:TYR:CE2	2.82	0.40
2:B:306:LEU:O	2:B:309:LEU:HB2	2.21	0.40
2:B:363:ILE:CG2	2:B:398:ILE:HD11	2.47	0.40
2:B:566:ALA:HB2	2:B:581:TYR:CD2	2.56	0.40
3:M:249:TYR:HE1	3:M:467:TYR:CZ	2.37	0.40
3:M:320:ILE:CG2	3:M:439:TYR:OH	2.70	0.40
3:M:376:ILE:HG22	3:M:379:LEU:CD1	2.51	0.40
4:S:85:PHE:HZ	4:S:109:LEU:HD23	1.85	0.40
1:A:84:MET:O	1:A:85:ALA:O	2.36	0.40
1:A:125:THR:HA	1:A:158:LEU:HD13	2.04	0.40
1:A:141:VAL:HG21	4:S:157:ASN:N	2.36	0.40
1:A:149:GLY:C	1:A:151:SER:N	2.72	0.40
1:A:150:LEU:CG	1:A:162:ILE:HG12	2.52	0.40
1:A:174:ARG:HA	1:A:175:PRO:HD2	1.66	0.40
1:A:175:PRO:CG	1:A:211:ASP:OD2	2.69	0.40
1:A:259:LEU:C	1:A:259:LEU:HD23	2.41	0.40
1:A:260:PHE:CZ	1:A:274:LEU:CG	3.04	0.40
1:A:288:THR:HB	1:A:322:PHE:CZ	2.42	0.40
1:A:313:MET:CA	1:A:348:PHE:CZ	3.02	0.40
1:A:326:GLN:HA	1:A:331:ARG:NE	2.36	0.40
1:A:359:LEU:HD22	1:A:367:ILE:HG21	2.03	0.40
1:A:504:ILE:O	1:A:506:LYS:N	2.52	0.40
1:A:557:LYS:HG2	2:B:606:ASP:N	2.36	0.40
2:B:17:VAL:CB	3:M:119:ASP:OD2	2.69	0.40
2:B:41:ASN:HD22	2:B:43:ASN:HA	1.87	0.40
2:B:105:LEU:C	2:B:145:MET:CE	2.90	0.40
2:B:132:SER:C	2:B:169:VAL:HG23	2.41	0.40
2:B:171:GLY:O	2:B:174:ALA:HB3	2.22	0.40
2:B:257:LYS:CA	2:B:260:LEU:HG	2.51	0.40
2:B:278:PRO:HB3	2:B:289:PRO:O	2.22	0.40
2:B:306:LEU:HD22	2:B:321:CYS:CB	2.52	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:319:LEU:O	2:B:321:CYS:N	2.55	0.40
2:B:353:GLN:HE21	3:M:47:SER:CB	1.91	0.40
2:B:542:PRO:O	2:B:607:ILE:HD13	2.22	0.40
3:M:7:ILE:HG22	3:M:75:TRP:O	2.17	0.40
3:M:104:PHE:O	3:M:105:ASP:CB	2.69	0.40
4:S:87:PHE:CD1	4:S:87:PHE:N	2.89	0.40
1:A:192:TYR:CD2	1:A:192:TYR:O	2.75	0.40
1:A:220:SER:CB	4:S:142:ILE:CA	3.00	0.40
1:A:220:SER:CB	4:S:142:ILE:HA	2.49	0.40
1:A:566:PHE:CZ	1:A:618:THR:HB	2.53	0.40
1:A:637:GLU:HB2	2:B:513:TRP:CG	2.57	0.40
2:B:38:TYR:CA	2:B:42:ILE:H	2.13	0.40
2:B:64:LYS:CG	2:B:100:LEU:CD1	2.98	0.40
2:B:70:MET:CB	2:B:104:TYR:CE1	2.99	0.40
2:B:84:ALA:O	2:B:87:VAL:HB	2.21	0.40
2:B:200:MET:HE3	2:B:232:ARG:N	2.24	0.40
2:B:361:GLN:HG2	2:B:394:TRP:CE2	2.56	0.40
2:B:451:MET:CE	2:B:489:ILE:CG1	2.98	0.40
3:M:220:GLU:HA	3:M:474:THR:CG2	2.50	0.40
3:M:270:PRO:N	3:M:302:TYR:CD1	2.90	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	A	574/964 (60%)	511 (89%)	39 (7%)	24 (4%)	2	17
2	B	619/809 (76%)	474 (77%)	79 (13%)	66 (11%)	0	6
3	M	385/483 (80%)	315 (82%)	52 (14%)	18 (5%)	2	16
4	S	166/194 (86%)	150 (90%)	12 (7%)	4 (2%)	5	27
All	All	1744/2450 (71%)	1450 (83%)	182 (10%)	112 (6%)	2	13

