



## Full wwPDB EM Validation Report ⓘ

Jun 25, 2025 – 01:56 pm BST

PDB ID : 6FKI / pdb\_00006fki  
EMDB ID : EMD-4272  
Title : Chloroplast F1Fo conformation 3  
Authors : Hahn, A.; Vonck, J.; Mills, D.J.; Meier, T.; Kuehlbrandt, W.  
Deposited on : 2018-01-24  
Resolution : 4.30 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev118  
Mogul : 1.8.4, CSD as541be (2020)  
MolProbity : 4-5-2 with Phenix2.0rc1  
buster-report : 1.1.7 (2018)  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
MapQ : 1.9.13  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.44

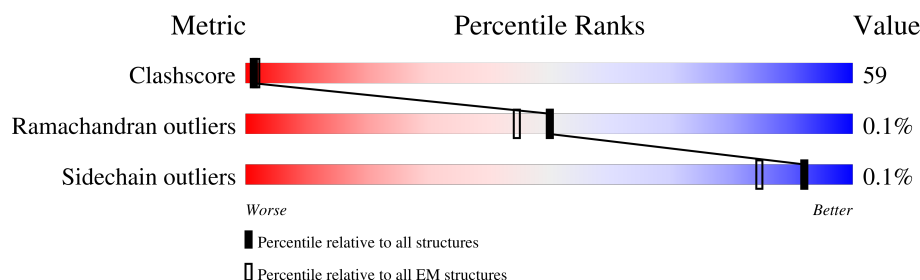
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*ELECTRON MICROSCOPY*

The reported resolution of this entry is 4.30 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	247	
2	b	184	
3	d	257	
4	p	222	
5	e	134	
6	g	364	
7	G	81	
7	H	81	

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Mol	Chain	Length	Quality of chain
7	I	81	
7	J	81	
7	K	81	
7	L	81	
7	M	81	
7	N	81	
7	O	81	
7	P	81	
7	Q	81	
7	R	81	
7	S	81	
7	T	81	
8	A	507	
8	C	507	
8	E	507	
9	B	498	
9	D	498	
9	F	498	

## 2 Entry composition

There are 12 unique types of molecules in this entry. The entry contains 39231 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 ATP synthase subunit a, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	a	223	Total	C	N	O	S	0	0
			1741	1172	268	299	2		

- Molecule 2 is a protein called ATP synthase subunit b, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	b	161	Total	C	N	O	S	0	0
			1299	812	236	246	5		

- Molecule 3 is a protein called ATP synthase delta chain, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	d	179	Total	C	N	O	S	0	0
			1383	876	229	277	1		

- Molecule 4 is a protein called ATP synthase subunit b', chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	p	143	Total	C	N	O	S	0	0
			1124	713	182	224	5		

- Molecule 5 is a protein called ATP synthase epsilon chain, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	e	131	Total	C	N	O	S	0	0
			1010	622	190	195	3		

- Molecule 6 is a protein called ATP synthase gamma chain, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	g	321	Total	C	N	O	S	0	0
			2497	1574	429	482	12		

- Molecule 7 is a protein called ATP synthase subunit c, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	S	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	N	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	O	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	P	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	Q	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	R	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	M	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	T	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	G	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	H	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	L	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	K	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	J	79	Total	C	N	O	S	0	0
			544	355	90	98	1		
7	I	79	Total	C	N	O	S	0	0
			544	355	90	98	1		

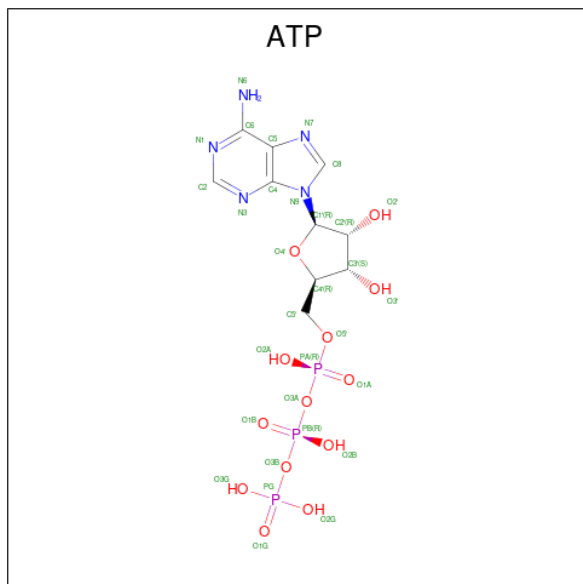
- Molecule 8 is a protein called ATP synthase subunit alpha, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	E	498	Total	C	N	O	S	0	0
			3827	2408	661	745	13		
8	A	501	Total	C	N	O	S	0	0
			3849	2422	666	748	13		
8	C	501	Total	C	N	O	S	0	0
			3851	2423	667	748	13		

- Molecule 9 is a protein called ATP synthase subunit beta, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	F	479	Total	C	N	O	S	0	0
			3627	2288	623	700	16		
9	B	478	Total	C	N	O	S	0	0
			3619	2282	622	699	16		
9	D	480	Total	C	N	O	S	0	0
			3636	2294	625	701	16		

- Molecule 10 is ADENOSINE-5'-TRIPHOSPHATE (CCD ID: ATP) (formula:  $C_{10}H_{16}N_5O_{13}P_3$ ).



Mol	Chain	Residues	Atoms					AltConf
10	E	1	Total	C	N	O	P	0
			31	10	5	13	3	
10	A	1	Total	C	N	O	P	0
			31	10	5	13	3	
10	C	1	Total	C	N	O	P	0
			31	10	5	13	3	

- Molecule 11 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

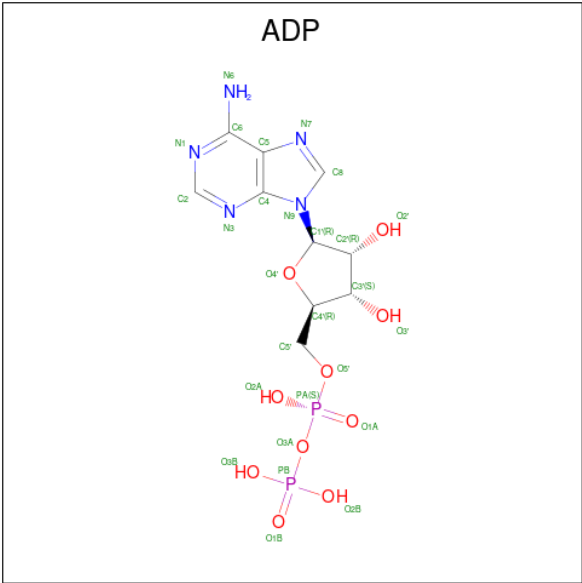
Mol	Chain	Residues	Atoms		AltConf
11	E	1	Total	Mg	0
			1	1	
11	A	1	Total	Mg	0
			1	1	
11	B	1	Total	Mg	0
			1	1	
11	C	1	Total	Mg	0
			1	1	

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Mol	Chain	Residues	Atoms		AltConf
11	D	1	Total	Mg	0
			1	1	

- Molecule 12 is ADENOSINE-5'-DIPHOSPHATE (CCD ID: ADP) (formula: C<sub>10</sub>H<sub>15</sub>N<sub>5</sub>O<sub>10</sub>P<sub>2</sub>).

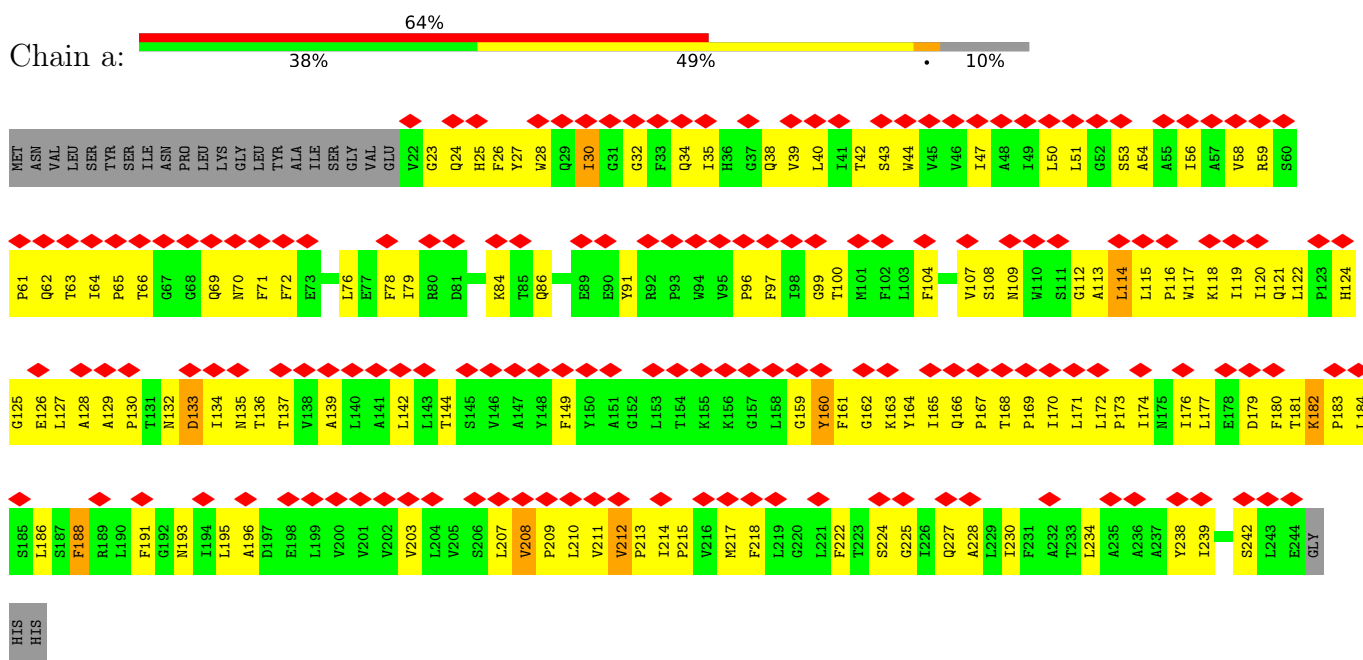


Mol	Chain	Residues	Atoms					AltConf
12	B	1	Total	C	N	O	P	0
			27	10	5	10	2	
12	D	1	Total	C	N	O	P	0
			27	10	5	10	2	

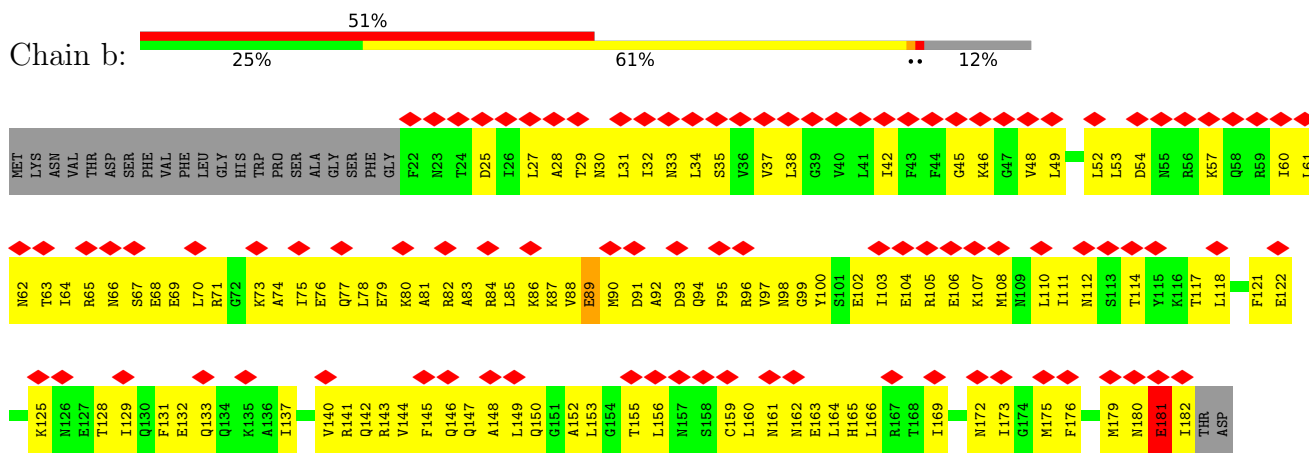
### 3 Residue-property plots

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: ATP synthase subunit a, chloroplastic



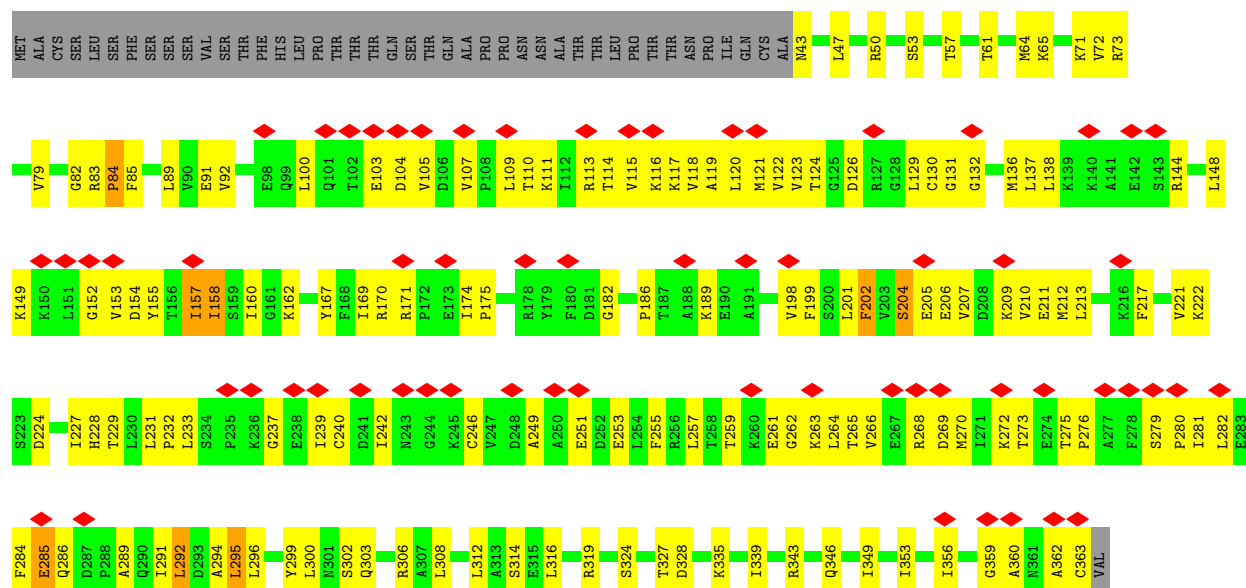
- Molecule 2: ATP synthase subunit b, chloroplastic



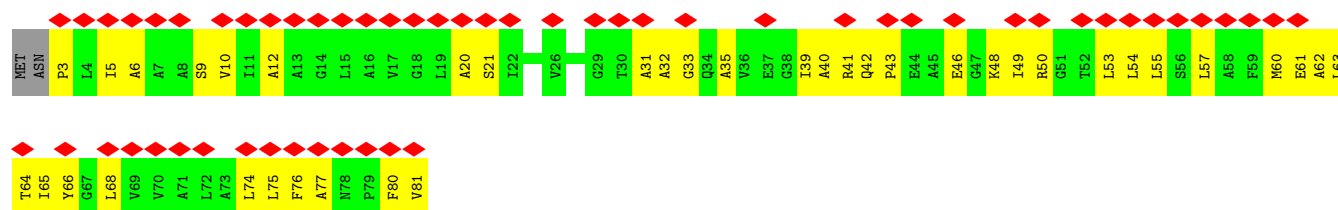
- Molecule 3: ATP synthase delta chain, chloroplastic



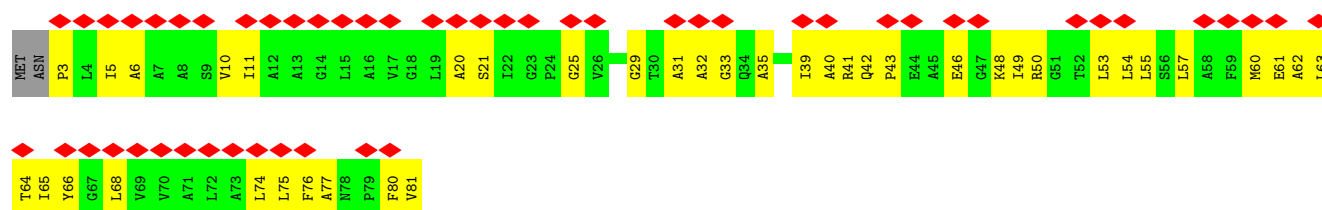




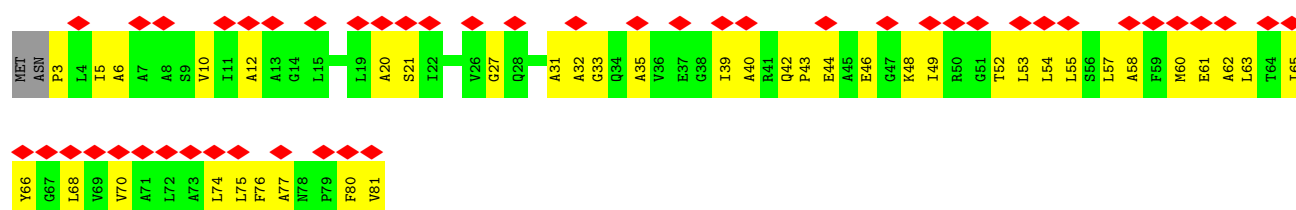
• Molecule 7: ATP synthase subunit c, chloroplastic



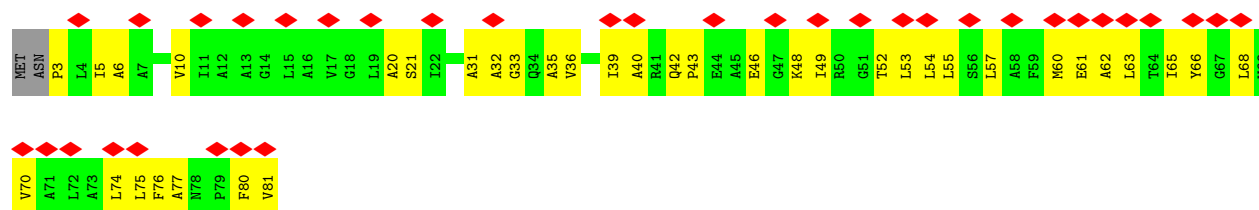

• Molecule 7: ATP synthase subunit c, chloroplastic



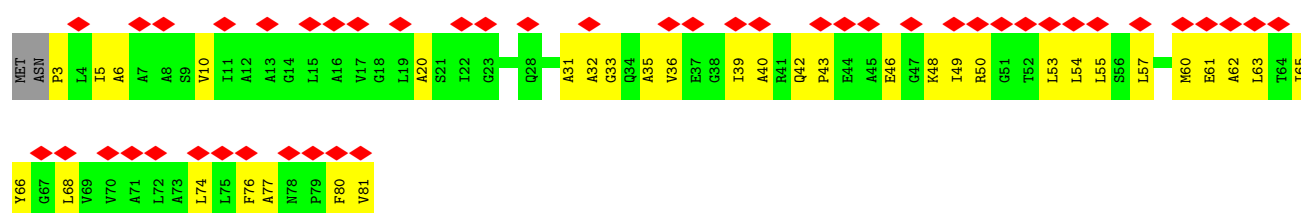
• Molecule 7: ATP synthase subunit c, chloroplastic



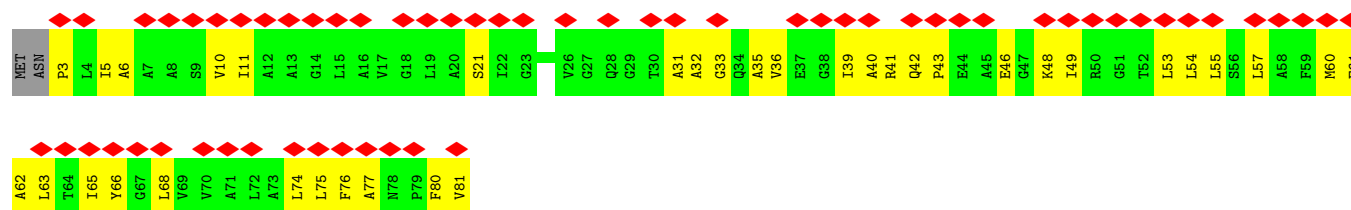

## • Molecule 7: ATP synthase subunit c, chloroplastic

Chain P: 

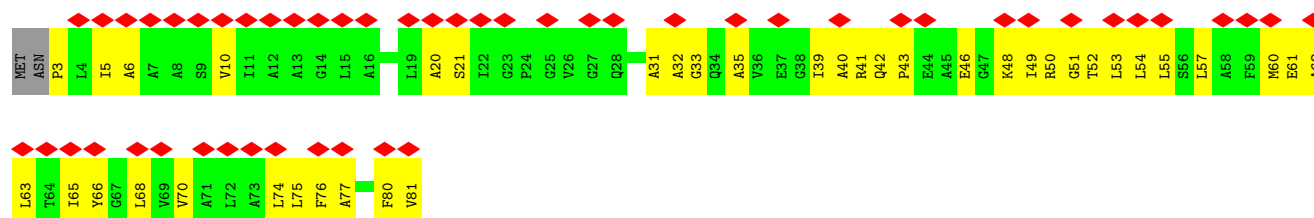
## • Molecule 7: ATP synthase subunit c, chloroplastic

Chain Q: 

## • Molecule 7: ATP synthase subunit c, chloroplastic

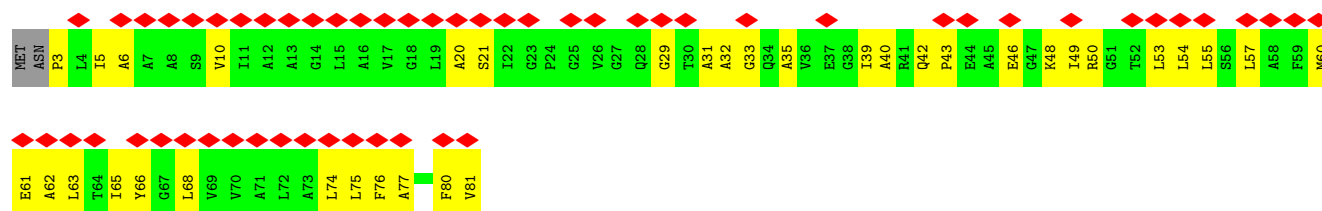
Chain R: 

## • Molecule 7: ATP synthase subunit c, chloroplastic

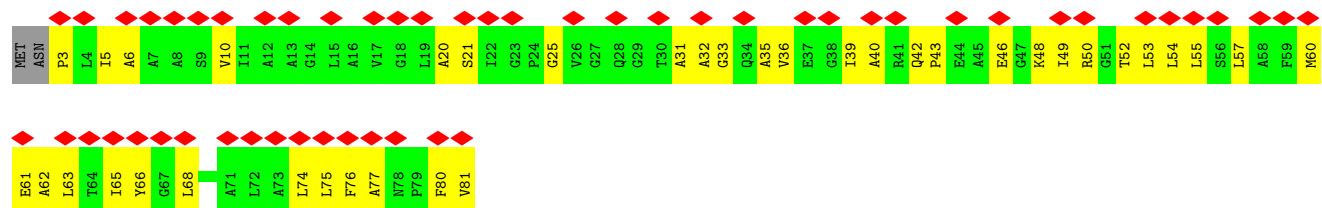
Chain M: 

## • Molecule 7: ATP synthase subunit c, chloroplastic

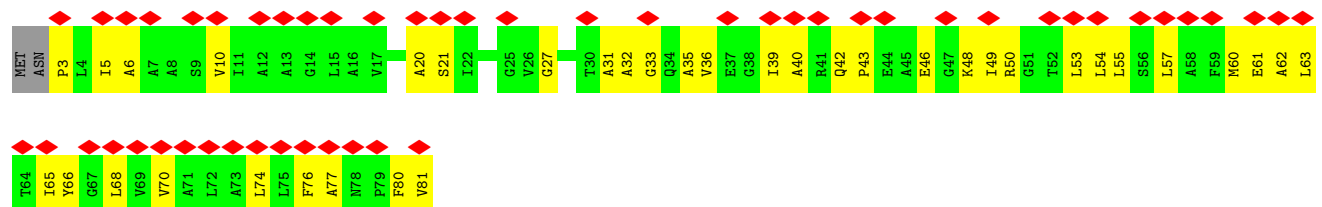
Chain T: 



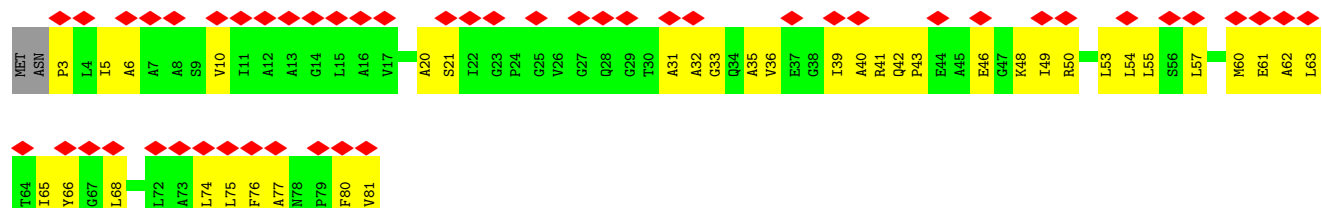
- Molecule 7: ATP synthase subunit c, chloroplastic



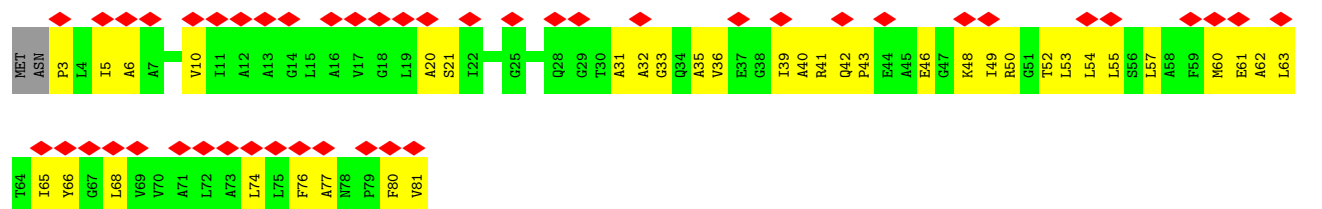
- Molecule 7: ATP synthase subunit c, chloroplastic



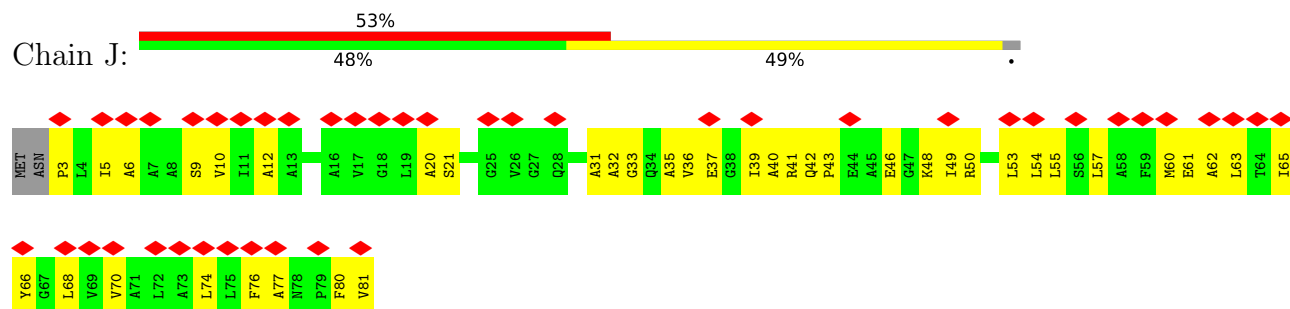
- Molecule 7: ATP synthase subunit c, chloroplastic



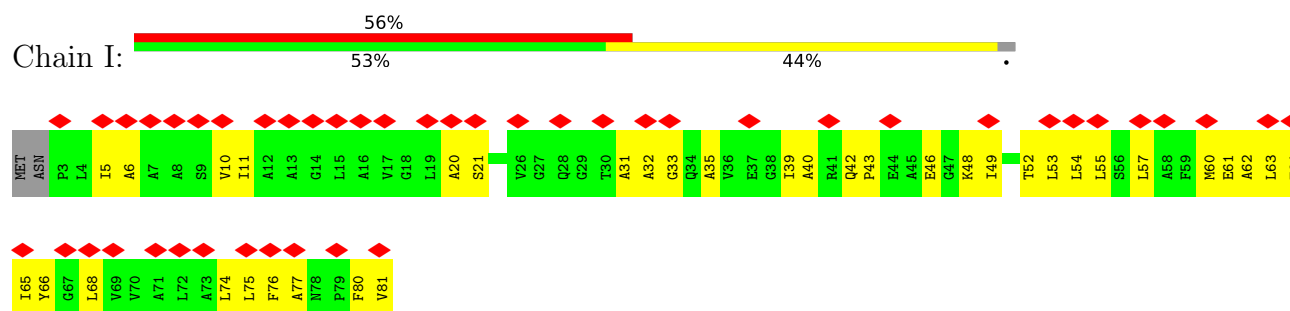
- Molecule 7: ATP synthase subunit c, chloroplastic



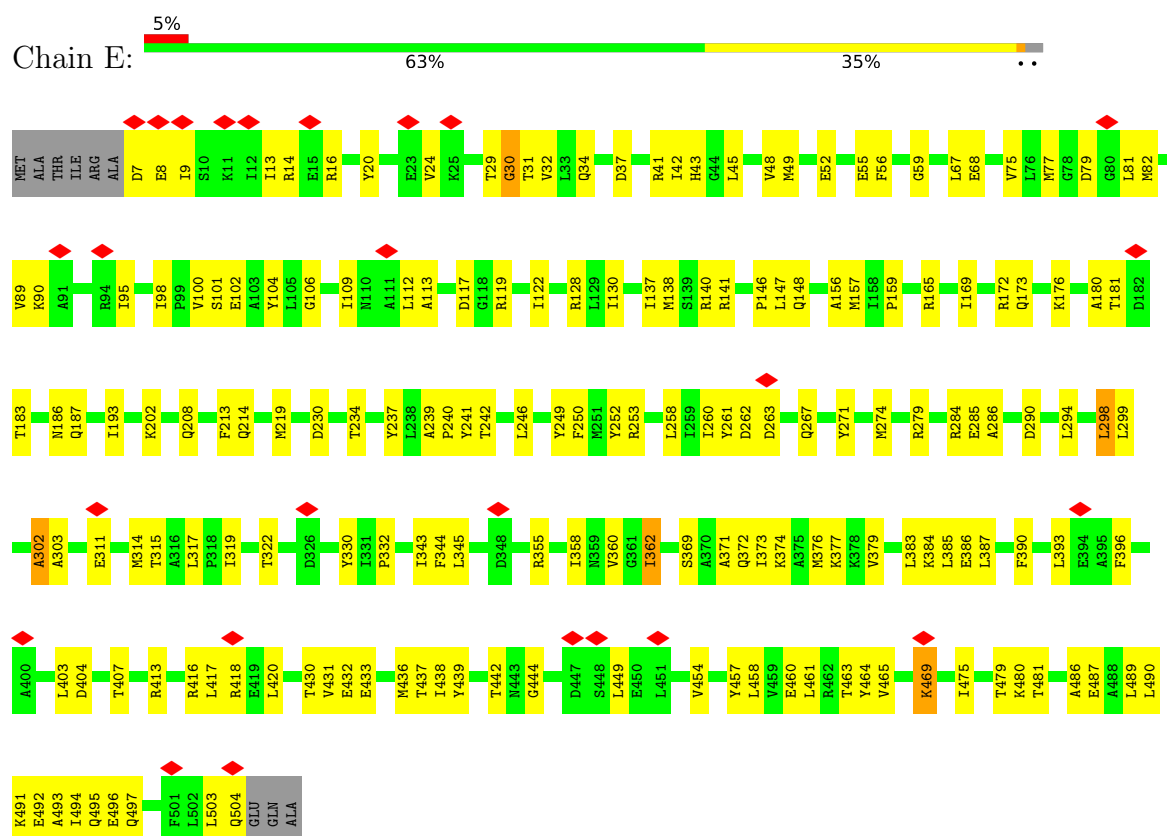
- Molecule 7: ATP synthase subunit c, chloroplastic



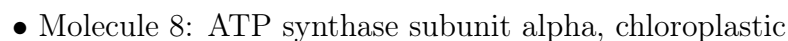
- Molecule 7: ATP synthase subunit c, chloroplastic



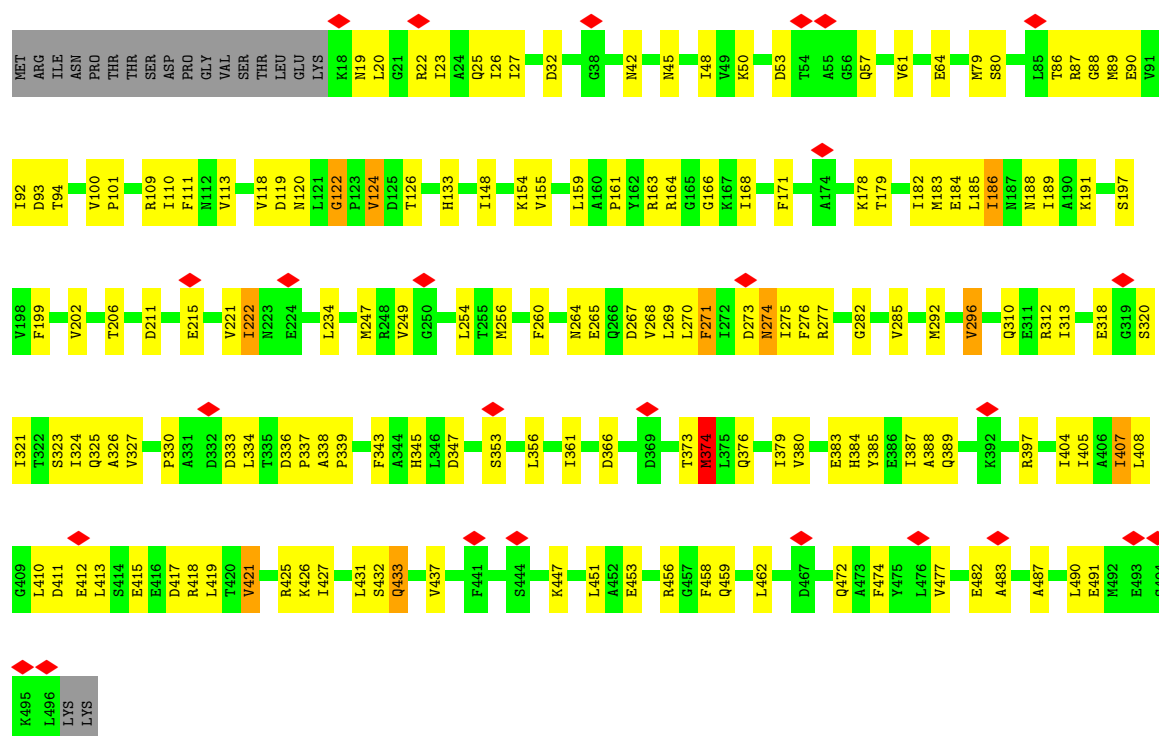
- Molecule 8: ATP synthase subunit alpha, chloroplastic



- Molecule 8: ATP synthase subunit alpha, chloroplastic

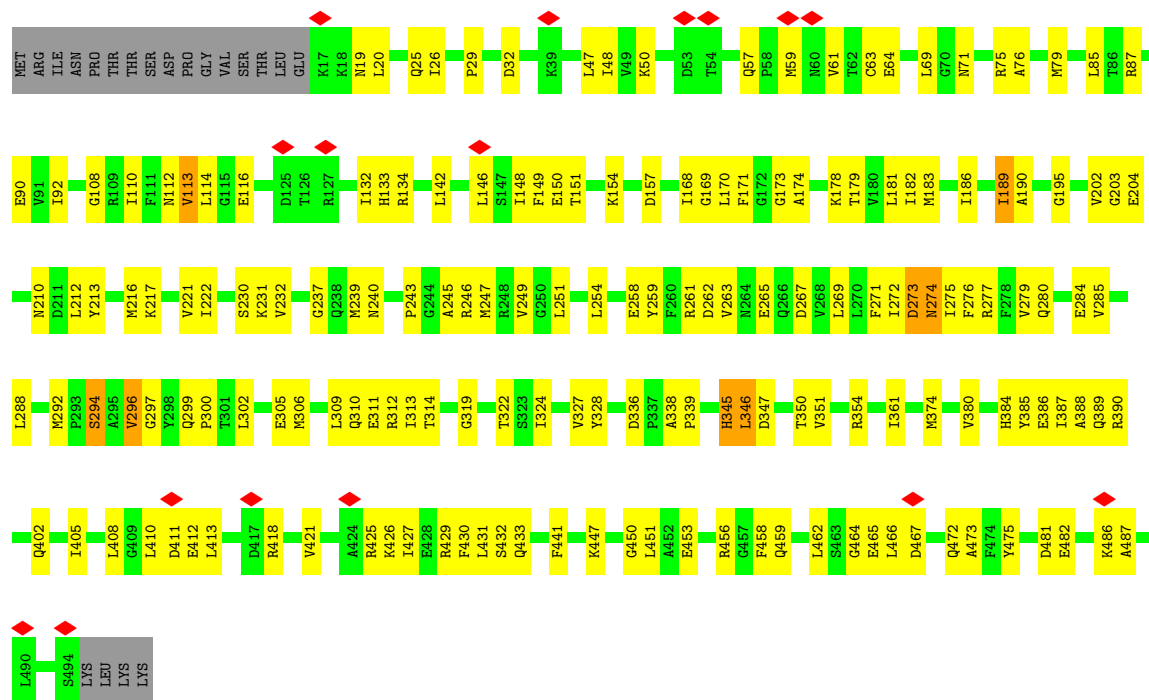


Chain F: 

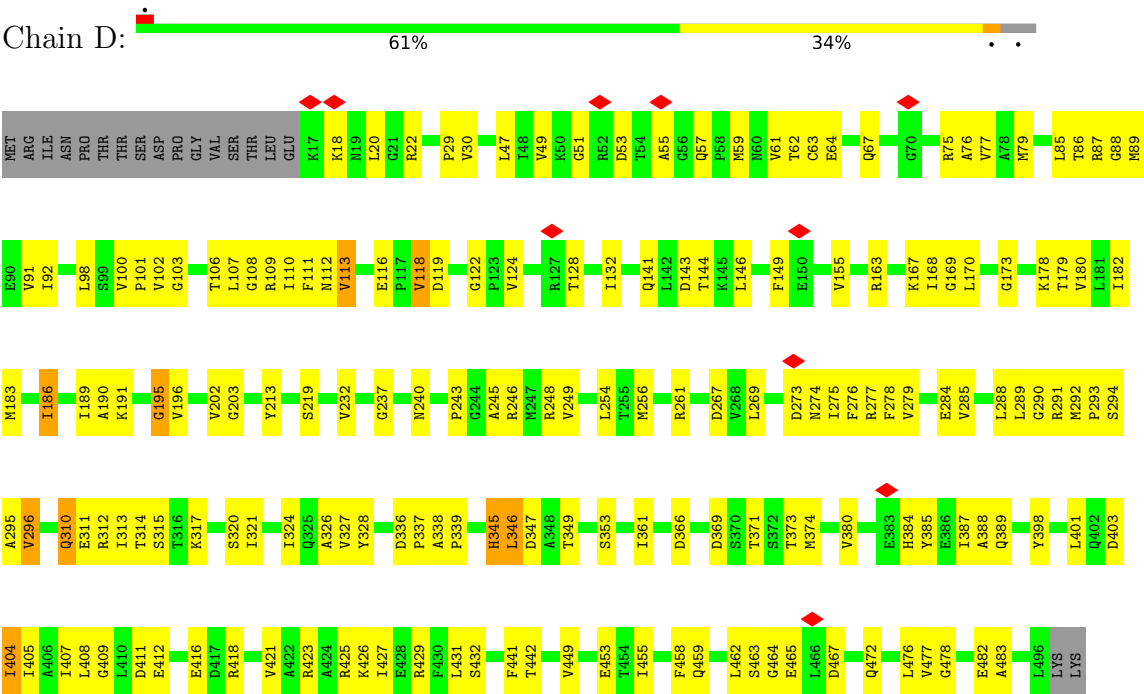


- Molecule 9: ATP synthase subunit beta, chloroplastic

Chain B: 



- Molecule 9: ATP synthase subunit beta, chloroplastic





## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	14409	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING ONLY	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	25	Depositor
Minimum defocus (nm)	1500	Depositor
Maximum defocus (nm)	2500	Depositor
Magnification	132953	Depositor
Image detector	FEI FALCON III (4k x 4k)	Depositor
Maximum map value	0.225	Depositor
Minimum map value	-0.142	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.013	Depositor
Recommended contour level	0.05	Depositor
Map size ( $\text{\AA}$ )	368.55, 368.55, 368.55	wwPDB
Map dimensions	350, 350, 350	wwPDB
Map angles ( $^\circ$ )	90.0, 90.0, 90.0	wwPDB
Pixel spacing ( $\text{\AA}$ )	1.053, 1.053, 1.053	Depositor

## 5 Model quality

### 5.1 Standard geometry

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

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	0.53	0/1792	1.14	19/2454 (0.8%)
2	b	0.44	0/1309	0.89	2/1754 (0.1%)
3	d	0.63	0/1399	1.00	7/1898 (0.4%)
4	p	0.62	1/1134 (0.1%)	0.98	6/1519 (0.4%)
5	e	0.47	0/1019	0.82	0/1381
6	g	0.76	1/2526 (0.0%)	0.94	8/3412 (0.2%)
7	G	0.59	0/551	1.00	2/750 (0.3%)
7	H	0.59	0/551	0.99	2/750 (0.3%)
7	I	0.58	0/551	1.00	2/750 (0.3%)
7	J	0.58	0/551	1.00	2/750 (0.3%)
7	K	0.58	0/551	1.00	2/750 (0.3%)
7	L	0.58	0/551	0.99	2/750 (0.3%)
7	M	0.58	0/551	0.99	2/750 (0.3%)
7	N	0.58	0/551	1.00	2/750 (0.3%)
7	O	0.58	0/551	1.00	2/750 (0.3%)
7	P	0.58	0/551	1.00	2/750 (0.3%)
7	Q	0.58	0/551	1.00	2/750 (0.3%)
7	R	0.58	0/551	1.00	2/750 (0.3%)
7	S	0.59	0/551	0.99	2/750 (0.3%)
7	T	0.58	0/551	0.99	2/750 (0.3%)
8	A	0.90	1/3898 (0.0%)	0.94	10/5274 (0.2%)
8	C	0.94	2/3900 (0.1%)	1.00	12/5276 (0.2%)
8	E	0.86	0/3876	0.93	6/5244 (0.1%)
9	B	0.79	1/3676 (0.0%)	0.92	11/4983 (0.2%)
9	D	0.82	0/3693	0.86	8/5005 (0.2%)
9	F	1.00	2/3684 (0.1%)	0.94	18/4994 (0.4%)
All	All	0.78	8/39620 (0.0%)	0.96	135/53694 (0.3%)

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
2	b	0	1
4	p	0	1
7	G	0	1
7	H	0	1
7	I	0	1
7	J	0	1
7	K	0	1
7	L	0	1
7	M	0	1
7	N	0	1
7	O	0	1
7	P	0	1
7	Q	0	1
7	R	0	1
7	S	0	1
7	T	0	1
8	A	0	2
8	C	0	3
8	E	0	2
9	D	0	1
All	All	0	24

All (8) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
9	F	276	PHE	C-N	-27.86	0.97	1.33
8	A	30	GLY	C-N	14.79	1.52	1.33
6	g	84	PRO	N-CD	12.76	1.65	1.47
4	p	105	ASP	C-N	-11.84	1.17	1.33
9	B	271	PHE	C-N	7.57	1.43	1.33
9	F	271	PHE	C-N	-7.27	1.23	1.33
8	C	30	GLY	C-N	6.36	1.42	1.33
8	C	296	SER	C-O	-5.15	1.18	1.24

All (135) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	a	212	VAL	CA-C-N	13.34	133.74	119.87
1	a	212	VAL	C-N-CA	13.34	133.74	119.87
8	C	30	GLY	CA-C-N	13.07	140.12	121.42
8	C	30	GLY	C-N-CA	13.07	140.12	121.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	b	181	GLU	O-C-N	-11.92	109.79	122.07
8	C	30	GLY	O-C-N	-10.54	109.00	122.70
8	A	30	GLY	O-C-N	10.49	133.32	123.57
1	a	208	VAL	CA-C-N	10.38	130.18	120.21
1	a	208	VAL	C-N-CA	10.38	130.18	120.21
9	B	189	ILE	N-CA-C	9.97	120.79	110.72
8	E	30	GLY	O-C-N	-9.87	109.87	122.70
1	a	30	ILE	N-CA-C	9.29	121.12	108.11
3	d	216	SER	N-CA-C	9.04	121.13	111.28
9	B	294	SER	N-CA-C	8.56	120.39	111.14
4	p	109	TYR	N-CA-C	8.49	120.16	111.07
9	B	190	ALA	N-CA-C	8.41	120.07	111.07
3	d	117	ASN	CA-C-N	8.21	128.16	119.05
3	d	117	ASN	C-N-CA	8.21	128.16	119.05
1	a	59	ARG	N-CA-C	7.94	119.94	111.28
9	F	191	LYS	N-CA-C	7.88	119.65	111.14
8	E	469	LYS	CA-C-N	7.79	127.85	119.28
8	E	469	LYS	C-N-CA	7.79	127.85	119.28
8	A	467	THR	N-CA-C	7.77	119.53	111.14
8	E	59	GLY	N-CA-C	7.73	125.65	115.36
9	F	271	PHE	O-C-N	-7.55	114.36	123.27
9	D	345	HIS	N-CA-C	7.50	119.46	111.28
8	A	21	ASN	N-CA-C	7.43	119.46	111.36
1	a	212	VAL	CB-CA-C	-7.29	106.63	114.35
1	a	188	PHE	N-CA-C	7.25	119.27	111.36
6	g	162	LYS	N-CA-C	7.24	119.25	111.36
3	d	116	ALA	N-CA-C	7.24	119.25	111.36
1	a	211	VAL	N-CA-C	7.22	117.98	110.62
9	B	273	ASP	CA-CB-CG	7.10	119.70	112.60
9	F	122	GLY	CA-C-N	7.06	127.03	119.76
9	F	122	GLY	C-N-CA	7.06	127.03	119.76
9	B	345	HIS	N-CA-C	7.03	119.02	111.36
6	g	285	GLU	N-CA-C	6.91	118.81	111.28
8	A	190	GLN	N-CA-C	6.70	118.94	110.24
8	A	371	ALA	N-CA-C	6.62	118.58	111.36
1	a	129	ALA	CA-C-N	6.56	126.00	118.85
1	a	129	ALA	C-N-CA	6.56	126.00	118.85
8	C	379	VAL	N-CA-C	6.55	116.69	110.53
9	F	285	VAL	N-CA-CB	6.53	120.38	110.58
8	C	113	ALA	N-CA-C	6.51	120.52	112.58
8	C	231	SER	CA-C-N	6.50	126.47	119.78
8	C	231	SER	C-N-CA	6.50	126.47	119.78

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
9	B	113	VAL	CB-CA-C	-6.50	103.37	112.14
1	a	210	LEU	N-CA-C	6.48	119.16	111.71
4	p	200	ILE	N-CA-C	-6.46	104.03	110.62
6	g	158	ILE	N-CA-C	-6.46	98.56	107.99
3	d	119	VAL	N-CA-C	6.45	117.20	110.62
3	d	214	ASP	CA-C-N	6.45	125.88	118.85
3	d	214	ASP	C-N-CA	6.45	125.88	118.85
7	P	76	PHE	CA-C-N	-6.43	113.34	122.41
7	P	76	PHE	C-N-CA	-6.43	113.34	122.41
7	G	76	PHE	CA-C-N	-6.40	113.38	122.41
7	G	76	PHE	C-N-CA	-6.40	113.38	122.41
7	M	76	PHE	CA-C-N	-6.40	113.39	122.41
7	M	76	PHE	C-N-CA	-6.40	113.39	122.41
7	I	76	PHE	CA-C-N	-6.40	113.39	122.41
7	I	76	PHE	C-N-CA	-6.40	113.39	122.41
7	R	76	PHE	CA-C-N	-6.39	113.40	122.41
7	R	76	PHE	C-N-CA	-6.39	113.40	122.41
7	K	76	PHE	CA-C-N	-6.39	113.40	122.41
7	K	76	PHE	C-N-CA	-6.39	113.40	122.41
7	O	76	PHE	CA-C-N	-6.37	113.42	122.41
7	O	76	PHE	C-N-CA	-6.37	113.42	122.41
7	L	76	PHE	CA-C-N	-6.37	113.43	122.41
7	L	76	PHE	C-N-CA	-6.37	113.43	122.41
7	N	76	PHE	CA-C-N	-6.37	113.43	122.41
7	N	76	PHE	C-N-CA	-6.37	113.43	122.41
7	H	76	PHE	CA-C-N	-6.36	113.44	122.41
7	H	76	PHE	C-N-CA	-6.36	113.44	122.41
7	Q	76	PHE	CA-C-N	-6.36	113.45	122.41
7	Q	76	PHE	C-N-CA	-6.36	113.45	122.41
7	T	76	PHE	CA-C-N	-6.35	113.46	122.41
7	T	76	PHE	C-N-CA	-6.35	113.46	122.41
7	J	76	PHE	CA-C-N	-6.35	113.46	122.41
7	J	76	PHE	C-N-CA	-6.35	113.46	122.41
7	S	76	PHE	CA-C-N	-6.33	113.48	122.41
7	S	76	PHE	C-N-CA	-6.33	113.48	122.41
9	D	113	VAL	CB-CA-C	-6.26	103.69	112.14
6	g	204	SER	N-CA-C	-6.26	101.68	110.50
9	D	346	LEU	N-CA-C	6.25	119.54	110.28
2	b	89	GLU	N-CA-C	-6.24	103.76	111.75
9	B	262	ASP	N-CA-C	6.21	118.12	111.36
9	B	346	LEU	N-CA-C	6.03	119.21	110.28
9	F	379	ILE	N-CA-C	6.02	117.05	111.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
8	A	11	LYS	N-CA-C	-5.99	105.63	113.12
9	F	124	VAL	N-CA-C	-5.94	99.75	108.54
9	D	186	ILE	CB-CA-C	-5.92	104.40	111.97
8	C	376	MET	N-CA-C	5.88	117.77	111.36
1	a	181	THR	N-CA-C	5.86	117.67	111.28
9	F	374	MET	N-CA-C	5.86	120.29	113.20
4	p	105	ASP	CA-C-N	-5.85	109.17	121.80
4	p	105	ASP	C-N-CA	-5.85	109.17	121.80
1	a	114	LEU	CA-C-N	-5.84	113.49	121.61
1	a	114	LEU	C-N-CA	-5.84	113.49	121.61
9	F	379	ILE	CB-CA-C	-5.80	104.32	111.92
6	g	263	LYS	N-CA-C	5.78	118.17	110.53
4	p	106	LYS	CA-C-N	-5.76	113.30	120.56
4	p	106	LYS	C-N-CA	-5.76	113.30	120.56
9	B	263	VAL	N-CA-C	5.74	116.75	111.81
1	a	160	TYR	N-CA-C	5.72	117.97	108.99
8	A	449	LEU	N-CA-C	5.69	118.09	110.35
9	D	118	VAL	N-CA-C	5.67	117.56	112.17
9	D	310	GLN	N-CA-C	5.66	117.53	111.36
8	E	362	ILE	N-CA-C	5.65	116.43	110.72
9	B	271	PHE	O-C-N	-5.57	116.44	123.01
8	C	179	VAL	CB-CA-C	-5.52	104.90	111.97
9	D	195	GLY	N-CA-C	5.46	120.21	111.64
9	F	271	PHE	CA-C-N	5.43	130.49	123.11
9	F	271	PHE	C-N-CA	5.43	130.49	123.11
9	F	274	ASN	CA-CB-CG	5.43	118.03	112.60
8	A	30	GLY	CA-C-N	-5.42	115.52	123.00
8	A	30	GLY	C-N-CA	-5.42	115.52	123.00
9	F	171	PHE	CA-C-N	-5.41	117.45	121.61
9	F	171	PHE	C-N-CA	-5.41	117.45	121.61
9	F	407	ILE	N-CA-C	5.40	116.18	110.72
8	E	302	ALA	N-CA-C	5.40	117.10	108.41
9	F	421	VAL	N-CA-C	-5.40	105.86	111.58
6	g	295	LEU	N-CA-C	5.35	117.11	111.28
9	F	186	ILE	CB-CA-C	-5.32	104.97	112.14
6	g	202	PHE	N-CA-C	5.29	117.13	111.36
8	C	298	LEU	N-CA-C	5.27	117.03	111.28
6	g	157	ILE	N-CA-C	5.22	115.48	108.17
1	a	182	LYS	CA-C-N	-5.19	113.35	119.84
1	a	182	LYS	C-N-CA	-5.19	113.35	119.84
8	C	108	VAL	N-CA-C	-5.16	102.08	108.89
8	C	15	GLU	N-CA-C	-5.15	105.66	111.28

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
9	D	404	ILE	N-CA-C	5.14	115.91	110.72
9	B	274	ASN	OD1-CG-ND2	5.11	127.71	122.60
1	a	133	ASP	N-CA-C	-5.07	102.78	110.28
9	F	222	ILE	N-CA-C	-5.06	101.09	108.17
8	A	130	ILE	N-CA-C	5.03	115.80	110.72

There are no chirality outliers.

All (24) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
8	A	226	ALA	Peptide
8	A	262	ASP	Peptide
8	C	10	SER	Peptide
8	C	262	ASP	Peptide
8	C	30	GLY	Mainchain
9	D	273	ASP	Peptide
8	E	262	ASP	Peptide
8	E	30	GLY	Mainchain
7	G	77	ALA	Mainchain
7	H	77	ALA	Mainchain
7	I	77	ALA	Mainchain
7	J	77	ALA	Mainchain
7	K	77	ALA	Mainchain
7	L	77	ALA	Mainchain
7	M	77	ALA	Mainchain
7	N	77	ALA	Mainchain
7	O	77	ALA	Mainchain
7	P	77	ALA	Mainchain
7	Q	77	ALA	Mainchain
7	R	77	ALA	Mainchain
7	S	77	ALA	Mainchain
7	T	77	ALA	Mainchain
2	b	181	GLU	Mainchain
4	p	105	ASP	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	1741	0	1781	606	0
2	b	1299	0	1310	912	0
3	d	1383	0	1406	357	0
4	p	1124	0	1145	922	0
5	e	1010	0	1040	150	0
6	g	2497	0	2585	398	0
7	G	544	0	581	113	0
7	H	544	0	581	111	0
7	I	544	0	581	110	0
7	J	544	0	581	112	0
7	K	544	0	581	105	0
7	L	544	0	581	118	0
7	M	544	0	581	121	0
7	N	544	0	581	120	0
7	O	544	0	581	119	0
7	P	544	0	581	106	0
7	Q	544	0	581	113	0
7	R	544	0	581	115	0
7	S	544	0	581	118	0
7	T	544	0	581	114	0
8	A	3849	0	3922	569	0
8	C	3851	0	3928	332	0
8	E	3827	0	3900	301	0
9	B	3619	0	3668	212	0
9	D	3636	0	3692	294	0
9	F	3627	0	3679	193	0
10	A	31	0	12	1	0
10	C	31	0	12	1	0
10	E	31	0	12	1	0
11	A	1	0	0	0	0
11	B	1	0	0	0	0
11	C	1	0	0	0	0
11	D	1	0	0	0	0
11	E	1	0	0	0	0
12	B	27	0	12	0	0
12	D	27	0	12	2	0
All	All	39231	0	40250	4676	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 59.