All (112) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	267	GLU
1	A	278	ILE
1	A	279	LEU
1	A	305	GLU
1	A	401	VAL
1	A	405	THR
1	A	440	ASN
1	A	441	TYR
1	A	449	TRP
1	A	507	GLN
2	B	14	THR
2	B	18	ILE
2	B	19	THR
2	B	43	ASN
2	B	78	ASP
2	B	154	ILE
2	B	156	HIS
2	B	176	ALA
2	B	177	ILE
2	B	179	LYS
2	B	192	LEU
2	B	194	ASP
2	B	196	LEU
2	B	197	LYS
2	B	198	GLU
2	B	200	MET
2	B	215	TYR
2	B	218	CYS
2	B	228	GLY
2	B	261	PRO
2	B	274	PRO
2	B	287	GLU
2	B	296	ASP
2	B	436	LEU
2	B	558	TYR
2	B	560	ILE
2	B	561	ASP
2	B	564	LYS
2	B	565	GLN
2	B	568	VAL
2	B	578	PRO
2	B	583	PHE

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Mol	Chain	Res	Type
3	M	105	ASP
3	M	282	VAL
3	M	352	GLN
3	M	447	ILE
3	M	470	ALA
4	S	50	PHE
1	A	116	LYS
1	A	126	ASN
1	A	174	ARG
1	A	536	MET
2	B	3	ASP
2	B	29	LYS
2	B	135	ARG
2	B	144	ASP
2	B	158	VAL
2	B	178	ILE
2	B	199	LEU
2	B	227	HIS
2	B	332	LEU
2	B	383	VAL
2	B	432	ALA
3	M	45	SER
3	M	72	LEU
3	M	82	LYS
3	M	316	ARG
3	M	317	MET
3	M	429	ASP
1	A	256	LEU
1	A	421	PRO
1	A	571	ARG
1	A	608	ARG
2	B	15	ALA
2	B	38	TYR
2	B	232	ARG
2	B	337	THR
2	B	503	LEU
2	B	538	SER
2	B	590	GLN
3	M	86	PRO
3	M	254	PRO
3	M	355	ASP
1	A	306	GLU

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Mol	Chain	Res	Type
1	A	406	GLY
2	B	213	LEU
2	B	222	HIS
2	B	260	LEU
2	B	523	PHE
2	B	600	LYS
3	M	118	TYR
3	M	363	ASN
4	S	15	ARG
4	S	81	ALA
1	A	419	ILE
2	B	20	ARG
2	B	142	LEU
2	B	535	GLN
2	B	573	GLU
3	M	57	GLY
1	A	302	ASN
1	A	439	ASP
2	B	37	TYR
2	B	155	LEU
2	B	372	THR
2	B	42	ILE
2	B	542	PRO
3	M	268	GLY
2	B	8	ILE
2	B	347	VAL
4	S	26	PRO
1	A	601	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	A	536/898 (60%)	527 (98%)	9 (2%)	56	72
2	B	564/738 (76%)	548 (97%)	16 (3%)	38	57
3	M	353/441 (80%)	342 (97%)	11 (3%)	35	54

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	S	157/175 (90%)	155 (99%)	2 (1%)	65	77
All	All	1610/2252 (72%)	1572 (98%)	38 (2%)	45	62

All (38) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	175	PRO
1	A	186	PHE
1	A	196	LEU
1	A	241	TYR
1	A	288	THR
1	A	292	TYR
1	A	376	GLU
1	A	418	ILE
1	A	439	ASP
2	B	1	MET
2	B	4	SER
2	B	10	SER
2	B	12	LEU
2	B	14	THR
2	B	16	LYS
2	B	19	THR
2	B	29	LYS
2	B	30	LEU
2	B	233	TYR
2	B	261	PRO
2	B	285	GLU
2	B	352	ASN
2	B	418	TYR
2	B	453	TRP
2	B	592	TYR
3	M	68	VAL
3	M	100	LEU
3	M	118	TYR
3	M	214	LEU
3	M	293	PRO
3	M	343	ASN
3	M	352	GLN
3	M	437	TYR
3	M	439	TYR
3	M	472	TYR
3	M	479	PHE

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Mol	Chain	Res	Type
4	S	38	LEU
4	S	47	GLN

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (26) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	101	GLN
1	A	102	GLN
1	A	199	ASN
1	A	235	GLN
1	A	345	ASN
1	A	440	ASN
1	A	462	GLN
1	A	565	ASN
2	B	41	ASN
2	B	156	HIS
2	B	307	ASN
2	B	333	GLN
2	B	352	ASN
2	B	441	GLN
2	B	486	HIS
2	B	547	GLN
2	B	567	GLN
2	B	574	ASN
3	M	17	GLN
3	M	117	ASN
3	M	241	HIS
3	M	326	HIS
3	M	343	ASN
3	M	346	ASN
4	S	47	GLN
4	S	103	GLN

### 5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

## 5.4 Non-standard residues in protein, DNA, RNA chains ⓘ

There are no non-standard protein/DNA/RNA residues in this entry.

## 5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

There are no chain breaks in this entry.

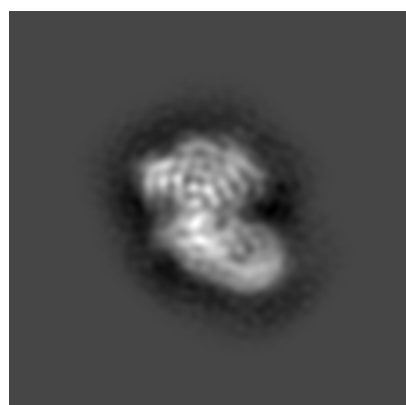
## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-13189. These allow visual inspection of the internal detail of the map and identification of artifacts.

No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

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

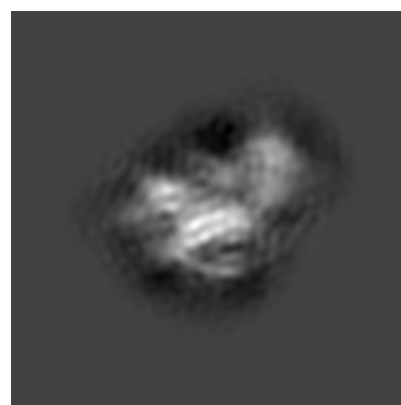
#### 6.1.1 Primary map



X



Y

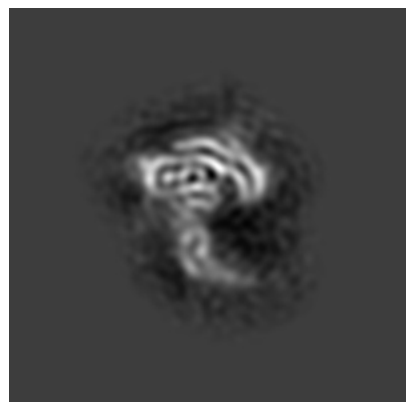


Z

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

### 6.2 Central slices [i](#)

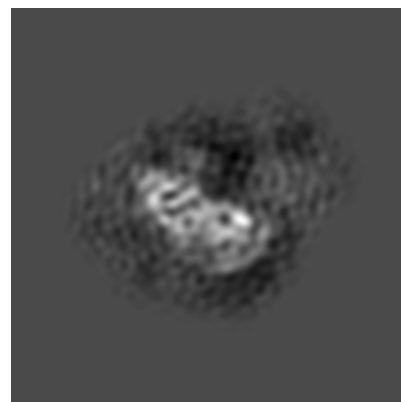
#### 6.2.1 Primary map



X Index: 132



Y Index: 132



Z Index: 132

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

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

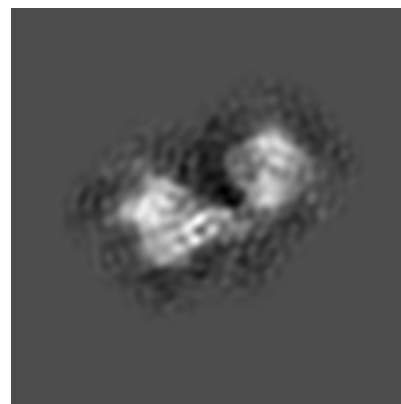
### 6.3.1 Primary map



X Index: 134



Y Index: 123



Z Index: 112

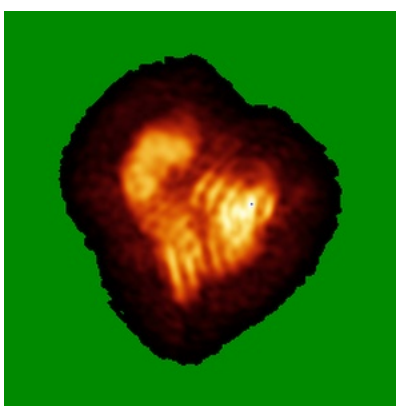
The images above show the largest variance slices of the map in three orthogonal directions.