All (4676) 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)
1:a:51:LEU:CD2	4:p:100:LEU:CD1	1.75	1.64
8:A:13:ILE:CD1	8:A:16:ARG:HH22	1.09	1.63
1:a:161:PHE:HE1	1:a:163:LYS:CG	1.07	1.61
1:a:65:PRO:CB	1:a:70:ASN:HD21	0.96	1.61
1:a:24:GLN:HB3	4:p:84:PHE:CD2	1.32	1.60
1:a:161:PHE:CE1	1:a:163:LYS:CG	1.83	1.59
1:a:161:PHE:CZ	1:a:163:LYS:HD2	1.29	1.59
2:b:144:VAL:CG1	4:p:210:LEU:CD2	1.76	1.59
2:b:160:LEU:HD11	2:b:164:LEU:CA	1.13	1.59
2:b:144:VAL:CG1	4:p:210:LEU:HD23	1.17	1.59
1:a:119:ILE:CG2	2:b:28:ALA:HA	1.18	1.57
3:d:222:THR:CG2	3:d:234:ASP:HB2	1.23	1.57
1:a:119:ILE:HG23	2:b:28:ALA:CA	1.23	1.55
4:p:214:ILE:HA	8:A:17:ILE:CD1	1.13	1.55
5:e:21:GLU:HG2	5:e:34:LEU:CD2	1.35	1.55
1:a:161:PHE:CE1	1:a:163:LYS:HG3	1.41	1.55
4:p:195:GLN:CG	8:A:4:ILE:CD1	1.83	1.54
2:b:68:GLU:CG	4:p:126:GLN:HB3	1.14	1.54
2:b:141:ARG:CD	8:A:12:ILE:HG21	1.37	1.53
2:b:142:GLN:CD	3:d:248:LEU:HD23	1.25	1.52
1:a:134:ILE:HG23	1:a:191:PHE:CZ	1.42	1.52
2:b:133:GLN:CG	8:A:3:THR:HG22	1.39	1.51
2:b:165:HIS:CD2	3:d:219:ALA:HB1	1.45	1.51
2:b:88:VAL:CA	4:p:148:MET:HG3	1.05	1.51
8:C:77:MET:CE	8:C:112:LEU:HD21	1.37	1.51
2:b:103:ILE:CB	4:p:159:LEU:HA	1.14	1.50
2:b:82:ARG:HB3	4:p:140:LEU:CD2	1.01	1.49
2:b:103:ILE:HB	4:p:159:LEU:CA	1.15	1.49
2:b:176:PHE:CE1	3:d:233:VAL:CG2	1.90	1.48
1:a:115:LEU:CD2	2:b:38:LEU:HD13	1.40	1.48
2:b:88:VAL:CB	4:p:148:MET:HG3	1.38	1.48
2:b:82:ARG:CB	4:p:140:LEU:HD23	0.99	1.47
2:b:107:LYS:HA	4:p:166:THR:CG2	1.13	1.47
2:b:133:GLN:HG2	8:A:3:THR:CG2	1.45	1.47
1:a:135:ASN:HD21	4:p:86:PHE:CA	1.23	1.46
2:b:88:VAL:HA	4:p:148:MET:CG	0.99	1.46
6:g:281:ILE:HA	7:N:41:ARG:NH2	1.27	1.46
2:b:125:LYS:HD2	4:p:185:LEU:CD2	1.42	1.45
2:b:160:LEU:CD1	2:b:163:GLU:C	1.85	1.45
1:a:119:ILE:CD1	2:b:32:ILE:H	1.26	1.45
3:d:154:ARG:HH21	8:C:5:ARG:CZ	1.26	1.45
1:a:139:ALA:HB1	4:p:94:MET:SD	1.57	1.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:107:LYS:CA	4:p:166:THR:HG21	1.26	1.44
2:b:176:PHE:CE1	3:d:233:VAL:HG21	0.93	1.44
2:b:68:GLU:HB2	4:p:126:GLN:CB	1.38	1.44
1:a:124:HIS:NE2	4:p:79:GLU:CB	1.79	1.44
2:b:125:LYS:CD	4:p:185:LEU:CD2	1.91	1.44
1:a:104:PHE:HB2	4:p:97:PHE:CE1	1.52	1.43
1:a:222:PHE:CZ	7:Q:54:LEU:HD23	1.51	1.43
3:d:222:THR:HG22	3:d:234:ASP:CB	0.99	1.43
2:b:70:LEU:HD12	4:p:127:LEU:CD2	1.47	1.43
2:b:70:LEU:CD1	4:p:127:LEU:HD21	1.45	1.43
8:A:464:TYR:CE1	8:A:468:ASN:ND2	1.84	1.43
2:b:141:ARG:HD2	8:A:12:ILE:CG2	1.47	1.43
6:g:239:ILE:HD12	6:g:303:GLN:NE2	1.16	1.42
8:A:13:ILE:HD13	8:A:16:ARG:NH2	1.13	1.42
1:a:40:LEU:CG	4:p:89:THR:OG1	1.66	1.42
2:b:122:GLU:OE1	4:p:181:ILE:CG2	1.66	1.42
1:a:35:ILE:HG22	4:p:88:LEU:N	1.32	1.41
1:a:116:PRO:HD3	2:b:35:SER:N	1.34	1.41
1:a:124:HIS:NE2	4:p:79:GLU:HB3	1.24	1.41
2:b:70:LEU:N	4:p:130:VAL:CG1	1.79	1.41
2:b:77:GLN:CG	4:p:138:LYS:HG3	1.51	1.41
1:a:135:ASN:ND2	4:p:86:PHE:HA	1.12	1.40
6:g:103:GLU:CG	7:R:41:ARG:HE	1.30	1.40
8:E:344:PHE:CD2	8:E:362:ILE:HG21	1.52	1.40
1:a:63:THR:CA	4:p:119:ARG:HH11	1.32	1.40
8:A:214:GLN:NE2	8:A:219:MET:HG2	1.29	1.40
2:b:103:ILE:HD11	4:p:162:MET:N	1.34	1.39
9:B:20:LEU:CD1	9:B:90:GLU:OE2	1.70	1.38
2:b:85:LEU:O	4:p:147:VAL:CG1	1.72	1.38
2:b:144:VAL:HG21	8:A:9:ILE:CG2	1.52	1.38
6:g:113:ARG:NH1	6:g:205:GLU:HB3	1.37	1.38
2:b:68:GLU:CB	4:p:126:GLN:HB3	1.40	1.38
8:E:355:ARG:NH2	9:D:389:GLN:HE22	0.96	1.38
3:d:222:THR:CG2	3:d:234:ASP:CB	1.79	1.37
1:a:161:PHE:CZ	1:a:163:LYS:CD	2.07	1.36
8:A:464:TYR:HE1	8:A:468:ASN:ND2	1.16	1.36
2:b:82:ARG:CG	2:b:85:LEU:HD21	1.54	1.36
2:b:145:PHE:HB2	8:A:16:ARG:NH2	1.35	1.36
2:b:69:GLU:C	4:p:130:VAL:CG1	1.95	1.36
2:b:73:LYS:H	4:p:130:VAL:CG2	1.39	1.36
1:a:24:GLN:HB3	4:p:84:PHE:CE2	1.60	1.35

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:191:LYS:CE	9:D:219:SER:HB2	1.56	1.35
1:a:104:PHE:HB2	4:p:97:PHE:CZ	1.60	1.35
1:a:165:ILE:CG2	1:a:168:THR:O	1.75	1.35
1:a:208:VAL:CG2	1:a:209:PRO:HD2	1.55	1.35
1:a:119:ILE:HA	2:b:29:THR:N	1.38	1.34
2:b:70:LEU:CA	4:p:130:VAL:HG13	1.56	1.34
1:a:65:PRO:HB2	1:a:70:ASN:ND2	1.03	1.34
6:g:359:GLY:C	9:D:293:PRO:HD3	1.51	1.34
8:E:344:PHE:CD2	8:E:362:ILE:CG2	2.10	1.34
2:b:69:GLU:C	4:p:130:VAL:HG11	1.53	1.34
2:b:85:LEU:O	4:p:147:VAL:HG11	1.21	1.34
1:a:35:ILE:CG2	4:p:88:LEU:H	1.41	1.33
2:b:70:LEU:C	4:p:130:VAL:HG13	1.50	1.33
4:p:210:LEU:HD21	8:A:13:ILE:CG1	1.58	1.33
1:a:119:ILE:HG13	2:b:28:ALA:C	1.53	1.33
6:g:240:CYS:HB3	6:g:246:CYS:SG	1.69	1.33
6:g:261:GLU:HG3	9:F:397:ARG:NH2	1.41	1.33
2:b:103:ILE:HB	4:p:159:LEU:CB	1.56	1.32
9:F:50:LYS:HE2	9:F:92:ILE:CD1	1.59	1.32
1:a:51:LEU:CD2	4:p:100:LEU:HD12	1.40	1.32
4:p:214:ILE:CA	8:A:17:ILE:CD1	2.03	1.32
2:b:85:LEU:HD22	4:p:143:GLN:CG	1.36	1.32
2:b:125:LYS:CD	4:p:185:LEU:HD23	1.52	1.32
1:a:164:TYR:HD2	1:a:166:GLN:CG	1.40	1.31
5:e:34:LEU:HB2	7:L:42:GLN:NE2	1.46	1.31
5:e:119:LEU:HD11	5:e:123:ARG:NE	1.45	1.31
2:b:160:LEU:CG	2:b:164:LEU:H	1.43	1.31
6:g:240:CYS:CB	6:g:246:CYS:SG	2.18	1.31
2:b:133:GLN:CG	8:A:3:THR:CG2	2.01	1.30
1:a:43:SER:CB	4:p:93:ILE:HG12	1.60	1.30
1:a:230:ILE:HD11	7:Q:55:LEU:CD1	1.62	1.30
1:a:115:LEU:HD21	2:b:38:LEU:CD1	1.61	1.30
3:d:224:ARG:HH11	3:d:232:LEU:CD1	1.44	1.29
2:b:77:GLN:HB3	4:p:134:SER:O	1.22	1.29
3:d:224:ARG:NH1	3:d:232:LEU:CD1	1.95	1.29
1:a:119:ILE:CD1	2:b:32:ILE:N	1.93	1.29
1:a:134:ILE:CG2	1:a:191:PHE:CZ	2.14	1.29
1:a:161:PHE:CE1	1:a:163:LYS:CB	2.15	1.29
4:p:195:GLN:CG	8:A:4:ILE:HD11	1.47	1.28
8:A:214:GLN:NE2	8:A:219:MET:CG	1.97	1.28
5:e:21:GLU:CG	5:e:34:LEU:CD2	2.11	1.28

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:144:VAL:CG2	8:A:9:ILE:HG21	1.62	1.28
5:e:119:LEU:CD1	5:e:123:ARG:NE	1.95	1.28
8:E:169:ILE:CG2	8:E:344:PHE:CD1	2.16	1.28
1:a:40:LEU:CD2	4:p:89:THR:OG1	1.81	1.28
6:g:360:ALA:N	9:D:293:PRO:CD	1.95	1.28
1:a:134:ILE:CG2	1:a:191:PHE:CE2	2.16	1.27
2:b:82:ARG:O	2:b:85:LEU:HG	1.22	1.27
2:b:141:ARG:O	8:A:13:ILE:HD11	1.30	1.27
8:E:355:ARG:NH2	9:D:389:GLN:NE2	1.80	1.27
1:a:134:ILE:HG22	1:a:191:PHE:CD2	1.68	1.27
2:b:160:LEU:HD13	2:b:163:GLU:CA	1.63	1.27
3:d:228:GLU:OE1	8:A:43:HIS:NE2	1.65	1.27
2:b:88:VAL:HG22	4:p:148:MET:SD	1.73	1.27
2:b:103:ILE:HD13	4:p:158:ALA:O	1.33	1.27
2:b:160:LEU:HD13	2:b:163:GLU:C	1.53	1.27
1:a:119:ILE:CD1	2:b:32:ILE:HG12	1.65	1.26
1:a:51:LEU:CG	4:p:100:LEU:HD13	1.63	1.26
3:d:224:ARG:NH1	3:d:232:LEU:HD12	1.49	1.26
8:C:464:TYR:OH	8:C:496:GLU:OE1	1.53	1.26
1:a:51:LEU:CG	4:p:100:LEU:CD1	2.14	1.26
1:a:164:TYR:CD2	1:a:166:GLN:HG3	1.69	1.26
2:b:71:ARG:NH1	4:p:126:GLN:OE1	1.66	1.26
2:b:142:GLN:CD	3:d:248:LEU:CD2	2.09	1.26
4:p:212:ASP:O	4:p:215:VAL:HG12	1.17	1.25
8:E:37:ASP:OD2	9:D:291:ARG:NE	1.67	1.25
6:g:103:GLU:CG	7:R:41:ARG:NE	1.98	1.25
8:A:111:ALA:HB2	8:A:227:GLU:OE2	1.29	1.25
1:a:124:HIS:NE2	4:p:79:GLU:CA	1.87	1.25
6:g:261:GLU:CG	9:F:397:ARG:NH2	1.99	1.25
1:a:100:THR:HG21	4:p:101:MET:CG	1.68	1.24
5:e:34:LEU:HD12	7:K:43:PRO:CG	1.66	1.24
2:b:76:GLU:O	2:b:80:LYS:HG2	1.31	1.24
3:d:148:ILE:CD1	8:C:17:ILE:HA	1.65	1.24
1:a:40:LEU:HG	4:p:89:THR:OG1	1.07	1.24
6:g:359:GLY:CA	9:D:293:PRO:HG3	1.68	1.24
1:a:122:LEU:CD1	1:a:127:LEU:HD11	1.67	1.23
2:b:129:ILE:HG23	4:p:192:LEU:CD1	1.69	1.23
2:b:71:ARG:H	4:p:130:VAL:CG1	1.37	1.23
2:b:80:LYS:CB	4:p:137:VAL:HB	1.69	1.23
8:A:111:ALA:CB	8:A:227:GLU:OE2	1.87	1.23
2:b:103:ILE:CD1	4:p:162:MET:N	2.00	1.23

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:160:LEU:HG	2:b:164:LEU:CB	1.67	1.23
4:p:152:ARG:CD	8:A:503:LEU:HB2	1.67	1.23
2:b:68:GLU:CG	4:p:126:GLN:CB	2.11	1.23
6:g:137:LEU:HD11	6:g:213:LEU:CD1	1.69	1.23
6:g:360:ALA:HB1	9:D:292:MET:SD	1.79	1.23
2:b:82:ARG:HG3	2:b:85:LEU:CD2	1.68	1.23
6:g:123:VAL:CG1	6:g:308:LEU:HD23	1.67	1.23
2:b:176:PHE:CD1	3:d:233:VAL:HG21	1.74	1.22
8:A:13:ILE:CD1	8:A:16:ARG:NH2	1.77	1.22
2:b:73:LYS:N	4:p:130:VAL:HG23	1.55	1.22
2:b:144:VAL:HG12	4:p:210:LEU:CD2	1.44	1.22
3:d:233:VAL:CG1	8:A:26:VAL:HA	1.67	1.22
3:d:227:ASN:CB	8:A:46:ASP:OD2	1.87	1.22
4:p:193:GLU:HG3	4:p:196:LYS:CE	1.68	1.22
1:a:26:PHE:HB3	4:p:83:LEU:CA	1.69	1.21
2:b:107:LYS:C	4:p:166:THR:OG1	1.83	1.21
2:b:70:LEU:N	4:p:130:VAL:HG13	1.44	1.21
4:p:195:GLN:HG2	8:A:4:ILE:CD1	1.50	1.21
6:g:123:VAL:CG1	6:g:308:LEU:CD2	2.17	1.21
2:b:77:GLN:CD	4:p:138:LYS:HG3	1.58	1.20
8:E:208:GLN:NE2	9:D:144:THR:HG23	1.52	1.20
1:a:100:THR:CG2	4:p:101:MET:HG3	1.70	1.20
1:a:116:PRO:HD3	2:b:34:LEU:C	1.46	1.20
6:g:239:ILE:CD1	6:g:303:GLN:NE2	2.05	1.20
1:a:118:LYS:HB2	2:b:29:THR:O	1.41	1.20
1:a:164:TYR:CD2	1:a:166:GLN:CG	2.23	1.20
2:b:68:GLU:CB	4:p:126:GLN:CB	1.97	1.20
8:C:9:ILE:CG2	8:C:12:ILE:HD12	1.71	1.20
2:b:142:GLN:OE1	3:d:248:LEU:CD2	1.89	1.19
6:g:113:ARG:HH11	6:g:205:GLU:CB	1.55	1.19
1:a:118:LYS:O	2:b:29:THR:OG1	1.57	1.19
1:a:134:ILE:HD12	1:a:135:ASN:N	1.55	1.19
2:b:71:ARG:N	4:p:130:VAL:CG1	1.79	1.19
2:b:125:LYS:CB	4:p:185:LEU:HD22	1.73	1.19
2:b:160:LEU:HD12	2:b:160:LEU:O	1.42	1.19
2:b:81:ALA:N	4:p:140:LEU:HB2	1.54	1.19
2:b:144:VAL:CG1	8:A:13:ILE:HG13	1.70	1.19
3:d:242:GLU:HG2	8:A:20:TYR:OH	1.43	1.19
7:G:62:ALA:O	7:G:65:ILE:HG22	1.43	1.19
8:A:494:ILE:CG2	8:A:498:MET:HE2	1.70	1.19
2:b:80:LYS:C	4:p:140:LEU:HB2	1.66	1.19

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:152:ARG:HB3	8:A:503:LEU:CD2	1.71	1.19
7:J:62:ALA:O	7:J:65:ILE:HG22	1.43	1.19
7:M:62:ALA:O	7:M:65:ILE:HG22	1.43	1.19
9:B:411:ASP:CA	9:B:418:ARG:HH22	1.55	1.19
2:b:64:ILE:HD13	4:p:123:ILE:HG12	1.24	1.18
7:R:62:ALA:O	7:R:65:ILE:HG22	1.43	1.18
2:b:85:LEU:CD2	4:p:143:GLN:CG	2.09	1.18
2:b:88:VAL:HG21	4:p:145:ASN:CA	1.64	1.18
3:d:154:ARG:NH2	8:C:5:ARG:CZ	2.06	1.18
7:I:62:ALA:O	7:I:65:ILE:HG22	1.43	1.18
8:E:386:GLU:OE1	8:E:413:ARG:HD2	1.43	1.18
2:b:103:ILE:CG1	4:p:162:MET:HB2	1.72	1.18
7:S:62:ALA:O	7:S:65:ILE:HG22	1.43	1.18
8:E:355:ARG:HH22	9:D:389:GLN:NE2	1.37	1.18
4:p:213:ASP:OD2	8:A:14:ARG:HG2	1.42	1.18
2:b:64:ILE:HD11	4:p:120:ASP:HA	1.26	1.18
2:b:71:ARG:N	4:p:130:VAL:HG13	1.38	1.17
7:T:62:ALA:O	7:T:65:ILE:HG22	1.43	1.17
7:K:62:ALA:O	7:K:65:ILE:HG22	1.43	1.17
1:a:51:LEU:HD21	4:p:100:LEU:CD1	1.44	1.17
2:b:84:ARG:CB	4:p:141:GLU:HA	1.74	1.17
2:b:165:HIS:CD2	3:d:219:ALA:CB	2.10	1.17
6:g:359:GLY:C	9:D:293:PRO:CD	2.16	1.17
2:b:88:VAL:HB	4:p:147:VAL:HG12	1.22	1.17
2:b:141:ARG:CD	8:A:12:ILE:CG2	2.11	1.17
4:p:210:LEU:HD21	8:A:13:ILE:CB	1.74	1.17
5:e:21:GLU:CD	5:e:34:LEU:HD21	1.69	1.17
6:g:124:THR:HG22	6:g:137:LEU:CD2	1.75	1.17
7:Q:62:ALA:O	7:Q:65:ILE:HG22	1.43	1.17
7:L:62:ALA:O	7:L:65:ILE:HG22	1.43	1.17
1:a:115:LEU:CD2	2:b:38:LEU:CD1	2.20	1.17
2:b:103:ILE:CD1	4:p:162:MET:H	1.56	1.17
1:a:24:GLN:CB	4:p:84:PHE:CD2	2.28	1.16
6:g:360:ALA:CA	9:D:293:PRO:HD2	1.76	1.16
1:a:134:ILE:HG22	1:a:191:PHE:CG	1.80	1.16
2:b:144:VAL:HG11	4:p:210:LEU:CD2	1.54	1.16
3:d:222:THR:CG2	3:d:234:ASP:HB3	1.69	1.16
4:p:210:LEU:HD11	8:A:13:ILE:CG2	1.76	1.16
7:P:62:ALA:O	7:P:65:ILE:HG22	1.43	1.16
8:A:344:PHE:CD2	8:A:362:ILE:HG22	1.80	1.16
9:D:401:LEU:HD22	9:D:404:ILE:HD11	1.26	1.16

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:148:ILE:HD11	8:C:17:ILE:CA	1.74	1.16
1:a:208:VAL:CG2	1:a:209:PRO:CD	2.25	1.15
2:b:80:LYS:HB2	4:p:137:VAL:HB	1.19	1.15
7:O:62:ALA:O	7:O:65:ILE:HG22	1.43	1.15
7:H:62:ALA:O	7:H:65:ILE:HG22	1.43	1.15
9:F:100:VAL:CG1	9:F:256:MET:HE3	1.76	1.15
2:b:70:LEU:N	4:p:130:VAL:HG11	1.51	1.15
8:E:169:ILE:HG21	8:E:344:PHE:CD1	1.75	1.15
9:B:431:LEU:CD2	9:B:458:PHE:HZ	1.60	1.15
8:C:6:ALA:O	8:C:7:ASP:OD1	1.61	1.15
1:a:136:THR:N	4:p:90:LEU:HD22	1.60	1.15
6:g:73:ARG:HH11	9:F:412:GLU:CD	1.54	1.15
9:F:100:VAL:HG11	9:F:256:MET:HE3	1.27	1.15
1:a:26:PHE:CB	4:p:83:LEU:HA	1.33	1.15
1:a:51:LEU:HD23	4:p:100:LEU:CD1	1.76	1.15
2:b:137:ILE:HG13	8:A:5:ARG:HG2	1.16	1.15
2:b:61:LEU:HD11	2:b:65:ARG:HD2	1.20	1.15
2:b:67:SER:OG	4:p:124:LYS:O	1.64	1.15
8:E:386:GLU:OE1	8:E:413:ARG:CD	1.94	1.15
9:D:431:LEU:CD2	9:D:458:PHE:HZ	1.59	1.15
9:D:387:ILE:CD1	9:D:455:ILE:HG23	1.76	1.14
1:a:163:LYS:CE	1:a:179:ASP:OD2	1.94	1.14
4:p:214:ILE:HA	8:A:17:ILE:HD12	1.16	1.14
7:N:62:ALA:O	7:N:65:ILE:HG22	1.43	1.14
1:a:230:ILE:CD1	7:Q:55:LEU:HD11	1.77	1.14
6:g:123:VAL:HG13	6:g:308:LEU:CD2	1.78	1.14
9:B:20:LEU:HD13	9:B:90:GLU:OE2	1.46	1.14
8:C:9:ILE:HG22	8:C:12:ILE:HD12	1.21	1.14
1:a:51:LEU:HG	4:p:100:LEU:CD1	1.75	1.14
6:g:356:ILE:HG21	8:E:286:ALA:HB2	1.20	1.14
1:a:58:VAL:HG11	4:p:107:ILE:HG22	1.14	1.14
3:d:154:ARG:HH21	8:C:5:ARG:NH1	1.43	1.14
4:p:206:GLN:O	4:p:210:LEU:CB	1.96	1.14
8:E:82:MET:HE2	8:E:82:MET:HA	1.26	1.14
8:E:469:LYS:HD2	8:E:493:ALA:HB2	1.18	1.14
1:a:35:ILE:HG21	4:p:88:LEU:HB2	1.21	1.13
2:b:142:GLN:NE2	3:d:248:LEU:HA	1.61	1.13
1:a:40:LEU:HD21	4:p:89:THR:HA	1.26	1.13
2:b:64:ILE:HD13	4:p:123:ILE:CG1	1.76	1.13
2:b:83:ALA:N	4:p:140:LEU:HB3	1.44	1.13
4:p:210:LEU:CD2	8:A:13:ILE:HG13	1.78	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:B:299:GLN:NE2	9:B:305:GLU:OE1	1.80	1.13
1:a:119:ILE:CB	2:b:28:ALA:HA	1.77	1.13
2:b:78:LEU:O	4:p:140:LEU:HG	1.43	1.13
5:e:19:VAL:CG2	5:e:53:LEU:HB2	1.79	1.13
6:g:281:ILE:HD12	7:M:41:ARG:CG	1.77	1.13
1:a:39:VAL:HG21	1:a:133:ASP:OD2	1.44	1.13
8:E:37:ASP:OD1	9:D:291:ARG:NH2	1.81	1.13
9:D:411:ASP:HA	9:D:418:ARG:HH21	1.11	1.13
1:a:26:PHE:CB	4:p:83:LEU:CA	2.26	1.12
1:a:122:LEU:HD12	1:a:127:LEU:CD1	1.79	1.13
1:a:122:LEU:HD23	1:a:124:HIS:H	1.09	1.12
2:b:88:VAL:HB	4:p:147:VAL:CG1	1.79	1.13
2:b:125:LYS:CD	4:p:185:LEU:HD22	1.62	1.12
2:b:169:ILE:HG23	3:d:221:PHE:CE2	1.84	1.12
8:E:169:ILE:HG21	8:E:344:PHE:CE1	1.83	1.12
9:F:50:LYS:HE2	9:F:92:ILE:HD12	1.28	1.12
1:a:70:ASN:O	2:b:53:LEU:CD1	1.92	1.12
1:a:100:THR:HG21	4:p:101:MET:CB	1.80	1.12
4:p:152:ARG:HD3	8:A:503:LEU:HD22	1.32	1.12
4:p:195:GLN:HG3	8:A:4:ILE:CD1	1.59	1.12
4:p:210:LEU:HD11	8:A:13:ILE:CB	1.79	1.12
9:F:50:LYS:CE	9:F:92:ILE:CD1	2.27	1.12
8:A:492:GLU:O	8:A:496:GLU:HG2	1.48	1.12
9:D:387:ILE:HD13	9:D:455:ILE:HG23	1.23	1.12
1:a:43:SER:OG	4:p:93:ILE:HG12	1.44	1.12
2:b:74:ALA:N	4:p:130:VAL:O	1.82	1.12
1:a:115:LEU:HD23	2:b:38:LEU:HD13	1.28	1.12
4:p:212:ASP:O	4:p:215:VAL:CG1	1.97	1.12
6:g:103:GLU:HG3	7:R:41:ARG:HE	0.95	1.12
8:C:77:MET:CE	8:C:112:LEU:CD2	2.27	1.12
1:a:40:LEU:HG	4:p:89:THR:CB	1.78	1.11
2:b:129:ILE:CG2	4:p:192:LEU:HD11	1.80	1.11
3:d:233:VAL:HG12	8:A:26:VAL:HA	1.21	1.11
8:E:42:ILE:HD13	8:E:89:VAL:HG11	1.32	1.11
9:D:191:LYS:HE2	9:D:219:SER:HB2	1.25	1.11
2:b:114:THR:O	4:p:174:LEU:CD1	1.84	1.11
1:a:70:ASN:OD1	2:b:54:ASP:OD1	1.68	1.11
2:b:144:VAL:HG12	4:p:210:LEU:HD21	1.32	1.11
2:b:160:LEU:HD11	2:b:163:GLU:C	1.58	1.11
8:E:169:ILE:CG2	8:E:344:PHE:HD1	1.58	1.11
2:b:103:ILE:CG2	4:p:159:LEU:HA	1.78	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:160:LEU:HD13	2:b:163:GLU:CB	1.78	1.11
3:d:222:THR:HG22	3:d:234:ASP:HB3	1.25	1.11
9:D:112:ASN:HD22	9:D:116:GLU:HB2	1.07	1.11
9:D:431:LEU:HD23	9:D:458:PHE:CZ	1.83	1.11
1:a:70:ASN:O	2:b:53:LEU:CG	1.98	1.11
2:b:88:VAL:CG2	4:p:148:MET:HG3	1.79	1.11
2:b:117:THR:CG2	4:p:174:LEU:HD11	1.81	1.11
2:b:142:GLN:NE2	3:d:248:LEU:HD23	1.65	1.11
4:p:108:TYR:CE1	4:p:112:LEU:CD1	2.33	1.11
1:a:119:ILE:HD11	2:b:32:ILE:H	1.04	1.10
2:b:77:GLN:HG2	4:p:138:LYS:CG	1.79	1.10
2:b:82:ARG:HA	2:b:85:LEU:HD23	1.24	1.10
2:b:80:LYS:N	4:p:137:VAL:HA	1.66	1.10
2:b:81:ALA:HB3	4:p:140:LEU:N	1.65	1.10
8:C:77:MET:HE1	8:C:112:LEU:CD2	1.81	1.10
9:D:112:ASN:OD1	9:D:118:VAL:HG21	1.50	1.10
1:a:161:PHE:CZ	1:a:163:LYS:HB2	1.85	1.10
2:b:77:GLN:HG2	4:p:138:LYS:HG3	1.11	1.10
2:b:129:ILE:HG23	4:p:192:LEU:HD11	1.13	1.10
9:B:221:VAL:HG11	9:B:232:VAL:HG21	1.25	1.10
8:C:77:MET:HE1	8:C:112:LEU:HD21	1.23	1.10
9:D:112:ASN:ND2	9:D:116:GLU:HB2	1.65	1.10
1:a:116:PRO:CD	2:b:35:SER:N	2.12	1.10
2:b:62:ASN:O	2:b:66:ASN:ND2	1.84	1.10
2:b:103:ILE:CD1	4:p:158:ALA:O	1.99	1.10
1:a:51:LEU:HD23	4:p:100:LEU:HD11	1.30	1.10
2:b:68:GLU:HG3	4:p:126:GLN:HB3	1.18	1.10
4:p:199:THR:C	8:A:6:ALA:CB	2.23	1.10
8:A:494:ILE:HG22	8:A:498:MET:HE2	1.12	1.10
2:b:71:ARG:H	4:p:130:VAL:HG12	1.12	1.09
2:b:88:VAL:HG21	4:p:145:ASN:HA	1.12	1.09
2:b:107:LYS:CA	4:p:166:THR:CG2	1.92	1.09
2:b:117:THR:HG22	4:p:174:LEU:HD11	1.17	1.09
2:b:180:ASN:ND2	3:d:201:ILE:HG21	1.64	1.09
5:e:60:LEU:HD12	5:e:60:LEU:O	1.50	1.09
9:B:294:SER:O	8:C:286:ALA:CB	2.00	1.09
3:d:179:VAL:HG22	3:d:210:LYS:HB2	1.30	1.09
4:p:193:GLU:CG	4:p:196:LYS:HE2	1.82	1.09
5:e:19:VAL:HA	5:e:53:LEU:HD13	1.30	1.09
1:a:119:ILE:HD11	2:b:32:ILE:HG12	1.12	1.09
1:a:119:ILE:HD12	2:b:31:LEU:H	0.92	1.09

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:135:ASN:ND2	4:p:86:PHE:CD1	2.21	1.09
2:b:84:ARG:HB3	4:p:141:GLU:CA	1.83	1.09
2:b:125:LYS:HD3	4:p:185:LEU:CD2	1.77	1.09
6:g:124:THR:CG2	6:g:137:LEU:HD23	1.83	1.09
7:O:43:PRO:HG2	7:P:42:GLN:HE22	1.18	1.09
9:F:50:LYS:CE	9:F:92:ILE:HD11	1.82	1.09
9:B:431:LEU:HD23	9:B:458:PHE:HZ	1.11	1.09
1:a:43:SER:HB2	4:p:93:ILE:HG12	1.28	1.09
1:a:163:LYS:NZ	1:a:179:ASP:OD2	1.83	1.09
2:b:133:GLN:HG3	8:A:3:THR:HG22	1.17	1.09
8:A:383:LEU:HD22	8:A:438:ILE:HD12	1.33	1.09
9:D:431:LEU:CD2	9:D:458:PHE:CZ	2.33	1.09
1:a:119:ILE:HD11	2:b:32:ILE:CG1	1.82	1.09
1:a:119:ILE:HD12	2:b:31:LEU:N	1.66	1.09
1:a:134:ILE:HA	1:a:191:PHE:CD1	1.87	1.09
2:b:145:PHE:CE2	2:b:149:LEU:HD11	1.87	1.09
2:b:161:ASN:O	2:b:165:HIS:HB3	1.49	1.09
2:b:169:ILE:HG23	3:d:221:PHE:HE2	1.02	1.09
6:g:149:LYS:HE2	6:g:155:TYR:HE1	1.18	1.09
1:a:119:ILE:HB	2:b:31:LEU:HG	1.16	1.08
2:b:77:GLN:CD	4:p:138:LYS:CG	2.06	1.08
2:b:165:HIS:CG	3:d:219:ALA:HB1	1.87	1.08
4:p:193:GLU:HG3	4:p:196:LYS:HE2	1.13	1.08
8:C:138:MET:HG3	9:D:119:ASP:HA	1.31	1.08
1:a:230:ILE:HD11	7:Q:55:LEU:HD11	1.23	1.08
2:b:125:LYS:CA	4:p:185:LEU:HD22	1.83	1.08
2:b:141:ARG:HD3	8:A:12:ILE:HG21	1.16	1.08
5:e:96:LEU:HD11	5:e:123:ARG:HG2	1.32	1.08
6:g:119:ALA:HB2	6:g:207:VAL:HG11	1.27	1.08
9:B:410:LEU:O	9:B:418:ARG:NH2	1.86	1.08
8:C:27:VAL:CB	8:C:47:GLU:HB2	1.84	1.08
9:D:112:ASN:OD1	9:D:118:VAL:CG2	2.01	1.08
1:a:118:LYS:HA	2:b:30:ASN:OD1	1.27	1.08
2:b:88:VAL:HG11	4:p:145:ASN:O	1.50	1.08
6:g:281:ILE:HG21	7:M:41:ARG:HA	1.35	1.08
1:a:122:LEU:HD22	1:a:125:GLY:H	1.10	1.08
1:a:161:PHE:HZ	1:a:163:LYS:CD	1.52	1.08
1:a:208:VAL:HG22	1:a:209:PRO:HD2	1.18	1.08
2:b:73:LYS:N	4:p:130:VAL:CG2	2.14	1.08
2:b:75:ILE:O	4:p:137:VAL:HG13	1.54	1.08
6:g:261:GLU:HG3	9:F:397:ARG:CZ	1.82	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:24:GLN:HA	4:p:83:LEU:CD1	1.83	1.08
2:b:73:LYS:H	4:p:130:VAL:HG23	1.00	1.08
2:b:82:ARG:HA	2:b:85:LEU:CD2	1.83	1.08
4:p:210:LEU:CD1	8:A:13:ILE:HB	1.84	1.08
5:e:22:ILE:HG22	5:e:51:ILE:HG13	1.15	1.08
9:F:275:ILE:HG22	9:F:326:ALA:O	1.50	1.08
2:b:77:GLN:HB3	4:p:134:SER:C	1.78	1.07
2:b:133:GLN:O	2:b:137:ILE:HD12	1.52	1.07
2:b:144:VAL:HG11	4:p:210:LEU:HD22	1.36	1.07
2:b:161:ASN:O	2:b:165:HIS:CB	2.02	1.07
3:d:152:SER:C	3:d:153:GLU:OE1	1.97	1.07
4:p:195:GLN:CG	8:A:4:ILE:HD13	1.83	1.07
8:E:104:TYR:CD1	8:E:122:ILE:HD13	1.87	1.07
3:d:199:GLN:NE2	3:d:207:VAL:HG12	1.68	1.07
1:a:51:LEU:CD2	4:p:100:LEU:HD11	1.76	1.07
1:a:161:PHE:CE1	1:a:163:LYS:CD	2.31	1.07
6:g:359:GLY:HA3	9:D:293:PRO:HG3	1.19	1.07
8:C:27:VAL:HB	8:C:47:GLU:CB	1.85	1.07
9:D:431:LEU:HD23	9:D:458:PHE:HZ	0.93	1.07
1:a:24:GLN:HA	4:p:83:LEU:HD11	1.28	1.07
2:b:88:VAL:CA	4:p:148:MET:CG	1.80	1.07
2:b:160:LEU:CG	2:b:164:LEU:CB	2.33	1.07
4:p:192:LEU:HD13	8:A:3:THR:HG21	1.08	1.07
1:a:63:THR:CA	4:p:119:ARG:NH1	1.74	1.07
2:b:71:ARG:N	4:p:127:LEU:O	1.88	1.07
2:b:118:LEU:HD11	4:p:177:GLY:C	1.79	1.07
3:d:87:ASP:CG	3:d:172:ILE:HD11	1.80	1.07
3:d:227:ASN:HB2	8:A:46:ASP:OD2	0.91	1.07
5:e:22:ILE:HG22	5:e:51:ILE:CG1	1.84	1.07
6:g:251:GLU:OE2	6:g:270:MET:HB2	1.54	1.07
8:A:214:GLN:NE2	8:A:219:MET:CB	2.17	1.07
1:a:51:LEU:HG	4:p:100:LEU:HD13	1.07	1.06
2:b:137:ILE:CG1	8:A:5:ARG:HG2	1.84	1.06
3:d:247:GLN:HE22	8:A:13:ILE:HG23	1.14	1.06
8:A:433:GLU:OE1	8:A:462:ARG:HD3	1.53	1.06
1:a:116:PRO:HB3	2:b:34:LEU:H	1.18	1.06
1:a:193:ASN:ND2	1:a:227:GLN:HG3	1.69	1.06
6:g:103:GLU:HG3	7:R:41:ARG:NE	1.58	1.06
1:a:114:LEU:HD13	2:b:37:VAL:O	1.55	1.06
1:a:165:ILE:HG22	1:a:168:THR:O	1.51	1.06
1:a:104:PHE:CB	4:p:97:PHE:CE1	2.38	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:88:VAL:HG22	4:p:148:MET:CG	1.84	1.06
2:b:125:LYS:HB3	4:p:185:LEU:HB2	1.34	1.06
1:a:137:THR:HG21	1:a:191:PHE:HB2	1.35	1.06
2:b:141:ARG:HD2	8:A:12:ILE:HG22	1.26	1.06
6:g:356:ILE:CD1	9:D:295:ALA:HA	1.85	1.06
1:a:118:LYS:CA	2:b:30:ASN:OD1	1.99	1.05
2:b:144:VAL:HG13	4:p:210:LEU:HD23	1.28	1.05
4:p:192:LEU:HD22	8:A:3:THR:OG1	1.52	1.05
5:e:119:LEU:CD1	5:e:123:ARG:CZ	2.34	1.05
6:g:356:ILE:HD11	9:D:295:ALA:HA	1.12	1.05
8:A:98:ILE:HD13	8:A:245:ALA:HB3	1.37	1.05
2:b:69:GLU:CA	4:p:130:VAL:HG11	1.87	1.05
2:b:85:LEU:HD22	4:p:143:GLN:HG3	1.07	1.05
2:b:103:ILE:CD1	4:p:162:MET:HB2	1.86	1.05
4:p:214:ILE:CA	8:A:17:ILE:HD12	1.78	1.05
5:e:19:VAL:HG23	5:e:53:LEU:CB	1.85	1.05
8:A:359:ASN:ND2	8:A:362:ILE:HD12	1.70	1.05
2:b:83:ALA:H	4:p:140:LEU:CB	1.68	1.05
5:e:38:ALA:HB2	7:L:41:ARG:HA	1.07	1.05
8:A:382:LYS:HD3	8:A:442:THR:CG2	1.86	1.05
1:a:208:VAL:HG23	1:a:209:PRO:HD2	1.35	1.05
2:b:122:GLU:CD	4:p:181:ILE:HG23	1.80	1.05
4:p:213:ASP:OD2	8:A:14:ARG:HA	1.55	1.05
5:e:21:GLU:CG	5:e:34:LEU:HD21	1.81	1.05
6:g:103:GLU:HG2	7:R:41:ARG:HE	1.18	1.05
2:b:78:LEU:HD11	4:p:139:GLN:OE1	1.57	1.05
2:b:114:THR:O	4:p:174:LEU:HD11	1.46	1.04
9:B:431:LEU:CD2	9:B:458:PHE:CZ	2.39	1.04
1:a:65:PRO:CB	1:a:70:ASN:ND2	1.73	1.04
1:a:119:ILE:CG1	2:b:28:ALA:C	2.28	1.04
2:b:92:ALA:CB	4:p:147:VAL:O	2.04	1.04
2:b:132:GLU:HG3	4:p:189:LEU:HD22	1.35	1.04
7:M:48:LYS:HB3	7:L:46:GLU:OE1	1.57	1.04
8:C:77:MET:HE3	8:C:112:LEU:HD21	1.38	1.04
1:a:119:ILE:HG23	2:b:28:ALA:N	1.70	1.04
2:b:141:ARG:O	8:A:13:ILE:CD1	2.05	1.04
2:b:160:LEU:HG	2:b:164:LEU:HB3	1.38	1.04
2:b:160:LEU:HD12	2:b:164:LEU:N	1.38	1.04
6:g:118:VAL:HG11	6:g:148:LEU:CD1	1.86	1.04
1:a:139:ALA:HB1	4:p:94:MET:CE	1.87	1.04
1:a:161:PHE:CE1	1:a:163:LYS:HB2	1.81	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:174:ILE:HG21	7:O:58:ALA:HB1	1.39	1.04
5:e:38:ALA:HB3	7:L:41:ARG:HG3	1.07	1.04
9:D:20:LEU:CD2	9:D:92:ILE:HG22	1.88	1.04
1:a:163:LYS:HE2	1:a:179:ASP:OD2	1.56	1.03
1:a:208:VAL:HG22	1:a:209:PRO:CD	1.86	1.03
2:b:77:GLN:CB	4:p:134:SER:O	2.06	1.03
4:p:210:LEU:HD21	8:A:13:ILE:HG13	1.07	1.03
8:C:432:GLU:HG3	8:C:476:ILE:CG2	1.88	1.03
6:g:349:ILE:HD13	8:E:284:ARG:HH21	1.16	1.03
1:a:114:LEU:HB3	2:b:38:LEU:N	1.72	1.03
1:a:119:ILE:CA	2:b:29:THR:N	2.22	1.03
2:b:160:LEU:HG	2:b:164:LEU:HB2	1.38	1.03
3:d:224:ARG:HH11	3:d:232:LEU:HD12	1.01	1.03
9:F:100:VAL:CG1	9:F:256:MET:CE	2.36	1.03
9:B:411:ASP:HA	9:B:418:ARG:HH22	0.89	1.03
1:a:58:VAL:CG1	4:p:107:ILE:HG22	1.89	1.03
2:b:125:LYS:HA	4:p:185:LEU:HD22	1.37	1.03
2:b:142:GLN:HE22	3:d:248:LEU:HA	1.07	1.03
3:d:151:ASP:HB2	8:C:20:TYR:OH	1.58	1.03
1:a:64:ILE:CG2	2:b:57:LYS:HE2	1.89	1.02
1:a:165:ILE:HG23	1:a:168:THR:O	1.58	1.02
2:b:61:LEU:HD11	2:b:65:ARG:CD	1.87	1.02
2:b:88:VAL:CG2	4:p:148:MET:CG	2.35	1.02
2:b:114:THR:O	2:b:117:THR:HG22	1.58	1.02
2:b:129:ILE:HD11	4:p:188:ALA:CB	1.89	1.02
6:g:281:ILE:CA	7:N:41:ARG:NH2	2.22	1.02
6:g:360:ALA:N	9:D:293:PRO:HD2	1.66	1.02
8:A:386:GLU:OE2	8:A:442:THR:HG23	1.57	1.02
9:B:20:LEU:HD11	9:B:90:GLU:OE2	1.59	1.02
1:a:135:ASN:ND2	4:p:86:PHE:CA	1.97	1.02
2:b:103:ILE:CG1	4:p:163:LYS:N	2.22	1.02
1:a:70:ASN:O	2:b:53:LEU:HB3	1.58	1.02
2:b:68:GLU:CD	4:p:126:GLN:HB3	1.85	1.02
7:G:43:PRO:HG2	7:H:42:GLN:HE22	1.18	1.02
7:H:43:PRO:HG2	7:I:42:GLN:HE22	1.23	1.02
1:a:135:ASN:OD1	4:p:86:PHE:CG	2.12	1.02
2:b:77:GLN:CG	4:p:138:LYS:CG	2.28	1.02
9:B:411:ASP:HA	9:B:418:ARG:NH2	1.73	1.02
9:D:401:LEU:CD2	9:D:404:ILE:HD11	1.88	1.02
4:p:152:ARG:HG2	8:A:503:LEU:HB3	1.39	1.02
7:K:42:GLN:OE1	7:J:40:ALA:O	1.77	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:122:LEU:CD1	1:a:127:LEU:CD1	2.37	1.01
3:d:74:THR:HG21	8:C:4:ILE:HG22	1.39	1.01
1:a:134:ILE:CG2	1:a:191:PHE:CE1	2.43	1.01
1:a:184:LEU:HD23	1:a:188:PHE:HD2	1.24	1.01
2:b:103:ILE:HG12	4:p:162:MET:HB2	1.40	1.01
2:b:107:LYS:O	4:p:166:THR:OG1	1.68	1.01
2:b:160:LEU:CG	2:b:164:LEU:N	2.11	1.01
3:d:148:ILE:HD11	8:C:17:ILE:HA	1.02	1.01
4:p:152:ARG:HB3	8:A:503:LEU:HD23	1.02	1.01
6:g:123:VAL:HG11	6:g:308:LEU:HD23	1.02	1.01
9:B:237:GLY:HA3	9:B:249:VAL:HG21	1.42	1.01
8:C:430:THR:HG22	8:C:432:GLU:OE1	1.60	1.01
1:a:182:LYS:HD2	1:a:238:TYR:CE1	1.95	1.01
2:b:145:PHE:HE2	3:d:244:ILE:HA	1.20	1.01
4:p:152:ARG:NE	8:A:503:LEU:HB2	1.75	1.01
5:e:34:LEU:CD1	7:K:43:PRO:HG2	1.89	1.01
7:S:68:LEU:HD12	7:T:66:TYR:CG	1.95	1.01
9:B:20:LEU:CD2	9:B:92:ILE:HG12	1.91	1.01
1:a:35:ILE:CG2	4:p:88:LEU:HB2	1.91	1.01
1:a:114:LEU:HA	2:b:37:VAL:HB	1.38	1.01
2:b:144:VAL:CG2	8:A:9:ILE:CG2	2.31	1.01
7:S:43:PRO:CB	7:T:48:LYS:NZ	2.23	1.01
8:E:104:TYR:CE1	8:E:122:ILE:HD13	1.96	1.01
1:a:70:ASN:O	2:b:53:LEU:CB	2.09	1.01
1:a:134:ILE:HG22	1:a:191:PHE:CE2	1.88	1.01
2:b:118:LEU:CD1	4:p:177:GLY:C	2.33	1.01
3:d:154:ARG:HE	8:C:5:ARG:HH12	1.09	1.01
6:g:281:ILE:CD1	7:M:41:ARG:HG3	1.90	1.01
9:D:179:THR:O	9:D:182:ILE:HG22	1.60	1.01
2:b:176:PHE:CZ	3:d:233:VAL:HG21	1.94	1.00
3:d:74:THR:CG2	8:C:4:ILE:HG22	1.92	1.00
5:e:38:ALA:CB	7:L:41:ARG:HG3	1.91	1.00
6:g:149:LYS:HE2	6:g:155:TYR:CE1	1.95	1.00
8:E:344:PHE:CE2	8:E:362:ILE:CG2	2.43	1.00
8:C:237:TYR:CD1	8:C:271:TYR:HB2	1.96	1.00
4:p:214:ILE:CA	8:A:17:ILE:HD13	1.79	1.00
7:T:40:ALA:O	7:G:42:GLN:OE1	1.77	1.00
8:E:208:GLN:HE22	9:D:144:THR:CG2	1.74	1.00
2:b:71:ARG:HH11	4:p:126:GLN:HA	1.24	1.00
2:b:82:ARG:O	2:b:85:LEU:CG	2.09	1.00
2:b:103:ILE:HG12	4:p:162:MET:CB	1.90	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:B:447:LYS:NZ	9:B:482:GLU:OE2	1.93	1.00
4:p:206:GLN:O	4:p:210:LEU:N	1.95	1.00
9:F:110:ILE:HG22	9:F:118:VAL:CG2	1.90	1.00
3:d:131:GLU:C	8:E:13:ILE:HD11	1.85	1.00
4:p:196:LYS:O	4:p:199:THR:HG22	1.62	1.00
1:a:100:THR:HG21	4:p:101:MET:HB2	1.37	1.00
2:b:118:LEU:HD13	4:p:178:ARG:N	1.75	1.00
3:d:199:GLN:HE21	3:d:207:VAL:HG12	1.27	1.00
1:a:135:ASN:CG	4:p:86:PHE:CG	2.40	0.99
1:a:116:PRO:CD	2:b:34:LEU:C	2.33	0.99
2:b:103:ILE:CD1	4:p:162:MET:CA	2.40	0.99
2:b:144:VAL:HG21	8:A:9:ILE:HG21	1.00	0.99
7:G:40:ALA:O	7:H:42:GLN:OE1	1.80	0.99
1:a:63:THR:HA	4:p:119:ARG:NH1	1.14	0.99
2:b:103:ILE:HD11	4:p:162:MET:CA	1.93	0.99
4:p:192:LEU:HD13	8:A:3:THR:CG2	1.91	0.99
7:T:40:ALA:HB2	7:G:49:ILE:HD11	1.43	0.99
6:g:251:GLU:OE2	6:g:270:MET:CB	2.10	0.99
7:S:43:PRO:CB	7:T:48:LYS:HZ3	1.74	0.99
8:A:214:GLN:HE22	8:A:219:MET:CG	1.65	0.99
6:g:353:ILE:HD11	8:E:284:ARG:HG2	1.44	0.99
9:B:221:VAL:HG11	9:B:232:VAL:CG2	1.93	0.99
1:a:119:ILE:CB	2:b:31:LEU:HG	1.93	0.99
1:a:122:LEU:HD12	1:a:127:LEU:HD11	1.00	0.99
7:P:43:PRO:CB	7:Q:48:LYS:HZ3	1.76	0.99
8:E:344:PHE:CD2	8:E:362:ILE:HG22	1.95	0.99
1:a:24:GLN:NE2	1:a:195:LEU:HD11	1.77	0.99
1:a:116:PRO:CB	2:b:34:LEU:H	1.76	0.99
4:p:152:ARG:CB	8:A:503:LEU:HD23	1.92	0.99
6:g:118:VAL:HG11	6:g:148:LEU:HD11	1.44	0.99
9:B:221:VAL:CG1	9:B:232:VAL:CG2	2.40	0.99
1:a:134:ILE:HG23	1:a:191:PHE:CE2	1.91	0.98
7:O:40:ALA:O	7:P:42:GLN:OE1	1.81	0.98
1:a:115:LEU:HD21	2:b:38:LEU:HD13	1.03	0.98
2:b:144:VAL:HG12	8:A:13:ILE:HG13	1.43	0.98
7:T:43:PRO:CB	7:G:48:LYS:NZ	2.26	0.98
8:A:8:GLU:O	8:A:12:ILE:HG13	1.63	0.98
3:d:115:PHE:HB3	3:d:155:ILE:HD11	1.45	0.98
1:a:119:ILE:HB	2:b:31:LEU:CG	1.94	0.98
4:p:203:LEU:HD23	8:A:6:ALA:O	1.64	0.98
5:e:21:GLU:HG2	5:e:34:LEU:HD23	1.01	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:K:48:LYS:NZ	7:J:43:PRO:CB	2.27	0.98
8:E:469:LYS:CD	8:E:493:ALA:HB2	1.93	0.98
1:a:40:LEU:HD21	4:p:89:THR:CA	1.93	0.98
1:a:43:SER:OG	4:p:93:ILE:CG1	2.11	0.98
1:a:79:ILE:HD13	1:a:99:GLY:CA	1.93	0.98
6:g:362:ALA:HB1	8:C:281:PRO:HB3	1.45	0.98
7:S:40:ALA:O	7:T:42:GLN:OE1	1.81	0.98
7:H:40:ALA:O	7:I:42:GLN:OE1	1.81	0.98
9:F:387:ILE:HD11	9:F:458:PHE:HB2	1.41	0.98
8:A:28:ASN:HB2	8:A:91:ALA:HB3	1.42	0.98
3:d:186:LEU:HD11	3:d:191:LEU:CG	1.92	0.98
5:e:21:GLU:OE2	5:e:34:LEU:HD21	1.64	0.98
6:g:83:ARG:NH2	6:g:273:THR:CG2	2.27	0.98
6:g:148:LEU:O	6:g:152:GLY:N	1.96	0.98
2:b:70:LEU:HB3	4:p:131:LYS:HB2	1.44	0.98
8:C:392:GLU:CD	9:D:429:ARG:HH22	1.71	0.98
1:a:100:THR:CG2	4:p:101:MET:CG	2.34	0.98
4:p:108:TYR:CE1	4:p:112:LEU:HD12	1.99	0.98
6:g:261:GLU:CD	9:F:397:ARG:NH2	2.22	0.98
2:b:144:VAL:CG1	4:p:210:LEU:HD21	1.90	0.97
5:e:21:GLU:HG2	5:e:34:LEU:HD21	1.40	0.97
6:g:281:ILE:HD12	7:M:41:ARG:HG2	1.43	0.97
7:S:68:LEU:HD12	7:T:66:TYR:HB3	1.45	0.97
3:d:186:LEU:HD11	3:d:191:LEU:CD2	1.93	0.97
4:p:210:LEU:HD11	8:A:13:ILE:HG22	1.44	0.97
3:d:115:PHE:HB3	3:d:155:ILE:CD1	1.95	0.97
8:E:42:ILE:HD11	8:E:56:PHE:CZ	1.98	0.97
1:a:139:ALA:CB	4:p:94:MET:SD	2.52	0.97
6:g:221:VAL:HG11	8:A:401:SER:HB2	1.46	0.97
1:a:134:ILE:HG23	1:a:191:PHE:CE1	1.99	0.97
6:g:261:GLU:HG2	6:g:262:GLY:H	1.30	0.97
7:P:40:ALA:O	7:Q:42:GLN:OE1	1.80	0.97
9:B:294:SER:O	8:C:286:ALA:HB3	1.64	0.97
9:D:163:ARG:HD2	9:D:374:MET:HE3	1.45	0.97
8:E:355:ARG:CZ	9:D:389:GLN:HE22	1.77	0.97
2:b:64:ILE:CD1	4:p:123:ILE:CG1	2.33	0.97
5:e:38:ALA:HB2	7:L:41:ARG:CA	1.94	0.97
9:B:431:LEU:HD21	9:B:458:PHE:CZ	1.98	0.96
1:a:53:SER:O	1:a:56:ILE:HG22	1.65	0.96
2:b:84:ARG:HB3	4:p:141:GLU:HA	0.97	0.96
7:T:43:PRO:CB	7:G:48:LYS:HZ3	1.77	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:164:TYR:CD2	1:a:166:GLN:HG2	1.98	0.96
3:d:242:GLU:CG	8:A:20:TYR:OH	2.13	0.96
6:g:281:ILE:HB	7:M:41:ARG:HG3	1.45	0.96
8:A:159:PRO:HG2	8:A:368:GLY:O	1.63	0.96
2:b:76:GLU:O	2:b:80:LYS:CG	2.14	0.96
2:b:85:LEU:CD2	4:p:143:GLN:HG3	1.81	0.96
6:g:118:VAL:CG1	6:g:148:LEU:CD1	2.42	0.96
7:J:74:LEU:CD2	7:I:10:VAL:CG2	2.44	0.96
1:a:65:PRO:HB2	1:a:70:ASN:CG	1.90	0.96
1:a:119:ILE:CD1	2:b:31:LEU:H	1.77	0.96
1:a:214:ILE:HG13	1:a:215:PRO:HD3	1.47	0.96
2:b:67:SER:O	4:p:130:VAL:HG12	1.66	0.96
2:b:88:VAL:CB	4:p:148:MET:CG	2.21	0.96
5:e:119:LEU:HD11	5:e:123:ARG:CZ	1.91	0.96
8:A:382:LYS:CD	8:A:442:THR:HG21	1.94	0.96
1:a:100:THR:HG21	4:p:101:MET:HG3	1.22	0.96
2:b:118:LEU:HD11	4:p:177:GLY:O	1.64	0.96
4:p:108:TYR:CD1	4:p:112:LEU:HD12	2.00	0.96
2:b:107:LYS:CA	4:p:166:THR:OG1	2.07	0.96
4:p:192:LEU:CD1	8:A:3:THR:HG21	1.95	0.96
5:e:119:LEU:HD11	5:e:123:ARG:HE	0.99	0.96
7:P:43:PRO:CB	7:Q:48:LYS:NZ	2.27	0.96
8:E:344:PHE:HD2	8:E:362:ILE:CG2	1.63	0.96
6:g:359:GLY:HA3	9:D:293:PRO:CG	1.95	0.96
1:a:161:PHE:CE1	1:a:163:LYS:HD2	1.99	0.96
6:g:251:GLU:OE2	6:g:270:MET:SD	2.23	0.96
6:g:221:VAL:HG13	8:A:401:SER:OG	1.64	0.95
2:b:133:GLN:HG2	8:A:3:THR:HG23	1.48	0.95
6:g:356:ILE:HD11	9:D:295:ALA:CA	1.95	0.95
7:J:74:LEU:HD22	7:I:10:VAL:CG2	1.96	0.95
9:F:361:ILE:HG23	9:F:432:SER:HB3	1.44	0.95
1:a:222:PHE:HZ	7:Q:54:LEU:CD2	1.78	0.95
2:b:95:PHE:CD2	4:p:151:ALA:O	2.18	0.95
2:b:133:GLN:CG	8:A:3:THR:HG23	1.96	0.95
9:F:120:ASN:O	9:F:120:ASN:OD1	1.82	0.95
8:A:469:LYS:NZ	8:A:492:GLU:HB2	1.81	0.95
8:C:49:MET:SD	8:C:95:ILE:HD11	2.06	0.95
9:D:189:ILE:HG22	9:D:269:LEU:HD11	1.49	0.95
1:a:222:PHE:CZ	7:Q:54:LEU:CD2	2.47	0.95
3:d:195:ALA:O	3:d:199:GLN:HG3	1.66	0.95
3:d:228:GLU:CD	8:A:43:HIS:CE1	2.44	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:172:ARG:HB3	8:A:321:GLU:OE2	1.65	0.95
1:a:39:VAL:HG21	1:a:133:ASP:CG	1.89	0.95
3:d:186:LEU:HD11	3:d:191:LEU:HD21	1.48	0.95
7:Q:43:PRO:HG2	7:R:42:GLN:HE22	1.27	0.95
2:b:92:ALA:HB2	4:p:147:VAL:O	1.64	0.95
2:b:148:ALA:HA	4:p:211:SER:HA	1.46	0.95
8:E:208:GLN:HE22	9:D:144:THR:HG23	1.13	0.95
8:E:355:ARG:CZ	9:D:389:GLN:NE2	2.30	0.95
2:b:142:GLN:OE1	3:d:248:LEU:HD23	1.56	0.95
9:B:251:LEU:CD2	9:B:309:LEU:HD13	1.96	0.95
1:a:104:PHE:CB	4:p:97:PHE:CZ	2.49	0.95
1:a:122:LEU:HD22	1:a:125:GLY:N	1.81	0.95
2:b:68:GLU:HB2	4:p:126:GLN:HB2	0.96	0.95
2:b:81:ALA:CB	4:p:140:LEU:N	2.24	0.95
4:p:210:LEU:HD11	8:A:13:ILE:HB	1.41	0.95
2:b:125:LYS:CG	4:p:185:LEU:HD22	1.96	0.95
2:b:144:VAL:HG21	8:A:9:ILE:HG22	1.49	0.95
7:K:48:LYS:HZ3	7:J:43:PRO:CB	1.80	0.95
8:A:98:ILE:CD1	8:A:245:ALA:CB	2.45	0.95
9:B:347:ASP:OD1	8:C:202:LYS:NZ	2.00	0.95
8:C:392:GLU:CD	9:D:429:ARG:NH2	2.24	0.95
1:a:134:ILE:N	1:a:191:PHE:CE1	2.35	0.94
2:b:79:GLU:N	4:p:136:GLU:HG3	1.81	0.94
7:J:42:GLN:HE22	7:I:43:PRO:HG2	1.31	0.94
7:Q:40:ALA:O	7:R:42:GLN:OE1	1.85	0.94
8:A:98:ILE:CD1	8:A:245:ALA:HB3	1.96	0.94
8:E:169:ILE:HD11	8:E:322:THR:HG21	1.45	0.94
1:a:104:PHE:CD1	4:p:97:PHE:CD1	2.56	0.94
2:b:125:LYS:HB3	4:p:185:LEU:CB	1.96	0.94
3:d:224:ARG:NH1	3:d:232:LEU:HD13	1.78	0.94
6:g:281:ILE:HD12	7:M:41:ARG:HG3	1.45	0.94
2:b:73:LYS:H	4:p:130:VAL:HG22	1.32	0.94
4:p:195:GLN:HG3	8:A:4:ILE:HD13	1.37	0.94
7:M:66:TYR:HB3	7:L:68:LEU:HD13	1.48	0.94
8:C:165:ARG:HD3	8:C:299:LEU:O	1.67	0.94
2:b:64:ILE:CD1	4:p:123:ILE:HG13	1.98	0.94
4:p:152:ARG:HG2	8:A:503:LEU:CB	1.97	0.94
9:F:179:THR:HG21	9:F:215:GLU:OE1	1.68	0.94
1:a:65:PRO:HB3	1:a:70:ASN:ND2	1.82	0.94
2:b:165:HIS:HD2	3:d:219:ALA:HB1	1.32	0.94
5:e:22:ILE:CG1	5:e:33:VAL:HG21	1.98	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:L:42:GLN:HE22	7:K:43:PRO:HG2	1.30	0.94
8:E:169:ILE:HG22	8:E:344:PHE:HD1	1.33	0.94
8:E:344:PHE:CE2	8:E:362:ILE:HG21	2.02	0.94
8:A:111:ALA:N	8:A:227:GLU:OE2	2.00	0.94
2:b:103:ILE:HD13	4:p:162:MET:H	1.31	0.94
5:e:37:HIS:CE1	7:L:41:ARG:HE	1.85	0.94
6:g:137:LEU:CD1	6:g:213:LEU:CD1	2.46	0.94
6:g:281:ILE:CB	7:M:41:ARG:HG3	1.97	0.94
2:b:160:LEU:HD13	2:b:163:GLU:HB2	1.49	0.94
2:b:173:ILE:HG12	3:d:221:PHE:HZ	1.32	0.94
3:d:148:ILE:CD1	8:C:17:ILE:CA	2.35	0.94
4:p:213:ASP:OD2	8:A:14:ARG:CG	2.15	0.94
2:b:60:ILE:O	2:b:63:THR:HG22	1.68	0.93
2:b:122:GLU:HA	4:p:181:ILE:CG2	1.98	0.93
3:d:227:ASN:HB2	8:A:46:ASP:CG	1.93	0.93
6:g:149:LYS:CE	6:g:155:TYR:HE1	1.79	0.93
8:A:159:PRO:CG	8:A:368:GLY:O	2.16	0.93
9:B:251:LEU:HD23	9:B:309:LEU:HD13	1.48	0.93
2:b:133:GLN:HG2	8:A:3:THR:CA	1.98	0.93
2:b:160:LEU:CG	2:b:164:LEU:HB2	1.96	0.93
6:g:281:ILE:HA	7:N:41:ARG:HH22	1.15	0.93
1:a:137:THR:HG21	1:a:191:PHE:CB	1.97	0.93
4:p:159:LEU:HD12	4:p:160:ASN:N	1.82	0.93
1:a:122:LEU:CD2	1:a:124:HIS:H	1.81	0.93
7:K:42:GLN:HE22	7:J:43:PRO:HG2	1.33	0.93
7:K:49:ILE:HD11	7:J:40:ALA:HB2	1.46	0.93
7:O:43:PRO:CB	7:P:48:LYS:HZ3	1.82	0.93
2:b:122:GLU:OE1	4:p:181:ILE:HG23	0.76	0.93
2:b:160:LEU:HD12	2:b:164:LEU:H	0.98	0.93
4:p:152:ARG:HD3	8:A:503:LEU:CD2	1.97	0.93
7:J:74:LEU:CD2	7:I:10:VAL:HG23	1.98	0.93
8:E:284:ARG:HH11	8:E:330:TYR:HD1	1.13	0.93
9:B:431:LEU:HD23	9:B:458:PHE:CZ	2.00	0.93
1:a:208:VAL:HG23	1:a:209:PRO:CD	1.96	0.93
6:g:123:VAL:HG11	6:g:308:LEU:CD2	1.84	0.93
8:A:109:ILE:HD11	8:A:113:ALA:HA	1.50	0.93
2:b:85:LEU:O	4:p:147:VAL:HG12	1.66	0.93
7:T:43:PRO:HG2	7:G:42:GLN:HE22	1.34	0.93
7:J:42:GLN:OE1	7:I:40:ALA:O	1.85	0.93
8:A:141:ARG:NH1	8:A:311:GLU:OE2	2.02	0.93
8:A:494:ILE:HG22	8:A:498:MET:CE	1.98	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:103:ILE:CG1	4:p:163:LYS:H	1.82	0.93
6:g:137:LEU:HD11	6:g:213:LEU:HD12	1.47	0.93
9:D:275:ILE:CG2	9:D:327:VAL:HG22	1.99	0.93
1:a:119:ILE:HG22	2:b:27:LEU:O	1.66	0.93
2:b:85:LEU:HD12	2:b:86:LYS:N	1.84	0.93
2:b:142:GLN:OE1	3:d:248:LEU:HD21	1.68	0.92
6:g:253:GLU:HB3	6:g:270:MET:HB3	1.52	0.92
8:C:157:MET:HB3	8:C:384:LYS:HD3	1.51	0.92
1:a:24:GLN:O	4:p:83:LEU:C	2.12	0.92
2:b:145:PHE:CE2	3:d:244:ILE:HA	2.04	0.92
4:p:193:GLU:O	4:p:196:LYS:HG2	1.69	0.92
6:g:261:GLU:OE2	9:F:397:ARG:NH2	2.02	0.92
9:F:50:LYS:CE	9:F:92:ILE:HD12	1.96	0.92
8:A:13:ILE:HD13	8:A:16:ARG:CZ	1.98	0.92
1:a:35:ILE:CG2	4:p:88:LEU:N	2.14	0.92
2:b:103:ILE:CB	4:p:159:LEU:CA	1.94	0.92
5:e:22:ILE:HG13	5:e:33:VAL:CG2	1.98	0.92
7:S:48:LYS:NZ	7:R:43:PRO:CB	2.33	0.92
8:C:54:VAL:CG1	8:C:89:VAL:HG22	1.99	0.92
1:a:119:ILE:HD12	2:b:32:ILE:H	1.32	0.92
1:a:164:TYR:HD2	1:a:166:GLN:HG3	0.76	0.92
9:B:294:SER:O	8:C:286:ALA:HB2	1.67	0.92
8:C:432:GLU:HG3	8:C:476:ILE:HG21	1.49	0.92
2:b:88:VAL:CB	4:p:147:VAL:CG1	2.43	0.92
7:S:43:PRO:HB2	7:T:48:LYS:NZ	1.84	0.92
7:O:46:GLU:OE1	7:P:48:LYS:HB3	1.68	0.92
1:a:119:ILE:CG2	2:b:27:LEU:O	2.04	0.92
2:b:103:ILE:HD13	4:p:162:MET:HB2	1.51	0.92
3:d:222:THR:HG21	3:d:234:ASP:CB	1.98	0.92
7:N:10:VAL:CG2	7:O:74:LEU:CD2	2.46	0.92
1:a:25:HIS:O	4:p:83:LEU:CB	2.17	0.92
1:a:64:ILE:HG23	2:b:57:LYS:HE2	1.50	0.92
1:a:135:ASN:ND2	4:p:86:PHE:CG	2.38	0.92
1:a:135:ASN:C	4:p:90:LEU:HD22	1.94	0.92
2:b:142:GLN:HE22	3:d:248:LEU:CA	1.83	0.92
3:d:154:ARG:NH2	8:C:5:ARG:NH1	2.14	0.92
1:a:119:ILE:CG1	2:b:32:ILE:HG12	1.98	0.92
2:b:103:ILE:CD1	4:p:162:MET:CB	2.46	0.92
2:b:107:LYS:HE3	4:p:166:THR:HA	1.51	0.92
1:a:122:LEU:O	2:b:25:ASP:OD2	1.87	0.92
4:p:210:LEU:CD2	8:A:13:ILE:CB	2.48	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:146:LEU:HD22	9:D:374:MET:HE1	1.49	0.92
1:a:119:ILE:HA	2:b:29:THR:H	0.98	0.92
6:g:221:VAL:CG2	8:A:398:GLN:O	2.17	0.92
1:a:24:GLN:O	4:p:83:LEU:O	1.88	0.91
2:b:160:LEU:CD1	2:b:164:LEU:CA	1.94	0.91
6:g:281:ILE:HA	7:N:41:ARG:HH21	1.14	0.91
2:b:80:LYS:H	4:p:137:VAL:CB	1.83	0.91
4:p:152:ARG:HD3	8:A:503:LEU:HB2	1.51	0.91
7:N:43:PRO:CB	7:O:48:LYS:HZ3	1.82	0.91
7:N:46:GLU:OE1	7:O:48:LYS:HB3	1.70	0.91
7:H:43:PRO:CB	7:I:48:LYS:HZ3	1.81	0.91
8:C:54:VAL:HG12	8:C:89:VAL:CG2	2.00	0.91
2:b:81:ALA:CB	4:p:139:GLN:C	2.44	0.91
6:g:73:ARG:NH1	9:F:412:GLU:CD	2.28	0.91
7:S:68:LEU:HD12	7:T:66:TYR:CB	1.99	0.91
7:N:40:ALA:HB2	7:O:49:ILE:HD11	1.53	0.91
1:a:40:LEU:HD23	4:p:89:THR:OG1	1.68	0.91
5:e:34:LEU:HD12	7:K:43:PRO:HG2	0.93	0.91
8:E:379:VAL:HG12	8:E:438:ILE:CG2	2.00	0.91
5:e:119:LEU:HD13	5:e:123:ARG:NE	1.84	0.91
6:g:239:ILE:HD12	6:g:303:GLN:HE21	1.19	0.91
7:O:10:VAL:CG2	7:P:74:LEU:CD2	2.47	0.91
7:G:43:PRO:HG2	7:H:42:GLN:NE2	1.86	0.91
5:e:22:ILE:CG2	5:e:51:ILE:HG13	2.01	0.91
6:g:53:SER:HB3	9:D:403:ASP:OD2	1.69	0.91
7:O:43:PRO:HG2	7:P:42:GLN:NE2	1.85	0.91
9:F:110:ILE:HG22	9:F:118:VAL:HG22	1.50	0.91
1:a:79:ILE:HD13	1:a:99:GLY:HA2	1.50	0.91
2:b:88:VAL:HA	4:p:148:MET:HG2	0.93	0.91
6:g:251:GLU:OE2	6:g:270:MET:CG	2.18	0.91
1:a:124:HIS:NE2	4:p:79:GLU:N	2.18	0.91
1:a:230:ILE:HD11	7:Q:55:LEU:HD13	1.50	0.91
4:p:195:GLN:CG	8:A:4:ILE:HD12	2.00	0.91
2:b:180:ASN:HB3	3:d:201:ILE:HG22	1.52	0.91
6:g:356:ILE:O	9:D:293:PRO:HG2	1.70	0.91
7:H:46:GLU:OE1	7:I:48:LYS:HB3	1.69	0.91
2:b:118:LEU:CD1	4:p:178:ARG:N	2.33	0.91
2:b:145:PHE:CB	8:A:16:ARG:NH2	2.30	0.91
5:e:19:VAL:HG23	5:e:53:LEU:HB2	0.93	0.91
7:P:43:PRO:HG2	7:Q:42:GLN:HE22	1.35	0.91
2:b:88:VAL:CG2	4:p:148:MET:SD	2.60	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:90:LEU:HB2	4:p:91:PRO:HD3	1.52	0.90
8:A:98:ILE:HD11	8:A:245:ALA:CB	2.01	0.90
8:C:248:GLU:OE2	8:C:301:ARG:NE	2.03	0.90
2:b:82:ARG:C	2:b:85:LEU:HG	1.94	0.90
3:d:186:LEU:CD1	3:d:191:LEU:HG	2.00	0.90
7:N:48:LYS:HZ3	7:M:43:PRO:CB	1.84	0.90
7:M:48:LYS:CB	7:L:46:GLU:OE1	2.19	0.90
8:A:109:ILE:HD11	8:A:113:ALA:CA	2.02	0.90
4:p:152:ARG:CD	8:A:503:LEU:CB	2.49	0.90
7:M:74:LEU:CD2	7:L:10:VAL:CG2	2.49	0.90
8:A:13:ILE:HD12	8:A:16:ARG:NH2	1.85	0.90
8:A:111:ALA:HB2	8:A:227:GLU:CD	1.96	0.90
1:a:134:ILE:HG22	1:a:191:PHE:CD1	2.06	0.90
2:b:103:ILE:CG1	4:p:162:MET:CB	2.48	0.90
2:b:160:LEU:CD1	2:b:164:LEU:CB	2.50	0.90
8:A:98:ILE:HD11	8:A:245:ALA:HB1	1.51	0.90
8:C:104:TYR:CD1	8:C:122:ILE:HD13	2.07	0.90
1:a:161:PHE:CZ	1:a:163:LYS:CG	2.40	0.90
9:F:447:LYS:NZ	9:F:482:GLU:OE2	2.03	0.90
4:p:195:GLN:HG3	8:A:4:ILE:HD12	1.53	0.90
7:M:74:LEU:CD2	7:L:10:VAL:HG23	2.02	0.90
8:A:382:LYS:HD3	8:A:442:THR:HG21	1.51	0.90
9:D:109:ARG:NH1	9:D:119:ASP:OD1	2.05	0.90
1:a:120:ILE:HG13	2:b:25:ASP:O	1.70	0.90
2:b:99:GLY:O	4:p:159:LEU:HB3	1.71	0.90
3:d:247:GLN:NE2	8:A:13:ILE:HA	1.87	0.90
2:b:82:ARG:CG	4:p:140:LEU:HD23	2.00	0.90
3:d:154:ARG:HE	8:C:5:ARG:NH1	1.68	0.90
7:N:10:VAL:CG2	7:O:74:LEU:HD22	2.01	0.90
1:a:119:ILE:HD13	2:b:32:ILE:N	1.83	0.90
2:b:82:ARG:HB3	4:p:140:LEU:CG	2.02	0.90
4:p:206:GLN:O	4:p:210:LEU:HB2	1.70	0.90
7:N:10:VAL:HG23	7:O:74:LEU:CD2	2.02	0.90
1:a:79:ILE:CD1	1:a:99:GLY:HA2	2.02	0.90
2:b:106:GLU:OE1	4:p:163:LYS:HD2	1.71	0.90
6:g:281:ILE:CD1	7:M:41:ARG:CG	2.48	0.90
7:S:40:ALA:HB2	7:T:49:ILE:HD11	1.54	0.90
7:G:10:VAL:CG2	7:H:74:LEU:CD2	2.50	0.90
1:a:135:ASN:OD1	4:p:86:PHE:CB	2.20	0.89
6:g:103:GLU:HG2	7:R:41:ARG:NE	1.72	0.89
6:g:119:ALA:CB	6:g:207:VAL:HG11	2.01	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:70:ASN:C	2:b:53:LEU:HB3	1.97	0.89
4:p:152:ARG:CD	8:A:503:LEU:HD22	2.01	0.89
2:b:78:LEU:HA	4:p:135:SER:O	1.72	0.89
3:d:243:GLU:CD	4:p:218:VAL:HG12	1.98	0.89
7:K:48:LYS:NZ	7:J:43:PRO:HB2	1.87	0.89
9:B:425:ARG:NH1	9:B:429:ARG:HH22	1.71	0.89
8:C:271:TYR:CD2	8:C:294:LEU:HD22	2.08	0.89
2:b:173:ILE:CG1	3:d:221:PHE:HZ	1.86	0.89
4:p:213:ASP:CG	8:A:14:ARG:HG2	1.96	0.89
1:a:118:LYS:CB	2:b:29:THR:O	2.20	0.89
1:a:124:HIS:CD2	4:p:79:GLU:HB3	2.05	0.89
2:b:125:LYS:O	4:p:185:LEU:CD1	2.21	0.89
5:e:96:LEU:HD11	5:e:123:ARG:CG	2.03	0.89
6:g:353:ILE:HD11	8:E:284:ARG:HA	1.54	0.89
8:E:140:ARG:HE	9:F:206:THR:CG2	1.85	0.89
4:p:199:THR:O	8:A:6:ALA:HB1	1.71	0.89
6:g:104:ASP:HA	7:S:41:ARG:NH2	1.88	0.89
2:b:133:GLN:HG3	8:A:3:THR:CG2	1.82	0.89
4:p:167:GLN:O	4:p:171:GLU:HB2	1.73	0.89
4:p:199:THR:HA	4:p:202:SER:OG	1.73	0.89
6:g:251:GLU:HB2	6:g:272:LYS:HG2	1.52	0.89
9:B:149:PHE:CZ	9:B:189:ILE:HD13	2.08	0.89
6:g:261:GLU:CD	9:F:397:ARG:HH21	1.81	0.89
6:g:343:ARG:NH2	8:A:326:ASP:HB2	1.87	0.89
3:d:132:ILE:N	8:E:13:ILE:HD11	1.88	0.89
5:e:22:ILE:CG1	5:e:33:VAL:CG2	2.52	0.88
6:g:137:LEU:CD1	6:g:213:LEU:HD13	2.02	0.88
6:g:359:GLY:O	9:D:293:PRO:HD3	1.72	0.88
7:T:43:PRO:HB2	7:G:48:LYS:NZ	1.86	0.88
9:D:431:LEU:HD21	9:D:458:PHE:CZ	2.06	0.88
1:a:54:ALA:CB	4:p:104:LEU:HD11	2.03	0.88
2:b:145:PHE:N	8:A:13:ILE:HD12	1.87	0.88
7:H:43:PRO:HG2	7:I:42:GLN:NE2	1.87	0.88
2:b:137:ILE:HG13	8:A:5:ARG:CG	2.03	0.88
8:E:460:GLU:O	8:E:463:THR:HG22	1.72	0.88
1:a:124:HIS:CE1	4:p:79:GLU:HB3	2.07	0.88
2:b:103:ILE:CD1	4:p:163:LYS:H	1.86	0.88
7:N:43:PRO:CB	7:O:48:LYS:NZ	2.37	0.88
2:b:103:ILE:HG13	4:p:163:LYS:N	1.87	0.88
2:b:180:ASN:CG	3:d:201:ILE:HG21	1.97	0.88
1:a:54:ALA:CB	4:p:104:LEU:CD1	2.52	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:J:48:LYS:NZ	7:I:43:PRO:CB	2.37	0.88
2:b:133:GLN:HG2	8:A:3:THR:CB	2.02	0.88
4:p:214:ILE:HA	8:A:17:ILE:HD13	0.90	0.88
6:g:103:GLU:HG3	7:R:41:ARG:CG	2.04	0.88
6:g:221:VAL:HG21	8:A:401:SER:HB3	1.52	0.88
9:B:149:PHE:HZ	9:B:189:ILE:HD13	1.37	0.88
2:b:84:ARG:CG	4:p:144:ALA:HB3	2.04	0.88
2:b:129:ILE:HD11	4:p:188:ALA:HB3	1.52	0.88
6:g:324:SER:O	6:g:327:THR:HG22	1.73	0.88
6:g:360:ALA:CA	9:D:293:PRO:CD	2.47	0.88
9:B:411:ASP:CA	9:B:418:ARG:NH2	2.30	0.88
2:b:107:LYS:CE	4:p:166:THR:HA	2.02	0.88
4:p:203:LEU:HD22	8:A:7:ASP:O	1.75	0.88
4:p:210:LEU:CD2	8:A:13:ILE:CG1	2.46	0.87
7:P:43:PRO:HB2	7:Q:48:LYS:NZ	1.90	0.87
1:a:62:GLN:O	4:p:119:ARG:NH1	2.06	0.87
4:p:219:LEU:HB2	4:p:220:PRO:HD2	1.55	0.87
9:F:50:LYS:HE3	9:F:92:ILE:HD11	1.56	0.87
2:b:125:LYS:HA	4:p:185:LEU:CD2	2.05	0.87
4:p:210:LEU:O	4:p:213:ASP:HB3	1.75	0.87
1:a:35:ILE:HG21	4:p:88:LEU:CB	2.04	0.87
1:a:47:ILE:HG13	4:p:96:GLU:HG2	1.57	0.87
1:a:212:VAL:HB	1:a:213:PRO:HD3	1.56	0.87
2:b:88:VAL:CG2	4:p:145:ASN:HA	2.02	0.87
6:g:356:ILE:CG2	8:E:286:ALA:HB2	2.03	0.87
6:g:363:CYS:HB3	9:D:290:GLY:O	1.75	0.87
7:M:48:LYS:CG	7:L:46:GLU:OE1	2.22	0.87
1:a:136:THR:N	4:p:90:LEU:CD2	2.36	0.87
4:p:203:LEU:HD22	8:A:7:ASP:C	1.98	0.87
6:g:239:ILE:HD12	6:g:303:GLN:HE22	1.28	0.87
7:S:68:LEU:CD1	7:T:66:TYR:CG	2.58	0.87
7:G:46:GLU:OE1	7:H:48:LYS:HB3	1.74	0.87
1:a:184:LEU:HD23	1:a:188:PHE:CD2	2.09	0.87
6:g:123:VAL:HG13	6:g:308:LEU:HD22	1.56	0.87
6:g:221:VAL:CG2	8:A:401:SER:HB3	2.04	0.87
7:M:74:LEU:HD22	7:L:10:VAL:CG2	2.02	0.87
7:M:80:PHE:CE2	7:L:81:VAL:HG21	2.09	0.87
9:D:275:ILE:HG22	9:D:327:VAL:HG22	1.57	0.87
1:a:53:SER:O	1:a:56:ILE:CG2	2.22	0.87
4:p:214:ILE:CB	8:A:17:ILE:HD12	2.03	0.87
7:O:10:VAL:HG23	7:P:74:LEU:CD2	2.05	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:144:ALA:O	4:p:147:VAL:HG12	1.73	0.87
1:a:135:ASN:HD21	4:p:86:PHE:CB	1.88	0.86
2:b:133:GLN:O	2:b:137:ILE:CD1	2.22	0.86
7:H:40:ALA:HB2	7:I:49:ILE:HD11	1.57	0.86
8:A:111:ALA:CA	8:A:227:GLU:OE2	2.22	0.86
2:b:103:ILE:HD13	4:p:162:MET:CB	2.04	0.86
7:Q:10:VAL:CG2	7:R:74:LEU:CD2	2.52	0.86
7:L:42:GLN:OE1	7:K:40:ALA:O	1.93	0.86
8:A:469:LYS:HD2	8:A:493:ALA:HB2	1.57	0.86
3:d:228:GLU:CD	8:A:43:HIS:NE2	2.32	0.86
9:D:113:VAL:HG22	9:D:249:VAL:HG13	1.57	0.86
1:a:161:PHE:CZ	1:a:163:LYS:CB	2.47	0.86
1:a:44:TRP:CD1	4:p:96:GLU:OE1	2.28	0.86
1:a:119:ILE:HD11	2:b:32:ILE:N	1.70	0.86
2:b:125:LYS:HD3	4:p:185:LEU:HD22	1.46	0.86
3:d:247:GLN:NE2	8:A:13:ILE:HD13	1.90	0.86
7:N:74:LEU:CD2	7:M:10:VAL:CG2	2.53	0.86
2:b:107:LYS:HA	4:p:166:THR:HG23	1.55	0.86
4:p:210:LEU:CD1	8:A:13:ILE:CG2	2.54	0.86
6:g:104:ASP:HA	7:S:41:ARG:HH21	1.41	0.86
7:Q:43:PRO:CB	7:R:48:LYS:NZ	2.38	0.86
7:K:48:LYS:HB3	7:J:46:GLU:OE1	1.75	0.86
8:C:77:MET:HE2	8:C:112:LEU:HD21	1.56	0.86
9:D:100:VAL:CG1	9:D:256:MET:HE3	2.06	0.86
1:a:24:GLN:C	4:p:83:LEU:HG	2.00	0.86
1:a:26:PHE:HB3	4:p:83:LEU:HA	0.86	0.86
1:a:100:THR:HB	4:p:97:PHE:CE2	2.09	0.86
1:a:214:ILE:CG1	1:a:215:PRO:HD3	2.05	0.86
2:b:92:ALA:HB3	4:p:147:VAL:O	1.73	0.86
2:b:141:ARG:CA	8:A:9:ILE:HD12	2.06	0.86
2:b:144:VAL:HG22	4:p:207:ILE:HD11	1.55	0.86
9:F:202:VAL:HG22	9:F:249:VAL:HG23	1.57	0.86
1:a:119:ILE:CG1	2:b:28:ALA:HA	2.04	0.86
2:b:125:LYS:HD2	4:p:185:LEU:HD23	0.88	0.86
6:g:221:VAL:CG2	8:A:401:SER:CB	2.53	0.86
9:B:410:LEU:C	9:B:418:ARG:NH2	2.33	0.86
9:D:338:ALA:HB3	9:D:339:PRO:HD3	1.57	0.86
1:a:70:ASN:C	2:b:53:LEU:CB	2.48	0.86
3:d:242:GLU:HG2	8:A:20:TYR:CZ	2.11	0.86
7:S:43:PRO:HG2	7:T:42:GLN:HE22	1.41	0.86
7:N:68:LEU:HD13	7:O:66:TYR:HB3	1.57	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:O:10:VAL:CG2	7:P:74:LEU:HD22	2.04	0.86
7:P:40:ALA:HB2	7:Q:49:ILE:HD11	1.57	0.86
1:a:119:ILE:CG2	2:b:28:ALA:CA	2.07	0.86
8:E:386:GLU:OE1	8:E:413:ARG:HD3	1.73	0.86
8:E:464:TYR:OH	8:E:496:GLU:OE1	1.94	0.86
8:A:432:GLU:O	8:A:435:VAL:HG12	1.76	0.86
1:a:118:LYS:CB	2:b:29:THR:C	2.47	0.85
6:g:359:GLY:CA	9:D:293:PRO:CG	2.53	0.85
7:N:48:LYS:HB3	7:M:46:GLU:OE1	1.76	0.85
7:Q:40:ALA:HB2	7:R:49:ILE:HD11	1.58	0.85
2:b:95:PHE:CD2	4:p:151:ALA:C	2.18	0.85
7:T:46:GLU:OE1	7:G:48:LYS:HB3	1.74	0.85
7:H:43:PRO:CB	7:I:48:LYS:NZ	2.38	0.85
8:E:344:PHE:HD2	8:E:362:ILE:HG21	0.90	0.85
8:E:355:ARG:HD3	10:E:600:ATP:C2	2.11	0.85
2:b:100:TYR:O	2:b:103:ILE:HG22	1.76	0.85
4:p:195:GLN:HG2	8:A:4:ILE:HD11	0.87	0.85
8:E:82:MET:HA	8:E:82:MET:CE	2.05	0.85
8:E:413:ARG:HH12	8:E:444:GLY:N	1.74	0.85
8:C:237:TYR:CE1	8:C:271:TYR:CD1	2.64	0.85
4:p:156:SER:O	4:p:159:LEU:HG	1.77	0.85
7:G:40:ALA:HB2	7:H:49:ILE:HD11	1.58	0.85
1:a:39:VAL:CG2	1:a:133:ASP:OD2	2.25	0.85
2:b:80:LYS:CG	4:p:137:VAL:HB	2.06	0.85
4:p:206:GLN:O	4:p:210:LEU:HB3	1.77	0.85
1:a:116:PRO:HB3	2:b:34:LEU:N	1.92	0.85
3:d:72:ASP:OD1	3:d:77:ARG:NE	2.09	0.85
4:p:127:LEU:O	4:p:130:VAL:HG12	1.75	0.85
6:g:360:ALA:HA	9:D:293:PRO:HD2	1.56	0.85
2:b:87:LYS:O	2:b:90:MET:HB3	1.76	0.85
5:e:19:VAL:CA	5:e:53:LEU:HD13	2.06	0.85
7:G:10:VAL:CG2	7:H:74:LEU:HD22	2.07	0.85
8:A:464:TYR:CD1	8:A:468:ASN:ND2	2.45	0.85
2:b:61:LEU:CD1	2:b:65:ARG:CD	2.55	0.85
5:e:38:ALA:HB3	7:L:41:ARG:CG	2.01	0.85
7:S:42:GLN:HE22	7:R:43:PRO:HG2	1.42	0.85
7:S:42:GLN:OE1	7:R:40:ALA:O	1.94	0.85
9:F:50:LYS:HD2	9:F:90:GLU:CD	2.02	0.85
2:b:176:PHE:CD1	3:d:233:VAL:CG2	2.44	0.84
1:a:193:ASN:ND2	1:a:227:GLN:CG	2.40	0.84
2:b:122:GLU:HA	4:p:181:ILE:HG21	1.58	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:136:SER:OG	3:d:138:LEU:HD21	1.76	0.84
1:a:24:GLN:CB	4:p:84:PHE:CE2	2.20	0.84
2:b:128:THR:HG22	4:p:189:LEU:HD21	1.56	0.84
7:N:48:LYS:NZ	7:M:43:PRO:CB	2.39	0.84
8:C:54:VAL:HG12	8:C:89:VAL:HG22	1.59	0.84
2:b:125:LYS:HB3	4:p:185:LEU:CG	2.06	0.84
2:b:125:LYS:O	4:p:185:LEU:HD13	1.76	0.84
7:S:74:LEU:CD2	7:R:10:VAL:CG2	2.55	0.84
7:N:46:GLU:OE1	7:O:48:LYS:CG	2.25	0.84
8:A:214:GLN:NE2	8:A:219:MET:HB3	1.92	0.84
1:a:134:ILE:CA	1:a:191:PHE:CE1	2.59	0.84
2:b:69:GLU:N	4:p:130:VAL:HG11	1.65	0.84
2:b:121:PHE:CZ	2:b:125:LYS:HE2	2.12	0.84
4:p:163:LYS:O	4:p:166:THR:HG22	1.78	0.84
6:g:129:LEU:HD23	9:B:412:GLU:CD	2.02	0.84
7:G:68:LEU:HD13	7:H:66:TYR:HB3	1.59	0.84
8:E:369:SER:O	8:E:377:LYS:HE2	1.77	0.84
8:C:104:TYR:CE1	8:C:122:ILE:HD13	2.12	0.84