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

### 6.4.1 Primary map



X



Y

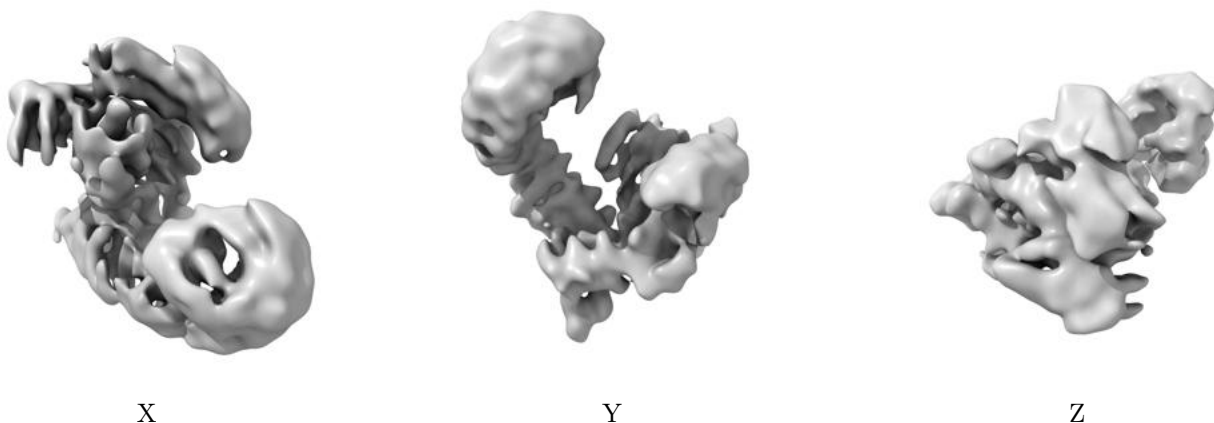


Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

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

### 6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.0155. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

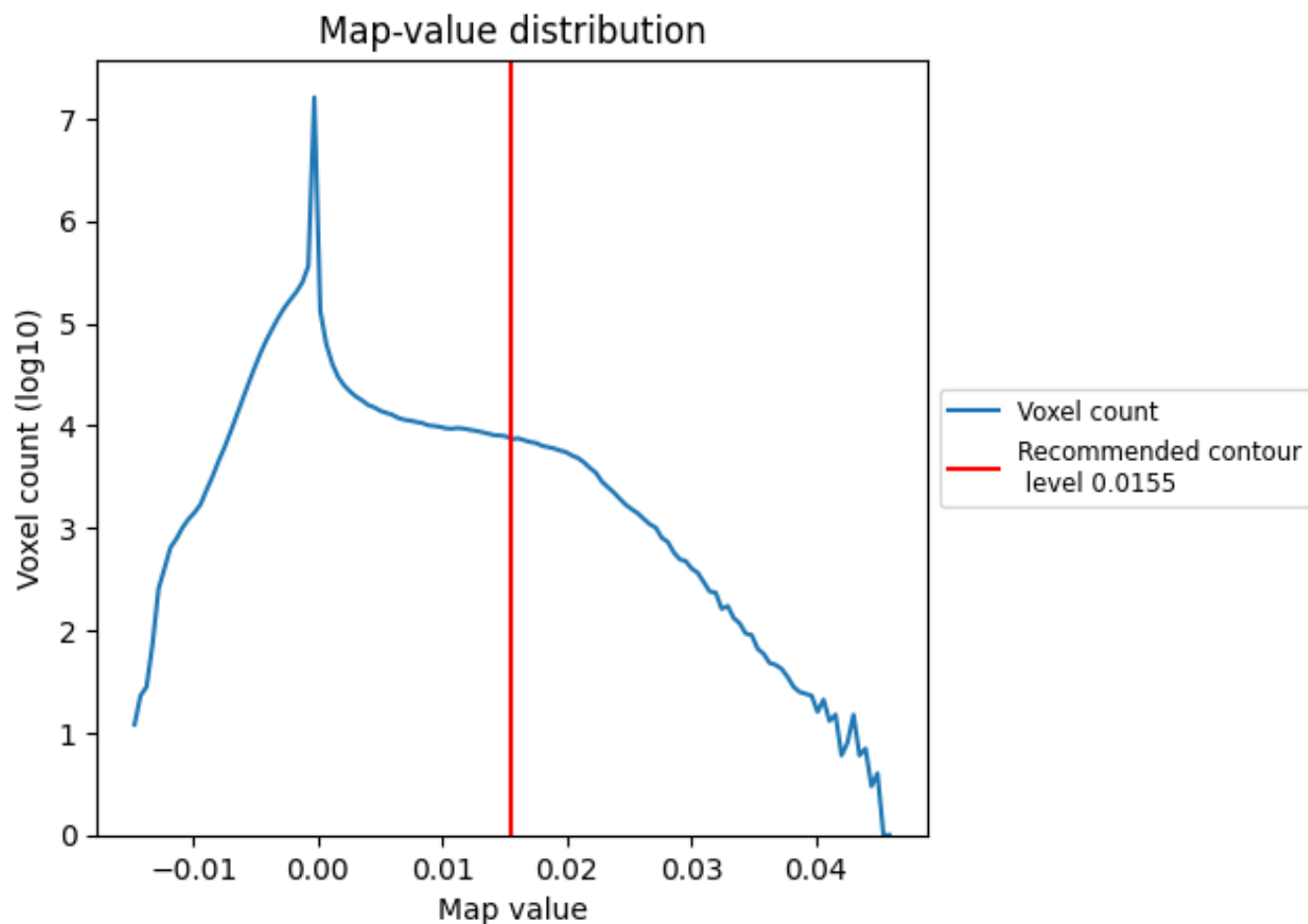
## 6.6 Mask visualisation [i](#)