3:d:76:SER:HA	3:d:164:GLU:HG2	1.59	0.84
6:g:104:ASP:CA	7:S:41:ARG:NH2	2.41	0.84
7:S:48:LYS:NZ	7:R:43:PRO:HB2	1.93	0.84
2:b:73:LYS:O	4:p:134:SER:HA	1.74	0.84
8:C:237:TYR:HD1	8:C:271:TYR:HB2	1.39	0.84
2:b:61:LEU:CD1	2:b:65:ARG:HD2	2.07	0.84
2:b:148:ALA:CA	4:p:211:SER:HA	2.08	0.84
3:d:152:SER:O	3:d:153:GLU:CD	2.20	0.84
7:O:40:ALA:HB2	7:P:49:ILE:HD11	1.60	0.84
7:J:48:LYS:HZ3	7:I:43:PRO:CB	1.90	0.84
1:a:137:THR:CG2	1:a:191:PHE:HB2	2.08	0.84
2:b:132:GLU:HG3	4:p:189:LEU:CD2	2.06	0.84
6:g:221:VAL:HG21	8:A:401:SER:CB	2.08	0.84
9:F:385:TYR:CE1	9:F:389:GLN:NE2	2.46	0.83
1:a:119:ILE:HD13	2:b:31:LEU:CD1	2.08	0.83
1:a:122:LEU:HD23	1:a:124:HIS:N	1.91	0.83
2:b:84:ARG:CB	4:p:144:ALA:HB3	1.89	0.83
3:d:235:MET:HE2	8:A:22:ARG:HH21	1.43	0.83
6:g:353:ILE:CD1	8:E:284:ARG:HA	2.07	0.83
7:H:10:VAL:CG2	7:I:74:LEU:CD2	2.56	0.83
9:F:336:ASP:OD1	9:F:337:PRO:HD2	1.79	0.83
1:a:54:ALA:HB1	4:p:104:LEU:CD1	2.08	0.83
1:a:119:ILE:HA	2:b:28:ALA:C	2.02	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:122:GLU:CA	4:p:181:ILE:HG21	2.08	0.83
2:b:125:LYS:CB	4:p:185:LEU:CD2	2.56	0.83
6:g:356:ILE:HG21	8:E:286:ALA:CB	2.05	0.83
1:a:119:ILE:HD13	2:b:31:LEU:CG	2.08	0.83
2:b:80:LYS:C	4:p:140:LEU:CB	2.49	0.83
2:b:148:ALA:CB	4:p:211:SER:HA	2.09	0.83
7:G:10:VAL:HG23	7:H:74:LEU:CD2	2.08	0.83
9:B:345:HIS:O	8:C:202:LYS:HD3	1.77	0.83
2:b:78:LEU:CD1	4:p:139:GLN:OE1	2.26	0.83
8:A:152:ILE:H	8:A:423:GLN:HE22	1.26	0.83
1:a:70:ASN:O	2:b:53:LEU:CD2	2.25	0.83
2:b:82:ARG:CA	4:p:140:LEU:HD23	2.08	0.83
2:b:107:LYS:N	4:p:166:THR:HG21	1.94	0.83
2:b:144:VAL:HG12	4:p:210:LEU:HD23	1.04	0.83
4:p:192:LEU:HD22	8:A:3:THR:CG2	2.08	0.83
6:g:202:PHE:CG	6:g:231:LEU:HD13	2.13	0.83
8:A:157:MET:SD	8:A:387:LEU:HD12	2.18	0.83
1:a:24:GLN:CA	4:p:83:LEU:CD1	2.56	0.83
1:a:65:PRO:CG	1:a:70:ASN:HD21	1.91	0.83
5:e:110:ARG:NH2	6:g:201:LEU:HD13	1.93	0.83
6:g:279:SER:OG	6:g:282:LEU:HB2	1.78	0.83
7:S:74:LEU:HD22	7:R:10:VAL:CG2	2.08	0.83
7:Q:43:PRO:CB	7:R:48:LYS:HZ3	1.91	0.83
8:A:214:GLN:HE21	8:A:219:MET:HG2	0.98	0.83
7:J:66:TYR:HB3	7:I:68:LEU:HD13	1.60	0.83
1:a:119:ILE:HG13	2:b:28:ALA:O	1.78	0.83
3:d:181:THR:O	3:d:221:PHE:CB	2.26	0.83
3:d:198:VAL:HA	3:d:201:ILE:HD11	1.59	0.83
4:p:195:GLN:CD	8:A:4:ILE:HD11	2.04	0.83
2:b:64:ILE:CD1	4:p:120:ASP:HA	2.07	0.83
2:b:125:LYS:C	4:p:185:LEU:HD13	2.03	0.83
7:S:43:PRO:HB3	7:T:48:LYS:NZ	1.94	0.83
8:E:37:ASP:CG	9:D:291:ARG:HH21	1.86	0.83
8:A:420:LEU:HD11	8:A:441:GLY:HA3	1.59	0.83
1:a:114:LEU:CA	2:b:37:VAL:HB	2.06	0.82
1:a:119:ILE:HD13	2:b:31:LEU:HD12	1.60	0.82
2:b:80:LYS:H	4:p:137:VAL:HA	1.44	0.82
2:b:82:ARG:CA	2:b:85:LEU:CD2	2.57	0.82
4:p:210:LEU:CD2	8:A:13:ILE:HB	2.08	0.82
7:Q:46:GLU:OE1	7:R:48:LYS:HB3	1.78	0.82
7:K:42:GLN:NE2	7:J:43:PRO:HG2	1.94	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:76:LEU:HD13	4:p:109:TYR:CD1	2.14	0.82
3:d:186:LEU:HD11	3:d:191:LEU:HG	1.61	0.82
6:g:292:LEU:HD12	6:g:292:LEU:O	1.79	0.82
8:E:37:ASP:CG	9:D:291:ARG:NH2	2.36	0.82
1:a:119:ILE:CG2	2:b:27:LEU:C	2.52	0.82
6:g:251:GLU:CG	6:g:272:LYS:HG2	2.09	0.82
1:a:114:LEU:HA	2:b:37:VAL:CB	2.01	0.82
1:a:119:ILE:CD1	2:b:31:LEU:HG	2.09	0.82
3:d:247:GLN:O	8:A:16:ARG:HD3	1.79	0.82
4:p:152:ARG:CG	8:A:503:LEU:CB	2.58	0.82
6:g:53:SER:CB	9:D:403:ASP:OD2	2.27	0.82
9:B:169:GLY:HA3	9:B:346:LEU:HD13	1.59	0.82
1:a:119:ILE:HD13	2:b:31:LEU:HG	1.61	0.82
1:a:119:ILE:HD11	2:b:32:ILE:CB	2.10	0.82
4:p:213:ASP:OD2	8:A:14:ARG:CA	2.28	0.82
5:e:34:LEU:HB2	7:L:42:GLN:HE22	1.43	0.82
7:O:68:LEU:HD13	7:P:66:TYR:HB3	1.60	0.82
7:Q:43:PRO:HG2	7:R:42:GLN:NE2	1.93	0.82
1:a:119:ILE:CG1	2:b:28:ALA:CA	2.57	0.82
7:J:80:PHE:CE2	7:I:81:VAL:HG21	2.14	0.82
9:B:20:LEU:HD13	9:B:90:GLU:CD	2.04	0.82
8:C:237:TYR:OH	8:C:294:LEU:HD13	1.80	0.82
1:a:35:ILE:CG2	4:p:88:LEU:CB	2.57	0.82
2:b:141:ARG:N	8:A:9:ILE:HD12	1.95	0.82
3:d:86:ALA:CB	3:d:95:THR:HG21	2.09	0.82
3:d:154:ARG:HH21	8:C:5:ARG:NE	1.77	0.82
7:N:81:VAL:HG21	7:O:80:PHE:CE2	2.15	0.82
1:a:39:VAL:HG11	1:a:133:ASP:OD2	1.80	0.81
1:a:164:TYR:CE2	1:a:166:GLN:HG2	2.13	0.81
1:a:135:ASN:OD1	4:p:86:PHE:HB3	1.79	0.81
4:p:192:LEU:CD2	8:A:3:THR:OG1	2.28	0.81
6:g:360:ALA:N	9:D:293:PRO:CG	2.42	0.81
7:N:74:LEU:CD2	7:M:10:VAL:HG23	2.10	0.81
7:J:62:ALA:C	7:J:65:ILE:HG22	2.05	0.81
8:A:154:ILE:CD1	8:A:358:ILE:HD11	2.10	0.81
1:a:25:HIS:O	4:p:83:LEU:HB3	1.80	0.81
1:a:70:ASN:O	2:b:53:LEU:HD22	1.80	0.81
1:a:122:LEU:HA	2:b:25:ASP:HB2	1.61	0.81
1:a:222:PHE:CE1	7:Q:54:LEU:HD23	2.15	0.81
2:b:48:VAL:O	2:b:52:LEU:HD13	1.81	0.81
2:b:68:GLU:CD	4:p:126:GLN:HG3	2.05	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:129:ILE:HD11	4:p:188:ALA:HB1	1.59	0.81
7:S:49:ILE:HD11	7:R:40:ALA:HB2	1.62	0.81
7:R:62:ALA:C	7:R:65:ILE:HG22	2.05	0.81
7:T:43:PRO:C	7:G:48:LYS:HZ3	1.89	0.81
7:J:48:LYS:HB3	7:I:46:GLU:OE1	1.81	0.81
9:F:387:ILE:HD11	9:F:458:PHE:CB	2.09	0.81
2:b:81:ALA:HB3	4:p:139:GLN:C	1.94	0.81
2:b:84:ARG:HG2	4:p:144:ALA:HB3	1.61	0.81
6:g:353:ILE:HD11	8:E:284:ARG:CG	2.10	0.81
7:S:74:LEU:CD2	7:R:10:VAL:HG23	2.10	0.81
7:M:74:LEU:HD21	7:L:10:VAL:HG23	1.61	0.81
7:T:43:PRO:HG2	7:G:42:GLN:NE2	1.94	0.81
1:a:54:ALA:HB1	4:p:104:LEU:HD11	1.60	0.81
1:a:134:ILE:CG2	1:a:191:PHE:CD2	2.47	0.81
6:g:221:VAL:HG21	8:A:398:GLN:O	1.80	0.81
8:E:42:ILE:HD11	8:E:56:PHE:CE1	2.16	0.81
9:B:20:LEU:HD22	9:B:92:ILE:HG12	1.60	0.81
1:a:124:HIS:CD2	4:p:80:LYS:O	2.33	0.81
7:S:64:THR:HG21	7:T:63:LEU:CD2	2.10	0.81
7:P:62:ALA:C	7:P:65:ILE:HG22	2.06	0.81
7:Q:62:ALA:C	7:Q:65:ILE:HG22	2.05	0.81
3:d:132:ILE:HA	8:E:13:ILE:CD1	2.10	0.81
4:p:87:ASN:OD1	4:p:89:THR:HG22	1.79	0.81
7:H:62:ALA:C	7:H:65:ILE:HG22	2.06	0.81
7:K:48:LYS:HZ3	7:J:43:PRO:C	1.88	0.81
9:F:50:LYS:HE2	9:F:92:ILE:HD11	1.45	0.81
9:D:53:ASP:HB3	9:D:59:MET:CE	2.10	0.81
2:b:85:LEU:HD23	4:p:143:GLN:CG	2.10	0.81
2:b:103:ILE:CG2	4:p:162:MET:HB2	2.11	0.81
4:p:167:GLN:O	4:p:171:GLU:CB	2.27	0.81
6:g:240:CYS:HB2	6:g:246:CYS:SG	2.20	0.81
7:O:46:GLU:OE1	7:P:48:LYS:CG	2.28	0.81
7:O:62:ALA:C	7:O:65:ILE:HG22	2.06	0.81
7:T:62:ALA:C	7:T:65:ILE:HG22	2.06	0.81
7:H:46:GLU:OE1	7:I:48:LYS:CG	2.28	0.81
8:E:106:GLY:HA2	8:E:219:MET:HE3	1.63	0.81
2:b:57:LYS:O	2:b:60:ILE:HG22	1.81	0.81
2:b:145:PHE:CA	8:A:13:ILE:HD12	2.10	0.81
2:b:147:GLN:OE1	4:p:207:ILE:CG2	2.29	0.81
5:e:46:ILE:HG22	5:e:63:MET:C	2.06	0.81
7:M:62:ALA:C	7:M:65:ILE:HG22	2.06	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:L:48:LYS:HB3	7:K:46:GLU:OE1	1.81	0.81
7:I:62:ALA:C	7:I:65:ILE:HG22	2.05	0.81
7:L:62:ALA:C	7:L:65:ILE:HG22	2.06	0.81
9:B:20:LEU:HD21	9:B:92:ILE:HG12	1.62	0.81
8:C:78:GLY:HA2	8:C:232:PRO:HG3	1.63	0.81
3:d:179:VAL:HG22	3:d:210:LYS:CB	2.10	0.80
8:A:193:ILE:O	8:A:193:ILE:HG22	1.77	0.80
1:a:112:GLY:HA2	1:a:128:ALA:O	1.81	0.80
2:b:161:ASN:O	2:b:165:HIS:HB2	1.80	0.80
3:d:177:VAL:HG11	3:d:210:LYS:HD2	1.62	0.80
4:p:152:ARG:CB	8:A:503:LEU:CD2	2.52	0.80
7:N:46:GLU:OE1	7:O:48:LYS:CB	2.29	0.80
7:G:62:ALA:C	7:G:65:ILE:HG22	2.06	0.80
1:a:139:ALA:CB	4:p:94:MET:CE	2.59	0.80
2:b:141:ARG:HA	8:A:9:ILE:HD12	1.63	0.80
5:e:50:ARG:HG2	5:e:59:THR:HG22	1.61	0.80
1:a:184:LEU:CD2	1:a:188:PHE:CD2	2.64	0.80
4:p:93:ILE:HD12	4:p:96:GLU:HB3	1.62	0.80
5:e:38:ALA:N	7:L:41:ARG:O	2.14	0.80
7:S:48:LYS:HZ2	7:R:43:PRO:CB	1.94	0.80
4:p:219:LEU:HD12	4:p:220:PRO:O	1.82	0.80
7:O:46:GLU:OE1	7:P:48:LYS:CB	2.29	0.80
7:H:46:GLU:OE1	7:I:48:LYS:CB	2.30	0.80
7:K:62:ALA:C	7:K:65:ILE:HG22	2.06	0.80
2:b:145:PHE:CZ	3:d:243:GLU:O	2.34	0.80
5:e:34:LEU:CB	7:L:42:GLN:NE2	2.39	0.80
6:g:359:GLY:C	9:D:293:PRO:CG	2.54	0.80
8:C:432:GLU:HG3	8:C:476:ILE:HG23	1.61	0.80
1:a:53:SER:C	1:a:56:ILE:HG22	2.05	0.80
3:d:242:GLU:CG	8:A:20:TYR:CZ	2.64	0.80
4:p:195:GLN:CD	8:A:4:ILE:CD1	2.55	0.80
4:p:212:ASP:C	4:p:215:VAL:HG12	2.07	0.80
6:g:281:ILE:CA	7:N:41:ARG:HH21	1.90	0.80
6:g:362:ALA:CB	8:C:281:PRO:HB3	2.12	0.80
7:N:74:LEU:HD22	7:M:10:VAL:CG2	2.10	0.80
2:b:80:LYS:HB2	4:p:137:VAL:CB	2.07	0.80
6:g:119:ALA:HB3	6:g:210:VAL:HG22	1.63	0.80
6:g:251:GLU:CB	6:g:272:LYS:HG2	2.12	0.80
7:S:62:ALA:C	7:S:65:ILE:HG22	2.06	0.80
7:L:49:ILE:HD11	7:K:40:ALA:HB2	1.62	0.80
4:p:159:LEU:O	4:p:163:LYS:N	2.11	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:N:62:ALA:C	7:N:65:ILE:HG22	2.06	0.80
8:A:492:GLU:O	8:A:496:GLU:CG	2.30	0.80
9:B:71:ASN:HD21	8:C:9:ILE:HA	1.47	0.80
9:B:245:ALA:O	9:B:249:VAL:HG23	1.82	0.80
1:a:35:ILE:HG22	4:p:88:LEU:CA	2.11	0.80
2:b:78:LEU:O	4:p:140:LEU:CG	2.28	0.80
7:O:43:PRO:CB	7:P:48:LYS:NZ	2.44	0.80
7:L:42:GLN:NE2	7:K:43:PRO:HG2	1.97	0.80
8:E:37:ASP:OD2	9:D:291:ARG:CZ	2.30	0.80
8:C:100:VAL:HG23	8:C:246:LEU:HD23	1.62	0.80
3:d:148:ILE:HG12	8:C:16:ARG:HG2	1.62	0.79
7:S:10:VAL:CG2	7:T:74:LEU:CD2	2.59	0.79
7:T:10:VAL:CG2	7:G:74:LEU:CD2	2.61	0.79
8:E:202:LYS:HD3	9:D:345:HIS:O	1.82	0.79
8:E:355:ARG:NH1	9:D:385:TYR:OH	2.15	0.79
1:a:34:GLN:O	4:p:87:ASN:CB	2.26	0.79
1:a:135:ASN:CG	4:p:86:PHE:HA	2.05	0.79
4:p:210:LEU:CD1	8:A:13:ILE:CB	2.52	0.79
7:N:10:VAL:HG23	7:O:74:LEU:HD21	1.63	0.79
8:A:98:ILE:HD13	8:A:245:ALA:CB	2.09	0.79
9:D:191:LYS:NZ	9:D:219:SER:HB2	1.97	0.79
2:b:71:ARG:O	4:p:133:THR:OG1	1.99	0.79
2:b:80:LYS:N	4:p:137:VAL:CA	2.45	0.79
3:d:180:VAL:HG22	3:d:223:ILE:HG22	1.63	0.79
6:g:132:GLY:O	6:g:136:MET:HG2	1.80	0.79
8:A:382:LYS:HD3	8:A:442:THR:HG22	1.63	0.79
9:B:311:GLU:HG3	8:C:230:ASP:HB3	1.64	0.79
2:b:125:LYS:CB	4:p:185:LEU:HB2	2.10	0.79
6:g:229:THR:HG23	6:g:229:THR:O	1.82	0.79
7:P:46:GLU:OE1	7:Q:48:LYS:HB3	1.80	0.79
7:Q:6:ALA:HB1	7:R:80:PHE:HD2	1.45	0.79
9:B:71:ASN:HD21	8:C:9:ILE:C	1.90	0.79
1:a:61:PRO:HB3	4:p:112:LEU:CD2	2.12	0.79
2:b:68:GLU:HG3	4:p:126:GLN:CB	1.92	0.79
2:b:73:LYS:O	4:p:134:SER:CA	2.28	0.79
3:d:72:ASP:N	8:A:70:ASN:HD21	1.80	0.79
3:d:181:THR:O	3:d:221:PHE:HB2	1.82	0.79
5:e:38:ALA:CB	7:L:41:ARG:HA	2.03	0.79
7:P:68:LEU:HD13	7:Q:66:TYR:HB3	1.65	0.79
8:C:439:TYR:CD2	8:C:490:LEU:HD23	2.17	0.79
9:D:100:VAL:HG11	9:D:256:MET:HE3	1.63	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:114:LEU:HD13	2:b:37:VAL:C	2.08	0.79
2:b:147:GLN:OE1	4:p:207:ILE:HG23	1.83	0.79
6:g:137:LEU:HD11	6:g:213:LEU:HD13	1.57	0.79
7:M:48:LYS:HB3	7:L:46:GLU:CD	2.07	0.79
7:K:48:LYS:HZ2	7:J:43:PRO:HB2	1.48	0.79
8:E:104:TYR:CE1	8:E:122:ILE:CD1	2.66	0.79
8:A:494:ILE:CG2	8:A:498:MET:CE	2.57	0.79
1:a:135:ASN:OD1	4:p:86:PHE:CD2	2.36	0.79
2:b:75:ILE:CG1	4:p:132:ASP:C	2.42	0.79
2:b:103:ILE:HD11	4:p:163:LYS:H	1.46	0.79
2:b:144:VAL:HG11	8:A:13:ILE:HG13	1.63	0.79
3:d:222:THR:HG22	3:d:234:ASP:CA	2.09	0.79
6:g:362:ALA:HB1	8:C:281:PRO:CB	2.12	0.79
7:T:46:GLU:OE1	7:G:48:LYS:CG	2.31	0.79
7:K:48:LYS:CG	7:J:46:GLU:OE1	2.31	0.79
5:e:22:ILE:HG12	5:e:33:VAL:HG21	1.64	0.79
6:g:356:ILE:HD13	8:E:286:ALA:HB2	1.65	0.79
8:E:82:MET:HE2	8:E:82:MET:CA	2.12	0.79
8:E:302:ALA:HB2	8:E:314:MET:HE2	1.64	0.79
8:A:440:THR:HG22	8:A:494:ILE:HD11	1.65	0.79
1:a:116:PRO:CG	2:b:35:SER:H	1.95	0.79
3:d:148:ILE:CD1	8:C:17:ILE:HG12	2.13	0.79
3:d:228:GLU:OE1	8:A:43:HIS:CE1	2.35	0.79
4:p:152:ARG:HD3	8:A:503:LEU:CB	2.13	0.79
6:g:239:ILE:CD1	6:g:303:GLN:HE22	1.86	0.79
6:g:363:CYS:SG	9:D:291:ARG:O	2.41	0.79
7:J:49:ILE:HD11	7:I:40:ALA:HB2	1.65	0.79
9:D:478:GLY:N	9:D:482:GLU:OE1	2.15	0.79
8:E:104:TYR:HD1	8:E:122:ILE:HD13	1.42	0.78
8:C:237:TYR:HE1	8:C:294:LEU:CD2	1.96	0.78
2:b:75:ILE:O	4:p:137:VAL:CG1	2.30	0.78
3:d:169:PHE:O	3:d:172:ILE:HG22	1.83	0.78
6:g:103:GLU:HG2	7:R:41:ARG:CZ	2.12	0.78
9:D:191:LYS:HE3	9:D:219:SER:HB2	1.64	0.78
7:G:43:PRO:CB	7:H:48:LYS:HZ3	1.97	0.78
9:F:385:TYR:HE1	9:F:389:GLN:NE2	1.80	0.78
9:B:221:VAL:CG1	9:B:232:VAL:HG23	2.13	0.78
2:b:77:GLN:HB3	4:p:134:SER:CA	2.12	0.78
2:b:78:LEU:HG	4:p:139:GLN:HB2	1.62	0.78
7:N:49:ILE:HD11	7:M:40:ALA:HB2	1.65	0.78
7:P:43:PRO:HB3	7:Q:48:LYS:NZ	1.98	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:344:PHE:CD2	8:A:362:ILE:CG2	2.66	0.78
8:C:382:LYS:HG2	9:D:442:THR:O	1.84	0.78
1:a:134:ILE:CA	1:a:191:PHE:CD1	2.67	0.78
1:a:230:ILE:CD1	7:Q:55:LEU:CD1	2.44	0.78
3:d:235:MET:CE	3:d:239:LYS:HG2	2.13	0.78
2:b:160:LEU:HD13	2:b:164:LEU:H	1.01	0.78
5:e:34:LEU:CD1	7:K:43:PRO:HB2	2.14	0.78
6:g:240:CYS:HB3	6:g:246:CYS:CB	2.13	0.78
7:S:46:GLU:OE1	7:T:48:LYS:HB3	1.83	0.78
1:a:116:PRO:CG	2:b:35:SER:N	2.47	0.78
7:P:10:VAL:CG2	7:Q:74:LEU:CD2	2.60	0.78
8:A:109:ILE:CD1	8:A:113:ALA:HA	2.13	0.78
1:a:114:LEU:HB3	2:b:37:VAL:C	2.08	0.78
2:b:67:SER:OG	4:p:128:SER:N	2.17	0.78
6:g:91:GLU:HG3	6:g:276:PRO:HD2	1.64	0.78
7:L:48:LYS:NZ	7:K:43:PRO:CB	2.47	0.78
8:E:461:LEU:HD23	8:E:461:LEU:O	1.82	0.78
8:A:13:ILE:HD11	8:A:16:ARG:HH22	1.41	0.78
8:A:154:ILE:HD12	8:A:358:ILE:CD1	2.14	0.78
9:B:216:MET:HE2	9:B:232:VAL:HG11	1.66	0.78
3:d:132:ILE:CG1	8:E:13:ILE:HD13	2.14	0.78
7:T:43:PRO:HB3	7:G:48:LYS:HD2	1.65	0.78
9:F:383:GLU:OE2	9:F:459:GLN:NE2	2.16	0.78
8:C:104:TYR:CE1	8:C:122:ILE:HG21	2.18	0.78
2:b:70:LEU:CB	4:p:131:LYS:HB2	2.14	0.77
7:G:81:VAL:HG21	7:H:80:PHE:CE2	2.19	0.77
1:a:165:ILE:HG22	1:a:165:ILE:O	1.84	0.77
2:b:60:ILE:HG13	4:p:120:ASP:OD2	1.84	0.77
7:S:10:VAL:CG2	7:T:74:LEU:HD22	2.14	0.77
7:O:81:VAL:HG21	7:P:80:PHE:CE2	2.19	0.77
7:T:54:LEU:HD11	7:G:55:LEU:CD2	2.13	0.77
7:G:10:VAL:HG23	7:H:74:LEU:HD21	1.66	0.77
7:K:48:LYS:HD2	7:J:43:PRO:HB3	1.67	0.77
1:a:26:PHE:HB3	4:p:83:LEU:CB	2.14	0.77
1:a:104:PHE:CD1	4:p:97:PHE:CE1	2.71	0.77
7:S:48:LYS:HB3	7:R:46:GLU:OE1	1.84	0.77
1:a:100:THR:HB	4:p:97:PHE:HE2	1.47	0.77
1:a:118:LYS:HB2	2:b:29:THR:C	2.08	0.77
1:a:119:ILE:CA	2:b:29:THR:H	1.88	0.77
1:a:122:LEU:CD2	1:a:125:GLY:H	1.95	0.77
2:b:107:LYS:C	4:p:166:THR:HG1	1.79	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:136:SER:OG	3:d:138:LEU:CD2	2.32	0.77
7:S:64:THR:HG21	7:T:63:LEU:HD22	1.66	0.77
8:E:140:ARG:HE	9:F:206:THR:HG22	1.50	0.77
9:F:113:VAL:HG22	9:F:249:VAL:CG1	2.15	0.77
9:B:336:ASP:HB3	9:B:339:PRO:HD2	1.64	0.77
1:a:120:ILE:H	2:b:30:ASN:HB2	1.50	0.77
2:b:79:GLU:HG3	4:p:136:GLU:HG2	1.65	0.77
3:d:72:ASP:N	8:A:70:ASN:ND2	2.32	0.77
3:d:247:GLN:HE22	8:A:13:ILE:CG2	1.95	0.77
7:N:66:TYR:HB3	7:M:68:LEU:HD13	1.64	0.77
7:T:43:PRO:HB2	7:G:48:LYS:HZ2	1.49	0.77
4:p:152:ARG:CG	8:A:503:LEU:HB2	2.15	0.77
7:G:43:PRO:CB	7:H:48:LYS:NZ	2.46	0.77
7:J:42:GLN:NE2	7:I:43:PRO:HG2	2.00	0.77
9:B:402:GLN:HA	9:B:405:ILE:HG12	1.66	0.77
2:b:71:ARG:HB3	4:p:126:GLN:O	1.85	0.77
4:p:192:LEU:HD22	8:A:3:THR:CB	2.15	0.77
6:g:124:THR:HG22	6:g:137:LEU:HD23	0.88	0.77
7:O:10:VAL:HG23	7:P:74:LEU:HD21	1.65	0.77
2:b:88:VAL:HG11	4:p:145:ASN:C	2.10	0.77
5:e:34:LEU:HB2	7:L:42:GLN:HE21	1.47	0.77
7:K:48:LYS:CB	7:J:46:GLU:OE1	2.33	0.77
8:E:49:MET:HE3	8:E:95:ILE:HG23	1.67	0.77
1:a:118:LYS:NZ	1:a:126:GLU:HB2	2.00	0.77
3:d:195:ALA:O	3:d:207:VAL:HG11	1.84	0.77
6:g:111:LYS:HE3	6:g:113:ARG:CG	2.14	0.77
9:B:425:ARG:HH12	9:B:429:ARG:HH22	1.32	0.77
1:a:116:PRO:CA	2:b:34:LEU:H	1.97	0.77
3:d:148:ILE:HD11	8:C:17:ILE:N	1.98	0.77
6:g:221:VAL:CG1	8:A:401:SER:CB	2.63	0.77
7:J:74:LEU:HD21	7:I:10:VAL:HG23	1.63	0.77
9:B:212:LEU:HD12	9:B:212:LEU:O	1.84	0.77
9:B:296:VAL:HG12	9:B:297:GLY:H	1.50	0.77
1:a:195:LEU:HD22	4:p:84:PHE:HE1	1.49	0.76
2:b:103:ILE:HG13	4:p:163:LYS:H	1.46	0.76
8:E:379:VAL:CG1	8:E:438:ILE:HG22	2.14	0.76
2:b:141:ARG:HG2	8:A:9:ILE:CD1	2.15	0.76
1:a:171:LEU:O	1:a:171:LEU:HD23	1.84	0.76
5:e:36:ASN:O	7:L:43:PRO:HD2	1.86	0.76
8:A:382:LYS:CD	8:A:442:THR:CG2	2.58	0.76
8:A:494:ILE:HG23	8:A:498:MET:HE2	1.67	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:191:LYS:CE	9:D:219:SER:CB	2.52	0.76
1:a:121:GLN:HG3	2:b:25:ASP:OD2	1.84	0.76
2:b:148:ALA:HA	4:p:211:SER:CA	2.15	0.76
4:p:210:LEU:HD21	8:A:13:ILE:CG2	2.16	0.76
6:g:117:LYS:HE3	6:g:154:ASP:OD2	1.85	0.76
6:g:221:VAL:CG1	8:A:401:SER:HB2	2.15	0.76
7:O:6:ALA:HB1	7:P:80:PHE:HD2	1.50	0.76
9:B:216:MET:HB3	9:B:222:ILE:HG12	1.68	0.76
9:B:311:GLU:CG	8:C:230:ASP:HB3	2.15	0.76
8:C:383:LEU:HD12	8:C:383:LEU:O	1.85	0.76
2:b:80:LYS:H	4:p:137:VAL:CA	1.98	0.76
4:p:149:ARG:HA	8:A:503:LEU:HD21	1.68	0.76
8:E:56:PHE:CD1	8:E:89:VAL:HB	2.21	0.76
1:a:79:ILE:HD13	1:a:99:GLY:HA3	1.64	0.76
2:b:103:ILE:HD11	4:p:163:LYS:N	1.99	0.76
2:b:144:VAL:HG22	4:p:207:ILE:CD1	2.14	0.76
6:g:103:GLU:HG3	7:R:41:ARG:CD	2.16	0.76
8:A:469:LYS:HZ1	8:A:492:GLU:HB2	1.48	0.76
3:d:154:ARG:NE	8:C:5:ARG:NH1	2.33	0.76
7:P:43:PRO:HG2	7:Q:42:GLN:NE2	2.00	0.76
9:B:221:VAL:O	9:B:230:SER:O	2.03	0.76
8:C:308:LEU:HD23	8:C:309:LEU:HD12	1.66	0.76
1:a:184:LEU:HD21	1:a:188:PHE:CE2	2.21	0.76
3:d:235:MET:HE3	3:d:239:LYS:HG2	1.68	0.76
4:p:193:GLU:HA	4:p:196:LYS:HE2	1.68	0.76
7:H:43:PRO:HB2	7:I:48:LYS:NZ	2.00	0.76
8:C:308:LEU:CD2	8:C:309:LEU:HD12	2.15	0.76
2:b:88:VAL:O	4:p:148:MET:HA	1.86	0.76
4:p:108:TYR:CD1	4:p:112:LEU:CD1	2.66	0.76
4:p:203:LEU:HD11	4:p:206:GLN:HB2	1.66	0.76
7:S:10:VAL:HG23	7:T:74:LEU:CD2	2.15	0.76
7:N:43:PRO:HB2	7:O:48:LYS:NZ	2.00	0.76
7:P:10:VAL:CG2	7:Q:74:LEU:HD22	2.16	0.76
7:T:46:GLU:OE1	7:G:48:LYS:CB	2.33	0.76
8:C:430:THR:CG2	8:C:432:GLU:OE1	2.33	0.76
1:a:134:ILE:O	1:a:137:THR:N	2.18	0.75
7:K:74:LEU:CD2	7:J:10:VAL:CG2	2.64	0.75
8:A:382:LYS:HB3	8:A:442:THR:HG21	1.67	0.75
3:d:198:VAL:O	3:d:202:THR:OG1	2.04	0.75
4:p:214:ILE:HG13	4:p:215:VAL:N	2.01	0.75
6:g:359:GLY:C	9:D:293:PRO:HG3	2.11	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Q:10:VAL:CG2	7:R:74:LEU:HD22	2.16	0.75
7:S:43:PRO:HB2	7:T:48:LYS:HZ2	1.49	0.75
7:T:43:PRO:HB3	7:G:48:LYS:NZ	2.01	0.75
7:H:10:VAL:HG23	7:I:74:LEU:CD2	2.16	0.75
9:F:22:ARG:NH1	9:F:90:GLU:HB3	2.00	0.75
9:F:148:ILE:HD13	9:F:451:LEU:HD11	1.68	0.75
9:B:20:LEU:HD13	9:B:90:GLU:CG	2.16	0.75
9:D:455:ILE:O	9:D:459:GLN:HG3	1.86	0.75
2:b:68:GLU:CD	4:p:126:GLN:CB	2.54	0.75
2:b:80:LYS:HG2	4:p:137:VAL:CG1	2.10	0.75
2:b:81:ALA:HB3	4:p:140:LEU:H	1.49	0.75
2:b:98:ASN:N	4:p:155:ILE:HD13	2.02	0.75
7:K:55:LEU:CD2	7:J:54:LEU:HD11	2.16	0.75
9:D:191:LYS:HE2	9:D:219:SER:CB	2.12	0.75
1:a:119:ILE:HG23	2:b:27:LEU:C	2.11	0.75
2:b:118:LEU:HD22	4:p:177:GLY:HA3	1.69	0.75
2:b:129:ILE:CD1	4:p:188:ALA:HB1	2.14	0.75
5:e:34:LEU:CD1	7:K:43:PRO:CG	2.57	0.75
6:g:261:GLU:CG	9:F:397:ARG:CZ	2.54	0.75
8:A:445:TYR:CD2	8:A:494:ILE:HG21	2.21	0.75
8:C:9:ILE:HG21	8:C:12:ILE:HD12	1.67	0.75
9:D:64:GLU:OE1	9:D:248:ARG:NE	2.19	0.75
6:g:91:GLU:CG	6:g:276:PRO:HD2	2.17	0.75
7:S:48:LYS:HZ2	7:R:43:PRO:HB2	1.48	0.75
1:a:134:ILE:HA	1:a:191:PHE:CE1	2.22	0.75
3:d:247:GLN:C	8:A:16:ARG:CD	2.51	0.75
7:Q:10:VAL:HG23	7:R:74:LEU:CD2	2.17	0.75
7:H:68:LEU:HD13	7:I:66:TYR:HB3	1.67	0.75
7:K:48:LYS:NZ	7:J:43:PRO:HB3	2.01	0.75
8:A:98:ILE:CD1	8:A:245:ALA:HB1	2.14	0.75
8:A:109:ILE:HD11	8:A:113:ALA:CB	2.16	0.75
8:E:214:GLN:HG3	8:E:219:MET:HG3	1.68	0.75
8:C:432:GLU:CG	8:C:476:ILE:CG2	2.65	0.75
5:e:34:LEU:HD12	7:K:43:PRO:CB	2.16	0.75
7:S:48:LYS:NZ	7:R:43:PRO:HB3	2.01	0.75
7:O:43:PRO:CG	7:P:42:GLN:HE22	1.98	0.75
8:E:390:PHE:CE2	8:E:418:ARG:HD2	2.22	0.75
1:a:135:ASN:C	4:p:90:LEU:CD2	2.60	0.74
3:d:198:VAL:HA	3:d:201:ILE:CD1	2.15	0.74
6:g:111:LYS:HE3	6:g:113:ARG:HG3	1.69	0.74
7:S:54:LEU:HD11	7:T:55:LEU:CD2	2.17	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:163:LYS:HE2	1:a:179:ASP:CG	2.12	0.74
2:b:82:ARG:CA	2:b:85:LEU:HD23	2.12	0.74
7:N:48:LYS:NZ	7:M:43:PRO:HB2	2.02	0.74
8:E:469:LYS:HE3	8:E:493:ALA:N	2.01	0.74
9:F:197:SER:HG	9:F:271:PHE:HE1	1.35	0.74
8:C:164:GLN:HG2	8:C:165:ARG:N	2.01	0.74
8:C:169:ILE:HG13	8:C:320:VAL:O	1.87	0.74
1:a:24:GLN:HB3	4:p:84:PHE:HD2	0.96	0.74
2:b:103:ILE:HB	4:p:159:LEU:HB2	1.63	0.74
2:b:169:ILE:CG2	3:d:221:PHE:HE2	1.93	0.74
6:g:261:GLU:HG2	6:g:262:GLY:N	2.02	0.74
7:G:46:GLU:OE1	7:H:48:LYS:CG	2.35	0.74
9:B:110:ILE:HD11	9:B:213:TYR:CD1	2.22	0.74
8:C:157:MET:SD	8:C:387:LEU:HD12	2.27	0.74
2:b:71:ARG:NE	4:p:125:GLU:O	2.20	0.74
2:b:79:GLU:CA	4:p:136:GLU:HG3	2.16	0.74
7:P:10:VAL:HG23	7:Q:74:LEU:CD2	2.16	0.74
7:H:6:ALA:HB1	7:I:80:PHE:HD2	1.52	0.74
8:E:104:TYR:HE1	8:E:122:ILE:CD1	2.00	0.74
8:A:13:ILE:HD13	8:A:16:ARG:HH22	0.62	0.74
9:B:361:ILE:HG23	9:B:432:SER:HB3	1.69	0.74
8:C:9:ILE:HG22	8:C:12:ILE:CD1	2.09	0.74
9:D:336:ASP:OD1	9:D:337:PRO:HD2	1.87	0.74
2:b:176:PHE:HE1	3:d:233:VAL:HG21	0.94	0.74
4:p:127:LEU:O	4:p:130:VAL:CG1	2.36	0.74
2:b:75:ILE:C	4:p:137:VAL:HG13	2.06	0.74
8:E:208:GLN:NE2	9:D:144:THR:CG2	2.37	0.74
1:a:124:HIS:HB3	4:p:80:LYS:O	1.87	0.74
2:b:78:LEU:CA	4:p:135:SER:O	2.28	0.74
6:g:118:VAL:HG11	6:g:148:LEU:HD13	1.70	0.74
7:P:46:GLU:OE1	7:Q:48:LYS:CG	2.36	0.74
9:F:183:MET:HE1	9:F:215:GLU:HB3	1.68	0.74
6:g:349:ILE:HD13	8:E:284:ARG:NH2	1.98	0.74
7:S:43:PRO:C	7:T:48:LYS:HZ3	1.96	0.74
7:N:54:LEU:HD11	7:O:55:LEU:CD2	2.17	0.74
7:N:80:PHE:CE2	7:M:81:VAL:HG21	2.23	0.74
7:P:81:VAL:HG21	7:Q:80:PHE:CE2	2.23	0.74
7:Q:43:PRO:HB2	7:R:48:LYS:NZ	2.01	0.74
7:L:48:LYS:HZ3	7:K:43:PRO:CB	2.01	0.74
7:K:66:TYR:HB3	7:J:68:LEU:HD13	1.70	0.74
9:D:143:ASP:OD2	9:D:315:SER:HB2	1.86	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:70:ASN:HB3	2:b:53:LEU:HB3	1.68	0.74
6:g:113:ARG:NH1	6:g:205:GLU:CB	2.29	0.74
7:S:48:LYS:HZ3	7:R:43:PRO:CB	1.98	0.74
7:G:6:ALA:HB1	7:H:80:PHE:HD2	1.53	0.74
7:H:10:VAL:CG2	7:I:74:LEU:HD22	2.16	0.74
8:E:173:GLN:HG3	9:D:371:THR:HG21	1.68	0.74
1:a:70:ASN:HB3	2:b:53:LEU:CB	2.18	0.73
2:b:82:ARG:CA	4:p:140:LEU:CD2	2.61	0.73
7:N:74:LEU:HD21	7:M:10:VAL:HG23	1.70	0.73
8:E:137:ILE:HD13	9:F:110:ILE:HD13	1.70	0.73
1:a:35:ILE:CG2	4:p:88:LEU:CA	2.64	0.73
2:b:122:GLU:HB2	4:p:181:ILE:HD12	1.70	0.73
4:p:108:TYR:CE1	4:p:112:LEU:HD11	2.21	0.73
4:p:219:LEU:CD1	4:p:220:PRO:O	2.36	0.73
8:A:153:ALA:HB3	8:A:358:ILE:HD12	1.68	0.73
8:A:185:LEU:HD13	8:A:216:ARG:HD2	1.70	0.73
8:A:494:ILE:O	8:A:498:MET:HG3	1.88	0.73
5:e:19:VAL:HG21	5:e:51:ILE:HG12	1.70	0.73
6:g:253:GLU:HB2	6:g:269:ASP:C	2.14	0.73
7:J:48:LYS:NZ	7:I:43:PRO:HB3	2.02	0.73
8:E:9:ILE:O	8:E:14:ARG:NH1	2.21	0.73
9:F:101:PRO:CB	9:F:126:THR:HG21	2.18	0.73
8:A:416:ARG:NH2	8:A:447:ASP:O	2.21	0.73
9:D:20:LEU:HD22	9:D:92:ILE:HG22	1.70	0.73
1:a:104:PHE:CG	4:p:97:PHE:CE1	2.77	0.73
2:b:82:ARG:CB	4:p:140:LEU:CD2	1.92	0.73
2:b:107:LYS:HE3	4:p:166:THR:CA	2.17	0.73
2:b:145:PHE:N	8:A:13:ILE:CD1	2.51	0.73
2:b:145:PHE:CD2	2:b:149:LEU:HD11	2.22	0.73
3:d:243:GLU:OE2	4:p:218:VAL:HG12	1.87	0.73
5:e:22:ILE:HG13	5:e:33:VAL:HG22	1.69	0.73
8:E:49:MET:HE3	8:E:95:ILE:CG2	2.18	0.73
1:a:135:ASN:ND2	4:p:86:PHE:CB	2.48	0.73
6:g:221:VAL:CG1	8:A:401:SER:OG	2.36	0.73
8:E:379:VAL:CG1	8:E:438:ILE:CG2	2.65	0.73
8:A:140:ARG:HB2	9:B:210:ASN:HD22	1.53	0.73
8:C:237:TYR:CE1	8:C:271:TYR:CG	2.76	0.73
1:a:124:HIS:CD2	4:p:79:GLU:CB	2.60	0.73
1:a:230:ILE:CG1	7:Q:55:LEU:HD11	2.18	0.73
2:b:173:ILE:CG1	3:d:221:PHE:CZ	2.71	0.73
3:d:235:MET:SD	8:A:24:VAL:HG12	2.29	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:152:ARG:HD3	8:A:503:LEU:CG	2.18	0.73
8:A:362:ILE:HG22	8:A:362:ILE:O	1.88	0.73
1:a:34:GLN:O	4:p:87:ASN:HB2	1.78	0.73
1:a:40:LEU:CD2	4:p:89:THR:CB	2.66	0.73
3:d:114:PHE:CD1	3:d:120:ILE:HD13	2.24	0.73
7:N:48:LYS:CG	7:M:46:GLU:OE1	2.36	0.73
9:F:110:ILE:CG2	9:F:118:VAL:CG2	2.65	0.73
1:a:165:ILE:HG22	1:a:168:THR:C	2.13	0.73
2:b:67:SER:O	4:p:130:VAL:CG1	2.37	0.73
2:b:103:ILE:CD1	4:p:163:LYS:N	2.50	0.73
2:b:141:ARG:HB3	8:A:16:ARG:NH1	2.03	0.73
6:g:360:ALA:CB	9:D:292:MET:HB3	2.18	0.73
7:S:55:LEU:CD2	7:R:54:LEU:HD11	2.18	0.73
7:H:54:LEU:HD11	7:I:55:LEU:CD2	2.19	0.73
4:p:210:LEU:CG	8:A:13:ILE:HG21	2.19	0.73
8:A:152:ILE:H	8:A:423:GLN:NE2	1.87	0.73
1:a:100:THR:HG23	4:p:101:MET:HG3	1.65	0.73
2:b:141:ARG:CG	8:A:12:ILE:HG21	2.17	0.73
4:p:206:GLN:O	4:p:210:LEU:CA	2.36	0.73
6:g:346:GLN:NE2	9:F:333:ASP:OD2	2.22	0.72
6:g:353:ILE:HD11	8:E:284:ARG:CA	2.19	0.72
7:N:43:PRO:C	7:O:48:LYS:HZ3	1.96	0.72
7:T:10:VAL:HG23	7:G:74:LEU:CD2	2.19	0.72
9:F:405:ILE:HG12	9:F:413:LEU:HD11	1.71	0.72
1:a:122:LEU:HA	2:b:25:ASP:CB	2.19	0.72
2:b:61:LEU:HD13	2:b:61:LEU:C	2.14	0.72
2:b:68:GLU:CD	4:p:126:GLN:CG	2.62	0.72
2:b:79:GLU:HG3	4:p:136:GLU:CG	2.18	0.72
2:b:82:ARG:CB	2:b:85:LEU:HD21	2.19	0.72
3:d:148:ILE:HD12	8:C:17:ILE:HG12	1.70	0.72
4:p:150:ALA:O	4:p:154:GLU:HG3	1.89	0.72
7:M:80:PHE:HD2	7:L:6:ALA:HB1	1.53	0.72
7:G:46:GLU:OE1	7:H:48:LYS:CB	2.37	0.72
8:A:214:GLN:HE21	8:A:219:MET:CG	1.80	0.72
2:b:180:ASN:CG	3:d:201:ILE:CG2	2.62	0.72
3:d:151:ASP:HB2	8:C:20:TYR:HH	1.53	0.72
3:d:183:VAL:HG22	3:d:214:ASP:O	1.90	0.72
5:e:22:ILE:HD11	5:e:74:ILE:HD13	1.71	0.72
7:N:6:ALA:HB1	7:O:80:PHE:HD2	1.54	0.72
7:G:43:PRO:CG	7:H:42:GLN:HE22	1.98	0.72
7:J:48:LYS:NZ	7:I:43:PRO:HB2	2.02	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:149:PHE:HE2	9:D:189:ILE:CD1	2.01	0.72
1:a:134:ILE:N	1:a:191:PHE:HE1	1.85	0.72
2:b:80:LYS:H	4:p:137:VAL:CG1	2.01	0.72
7:S:43:PRO:HB3	7:T:48:LYS:HD2	1.71	0.72
7:Q:46:GLU:OE1	7:R:48:LYS:CG	2.37	0.72
9:D:401:LEU:HD22	9:D:404:ILE:CD1	2.13	0.72
1:a:214:ILE:HG13	1:a:215:PRO:CD	2.20	0.72
8:E:98:ILE:HD13	8:E:242:THR:HG23	1.69	0.72
9:B:71:ASN:HD21	8:C:9:ILE:CA	2.02	0.72
1:a:115:LEU:HD21	2:b:38:LEU:HD11	1.69	0.72
2:b:83:ALA:H	4:p:140:LEU:HB3	0.73	0.72
4:p:196:LYS:HD2	8:A:5:ARG:HH22	1.53	0.72
7:K:62:ALA:O	7:K:65:ILE:CG2	2.33	0.72
8:A:419:GLU:OE2	8:A:422:LYS:HD2	1.88	0.72
2:b:176:PHE:CE1	3:d:233:VAL:CB	2.73	0.72
7:S:81:VAL:HG21	7:T:80:PHE:CE2	2.25	0.72
7:O:54:LEU:HD11	7:P:55:LEU:CD2	2.20	0.72
8:A:214:GLN:HE22	8:A:219:MET:CB	1.90	0.72
6:g:118:VAL:CG1	6:g:148:LEU:HD13	2.20	0.72
8:E:489:LEU:O	8:E:489:LEU:HD23	1.88	0.72
8:C:54:VAL:CG1	8:C:89:VAL:CG2	2.63	0.72
1:a:28:TRP:HB2	1:a:34:GLN:HB2	1.71	0.72
7:P:43:PRO:C	7:Q:48:LYS:HZ3	1.97	0.72
9:F:25:GLN:HG2	9:F:32:ASP:HB2	1.72	0.72
2:b:144:VAL:CB	8:A:13:ILE:HG13	2.20	0.72
8:E:379:VAL:HG12	8:E:438:ILE:HG22	1.70	0.72
8:C:469:LYS:NZ	8:C:492:GLU:HB2	2.05	0.72
2:b:118:LEU:CD1	4:p:178:ARG:HA	2.20	0.71
6:g:360:ALA:HA	9:D:293:PRO:CD	2.15	0.71
7:M:55:LEU:CD2	7:L:54:LEU:HD11	2.19	0.71
9:B:221:VAL:CG1	9:B:232:VAL:HG21	2.03	0.71
2:b:95:PHE:O	4:p:155:ILE:HG21	1.90	0.71
3:d:148:ILE:HD12	8:C:17:ILE:HA	1.69	0.71
7:P:54:LEU:HD11	7:Q:55:LEU:CD2	2.19	0.71
8:A:416:ARG:CZ	8:A:447:ASP:O	2.37	0.71
1:a:117:TRP:O	1:a:120:ILE:HG22	1.90	0.71
3:d:119:VAL:HG21	8:E:24:VAL:HG11	1.71	0.71
5:e:119:LEU:HD11	5:e:123:ARG:NH2	2.04	0.71
7:O:62:ALA:O	7:O:65:ILE:CG2	2.33	0.71
7:H:43:PRO:CG	7:I:42:GLN:HE22	2.02	0.71
8:E:492:GLU:O	8:E:495:GLN:HG2	1.90	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:111:ALA:HB3	8:A:235:LEU:HD22	1.72	0.71
8:A:154:ILE:HD12	8:A:358:ILE:HD11	1.72	0.71
9:B:311:GLU:CB	8:C:230:ASP:HB3	2.21	0.71
1:a:86:GLN:NE2	1:a:86:GLN:HA	2.05	0.71
1:a:118:LYS:NZ	1:a:126:GLU:CB	2.54	0.71
7:S:43:PRO:HG2	7:T:42:GLN:NE2	2.05	0.71
7:N:80:PHE:HD2	7:M:6:ALA:HB1	1.55	0.71
7:T:68:LEU:HD13	7:G:66:TYR:HB3	1.71	0.71
9:D:143:ASP:O	9:D:143:ASP:OD1	2.08	0.71
1:a:118:LYS:C	2:b:30:ASN:OD1	2.21	0.71
2:b:152:ALA:HB2	4:p:214:ILE:CD1	2.20	0.71
2:b:172:ASN:CG	3:d:235:MET:HB2	2.16	0.71
7:N:48:LYS:CB	7:M:46:GLU:OE1	2.38	0.71
7:M:49:ILE:HD11	7:L:40:ALA:HB2	1.71	0.71
7:T:10:VAL:CG2	7:G:74:LEU:HD22	2.18	0.71
1:a:116:PRO:HB3	2:b:32:ILE:C	2.15	0.71
7:Q:43:PRO:HB3	7:R:48:LYS:NZ	2.06	0.71
1:a:122:LEU:HD13	1:a:127:LEU:CD1	2.20	0.71
2:b:125:LYS:O	4:p:185:LEU:HD11	1.90	0.71
9:D:401:LEU:CD2	9:D:404:ILE:CD1	2.67	0.71
2:b:98:ASN:H	4:p:155:ILE:HD13	1.54	0.71
2:b:125:LYS:HB3	4:p:185:LEU:HD22	1.71	0.71
7:J:48:LYS:CG	7:I:46:GLU:OE1	2.39	0.71
2:b:152:ALA:HB2	4:p:214:ILE:HD11	1.71	0.71
3:d:247:GLN:NE2	8:A:13:ILE:CD1	2.49	0.71
4:p:193:GLU:CA	4:p:196:LYS:HE2	2.20	0.71
4:p:193:GLU:CB	4:p:196:LYS:HE2	2.20	0.71
5:e:22:ILE:HG12	5:e:33:VAL:CG2	2.21	0.71
5:e:34:LEU:HD11	7:K:43:PRO:HB2	1.72	0.71
7:H:43:PRO:C	7:I:48:LYS:HZ3	1.98	0.71
8:E:469:LYS:HD3	8:E:489:LEU:HD23	1.73	0.71
4:p:214:ILE:O	8:A:17:ILE:HD11	1.91	0.71
5:e:119:LEU:CD1	5:e:123:ARG:HE	1.73	0.71
6:g:110:THR:HG23	6:g:296:LEU:CD1	2.20	0.71
6:g:222:LYS:HE2	6:g:224:ASP:OD2	1.89	0.71
9:F:109:ARG:NH2	9:F:122:GLY:O	2.24	0.71
8:A:424:PRO:HB3	8:A:455:ARG:NH1	2.06	0.71
1:a:119:ILE:HG13	2:b:29:THR:N	2.05	0.70
1:a:134:ILE:CG2	1:a:191:PHE:CD1	2.71	0.70
1:a:225:GLY:O	1:a:228:ALA:HB3	1.90	0.70
6:g:300:LEU:HD23	6:g:300:LEU:C	2.14	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S:46:GLU:OE1	7:T:48:LYS:CG	2.39	0.70
7:T:43:PRO:HB3	7:G:48:LYS:CD	2.21	0.70
7:L:66:TYR:HB3	7:K:68:LEU:HD13	1.73	0.70
8:A:390:PHE:CD1	8:A:414:GLY:HA3	2.26	0.70
8:A:471:GLU:HB2	8:A:489:LEU:HD11	1.73	0.70
1:a:40:LEU:CG	4:p:89:THR:CB	2.51	0.70
1:a:96:PRO:HB2	4:p:101:MET:SD	2.31	0.70
2:b:118:LEU:CD1	4:p:178:ARG:CA	2.69	0.70
3:d:98:ASP:OD2	3:d:139:GLN:N	2.25	0.70
6:g:221:VAL:HG11	8:A:401:SER:CB	2.21	0.70
6:g:239:ILE:CG1	6:g:303:GLN:HE22	2.04	0.70
8:E:77:MET:HE2	8:E:112:LEU:HD21	1.73	0.70
8:C:27:VAL:HG23	8:C:47:GLU:OE1	1.91	0.70
1:a:35:ILE:HB	4:p:89:THR:H	1.54	0.70
2:b:160:LEU:CD2	2:b:164:LEU:HB2	2.21	0.70
8:E:230:ASP:HB3	9:D:311:GLU:CG	2.21	0.70
9:B:48:ILE:HG21	9:B:59:MET:HE2	1.73	0.70
2:b:67:SER:CB	4:p:124:LYS:O	2.39	0.70
6:g:129:LEU:HA	9:B:412:GLU:OE2	1.92	0.70
8:A:157:MET:HA	8:A:384:LYS:HE2	1.73	0.70
2:b:103:ILE:CB	4:p:159:LEU:CB	2.49	0.70
2:b:141:ARG:HB3	8:A:16:ARG:HH12	1.54	0.70
6:g:103:GLU:HG2	7:R:41:ARG:NH2	2.06	0.70
7:S:62:ALA:O	7:S:65:ILE:CG2	2.33	0.70
2:b:141:ARG:HG2	8:A:9:ILE:HD13	1.74	0.70
8:A:28:ASN:HB2	8:A:91:ALA:CB	2.18	0.70
8:A:134:ALA:HB3	9:B:240:ASN:HD22	1.55	0.70
9:D:20:LEU:HD21	9:D:92:ILE:HG22	1.71	0.70
7:S:66:TYR:HB3	7:R:68:LEU:HD13	1.73	0.70
7:P:46:GLU:OE1	7:Q:48:LYS:CB	2.39	0.70
1:a:124:HIS:HD2	4:p:80:LYS:O	1.73	0.70
2:b:160:LEU:CD1	2:b:160:LEU:O	2.32	0.70
6:g:109:LEU:HD21	6:g:199:PHE:CD1	2.26	0.70
7:S:43:PRO:HB3	7:T:48:LYS:CD	2.22	0.70
2:b:103:ILE:HD13	4:p:162:MET:N	1.96	0.70
4:p:88:LEU:O	4:p:92:ILE:HG12	1.92	0.70
7:H:10:VAL:HG23	7:I:74:LEU:HD21	1.72	0.70
8:A:65:LEU:HD12	8:A:75:VAL:CG2	2.22	0.70
1:a:134:ILE:HD12	1:a:135:ASN:CA	2.21	0.70
6:g:117:LYS:HG3	6:g:154:ASP:O	1.92	0.70
8:E:302:ALA:HB1	8:E:314:MET:HG2	1.74	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:433:GLU:OE1	8:A:462:ARG:CD	2.37	0.70
1:a:116:PRO:HG3	2:b:35:SER:H	1.54	0.69
1:a:132:ASN:O	1:a:191:PHE:CE1	2.45	0.69
1:a:195:LEU:HD21	4:p:84:PHE:CZ	2.26	0.69
2:b:61:LEU:CD1	2:b:65:ARG:HD3	2.22	0.69
5:e:19:VAL:HA	5:e:53:LEU:CD1	2.17	0.69
5:e:37:HIS:HE1	7:L:41:ARG:HE	1.39	0.69
6:g:104:ASP:CB	7:S:41:ARG:NH2	2.55	0.69
8:C:392:GLU:OE2	9:D:429:ARG:NH1	2.25	0.69
6:g:363:CYS:HB2	8:E:279:ARG:HH12	1.55	0.69
7:S:80:PHE:CE2	7:R:81:VAL:HG21	2.27	0.69
7:H:81:VAL:HG21	7:I:80:PHE:CE2	2.27	0.69
7:J:80:PHE:HD2	7:I:6:ALA:HB1	1.57	0.69
8:E:449:LEU:CD1	8:E:457:TYR:CD2	2.75	0.69
1:a:27:TYR:HB3	1:a:124:HIS:HB2	1.73	0.69
4:p:199:THR:C	8:A:6:ALA:HB1	2.04	0.69
7:Q:10:VAL:HG23	7:R:74:LEU:HD21	1.74	0.69
8:E:9:ILE:HG23	8:E:13:ILE:CG2	2.22	0.69
8:E:439:TYR:OH	8:E:487:GLU:OE2	2.06	0.69
8:E:469:LYS:HD3	8:E:489:LEU:CD2	2.23	0.69
9:F:100:VAL:CG1	9:F:256:MET:HE2	2.23	0.69
9:F:124:VAL:HG12	9:F:126:THR:HG22	1.74	0.69
9:F:268:VAL:HB	9:F:321:ILE:HG22	1.73	0.69
2:b:88:VAL:C	4:p:148:MET:N	2.47	0.69
2:b:144:VAL:HG13	4:p:207:ILE:HD13	1.74	0.69
6:g:222:LYS:HE2	6:g:224:ASP:CG	2.16	0.69
7:P:43:PRO:HB3	7:Q:48:LYS:HD2	1.74	0.69
7:Q:46:GLU:OE1	7:R:48:LYS:CB	2.39	0.69
7:K:74:LEU:CD2	7:J:10:VAL:HG23	2.23	0.69
8:A:469:LYS:HZ3	8:A:492:GLU:HB2	1.55	0.69
7:P:10:VAL:HG23	7:Q:74:LEU:HD21	1.74	0.69
8:E:230:ASP:HB3	9:D:311:GLU:HG3	1.73	0.69
8:E:449:LEU:HD11	8:E:457:TYR:CD2	2.27	0.69
1:a:58:VAL:HG11	4:p:107:ILE:CG2	2.09	0.69
2:b:165:HIS:HD2	3:d:219:ALA:CB	1.85	0.69
5:e:4:ASN:HB2	5:e:18:GLU:HG2	1.75	0.69
7:P:43:PRO:HB2	7:Q:48:LYS:HZ2	1.55	0.69
2:b:122:GLU:CA	4:p:181:ILE:CG2	2.68	0.69
3:d:239:LYS:HG2	8:A:22:ARG:HD2	1.75	0.69
4:p:108:TYR:HE1	4:p:112:LEU:CD1	2.02	0.69
7:N:55:LEU:CD2	7:M:54:LEU:HD11	2.22	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:M:48:LYS:HD3	7:L:46:GLU:OE1	1.92	0.69
7:K:48:LYS:CD	7:J:43:PRO:HB3	2.22	0.69
9:D:149:PHE:HE2	9:D:189:ILE:HD11	1.58	0.69
9:D:361:ILE:HG23	9:D:432:SER:HB3	1.74	0.69
1:a:134:ILE:HD12	1:a:134:ILE:C	2.17	0.69
1:a:222:PHE:HZ	7:Q:54:LEU:HD23	0.95	0.69
2:b:175:MET:HB3	8:A:24:VAL:HG22	1.74	0.69
7:J:55:LEU:CD2	7:I:54:LEU:HD11	2.23	0.69
8:A:329:ALA:HB3	8:A:332:PRO:HD2	1.75	0.69
8:C:439:TYR:CD2	8:C:490:LEU:CD2	2.75	0.69
2:b:71:ARG:HA	4:p:132:ASP:N	2.06	0.69
3:d:247:GLN:O	8:A:16:ARG:CD	2.41	0.69
6:g:50:ARG:HG2	6:g:50:ARG:HH11	1.56	0.69
7:M:48:LYS:HZ3	7:L:43:PRO:C	2.00	0.69
7:J:62:ALA:O	7:J:65:ILE:CG2	2.33	0.69
8:E:302:ALA:CB	8:E:314:MET:HG2	2.23	0.69
8:A:495:GLN:OE1	8:A:498:MET:HE3	1.93	0.69
8:C:138:MET:CG	9:D:119:ASP:HA	2.15	0.69
7:T:62:ALA:O	7:T:65:ILE:CG2	2.33	0.69
9:F:338:ALA:HB3	9:F:339:PRO:HD3	1.73	0.69
3:d:91:THR:O	3:d:95:THR:HG22	1.92	0.68
9:F:113:VAL:HG22	9:F:249:VAL:HG13	1.75	0.68
8:A:105:LEU:HD21	8:A:193:ILE:HG23	1.75	0.68
9:D:111:PHE:HE1	9:D:124:VAL:HG21	1.57	0.68
1:a:23:GLY:O	4:p:83:LEU:HD21	1.92	0.68
1:a:182:LYS:CD	1:a:238:TYR:CE1	2.74	0.68
6:g:353:ILE:CG1	8:E:284:ARG:HA	2.23	0.68
7:N:43:PRO:HB3	7:O:48:LYS:NZ	2.07	0.68
7:M:55:LEU:HD23	7:L:54:LEU:HD11	1.75	0.68
8:A:154:ILE:CD1	8:A:358:ILE:CD1	2.71	0.68
9:D:261:ARG:NH1	9:D:314:THR:O	2.26	0.68
1:a:79:ILE:CD1	1:a:99:GLY:CA	2.64	0.68
2:b:80:LYS:H	4:p:137:VAL:HG12	1.56	0.68
2:b:118:LEU:CD2	4:p:177:GLY:HA3	2.23	0.68
5:e:21:GLU:OE2	5:e:34:LEU:CD2	2.39	0.68
5:e:34:LEU:CD1	7:K:43:PRO:CB	2.70	0.68
6:g:117:LYS:HA	6:g:153:VAL:O	1.92	0.68
6:g:222:LYS:HE2	6:g:224:ASP:OD1	1.93	0.68
7:N:48:LYS:NZ	7:M:43:PRO:HB3	2.06	0.68
7:O:43:PRO:HB2	7:P:48:LYS:NZ	2.07	0.68
7:M:48:LYS:NZ	7:L:43:PRO:CB	2.57	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:E:379:VAL:HG12	8:E:438:ILE:HG21	1.74	0.68
8:A:42:ILE:HD13	8:A:89:VAL:HG11	1.76	0.68
2:b:92:ALA:HB2	4:p:148:MET:C	2.18	0.68
3:d:177:VAL:HG11	3:d:210:LYS:CD	2.23	0.68
6:g:121:MET:HE3	6:g:160:ILE:HD11	1.74	0.68
7:Q:68:LEU:HD13	7:R:66:TYR:HB3	1.76	0.68
8:A:214:GLN:HE22	8:A:219:MET:HB3	1.55	0.68
8:A:327:VAL:HG21	8:A:344:PHE:HE1	1.57	0.68
8:C:260:ILE:CD1	8:C:317:LEU:HD12	2.23	0.68
1:a:24:GLN:HA	4:p:83:LEU:CG	2.22	0.68
3:d:233:VAL:HG12	8:A:26:VAL:CA	2.14	0.68
6:g:104:ASP:HB2	7:S:41:ARG:NH2	2.09	0.68
6:g:137:LEU:HD12	6:g:213:LEU:HD13	1.76	0.68
7:Q:43:PRO:C	7:R:48:LYS:HZ3	2.01	0.68
8:E:169:ILE:HD11	8:E:322:THR:CG2	2.22	0.68
8:E:469:LYS:HE3	8:E:493:ALA:CA	2.24	0.68
9:F:189:ILE:HD11	9:F:267:ASP:O	1.94	0.68
9:F:447:LYS:HZ3	9:F:482:GLU:CD	2.00	0.68
8:A:152:ILE:N	8:A:423:GLN:HE22	1.90	0.68
8:C:169:ILE:HD11	8:C:322:THR:CG2	2.24	0.68
1:a:119:ILE:HG12	2:b:32:ILE:HG12	1.75	0.68
2:b:79:GLU:C	4:p:137:VAL:HA	2.18	0.68
2:b:103:ILE:HG12	4:p:162:MET:C	2.18	0.68
6:g:284:PHE:HD1	6:g:291:ILE:CD1	2.06	0.68
1:a:139:ALA:CB	4:p:94:MET:HE2	2.23	0.68
1:a:193:ASN:HD22	1:a:227:GLN:HG3	1.54	0.68
8:E:100:VAL:HG23	8:E:246:LEU:HD23	1.76	0.68
1:a:24:GLN:C	4:p:83:LEU:CG	2.67	0.68
2:b:125:LYS:HD2	4:p:185:LEU:CG	2.22	0.68
2:b:176:PHE:HE1	3:d:233:VAL:CG2	1.60	0.68
5:e:119:LEU:CD1	5:e:123:ARG:NH2	2.57	0.68
6:g:85:PHE:CE1	6:g:302:SER:OG	2.45	0.68
1:a:24:GLN:CB	4:p:84:PHE:HD2	1.84	0.68
6:g:251:GLU:HG3	6:g:272:LYS:HG2	1.75	0.68
7:H:43:PRO:HB3	7:I:48:LYS:NZ	2.09	0.68
7:K:74:LEU:HD22	7:J:10:VAL:CG2	2.23	0.68
9:F:275:ILE:HG21	9:F:327:VAL:HG22	1.74	0.68
6:g:353:ILE:CD1	8:E:284:ARG:HG2	2.23	0.68
7:L:48:LYS:NZ	7:K:43:PRO:HB2	2.08	0.68
8:E:237:TYR:CE1	8:E:294:LEU:HD11	2.28	0.68
1:a:119:ILE:CB	2:b:28:ALA:CA	2.54	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:182:LYS:HD2	1:a:238:TYR:CD1	2.27	0.67
2:b:125:LYS:CA	4:p:185:LEU:CD2	2.67	0.67
6:g:83:ARG:NH2	6:g:273:THR:HG21	2.07	0.67
6:g:253:GLU:HB3	6:g:270:MET:CB	2.24	0.67
7:S:46:GLU:OE1	7:T:48:LYS:CB	2.43	0.67
3:d:132:ILE:CA	8:E:13:ILE:CD1	2.72	0.67
6:g:115:VAL:O	6:g:115:VAL:HG12	1.92	0.67
6:g:118:VAL:CG1	6:g:148:LEU:HD11	2.18	0.67
7:R:62:ALA:O	7:R:65:ILE:CG2	2.33	0.67
8:E:490:LEU:C	8:E:490:LEU:HD23	2.19	0.67
9:F:347:ASP:O	9:F:373:THR:HG23	1.94	0.67
8:A:416:ARG:CD	8:A:447:ASP:HA	2.24	0.67
8:C:227:GLU:HG3	8:C:239:ALA:HB2	1.76	0.67
1:a:64:ILE:CG2	2:b:57:LYS:CE	2.69	0.67
8:A:109:ILE:CG1	8:A:113:ALA:HA	2.23	0.67
8:A:159:PRO:HG3	8:A:368:GLY:O	1.92	0.67
9:B:195:GLY:HA3	9:B:267:ASP:O	1.94	0.67
2:b:114:THR:O	2:b:117:THR:CG2	2.41	0.67
2:b:114:THR:C	2:b:117:THR:HG22	2.19	0.67
4:p:210:LEU:O	4:p:213:ASP:CB	2.41	0.67
9:F:22:ARG:HH12	9:F:90:GLU:HB3	1.59	0.67
8:A:216:ARG:CZ	8:A:426:SER:OG	2.41	0.67
1:a:116:PRO:HG3	2:b:35:SER:CB	2.23	0.67
1:a:195:LEU:HD22	4:p:84:PHE:CE1	2.30	0.67
1:a:217:MET:HE1	2:b:33:ASN:HD21	1.58	0.67
2:b:80:LYS:N	4:p:137:VAL:CB	2.57	0.67
7:J:48:LYS:CB	7:I:46:GLU:OE1	2.42	0.67
9:F:100:VAL:HG12	9:F:256:MET:HE2	1.76	0.67
9:F:310:GLN:HE22	9:F:325:GLN:HE22	1.43	0.67
1:a:120:ILE:N	2:b:30:ASN:HB2	2.10	0.67
2:b:92:ALA:H	4:p:148:MET:HA	1.59	0.67
7:Q:62:ALA:O	7:Q:65:ILE:CG2	2.33	0.67
8:E:373:ILE:HG13	8:E:376:MET:H	1.58	0.67
1:a:25:HIS:N	4:p:83:LEU:HG	2.08	0.67
1:a:35:ILE:HG22	4:p:88:LEU:H	0.56	0.67
1:a:62:GLN:OE1	1:a:66:THR:HB	1.94	0.67
1:a:137:THR:HG21	1:a:191:PHE:CA	2.24	0.67
2:b:114:THR:O	4:p:174:LEU:CG	2.42	0.67
3:d:233:VAL:HG11	8:A:26:VAL:HA	1.71	0.67
5:e:4:ASN:HB2	5:e:18:GLU:CD	2.20	0.67
6:g:123:VAL:HG22	6:g:160:ILE:HD12	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:284:PHE:CD1	6:g:291:ILE:HD12	2.30	0.67
7:Q:54:LEU:HD11	7:R:55:LEU:CD2	2.25	0.67
7:M:62:ALA:O	7:M:65:ILE:CG2	2.34	0.67
3:d:71:VAL:N	8:A:69:SER:O	2.28	0.67
8:E:490:LEU:HD23	8:E:490:LEU:O	1.93	0.67
1:a:122:LEU:CA	2:b:25:ASP:CB	2.63	0.67
2:b:60:ILE:CD1	4:p:120:ASP:OD2	2.43	0.67
3:d:152:SER:C	3:d:153:GLU:CD	2.63	0.67
3:d:224:ARG:CZ	3:d:232:LEU:HD12	2.24	0.67
3:d:235:MET:HE3	3:d:239:LYS:CG	2.25	0.67
4:p:203:LEU:HG	4:p:203:LEU:O	1.95	0.67
7:S:10:VAL:HG23	7:T:74:LEU:HD21	1.75	0.67
7:S:42:GLN:NE2	7:R:43:PRO:HG2	2.09	0.67
9:F:101:PRO:HB3	9:F:126:THR:HG21	1.77	0.67
9:B:71:ASN:ND2	8:C:9:ILE:HA	2.09	0.67
1:a:135:ASN:CG	4:p:86:PHE:CB	2.66	0.67
2:b:128:THR:HG21	4:p:189:LEU:HD11	1.77	0.67
8:E:355:ARG:NH1	9:D:389:GLN:NE2	2.41	0.67
1:a:70:ASN:O	2:b:53:LEU:HD13	1.34	0.66
1:a:116:PRO:HG3	2:b:35:SER:HB2	1.76	0.66
2:b:75:ILE:HD11	4:p:132:ASP:OD2	1.94	0.66
3:d:154:ARG:CZ	8:C:5:ARG:NH1	2.58	0.66
4:p:127:LEU:C	4:p:130:VAL:HG12	2.20	0.66
5:e:109:LYS:O	5:e:113:ILE:HG12	1.95	0.66
6:g:281:ILE:CG1	7:M:41:ARG:HG3	2.24	0.66
7:R:62:ALA:O	7:R:65:ILE:N	2.29	0.66
7:M:62:ALA:O	7:M:65:ILE:N	2.28	0.66
7:H:62:ALA:O	7:H:65:ILE:N	2.29	0.66
7:L:62:ALA:O	7:L:65:ILE:CG2	2.33	0.66
1:a:161:PHE:HE1	1:a:163:LYS:HG3	0.51	0.66
7:S:74:LEU:HD21	7:R:10:VAL:HG23	1.77	0.66
7:N:46:GLU:OE1	7:O:48:LYS:HG2	1.95	0.66
7:J:62:ALA:O	7:J:65:ILE:N	2.29	0.66
7:I:62:ALA:O	7:I:65:ILE:CG2	2.33	0.66
8:A:464:TYR:HE1	8:A:468:ASN:HD21	1.32	0.66
9:B:410:LEU:HD23	9:B:413:LEU:HD12	1.77	0.66
8:C:435:VAL:HG21	8:C:476:ILE:HD11	1.76	0.66
1:a:43:SER:HB2	4:p:93:ILE:CG1	2.15	0.66
6:g:343:ARG:NH2	8:A:326:ASP:CB	2.57	0.66
8:A:238:LEU:HG	8:A:238:LEU:O	1.93	0.66
8:A:296:SER:HB2	9:B:239:MET:HB3	1.77	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:430:THR:HG22	8:A:432:GLU:OE1	1.95	0.66
8:C:184:ILE:HD11	8:C:260:ILE:HG13	1.78	0.66
9:D:64:GLU:OE1	9:D:248:ARG:NH2	2.29	0.66
2:b:75:ILE:HG13	4:p:132:ASP:C	1.70	0.66
2:b:79:GLU:N	4:p:137:VAL:HA	2.03	0.66
3:d:86:ALA:HB1	3:d:95:THR:HG21	1.77	0.66
3:d:119:VAL:CG2	8:E:24:VAL:HG11	2.24	0.66
7:N:62:ALA:O	7:N:65:ILE:N	2.28	0.66
7:T:6:ALA:HB1	7:G:80:PHE:HD2	1.61	0.66
7:L:74:LEU:CD2	7:K:10:VAL:CG2	2.73	0.66
7:K:48:LYS:HZ2	7:J:43:PRO:CB	2.02	0.66
8:A:39:ILE:CD1	8:A:278:LEU:HD21	2.25	0.66
9:B:112:ASN:HD22	9:B:116:GLU:HB2	1.60	0.66
9:D:53:ASP:HB3	9:D:59:MET:HE3	1.76	0.66
1:a:76:LEU:HD13	4:p:109:TYR:HD1	1.60	0.66
2:b:160:LEU:CD1	2:b:164:LEU:H	0.10	0.66
2:b:160:LEU:HB2	2:b:163:GLU:HB2	1.78	0.66
7:N:62:ALA:O	7:N:65:ILE:CG2	2.33	0.66
7:O:46:GLU:CD	7:P:48:LYS:HB3	2.20	0.66
7:T:81:VAL:HG21	7:G:80:PHE:CE2	2.30	0.66
1:a:122:LEU:CA	2:b:25:ASP:HB2	2.25	0.66
2:b:77:GLN:O	2:b:80:LYS:HB2	1.95	0.66
2:b:175:MET:HB3	8:A:24:VAL:CG2	2.25	0.66
4:p:210:LEU:HG	8:A:13:ILE:HG21	1.78	0.66
7:Q:10:VAL:CG2	7:R:74:LEU:HD21	2.26	0.66
7:G:62:ALA:O	7:G:65:ILE:N	2.29	0.66
7:L:62:ALA:O	7:L:65:ILE:N	2.29	0.66
1:a:61:PRO:HB3	4:p:112:LEU:HD21	1.78	0.66
3:d:222:THR:HG23	3:d:234:ASP:HB2	1.60	0.66
3:d:233:VAL:CG1	8:A:26:VAL:CA	2.60	0.66
7:S:48:LYS:HD2	7:R:43:PRO:HB3	1.77	0.66
7:S:62:ALA:O	7:S:65:ILE:N	2.28	0.66
7:P:62:ALA:O	7:P:65:ILE:N	2.28	0.66
7:K:80:PHE:CE2	7:J:81:VAL:HG21	2.30	0.66
8:A:344:PHE:CE2	8:A:362:ILE:HG22	2.30	0.66
8:C:237:TYR:CE1	8:C:294:LEU:HD21	2.31	0.66
1:a:135:ASN:CG	4:p:86:PHE:CD1	2.71	0.66
3:d:119:VAL:HG21	8:E:24:VAL:CG1	2.25	0.66
4:p:193:GLU:O	4:p:196:LYS:CG	2.44	0.66
6:g:356:ILE:O	9:D:293:PRO:CG	2.43	0.66
7:P:43:PRO:CB	7:Q:48:LYS:HZ2	2.07	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:T:33:GLY:HA3	7:G:31:ALA:HA	1.77	0.66
7:H:62:ALA:O	7:H:65:ILE:CG2	2.33	0.66
3:d:242:GLU:HG3	8:A:20:TYR:CZ	2.31	0.66
7:T:40:ALA:CB	7:G:49:ILE:HD11	2.23	0.66
7:T:62:ALA:O	7:T:65:ILE:N	2.28	0.66
8:A:157:MET:HG2	8:A:384:LYS:HG2	1.77	0.66
9:D:111:PHE:CE1	9:D:124:VAL:HG21	2.31	0.66
2:b:79:GLU:N	4:p:136:GLU:CG	2.58	0.66
3:d:148:ILE:CD1	8:C:17:ILE:CG1	2.74	0.66
3:d:152:SER:O	3:d:153:GLU:OE1	2.14	0.66
4:p:212:ASP:OD1	4:p:215:VAL:HG11	1.96	0.66
7:P:62:ALA:O	7:P:65:ILE:CG2	2.33	0.66
9:B:296:VAL:HG12	9:B:297:GLY:N	2.10	0.66
9:B:311:GLU:HG3	8:C:230:ASP:CB	2.26	0.66
1:a:62:GLN:C	4:p:119:ARG:NH1	2.54	0.65
1:a:134:ILE:CD1	1:a:135:ASN:N	2.47	0.65
6:g:107:VAL:HG11	6:g:296:LEU:HD12	1.77	0.65
6:g:363:CYS:HB3	9:D:290:GLY:C	2.20	0.65
8:E:469:LYS:HB3	8:E:489:LEU:HD21	1.77	0.65
2:b:160:LEU:CD1	2:b:164:LEU:N	0.78	0.65
3:d:148:ILE:HG23	8:C:16:ARG:CD	2.25	0.65
4:p:199:THR:O	8:A:6:ALA:CB	2.36	0.65
8:E:284:ARG:NH1	8:E:330:TYR:HD1	1.91	0.65
2:b:121:PHE:CZ	2:b:125:LYS:CE	2.79	0.65
3:d:148:ILE:HG23	8:C:16:ARG:HD2	1.77	0.65
5:e:119:LEU:HD12	5:e:123:ARG:CZ	2.25	0.65
6:g:356:ILE:HD13	8:E:286:ALA:CB	2.25	0.65
7:S:43:PRO:CB	7:T:48:LYS:HZ2	2.00	0.65
7:G:43:PRO:HB2	7:H:48:LYS:NZ	2.11	0.65
8:A:464:TYR:CD2	8:A:497:GLN:HB2	2.31	0.65
9:B:275:ILE:HD13	9:B:306:MET:HE1	1.76	0.65
8:C:30:GLY:HA3	8:C:45:LEU:HG	1.79	0.65
4:p:107:ILE:O	4:p:111:PRO:HG2	1.96	0.65
7:N:46:GLU:CD	7:O:48:LYS:HB3	2.21	0.65
7:Q:6:ALA:HB1	7:R:80:PHE:CD2	2.30	0.65
8:E:237:TYR:HE1	8:E:294:LEU:HD11	1.61	0.65
3:d:114:PHE:HD1	3:d:120:ILE:HD13	1.60	0.65
3:d:222:THR:CB	3:d:234:ASP:HB3	2.27	0.65
5:e:60:LEU:O	5:e:60:LEU:CD1	2.36	0.65
6:g:83:ARG:HH22	6:g:273:THR:CG2	2.09	0.65
6:g:149:LYS:NZ	6:g:155:TYR:HE1	1.93	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Q:43:PRO:CG	7:R:42:GLN:HE22	2.07	0.65
7:K:48:LYS:HG2	7:J:46:GLU:OE1	1.96	0.65
8:E:109:ILE:HD12	8:E:113:ALA:HA	1.77	0.65
8:A:109:ILE:HG23	8:A:225:VAL:HA	1.77	0.65
9:D:411:ASP:HA	9:D:418:ARG:NH2	1.97	0.65
3:d:132:ILE:N	8:E:13:ILE:CD1	2.60	0.65
4:p:210:LEU:CD1	8:A:13:ILE:HG22	2.24	0.65
7:T:46:GLU:OE1	7:G:48:LYS:HG2	1.96	0.65
7:K:42:GLN:HE22	7:J:43:PRO:CG	2.09	0.65
7:I:62:ALA:O	7:I:65:ILE:N	2.28	0.65
9:B:71:ASN:ND2	8:C:9:ILE:C	2.54	0.65
5:e:3:LEU:HD23	5:e:35:PRO:HA	1.78	0.65
6:g:221:VAL:HG23	8:A:398:GLN:O	1.93	0.65
2:b:88:VAL:C	2:b:90:MET:N	2.53	0.65
3:d:87:ASP:CB	3:d:172:ILE:HD11	2.26	0.65
5:e:96:LEU:HD13	5:e:123:ARG:HA	1.78	0.65
7:O:42:GLN:HE21	7:O:44:GLU:HB3	1.62	0.65
9:B:113:VAL:HG22	9:B:249:VAL:HG22	1.79	0.65
8:C:166:GLU:O	8:C:318:PRO:HD2	1.96	0.65
7:K:80:PHE:HD2	7:J:6:ALA:HB1	1.62	0.65
1:a:149:PHE:HZ	1:a:163:LYS:NZ	1.95	0.65
2:b:107:LYS:CE	4:p:166:THR:CA	2.73	0.65
4:p:152:ARG:NH2	8:A:502:LEU:HB3	2.12	0.65
7:H:43:PRO:HB3	7:I:48:LYS:HD2	1.78	0.65
8:A:436:MET:HG3	8:A:490:LEU:CD1	2.27	0.65
9:D:261:ARG:HD3	9:D:321:ILE:CD1	2.27	0.65
1:a:134:ILE:N	4:p:86:PHE:CZ	2.61	0.64
2:b:71:ARG:NH1	4:p:126:GLN:HA	2.04	0.64
5:e:91:GLU:N	5:e:91:GLU:OE1	2.30	0.64
6:g:57:THR:O	6:g:61:THR:HG23	1.97	0.64
7:K:62:ALA:O	7:K:65:ILE:N	2.28	0.64
8:A:497:GLN:HA	8:A:497:GLN:NE2	2.12	0.64
9:B:338:ALA:HB3	9:B:339:PRO:HD3	1.79	0.64
1:a:63:THR:N	4:p:119:ARG:NH1	2.45	0.64
6:g:83:ARG:HH22	6:g:273:THR:HG23	1.62	0.64
7:P:43:PRO:HB3	7:Q:48:LYS:CD	2.27	0.64
7:T:10:VAL:HG23	7:G:74:LEU:HD21	1.78	0.64
8:E:430:THR:HG22	8:E:432:GLU:OE1	1.97	0.64
8:A:154:ILE:HD11	8:A:358:ILE:HD11	1.79	0.64
8:C:12:ILE:HD11	8:C:33:LEU:HD21	1.78	0.64
1:a:51:LEU:HD21	4:p:100:LEU:HD12	0.68	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:82:ARG:CG	2:b:85:LEU:CD2	2.48	0.64
6:g:360:ALA:CB	9:D:293:PRO:HD2	2.27	0.64
7:O:43:PRO:CG	7:P:42:GLN:NE2	2.59	0.64
7:O:43:PRO:HB3	7:P:48:LYS:NZ	2.12	0.64
7:T:43:PRO:CA	7:G:48:LYS:HZ3	2.11	0.64
8:C:130:ILE:HG23	8:C:241:TYR:HB3	1.80	0.64
2:b:125:LYS:HB3	4:p:185:LEU:CD2	2.23	0.64
2:b:133:GLN:HG2	8:A:3:THR:HA	1.80	0.64
7:M:48:LYS:CD	7:L:46:GLU:OE1	2.45	0.64
7:G:10:VAL:CG2	7:H:74:LEU:HD21	2.25	0.64
9:B:20:LEU:HD12	9:B:90:GLU:OE2	1.92	0.64
9:B:64:GLU:OE2	9:B:133:HIS:NE2	2.30	0.64
8:C:181:THR:HG23	8:C:213:PHE:CE1	2.33	0.64
1:a:126:GLU:CD	4:p:83:LEU:CD2	2.71	0.64
2:b:88:VAL:CB	4:p:147:VAL:HG13	1.94	0.64
3:d:109:GLU:N	3:d:110:PRO:HD2	2.13	0.64
3:d:201:ILE:HG13	3:d:202:THR:N	2.12	0.64
4:p:210:LEU:CG	8:A:13:ILE:CG2	2.74	0.64
6:g:202:PHE:CB	6:g:231:LEU:HD13	2.28	0.64
8:C:308:LEU:HD22	8:C:309:LEU:CD1	2.28	0.64
3:d:197:GLY:O	3:d:201:ILE:HG12	1.97	0.64
7:T:43:PRO:CB	7:G:48:LYS:HZ2	2.04	0.64
8:A:495:GLN:OE1	8:A:498:MET:CE	2.46	0.64
9:B:216:MET:CB	9:B:222:ILE:HG12	2.27	0.64
8:C:14:ARG:O	8:C:18:GLU:HG3	1.98	0.64
8:C:251:MET:HE3	8:C:252:TYR:CE1	2.32	0.64
9:D:167:LYS:NZ	9:D:310:GLN:OE1	2.31	0.64
9:D:387:ILE:HD13	9:D:455:ILE:CG2	2.15	0.64
1:a:124:HIS:CD2	4:p:79:GLU:OE1	2.51	0.64
2:b:159:CYS:HB3	2:b:164:LEU:HD23	1.80	0.64
3:d:198:VAL:HA	3:d:201:ILE:CG1	2.27	0.64
4:p:90:LEU:HB2	4:p:91:PRO:CD	2.26	0.64
5:e:42:THR:HG22	6:g:285:GLU:HB2	1.79	0.64
6:g:281:ILE:HD13	7:M:41:ARG:O	1.97	0.64
7:H:46:GLU:OE1	7:I:48:LYS:HG2	1.98	0.64
7:K:31:ALA:HA	7:J:33:GLY:HA3	1.80	0.64
8:A:446:LEU:HD13	8:A:454:VAL:HG13	1.80	0.64
8:A:469:LYS:NZ	8:A:492:GLU:CB	2.59	0.64
9:D:423:ARG:HH21	9:D:464:GLY:HA3	1.62	0.64
1:a:35:ILE:HB	4:p:89:THR:N	2.12	0.64
2:b:148:ALA:O	4:p:214:ILE:HD11	1.96	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:148:ILE:CD1	8:C:17:ILE:N	2.60	0.64
7:N:48:LYS:HZ3	7:M:43:PRO:C	2.06	0.64
7:O:62:ALA:O	7:O:65:ILE:N	2.29	0.64
7:G:62:ALA:O	7:G:65:ILE:CG2	2.33	0.64
8:A:475:ILE:HG21	8:A:486:ALA:HB2	1.79	0.64
9:D:67:GLN:HE21	9:D:75:ARG:HD2	1.63	0.64
1:a:119:ILE:HD11	2:b:32:ILE:CA	2.27	0.64
1:a:135:ASN:ND2	4:p:85:ASP:O	2.31	0.64
6:g:144:ARG:NH2	6:g:242:ILE:HD12	2.13	0.64
6:g:360:ALA:HA	9:D:292:MET:CB	2.28	0.64
7:H:46:GLU:CD	7:I:48:LYS:HB3	2.23	0.64
8:C:75:VAL:HG11	8:C:274:MET:HE3	1.80	0.64
9:D:62:THR:HG21	9:D:98:LEU:HD21	1.79	0.64
1:a:118:LYS:HZ2	1:a:126:GLU:HB2	1.62	0.64
1:a:119:ILE:HG23	2:b:28:ALA:CB	2.22	0.64
2:b:160:LEU:CD1	2:b:163:GLU:HB2	2.24	0.64
3:d:132:ILE:HG12	8:E:13:ILE:HD13	1.79	0.64
5:e:25:SER:O	5:e:44:VAL:HG23	1.97	0.64
7:G:43:PRO:HB3	7:H:48:LYS:NZ	2.12	0.64
8:E:9:ILE:CG2	8:E:14:ARG:HG3	2.27	0.64
8:C:164:GLN:HG2	8:C:165:ARG:H	1.62	0.64
1:a:40:LEU:CD2	4:p:89:THR:CA	2.73	0.63
1:a:44:TRP:NE1	4:p:96:GLU:OE1	2.29	0.63
2:b:74:ALA:CB	4:p:130:VAL:O	2.46	0.63
2:b:144:VAL:HG23	8:A:9:ILE:HG21	1.71	0.63
3:d:177:VAL:CG1	3:d:210:LYS:HD2	2.27	0.63
6:g:251:GLU:HG3	6:g:272:LYS:CG	2.28	0.63
7:O:43:PRO:C	7:P:48:LYS:HZ3	2.05	0.63
7:L:48:LYS:NZ	7:K:43:PRO:HB3	2.12	0.63
8:E:461:LEU:HD23	8:E:461:LEU:C	2.23	0.63
8:C:54:VAL:HG13	8:C:90:LYS:O	1.97	0.63
9:D:53:ASP:HB3	9:D:59:MET:HE2	1.80	0.63
2:b:60:ILE:CG1	4:p:120:ASP:OD2	2.45	0.63
2:b:95:PHE:O	4:p:155:ILE:CG2	2.32	0.63
2:b:125:LYS:HB3	4:p:185:LEU:HD13	1.78	0.63
2:b:180:ASN:CB	3:d:201:ILE:HG22	2.26	0.63
3:d:154:ARG:NH2	8:C:8:GLU:OE2	2.32	0.63
4:p:126:GLN:OE1	4:p:126:GLN:HA	1.96	0.63
4:p:192:LEU:HD22	8:A:3:THR:HG23	1.81	0.63
6:g:83:ARG:NH2	6:g:273:THR:HG23	2.11	0.63
2:b:71:ARG:HA	4:p:132:ASP:H	1.62	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:N:43:PRO:HB3	7:O:48:LYS:HD2	1.79	0.63
7:G:33:GLY:HA3	7:H:31:ALA:HA	1.81	0.63
9:F:385:TYR:HE1	9:F:389:GLN:HE21	1.36	0.63
9:D:411:ASP:CA	9:D:418:ARG:HH21	2.02	0.63
7:S:48:LYS:CD	7:R:43:PRO:HB3	2.28	0.63
8:A:104:TYR:HE2	8:A:109:ILE:HD12	1.63	0.63
8:A:432:GLU:O	8:A:435:VAL:CG1	2.46	0.63
1:a:116:PRO:CB	2:b:34:LEU:N	2.56	0.63
2:b:169:ILE:HG12	3:d:220:GLY:HA2	1.80	0.63
4:p:85:ASP:C	4:p:86:PHE:HD1	2.06	0.63
6:g:280:PRO:C	7:N:41:ARG:HH21	2.07	0.63
7:N:54:LEU:HD11	7:O:55:LEU:HD23	1.79	0.63
9:F:408:LEU:O	9:F:412:GLU:HB2	1.99	0.63
1:a:184:LEU:CD2	1:a:188:PHE:CE2	2.82	0.63
2:b:88:VAL:HA	4:p:148:MET:CB	2.15	0.63
4:p:177:GLY:O	4:p:181:ILE:HG12	1.99	0.63
7:M:80:PHE:CE2	7:L:81:VAL:CG2	2.80	0.63
8:E:100:VAL:HG23	8:E:246:LEU:CD2	2.29	0.63
8:E:345:LEU:HD21	8:E:358:ILE:HD13	1.79	0.63
9:F:384:HIS:O	9:F:387:ILE:HG22	1.98	0.63
1:a:53:SER:HA	1:a:56:ILE:HG22	1.79	0.63
1:a:121:GLN:CB	2:b:25:ASP:OD1	2.47	0.63
2:b:103:ILE:CG1	4:p:162:MET:C	2.72	0.63
5:e:19:VAL:HG11	5:e:51:ILE:HD11	1.81	0.63
7:S:48:LYS:HZ3	7:R:43:PRO:C	2.06	0.63
7:S:68:LEU:CD1	7:T:66:TYR:CD1	2.82	0.63
9:F:100:VAL:HG12	9:F:256:MET:CE	2.24	0.63
9:D:100:VAL:CG1	9:D:256:MET:CE	2.76	0.63
1:a:116:PRO:HD3	2:b:35:SER:CA	2.27	0.63
7:G:43:PRO:C	7:H:48:LYS:HZ3	2.07	0.63
9:B:20:LEU:HD22	9:B:92:ILE:CG1	2.29	0.63
1:a:119:ILE:HB	2:b:31:LEU:CD2	2.28	0.63
2:b:125:LYS:HB3	4:p:185:LEU:CD1	2.28	0.63
3:d:115:PHE:CB	3:d:155:ILE:HD11	2.27	0.63
3:d:217:LEU:HD23	3:d:236:SER:OG	1.99	0.63
7:O:54:LEU:HD11	7:P:55:LEU:HD23	1.81	0.63
7:K:74:LEU:HD21	7:J:10:VAL:HG23	1.78	0.63
9:F:296:VAL:O	9:F:296:VAL:HG12	1.99	0.63
8:A:382:LYS:CB	8:A:442:THR:HG21	2.29	0.63
8:C:176:LYS:HE3	10:C:601:ATP:O1B	1.99	0.63
1:a:30:ILE:CG2	1:a:32:GLY:H	2.12	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:K:48:LYS:HZ3	7:J:43:PRO:CA	2.12	0.62
8:C:35:VAL:HG21	8:C:83:ILE:CG2	2.28	0.62
8:C:260:ILE:HD12	8:C:317:LEU:HD12	1.81	0.62
1:a:118:LYS:O	2:b:29:THR:CB	2.47	0.62
2:b:122:GLU:OE1	4:p:181:ILE:CB	2.46	0.62
3:d:195:ALA:HB1	3:d:207:VAL:CG1	2.29	0.62
6:g:79:VAL:CG1	6:g:306:ARG:HH22	2.11	0.62
1:a:195:LEU:CD2	4:p:84:PHE:CE1	2.82	0.62
7:J:48:LYS:HZ2	7:I:43:PRO:CB	2.10	0.62
8:E:77:MET:CE	8:E:112:LEU:HD21	2.30	0.62
8:C:100:VAL:HG23	8:C:246:LEU:CD2	2.30	0.62
2:b:176:PHE:HE1	3:d:233:VAL:CB	2.10	0.62
3:d:131:GLU:OE2	8:E:16:ARG:NH1	2.31	0.62
5:e:24:LEU:CD1	5:e:25:SER:O	2.48	0.62
6:g:123:VAL:CG1	6:g:308:LEU:HD22	2.18	0.62
9:D:132:ILE:O	9:D:312:ARG:NH1	2.31	0.62
1:a:70:ASN:CG	2:b:54:ASP:OD1	2.40	0.62
2:b:146:GLN:HA	2:b:149:LEU:HD12	1.81	0.62
6:g:72:VAL:HG12	6:g:257:LEU:CD2	2.28	0.62
7:Q:43:PRO:HB2	7:R:48:LYS:HZ2	1.61	0.62
7:Q:62:ALA:O	7:Q:65:ILE:N	2.29	0.62
7:M:80:PHE:CZ	7:L:81:VAL:CG2	2.82	0.62
7:J:48:LYS:HZ3	7:I:43:PRO:C	2.06	0.62
9:F:336:ASP:HB3	9:F:339:PRO:HD2	1.80	0.62
8:C:48:VAL:HG22	8:C:49:MET:N	2.15	0.62
8:C:152:ILE:HG22	8:C:423:GLN:HE21	1.62	0.62
7:T:54:LEU:HD11	7:G:55:LEU:HD23	1.80	0.62
8:E:55:GLU:HG3	8:E:55:GLU:O	1.99	0.62
9:D:254:LEU:HD13	9:D:313:ILE:HG12	1.81	0.62
9:D:296:VAL:HG12	9:D:296:VAL:O	1.99	0.62
1:a:100:THR:HG23	4:p:101:MET:CG	2.24	0.62
2:b:78:LEU:N	4:p:134:SER:O	2.32	0.62
2:b:81:ALA:HB1	4:p:139:GLN:C	2.25	0.62
3:d:228:GLU:OE2	8:A:43:HIS:CE1	2.51	0.62
9:B:132:ILE:O	9:B:312:ARG:NH1	2.29	0.62
9:B:204:GLU:OE2	9:B:274:ASN:HB2	2.00	0.62
1:a:24:GLN:CA	4:p:83:LEU:HG	2.30	0.62
1:a:133:ASP:OD1	1:a:133:ASP:O	2.18	0.62
2:b:69:GLU:CG	2:b:73:LYS:HE3	2.29	0.62
2:b:73:LYS:N	4:p:130:VAL:HG22	2.00	0.62
2:b:148:ALA:HA	4:p:211:SER:CB	2.30	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:177:VAL:HG11	3:d:210:LYS:CE	2.30	0.62
3:d:247:GLN:NE2	8:A:13:ILE:CA	2.62	0.62
5:e:119:LEU:HD13	5:e:123:ARG:CD	2.29	0.62
7:S:6:ALA:HB1	7:T:80:PHE:HD2	1.64	0.62
7:S:48:LYS:CG	7:R:46:GLU:OE1	2.48	0.62
7:Q:43:PRO:HB3	7:R:48:LYS:HD2	1.80	0.62
7:J:42:GLN:HE22	7:I:43:PRO:CG	2.10	0.62
1:a:39:VAL:CG2	1:a:133:ASP:CG	2.71	0.62
2:b:133:GLN:CB	8:A:3:THR:HG22	2.23	0.62
3:d:152:SER:CA	3:d:153:GLU:OE1	2.48	0.62
4:p:193:GLU:C	4:p:196:LYS:HG2	2.24	0.62
6:g:53:SER:HB2	9:D:403:ASP:CG	2.25	0.62
8:A:436:MET:HG3	8:A:490:LEU:HD11	1.80	0.62
9:B:29:PRO:HD2	9:B:288:LEU:HD13	1.82	0.62
9:B:426:LYS:NZ	9:B:467:ASP:HA	2.15	0.62
8:C:110:ASN:HB2	8:C:114:LYS:O	1.99	0.62
4:p:210:LEU:HD21	8:A:13:ILE:HG21	1.82	0.62
7:S:61:GLU:O	7:S:64:THR:HG22	2.00	0.62
7:N:33:GLY:HA3	7:O:31:ALA:HA	1.81	0.62
8:E:355:ARG:HH22	9:D:389:GLN:HE22	0.62	0.62
9:B:410:LEU:HD21	9:B:421:VAL:HG11	1.81	0.62
9:D:47:LEU:HB2	9:D:63:CYS:HB2	1.82	0.62
1:a:35:ILE:CB	4:p:88:LEU:H	2.13	0.61
2:b:78:LEU:N	4:p:135:SER:C	2.49	0.61
2:b:176:PHE:HE1	3:d:233:VAL:CG1	2.13	0.61
6:g:71:LYS:HB3	6:g:316:LEU:HD13	1.82	0.61
7:S:48:LYS:CB	7:R:46:GLU:OE1	2.48	0.61
7:O:46:GLU:OE1	7:P:48:LYS:HG2	2.00	0.61
7:H:54:LEU:HD11	7:I:55:LEU:HD23	1.82	0.61
8:E:34:GLN:HE21	8:E:41:ARG:HG3	1.65	0.61
8:E:284:ARG:NH1	8:E:330:TYR:HB2	2.15	0.61
8:A:379:VAL:HG12	8:A:438:ILE:CG2	2.30	0.61
9:B:179:THR:O	9:B:182:ILE:HG22	2.00	0.61
8:C:164:GLN:CG	8:C:165:ARG:N	2.63	0.61
9:D:275:ILE:HG21	9:D:327:VAL:HG22	1.81	0.61
4:p:192:LEU:HD22	8:A:3:THR:HG1	1.65	0.61
6:g:126:ASP:OD1	6:g:182:GLY:HA3	1.99	0.61
7:P:6:ALA:HB1	7:Q:80:PHE:HD2	1.63	0.61
7:L:31:ALA:HA	7:K:33:GLY:HA3	1.81	0.61
7:J:80:PHE:CE2	7:I:81:VAL:CG2	2.83	0.61
8:E:438:ILE:O	8:E:442:THR:HG23	1.99	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:387:ILE:CD1	9:F:458:PHE:HB2	2.25	0.61
8:A:367:VAL:O	8:A:367:VAL:HG23	1.99	0.61
9:B:50:LYS:HD3	9:B:59:MET:HE3	1.81	0.61
2:b:92:ALA:HA	4:p:152:ARG:H	1.65	0.61
2:b:140:VAL:O	2:b:144:VAL:HG23	1.99	0.61
3:d:148:ILE:CG1	8:C:16:ARG:HG2	2.31	0.61
6:g:144:ARG:CZ	6:g:242:ILE:HD12	2.30	0.61
6:g:253:GLU:O	6:g:253:GLU:HG2	2.00	0.61
7:J:74:LEU:HD22	7:I:10:VAL:HG22	1.80	0.61
8:A:327:VAL:HG12	8:A:327:VAL:O	1.99	0.61
8:C:98:ILE:HD12	8:C:242:THR:HG23	1.82	0.61
8:C:104:TYR:CD2	8:C:246:LEU:HD21	2.35	0.61
8:C:385:LEU:HD23	9:D:476:LEU:HD11	1.82	0.61
9:D:179:THR:O	9:D:182:ILE:CG2	2.44	0.61
2:b:77:GLN:CB	4:p:134:SER:HA	2.30	0.61
7:H:43:PRO:CG	7:I:42:GLN:NE2	2.62	0.61
7:L:48:LYS:CB	7:K:46:GLU:OE1	2.48	0.61
8:C:231:SER:HB2	8:C:232:PRO:HD2	1.83	0.61
8:C:413:ARG:NH1	8:C:442:THR:O	2.33	0.61
1:a:119:ILE:CD1	2:b:31:LEU:C	2.71	0.61
2:b:92:ALA:HA	4:p:152:ARG:N	2.14	0.61
6:g:110:THR:HG23	6:g:296:LEU:HD13	1.81	0.61
8:E:169:ILE:HG23	8:E:344:PHE:CD1	2.26	0.61
9:F:426:LYS:HD3	9:F:474:PHE:HE2	1.64	0.61
8:C:143:VAL:HG13	8:C:162:ARG:O	2.00	0.61
8:C:237:TYR:CE1	8:C:294:LEU:CD2	2.81	0.61
2:b:42:ILE:HG23	2:b:46:LYS:HE3	1.82	0.61
2:b:92:ALA:HB2	4:p:148:MET:O	2.00	0.61
3:d:224:ARG:HH12	3:d:232:LEU:HD13	1.59	0.61
4:p:203:LEU:HD22	8:A:7:ASP:CA	2.31	0.61
7:K:55:LEU:HD23	7:J:54:LEU:HD11	1.82	0.61
8:A:39:ILE:HD13	8:A:278:LEU:HD21	1.81	0.61
9:B:261:ARG:NH1	9:B:314:THR:O	2.33	0.61
9:B:426:LYS:HZ3	9:B:467:ASP:HA	1.64	0.61
8:C:389:GLN:OE1	8:C:389:GLN:HA	2.00	0.61
1:a:27:TYR:CZ	1:a:122:LEU:HD11	2.08	0.61
1:a:39:VAL:CG1	1:a:133:ASP:OD2	2.48	0.61
2:b:172:ASN:ND2	3:d:235:MET:O	2.33	0.61
3:d:175:THR:O	3:d:176:GLU:OE1	2.18	0.61
7:S:43:PRO:CA	7:T:48:LYS:HZ3	2.13	0.61
7:O:43:PRO:HB3	7:P:48:LYS:HZ3	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:M:48:LYS:HG2	7:L:46:GLU:OE1	2.01	0.61
2:b:81:ALA:N	4:p:141:GLU:HG3	2.16	0.61
2:b:118:LEU:HD13	4:p:177:GLY:C	2.11	0.61
6:g:73:ARG:NH1	9:F:412:GLU:OE2	2.33	0.61
7:N:43:PRO:HB2	7:O:48:LYS:HZ2	1.66	0.61
9:F:42:ASN:OD1	9:F:45:ASN:ND2	2.34	0.61
8:C:151:LEU:HD22	8:C:356:PRO:HG3	1.82	0.61
1:a:184:LEU:HD21	1:a:188:PHE:HE2	1.63	0.61
2:b:77:GLN:HB3	4:p:134:SER:HA	1.82	0.61
6:g:269:ASP:OD1	6:g:269:ASP:O	2.18	0.61
6:g:281:ILE:HB	7:M:41:ARG:CG	2.25	0.61
7:J:48:LYS:HZ2	7:I:43:PRO:HB2	1.64	0.61
8:E:130:ILE:HG23	8:E:241:TYR:HB3	1.82	0.61
8:A:237:TYR:CD2	8:A:274:MET:HE1	2.35	0.61
9:B:254:LEU:HD21	9:B:312:ARG:HB2	1.82	0.61
9:D:30:VAL:HG22	9:D:77:VAL:HG22	1.82	0.61
1:a:58:VAL:CG1	4:p:107:ILE:CG2	2.74	0.61
1:a:193:ASN:HD22	1:a:227:GLN:CD	2.08	0.61
2:b:145:PHE:HZ	3:d:243:GLU:C	2.09	0.61
3:d:115:PHE:HB3	3:d:155:ILE:HD13	1.79	0.61
5:e:21:GLU:CG	5:e:34:LEU:HD22	2.24	0.61
6:g:103:GLU:HG2	7:R:41:ARG:HH21	1.66	0.61
7:S:80:PHE:HD2	7:R:6:ALA:HB1	1.66	0.61
7:M:48:LYS:HZ3	7:L:43:PRO:CB	2.14	0.61
8:E:9:ILE:CG2	8:E:13:ILE:HG22	2.31	0.61
8:A:42:ILE:CD1	8:A:89:VAL:HG11	2.31	0.61
8:C:420:LEU:HD23	8:C:458:LEU:HD11	1.82	0.61
1:a:116:PRO:HG3	2:b:35:SER:N	2.13	0.60
1:a:121:GLN:HG3	2:b:25:ASP:CG	2.25	0.60
2:b:68:GLU:OE1	4:p:126:GLN:HG3	2.00	0.60
2:b:74:ALA:HB3	4:p:130:VAL:O	2.01	0.60
2:b:79:GLU:CG	4:p:136:GLU:HG2	2.30	0.60
3:d:154:ARG:NH2	8:C:5:ARG:NE	2.44	0.60
5:e:4:ASN:HB2	5:e:18:GLU:CG	2.31	0.60
7:Q:81:VAL:HG21	7:R:80:PHE:CE2	2.35	0.60
8:E:202:LYS:NZ	9:D:347:ASP:OD1	2.33	0.60
9:F:183:MET:HE1	9:F:215:GLU:CB	2.31	0.60
8:A:104:TYR:HE2	8:A:109:ILE:CD1	2.13	0.60
1:a:119:ILE:CD1	2:b:32:ILE:CG1	2.54	0.60
1:a:119:ILE:CG1	2:b:31:LEU:HG	2.31	0.60
2:b:95:PHE:HZ	4:p:156:SER:HB2	1.66	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:141:ARG:O	8:A:16:ARG:NH2	2.34	0.60
3:d:235:MET:HG3	8:A:24:VAL:HG11	1.82	0.60
6:g:111:LYS:HE3	6:g:113:ARG:HG2	1.82	0.60
7:T:43:PRO:CG	7:G:42:GLN:HE22	2.10	0.60
8:E:45:LEU:O	9:F:87:ARG:NH2	2.34	0.60
8:E:75:VAL:HG12	8:E:234:THR:OG1	2.00	0.60
1:a:118:LYS:HZ2	1:a:126:GLU:CB	2.14	0.60
4:p:203:LEU:HD22	8:A:7:ASP:HA	1.83	0.60
5:e:60:LEU:HD12	5:e:60:LEU:C	2.22	0.60
6:g:221:VAL:HG22	8:A:401:SER:CB	2.30	0.60
6:g:356:ILE:CD1	9:D:295:ALA:CA	2.69	0.60
6:g:356:ILE:HG12	9:D:294:SER:O	2.01	0.60
7:G:46:GLU:CD	7:H:48:LYS:HB3	2.25	0.60
7:G:54:LEU:HD11	7:H:55:LEU:CD2	2.31	0.60
8:C:35:VAL:HG21	8:C:83:ILE:HG21	1.83	0.60
1:a:121:GLN:HB3	2:b:25:ASP:OD1	2.00	0.60
4:p:214:ILE:C	8:A:17:ILE:CD1	2.73	0.60
7:P:43:PRO:CA	7:Q:48:LYS:HZ3	2.14	0.60
7:L:74:LEU:HD22	7:K:10:VAL:CG2	2.32	0.60
8:C:469:LYS:HZ3	8:C:492:GLU:HB2	1.64	0.60
1:a:119:ILE:HD13	2:b:31:LEU:C	2.27	0.60
2:b:67:SER:HB3	4:p:124:LYS:HA	1.82	0.60
6:g:327:THR:HG23	6:g:328:ASP:N	2.15	0.60
7:S:46:GLU:OE1	7:T:48:LYS:HG2	2.02	0.60
7:N:48:LYS:HD2	7:M:43:PRO:HB3	1.84	0.60
8:E:469:LYS:CE	8:E:493:ALA:HB2	2.32	0.60
9:F:292:MET:HE3	8:A:279:ARG:HG3	1.84	0.60
8:A:65:LEU:HD12	8:A:75:VAL:HG21	1.83	0.60
8:C:237:TYR:HE1	8:C:294:LEU:HD21	1.65	0.60
1:a:86:GLN:HA	1:a:86:GLN:HE21	1.67	0.60
1:a:165:ILE:HG21	1:a:169:PRO:C	2.27	0.60
2:b:80:LYS:C	4:p:141:GLU:HG3	2.27	0.60
2:b:82:ARG:HG3	2:b:85:LEU:HD21	0.71	0.60
2:b:91:ASP:C	4:p:151:ALA:CB	2.64	0.60
4:p:127:LEU:HA	4:p:130:VAL:HG12	1.81	0.60
6:g:363:CYS:CB	9:D:290:GLY:O	2.49	0.60
7:G:43:PRO:CG	7:H:42:GLN:NE2	2.60	0.60
8:E:343:ILE:CG2	8:E:358:ILE:HD12	2.31	0.60
9:F:100:VAL:HG13	9:F:256:MET:CE	2.29	0.60
1:a:58:VAL:CG1	4:p:108:TYR:HA	2.31	0.60
2:b:64:ILE:HD11	4:p:120:ASP:CA	2.17	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:83:ALA:N	4:p:140:LEU:CB	2.33	0.60
6:g:360:ALA:HA	9:D:292:MET:HA	1.82	0.60
7:O:33:GLY:HA3	7:P:31:ALA:HA	1.84	0.60
7:K:49:ILE:HD11	7:J:40:ALA:CB	2.25	0.60
8:E:260:ILE:HD13	8:E:317:LEU:HB2	1.83	0.60
8:E:479:THR:HG22	8:E:481:THR:HB	1.84	0.60
8:A:439:TYR:OH	8:A:491:LYS:HE2	2.02	0.60
9:B:456:ARG:NH2	9:B:481:ASP:OD1	2.34	0.60
9:D:421:VAL:O	9:D:425:ARG:HG2	2.02	0.60
1:a:40:LEU:HG	4:p:89:THR:HB	1.77	0.60
1:a:118:LYS:O	2:b:30:ASN:OD1	2.19	0.60
2:b:81:ALA:HA	4:p:141:GLU:HG3	1.83	0.60
2:b:107:LYS:HA	4:p:166:THR:HG21	0.61	0.60
4:p:210:LEU:CD2	8:A:13:ILE:HG21	2.32	0.60
6:g:100:LEU:HD11	6:g:105:VAL:HG21	1.82	0.60
8:A:190:GLN:OE1	8:A:190:GLN:N	2.23	0.60
9:B:408:LEU:HD13	9:B:412:GLU:HG3	1.83	0.60
1:a:58:VAL:HA	4:p:112:LEU:HD11	1.84	0.60
1:a:64:ILE:HG22	2:b:57:LYS:HE2	1.82	0.60
1:a:137:THR:CB	1:a:191:PHE:HB2	2.31	0.60
1:a:212:VAL:N	1:a:213:PRO:CD	2.64	0.60
9:F:407:ILE:HG22	9:F:408:LEU:HD23	1.83	0.60
8:A:39:ILE:O	8:A:39:ILE:HG23	2.02	0.60
9:D:336:ASP:HB3	9:D:339:PRO:HD2	1.84	0.60
2:b:60:ILE:O	2:b:63:THR:CG2	2.47	0.60
3:d:148:ILE:HG13	8:C:20:TYR:CD2	2.36	0.60
7:M:66:TYR:CG	7:L:68:LEU:HD22	2.37	0.60
7:L:80:PHE:CE2	7:K:81:VAL:HG21	2.37	0.60
8:E:157:MET:HG2	8:E:384:LYS:HG2	1.84	0.60
8:A:9:ILE:HD13	8:A:12:ILE:HD12	1.83	0.60
8:C:104:TYR:CE2	8:C:246:LEU:HD21	2.37	0.60
9:D:385:TYR:O	9:D:389:GLN:HG2	2.02	0.60
4:p:152:ARG:HB3	8:A:503:LEU:HD22	1.79	0.59
5:e:4:ASN:CB	5:e:18:GLU:CD	2.75	0.59
7:Q:43:PRO:CB	7:R:48:LYS:HZ2	2.11	0.59
7:L:74:LEU:CD2	7:K:10:VAL:HG23	2.32	0.59
8:E:157:MET:SD	8:E:387:LEU:HD12	2.42	0.59
8:A:378:LYS:HD3	8:A:482:PHE:HB3	1.84	0.59
8:A:445:TYR:CE2	8:A:494:ILE:HG21	2.36	0.59
8:A:446:LEU:HD22	8:A:449:LEU:HD12	1.83	0.59
8:C:164:GLN:CG	8:C:165:ARG:H	2.15	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:24:GLN:C	4:p:83:LEU:CD1	2.74	0.59
1:a:40:LEU:HD21	4:p:89:THR:CB	2.31	0.59
1:a:133:ASP:C	1:a:191:PHE:HE1	2.10	0.59
1:a:193:ASN:HD22	1:a:227:GLN:CG	2.10	0.59
2:b:60:ILE:HD11	4:p:120:ASP:OD2	2.02	0.59
2:b:78:LEU:O	4:p:140:LEU:N	2.29	0.59
3:d:224:ARG:HG2	3:d:232:LEU:HD12	1.83	0.59
3:d:242:GLU:CG	8:A:20:TYR:CE1	2.84	0.59
4:p:214:ILE:HG13	4:p:215:VAL:H	1.63	0.59
5:e:34:LEU:HD12	7:L:42:GLN:HE22	1.66	0.59
6:g:353:ILE:HG12	8:E:284:ARG:HA	1.83	0.59
7:P:46:GLU:OE1	7:Q:48:LYS:HG2	2.01	0.59
8:E:42:ILE:HD11	8:E:56:PHE:HZ	1.61	0.59
8:A:106:GLY:HA2	8:A:219:MET:HG3	1.83	0.59
8:A:469:LYS:HZ1	8:A:492:GLU:CB	2.14	0.59
9:B:112:ASN:HB3	9:B:114:LEU:H	1.67	0.59
9:B:425:ARG:HB3	9:B:429:ARG:HH12	1.67	0.59
8:A:104:TYR:CE2	8:A:109:ILE:HD12	2.37	0.59
1:a:144:THR:HG21	1:a:186:LEU:HD21	1.84	0.59
2:b:77:GLN:HG2	4:p:138:LYS:N	2.17	0.59
2:b:95:PHE:HB3	4:p:151:ALA:C	2.18	0.59
4:p:213:ASP:OD2	8:A:14:ARG:CB	2.50	0.59
6:g:124:THR:CG2	6:g:137:LEU:CD2	2.61	0.59
7:Q:33:GLY:HA3	7:R:31:ALA:HA	1.85	0.59
8:E:413:ARG:HH12	8:E:444:GLY:H	1.49	0.59
8:A:373:ILE:O	8:A:377:LYS:HG3	2.02	0.59
8:C:237:TYR:HE1	8:C:294:LEU:HD22	1.64	0.59
9:D:449:VAL:HG13	9:D:453:GLU:OE1	2.01	0.59
1:a:172:LEU:N	1:a:173:PRO:HD2	2.18	0.59
2:b:60:ILE:HD11	4:p:120:ASP:HB2	1.83	0.59
7:G:46:GLU:OE1	7:H:48:LYS:HD3	2.01	0.59
7:J:74:LEU:HD21	7:I:75:LEU:HD21	1.85	0.59
8:A:382:LYS:HD2	8:A:442:THR:HG21	1.82	0.59
2:b:128:THR:O	2:b:132:GLU:HG2	2.02	0.59
5:e:96:LEU:CD1	5:e:123:ARG:CG	2.80	0.59
6:g:53:SER:CB	9:D:403:ASP:CG	2.75	0.59
7:O:46:GLU:OE1	7:P:48:LYS:HD3	2.02	0.59
8:A:22:ARG:HG2	8:A:22:ARG:O	2.03	0.59
2:b:77:GLN:CG	4:p:138:LYS:HG2	2.28	0.59
2:b:144:VAL:CB	8:A:9:ILE:CG2	2.80	0.59
4:p:179:LYS:O	4:p:179:LYS:HD3	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:253:GLU:HA	6:g:270:MET:HA	1.84	0.59
9:B:148:ILE:HD12	9:B:380:VAL:HG12	1.84	0.59
8:C:27:VAL:HB	8:C:47:GLU:HB2	0.88	0.59
9:D:22:ARG:NH1	9:D:88:GLY:O	2.36	0.59
6:g:231:LEU:HD21	6:g:300:LEU:HD12	1.83	0.59
9:F:405:ILE:HD11	9:F:421:VAL:HG21	1.84	0.59
8:C:392:GLU:OE1	9:D:429:ARG:NH2	2.35	0.59
1:a:53:SER:CA	1:a:56:ILE:HG22	2.32	0.59
1:a:212:VAL:CB	1:a:213:PRO:HD3	2.21	0.59
5:e:110:ARG:HD2	6:g:206:GLU:OE2	2.03	0.59
7:O:5:ILE:H	7:O:5:ILE:HD12	1.68	0.59
7:L:48:LYS:HD2	7:K:43:PRO:HB3	1.85	0.59
8:E:104:TYR:CE1	8:E:122:ILE:HG21	2.38	0.59
8:E:141:ARG:NH1	8:E:311:GLU:OE2	2.35	0.59
8:E:383:LEU:CD1	8:E:417:LEU:HD22	2.33	0.59
8:A:495:GLN:OE1	8:A:495:GLN:HA	2.03	0.59
9:B:134:ARG:NE	9:B:258:GLU:OE2	2.31	0.59
9:B:216:MET:CE	9:B:232:VAL:HG11	2.32	0.59
9:B:272:ILE:HG21	9:B:275:ILE:CG2	2.33	0.59