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

## 7 Map analysis [i](#)

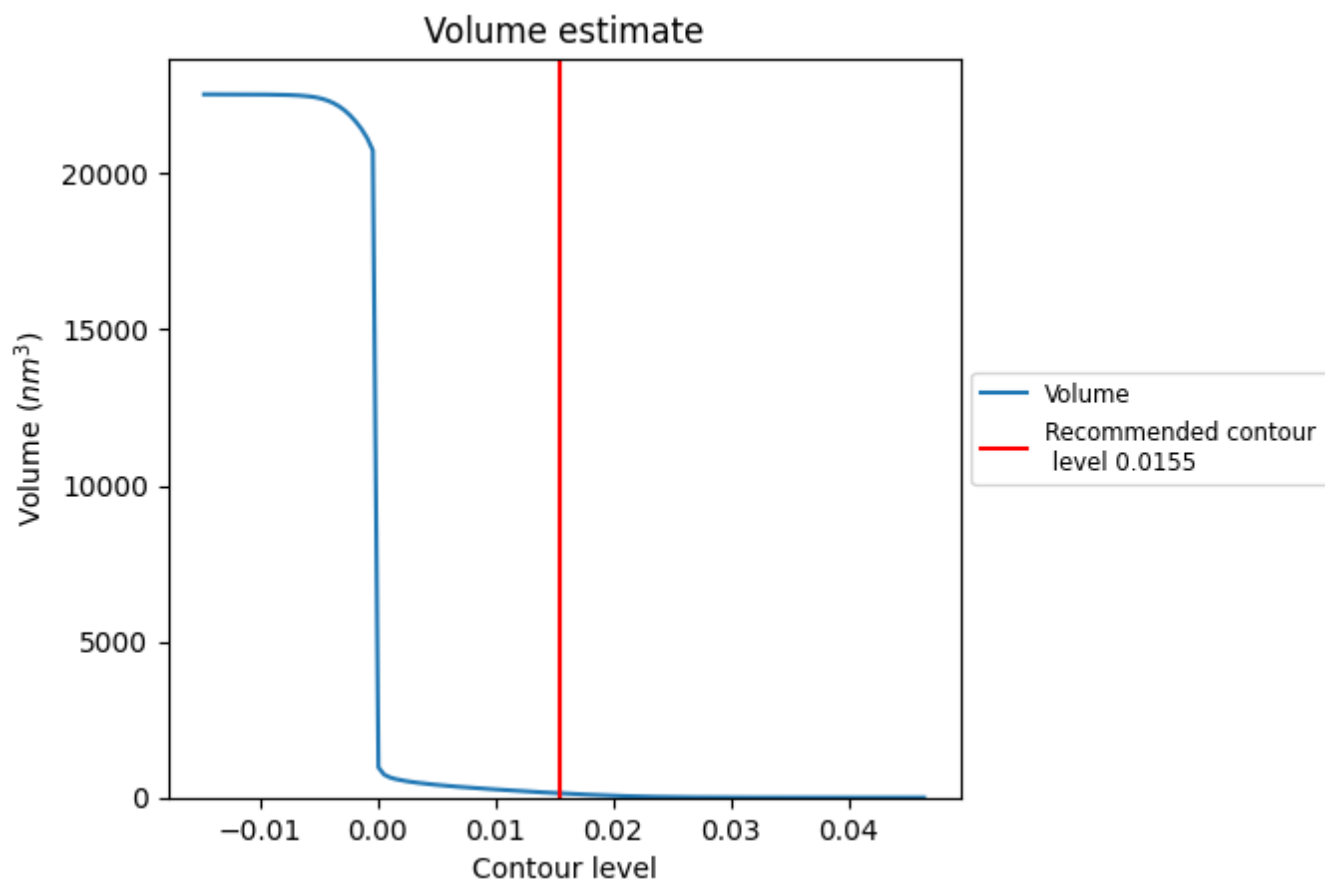
This section contains the results of statistical analysis of the map.