9:B:272:ILE:HG21	9:B:275:ILE:HG22	1.85	0.59
9:B:453:GLU:OE1	9:B:453:GLU:HA	2.03	0.59
2:b:145:PHE:H	8:A:13:ILE:CD1	2.15	0.59
4:p:122:SER:O	4:p:126:GLN:HG2	2.02	0.59
6:g:292:LEU:HD12	6:g:292:LEU:C	2.27	0.59
7:P:5:ILE:H	7:P:5:ILE:HD12	1.68	0.59
7:P:54:LEU:HD11	7:Q:55:LEU:HD23	1.85	0.59
7:K:5:ILE:H	7:K:5:ILE:HD12	1.68	0.59
9:B:410:LEU:HD23	9:B:413:LEU:CD1	2.33	0.59
1:a:70:ASN:CB	2:b:53:LEU:HB3	2.33	0.58
2:b:80:LYS:CA	4:p:137:VAL:HB	2.33	0.58
2:b:91:ASP:HB3	4:p:148:MET:HG2	1.84	0.58
7:O:43:PRO:HB3	7:P:48:LYS:HD2	1.84	0.58
7:H:5:ILE:H	7:H:5:ILE:HD12	1.68	0.58
9:B:169:GLY:HA3	9:B:346:LEU:CD1	2.29	0.58
1:a:24:GLN:CA	4:p:83:LEU:HD11	2.16	0.58
1:a:40:LEU:CD2	4:p:92:ILE:HB	2.34	0.58
1:a:116:PRO:CA	2:b:34:LEU:N	2.62	0.58
2:b:141:ARG:N	8:A:9:ILE:CD1	2.65	0.58
3:d:87:ASP:HB2	3:d:172:ILE:CD1	2.33	0.58
3:d:132:ILE:HA	8:E:13:ILE:HD13	1.83	0.58
5:e:110:ARG:HH21	6:g:201:LEU:HD13	1.66	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:N:5:ILE:H	7:N:5:ILE:HD12	1.68	0.58
7:N:81:VAL:CG2	7:O:80:PHE:CE2	2.86	0.58
7:M:32:ALA:HA	7:M:53:LEU:HD11	1.85	0.58
7:J:5:ILE:HD12	7:J:5:ILE:H	1.68	0.58
9:F:50:LYS:HD2	9:F:90:GLU:OE2	2.03	0.58
9:D:427:ILE:HD11	9:D:462:LEU:HD21	1.85	0.58
1:a:135:ASN:CG	4:p:86:PHE:CA	2.73	0.58
2:b:103:ILE:CB	4:p:162:MET:HB2	2.33	0.58
2:b:103:ILE:HD11	4:p:162:MET:C	2.28	0.58
6:g:227:ILE:HG23	6:g:242:ILE:HG12	1.85	0.58
7:S:31:ALA:HA	7:R:33:GLY:HA3	1.84	0.58
9:F:110:ILE:CG2	9:F:118:VAL:HG23	2.31	0.58
8:A:107:ARG:NH1	8:A:120:GLY:O	2.35	0.58
9:B:243:PRO:HB3	9:B:284:GLU:OE1	2.03	0.58
8:C:134:ALA:HB3	9:D:240:ASN:HD22	1.67	0.58
8:C:394:GLU:O	8:C:398:GLN:NE2	2.36	0.58
8:C:432:GLU:CG	8:C:476:ILE:HG21	2.26	0.58
9:D:102:VAL:HG23	9:D:128:THR:CG2	2.33	0.58
2:b:80:LYS:H	4:p:137:VAL:HB	1.65	0.58
2:b:180:ASN:ND2	3:d:201:ILE:CG2	2.54	0.58
3:d:84:ASP:O	3:d:88:VAL:HG23	2.04	0.58
5:e:100:GLU:CG	5:e:119:LEU:HD21	2.33	0.58
6:g:217:PHE:HB2	6:g:314:SER:OG	2.02	0.58
6:g:281:ILE:HG21	7:M:41:ARG:CA	2.24	0.58
7:O:10:VAL:CG2	7:P:74:LEU:HD21	2.25	0.58
2:b:107:LYS:HE3	4:p:166:THR:N	2.18	0.58
7:P:43:PRO:CG	7:Q:42:GLN:HE22	2.12	0.58
7:T:5:ILE:H	7:T:5:ILE:HD12	1.68	0.58
8:A:416:ARG:HD3	8:A:446:LEU:O	2.03	0.58
2:b:64:ILE:HD13	4:p:119:ARG:O	2.03	0.58
2:b:77:GLN:H	4:p:134:SER:HA	1.67	0.58
2:b:81:ALA:HB1	4:p:142:GLU:H	1.67	0.58
5:e:96:LEU:HD13	5:e:123:ARG:CA	2.33	0.58
6:g:121:MET:CE	6:g:160:ILE:HD11	2.33	0.58
9:B:216:MET:HE3	9:B:221:VAL:HG11	1.85	0.58
9:B:387:ILE:HD12	9:B:459:GLN:HG3	1.85	0.58
9:D:313:ILE:O	9:D:313:ILE:HG22	2.04	0.58
1:a:54:ALA:O	1:a:58:VAL:HG23	2.04	0.58
2:b:82:ARG:CB	2:b:85:LEU:CD2	2.80	0.58
2:b:82:ARG:HA	2:b:85:LEU:CG	2.34	0.58
3:d:75:ALA:CB	3:d:160:GLU:HG2	2.33	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:e:51:ILE:HG23	5:e:51:ILE:O	2.03	0.58
6:g:217:PHE:O	6:g:268:ARG:HD2	2.02	0.58
6:g:251:GLU:CG	6:g:272:LYS:CG	2.81	0.58
7:S:55:LEU:HD23	7:R:54:LEU:HD11	1.86	0.58
7:J:74:LEU:HD21	7:I:10:VAL:CG2	2.26	0.58
9:F:178:LYS:NZ	9:F:182:ILE:HD11	2.19	0.58
8:A:131:GLU:OE2	8:A:297:ARG:NH2	2.37	0.58
9:D:407:ILE:HG22	9:D:408:LEU:HD23	1.85	0.58
3:d:129:LEU:HD22	3:d:150:ILE:HD12	1.85	0.58
5:e:60:LEU:HB2	5:e:82:GLU:O	2.03	0.58
6:g:229:THR:HG22	6:g:240:CYS:O	2.04	0.58
7:S:54:LEU:HD11	7:T:55:LEU:HD23	1.85	0.58
7:H:32:ALA:HA	7:H:53:LEU:HD11	1.86	0.58
8:E:183:THR:O	8:E:187:GLN:OE1	2.21	0.58
8:C:170:GLY:HA3	8:C:174:THR:HG21	1.86	0.58
8:C:292:PHE:HE1	9:D:246:ARG:NH1	2.02	0.58
1:a:97:PHE:HA	4:p:101:MET:HE2	1.84	0.58
1:a:120:ILE:CG1	2:b:25:ASP:O	2.48	0.58
1:a:136:THR:CA	4:p:90:LEU:HD22	2.32	0.58
2:b:133:GLN:HG2	8:A:3:THR:N	2.18	0.58
2:b:141:ARG:HD3	8:A:12:ILE:CG2	2.03	0.58
3:d:227:ASN:OD1	8:A:46:ASP:HB3	2.03	0.58
7:N:43:PRO:CA	7:O:48:LYS:HZ3	2.16	0.58
7:R:5:ILE:HD12	7:R:5:ILE:H	1.69	0.58
7:G:32:ALA:HA	7:G:53:LEU:HD11	1.86	0.58
7:K:32:ALA:HA	7:K:53:LEU:HD11	1.85	0.58
8:C:104:TYR:CE1	8:C:122:ILE:CD1	2.86	0.58
1:a:121:GLN:CG	2:b:25:ASP:OD2	2.51	0.58
1:a:165:ILE:HG21	1:a:169:PRO:CA	2.33	0.58
2:b:103:ILE:HD12	4:p:158:ALA:O	1.99	0.58
2:b:145:PHE:CD2	2:b:149:LEU:CD1	2.87	0.58
3:d:175:THR:HG22	3:d:206:ASN:HB3	1.86	0.58
7:S:32:ALA:HA	7:S:53:LEU:HD11	1.86	0.58
7:N:32:ALA:HA	7:N:53:LEU:HD11	1.85	0.58
7:L:74:LEU:HD21	7:K:10:VAL:HG23	1.86	0.58
8:E:465:VAL:HG22	8:E:493:ALA:HB1	1.86	0.58
8:E:489:LEU:HD23	8:E:489:LEU:C	2.27	0.58
8:A:45:LEU:O	9:B:87:ARG:NH2	2.37	0.58
1:a:149:PHE:HZ	1:a:163:LYS:HZ2	1.49	0.57
1:a:208:VAL:HG22	1:a:209:PRO:N	2.18	0.57
5:e:46:ILE:HG22	5:e:63:MET:O	2.03	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:H:33:GLY:HA3	7:I:31:ALA:HA	1.86	0.57
7:H:43:PRO:HB2	7:I:48:LYS:HZ2	1.68	0.57
8:E:379:VAL:HG11	8:E:438:ILE:HB	1.84	0.57
9:B:384:HIS:O	9:B:387:ILE:HG22	2.04	0.57
2:b:88:VAL:O	2:b:89:GLU:C	2.46	0.57
4:p:152:ARG:CG	8:A:503:LEU:CD2	2.81	0.57
6:g:221:VAL:HG22	8:A:401:SER:HB3	1.84	0.57
7:N:10:VAL:HG22	7:O:74:LEU:HD22	1.85	0.57
7:P:32:ALA:HA	7:P:53:LEU:HD11	1.85	0.57
7:L:5:ILE:H	7:L:5:ILE:HD12	1.68	0.57
7:K:42:GLN:NE2	7:J:43:PRO:CG	2.67	0.57
7:J:32:ALA:HA	7:J:53:LEU:HD11	1.85	0.57
9:B:203:GLY:O	9:B:277:ARG:HG3	2.05	0.57
3:d:180:VAL:HG13	3:d:223:ILE:HG22	1.85	0.57
4:p:199:THR:HG23	4:p:200:ILE:N	2.19	0.57
6:g:280:PRO:HG2	7:N:42:GLN:NE2	2.20	0.57
7:O:32:ALA:HA	7:O:53:LEU:HD11	1.86	0.57
7:J:80:PHE:CZ	7:I:81:VAL:CG2	2.87	0.57
8:E:344:PHE:CE2	8:E:362:ILE:HG22	2.28	0.57
8:C:9:ILE:O	8:C:9:ILE:HG12	2.03	0.57
1:a:132:ASN:O	1:a:191:PHE:HE1	1.87	0.57
1:a:134:ILE:HD12	1:a:135:ASN:H	1.60	0.57
1:a:139:ALA:HB1	4:p:94:MET:HE2	1.77	0.57
3:d:242:GLU:HG2	8:A:20:TYR:CE1	2.38	0.57
6:g:229:THR:O	6:g:229:THR:CG2	2.50	0.57
7:N:46:GLU:OE1	7:O:48:LYS:HD3	2.04	0.57
7:T:32:ALA:HA	7:T:53:LEU:HD11	1.85	0.57
1:a:53:SER:HA	1:a:56:ILE:CG2	2.34	0.57
1:a:119:ILE:CB	2:b:28:ALA:C	2.78	0.57
2:b:103:ILE:HG12	4:p:162:MET:CA	2.34	0.57
2:b:144:VAL:HB	8:A:13:ILE:CG1	2.35	0.57
3:d:177:VAL:CG1	3:d:210:LYS:CD	2.83	0.57
3:d:186:LEU:C	3:d:186:LEU:HD12	2.30	0.57
7:L:32:ALA:HA	7:L:53:LEU:HD11	1.85	0.57
8:E:9:ILE:HD12	8:E:9:ILE:N	2.20	0.57
8:A:419:GLU:OE2	8:A:422:LYS:CE	2.52	0.57
9:D:274:ASN:ND2	9:D:276:PHE:HB3	2.20	0.57
1:a:195:LEU:HD11	4:p:84:PHE:CZ	2.40	0.57
2:b:70:LEU:HB2	4:p:127:LEU:CD2	2.33	0.57
6:g:89:LEU:HD21	6:g:299:TYR:HA	1.86	0.57
7:Q:5:ILE:H	7:Q:5:ILE:HD12	1.68	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:469:LYS:NZ	8:A:493:ALA:N	2.52	0.57
8:C:12:ILE:CD1	8:C:33:LEU:HD21	2.34	0.57
2:b:74:ALA:CA	4:p:130:VAL:O	2.52	0.57
4:p:204:ASP:O	4:p:207:ILE:HB	2.04	0.57
5:e:82:GLU:OE2	6:g:189:LYS:NZ	2.35	0.57
6:g:73:ARG:HH11	9:F:412:GLU:CG	2.17	0.57
6:g:360:ALA:HB2	9:D:292:MET:HB3	1.87	0.57
8:A:216:ARG:HH21	8:A:216:ARG:HG3	1.70	0.57
8:C:104:TYR:HD2	8:C:246:LEU:CD2	2.18	0.57
1:a:30:ILE:HG22	1:a:32:GLY:H	1.70	0.57
1:a:180:PHE:O	1:a:183:PRO:HD2	2.04	0.57
2:b:117:THR:CG2	4:p:174:LEU:HD21	2.34	0.57
2:b:131:PHE:HE1	8:A:119:ARG:C	2.12	0.57
2:b:145:PHE:HB2	8:A:13:ILE:HD12	1.86	0.57
3:d:148:ILE:HD11	8:C:16:ARG:C	2.30	0.57
6:g:362:ALA:HB1	8:C:281:PRO:CA	2.35	0.57
7:G:5:ILE:H	7:G:5:ILE:HD12	1.68	0.57
7:L:48:LYS:HZ2	7:K:43:PRO:CB	2.18	0.57
8:A:157:MET:HA	8:A:384:LYS:CE	2.34	0.57
8:A:374:LYS:HB2	8:A:480:LYS:O	2.05	0.57
8:C:98:ILE:CD1	8:C:242:THR:HG23	2.35	0.57
1:a:43:SER:OG	4:p:93:ILE:CB	2.53	0.57
1:a:119:ILE:HA	2:b:28:ALA:CA	2.34	0.57
2:b:96:ARG:HA	4:p:155:ILE:N	2.06	0.57
2:b:152:ALA:CB	4:p:214:ILE:HD11	2.33	0.57
3:d:195:ALA:HB1	3:d:207:VAL:HG13	1.87	0.57
7:S:5:ILE:H	7:S:5:ILE:HD12	1.69	0.57
7:H:10:VAL:CG2	7:I:74:LEU:HD21	2.29	0.57
7:H:43:PRO:HB3	7:I:48:LYS:CD	2.35	0.57
8:E:475:ILE:HG21	8:E:486:ALA:HB2	1.86	0.57
9:D:143:ASP:OD1	9:D:315:SER:OG	2.23	0.57
9:D:179:THR:C	9:D:182:ILE:HG22	2.27	0.57
3:d:132:ILE:HG13	8:E:13:ILE:HD13	1.87	0.57
3:d:148:ILE:HD13	8:C:17:ILE:HG12	1.85	0.57
6:g:50:ARG:HG2	6:g:50:ARG:NH1	2.20	0.57
7:N:81:VAL:CG2	7:O:80:PHE:CZ	2.88	0.57
7:M:5:ILE:HD12	7:M:5:ILE:H	1.68	0.57
8:E:284:ARG:NH1	8:E:330:TYR:CD1	2.66	0.57
9:F:50:LYS:NZ	9:F:92:ILE:HD12	2.19	0.57
8:C:240:PRO:HG3	8:C:267:GLN:CD	2.30	0.57
2:b:144:VAL:HB	8:A:13:ILE:HD11	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:198:VAL:HA	3:d:201:ILE:HG12	1.87	0.56
6:g:167:TYR:OH	6:g:171:ARG:NH2	2.38	0.56
7:S:33:GLY:HA3	7:T:31:ALA:HA	1.86	0.56
7:R:62:ALA:CA	7:R:65:ILE:HG22	2.35	0.56
7:T:46:GLU:CD	7:G:48:LYS:HB3	2.30	0.56
7:L:80:PHE:HD2	7:K:6:ALA:HB1	1.70	0.56
9:F:185:LEU:HD13	9:F:324:ILE:HD13	1.86	0.56
8:A:419:GLU:OE2	8:A:419:GLU:HA	2.03	0.56
8:A:432:GLU:C	8:A:435:VAL:HG12	2.30	0.56
9:B:71:ASN:H	8:C:9:ILE:HG13	1.70	0.56
8:C:9:ILE:HD11	8:C:85:GLU:OE2	2.04	0.56
8:C:247:ALA:HB3	8:C:314:MET:HE3	1.86	0.56
8:C:308:LEU:CD2	8:C:309:LEU:CD1	2.82	0.56
1:a:115:LEU:HD23	2:b:38:LEU:CD1	2.13	0.56
2:b:103:ILE:CG1	4:p:162:MET:CA	2.82	0.56
2:b:125:LYS:HD2	4:p:185:LEU:CB	2.35	0.56
2:b:180:ASN:HB3	3:d:201:ILE:CG2	2.30	0.56
7:G:62:ALA:CA	7:G:65:ILE:HG22	2.36	0.56
7:J:55:LEU:HD23	7:I:54:LEU:HD11	1.86	0.56
9:F:184:GLU:HG3	9:F:188:ASN:ND2	2.20	0.56
8:A:435:VAL:HG13	8:A:436:MET:N	2.19	0.56
9:B:221:VAL:HG13	9:B:232:VAL:HG23	1.84	0.56
1:a:165:ILE:CG2	1:a:169:PRO:HA	2.35	0.56
1:a:165:ILE:HG21	1:a:169:PRO:HA	1.86	0.56
2:b:70:LEU:H	4:p:130:VAL:HG11	1.60	0.56
2:b:122:GLU:OE1	4:p:181:ILE:HD12	2.05	0.56
2:b:176:PHE:HE1	3:d:233:VAL:HG11	1.68	0.56
3:d:91:THR:OG1	3:d:141:HIS:HE1	1.86	0.56
4:p:196:LYS:O	4:p:199:THR:CG2	2.46	0.56
6:g:261:GLU:HG3	9:F:397:ARG:HH22	1.56	0.56
6:g:284:PHE:HD1	6:g:291:ILE:HD12	1.69	0.56
7:S:40:ALA:CB	7:T:49:ILE:HD11	2.33	0.56
7:Q:32:ALA:HA	7:Q:53:LEU:HD11	1.85	0.56
7:Q:46:GLU:OE1	7:R:48:LYS:HG2	2.05	0.56
7:R:32:ALA:HA	7:R:53:LEU:HD11	1.86	0.56
7:T:43:PRO:CG	7:G:42:GLN:NE2	2.67	0.56
7:I:32:ALA:HA	7:I:53:LEU:HD11	1.86	0.56
9:B:50:LYS:NZ	9:B:92:ILE:HD12	2.20	0.56
9:B:402:GLN:HA	9:B:405:ILE:CG1	2.35	0.56
8:C:169:ILE:HD11	8:C:322:THR:HG21	1.87	0.56
1:a:120:ILE:HG22	2:b:30:ASN:HB2	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:145:PHE:HE2	2:b:149:LEU:HD11	1.62	0.56
2:b:160:LEU:CG	2:b:164:LEU:HB3	2.16	0.56
7:N:55:LEU:HD23	7:M:54:LEU:HD11	1.86	0.56
7:T:62:ALA:CA	7:T:65:ILE:HG22	2.36	0.56
7:L:62:ALA:CA	7:L:65:ILE:HG22	2.36	0.56
7:J:62:ALA:CA	7:J:65:ILE:HG22	2.36	0.56
8:A:61:ILE:HG22	8:A:77:MET:CE	2.35	0.56
8:A:109:ILE:HD11	8:A:113:ALA:HB1	1.85	0.56
8:A:362:ILE:CG2	8:A:362:ILE:O	2.52	0.56
8:A:446:LEU:HD21	8:A:457:TYR:CD2	2.40	0.56
8:C:181:THR:HG23	8:C:213:PHE:HE1	1.70	0.56
9:D:144:THR:O	9:D:144:THR:HG22	2.04	0.56
9:D:155:VAL:HG22	9:D:431:LEU:HB3	1.86	0.56
2:b:80:LYS:HG2	4:p:137:VAL:CB	2.36	0.56
2:b:131:PHE:CE1	8:A:119:ARG:C	2.84	0.56
4:p:219:LEU:HD12	4:p:219:LEU:C	2.31	0.56
6:g:343:ARG:HH21	8:A:326:ASP:HB2	1.71	0.56
7:S:74:LEU:HD21	7:R:75:LEU:HD21	1.88	0.56
7:N:48:LYS:HG2	7:M:46:GLU:OE1	2.06	0.56
7:L:48:LYS:CG	7:K:46:GLU:OE1	2.53	0.56
7:K:48:LYS:HB3	7:J:46:GLU:CD	2.30	0.56
8:E:140:ARG:HE	9:F:206:THR:HG21	1.66	0.56
8:E:487:GLU:O	8:E:491:LYS:HG3	2.06	0.56
9:F:336:ASP:OD1	9:F:337:PRO:CD	2.52	0.56
9:F:487:ALA:O	9:F:491:GLU:HG3	2.04	0.56
1:a:35:ILE:HG22	4:p:88:LEU:CB	2.31	0.56
1:a:184:LEU:CD2	1:a:188:PHE:HD2	1.97	0.56
3:d:132:ILE:CA	8:E:13:ILE:HD13	2.36	0.56
4:p:195:GLN:NE2	8:A:4:ILE:HD12	2.21	0.56
5:e:22:ILE:HG22	5:e:51:ILE:CD1	2.34	0.56
7:N:48:LYS:HB3	7:M:46:GLU:CD	2.29	0.56
8:E:9:ILE:HG22	8:E:14:ARG:HG3	1.87	0.56
8:A:131:GLU:CD	8:A:297:ARG:NH2	2.64	0.56
8:A:430:THR:CG2	8:A:432:GLU:OE1	2.53	0.56
9:B:247:MET:SD	9:B:285:VAL:HG21	2.46	0.56
8:C:165:ARG:CD	8:C:299:LEU:O	2.49	0.56
8:C:237:TYR:CE1	8:C:294:LEU:CD1	2.88	0.56
1:a:24:GLN:C	4:p:83:LEU:HD12	2.30	0.56
2:b:129:ILE:CD1	4:p:188:ALA:CB	2.72	0.56
3:d:76:SER:CA	3:d:164:GLU:HG2	2.33	0.56
3:d:148:ILE:HG12	8:C:16:ARG:CG	2.33	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:25:GLN:HE21	9:F:27:ILE:HD12	1.70	0.56
8:A:77:MET:SD	8:A:238:LEU:HD22	2.46	0.56
8:A:461:LEU:C	8:A:461:LEU:HD23	2.31	0.56
8:C:156:ALA:HB2	8:C:438:ILE:HD11	1.88	0.56
8:C:449:LEU:HD11	8:C:454:VAL:HG22	1.86	0.56
6:g:349:ILE:CD1	8:E:284:ARG:HH21	2.04	0.56
7:S:62:ALA:CA	7:S:65:ILE:HG22	2.35	0.56
7:Q:43:PRO:HB3	7:R:48:LYS:CD	2.36	0.56
7:H:43:PRO:CA	7:I:48:LYS:HZ3	2.18	0.56
9:D:86:THR:H	9:D:89:MET:HE2	1.70	0.56
1:a:195:LEU:HD21	4:p:84:PHE:HZ	1.70	0.56
6:g:85:PHE:HE1	6:g:302:SER:OG	1.87	0.56
8:E:159:PRO:HB2	8:E:371:ALA:HB3	1.86	0.56
8:A:98:ILE:HG13	8:A:98:ILE:O	2.05	0.56
8:A:359:ASN:ND2	8:A:362:ILE:CD1	2.57	0.56
1:a:54:ALA:CB	4:p:104:LEU:HD13	2.34	0.56
1:a:161:PHE:HZ	1:a:163:LYS:HD2	0.78	0.56
2:b:79:GLU:CG	4:p:136:GLU:CG	2.84	0.56
7:P:62:ALA:CA	7:P:65:ILE:HG22	2.35	0.56
7:M:74:LEU:HD21	7:L:75:LEU:HD21	1.87	0.56
7:J:48:LYS:HD2	7:I:43:PRO:HB3	1.87	0.56
8:E:267:GLN:NE2	8:E:294:LEU:HD21	2.21	0.56
8:E:497:GLN:OE1	8:E:497:GLN:HA	2.05	0.56
9:B:71:ASN:H	8:C:9:ILE:CD1	2.19	0.56
9:B:150:GLU:OE2	9:B:450:GLY:HA2	2.05	0.56
8:C:104:TYR:HD1	8:C:122:ILE:HD13	1.63	0.56
9:D:102:VAL:HG23	9:D:128:THR:HG21	1.87	0.56
9:D:169:GLY:HA3	9:D:346:LEU:HD13	1.88	0.56
1:a:64:ILE:HG22	2:b:57:LYS:CE	2.36	0.55
1:a:134:ILE:N	4:p:86:PHE:CE1	2.74	0.55
3:d:176:GLU:OE1	3:d:176:GLU:HA	2.06	0.55
4:p:152:ARG:CB	8:A:503:LEU:HD22	2.33	0.55
4:p:159:LEU:HD12	4:p:159:LEU:C	2.30	0.55
5:e:52:ARG:HG2	5:e:57:TRP:HB3	1.87	0.55
5:e:62:LEU:C	5:e:62:LEU:HD12	2.31	0.55
7:Q:46:GLU:CD	7:R:48:LYS:HB3	2.31	0.55
7:M:62:ALA:CA	7:M:65:ILE:HG22	2.36	0.55
7:H:62:ALA:CA	7:H:65:ILE:HG22	2.35	0.55
8:E:37:ASP:CG	9:D:291:ARG:CZ	2.78	0.55
8:A:65:LEU:HD12	8:A:75:VAL:HG23	1.86	0.55
8:C:488:ALA:O	8:C:492:GLU:HG3	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:135:ASN:N	4:p:86:PHE:CZ	2.73	0.55
7:G:39:ILE:HG13	7:G:49:ILE:HG21	1.89	0.55
7:I:62:ALA:CA	7:I:65:ILE:HG22	2.36	0.55
9:B:168:ILE:HD12	9:B:324:ILE:HG12	1.87	0.55
8:C:19:GLY:O	8:C:22:ARG:HB3	2.06	0.55
8:C:78:GLY:CA	8:C:232:PRO:HG3	2.34	0.55
1:a:35:ILE:HD12	1:a:40:LEU:HD11	1.88	0.55
1:a:118:LYS:O	2:b:29:THR:CA	2.52	0.55
4:p:209:ALA:HB1	8:A:14:ARG:NH2	2.21	0.55
6:g:104:ASP:OD1	6:g:104:ASP:O	2.23	0.55
6:g:343:ARG:CZ	8:A:326:ASP:CG	2.80	0.55
7:Q:62:ALA:CA	7:Q:65:ILE:HG22	2.35	0.55
7:M:74:LEU:HD21	7:L:10:VAL:CG2	2.26	0.55
7:K:62:ALA:CA	7:K:65:ILE:HG22	2.36	0.55
8:E:147:LEU:HD21	8:E:258:LEU:HD13	1.88	0.55
9:F:23:ILE:HG21	9:F:26:ILE:HD11	1.89	0.55
4:p:83:LEU:HD12	4:p:84:PHE:CD2	2.41	0.55
7:O:39:ILE:HG13	7:O:49:ILE:HG21	1.89	0.55
7:I:39:ILE:HG13	7:I:49:ILE:HG21	1.89	0.55
8:E:52:GLU:HA	8:E:95:ILE:HG22	1.88	0.55
8:E:463:THR:HG23	8:E:464:TYR:N	2.20	0.55
8:C:241:TYR:OH	8:C:294:LEU:HD12	2.07	0.55
2:b:129:ILE:HG23	4:p:192:LEU:HD13	1.79	0.55
6:g:149:LYS:HZ3	6:g:155:TYR:HE1	1.51	0.55
6:g:202:PHE:CZ	6:g:232:PRO:HG3	2.41	0.55
7:N:62:ALA:CA	7:N:65:ILE:HG22	2.36	0.55
7:O:10:VAL:HG22	7:P:74:LEU:HD22	1.87	0.55
8:E:29:THR:HG22	8:E:90:LYS:HB3	1.89	0.55
8:A:164:GLN:CD	8:A:367:VAL:HG22	2.32	0.55
9:B:79:MET:CE	9:B:113:VAL:HG11	2.35	0.55
2:b:78:LEU:C	4:p:140:LEU:HG	2.17	0.55
2:b:118:LEU:HD12	4:p:178:ARG:HA	1.89	0.55
2:b:145:PHE:CB	8:A:13:ILE:HD12	2.36	0.55
3:d:74:THR:HG23	8:C:4:ILE:HG22	1.86	0.55
4:p:203:LEU:HG	4:p:206:GLN:HB3	1.89	0.55
6:g:114:THR:HG23	6:g:116:LYS:HG3	1.88	0.55
6:g:129:LEU:HD23	9:B:412:GLU:OE1	2.07	0.55
6:g:280:PRO:HG2	7:N:42:GLN:HE22	1.71	0.55
7:N:74:LEU:HD21	7:M:10:VAL:CG2	2.30	0.55
7:Q:39:ILE:HG13	7:Q:49:ILE:HG21	1.89	0.55
7:H:43:PRO:HB3	7:I:48:LYS:HZ3	1.69	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:101:PRO:HB3	9:F:126:THR:CG2	2.36	0.55
8:C:30:GLY:O	8:C:89:VAL:HG12	2.05	0.55
8:C:77:MET:HE3	8:C:112:LEU:CD2	2.17	0.55
8:C:138:MET:SD	9:D:213:TYR:CE2	3.00	0.55
1:a:166:GLN:N	1:a:167:PRO:HD2	2.21	0.55
2:b:74:ALA:HB1	4:p:135:SER:H	1.70	0.55
2:b:103:ILE:HG12	4:p:163:LYS:N	2.12	0.55
2:b:147:GLN:CD	4:p:207:ILE:CG2	2.80	0.55
7:M:48:LYS:NZ	7:L:43:PRO:HB2	2.21	0.55
9:B:259:TYR:C	9:B:259:TYR:CD1	2.85	0.55
8:C:199:ILE:HG12	8:C:240:PRO:HD3	1.89	0.55
8:C:393:LEU:C	8:C:393:LEU:HD12	2.31	0.55
8:C:420:LEU:CD2	8:C:458:LEU:HD11	2.37	0.55
8:C:490:LEU:O	8:C:494:ILE:HG13	2.06	0.55
9:D:416:GLU:HA	9:D:416:GLU:OE1	2.06	0.55
2:b:91:ASP:HB3	4:p:148:MET:CB	2.37	0.55
5:e:58:LEU:C	5:e:58:LEU:HD12	2.31	0.55
5:e:62:LEU:HB2	5:e:78:VAL:HG21	1.87	0.55
6:g:103:GLU:HG3	7:R:41:ARG:HG3	1.85	0.55
7:S:74:LEU:HD22	7:R:10:VAL:HG22	1.88	0.55
7:N:43:PRO:HB3	7:O:48:LYS:CD	2.37	0.55
7:J:39:ILE:HG13	7:J:49:ILE:HG21	1.89	0.55
8:E:9:ILE:HG23	8:E:13:ILE:HG22	1.88	0.55
9:F:101:PRO:HB2	9:F:126:THR:HG21	1.87	0.55
9:B:171:PHE:HB2	9:B:351:VAL:HA	1.88	0.55
1:a:25:HIS:HA	4:p:83:LEU:O	2.07	0.55
1:a:58:VAL:HG11	4:p:108:TYR:HA	1.89	0.55
1:a:119:ILE:HG12	2:b:28:ALA:HA	1.89	0.55
1:a:234:LEU:HG	1:a:238:TYR:CE2	2.41	0.55
2:b:81:ALA:CA	4:p:141:GLU:HG3	2.37	0.55
3:d:217:LEU:HA	3:d:237:VAL:HG23	1.89	0.55
4:p:203:LEU:CD2	8:A:7:ASP:C	2.76	0.55
7:N:43:PRO:CB	7:O:48:LYS:HZ2	2.18	0.55
8:A:37:ASP:HB3	8:A:277:LEU:HB3	1.87	0.55
8:A:457:TYR:OH	8:A:498:MET:HG2	2.07	0.55
1:a:117:TRP:O	1:a:120:ILE:CG2	2.54	0.55
2:b:61:LEU:HD13	2:b:65:ARG:HD3	1.89	0.55
2:b:80:LYS:N	4:p:137:VAL:HB	2.21	0.55
2:b:172:ASN:CG	3:d:235:MET:CB	2.80	0.55
3:d:148:ILE:HD13	8:C:17:ILE:CG1	2.37	0.55
4:p:193:GLU:HA	4:p:196:LYS:CE	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:72:VAL:HG12	6:g:257:LEU:HD21	1.88	0.55
6:g:261:GLU:CG	6:g:262:GLY:H	2.04	0.55
7:N:39:ILE:HG13	7:N:49:ILE:HG21	1.89	0.55
7:O:46:GLU:OE1	7:P:48:LYS:CD	2.54	0.55
7:O:62:ALA:CA	7:O:65:ILE:HG22	2.36	0.55
9:B:50:LYS:HE2	9:B:92:ILE:HD12	1.89	0.55
1:a:168:THR:HG22	1:a:170:ILE:H	1.72	0.54
2:b:96:ARG:HD2	4:p:154:GLU:OE1	2.07	0.54
4:p:127:LEU:HA	4:p:130:VAL:CG1	2.37	0.54
4:p:152:ARG:CD	8:A:503:LEU:CD2	2.71	0.54
6:g:105:VAL:HG22	6:g:289:ALA:HB2	1.89	0.54
7:N:48:LYS:HZ2	7:M:43:PRO:HB2	1.69	0.54
7:G:54:LEU:HD11	7:H:55:LEU:HD23	1.89	0.54
8:E:156:ALA:HA	8:E:376:MET:HE2	1.89	0.54
8:E:345:LEU:CD2	8:E:358:ILE:HD13	2.37	0.54
9:F:154:LYS:NZ	9:F:477:VAL:O	2.36	0.54
9:D:195:GLY:HA3	9:D:267:ASP:O	2.07	0.54
9:D:261:ARG:HD3	9:D:321:ILE:HG13	1.89	0.54
1:a:118:LYS:HZ1	1:a:126:GLU:CB	2.18	0.54
2:b:79:GLU:N	4:p:137:VAL:CA	2.47	0.54
4:p:144:ALA:C	4:p:147:VAL:HG12	2.32	0.54
6:g:261:GLU:CG	9:F:397:ARG:HH21	1.99	0.54
7:T:39:ILE:HG13	7:T:49:ILE:HG21	1.89	0.54
7:G:43:PRO:HB3	7:H:48:LYS:HD2	1.89	0.54
7:J:48:LYS:HB3	7:I:46:GLU:CD	2.32	0.54
8:A:287:TYR:HB3	8:A:291:VAL:HG21	1.89	0.54
9:B:212:LEU:HD12	9:B:212:LEU:C	2.32	0.54
9:B:387:ILE:HG23	9:B:388:ALA:N	2.23	0.54
8:C:437:THR:HG23	8:C:458:LEU:HD13	1.90	0.54
1:a:38:GLN:HG3	1:a:39:VAL:HG23	1.89	0.54
1:a:54:ALA:HB3	4:p:104:LEU:HD13	1.89	0.54
1:a:195:LEU:CD2	4:p:84:PHE:CZ	2.90	0.54
4:p:210:LEU:CG	8:A:13:ILE:HB	2.35	0.54
5:e:39:PRO:O	7:M:41:ARG:NH2	2.41	0.54
6:g:113:ARG:HH11	6:g:205:GLU:HB3	0.62	0.54
7:Q:43:PRO:CG	7:R:42:GLN:NE2	2.68	0.54
7:L:55:LEU:CD2	7:K:54:LEU:HD11	2.37	0.54
9:F:275:ILE:CG2	9:F:327:VAL:HG22	2.37	0.54
8:A:419:GLU:OE2	8:A:422:LYS:CD	2.55	0.54
8:C:449:LEU:HD11	8:C:454:VAL:CG2	2.37	0.54
3:d:165:PHE:C	3:d:165:PHE:CD1	2.85	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:195:GLN:CD	8:A:4:ILE:HD12	2.30	0.54
7:H:46:GLU:OE1	7:I:48:LYS:HD3	2.08	0.54
9:F:330:PRO:HD2	9:F:339:PRO:CG	2.37	0.54
8:A:61:ILE:O	8:A:77:MET:HB2	2.06	0.54
8:A:450:GLU:HB2	8:A:453:GLN:CD	2.33	0.54
9:B:472:GLN:O	9:B:486:LYS:HE2	2.07	0.54
9:D:203:GLY:O	9:D:277:ARG:HG3	2.06	0.54
1:a:78:PHE:CD1	1:a:78:PHE:C	2.85	0.54
2:b:68:GLU:HG3	4:p:126:GLN:CA	2.37	0.54
2:b:131:PHE:HE1	8:A:119:ARG:O	1.91	0.54
3:d:86:ALA:HB2	3:d:95:THR:HG21	1.89	0.54
5:e:119:LEU:O	5:e:123:ARG:HG3	2.07	0.54
7:L:39:ILE:HG13	7:L:49:ILE:HG21	1.89	0.54
7:L:48:LYS:HZ2	7:K:43:PRO:HB2	1.70	0.54
7:J:48:LYS:HG2	7:I:46:GLU:OE1	2.07	0.54
9:D:85:LEU:HD22	9:D:89:MET:HE1	1.90	0.54
9:D:401:LEU:HD23	9:D:404:ILE:HD11	1.84	0.54
3:d:148:ILE:CD1	8:C:16:ARG:HG2	2.37	0.54
6:g:109:LEU:HD11	6:g:199:PHE:CD1	2.42	0.54
6:g:149:LYS:CE	6:g:155:TYR:CE1	2.69	0.54
7:O:81:VAL:CG2	7:P:80:PHE:CZ	2.90	0.54
7:G:46:GLU:OE1	7:H:48:LYS:HG2	2.08	0.54
7:L:48:LYS:HZ3	7:K:43:PRO:C	2.16	0.54
7:J:31:ALA:HA	7:I:33:GLY:HA3	1.88	0.54
9:F:179:THR:CG2	9:F:215:GLU:OE1	2.50	0.54
8:C:195:VAL:HB	8:C:259:ILE:HG13	1.90	0.54
1:a:51:LEU:HG	4:p:100:LEU:HD11	1.80	0.54
1:a:134:ILE:O	1:a:135:ASN:C	2.51	0.54
2:b:79:GLU:HA	4:p:136:GLU:HG3	1.89	0.54
2:b:117:THR:CG2	4:p:174:LEU:CD1	2.72	0.54
6:g:359:GLY:N	9:D:293:PRO:HG3	2.20	0.54
7:N:46:GLU:OE1	7:O:48:LYS:CD	2.55	0.54
8:E:9:ILE:HB	8:E:14:ARG:CZ	2.38	0.54
1:a:24:GLN:CA	4:p:83:LEU:CG	2.84	0.54
2:b:69:GLU:HG2	2:b:73:LYS:HE3	1.89	0.54
2:b:88:VAL:C	4:p:147:VAL:C	2.75	0.54
2:b:122:GLU:CD	4:p:181:ILE:HD12	2.32	0.54
3:d:243:GLU:OE2	4:p:218:VAL:CG1	2.56	0.54
4:p:203:LEU:CD1	4:p:206:GLN:HB2	2.35	0.54
4:p:214:ILE:CG2	8:A:17:ILE:HD12	2.37	0.54
6:g:85:PHE:C	6:g:85:PHE:CD1	2.85	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:N:21:SER:OG	7:O:20:ALA:HA	2.08	0.54
7:G:43:PRO:HB2	7:H:48:LYS:HZ2	1.73	0.54
9:F:387:ILE:HG23	9:F:388:ALA:N	2.23	0.54
8:A:101:SER:HB3	8:A:124:ALA:HA	1.90	0.54
8:A:105:LEU:HD21	8:A:193:ILE:CG2	2.38	0.54
8:A:157:MET:SD	8:A:387:LEU:CD1	2.95	0.54
8:C:53:LEU:HD22	8:C:61:ILE:HG22	1.89	0.54
9:D:149:PHE:CE2	9:D:189:ILE:CD1	2.89	0.54
1:a:112:GLY:CA	1:a:128:ALA:O	2.54	0.54
2:b:100:TYR:O	2:b:103:ILE:CG2	2.53	0.54
2:b:103:ILE:CD1	4:p:158:ALA:C	2.80	0.54
2:b:145:PHE:CE2	2:b:149:LEU:CD1	2.77	0.54
3:d:243:GLU:CD	4:p:218:VAL:CG1	2.77	0.54
4:p:210:LEU:CD2	8:A:13:ILE:CG2	2.83	0.54
7:M:39:ILE:HG13	7:M:49:ILE:HG21	1.89	0.54
9:D:107:LEU:HD21	9:D:196:VAL:HG11	1.89	0.54
9:D:149:PHE:CE2	9:D:189:ILE:HD13	2.43	0.54
1:a:134:ILE:HG22	1:a:191:PHE:CE1	2.23	0.54
2:b:144:VAL:HB	8:A:13:ILE:HG13	1.89	0.54
3:d:195:ALA:O	3:d:199:GLN:CG	2.50	0.54
5:e:3:LEU:C	5:e:3:LEU:HD12	2.32	0.54
6:g:202:PHE:CD1	6:g:202:PHE:C	2.86	0.54
6:g:362:ALA:CB	8:C:281:PRO:CB	2.81	0.54
7:P:39:ILE:HG13	7:P:49:ILE:HG21	1.89	0.54
8:A:420:LEU:CD2	8:A:458:LEU:HD11	2.38	0.54
8:C:82:MET:HE2	8:C:82:MET:HA	1.90	0.54
9:D:347:ASP:O	9:D:373:THR:HG23	2.07	0.54
2:b:141:ARG:CG	8:A:9:ILE:CD1	2.85	0.53
2:b:146:GLN:O	2:b:149:LEU:HB2	2.08	0.53
2:b:160:LEU:HD21	2:b:164:LEU:HB2	1.91	0.53
3:d:186:LEU:CD1	3:d:191:LEU:HD21	2.32	0.53
4:p:212:ASP:OD1	4:p:215:VAL:CG1	2.55	0.53
6:g:83:ARG:HH21	6:g:273:THR:CG2	2.19	0.53
6:g:360:ALA:HA	9:D:292:MET:CA	2.37	0.53
7:N:10:VAL:CG2	7:O:74:LEU:HD21	2.25	0.53
7:N:48:LYS:HD3	7:M:46:GLU:OE1	2.08	0.53
7:L:57:LEU:O	7:L:61:GLU:N	2.35	0.53
8:E:479:THR:HG22	8:E:481:THR:CB	2.39	0.53
8:A:169:ILE:HD11	8:A:332:PRO:HB3	1.89	0.53
9:D:118:VAL:O	9:D:118:VAL:HG12	2.08	0.53
1:a:35:ILE:CB	4:p:88:LEU:N	2.70	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:P:46:GLU:CD	7:Q:48:LYS:HB3	2.34	0.53
7:L:42:GLN:HE22	7:K:43:PRO:CG	2.13	0.53
7:J:20:ALA:HA	7:I:21:SER:OG	2.08	0.53
9:F:48:ILE:CD1	9:F:94:THR:CG2	2.86	0.53
9:B:47:LEU:HB2	9:B:63:CYS:HB2	1.90	0.53
9:B:113:VAL:HG21	9:B:245:ALA:HB1	1.90	0.53
8:C:237:TYR:CE1	8:C:294:LEU:HD11	2.43	0.53
8:C:432:GLU:CG	8:C:476:ILE:HG23	2.32	0.53
1:a:195:LEU:HD11	4:p:84:PHE:HZ	1.69	0.53
2:b:156:LEU:HD21	4:p:219:LEU:HD23	1.89	0.53
3:d:148:ILE:CG2	8:C:16:ARG:HD2	2.38	0.53
3:d:186:LEU:CD1	3:d:191:LEU:CD2	2.77	0.53
6:g:363:CYS:HG	9:D:291:ARG:C	2.08	0.53
7:K:74:LEU:HD21	7:J:10:VAL:CG2	2.37	0.53
9:F:48:ILE:CD1	9:F:94:THR:HG21	2.38	0.53
8:C:104:TYR:CD1	8:C:122:ILE:HG21	2.43	0.53
8:C:457:TYR:HE1	8:C:497:GLN:HG3	1.73	0.53
9:D:55:ALA:HB3	9:D:57:GLN:HG2	1.89	0.53
1:a:76:LEU:HD13	4:p:109:TYR:CE1	2.43	0.53
2:b:180:ASN:CB	3:d:201:ILE:CG2	2.86	0.53
7:S:39:ILE:HG13	7:S:49:ILE:HG21	1.89	0.53
7:P:57:LEU:O	7:P:61:GLU:N	2.35	0.53
7:R:39:ILE:HG13	7:R:49:ILE:HG21	1.89	0.53
7:H:39:ILE:HG13	7:H:49:ILE:HG21	1.89	0.53
7:H:57:LEU:O	7:H:61:GLU:N	2.35	0.53
9:F:48:ILE:HD11	9:F:94:THR:HG21	1.90	0.53
9:F:50:LYS:HD2	9:F:90:GLU:CG	2.36	0.53
9:F:86:THR:H	9:F:89:MET:HE2	1.73	0.53
8:A:98:ILE:HG23	8:A:130:ILE:CG1	2.39	0.53
8:A:251:MET:HE2	8:A:314:MET:HB2	1.90	0.53
9:B:433:GLN:NE2	9:B:447:LYS:O	2.40	0.53
9:D:107:LEU:HD21	9:D:196:VAL:CG1	2.39	0.53
9:D:243:PRO:HB2	9:D:285:VAL:HG13	1.91	0.53
3:d:177:VAL:CG1	3:d:210:LYS:HE3	2.39	0.53
7:S:43:PRO:CG	7:T:42:GLN:HE22	2.17	0.53
7:M:70:VAL:HG13	7:L:75:LEU:HD11	1.91	0.53
7:K:39:ILE:HG13	7:K:49:ILE:HG21	1.89	0.53
8:E:169:ILE:HG22	8:E:344:PHE:CD1	2.17	0.53
8:C:176:LYS:HZ3	8:C:321:GLU:HB3	1.73	0.53
1:a:64:ILE:HG22	2:b:57:LYS:NZ	2.22	0.53
2:b:152:ALA:CB	4:p:214:ILE:CD1	2.87	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:160:LEU:HD13	2:b:163:GLU:N	2.20	0.53
5:e:18:GLU:O	5:e:53:LEU:HD22	2.09	0.53
6:g:353:ILE:HD11	8:E:284:ARG:CB	2.38	0.53
7:N:75:LEU:HD21	7:O:74:LEU:HD21	1.90	0.53
7:M:31:ALA:HA	7:L:33:GLY:HA3	1.90	0.53
8:A:495:GLN:O	8:A:499:GLU:HG2	2.09	0.53
9:D:146:LEU:HD23	9:D:163:ARG:HE	1.73	0.53
2:b:145:PHE:HB2	8:A:13:ILE:CD1	2.38	0.53
4:p:152:ARG:HH21	8:A:500:ARG:C	2.16	0.53
4:p:193:GLU:HG3	4:p:196:LYS:NZ	2.21	0.53
6:g:281:ILE:CD1	7:M:41:ARG:HG2	2.23	0.53
8:E:284:ARG:NH1	8:E:330:TYR:CB	2.72	0.53
8:A:39:ILE:HD13	8:A:278:LEU:CD2	2.38	0.53
8:A:416:ARG:HD2	8:A:447:ASP:HA	1.90	0.53
1:a:165:ILE:CG2	1:a:168:THR:C	2.71	0.53
2:b:118:LEU:HD11	4:p:181:ILE:HG12	1.90	0.53
2:b:173:ILE:HG12	3:d:221:PHE:CZ	2.25	0.53
2:b:181:GLU:O	2:b:182:ILE:C	2.51	0.53
3:d:172:ILE:HG23	3:d:173:THR:N	2.22	0.53
4:p:150:ALA:O	4:p:154:GLU:CG	2.55	0.53
6:g:109:LEU:HD21	6:g:199:PHE:HD1	1.71	0.53
6:g:264:LEU:HD21	9:F:408:LEU:HD11	1.91	0.53
7:N:48:LYS:CD	7:M:43:PRO:HB3	2.38	0.53
8:E:479:THR:CG2	8:E:481:THR:HB	2.39	0.53
8:C:58:GLU:HG3	8:C:82:MET:HB3	1.91	0.53
3:d:176:GLU:HG3	3:d:202:THR:HG21	1.90	0.53
4:p:179:LYS:HD3	4:p:179:LYS:C	2.33	0.53
6:g:356:ILE:CG1	9:D:295:ALA:HA	2.39	0.53
7:J:48:LYS:HD3	7:I:46:GLU:OE1	2.08	0.53
9:F:111:PHE:CZ	9:F:124:VAL:HG21	2.44	0.53
8:C:56:PHE:CE1	8:C:76:LEU:CD2	2.92	0.53
8:C:237:TYR:CZ	8:C:294:LEU:CD1	2.92	0.53
9:D:112:ASN:ND2	9:D:116:GLU:CB	2.56	0.53
9:D:254:LEU:CD1	9:D:313:ILE:HG12	2.39	0.53
2:b:107:LYS:HE3	4:p:165:GLU:C	2.34	0.53
2:b:129:ILE:HG21	4:p:192:LEU:HD11	1.82	0.53
2:b:145:PHE:HZ	3:d:243:GLU:O	1.88	0.53
5:e:37:HIS:CE1	7:L:41:ARG:NE	2.69	0.53
7:M:48:LYS:HZ2	7:L:43:PRO:CB	2.22	0.53
7:L:48:LYS:CD	7:K:43:PRO:HB3	2.39	0.53
7:K:57:LEU:HD23	7:K:57:LEU:N	2.24	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:J:48:LYS:CD	7:I:43:PRO:HB3	2.39	0.53
9:F:431:LEU:HD23	9:F:458:PHE:HZ	1.74	0.53
9:D:401:LEU:HD23	9:D:404:ILE:CD1	2.39	0.53
1:a:159:GLY:O	1:a:160:TYR:CG	2.62	0.52
3:d:152:SER:HA	3:d:153:GLU:OE1	2.09	0.52
3:d:222:THR:CG2	3:d:234:ASP:CG	2.75	0.52
5:e:24:LEU:HD12	5:e:25:SER:O	2.08	0.52
7:N:31:ALA:HA	7:M:33:GLY:HA3	1.90	0.52
7:N:57:LEU:N	7:N:57:LEU:HD23	2.24	0.52
7:P:33:GLY:HA3	7:Q:31:ALA:HA	1.89	0.52
7:G:46:GLU:OE1	7:H:48:LYS:CD	2.57	0.52
7:H:6:ALA:HB1	7:I:80:PHE:CD2	2.39	0.52
8:E:104:TYR:HE1	8:E:122:ILE:HD12	1.73	0.52
8:E:355:ARG:HH22	9:D:389:GLN:CD	2.10	0.52
8:E:469:LYS:HD2	8:E:493:ALA:CB	2.13	0.52
9:F:155:VAL:HG22	9:F:431:LEU:HB3	1.91	0.52
9:F:270:LEU:O	9:F:323:SER:HA	2.09	0.52
9:B:50:LYS:CE	9:B:92:ILE:HD12	2.39	0.52
9:B:411:ASP:N	9:B:418:ARG:NH2	2.57	0.52
8:C:292:PHE:CE1	9:D:246:ARG:NH1	2.76	0.52
8:C:432:GLU:HA	8:C:476:ILE:HD13	1.90	0.52
9:D:336:ASP:OD1	9:D:337:PRO:CD	2.55	0.52
2:b:64:ILE:O	4:p:123:ILE:O	2.27	0.52
7:O:6:ALA:HB1	7:P:80:PHE:CD2	2.38	0.52
9:F:274:ASN:OD1	9:F:277:ARG:HG3	2.10	0.52
8:C:237:TYR:CD1	8:C:271:TYR:CB	2.84	0.52
9:D:53:ASP:CB	9:D:59:MET:CE	2.86	0.52
9:D:102:VAL:CG2	9:D:128:THR:CG2	2.86	0.52
1:a:119:ILE:CA	2:b:28:ALA:C	2.75	0.52
1:a:230:ILE:HG13	7:Q:55:LEU:HD11	1.90	0.52
2:b:133:GLN:CA	8:A:3:THR:HG22	2.39	0.52
3:d:87:ASP:HB2	3:d:172:ILE:HD11	1.91	0.52
5:e:22:ILE:HA	5:e:51:ILE:HA	1.92	0.52
6:g:237:GLY:O	6:g:249:ALA:HB2	2.10	0.52
7:Q:10:VAL:HG22	7:R:74:LEU:CD2	2.38	0.52
9:F:79:MET:HE3	9:F:113:VAL:HG11	1.91	0.52
1:a:119:ILE:CA	2:b:28:ALA:CA	2.87	0.52
1:a:124:HIS:CB	4:p:80:LYS:O	2.56	0.52
1:a:230:ILE:CG1	7:Q:55:LEU:CD1	2.85	0.52
2:b:80:LYS:CG	4:p:137:VAL:CB	2.84	0.52
2:b:103:ILE:HG21	4:p:159:LEU:HA	1.79	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:187:GLU:N	3:d:187:GLU:OE2	2.42	0.52
5:e:19:VAL:C	5:e:53:LEU:HD13	2.34	0.52
7:L:57:LEU:N	7:L:57:LEU:HD23	2.24	0.52
9:F:453:GLU:OE1	9:F:456:ARG:NH2	2.25	0.52
8:A:67:LEU:HD12	9:B:26:ILE:HD13	1.92	0.52
9:B:411:ASP:C	9:B:418:ARG:NH2	2.67	0.52
8:C:237:TYR:CE1	8:C:271:TYR:HB2	2.42	0.52
9:D:75:ARG:HD3	9:D:289:LEU:HD23	1.91	0.52
9:D:141:GLN:O	9:D:317:LYS:HG3	2.09	0.52
3:d:92:LEU:HD21	3:d:172:ILE:HD13	1.90	0.52
3:d:224:ARG:CD	3:d:232:LEU:HD12	2.39	0.52
4:p:193:GLU:HA	4:p:196:LYS:HG2	1.90	0.52
5:e:8:LEU:HD13	6:g:85:PHE:HB2	1.92	0.52
5:e:24:LEU:HD13	5:e:25:SER:O	2.08	0.52
6:g:72:VAL:HG23	6:g:316:LEU:HB3	1.92	0.52
6:g:362:ALA:HB1	8:C:281:PRO:HA	1.91	0.52
7:S:57:LEU:HD23	7:S:57:LEU:N	2.24	0.52
7:P:57:LEU:HD23	7:P:57:LEU:N	2.24	0.52
7:Q:46:GLU:OE1	7:R:48:LYS:HD3	2.09	0.52
7:Q:54:LEU:HD11	7:R:55:LEU:HD23	1.90	0.52
7:Q:57:LEU:N	7:Q:57:LEU:HD23	2.24	0.52
9:F:50:LYS:HE2	9:F:92:ILE:CG1	2.34	0.52
9:F:384:HIS:C	9:F:384:HIS:CD2	2.87	0.52
8:A:193:ILE:O	8:A:193:ILE:CG2	2.49	0.52
3:d:175:THR:O	3:d:176:GLU:CD	2.52	0.52
3:d:220:GLY:H	3:d:236:SER:HB2	1.75	0.52
7:M:48:LYS:HZ2	7:L:43:PRO:HB2	1.75	0.52
7:T:57:LEU:N	7:T:57:LEU:HD23	2.24	0.52
7:J:57:LEU:N	7:J:57:LEU:HD23	2.24	0.52
7:I:57:LEU:N	7:I:57:LEU:HD23	2.24	0.52
9:B:292:MET:HE1	8:C:282:PRO:HD3	1.91	0.52
9:B:411:ASP:N	9:B:418:ARG:HH22	2.05	0.52
1:a:119:ILE:CD1	2:b:31:LEU:N	2.50	0.52
2:b:137:ILE:HA	8:A:5:ARG:HB3	1.91	0.52
4:p:101:MET:O	4:p:105:ASP:HB2	2.10	0.52
7:T:57:LEU:O	7:T:61:GLU:N	2.35	0.52
8:E:9:ILE:HG23	8:E:13:ILE:HG21	1.91	0.52
8:E:290:ASP:O	8:E:294:LEU:N	2.36	0.52
2:b:160:LEU:CD1	2:b:163:GLU:CB	2.69	0.52
6:g:83:ARG:N	6:g:84:PRO:HD2	2.25	0.52
7:O:3:PRO:HD2	7:O:5:ILE:HD12	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:M:3:PRO:HD2	7:M:5:ILE:HD12	1.92	0.52
9:F:64:GLU:OE2	9:F:133:HIS:NE2	2.43	0.52
9:D:423:ARG:NH2	9:D:464:GLY:HA3	2.24	0.52
1:a:24:GLN:O	4:p:83:LEU:HD12	2.09	0.52
2:b:70:LEU:HB3	4:p:131:LYS:CB	2.29	0.52
2:b:111:ILE:HG13	2:b:112:ASN:N	2.24	0.52
2:b:145:PHE:CZ	3:d:243:GLU:C	2.84	0.52
4:p:108:TYR:HE1	4:p:112:LEU:HD11	1.65	0.52
4:p:192:LEU:CD2	8:A:3:THR:CG2	2.84	0.52
4:p:195:GLN:HG2	8:A:4:ILE:HD13	1.58	0.52
4:p:196:LYS:C	4:p:199:THR:HG22	2.32	0.52
6:g:209:LYS:HA	6:g:231:LEU:O	2.10	0.52
7:P:10:VAL:CG2	7:Q:74:LEU:HD21	2.35	0.52
7:R:3:PRO:HD2	7:R:5:ILE:HD12	1.92	0.52
7:G:3:PRO:HD2	7:G:5:ILE:HD12	1.92	0.52
8:E:420:LEU:CD2	8:E:458:LEU:HD11	2.39	0.52
9:B:71:ASN:ND2	8:C:9:ILE:O	2.43	0.52
9:B:306:MET:HE3	9:B:310:GLN:CG	2.40	0.52
2:b:122:GLU:CG	4:p:181:ILE:HD12	2.39	0.52
4:p:110:THR:HB	4:p:111:PRO:HD3	1.91	0.52
5:e:100:GLU:HG2	5:e:119:LEU:HD21	1.91	0.52
6:g:110:THR:CG2	6:g:296:LEU:CD1	2.86	0.52
7:N:3:PRO:HD2	7:N:5:ILE:HD12	1.92	0.52
8:A:497:GLN:NE2	8:A:497:GLN:CA	2.73	0.52
9:B:157:ASP:HB3	9:B:451:LEU:HD13	1.92	0.52
8:C:13:ILE:O	8:C:17:ILE:HG13	2.10	0.52
2:b:96:ARG:O	4:p:155:ILE:HG23	2.09	0.51
6:g:123:VAL:HG13	6:g:308:LEU:HD21	1.82	0.51
7:M:57:LEU:O	7:M:61:GLU:N	2.35	0.51
7:H:40:ALA:CB	7:I:49:ILE:HD11	2.37	0.51
7:H:57:LEU:HD23	7:H:57:LEU:N	2.24	0.51
9:F:427:ILE:HD11	9:F:462:LEU:HD21	1.92	0.51
9:B:189:ILE:HG22	9:B:269:LEU:HD11	1.91	0.51
9:B:294:SER:HB3	9:B:300:PRO:HA	1.92	0.51
8:C:104:TYR:CD2	8:C:246:LEU:CD2	2.93	0.51
4:p:196:LYS:CD	8:A:5:ARG:HH22	2.22	0.51
7:H:3:PRO:HD2	7:H:5:ILE:HD12	1.92	0.51
9:F:164:ARG:NH1	9:F:320:SER:HB3	2.26	0.51
1:a:27:TYR:OH	1:a:122:LEU:CD1	2.58	0.51
2:b:70:LEU:HD12	4:p:127:LEU:CG	2.35	0.51