### 7.1 Map-value distribution [i](#)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

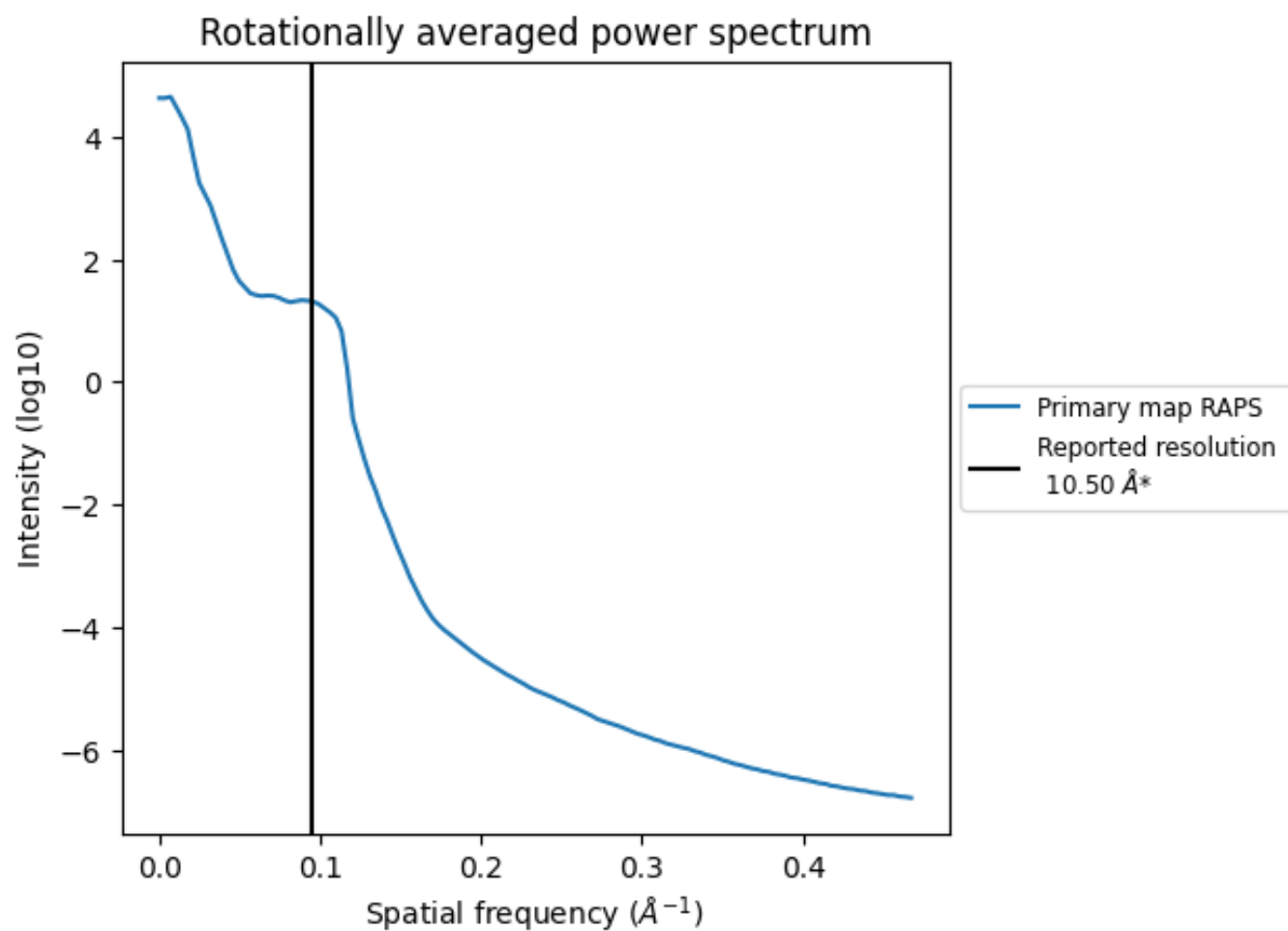
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 137 nm<sup>3</sup>; this corresponds to an approximate mass of 124 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.

### 7.3 Rotationally averaged power spectrum ⓘ



\*Reported resolution corresponds to spatial frequency of 0.095 Å<sup>-1</sup>

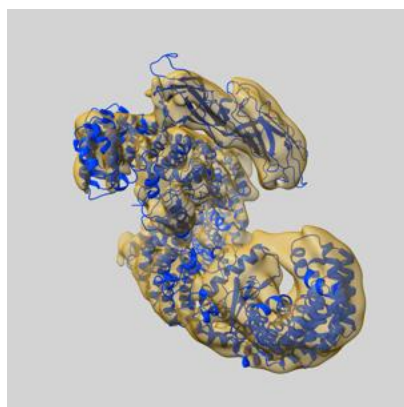
## 8 Fourier-Shell correlation ⓘ

This section was not generated. No FSC curve or half-maps provided.