2:b:122:GLU:N	4:p:181:ILE:HG21	2.26	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:217:LEU:HA	3:d:237:VAL:CG2	2.40	0.51
7:S:20:ALA:HA	7:R:21:SER:OG	2.10	0.51
7:O:57:LEU:N	7:O:57:LEU:HD23	2.24	0.51
7:P:3:PRO:HD2	7:P:5:ILE:HD12	1.92	0.51
7:Q:40:ALA:CB	7:R:49:ILE:HD11	2.37	0.51
7:H:46:GLU:OE1	7:I:48:LYS:CD	2.58	0.51
9:F:25:GLN:HE21	9:F:27:ILE:CD1	2.23	0.51
9:F:353:SER:HB2	9:F:366:ASP:HB2	1.93	0.51
9:B:221:VAL:HG12	9:B:232:VAL:HB	1.93	0.51
9:B:237:GLY:CA	9:B:249:VAL:HG21	2.29	0.51
1:a:174:ILE:CG2	7:O:58:ALA:HB1	2.25	0.51
1:a:184:LEU:HD21	1:a:188:PHE:CD2	2.41	0.51
6:g:169:ILE:HG23	6:g:170:ARG:N	2.25	0.51
6:g:281:ILE:CA	7:N:41:ARG:HH22	2.04	0.51
9:F:120:ASN:O	9:F:120:ASN:CG	2.51	0.51
8:A:412:ALA:HB1	8:A:416:ARG:HH12	1.74	0.51
8:A:424:PRO:CB	8:A:455:ARG:NH1	2.73	0.51
9:B:169:GLY:CA	9:B:346:LEU:HD13	2.34	0.51
1:a:208:VAL:HG21	1:a:212:VAL:HG21	1.92	0.51
2:b:144:VAL:HB	8:A:13:ILE:CD1	2.41	0.51
2:b:175:MET:CB	8:A:24:VAL:HG21	2.40	0.51
4:p:167:GLN:O	4:p:171:GLU:HB3	2.10	0.51
6:g:327:THR:CG2	6:g:328:ASP:N	2.74	0.51
7:G:43:PRO:CB	7:H:48:LYS:HZ2	2.21	0.51
8:E:37:ASP:OD2	9:D:291:ARG:NH2	2.42	0.51
8:E:284:ARG:HH12	8:E:330:TYR:HA	1.76	0.51
8:A:327:VAL:HG21	8:A:344:PHE:CE1	2.41	0.51
9:D:202:VAL:HG22	9:D:249:VAL:HG23	1.93	0.51
1:a:70:ASN:CB	2:b:53:LEU:CB	2.88	0.51
2:b:57:LYS:O	2:b:60:ILE:CG2	2.56	0.51
4:p:127:LEU:CA	4:p:130:VAL:HG12	2.36	0.51
6:g:360:ALA:CB	9:D:292:MET:SD	2.74	0.51
7:N:48:LYS:HZ3	7:M:43:PRO:CA	2.24	0.51
7:T:3:PRO:HD2	7:T:5:ILE:HD12	1.92	0.51
7:G:57:LEU:N	7:G:57:LEU:HD23	2.24	0.51
7:G:81:VAL:CG2	7:H:80:PHE:CZ	2.94	0.51
8:E:67:LEU:O	9:F:25:GLN:HB2	2.10	0.51
8:C:383:LEU:HD12	8:C:383:LEU:C	2.35	0.51
1:a:65:PRO:HB2	1:a:70:ASN:HD21	0.34	0.51
1:a:119:ILE:HG22	2:b:27:LEU:C	2.29	0.51
2:b:122:GLU:CB	4:p:181:ILE:HD12	2.37	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:188:ASN:HD22	3:d:188:ASN:N	2.09	0.51
4:p:206:GLN:HE21	4:p:210:LEU:HB2	1.76	0.51
7:O:43:PRO:CA	7:P:48:LYS:HZ3	2.22	0.51
7:R:57:LEU:N	7:R:57:LEU:HD23	2.24	0.51
7:G:81:VAL:CG2	7:H:80:PHE:CE2	2.92	0.51
7:L:3:PRO:HD2	7:L:5:ILE:HD12	1.92	0.51
9:F:334:LEU:HD22	9:F:343:PHE:HE2	1.76	0.51
9:F:361:ILE:HG23	9:F:432:SER:CB	2.31	0.51
8:A:359:ASN:HD22	8:A:362:ILE:HD12	1.64	0.51
2:b:114:THR:O	4:p:174:LEU:HG	2.10	0.51
2:b:125:LYS:HG3	4:p:181:ILE:HG22	1.91	0.51
2:b:142:GLN:O	2:b:143:ARG:C	2.52	0.51
3:d:224:ARG:CG	3:d:232:LEU:HD12	2.41	0.51
3:d:228:GLU:OE2	8:A:43:HIS:HE1	1.93	0.51
5:e:23:ILE:HG23	5:e:50:ARG:HB2	1.93	0.51
8:E:159:PRO:HB3	8:E:372:GLN:HG3	1.93	0.51
8:A:436:MET:O	8:A:440:THR:HG23	2.11	0.51
9:D:168:ILE:HD12	9:D:324:ILE:HG12	1.93	0.51
1:a:214:ILE:HG12	1:a:215:PRO:HD3	1.89	0.51
2:b:88:VAL:CA	4:p:148:MET:HG2	1.89	0.51
6:g:71:LYS:CB	6:g:316:LEU:HD13	2.41	0.51
7:N:68:LEU:HD22	7:O:66:TYR:CG	2.46	0.51
7:M:48:LYS:NZ	7:L:43:PRO:HB3	2.24	0.51
8:E:104:TYR:CD1	8:E:122:ILE:HG21	2.46	0.51
8:A:24:VAL:HG23	8:A:24:VAL:O	2.11	0.51
8:A:165:ARG:HB3	8:A:339:THR:HG23	1.92	0.51
8:C:394:GLU:OE1	8:C:418:ARG:NH2	2.39	0.51
2:b:79:GLU:N	4:p:136:GLU:CB	2.73	0.51
3:d:177:VAL:HA	3:d:208:ARG:O	2.11	0.51
3:d:181:THR:O	3:d:221:PHE:CA	2.58	0.51
6:g:155:TYR:HE2	6:g:174:ILE:HG23	1.76	0.51
7:N:43:PRO:HB3	7:O:48:LYS:HZ3	1.69	0.51
7:O:35:ALA:CB	7:O:53:LEU:HD13	2.41	0.51
7:O:81:VAL:CG2	7:P:80:PHE:CE2	2.92	0.51
7:M:57:LEU:N	7:M:57:LEU:HD23	2.24	0.51
7:M:74:LEU:HD22	7:L:10:VAL:HG22	1.92	0.51
8:E:430:THR:HG22	8:E:431:VAL:N	2.26	0.51
9:F:163:ARG:HG2	9:F:374:MET:HG3	1.92	0.51
8:A:413:ARG:NH2	8:A:444:GLY:HA3	2.26	0.51
8:C:237:TYR:CZ	8:C:294:LEU:HD13	2.44	0.51
1:a:121:GLN:HB2	2:b:27:LEU:HB2	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:88:VAL:CA	4:p:148:MET:CA	2.87	0.50
3:d:75:ALA:HB1	3:d:160:GLU:HG2	1.93	0.50
4:p:214:ILE:C	8:A:17:ILE:HD11	2.36	0.50
5:e:8:LEU:HD12	5:e:8:LEU:O	2.10	0.50
5:e:100:GLU:HG3	5:e:119:LEU:HD21	1.94	0.50
7:S:3:PRO:HD2	7:S:5:ILE:HD12	1.92	0.50
7:Q:3:PRO:HD2	7:Q:5:ILE:HD12	1.92	0.50
8:E:176:LYS:HD2	8:E:319:ILE:CG2	2.41	0.50
9:F:336:ASP:HB3	9:F:339:PRO:CD	2.41	0.50
8:A:177:THR:OG1	10:A:600:ATP:O2B	2.28	0.50
9:B:69:LEU:HD11	9:B:75:ARG:HB2	1.93	0.50
9:B:113:VAL:HG22	9:B:249:VAL:CG2	2.40	0.50
1:a:51:LEU:CG	4:p:100:LEU:HD11	2.16	0.50
1:a:63:THR:HA	4:p:119:ARG:HH11	0.44	0.50
1:a:121:GLN:HG3	2:b:25:ASP:OD1	2.11	0.50
6:g:202:PHE:CE2	6:g:231:LEU:HB3	2.46	0.50
7:Q:35:ALA:CB	7:Q:53:LEU:HD13	2.41	0.50
7:Q:57:LEU:O	7:Q:61:GLU:N	2.35	0.50
7:M:35:ALA:CB	7:M:53:LEU:HD13	2.41	0.50
8:E:172:ARG:NH2	9:D:369:ASP:OD1	2.44	0.50
8:A:20:TYR:CE2	8:A:22:ARG:HB2	2.47	0.50
8:A:216:ARG:HG3	8:A:216:ARG:NH2	2.26	0.50
9:D:64:GLU:OE1	9:D:248:ARG:CZ	2.59	0.50
9:D:109:ARG:NH1	9:D:119:ASP:CG	2.69	0.50
9:D:182:ILE:HG23	9:D:183:MET:N	2.26	0.50
1:a:70:ASN:CA	2:b:53:LEU:HB3	2.41	0.50
1:a:135:ASN:HD21	4:p:86:PHE:N	2.00	0.50
2:b:45:GLY:O	2:b:49:LEU:CB	2.59	0.50
2:b:103:ILE:HD11	4:p:161:LYS:C	2.23	0.50
2:b:144:VAL:HG11	8:A:9:ILE:HG22	1.92	0.50
3:d:157:LEU:HD12	3:d:157:LEU:O	2.12	0.50
3:d:179:VAL:HA	3:d:210:LYS:O	2.11	0.50
4:p:219:LEU:HB2	4:p:220:PRO:CD	2.35	0.50
6:g:122:VAL:HG13	6:g:213:LEU:HD11	1.93	0.50
6:g:360:ALA:HA	9:D:292:MET:HB3	1.93	0.50
7:P:81:VAL:CG2	7:Q:80:PHE:CE2	2.94	0.50
7:M:35:ALA:HA	7:L:36:VAL:HG11	1.93	0.50
7:H:35:ALA:CB	7:H:53:LEU:HD13	2.41	0.50
7:K:35:ALA:CB	7:K:53:LEU:HD13	2.41	0.50
8:A:432:GLU:HG3	8:A:476:ILE:HB	1.92	0.50
8:C:271:TYR:CD2	8:C:294:LEU:CD2	2.90	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:119:ILE:CD1	2:b:31:LEU:CA	2.90	0.50
1:a:124:HIS:CE1	4:p:79:GLU:N	2.80	0.50
2:b:85:LEU:CD1	2:b:86:LYS:N	2.68	0.50
2:b:122:GLU:OE2	4:p:184:GLU:OE1	2.29	0.50
8:E:383:LEU:HD13	8:E:417:LEU:HD22	1.93	0.50
8:A:65:LEU:CD1	8:A:75:VAL:HG21	2.41	0.50
8:C:100:VAL:CG2	8:C:246:LEU:HD23	2.35	0.50
9:D:102:VAL:HG22	9:D:128:THR:HG22	1.93	0.50
9:D:109:ARG:NH2	9:D:122:GLY:O	2.43	0.50
9:D:146:LEU:CD2	9:D:163:ARG:HE	2.25	0.50
9:D:401:LEU:HA	9:D:404:ILE:HG12	1.93	0.50
1:a:40:LEU:HD21	4:p:92:ILE:HB	1.94	0.50
2:b:88:VAL:N	4:p:148:MET:HG3	2.06	0.50
3:d:247:GLN:C	8:A:16:ARG:HD3	2.30	0.50
4:p:214:ILE:HG22	8:A:17:ILE:HD12	1.93	0.50
6:g:174:ILE:HG22	6:g:175:PRO:O	2.11	0.50
6:g:257:LEU:O	6:g:257:LEU:HD12	2.11	0.50
7:R:35:ALA:CB	7:R:53:LEU:HD13	2.41	0.50
7:G:21:SER:OG	7:H:20:ALA:HA	2.10	0.50
8:A:383:LEU:HD22	8:A:438:ILE:CD1	2.22	0.50
8:A:419:GLU:OE2	8:A:422:LYS:NZ	2.43	0.50
8:C:167:LEU:HA	8:C:318:PRO:HD2	1.93	0.50
8:C:259:ILE:HG23	8:C:259:ILE:O	2.12	0.50
9:D:168:ILE:HB	9:D:324:ILE:HA	1.93	0.50
3:d:151:ASP:CB	8:C:20:TYR:OH	2.46	0.50
4:p:152:ARG:CG	8:A:503:LEU:HB3	2.23	0.50
6:g:253:GLU:HB2	6:g:269:ASP:O	2.12	0.50
7:T:35:ALA:CB	7:T:53:LEU:HD13	2.41	0.50
8:E:202:LYS:CD	9:D:345:HIS:O	2.55	0.50
8:A:131:GLU:OE1	8:A:297:ARG:NH2	2.45	0.50
8:A:382:LYS:HB3	8:A:442:THR:CG2	2.38	0.50
9:D:61:VAL:O	9:D:61:VAL:HG23	2.11	0.50
9:D:110:ILE:HD11	9:D:213:TYR:CD1	2.47	0.50
1:a:53:SER:O	1:a:56:ILE:HG23	2.06	0.50
1:a:100:THR:CG2	4:p:101:MET:HB2	2.25	0.50
1:a:124:HIS:CE1	4:p:79:GLU:H	2.29	0.50
1:a:124:HIS:CG	4:p:80:LYS:O	2.65	0.50
2:b:70:LEU:HB2	4:p:127:LEU:HD23	1.94	0.50
3:d:74:THR:HG23	8:C:4:ILE:N	2.26	0.50
3:d:119:VAL:CG1	8:E:31:THR:OG1	2.59	0.50
3:d:198:VAL:O	3:d:202:THR:CB	2.59	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:p:203:LEU:HD21	8:A:10:SER:H	1.75	0.50
5:e:119:LEU:HD11	5:e:123:ARG:HH21	1.76	0.50
7:K:3:PRO:HD2	7:K:5:ILE:HD12	1.92	0.50
8:E:404:ASP:N	8:E:404:ASP:OD1	2.42	0.50
9:F:20:LEU:HD21	9:F:92:ILE:HG12	1.93	0.50
9:B:71:ASN:ND2	8:C:9:ILE:CA	2.69	0.50
8:C:164:GLN:NE2	8:C:166:GLU:OE1	2.45	0.50
8:C:247:ALA:HB3	8:C:314:MET:CE	2.41	0.50
8:C:410:GLN:HE22	9:D:472:GLN:HG3	1.77	0.50
2:b:87:LYS:O	2:b:90:MET:CB	2.53	0.50
2:b:91:ASP:CB	4:p:148:MET:HG2	2.42	0.50
6:g:261:GLU:CG	6:g:262:GLY:N	2.69	0.50
7:O:43:PRO:HB3	7:P:48:LYS:CD	2.42	0.50
7:J:3:PRO:HD2	7:J:5:ILE:HD12	1.92	0.50
7:I:35:ALA:CB	7:I:53:LEU:HD13	2.41	0.50
8:E:165:ARG:HD3	8:E:299:LEU:O	2.12	0.50
8:A:185:LEU:CD1	8:A:426:SER:OG	2.60	0.50
9:B:251:LEU:CD2	9:B:309:LEU:CD1	2.81	0.50
1:a:79:ILE:HD11	1:a:99:GLY:HA2	1.88	0.50
1:a:208:VAL:HG23	1:a:209:PRO:HD3	1.85	0.50
4:p:87:ASN:OD1	4:p:89:THR:N	2.29	0.50
5:e:19:VAL:HG21	5:e:51:ILE:CG1	2.39	0.50
7:N:48:LYS:HZ2	7:M:43:PRO:CB	2.20	0.50
8:E:285:GLU:O	9:D:295:ALA:HB2	2.12	0.50
8:C:251:MET:CE	8:C:252:TYR:CE1	2.94	0.50
8:C:254:GLU:HG2	8:C:310:GLY:HA3	1.93	0.50
3:d:217:LEU:HD23	3:d:236:SER:HG	1.76	0.49
3:d:235:MET:CG	8:A:24:VAL:HG11	2.42	0.49
4:p:179:LYS:C	4:p:179:LYS:CD	2.85	0.49
7:G:35:ALA:CB	7:G:53:LEU:HD13	2.42	0.49
7:L:35:ALA:CB	7:L:53:LEU:HD13	2.41	0.49
8:E:169:ILE:HG13	8:E:322:THR:HG23	1.93	0.49
8:A:100:VAL:HG23	8:A:246:LEU:HD23	1.94	0.49
8:C:22:ARG:HG3	8:C:23:GLU:N	2.26	0.49
8:C:239:ALA:N	8:C:240:PRO:HD2	2.26	0.49
9:D:113:VAL:HG22	9:D:249:VAL:CG1	2.37	0.49
9:D:173:GLY:O	9:D:178:LYS:NZ	2.45	0.49
1:a:159:GLY:O	1:a:160:TYR:CD2	2.65	0.49
3:d:195:ALA:C	3:d:207:VAL:HG11	2.38	0.49
5:e:15:TRP:NE1	5:e:17:SER:O	2.45	0.49
6:g:72:VAL:CG1	6:g:257:LEU:CD2	2.91	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:285:GLU:OE1	6:g:285:GLU:HA	2.12	0.49
7:S:57:LEU:O	7:S:61:GLU:N	2.35	0.49
7:N:10:VAL:HG22	7:O:74:LEU:CD2	2.40	0.49
7:T:10:VAL:CG2	7:G:74:LEU:HD21	2.35	0.49
8:E:489:LEU:CD2	8:E:489:LEU:C	2.85	0.49
2:b:88:VAL:O	2:b:90:MET:N	2.44	0.49
2:b:91:ASP:C	4:p:151:ALA:HB3	2.19	0.49
2:b:140:VAL:HG11	8:A:6:ALA:HA	1.93	0.49
3:d:129:LEU:HD23	3:d:147:ASN:OD1	2.11	0.49
6:g:107:VAL:CG1	6:g:296:LEU:HD12	2.42	0.49
6:g:110:THR:CG2	6:g:296:LEU:HD13	2.42	0.49
7:S:81:VAL:CG2	7:T:80:PHE:CE2	2.95	0.49
7:N:35:ALA:CB	7:N:53:LEU:HD13	2.41	0.49
8:E:208:GLN:HE22	9:D:144:THR:HG22	1.72	0.49
9:F:50:LYS:HD2	9:F:90:GLU:HG3	1.93	0.49
9:F:110:ILE:HB	9:F:119:ASP:HB3	1.94	0.49
9:F:433:GLN:NE2	9:F:447:LYS:O	2.46	0.49
9:D:189:ILE:CG2	9:D:269:LEU:HD11	2.33	0.49
1:a:119:ILE:HD12	2:b:31:LEU:CA	2.41	0.49
7:N:57:LEU:O	7:N:61:GLU:N	2.35	0.49
7:M:48:LYS:HD2	7:L:43:PRO:HB3	1.93	0.49
9:F:385:TYR:O	9:F:389:GLN:HG2	2.13	0.49
9:B:170:LEU:HD22	9:B:181:LEU:HD23	1.94	0.49
8:C:104:TYR:HD2	8:C:246:LEU:HD22	1.77	0.49
9:D:398:TYR:OH	9:D:425:ARG:HD3	2.12	0.49
2:b:103:ILE:CB	4:p:159:LEU:HB2	2.33	0.49
3:d:222:THR:HG22	3:d:234:ASP:HB2	0.49	0.49
5:e:58:LEU:HD12	5:e:58:LEU:O	2.11	0.49
8:A:58:GLU:OE1	8:A:58:GLU:HA	2.11	0.49
9:B:71:ASN:N	8:C:9:ILE:HG13	2.27	0.49
1:a:27:TYR:CZ	1:a:122:LEU:CD1	2.69	0.49
4:p:199:THR:CG2	4:p:200:ILE:N	2.76	0.49
5:e:5:LEU:CD2	5:e:51:ILE:HD12	2.42	0.49
6:g:227:ILE:CG2	6:g:242:ILE:CD1	2.89	0.49
7:S:35:ALA:CB	7:S:53:LEU:HD13	2.41	0.49
7:N:80:PHE:CZ	7:M:81:VAL:CG2	2.95	0.49
8:E:463:THR:CG2	8:E:464:TYR:N	2.76	0.49
9:F:124:VAL:HG12	9:F:126:THR:CG2	2.41	0.49
9:F:163:ARG:HG2	9:F:374:MET:CG	2.43	0.49
9:F:168:ILE:HB	9:F:324:ILE:HA	1.93	0.49
9:F:211:ASP:O	9:F:215:GLU:HG3	2.11	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:366:ARG:HD2	9:B:441:PHE:HE1	1.77	0.49
3:d:119:VAL:CG1	8:E:43:HIS:CD2	2.96	0.49
3:d:186:LEU:HD11	3:d:191:LEU:CD1	2.40	0.49
4:p:199:THR:HG21	8:A:6:ALA:H	1.78	0.49
5:e:45:ASP:OD2	5:e:120:ARG:NH2	2.46	0.49
7:S:48:LYS:HZ2	7:R:43:PRO:HB3	1.65	0.49
8:E:173:GLN:CG	9:D:371:THR:HG21	2.42	0.49
8:E:376:MET:HE1	8:E:438:ILE:HD12	1.93	0.49
8:A:469:LYS:HZ2	8:A:492:GLU:C	2.21	0.49
8:A:469:LYS:HZ2	8:A:493:ALA:N	2.11	0.49
8:C:98:ILE:HG22	8:C:128:ARG:O	2.12	0.49
8:C:104:TYR:HE1	8:C:122:ILE:CD1	2.25	0.49
1:a:139:ALA:HB2	4:p:94:MET:HE2	1.94	0.49
2:b:61:LEU:HD13	2:b:61:LEU:O	2.13	0.49
2:b:91:ASP:O	4:p:151:ALA:CB	2.60	0.49
3:d:228:GLU:HG3	8:A:44:GLY:HA2	1.94	0.49
6:g:103:GLU:CB	7:R:41:ARG:HG3	2.43	0.49
6:g:280:PRO:O	7:N:41:ARG:NH2	2.45	0.49
7:S:75:LEU:HD21	7:T:74:LEU:HD21	1.95	0.49
7:P:35:ALA:CB	7:P:53:LEU:HD13	2.42	0.49
7:J:35:ALA:CB	7:J:53:LEU:HD13	2.41	0.49
7:J:66:TYR:CG	7:I:68:LEU:HD22	2.48	0.49
8:E:176:LYS:HD2	8:E:319:ILE:HG23	1.95	0.49
9:F:199:PHE:HB3	9:F:234:LEU:HD23	1.94	0.49
8:C:176:LYS:NZ	8:C:321:GLU:HB3	2.27	0.49
9:D:404:ILE:HG13	9:D:405:ILE:N	2.28	0.49
2:b:77:GLN:CG	4:p:134:SER:O	2.60	0.49
2:b:103:ILE:HG21	4:p:162:MET:HB2	1.89	0.49
6:g:239:ILE:HG23	6:g:239:ILE:O	2.12	0.49
6:g:356:ILE:HG23	9:D:293:PRO:O	2.13	0.49
7:I:57:LEU:O	7:I:61:GLU:N	2.35	0.49
8:E:68:GLU:O	9:F:87:ARG:NH1	2.46	0.49
8:A:79:ASP:HB3	8:A:81:LEU:HG	1.95	0.49
8:A:214:GLN:HE22	8:A:219:MET:HG2	1.29	0.49
8:A:420:LEU:HD23	8:A:458:LEU:CD1	2.43	0.49
9:B:425:ARG:HH12	9:B:429:ARG:NH2	2.06	0.49
9:D:387:ILE:HD11	9:D:455:ILE:HG23	1.84	0.49
2:b:61:LEU:O	2:b:61:LEU:HD22	2.13	0.49
2:b:114:THR:HA	2:b:117:THR:HG22	1.94	0.49
3:d:148:ILE:HG23	8:C:16:ARG:HD3	1.95	0.49
3:d:198:VAL:CA	3:d:201:ILE:HG12	2.43	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:e:38:ALA:HB1	7:M:41:ARG:CZ	2.43	0.49
6:g:72:VAL:HG22	6:g:316:LEU:HB2	1.95	0.49
7:Q:43:PRO:CA	7:R:48:LYS:HZ3	2.26	0.49
8:A:158:ILE:HG12	8:A:363:SER:HB2	1.94	0.49
8:A:424:PRO:HB3	8:A:455:ARG:HH11	1.78	0.49
8:C:45:LEU:O	9:D:87:ARG:NH2	2.41	0.49
8:C:107:ARG:HD2	8:C:119:ARG:NH1	2.28	0.49
8:C:138:MET:HE1	9:D:213:TYR:CE2	2.48	0.49
2:b:95:PHE:CZ	4:p:156:SER:HB2	2.46	0.48
3:d:181:THR:HG21	3:d:214:ASP:OD2	2.13	0.48
5:e:32:GLY:O	7:K:41:ARG:O	2.30	0.48
6:g:202:PHE:CB	6:g:231:LEU:CD1	2.91	0.48
7:O:57:LEU:O	7:O:61:GLU:N	2.35	0.48
8:E:461:LEU:C	8:E:461:LEU:CD2	2.85	0.48
9:F:50:LYS:CD	9:F:90:GLU:OE2	2.60	0.48
9:D:102:VAL:CG2	9:D:128:THR:HG22	2.43	0.48
2:b:78:LEU:HA	4:p:139:GLN:H	1.77	0.48
2:b:118:LEU:HD13	4:p:178:ARG:CA	2.40	0.48
6:g:79:VAL:CG1	6:g:306:ARG:NH2	2.76	0.48
6:g:211:GLU:HB2	6:g:228:HIS:O	2.13	0.48
7:S:74:LEU:HD21	7:R:10:VAL:CG2	2.37	0.48
8:E:101:SER:OG	8:E:102:GLU:N	2.46	0.48
9:F:254:LEU:HD21	9:F:312:ARG:HB2	1.94	0.48
8:A:54:VAL:O	8:A:54:VAL:HG23	2.12	0.48
8:A:109:ILE:CG2	8:A:225:VAL:HA	2.41	0.48
8:A:430:THR:HG22	8:A:431:VAL:N	2.28	0.48
8:C:420:LEU:HD11	8:C:441:GLY:HA3	1.95	0.48
9:D:441:PHE:CD1	12:D:600:ADP:H2'	2.47	0.48
1:a:118:LYS:NZ	1:a:126:GLU:HB3	2.29	0.48
8:E:490:LEU:C	8:E:490:LEU:CD2	2.85	0.48
8:A:98:ILE:HG23	8:A:130:ILE:HG12	1.96	0.48
8:A:100:VAL:HG23	8:A:246:LEU:CD2	2.43	0.48
8:A:445:TYR:HD2	8:A:494:ILE:HG21	1.72	0.48
9:B:79:MET:HE3	9:B:113:VAL:HG11	1.96	0.48
8:C:232:PRO:HG2	8:C:235:LEU:HG	1.94	0.48
2:b:88:VAL:C	4:p:148:MET:CA	2.86	0.48
2:b:117:THR:HG23	4:p:174:LEU:HD21	1.94	0.48
2:b:145:PHE:H	8:A:13:ILE:HD11	1.78	0.48
3:d:186:LEU:HD12	3:d:186:LEU:O	2.13	0.48
4:p:193:GLU:HA	4:p:196:LYS:CD	2.44	0.48
6:g:123:VAL:HG22	6:g:308:LEU:HD21	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S:80:PHE:CE2	7:R:81:VAL:CG2	2.95	0.48
7:N:80:PHE:CE2	7:M:81:VAL:CG2	2.94	0.48
8:E:156:ALA:O	8:E:372:GLN:NE2	2.46	0.48
8:E:214:GLN:HG3	8:E:219:MET:CG	2.42	0.48
9:B:61:VAL:HG11	9:B:85:LEU:HD11	1.96	0.48
1:a:84:LYS:HA	1:a:91:TYR:HB3	1.96	0.48
7:N:74:LEU:HD22	7:M:10:VAL:HG22	1.93	0.48
7:M:20:ALA:HA	7:L:21:SER:OG	2.13	0.48
8:E:416:ARG:HG3	8:E:454:VAL:HG21	1.95	0.48
8:A:148:GLN:O	8:A:186:ASN:ND2	2.46	0.48
8:A:329:ALA:HB3	8:A:332:PRO:CD	2.44	0.48
8:A:497:GLN:HA	8:A:497:GLN:HE21	1.79	0.48
2:b:84:ARG:HB3	4:p:141:GLU:C	2.37	0.48
6:g:231:LEU:HD23	6:g:233:LEU:HD12	1.96	0.48
7:Q:10:VAL:HG22	7:R:74:LEU:HD22	1.91	0.48
9:F:163:ARG:HB2	9:F:166:GLY:HA3	1.95	0.48
8:A:344:PHE:CD2	8:A:362:ILE:O	2.67	0.48
1:a:104:PHE:O	1:a:108:SER:OG	2.27	0.48
1:a:165:ILE:CG2	1:a:169:PRO:CA	2.92	0.48
2:b:88:VAL:HG22	4:p:148:MET:CB	2.43	0.48
2:b:91:ASP:HB3	4:p:148:MET:CG	2.43	0.48
5:e:96:LEU:HD13	5:e:123:ARG:N	2.29	0.48
6:g:240:CYS:HB3	6:g:246:CYS:HA	1.96	0.48
8:A:432:GLU:HA	8:A:435:VAL:HG12	1.95	0.48
9:B:464:GLY:O	9:B:465:GLU:C	2.56	0.48
9:D:113:VAL:HG21	9:D:245:ALA:HB1	1.95	0.48
1:a:44:TRP:CD1	4:p:96:GLU:CD	2.91	0.48
2:b:147:GLN:HA	2:b:150:GLN:CD	2.38	0.48
3:d:74:THR:HB	3:d:154:ARG:NH1	2.28	0.48
4:p:159:LEU:CD1	4:p:160:ASN:N	2.68	0.48
7:O:43:PRO:HB2	7:P:48:LYS:HZ2	1.79	0.48
7:O:68:LEU:HD22	7:P:66:TYR:CG	2.49	0.48
7:J:12:ALA:HB2	7:I:11:ILE:HA	1.95	0.48
8:E:48:VAL:HG22	8:E:49:MET:N	2.27	0.48
8:A:464:TYR:HD2	8:A:497:GLN:HB2	1.78	0.48
9:B:146:LEU:HD22	9:B:374:MET:HE1	1.95	0.48
9:B:251:LEU:HD23	9:B:309:LEU:CD1	2.33	0.48
8:C:4:ILE:O	8:C:4:ILE:HG12	2.13	0.48
8:C:151:LEU:HB2	8:C:154:ILE:HD13	1.96	0.48
1:a:100:THR:CG2	4:p:101:MET:CB	2.71	0.48
1:a:122:LEU:HB3	1:a:125:GLY:O	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:98:ASN:O	2:b:102:GLU:HG2	2.14	0.48
2:b:103:ILE:CA	4:p:159:LEU:HB2	2.44	0.48
7:P:43:PRO:CG	7:Q:42:GLN:NE2	2.73	0.48
9:F:313:ILE:HG21	9:F:323:SER:HB3	1.95	0.48
8:A:52:GLU:HA	8:A:95:ILE:HA	1.96	0.48
8:A:403:LEU:HB3	8:A:408:GLN:HB2	1.95	0.48
9:B:50:LYS:HZ3	9:B:92:ILE:HD12	1.78	0.48
8:C:404:ASP:N	8:C:404:ASP:OD1	2.44	0.48
1:a:24:GLN:NE2	1:a:195:LEU:CD1	2.65	0.48
2:b:77:GLN:CG	4:p:137:VAL:HG23	2.32	0.48
6:g:138:LEU:HD13	6:g:167:TYR:CD1	2.49	0.48
6:g:356:ILE:HD11	9:D:295:ALA:CB	2.44	0.48
7:N:80:PHE:CD2	7:M:6:ALA:HB1	2.43	0.48
9:F:426:LYS:HG2	9:F:474:PHE:CD2	2.48	0.48
8:A:107:ARG:NH2	8:A:117:ASP:OD2	2.45	0.48
8:A:457:TYR:OH	8:A:498:MET:CG	2.61	0.48
1:a:116:PRO:HB3	2:b:33:ASN:N	2.27	0.47
1:a:130:PRO:O	1:a:133:ASP:OD1	2.32	0.47
1:a:137:THR:HG21	1:a:191:PHE:HA	1.94	0.47
2:b:85:LEU:HD12	2:b:85:LEU:C	2.39	0.47
2:b:114:THR:CA	2:b:117:THR:HG22	2.44	0.47
4:p:215:VAL:HG13	4:p:216:LYS:N	2.28	0.47
6:g:240:CYS:HB3	6:g:246:CYS:CA	2.43	0.47
6:g:335:LYS:O	6:g:339:ILE:HG12	2.14	0.47
7:L:49:ILE:HD11	7:K:40:ALA:CB	2.39	0.47
9:F:124:VAL:CG1	9:F:126:THR:HG22	2.42	0.47
8:A:413:ARG:CZ	8:A:444:GLY:CA	2.92	0.47
9:D:51:GLY:O	9:D:59:MET:O	2.31	0.47
3:d:235:MET:HE1	3:d:239:LYS:HG2	1.93	0.47
7:G:10:VAL:HG22	7:H:74:LEU:HD22	1.91	0.47
8:E:503:LEU:O	8:E:504:GLN:HG2	2.14	0.47
9:B:19:ASN:N	9:B:19:ASN:OD1	2.45	0.47
8:C:101:SER:HB3	8:C:124:ALA:HA	1.96	0.47
1:a:50:LEU:HD21	1:a:107:VAL:HG21	1.96	0.47
2:b:71:ARG:HB2	4:p:127:LEU:C	2.39	0.47
3:d:98:ASP:OD2	3:d:139:GLN:HG2	2.14	0.47
3:d:216:SER:O	3:d:217:LEU:O	2.32	0.47
4:p:196:LYS:HD2	8:A:5:ARG:NH2	2.27	0.47
5:e:20:LYS:HG3	5:e:53:LEU:HD12	1.96	0.47
6:g:209:LYS:O	6:g:209:LYS:HG2	2.14	0.47
7:O:75:LEU:HD21	7:P:74:LEU:HD21	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:148:ILE:HD11	9:F:161:PRO:HB3	1.96	0.47
8:A:36:GLY:O	8:A:39:ILE:HG22	2.14	0.47
1:a:130:PRO:O	1:a:136:THR:OG1	2.26	0.47
1:a:203:VAL:O	1:a:207:LEU:HD13	2.14	0.47
2:b:71:ARG:CB	4:p:127:LEU:O	2.62	0.47
6:g:360:ALA:HB2	9:D:293:PRO:HD2	1.95	0.47
7:S:48:LYS:HG2	7:R:46:GLU:OE1	2.13	0.47
7:O:21:SER:OG	7:P:20:ALA:HA	2.13	0.47
9:F:411:ASP:O	9:F:418:ARG:NH2	2.47	0.47
8:A:153:ALA:HB3	8:A:358:ILE:CD1	2.41	0.47
8:A:413:ARG:NH2	8:A:444:GLY:CA	2.76	0.47
1:a:109:ASN:HB3	1:a:224:SER:HB3	1.96	0.47
1:a:114:LEU:CD1	2:b:37:VAL:C	2.83	0.47
1:a:116:PRO:HA	2:b:34:LEU:H	1.79	0.47
2:b:45:GLY:O	2:b:49:LEU:HB2	2.14	0.47
2:b:68:GLU:O	4:p:126:GLN:O	2.32	0.47
6:g:222:LYS:CE	6:g:224:ASP:OD1	2.61	0.47
6:g:359:GLY:HA3	9:D:293:PRO:CB	2.43	0.47
7:N:48:LYS:CD	7:M:46:GLU:OE1	2.62	0.47
7:M:48:LYS:O	7:L:46:GLU:OE2	2.32	0.47
7:L:55:LEU:HD23	7:K:54:LEU:HD11	1.97	0.47
8:E:68:GLU:HA	9:F:25:GLN:HB2	1.97	0.47
8:E:147:LEU:HD22	8:E:315:THR:HG21	1.95	0.47
8:E:430:THR:HG22	8:E:431:VAL:H	1.80	0.47
9:D:274:ASN:HD21	9:D:328:TYR:CB	2.27	0.47
2:b:96:ARG:O	4:p:155:ILE:CG2	2.59	0.47
3:d:148:ILE:HD12	8:C:17:ILE:CG1	2.41	0.47
8:E:48:VAL:HG23	8:E:52:GLU:OE1	2.14	0.47
8:E:302:ALA:HB2	8:E:314:MET:CE	2.40	0.47
9:F:179:THR:HG22	9:F:183:MET:HE3	1.97	0.47
8:C:146:PRO:HG2	8:C:373:ILE:HG23	1.96	0.47
1:a:120:ILE:HG22	2:b:30:ASN:CB	2.44	0.47
1:a:173:PRO:O	1:a:177:LEU:HG	2.14	0.47
2:b:77:GLN:CB	4:p:134:SER:CA	2.86	0.47
2:b:153:LEU:HD13	3:d:240:GLN:HE22	1.80	0.47
2:b:160:LEU:CG	2:b:163:GLU:HB2	2.45	0.47
3:d:181:THR:O	3:d:221:PHE:HA	2.14	0.47
4:p:193:GLU:CA	4:p:196:LYS:HG2	2.45	0.47
4:p:206:GLN:HE21	4:p:210:LEU:HD13	1.79	0.47
5:e:62:LEU:HD13	5:e:78:VAL:HG21	1.96	0.47
6:g:79:VAL:HG13	6:g:306:ARG:NH2	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:221:VAL:HG21	8:A:401:SER:HB2	1.94	0.47
6:g:360:ALA:H	9:D:293:PRO:CG	2.26	0.47
7:S:10:VAL:CG2	7:T:74:LEU:HD21	2.36	0.47
7:N:53:LEU:CD2	7:O:52:THR:CG2	2.93	0.47
7:N:74:LEU:HD21	7:M:75:LEU:HD21	1.97	0.47
8:E:42:ILE:CD1	8:E:89:VAL:HG11	2.23	0.47
8:E:156:ALA:HB2	8:E:438:ILE:HD11	1.96	0.47
9:F:163:ARG:HG2	9:F:374:MET:SD	2.54	0.47
9:F:330:PRO:HD2	9:F:339:PRO:HG3	1.97	0.47
8:C:210:VAL:HG13	8:C:219:MET:HE3	1.97	0.47
8:C:247:ALA:CB	8:C:314:MET:HE3	2.44	0.47
2:b:75:ILE:HD12	4:p:133:THR:HG23	1.43	0.47
2:b:91:ASP:HB2	4:p:148:MET:HE3	1.96	0.47
2:b:122:GLU:HB2	4:p:181:ILE:CD1	2.42	0.47
3:d:177:VAL:CG1	3:d:210:LYS:CE	2.93	0.47
5:e:100:GLU:O	5:e:104:ARG:HG3	2.15	0.47
6:g:132:GLY:O	6:g:136:MET:CG	2.57	0.47
7:P:75:LEU:HD21	7:Q:74:LEU:HD21	1.97	0.47
7:T:21:SER:OG	7:G:20:ALA:HA	2.14	0.47
8:E:263:ASP:OD1	8:E:319:ILE:HG22	2.15	0.47
8:E:284:ARG:HH12	8:E:330:TYR:CA	2.27	0.47
8:A:469:LYS:NZ	8:A:492:GLU:C	2.73	0.47
8:C:171:ASP:O	8:C:176:LYS:HE2	2.15	0.47
8:C:227:GLU:CG	8:C:239:ALA:HB2	2.43	0.47
8:C:392:GLU:CG	9:D:429:ARG:HH22	2.26	0.47
9:D:101:PRO:O	9:D:256:MET:HE2	2.14	0.47
1:a:50:LEU:HB3	4:p:100:LEU:HD21	1.96	0.47
2:b:88:VAL:CA	4:p:148:MET:CB	2.75	0.47
3:d:195:ALA:HB1	3:d:207:VAL:HG11	1.97	0.47
5:e:5:LEU:HD22	5:e:22:ILE:HG21	1.96	0.47
5:e:91:GLU:N	5:e:91:GLU:CD	2.72	0.47
6:g:82:GLY:C	6:g:84:PRO:HD2	2.40	0.47
6:g:138:LEU:HD13	6:g:167:TYR:CG	2.49	0.47
7:S:46:GLU:CD	7:T:48:LYS:HB3	2.38	0.47
8:E:461:LEU:HA	8:E:497:GLN:HG2	1.96	0.47
8:A:154:ILE:HD12	8:A:358:ILE:HD13	1.93	0.47
9:B:269:LEU:HD23	9:B:322:THR:HB	1.97	0.47
2:b:94:GLN:O	4:p:155:ILE:CD1	2.62	0.47
5:e:19:VAL:HG22	5:e:52:ARG:O	2.15	0.47
6:g:107:VAL:HG21	6:g:292:LEU:HG	1.97	0.47
6:g:121:MET:HA	6:g:158:ILE:O	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:251:GLU:CD	6:g:270:MET:SD	2.97	0.47
8:E:56:PHE:CE1	8:E:89:VAL:HB	2.50	0.47
8:E:214:GLN:CG	8:E:219:MET:HG3	2.42	0.47
8:C:53:LEU:HD22	8:C:61:ILE:CG2	2.45	0.47
1:a:28:TRP:CB	1:a:34:GLN:HB2	2.43	0.46
2:b:125:LYS:CB	4:p:185:LEU:HD13	2.44	0.46
2:b:148:ALA:HB2	4:p:211:SER:HA	1.92	0.46
4:p:214:ILE:CG1	4:p:215:VAL:N	2.76	0.46
6:g:360:ALA:CA	9:D:292:MET:HB3	2.44	0.46
7:L:42:GLN:NE2	7:K:43:PRO:CG	2.74	0.46
7:J:57:LEU:O	7:J:61:GLU:N	2.35	0.46
8:E:137:ILE:HG23	8:E:138:MET:HG2	1.97	0.46
8:E:249:TYR:OH	8:E:253:ARG:NH1	2.48	0.46
8:E:437:THR:HG23	8:E:458:LEU:HD13	1.97	0.46
9:F:19:ASN:HB2	9:F:93:ASP:HB3	1.97	0.46
9:B:71:ASN:H	8:C:9:ILE:CG1	2.27	0.46
8:C:27:VAL:CG2	8:C:47:GLU:OE1	2.62	0.46
8:C:424:PRO:HG2	8:C:427:ALA:CB	2.45	0.46
5:e:34:LEU:HB3	5:e:35:PRO:CD	2.45	0.46
8:E:343:ILE:HG22	8:E:358:ILE:HD12	1.98	0.46
8:E:403:LEU:HD13	8:E:407:THR:CG2	2.45	0.46
9:F:265:GLU:HG2	9:F:318:GLU:HB3	1.96	0.46
8:A:393:LEU:HD22	8:A:407:THR:HG23	1.97	0.46
9:B:247:MET:CE	9:B:285:VAL:HG21	2.45	0.46
8:C:284:ARG:HH21	9:D:296:VAL:HG12	1.81	0.46
8:C:379:VAL:HG12	8:C:438:ILE:HG22	1.97	0.46
9:D:143:ASP:OD2	9:D:315:SER:CB	2.61	0.46
1:a:239:ILE:O	1:a:242:SER:OG	2.27	0.46
2:b:122:GLU:CB	4:p:181:ILE:HG23	2.45	0.46
3:d:75:ALA:CB	3:d:160:GLU:CG	2.92	0.46
5:e:43:ALA:HB1	6:g:294:ALA:CB	2.45	0.46
6:g:83:ARG:N	6:g:84:PRO:CD	2.77	0.46
7:P:46:GLU:OE1	7:Q:48:LYS:HD3	2.15	0.46
7:Q:46:GLU:OE1	7:R:48:LYS:CD	2.63	0.46
7:L:48:LYS:HD3	7:K:46:GLU:OE1	2.16	0.46
8:E:393:LEU:HD22	8:E:396:PHE:CD2	2.51	0.46
8:A:109:ILE:CG2	8:A:225:VAL:HG22	2.45	0.46
9:B:50:LYS:HE2	9:B:92:ILE:CD1	2.44	0.46
9:B:280:GLN:O	9:B:284:GLU:HG3	2.14	0.46
8:C:51:GLY:O	8:C:95:ILE:HG23	2.14	0.46
8:C:349:LEU:HD22	8:C:354:ILE:HD12	1.95	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:261:ARG:HD3	9:D:321:ILE:CG1	2.46	0.46
5:e:20:LYS:N	5:e:53:LEU:HD13	2.30	0.46
6:g:227:ILE:HG23	6:g:242:ILE:CD1	2.46	0.46
7:H:43:PRO:CB	7:I:48:LYS:HZ2	2.23	0.46
8:E:239:ALA:N	8:E:240:PRO:HD2	2.31	0.46
9:B:174:ALA:O	9:B:354:ARG:NH2	2.48	0.46
9:D:288:LEU:HD23	9:D:288:LEU:HA	1.70	0.46
9:D:398:TYR:CZ	9:D:425:ARG:HD3	2.51	0.46
1:a:121:GLN:CG	2:b:25:ASP:OD1	2.62	0.46
2:b:103:ILE:CG2	4:p:162:MET:CB	2.90	0.46
3:d:74:THR:HB	3:d:154:ARG:HH12	1.80	0.46
3:d:109:GLU:N	3:d:110:PRO:CD	2.76	0.46
3:d:148:ILE:CG1	8:C:20:TYR:CD2	2.98	0.46
3:d:155:ILE:HA	3:d:158:VAL:HG23	1.96	0.46
7:R:57:LEU:O	7:R:61:GLU:N	2.35	0.46
8:E:165:ARG:CD	8:E:299:LEU:HB3	2.45	0.46
8:E:379:VAL:CG1	8:E:438:ILE:HB	2.46	0.46
9:F:376:GLN:O	9:F:380:VAL:HG22	2.15	0.46
8:C:27:VAL:HG12	8:C:46:ASP:HB2	1.97	0.46
8:C:56:PHE:CZ	8:C:76:LEU:HD21	2.51	0.46
8:C:185:LEU:HD23	8:C:218:ALA:HB2	1.97	0.46
9:D:407:ILE:HG22	9:D:408:LEU:CD2	2.45	0.46
1:a:72:PHE:CB	4:p:108:TYR:OH	2.64	0.46
2:b:79:GLU:HB2	4:p:137:VAL:HG12	1.97	0.46
2:b:122:GLU:CB	4:p:181:ILE:CG2	2.93	0.46
2:b:147:GLN:O	2:b:150:GLN:N	2.49	0.46
4:p:214:ILE:HB	4:p:218:VAL:HG13	1.98	0.46
6:g:120:LEU:O	6:g:157:ILE:HA	2.16	0.46
6:g:144:ARG:HA	6:g:144:ARG:HD2	1.40	0.46
6:g:240:CYS:CB	6:g:246:CYS:HG	1.96	0.46
7:S:21:SER:OG	7:T:20:ALA:HA	2.16	0.46
8:E:140:ARG:NH1	8:E:303:ALA:HB3	2.31	0.46
9:F:20:LEU:CD2	9:F:92:ILE:HG12	2.46	0.46
9:B:237:GLY:H	9:B:249:VAL:HG11	1.80	0.46
9:B:292:MET:CE	8:C:276:LEU:HD23	2.46	0.46
9:B:462:LEU:HD23	9:B:462:LEU:HA	1.78	0.46
8:C:82:MET:HE2	8:C:82:MET:N	2.30	0.46
8:C:354:ILE:HG12	8:C:422:LYS:HE2	1.97	0.46
1:a:72:PHE:HB3	4:p:108:TYR:CE2	2.50	0.46
1:a:114:LEU:CG	2:b:37:VAL:C	2.89	0.46
1:a:134:ILE:HG21	1:a:191:PHE:CE2	2.38	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:70:LEU:HB2	4:p:127:LEU:O	2.15	0.46
2:b:78:LEU:N	4:p:138:LYS:H	2.14	0.46
2:b:117:THR:HG23	2:b:118:LEU:N	2.31	0.46
2:b:125:LYS:HD2	4:p:185:LEU:HB3	1.97	0.46
2:b:172:ASN:HB3	3:d:235:MET:HB2	1.98	0.46
3:d:177:VAL:HG13	3:d:210:LYS:HE3	1.98	0.46
8:E:469:LYS:CE	8:E:493:ALA:CA	2.92	0.46
9:B:313:ILE:HG22	9:B:313:ILE:O	2.15	0.46
9:B:345:HIS:HE1	8:C:229:ALA:HB1	1.81	0.46
8:C:63:ILE:HD11	8:C:238:LEU:HD11	1.96	0.46
8:C:152:ILE:HG22	8:C:423:GLN:NE2	2.30	0.46
8:C:439:TYR:CD2	8:C:490:LEU:HD21	2.51	0.46
1:a:51:LEU:CD1	4:p:100:LEU:HD13	2.38	0.46
1:a:104:PHE:HD1	4:p:97:PHE:CE1	2.32	0.46
3:d:74:THR:CG2	3:d:154:ARG:HH12	2.29	0.46
4:p:106:LYS:O	4:p:111:PRO:CD	2.64	0.46
5:e:21:GLU:O	5:e:52:ARG:O	2.34	0.46
6:g:103:GLU:CB	7:R:41:ARG:NE	2.74	0.46
6:g:186:PRO:HD3	6:g:312:LEU:CD2	2.46	0.46
6:g:202:PHE:HZ	6:g:232:PRO:HG3	1.79	0.46
7:H:81:VAL:CG2	7:I:80:PHE:CZ	2.99	0.46
7:J:48:LYS:HZ3	7:I:43:PRO:CA	2.27	0.46
7:J:48:LYS:CD	7:I:46:GLU:OE1	2.62	0.46
8:E:29:THR:HG22	8:E:90:LYS:CB	2.45	0.46
8:A:494:ILE:C	8:A:498:MET:HE2	2.40	0.46
8:C:152:ILE:H	8:C:423:GLN:NE2	2.13	0.46
9:D:261:ARG:HD3	9:D:321:ILE:HD11	1.98	0.46
1:a:71:PHE:HE1	2:b:49:LEU:HD13	1.21	0.46
1:a:104:PHE:CG	4:p:97:PHE:CD1	3.00	0.46
4:p:85:ASP:O	4:p:86:PHE:HD1	1.99	0.46
4:p:112:LEU:O	4:p:116:MET:HG2	2.16	0.46
5:e:42:THR:HG22	6:g:285:GLU:CB	2.46	0.46
6:g:360:ALA:O	9:D:292:MET:HG2	2.15	0.46
8:E:385:LEU:HD23	8:E:385:LEU:N	2.31	0.46
9:F:22:ARG:HD2	9:F:88:GLY:O	2.15	0.46
9:F:426:LYS:HG2	9:F:474:PHE:HD2	1.81	0.46
2:b:82:ARG:CG	4:p:140:LEU:CD2	2.78	0.46
2:b:160:LEU:HD22	2:b:163:GLU:CB	2.46	0.46
3:d:119:VAL:HG11	8:E:31:THR:OG1	2.16	0.46
3:d:230:SER:OG	8:A:45:LEU:HA	2.16	0.46
8:E:193:ILE:CD1	8:E:250:PHE:CD1	2.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:E:355:ARG:HH12	9:D:389:GLN:NE2	2.11	0.46
8:A:193:ILE:CD1	8:A:250:PHE:CD1	2.99	0.46
9:D:274:ASN:HB3	9:D:277:ARG:HG2	1.97	0.46
2:b:71:ARG:HB2	4:p:127:LEU:O	2.16	0.45
2:b:84:ARG:HD3	4:p:141:GLU:HB3	1.97	0.45
4:p:152:ARG:CG	8:A:503:LEU:HD22	2.45	0.45
6:g:300:LEU:C	6:g:300:LEU:CD2	2.85	0.45
7:K:57:LEU:O	7:K:61:GLU:N	2.35	0.45
8:A:193:ILE:HD13	8:A:250:PHE:CD1	2.51	0.45
9:B:108:GLY:HA2	9:B:222:ILE:HD12	1.98	0.45
8:C:271:TYR:CG	8:C:294:LEU:HD22	2.51	0.45
9:D:261:ARG:HH11	9:D:321:ILE:HD12	1.81	0.45
1:a:114:LEU:N	2:b:37:VAL:HG11	2.28	0.45
2:b:81:ALA:N	4:p:140:LEU:CB	2.44	0.45
2:b:149:LEU:HA	4:p:214:ILE:CD1	2.46	0.45
2:b:176:PHE:CE2	2:b:180:ASN:ND2	2.73	0.45
3:d:72:ASP:H	8:A:70:ASN:ND2	2.14	0.45
5:e:38:ALA:CB	7:L:41:ARG:CG	2.74	0.45
7:R:62:ALA:HA	7:R:65:ILE:CG2	2.47	0.45
9:F:273:ASP:HA	9:F:274:ASN:HA	1.64	0.45
8:A:109:ILE:HG12	8:A:113:ALA:HA	1.95	0.45
8:A:446:LEU:HD21	8:A:457:TYR:CE2	2.51	0.45
9:B:50:LYS:O	9:B:57:GLN:NE2	2.49	0.45
8:C:48:VAL:CG2	8:C:49:MET:N	2.79	0.45
8:C:109:ILE:CD1	8:C:246:LEU:HD11	2.46	0.45
2:b:33:ASN:OD1	2:b:33:ASN:O	2.35	0.45
2:b:78:LEU:C	4:p:136:GLU:HG3	2.39	0.45
2:b:88:VAL:O	2:b:91:ASP:N	2.50	0.45
2:b:107:LYS:NZ	4:p:165:GLU:HG2	2.32	0.45
3:d:136:SER:OG	3:d:138:LEU:HD23	2.15	0.45
4:p:166:THR:HG23	4:p:167:GLN:N	2.30	0.45
4:p:203:LEU:CD2	8:A:7:ASP:O	2.57	0.45
7:P:81:VAL:CG2	7:Q:80:PHE:CZ	2.99	0.45
7:G:43:PRO:HB3	7:H:48:LYS:CD	2.46	0.45
8:E:279:ARG:HA	9:D:292:MET:HE2	1.98	0.45
9:F:336:ASP:O	9:F:339:PRO:HD2	2.17	0.45
9:F:384:HIS:O	9:F:384:HIS:HD2	1.99	0.45
8:A:190:GLN:H	8:A:190:GLN:CD	2.15	0.45
9:B:25:GLN:HB2	9:B:32:ASP:HB2	1.99	0.45
8:C:141:ARG:HD3	8:C:304:LYS:HE3	1.98	0.45
9:D:254:LEU:HD21	9:D:312:ARG:HB2	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:24:GLN:O	4:p:84:PHE:HD2	1.99	0.45
1:a:215:PRO:O	1:a:218:PHE:HB2	2.17	0.45
4:p:108:TYR:CE1	4:p:112:LEU:HD13	2.43	0.45
8:E:390:PHE:HE2	8:E:418:ARG:HD2	1.76	0.45
9:F:260:PHE:HD1	9:F:264:ASN:HD22	1.63	0.45
8:A:30:GLY:O	8:A:89:VAL:N	2.39	0.45
8:A:180:ALA:HB2	8:A:319:ILE:HD11	1.97	0.45
8:C:393:LEU:HD12	8:C:411:LEU:HD21	1.98	0.45
2:b:160:LEU:CD1	2:b:164:LEU:HB3	2.42	0.45
3:d:114:PHE:CE1	3:d:120:ILE:HG21	2.52	0.45
4:p:209:ALA:C	8:A:14:ARG:NH2	2.75	0.45
5:e:24:LEU:HD12	5:e:24:LEU:C	2.41	0.45
6:g:148:LEU:N	6:g:148:LEU:HD23	2.31	0.45
7:S:62:ALA:HA	7:S:65:ILE:CG2	2.47	0.45
7:P:62:ALA:HA	7:P:65:ILE:CG2	2.47	0.45
7:T:62:ALA:HA	7:T:65:ILE:CG2	2.47	0.45
7:G:75:LEU:HD21	7:H:74:LEU:HD21	1.99	0.45
7:L:74:LEU:HD21	7:K:10:VAL:CG2	2.44	0.45
8:E:32:VAL:HG22	8:E:56:PHE:CE1	2.51	0.45
8:A:42:ILE:HD12	8:A:74:VAL:HG21	1.97	0.45
8:A:149:THR:C	8:A:186:ASN:HD22	2.24	0.45
8:A:176:LYS:HD2	8:A:319:ILE:HG23	1.99	0.45
8:C:57:GLU:OE1	8:C:87:SER:HB2	2.16	0.45
2:b:85:LEU:HD12	2:b:86:LYS:CA	2.47	0.45
2:b:85:LEU:N	4:p:140:LEU:O	2.33	0.45
2:b:88:VAL:CG2	4:p:148:MET:CB	2.94	0.45
2:b:131:PHE:CZ	8:A:119:ARG:HA	2.51	0.45
2:b:132:GLU:CG	4:p:189:LEU:HD22	2.25	0.45
2:b:147:GLN:OE1	4:p:207:ILE:HG21	2.11	0.45
2:b:160:LEU:CB	2:b:163:GLU:HB2	2.44	0.45
2:b:173:ILE:HG13	3:d:221:PHE:CZ	2.50	0.45
4:p:193:GLU:HG3	4:p:196:LYS:HE3	1.83	0.45
6:g:227:ILE:HG21	6:g:242:ILE:CD1	2.47	0.45
7:S:43:PRO:HB3	7:T:48:LYS:CE	2.46	0.45
7:G:62:ALA:HA	7:G:65:ILE:CG2	2.47	0.45
7:G:68:LEU:HD22	7:H:66:TYR:CG	2.51	0.45
7:J:80:PHE:CD2	7:I:6:ALA:HB1	2.46	0.45
8:E:322:THR:HG22	8:E:332:PRO:HG2	1.99	0.45
8:E:360:VAL:O	8:E:360:VAL:HG22	2.17	0.45
8:A:156:ALA:O	8:A:372:GLN:NE2	2.49	0.45
8:C:169:ILE:HD11	8:C:322:THR:HG23	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:161:PHE:CE1	1:a:163:LYS:N	2.81	0.45
1:a:164:TYR:CE2	1:a:166:GLN:HA	2.52	0.45
1:a:195:LEU:CD1	4:p:84:PHE:CZ	3.00	0.45
2:b:92:ALA:N	4:p:148:MET:HA	2.30	0.45
2:b:118:LEU:O	2:b:121:PHE:HB3	2.16	0.45
2:b:121:PHE:CE2	2:b:125:LYS:HE2	2.51	0.45
3:d:186:LEU:CD1	3:d:191:LEU:CG	2.64	0.45
3:d:187:GLU:HB2	3:d:190:HIS:CE1	2.52	0.45
6:g:158:ILE:CD1	6:g:198:VAL:HG22	2.47	0.45
7:I:62:ALA:HA	7:I:65:ILE:CG2	2.47	0.45
8:E:146:PRO:HB2	8:E:148:GLN:HE21	1.81	0.45
8:E:302:ALA:HB1	8:E:314:MET:O	2.17	0.45
8:A:416:ARG:NE	8:A:447:ASP:HA	2.32	0.45
8:A:420:LEU:HD23	8:A:458:LEU:HD11	1.97	0.45
9:B:173:GLY:O	9:B:178:LYS:NZ	2.50	0.45
9:B:427:ILE:HD11	9:B:462:LEU:HD21	1.99	0.45
8:C:68:GLU:O	9:D:87:ARG:NH1	2.49	0.45
1:a:172:LEU:HD11	1:a:176:ILE:HD11	1.98	0.45
2:b:128:THR:HB	4:p:185:LEU:HD11	1.98	0.45
2:b:142:GLN:OE1	3:d:248:LEU:CG	2.61	0.45
7:T:33:GLY:CA	7:G:31:ALA:HA	2.47	0.45