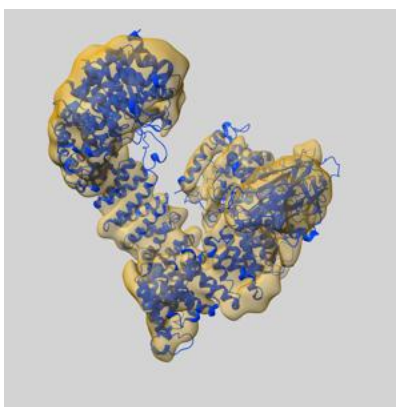
## 9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-13189 and PDB model 7P3Z. Per-residue inclusion information can be found in [section 3](#) on [page 5](#).

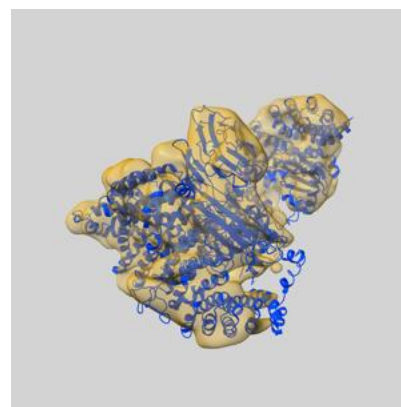
### 9.1 Map-model overlay [i](#)



X



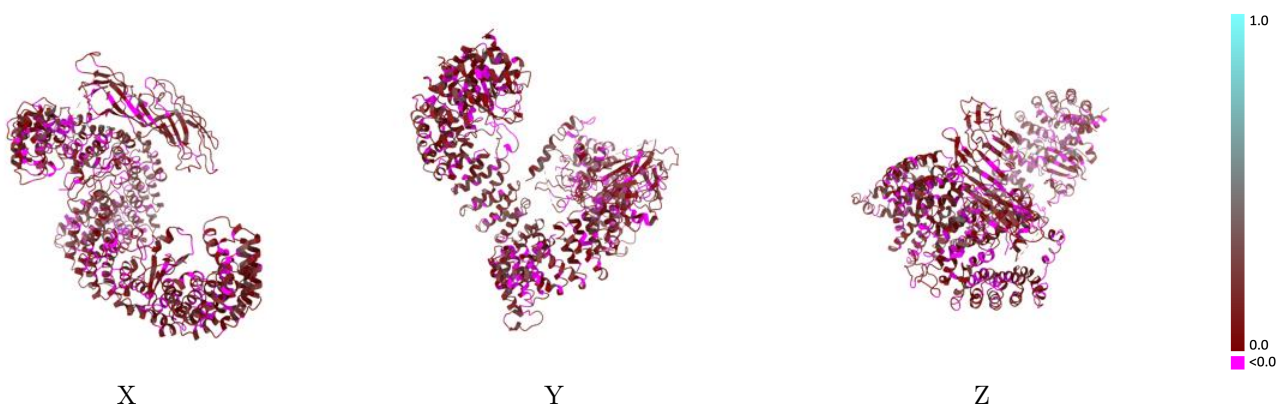
Y



Z

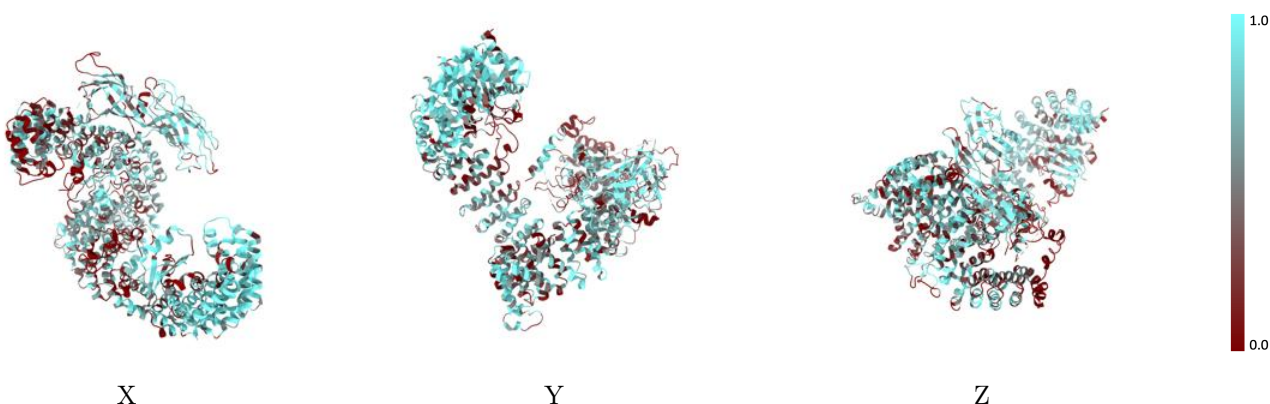
The images above show the 3D surface view of the map at the recommended contour level 0.0155 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



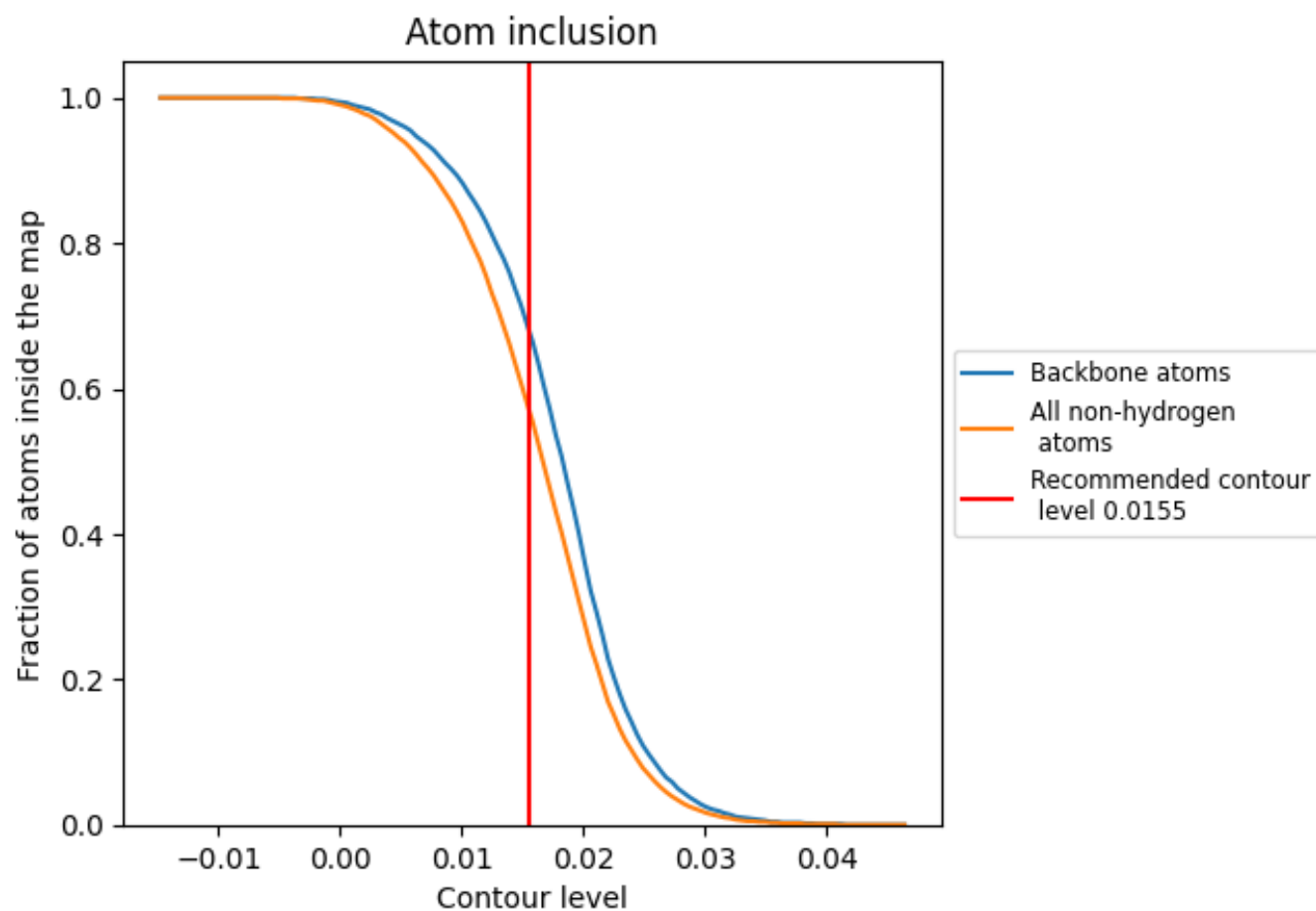
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.0155).

## 9.4 Atom inclusion [i](#)



At the recommended contour level, 68% of all backbone atoms, 57% 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 (0.0155) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	<div></div> 0.5720	<div></div> 0.0890
A	<div></div> 0.6260	<div></div> 0.1070
B	<div></div> 0.5360	<div></div> 0.0790
M	<div></div> 0.5570	<div></div> 0.0870
S	<div></div> 0.5510	<div></div> 0.0730