8:E:128:ARG:HH22	8:E:252:TYR:HE2	1.65	0.45
9:F:384:HIS:O	9:F:384:HIS:CD2	2.70	0.45
9:F:405:ILE:HD13	9:F:410:LEU:HD21	1.97	0.45
8:C:99:PRO:O	8:C:104:TYR:OH	2.31	0.45
1:a:50:LEU:CB	4:p:100:LEU:HD21	2.47	0.45
2:b:110:LEU:HD22	4:p:167:GLN:HG2	1.25	0.45
3:d:71:VAL:HA	8:A:69:SER:C	2.42	0.45
4:p:192:LEU:CD1	8:A:3:THR:CG2	2.72	0.45
4:p:203:LEU:CD2	8:A:6:ALA:O	2.51	0.45
6:g:231:LEU:HD23	6:g:233:LEU:CD1	2.47	0.45
6:g:255:PHE:O	6:g:255:PHE:CD1	2.70	0.45
7:L:48:LYS:HB3	7:K:46:GLU:CD	2.40	0.45
7:J:80:PHE:CZ	7:I:81:VAL:HG22	2.52	0.45
9:D:170:LEU:HB2	9:D:326:ALA:HA	1.99	0.45
1:a:72:PHE:HB3	4:p:108:TYR:HE2	1.82	0.45
1:a:124:HIS:CD2	4:p:79:GLU:CD	2.95	0.45
7:K:62:ALA:HA	7:K:65:ILE:CG2	2.47	0.45
7:J:37:GLU:OE2	7:J:41:ARG:NH2	2.50	0.45
8:E:147:LEU:CD2	8:E:258:LEU:HD13	2.47	0.45
9:F:53:ASP:OD2	9:F:57:GLN:NE2	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:C:449:LEU:C	8:C:449:LEU:HD12	2.42	0.45
8:C:457:TYR:CD1	8:C:457:TYR:O	2.70	0.45
9:D:346:LEU:HD12	9:D:349:THR:HG22	1.98	0.45
1:a:39:VAL:CB	1:a:133:ASP:OD2	2.65	0.44
1:a:65:PRO:O	2:b:57:LYS:NZ	2.38	0.44
2:b:88:VAL:O	4:p:148:MET:CA	2.60	0.44
3:d:169:PHE:O	3:d:169:PHE:CD1	2.70	0.44
3:d:233:VAL:HG11	8:A:26:VAL:CA	2.40	0.44
4:p:152:ARG:CZ	8:A:502:LEU:HB3	2.47	0.44
7:N:53:LEU:CD2	7:O:52:THR:HG23	2.47	0.44
7:O:62:ALA:HA	7:O:65:ILE:CG2	2.47	0.44
7:P:40:ALA:CB	7:Q:49:ILE:HD11	2.37	0.44
7:M:80:PHE:CD2	7:L:6:ALA:HB1	2.43	0.44
7:G:6:ALA:HB1	7:H:80:PHE:CD2	2.42	0.44
7:L:62:ALA:HA	7:L:65:ILE:CG2	2.47	0.44
8:A:247:ALA:CB	8:A:314:MET:SD	3.05	0.44
8:A:471:GLU:CB	8:A:489:LEU:HD11	2.46	0.44
9:B:134:ARG:HE	9:B:258:GLU:CD	2.22	0.44
8:C:109:ILE:HD11	8:C:225:VAL:HG22	1.99	0.44
8:C:390:PHE:CD1	8:C:390:PHE:O	2.70	0.44
9:D:18:LYS:O	9:D:18:LYS:HG2	2.17	0.44
1:a:65:PRO:CG	1:a:70:ASN:ND2	2.64	0.44
1:a:133:ASP:C	1:a:191:PHE:CE1	2.89	0.44
2:b:70:LEU:O	2:b:74:ALA:CB	2.64	0.44
2:b:172:ASN:CB	3:d:235:MET:HB2	2.47	0.44
3:d:148:ILE:CD1	8:C:17:ILE:CB	2.95	0.44
4:p:128:SER:O	4:p:132:ASP:N	2.40	0.44
6:g:103:GLU:CG	7:R:41:ARG:HG3	2.48	0.44
6:g:213:LEU:HD12	6:g:213:LEU:C	2.43	0.44
7:M:51:GLY:C	7:L:50:ARG:NH2	2.76	0.44
8:E:181:THR:HG23	8:E:213:PHE:HE1	1.82	0.44
9:F:183:MET:HB3	9:F:437:VAL:HG11	1.98	0.44
9:B:274:ASN:HB3	9:B:277:ARG:HG2	1.99	0.44
8:C:227:GLU:OE1	8:C:227:GLU:HA	2.16	0.44
9:D:477:VAL:HG21	9:D:483:ALA:HB2	1.98	0.44
3:d:133:ILE:HD13	3:d:143:ALA:HB1	1.98	0.44
3:d:235:MET:CG	8:A:24:VAL:CG1	2.95	0.44
6:g:123:VAL:CG2	6:g:308:LEU:HD21	2.48	0.44
7:S:12:ALA:HB2	7:R:11:ILE:HA	2.00	0.44
7:M:80:PHE:CZ	7:L:81:VAL:HG22	2.51	0.44
8:E:82:MET:CE	8:E:82:MET:CA	2.86	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:E:106:GLY:CA	8:E:219:MET:HE3	2.41	0.44
8:E:271:TYR:CD1	8:E:294:LEU:CD1	3.01	0.44
8:E:302:ALA:HB2	8:E:314:MET:HG2	1.97	0.44
9:F:23:ILE:HG21	9:F:26:ILE:CD1	2.47	0.44
8:A:105:LEU:CD2	8:A:193:ILE:HG23	2.45	0.44
8:A:435:VAL:CG1	8:A:436:MET:N	2.80	0.44
9:B:279:VAL:HG13	9:B:302:LEU:CD1	2.48	0.44
9:B:299:GLN:O	9:B:299:GLN:HG2	2.16	0.44
9:B:311:GLU:HB2	8:C:230:ASP:HB3	1.97	0.44
1:a:144:THR:HG21	1:a:186:LEU:CD2	2.46	0.44
6:g:275:THR:O	6:g:275:THR:OG1	2.34	0.44
7:N:62:ALA:HA	7:N:65:ILE:CG2	2.47	0.44
7:J:70:VAL:HG13	7:I:75:LEU:HD11	2.00	0.44
7:J:74:LEU:CD2	7:I:10:VAL:HG22	2.37	0.44
7:J:74:LEU:HD22	7:I:10:VAL:HG21	1.92	0.44
8:A:390:PHE:CD2	8:A:418:ARG:NH2	2.80	0.44
9:B:29:PRO:HD2	9:B:288:LEU:CD1	2.47	0.44
9:D:237:GLY:HA3	9:D:249:VAL:HG21	1.98	0.44
1:a:134:ILE:C	1:a:134:ILE:CD1	2.85	0.44
2:b:78:LEU:N	4:p:138:LYS:N	2.63	0.44
3:d:102:LEU:HD22	3:d:146:ILE:HD11	2.00	0.44
3:d:157:LEU:HD12	3:d:161:ILE:CD1	2.48	0.44
6:g:281:ILE:N	7:N:41:ARG:HH21	2.14	0.44
7:S:43:PRO:HB3	7:T:48:LYS:HZ3	1.65	0.44
7:N:75:LEU:HD11	7:O:70:VAL:HG13	1.98	0.44
7:Q:62:ALA:HA	7:Q:65:ILE:CG2	2.47	0.44
7:T:53:LEU:CD2	7:G:52:THR:HG23	2.48	0.44
7:H:62:ALA:HA	7:H:65:ILE:CG2	2.47	0.44
7:K:48:LYS:HD3	7:J:46:GLU:OE1	2.17	0.44
9:F:310:GLN:HE21	9:F:345:HIS:HB3	1.82	0.44
8:A:77:MET:HB3	8:A:112:LEU:HD21	2.00	0.44
8:C:106:GLY:HA2	8:C:219:MET:HG3	1.98	0.44
8:C:322:THR:HG22	8:C:332:PRO:CG	2.47	0.44
9:D:180:VAL:HG23	12:D:600:ADP:O1A	2.18	0.44
9:D:380:VAL:HB	9:D:384:HIS:HD1	1.82	0.44
1:a:168:THR:CG2	1:a:170:ILE:HG12	2.47	0.44
3:d:165:PHE:CE1	3:d:169:PHE:HB2	2.53	0.44
6:g:85:PHE:CD1	6:g:85:PHE:O	2.71	0.44
6:g:122:VAL:HG23	6:g:157:ILE:HG23	1.98	0.44
7:M:52:THR:CG2	7:L:53:LEU:CD2	2.95	0.44
7:G:57:LEU:O	7:G:61:GLU:N	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:E:95:ILE:O	8:E:95:ILE:HG13	2.17	0.44
8:E:299:LEU:HD23	8:E:299:LEU:HA	1.77	0.44
9:F:425:ARG:HG2	9:F:425:ARG:NH1	2.32	0.44
8:C:25:LYS:HB3	8:C:27:VAL:HG13	1.98	0.44
8:C:322:THR:HG22	8:C:332:PRO:HG2	1.99	0.44
9:D:118:VAL:O	9:D:118:VAL:CG1	2.66	0.44
9:D:426:LYS:NZ	9:D:467:ASP:HA	2.32	0.44
1:a:161:PHE:CD1	1:a:162:GLY:N	2.85	0.44
2:b:144:VAL:CG2	8:A:9:ILE:HG22	2.27	0.44
3:d:157:LEU:CD1	3:d:161:ILE:HD11	2.48	0.44
5:e:22:ILE:CG2	5:e:51:ILE:CD1	2.95	0.44
7:L:31:ALA:HA	7:K:33:GLY:CA	2.45	0.44
7:J:42:GLN:NE2	7:I:43:PRO:CG	2.73	0.44
8:A:390:PHE:CE1	8:A:414:GLY:HA3	2.51	0.44
1:a:34:GLN:O	4:p:87:ASN:HB3	2.14	0.44
1:a:114:LEU:CD1	2:b:37:VAL:O	2.44	0.44
1:a:135:ASN:ND2	4:p:86:PHE:HD1	2.02	0.44
2:b:61:LEU:CD1	2:b:61:LEU:C	2.86	0.44
2:b:82:ARG:O	2:b:85:LEU:CD1	2.62	0.44
2:b:122:GLU:OE1	4:p:181:ILE:CD1	2.65	0.44
2:b:176:PHE:CD1	3:d:233:VAL:CB	3.00	0.44
7:S:9:SER:CB	7:R:10:VAL:HG21	2.48	0.44
7:S:64:THR:CG2	7:T:63:LEU:HD22	2.44	0.44
7:T:43:PRO:HB3	7:G:48:LYS:CE	2.48	0.44
8:E:169:ILE:HG23	8:E:344:PHE:HA	1.98	0.44
8:E:457:TYR:O	8:E:457:TYR:CD1	2.71	0.44
8:A:247:ALA:HB3	8:A:314:MET:SD	2.57	0.44
2:b:60:ILE:C	2:b:63:THR:HG22	2.40	0.44
2:b:64:ILE:HD12	4:p:123:ILE:HG13	1.89	0.44
3:d:199:GLN:CD	3:d:207:VAL:HG12	2.39	0.44
3:d:224:ARG:NH2	3:d:229:GLY:O	2.50	0.44
5:e:52:ARG:HA	5:e:57:TRP:HB3	2.00	0.44
6:g:109:LEU:HA	6:g:109:LEU:HD23	1.75	0.44
6:g:253:GLU:HB3	6:g:270:MET:CA	2.48	0.44
7:O:62:ALA:O	7:O:63:LEU:C	2.61	0.44
7:H:62:ALA:O	7:H:63:LEU:C	2.61	0.44
7:J:62:ALA:O	7:J:63:LEU:C	2.61	0.44
8:E:117:ASP:OD2	8:E:119:ARG:HD3	2.17	0.44
8:E:157:MET:HA	8:E:384:LYS:HE2	2.00	0.44
9:F:178:LYS:O	9:F:182:ILE:HG13	2.18	0.44
8:A:424:PRO:HG2	8:A:427:ALA:HB3	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:A:450:GLU:HB2	8:A:453:GLN:CG	2.48	0.44
9:B:475:TYR:O	9:B:486:LYS:NZ	2.49	0.44
8:C:96:ALA:HB1	8:C:130:ILE:HD12	2.00	0.44
2:b:70:LEU:CG	4:p:127:LEU:HD21	2.36	0.43
2:b:95:PHE:HZ	4:p:156:SER:CB	2.30	0.43
2:b:142:GLN:NE2	3:d:248:LEU:CD2	2.58	0.43
2:b:160:LEU:HD22	2:b:163:GLU:HB3	1.98	0.43
4:p:107:ILE:O	4:p:111:PRO:CG	2.64	0.43
4:p:166:THR:O	4:p:170:VAL:HB	2.18	0.43
5:e:90:GLN:C	5:e:90:GLN:CD	2.85	0.43
7:J:62:ALA:HA	7:J:65:ILE:CG2	2.47	0.43
7:I:5:ILE:HG22	7:I:81:VAL:HG13	2.00	0.43
8:A:424:PRO:HB3	8:A:455:ARG:HH12	1.83	0.43
8:A:494:ILE:O	8:A:498:MET:CG	2.63	0.43
1:a:26:PHE:C	4:p:85:ASP:OD2	2.61	0.43
1:a:62:GLN:C	4:p:115:PHE:HE2	2.26	0.43
1:a:117:TRP:HA	2:b:31:LEU:HB3	1.99	0.43
2:b:75:ILE:HG12	4:p:132:ASP:C	2.38	0.43
2:b:84:ARG:HB2	4:p:141:GLU:HA	1.87	0.43
2:b:88:VAL:C	2:b:90:MET:H	2.24	0.43
2:b:100:TYR:C	2:b:103:ILE:HG22	2.41	0.43
2:b:176:PHE:CE1	3:d:233:VAL:HG11	2.52	0.43
3:d:102:LEU:CD2	3:d:146:ILE:HD11	2.48	0.43
5:e:119:LEU:HD12	5:e:123:ARG:NH2	2.29	0.43
6:g:121:MET:HB3	6:g:158:ILE:HB	2.00	0.43
6:g:259:THR:CG2	9:F:404:ILE:HG21	2.48	0.43
7:S:48:LYS:HB3	7:R:46:GLU:CD	2.42	0.43
7:S:64:THR:HG21	7:T:63:LEU:HD23	1.96	0.43
7:K:48:LYS:CD	7:J:46:GLU:OE1	2.66	0.43
7:K:62:ALA:O	7:K:63:LEU:C	2.61	0.43
8:E:376:MET:CE	8:E:438:ILE:HD12	2.48	0.43
9:B:50:LYS:CE	9:B:92:ILE:CD1	2.96	0.43
9:B:148:ILE:HG12	9:B:149:PHE:H	1.82	0.43
9:B:170:LEU:HD23	9:B:350:THR:HB	1.99	0.43
9:B:275:ILE:CG1	9:B:327:VAL:HG22	2.48	0.43
9:B:336:ASP:HB3	9:B:339:PRO:CD	2.43	0.43
9:B:464:GLY:C	9:B:466:LEU:N	2.75	0.43
9:B:473:ALA:HB1	9:B:487:ALA:HB2	1.99	0.43
8:C:184:ILE:HD11	8:C:260:ILE:CG1	2.46	0.43
9:D:79:MET:HE3	9:D:113:VAL:HG11	1.99	0.43
3:d:101:LYS:HE2	3:d:101:LYS:HB2	1.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:188:ASN:N	3:d:188:ASN:ND2	2.66	0.43
3:d:222:THR:HB	3:d:234:ASP:HB3	2.01	0.43
4:p:107:ILE:O	4:p:111:PRO:HD2	2.18	0.43
4:p:127:LEU:C	4:p:127:LEU:HD23	2.44	0.43
5:e:38:ALA:HA	7:M:42:GLN:NE2	2.33	0.43
7:Q:62:ALA:O	7:Q:63:LEU:C	2.61	0.43
7:M:62:ALA:HA	7:M:65:ILE:CG2	2.47	0.43
7:T:62:ALA:O	7:T:63:LEU:C	2.61	0.43
7:I:62:ALA:O	7:I:63:LEU:C	2.61	0.43
9:F:330:PRO:CD	9:F:339:PRO:HG3	2.48	0.43
9:D:423:ARG:HH21	9:D:464:GLY:CA	2.29	0.43
1:a:116:PRO:HA	2:b:34:LEU:N	2.32	0.43
2:b:160:LEU:HD12	2:b:160:LEU:C	2.34	0.43
4:p:214:ILE:HB	4:p:218:VAL:CG1	2.49	0.43
6:g:104:ASP:HB2	7:S:41:ARG:CZ	2.47	0.43
6:g:279:SER:HA	6:g:280:PRO:HD3	1.86	0.43
7:N:20:ALA:HA	7:M:21:SER:OG	2.18	0.43
8:E:140:ARG:NE	9:F:206:THR:HG21	2.32	0.43
9:F:353:SER:HB3	9:F:356:LEU:HD13	2.00	0.43
8:A:379:VAL:HG12	8:A:438:ILE:HG22	2.00	0.43
8:A:413:ARG:CZ	8:A:444:GLY:N	2.82	0.43
9:B:48:ILE:CG2	9:B:59:MET:HE2	2.45	0.43
9:B:217:LYS:HA	9:B:222:ILE:O	2.17	0.43
8:C:392:GLU:OE2	9:D:429:ARG:CZ	2.66	0.43
8:C:424:PRO:HG2	8:C:427:ALA:HB3	2.00	0.43
1:a:35:ILE:HG21	4:p:88:LEU:CA	2.42	0.43
2:b:82:ARG:CA	2:b:85:LEU:HG	2.46	0.43
5:e:10:PRO:HD3	6:g:85:PHE:CE2	2.53	0.43
6:g:130:CYS:HA	6:g:319:ARG:HA	2.01	0.43
6:g:186:PRO:HD3	6:g:312:LEU:HD22	2.00	0.43
6:g:360:ALA:CA	9:D:293:PRO:HD3	2.37	0.43
7:S:3:PRO:HD2	7:S:5:ILE:CD1	2.49	0.43
7:S:10:VAL:HG22	7:T:74:LEU:HD22	1.98	0.43
7:N:3:PRO:HD2	7:N:5:ILE:CD1	2.49	0.43
7:N:33:GLY:CA	7:O:31:ALA:HA	2.49	0.43
7:M:3:PRO:HD2	7:M:5:ILE:CD1	2.49	0.43
7:M:74:LEU:HD22	7:L:10:VAL:HG21	1.92	0.43
7:H:3:PRO:HD2	7:H:5:ILE:CD1	2.49	0.43
7:J:9:SER:CB	7:I:10:VAL:HG21	2.49	0.43
8:E:180:ALA:HB1	8:E:260:ILE:HG21	1.99	0.43
8:E:393:LEU:HD22	8:E:396:PHE:CE2	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:F:425:ARG:HG2	9:F:425:ARG:HH11	1.82	0.43
8:A:227:GLU:HG3	8:A:239:ALA:HB2	2.01	0.43
8:A:379:VAL:HG12	8:A:438:ILE:HG21	1.99	0.43
9:B:345:HIS:HE1	8:C:229:ALA:CB	2.31	0.43
9:D:387:ILE:HD11	9:D:455:ILE:HG12	2.00	0.43
9:D:401:LEU:O	9:D:404:ILE:HG12	2.19	0.43
1:a:193:ASN:ND2	1:a:227:GLN:CD	2.74	0.43
2:b:77:GLN:HG2	4:p:138:LYS:CA	2.48	0.43
3:d:183:VAL:CG2	3:d:214:ASP:O	2.63	0.43
3:d:241:LEU:HA	3:d:241:LEU:HD23	1.77	0.43
4:p:193:GLU:CG	4:p:196:LYS:CE	2.58	0.43
7:O:53:LEU:CD2	7:P:52:THR:CG2	2.97	0.43
7:J:3:PRO:HD2	7:J:5:ILE:CD1	2.49	0.43
8:A:154:ILE:CD1	8:A:358:ILE:HD13	2.48	0.43
8:A:183:THR:O	8:A:187:GLN:HG2	2.18	0.43
8:C:237:TYR:CZ	8:C:294:LEU:HD11	2.53	0.43
1:a:42:THR:HG23	1:a:117:TRP:HZ3	1.83	0.43
1:a:113:ALA:C	2:b:34:LEU:CD1	2.83	0.43
1:a:119:ILE:HG12	2:b:28:ALA:CA	2.42	0.43
1:a:165:ILE:HG21	1:a:169:PRO:O	2.18	0.43
2:b:48:VAL:HG22	2:b:52:LEU:HD13	2.00	0.43
2:b:75:ILE:CD1	4:p:132:ASP:OD2	2.63	0.43
2:b:131:PHE:CE1	8:A:119:ARG:HA	2.54	0.43
2:b:175:MET:O	2:b:179:MET:HG2	2.19	0.43
7:S:55:LEU:HD21	7:R:54:LEU:HD11	1.99	0.43
7:S:81:VAL:CG2	7:T:80:PHE:CZ	3.01	0.43
7:P:3:PRO:HD2	7:P:5:ILE:CD1	2.49	0.43
7:T:3:PRO:HD2	7:T:5:ILE:CD1	2.49	0.43
7:G:3:PRO:HD2	7:G:5:ILE:CD1	2.49	0.43
7:G:75:LEU:HD11	7:H:70:VAL:HG13	2.00	0.43
8:A:148:GLN:HG3	8:A:431:VAL:CG2	2.48	0.43
8:A:469:LYS:HZ3	8:A:493:ALA:N	2.16	0.43
8:C:82:MET:HE2	8:C:82:MET:CA	2.48	0.43
1:a:208:VAL:CG2	1:a:209:PRO:HD3	2.38	0.43
2:b:179:MET:HG3	8:A:24:VAL:HG23	2.01	0.43
3:d:117:ASN:HA	3:d:118:PRO:HD3	1.65	0.43
3:d:198:VAL:C	3:d:201:ILE:HG12	2.44	0.43
3:d:233:VAL:O	3:d:233:VAL:HG23	2.18	0.43
3:d:247:GLN:HA	8:A:16:ARG:CG	2.19	0.43
4:p:179:LYS:HD3	4:p:183:VAL:HG23	1.99	0.43
6:g:53:SER:HB3	9:D:403:ASP:CG	2.38	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:253:GLU:CA	6:g:270:MET:HA	2.49	0.43
7:S:80:PHE:CZ	7:R:81:VAL:CG2	3.01	0.43
7:N:10:VAL:HG21	7:O:74:LEU:HD22	1.93	0.43
7:R:62:ALA:O	7:R:63:LEU:C	2.61	0.43
7:M:62:ALA:O	7:M:63:LEU:C	2.61	0.43
7:M:66:TYR:CD1	7:L:68:LEU:HD22	2.54	0.43
7:L:57:LEU:HD23	7:L:60:MET:HE3	2.01	0.43
8:E:237:TYR:CD2	8:E:274:MET:HE1	2.54	0.43
8:E:387:LEU:HD23	8:E:387:LEU:HA	1.65	0.43
8:E:490:LEU:HD23	8:E:494:ILE:HG13	2.01	0.43
8:C:48:VAL:CG2	8:C:52:GLU:HB3	2.49	0.43
8:C:439:TYR:HB3	8:C:490:LEU:HD21	2.00	0.43
9:D:267:ASP:OD1	9:D:267:ASP:N	2.47	0.43
1:a:30:ILE:HG23	1:a:32:GLY:H	1.83	0.43
4:p:100:LEU:O	4:p:104:LEU:HB3	2.18	0.43
4:p:112:LEU:HD23	4:p:112:LEU:HA	1.87	0.43
4:p:192:LEU:CD2	8:A:3:THR:HG23	2.47	0.43
4:p:210:LEU:HD13	8:A:13:ILE:HB	1.87	0.43
4:p:214:ILE:HB	8:A:17:ILE:HD12	1.94	0.43
6:g:118:VAL:HG12	6:g:148:LEU:HD13	2.00	0.43
7:P:62:ALA:O	7:P:63:LEU:C	2.61	0.43
7:R:3:PRO:HD2	7:R:5:ILE:CD1	2.49	0.43
7:T:57:LEU:HD23	7:T:60:MET:HE3	2.01	0.43
7:G:43:PRO:CA	7:H:48:LYS:HZ3	2.32	0.43
7:K:3:PRO:HD2	7:K:5:ILE:CD1	2.49	0.43
7:J:57:LEU:HD23	7:J:60:MET:HE3	2.01	0.43
9:F:269:LEU:HD13	9:F:271:PHE:CZ	2.54	0.43
9:B:386:GLU:O	9:B:390:ARG:HG3	2.19	0.43
8:C:151:LEU:HD12	8:C:154:ILE:HD13	2.00	0.43
8:C:237:TYR:HE1	8:C:271:TYR:CG	2.34	0.43
9:D:408:LEU:HB3	9:D:412:GLU:HG3	1.99	0.43
1:a:116:PRO:HG2	2:b:31:LEU:HB2	2.01	0.43
5:e:110:ARG:CD	6:g:206:GLU:OE2	2.67	0.43
6:g:265:THR:OG1	6:g:266:VAL:N	2.52	0.43
7:P:46:GLU:OE1	7:Q:48:LYS:CD	2.66	0.43
7:R:57:LEU:HD23	7:R:60:MET:HE3	2.01	0.43
8:E:261:TYR:CZ	8:E:298:LEU:HD11	2.54	0.43
9:F:178:LYS:HZ2	9:F:182:ILE:HD11	1.84	0.43
9:F:474:PHE:CD1	9:F:483:ALA:HB1	2.54	0.43
8:C:249:TYR:CD1	8:C:249:TYR:C	2.96	0.43
9:D:108:GLY:N	9:D:232:VAL:O	2.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:126:GLU:O	1:a:126:GLU:HG2	2.19	0.42
1:a:172:LEU:N	1:a:173:PRO:CD	2.82	0.42
2:b:96:ARG:HA	4:p:155:ILE:H	1.80	0.42
2:b:117:THR:HG21	4:p:174:LEU:HD11	1.89	0.42
2:b:125:LYS:CB	4:p:185:LEU:CB	2.81	0.42
3:d:78:TYR:HE1	8:C:8:GLU:OE1	2.02	0.42
3:d:129:LEU:HD22	3:d:150:ILE:CD1	2.48	0.42
4:p:159:LEU:HD12	4:p:160:ASN:CA	2.47	0.42
5:e:22:ILE:HG13	5:e:33:VAL:HG21	1.70	0.42
7:N:6:ALA:HB1	7:O:80:PHE:CD2	2.43	0.42
7:O:81:VAL:HG21	7:P:80:PHE:CZ	2.53	0.42
7:Q:3:PRO:HD2	7:Q:5:ILE:CD1	2.49	0.42
7:M:55:LEU:HD21	7:L:54:LEU:CG	2.47	0.42
7:L:20:ALA:HA	7:K:21:SER:OG	2.18	0.42
7:L:63:LEU:HD21	7:K:61:GLU:OE1	2.19	0.42
8:A:359:ASN:O	8:A:363:SER:OG	2.32	0.42
9:B:202:VAL:N	9:B:273:ASP:O	2.45	0.42
9:B:296:VAL:CG1	9:B:297:GLY:N	2.80	0.42
9:D:186:ILE:O	9:D:190:ALA:CB	2.67	0.42
1:a:27:TYR:OH	1:a:122:LEU:HD11	2.18	0.42
1:a:196:ALA:HB1	7:Q:65:ILE:HD11	2.01	0.42
3:d:148:ILE:HA	8:C:20:TYR:CE2	2.54	0.42
5:e:43:ALA:O	6:g:286:GLN:NE2	2.52	0.42
7:S:49:ILE:HD11	7:R:40:ALA:CB	2.39	0.42
7:T:46:GLU:OE1	7:G:48:LYS:HD3	2.19	0.42
7:H:81:VAL:CG2	7:I:80:PHE:CE2	3.01	0.42
8:A:196:TYR:HD2	8:A:224:VAL:HG13	1.84	0.42
8:A:379:VAL:HG21	8:A:435:VAL:HG23	2.00	0.42
9:B:261:ARG:O	9:B:265:GLU:HA	2.19	0.42
1:a:79:ILE:CD1	1:a:99:GLY:HA3	2.41	0.42
2:b:89:GLU:O	2:b:93:ASP:CG	2.62	0.42
3:d:119:VAL:HG11	8:E:43:HIS:HD2	1.85	0.42
7:N:48:LYS:HZ3	7:M:43:PRO:HB3	1.68	0.42
7:N:49:ILE:HD11	7:M:40:ALA:CB	2.44	0.42
7:N:66:TYR:CG	7:M:68:LEU:HD22	2.53	0.42
7:O:3:PRO:HD2	7:O:5:ILE:CD1	2.49	0.42
7:K:20:ALA:HA	7:J:21:SER:OG	2.19	0.42
7:K:48:LYS:CE	7:J:43:PRO:HB3	2.49	0.42
9:F:178:LYS:HZ3	9:F:182:ILE:HD11	1.81	0.42
9:F:186:ILE:HG21	9:F:221:VAL:HG11	2.01	0.42
8:A:107:ARG:NH2	8:A:117:ASP:OD1	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:64:GLU:O	9:D:76:ALA:HA	2.19	0.42
9:D:261:ARG:NH1	9:D:321:ILE:HD12	2.34	0.42
2:b:92:ALA:CB	4:p:148:MET:O	2.67	0.42
2:b:107:LYS:CD	4:p:166:THR:N	2.81	0.42
2:b:133:GLN:O	2:b:137:ILE:CG1	2.66	0.42
5:e:8:LEU:HD12	6:g:85:PHE:HD2	1.84	0.42
6:g:83:ARG:HH21	6:g:273:THR:HG22	1.83	0.42
7:N:39:ILE:CG1	7:N:49:ILE:HG21	2.50	0.42
7:T:46:GLU:OE1	7:G:48:LYS:CD	2.67	0.42
7:T:53:LEU:CD2	7:G:52:THR:CG2	2.97	0.42
8:E:374:LYS:HB2	8:E:480:LYS:O	2.18	0.42
9:D:463:SER:OG	9:D:465:GLU:HG3	2.20	0.42
1:a:72:PHE:CD2	4:p:108:TYR:CE2	3.07	0.42
1:a:134:ILE:HD12	1:a:135:ASN:HA	2.01	0.42
2:b:67:SER:CB	4:p:124:LYS:HA	2.47	0.42
2:b:82:ARG:C	4:p:140:LEU:HD22	2.44	0.42
2:b:103:ILE:HD12	4:p:158:ALA:C	2.44	0.42
2:b:155:THR:O	2:b:159:CYS:SG	2.75	0.42
2:b:165:HIS:CB	3:d:219:ALA:HB1	2.44	0.42
3:d:114:PHE:CE1	8:E:20:TYR:CD2	3.07	0.42
5:e:50:ARG:HG2	5:e:59:THR:CG2	2.41	0.42
6:g:131:GLY:HA2	9:B:411:ASP:OD2	2.19	0.42
7:O:75:LEU:HD11	7:P:70:VAL:HG13	2.01	0.42
7:Q:39:ILE:CG1	7:Q:49:ILE:HG21	2.49	0.42
7:G:33:GLY:CA	7:H:31:ALA:HA	2.47	0.42
7:L:3:PRO:HD2	7:L:5:ILE:CD1	2.49	0.42
8:A:164:GLN:CD	8:A:367:VAL:CG2	2.92	0.42
8:A:241:TYR:OH	8:A:294:LEU:O	2.38	0.42
8:A:379:VAL:HG21	8:A:435:VAL:CG2	2.49	0.42
9:B:251:LEU:HD21	9:B:309:LEU:HD13	1.93	0.42
9:B:385:TYR:OH	9:B:389:GLN:NE2	2.52	0.42
8:C:106:GLY:HA2	8:C:219:MET:O	2.20	0.42
9:D:191:LYS:NZ	9:D:219:SER:CB	2.75	0.42
1:a:24:GLN:CA	4:p:83:LEU:HD12	2.45	0.42
1:a:166:GLN:N	1:a:167:PRO:CD	2.82	0.42
4:p:199:THR:CG2	8:A:6:ALA:H	2.32	0.42
5:e:52:ARG:CG	5:e:57:TRP:HB3	2.50	0.42
6:g:259:THR:HG21	9:F:404:ILE:HG21	2.01	0.42
7:N:11:ILE:HA	7:O:12:ALA:HB2	2.02	0.42
7:O:57:LEU:HD23	7:O:60:MET:HE3	2.01	0.42
7:P:21:SER:OG	7:Q:20:ALA:HA	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:R:39:ILE:CG1	7:R:49:ILE:HG21	2.50	0.42
7:G:57:LEU:HD23	7:G:60:MET:HE3	2.01	0.42
7:K:52:THR:HG23	7:J:53:LEU:CD2	2.50	0.42
7:I:57:LEU:HD23	7:I:60:MET:HE3	2.01	0.42
8:E:230:ASP:CB	9:D:311:GLU:HG3	2.46	0.42
8:E:413:ARG:NH1	8:E:444:GLY:H	2.17	0.42
9:D:57:GLN:HA	9:D:57:GLN:OE1	2.19	0.42
9:D:405:ILE:O	9:D:409:GLY:N	2.53	0.42
2:b:147:GLN:O	2:b:148:ALA:C	2.62	0.42
3:d:148:ILE:CG1	8:C:20:TYR:HD2	2.33	0.42
3:d:169:PHE:CD1	3:d:169:PHE:C	2.96	0.42
4:p:87:ASN:OD1	4:p:89:THR:CG2	2.61	0.42
5:e:62:LEU:HD12	5:e:62:LEU:O	2.20	0.42
6:g:43:ASN:O	6:g:47:LEU:HG	2.20	0.42
7:N:48:LYS:HZ3	7:M:43:PRO:HB2	1.69	0.42
7:T:6:ALA:HB1	7:G:80:PHE:CD2	2.48	0.42
7:G:10:VAL:HG21	7:H:74:LEU:HD22	1.96	0.42
7:G:39:ILE:CG1	7:G:49:ILE:HG21	2.50	0.42
7:H:10:VAL:HG22	7:I:74:LEU:CD2	2.47	0.42
7:H:39:ILE:CG1	7:H:49:ILE:HG21	2.50	0.42
7:I:39:ILE:CG1	7:I:49:ILE:HG21	2.50	0.42
9:F:48:ILE:HD12	9:F:94:THR:CG2	2.48	0.42
9:F:124:VAL:CG1	9:F:126:THR:CG2	2.98	0.42
8:A:39:ILE:HD12	8:A:278:LEU:HD21	1.97	0.42
8:A:132:SER:HB2	8:A:133:PRO:HD2	2.01	0.42
8:A:185:LEU:HA	8:A:185:LEU:HD23	1.78	0.42
8:A:247:ALA:HB1	8:A:257:THR:HG21	2.01	0.42
8:A:340:ASP:O	8:A:366:ARG:HB2	2.20	0.42
8:C:109:ILE:HD11	8:C:246:LEU:CD1	2.50	0.42
8:C:240:PRO:HG2	8:C:267:GLN:NE2	2.34	0.42
9:D:387:ILE:HG13	9:D:388:ALA:N	2.33	0.42
2:b:82:ARG:C	4:p:140:LEU:CD2	2.93	0.42
3:d:77:ARG:NH2	8:C:8:GLU:O	2.53	0.42
7:P:57:LEU:HD23	7:P:60:MET:HE3	2.01	0.42
7:Q:57:LEU:HD23	7:Q:60:MET:HE3	2.01	0.42
7:M:39:ILE:CG1	7:M:49:ILE:HG21	2.50	0.42
8:E:439:TYR:CZ	8:E:487:GLU:HG3	2.55	0.42
8:A:420:LEU:CD2	8:A:458:LEU:CD1	2.97	0.42
8:C:387:LEU:HD11	8:C:421:LEU:HD11	2.02	0.42
9:D:338:ALA:CB	9:D:339:PRO:HD3	2.37	0.42
2:b:91:ASP:HB3	4:p:148:MET:HB3	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:234:ASP:OD1	3:d:238:LYS:HD2	2.20	0.42
3:d:235:MET:CE	8:A:22:ARG:HH21	2.25	0.42
4:p:144:ALA:HA	4:p:147:VAL:HG12	2.02	0.42
4:p:210:LEU:CD1	8:A:14:ARG:HG3	2.49	0.42
5:e:62:LEU:C	5:e:63:MET:HE2	2.44	0.42
6:g:65:LYS:HD3	6:g:324:SER:HB2	2.02	0.42
6:g:212:MET:HE3	6:g:212:MET:HB2	1.78	0.42
7:N:53:LEU:HD23	7:O:52:THR:HG23	2.01	0.42
7:P:39:ILE:CG1	7:P:49:ILE:HG21	2.50	0.42
7:T:29:GLY:O	7:G:31:ALA:HB2	2.19	0.42
7:K:80:PHE:CD2	7:J:6:ALA:HB1	2.49	0.42
8:E:7:ASP:OD1	8:E:8:GLU:N	2.53	0.42
8:E:117:ASP:OD1	8:E:117:ASP:N	2.49	0.42
8:C:10:SER:O	8:C:10:SER:OG	2.38	0.42
9:D:182:ILE:CG2	9:D:183:MET:N	2.83	0.42
1:a:124:HIS:HB3	4:p:81:ALA:O	2.19	0.42
1:a:230:ILE:HD11	7:Q:55:LEU:CD2	2.49	0.42
2:b:80:LYS:N	4:p:137:VAL:HG12	2.29	0.42
2:b:160:LEU:HD11	2:b:164:LEU:N	0.39	0.42
7:S:57:LEU:HD23	7:S:60:MET:HE3	2.01	0.42
7:N:62:ALA:O	7:N:63:LEU:C	2.61	0.42
7:L:39:ILE:CG1	7:L:49:ILE:HG21	2.49	0.42
7:K:57:LEU:HD23	7:K:60:MET:HE3	2.01	0.42
8:E:148:GLN:O	8:E:186:ASN:ND2	2.49	0.42
8:A:282:PRO:HB2	8:A:286:ALA:HA	2.02	0.42
8:A:495:GLN:O	8:A:499:GLU:CG	2.68	0.42
9:B:221:VAL:HG12	9:B:232:VAL:CB	2.49	0.42
9:B:261:ARG:HG3	9:B:319:GLY:HA3	2.02	0.42
8:C:387:LEU:HD23	8:C:387:LEU:HA	1.83	0.42
8:C:392:GLU:OE2	9:D:429:ARG:NH2	2.49	0.42
3:d:149:LEU:HD13	3:d:158:VAL:HG22	2.01	0.41
3:d:198:VAL:O	3:d:202:THR:N	2.44	0.41
6:g:363:CYS:HB3	9:D:290:GLY:CA	2.50	0.41
9:F:417:ASP:O	9:F:421:VAL:HG23	2.19	0.41
8:C:81:LEU:C	8:C:82:MET:HE2	2.45	0.41
8:C:138:MET:HE1	9:D:213:TYR:HE2	1.85	0.41
9:D:49:VAL:HG22	9:D:91:VAL:HG22	2.01	0.41
9:D:353:SER:HB2	9:D:366:ASP:HB2	2.02	0.41
2:b:88:VAL:N	4:p:147:VAL:CG1	2.75	0.41
2:b:105:ARG:O	2:b:108:MET:HB2	2.19	0.41
2:b:149:LEU:HD21	3:d:244:ILE:HG13	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:86:ALA:HB1	3:d:95:THR:CG2	2.48	0.41
5:e:4:ASN:HB2	5:e:18:GLU:OE2	2.19	0.41
6:g:64:MET:SD	9:D:408:LEU:HD13	2.60	0.41
6:g:72:VAL:CG2	6:g:316:LEU:HB2	2.50	0.41
7:O:39:ILE:CG1	7:O:49:ILE:HG21	2.50	0.41
7:T:43:PRO:C	7:G:48:LYS:NZ	2.71	0.41
7:T:75:LEU:HD21	7:G:74:LEU:HD21	2.02	0.41
9:F:61:VAL:HG13	9:F:80:SER:HB2	2.01	0.41
9:F:334:LEU:HD22	9:F:343:PHE:CE2	2.56	0.41
9:F:387:ILE:CG2	9:F:388:ALA:N	2.83	0.41
9:F:405:ILE:HD11	9:F:421:VAL:CG2	2.47	0.41
8:A:420:LEU:HD22	8:A:437:THR:CG2	2.50	0.41
9:B:154:LYS:NZ	9:B:430:PHE:O	2.51	0.41
9:B:183:MET:HE1	9:B:212:LEU:HD22	2.02	0.41
8:C:237:TYR:CZ	8:C:271:TYR:CD1	3.06	0.41
8:C:457:TYR:CD1	8:C:457:TYR:C	2.94	0.41
9:D:167:LYS:CE	9:D:310:GLN:OE1	2.67	0.41
9:D:202:VAL:HG11	9:D:278:PHE:HB2	2.03	0.41
1:a:42:THR:HG23	1:a:117:TRP:CZ3	2.55	0.41
1:a:119:ILE:N	2:b:31:LEU:N	2.65	0.41
2:b:70:LEU:C	4:p:130:VAL:HG22	2.45	0.41
2:b:104:GLU:O	2:b:108:MET:HG3	2.20	0.41
2:b:122:GLU:HA	4:p:181:ILE:HG23	1.91	0.41
2:b:160:LEU:CD2	2:b:164:LEU:N	2.79	0.41
6:g:221:VAL:HG22	8:A:401:SER:OG	2.20	0.41
7:S:46:GLU:OE1	7:T:48:LYS:HD3	2.20	0.41
7:S:62:ALA:O	7:S:63:LEU:C	2.61	0.41
7:N:29:GLY:O	7:O:31:ALA:HB2	2.20	0.41
7:O:40:ALA:CB	7:P:49:ILE:HD11	2.41	0.41
7:M:57:LEU:HD23	7:M:60:MET:HE3	2.01	0.41
7:L:62:ALA:O	7:L:63:LEU:C	2.61	0.41
7:K:52:THR:CG2	7:J:53:LEU:CD2	2.99	0.41
8:E:246:LEU:HD23	8:E:246:LEU:HA	1.79	0.41
8:A:140:ARG:HB2	9:B:210:ASN:ND2	2.28	0.41
9:B:387:ILE:CG2	9:B:388:ALA:N	2.82	0.41
9:B:412:GLU:O	9:B:413:LEU:C	2.64	0.41
8:C:170:GLY:O	8:C:321:GLU:HA	2.21	0.41
8:C:436:MET:CE	8:C:462:ARG:HA	2.50	0.41
2:b:31:LEU:HD12	2:b:32:ILE:HD13	2.03	0.41
2:b:133:GLN:CG	8:A:3:THR:HA	2.47	0.41
2:b:149:LEU:CD2	3:d:244:ILE:CG1	2.99	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:149:LEU:HD22	3:d:244:ILE:CG1	2.50	0.41
2:b:155:THR:O	2:b:159:CYS:N	2.53	0.41
5:e:88:ASP:HA	5:e:89:PRO:HD3	1.95	0.41
6:g:121:MET:HE3	6:g:121:MET:HB2	1.85	0.41
7:R:62:ALA:HA	7:R:65:ILE:HG22	2.03	0.41
9:B:63:CYS:HB3	9:B:76:ALA:HB1	2.03	0.41
9:B:402:GLN:HG2	9:B:405:ILE:HD11	2.02	0.41
2:b:92:ALA:CA	4:p:148:MET:O	2.67	0.41
2:b:162:ASN:O	2:b:166:LEU:CB	2.69	0.41
3:d:92:LEU:HD11	3:d:172:ILE:CD1	2.50	0.41
3:d:239:LYS:HE2	8:A:22:ARG:NH2	2.35	0.41
4:p:203:LEU:CD2	8:A:7:ASP:HA	2.49	0.41
6:g:232:PRO:O	6:g:233:LEU:C	2.63	0.41
7:T:39:ILE:CG1	7:T:49:ILE:HG21	2.50	0.41
7:G:62:ALA:O	7:G:63:LEU:C	2.61	0.41
7:H:53:LEU:CD2	7:I:52:THR:CG2	2.99	0.41
7:J:50:ARG:HG2	7:J:54:LEU:HD13	2.03	0.41
8:E:469:LYS:CE	8:E:493:ALA:HA	2.51	0.41
9:B:168:ILE:HB	9:B:324:ILE:HA	2.03	0.41
9:B:276:PHE:HB3	9:B:328:TYR:HB3	2.03	0.41
8:C:237:TYR:CE1	8:C:271:TYR:CB	3.03	0.41
8:C:376:MET:O	8:C:376:MET:HG3	2.20	0.41
9:D:64:GLU:CD	9:D:248:ARG:HE	2.25	0.41
9:D:168:ILE:CD1	9:D:324:ILE:HG12	2.51	0.41
7:S:43:PRO:CG	7:T:42:GLN:NE2	2.77	0.41
7:Q:81:VAL:CG2	7:R:80:PHE:CZ	3.04	0.41
7:K:31:ALA:HA	7:J:33:GLY:CA	2.49	0.41
7:J:39:ILE:CG1	7:J:49:ILE:HG21	2.49	0.41
8:E:379:VAL:CG1	8:E:438:ILE:CB	2.97	0.41
9:F:119:ASP:O	9:F:120:ASN:HB3	2.19	0.41
9:F:405:ILE:CG1	9:F:413:LEU:HD11	2.47	0.41
8:A:495:GLN:HA	8:A:498:MET:HE3	2.03	0.41
9:B:151:THR:HG22	9:B:189:ILE:HD11	2.02	0.41
9:B:182:ILE:O	9:B:186:ILE:HG13	2.21	0.41
9:B:251:LEU:HD23	9:B:251:LEU:N	2.34	0.41
9:D:29:PRO:HD2	9:D:288:LEU:CD1	2.51	0.41
2:b:60:ILE:HD11	4:p:120:ASP:CB	2.49	0.41
4:p:147:VAL:HG13	4:p:148:MET:N	2.34	0.41
6:g:129:LEU:HD23	9:B:412:GLU:OE2	2.18	0.41
6:g:363:CYS:HB3	9:D:290:GLY:HA2	2.02	0.41
7:N:57:LEU:HD23	7:N:60:MET:HE3	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:P:68:LEU:HD22	7:Q:66:TYR:CG	2.55	0.41
7:M:65:ILE:HD12	7:M:68:LEU:HD23	2.03	0.41
7:G:10:VAL:HG22	7:H:74:LEU:CD2	2.43	0.41
7:H:57:LEU:HD23	7:H:60:MET:HE3	2.01	0.41
8:E:29:THR:HA	8:E:89:VAL:O	2.20	0.41
8:E:90:LYS:HG3	8:E:90:LYS:O	2.19	0.41
8:A:344:PHE:HD2	8:A:362:ILE:O	2.04	0.41
8:C:126:GLU:OE2	8:C:253:ARG:NH2	2.54	0.41
8:C:342:GLN:HG3	8:C:344:PHE:CE2	2.55	0.41
9:D:112:ASN:CG	9:D:118:VAL:CG2	2.87	0.41
3:d:172:ILE:CG2	3:d:173:THR:N	2.83	0.41
4:p:87:ASN:OD1	4:p:87:ASN:C	2.64	0.41
5:e:51:ILE:O	5:e:51:ILE:CG2	2.67	0.41
6:g:253:GLU:OE1	6:g:253:GLU:N	2.53	0.41
6:g:253:GLU:CB	6:g:270:MET:CA	2.99	0.41
7:N:25:GLY:O	7:O:27:GLY:HA3	2.21	0.41
7:N:50:ARG:HG2	7:N:54:LEU:HD13	2.03	0.41
7:O:42:GLN:NE2	7:O:44:GLU:HB3	2.32	0.41
7:P:62:ALA:HA	7:P:65:ILE:HG22	2.03	0.41
7:R:65:ILE:HD12	7:R:68:LEU:HD23	2.03	0.41
7:M:31:ALA:HA	7:L:33:GLY:CA	2.51	0.41
7:M:52:THR:HG23	7:L:53:LEU:HD23	2.02	0.41
8:E:230:ASP:HB3	9:D:311:GLU:CB	2.51	0.41
8:E:433:GLU:HA	8:E:436:MET:HE3	2.02	0.41
9:F:163:ARG:CG	9:F:374:MET:HG3	2.51	0.41
8:A:151:LEU:N	8:A:151:LEU:HD12	2.36	0.41
8:C:439:TYR:OH	8:C:487:GLU:OE2	2.36	0.41
9:D:61:VAL:HG21	9:D:85:LEU:HD11	2.03	0.41
9:D:267:ASP:HA	9:D:320:SER:O	2.21	0.41
1:a:43:SER:OG	4:p:93:ILE:CD1	2.68	0.41
1:a:69:GLN:NE2	4:p:112:LEU:HD22	2.36	0.41
1:a:117:TRP:C	1:a:120:ILE:HG22	2.44	0.41
1:a:124:HIS:HD2	4:p:80:LYS:C	2.29	0.41
2:b:85:LEU:HD13	4:p:143:GLN:HB3	1.99	0.41
3:d:77:ARG:HG2	8:C:8:GLU:O	2.20	0.41
3:d:119:VAL:HG11	8:E:43:HIS:CD2	2.56	0.41
3:d:129:LEU:CD2	3:d:147:ASN:OD1	2.69	0.41
3:d:222:THR:HG21	3:d:234:ASP:CG	2.42	0.41
3:d:241:LEU:HD23	3:d:244:ILE:HD12	2.02	0.41
5:e:110:ARG:HB3	6:g:204:SER:OG	2.21	0.41
6:g:71:LYS:HB3	6:g:316:LEU:CD1	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:g:92:VAL:HG12	6:g:295:LEU:HD11	2.03	0.41
6:g:206:GLU:HG3	6:g:207:VAL:HG23	2.02	0.41
6:g:360:ALA:HB1	9:D:292:MET:CG	2.48	0.41
7:O:44:GLU:O	7:O:44:GLU:CD	2.63	0.41
7:Q:50:ARG:HG2	7:Q:54:LEU:HD13	2.03	0.41
7:M:50:ARG:HG2	7:M:54:LEU:HD13	2.03	0.41
7:M:55:LEU:HD21	7:L:54:LEU:HG	2.03	0.41
7:G:25:GLY:O	7:H:27:GLY:HA3	2.20	0.41
7:G:50:ARG:HG2	7:G:54:LEU:HD13	2.03	0.41
7:G:62:ALA:HA	7:G:65:ILE:HG22	2.03	0.41
7:G:81:VAL:HG21	7:H:80:PHE:CZ	2.56	0.41
7:L:65:ILE:HD12	7:L:68:LEU:HD23	2.03	0.41
7:K:32:ALA:O	7:K:36:VAL:HG23	2.21	0.41
9:F:415:GLU:O	9:F:419:LEU:HG	2.21	0.41
8:A:436:MET:HG3	8:A:490:LEU:HD13	1.99	0.41
8:A:439:TYR:CD2	8:A:490:LEU:HD23	2.56	0.41
8:A:460:GLU:OE1	8:A:460:GLU:HA	2.21	0.41
9:B:142:LEU:HD21	9:B:314:THR:HG21	2.02	0.41
9:B:231:LYS:O	9:B:232:VAL:HG23	2.21	0.41
1:a:70:ASN:HB3	2:b:53:LEU:C	2.45	0.41
2:b:96:ARG:HB2	4:p:154:GLU:HB2	2.03	0.41
3:d:132:ILE:HG12	8:E:13:ILE:HG21	2.03	0.41
5:e:48:ILE:HA	5:e:61:ALA:HA	2.02	0.41
6:g:123:VAL:CG2	6:g:308:LEU:CD2	2.98	0.41
7:N:40:ALA:CB	7:O:49:ILE:HD11	2.35	0.41
7:N:64:THR:HG22	7:O:66:TYR:HD2	1.86	0.41
7:O:10:VAL:HG21	7:P:74:LEU:HD22	1.95	0.41
7:O:62:ALA:HA	7:O:65:ILE:HG22	2.03	0.41
7:P:65:ILE:HD12	7:P:68:LEU:HD23	2.03	0.41
7:T:50:ARG:HG2	7:T:54:LEU:HD13	2.03	0.41
7:K:39:ILE:CG1	7:K:49:ILE:HG21	2.50	0.41
7:J:32:ALA:O	7:J:36:VAL:HG23	2.21	0.41
9:B:149:PHE:CZ	9:B:189:ILE:CD1	2.92	0.41
9:B:246:ARG:NH2	9:B:284:GLU:OE1	2.53	0.41
9:B:292:MET:HE2	8:C:276:LEU:HD23	2.03	0.41
8:C:178:ALA:O	8:C:182:ASP:CG	2.63	0.41
8:C:432:GLU:H	8:C:432:GLU:CD	2.28	0.41
9:D:79:MET:HE1	9:D:248:ARG:HG3	2.02	0.41
1:a:136:THR:H	4:p:90:LEU:CD2	2.29	0.40
2:b:87:LYS:O	2:b:90:MET:N	2.53	0.40
3:d:104:ARG:NH1	8:E:7:ASP:HB3	2.36	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:d:119:VAL:HG23	8:E:24:VAL:HG11	2.02	0.40
3:d:199:GLN:HE21	3:d:207:VAL:N	2.19	0.40
6:g:227:ILE:CG2	6:g:242:ILE:HD11	2.51	0.40
7:N:54:LEU:CG	7:O:55:LEU:HD21	2.51	0.40
7:T:54:LEU:HD11	7:G:55:LEU:HD21	1.99	0.40
7:H:10:VAL:HG22	7:I:74:LEU:HD22	1.97	0.40
7:H:50:ARG:HG2	7:H:54:LEU:HD13	2.03	0.40
7:K:50:ARG:HG2	7:K:54:LEU:HD13	2.03	0.40
7:J:62:ALA:HA	7:J:65:ILE:HG22	2.03	0.40
7:J:66:TYR:HD2	7:I:64:THR:HG22	1.86	0.40
8:E:322:THR:HG22	8:E:332:PRO:CG	2.51	0.40
8:E:403:LEU:HD13	8:E:407:THR:HG21	2.03	0.40
9:F:48:ILE:HD12	9:F:94:THR:HG22	2.02	0.40
9:F:477:VAL:HG21	9:F:483:ALA:HB2	2.03	0.40
8:A:104:TYR:CE2	8:A:109:ILE:CD1	2.99	0.40
8:A:176:LYS:HD2	8:A:319:ILE:CG2	2.51	0.40
8:A:227:GLU:CG	8:A:239:ALA:HB2	2.52	0.40
8:A:416:ARG:NE	8:A:447:ASP:O	2.54	0.40
9:D:246:ARG:NH2	9:D:284:GLU:OE1	2.54	0.40
9:D:427:ILE:HD11	9:D:462:LEU:CD2	2.49	0.40
1:a:122:LEU:HD13	1:a:125:GLY:O	2.22	0.40
3:d:201:ILE:HG13	3:d:202:THR:H	1.86	0.40
3:d:222:THR:CB	3:d:234:ASP:CB	2.81	0.40
7:S:42:GLN:HE22	7:R:43:PRO:CG	2.22	0.40
7:S:62:ALA:HA	7:S:65:ILE:HG22	2.03	0.40
7:N:65:ILE:HD12	7:N:68:LEU:HD23	2.03	0.40
7:R:32:ALA:O	7:R:36:VAL:HG23	2.21	0.40
7:M:51:GLY:C	7:L:50:ARG:HH21	2.28	0.40
7:T:65:ILE:HD12	7:T:68:LEU:HD23	2.03	0.40
7:H:21:SER:OG	7:I:20:ALA:HA	2.21	0.40
7:J:65:ILE:HD12	7:J:68:LEU:HD23	2.03	0.40
9:F:247:MET:SD	9:F:282:GLY:HA2	2.60	0.40
8:A:194:CYS:O	8:A:222:THR:HA	2.21	0.40
8:C:22:ARG:CG	8:C:23:GLU:N	2.84	0.40
1:a:142:LEU:HD23	1:a:142:LEU:HA	1.90	0.40
1:a:212:VAL:N	1:a:213:PRO:HD2	2.36	0.40
2:b:103:ILE:HG13	4:p:163:LYS:CA	2.51	0.40
2:b:169:ILE:HG23	3:d:221:PHE:CD2	2.49	0.40
5:e:4:ASN:HB3	5:e:18:GLU:CD	2.44	0.40
6:g:85:PHE:CE1	6:g:89:LEU:HD22	2.57	0.40
7:S:54:LEU:HD11	7:T:55:LEU:HD21	2.00	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:P:32:ALA:O	7:P:36:VAL:HG23	2.21	0.40
7:Q:32:ALA:O	7:Q:36:VAL:HG23	2.21	0.40
7:G:32:ALA:O	7:G:36:VAL:HG23	2.21	0.40
7:H:3:PRO:HA	7:I:5:ILE:HD11	2.03	0.40
7:K:80:PHE:CE2	7:J:81:VAL:CG2	3.04	0.40
7:I:62:ALA:HA	7:I:65:ILE:HG22	2.03	0.40
8:E:137:ILE:HG23	8:E:138:MET:N	2.37	0.40
8:A:417:LEU:HD23	8:A:417:LEU:HA	1.79	0.40
8:A:432:GLU:CA	8:A:435:VAL:HG12	2.51	0.40
8:C:65:LEU:HD23	8:C:65:LEU:HA	1.94	0.40
1:a:164:TYR:CE2	1:a:166:GLN:CG	2.79	0.40
2:b:71:ARG:O	4:p:133:THR:CB	2.68	0.40
2:b:71:ARG:HG3	4:p:132:ASP:HB3	2.03	0.40
2:b:97:VAL:H	4:p:155:ILE:HD11	1.30	0.40
2:b:149:LEU:CD2	3:d:244:ILE:HG13	2.52	0.40
3:d:180:VAL:HG13	3:d:223:ILE:CG2	2.49	0.40
3:d:222:THR:HG21	3:d:234:ASP:HB3	1.74	0.40
4:p:90:LEU:CB	4:p:91:PRO:CD	2.96	0.40
4:p:205:SER:O	4:p:206:GLN:C	2.64	0.40
7:S:74:LEU:CD2	7:R:10:VAL:HG22	2.45	0.40
7:N:62:ALA:HA	7:N:65:ILE:HG22	2.03	0.40
7:H:32:ALA:O	7:H:36:VAL:HG23	2.21	0.40
7:H:68:LEU:HD22	7:I:66:TYR:CG	2.57	0.40
7:L:32:ALA:O	7:L:36:VAL:HG23	2.21	0.40
7:I:65:ILE:HD12	7:I:68:LEU:HD23	2.03	0.40
8:A:250:PHE:HB3	8:A:255:ARG:HB2	2.04	0.40
8:C:403:LEU:C	8:C:403:LEU:HD12	2.47	0.40
9:D:103:GLY:O	9:D:106:THR:OG1	2.32	0.40
3:d:74:THR:CG2	8:C:4:ILE:N	2.85	0.40
6:g:227:ILE:HG21	6:g:242:ILE:HD11	2.03	0.40
7:S:35:ALA:HB2	7:S:53:LEU:HD13	2.04	0.40
7:S:39:ILE:CG1	7:S:49:ILE:HG21	2.50	0.40
7:S:50:ARG:HG2	7:S:54:LEU:HD13	2.03	0.40
7:G:35:ALA:HB2	7:G:53:LEU:HD13	2.04	0.40
8:E:79:ASP:HB3	8:E:81:LEU:HG	2.03	0.40
8:E:469:LYS:CE	8:E:493:ALA:CB	3.00	0.40
9:F:159:LEU:HA	9:F:159:LEU:HD23	1.91	0.40
9:F:321:ILE:HD13	9:F:321:ILE:HG21	1.83	0.40
9:F:472:GLN:O	9:F:490:LEU:HD11	2.22	0.40
8:A:364:VAL:O	8:A:364:VAL:HG23	2.21	0.40
9:B:216:MET:HE3	9:B:221:VAL:CG1	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:D:401:LEU:O	9:D:404:ILE:CG1	2.70	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	221/247 (90%)	211 (96%)	10 (4%)	0	100	100
2	b	157/184 (85%)	151 (96%)	6 (4%)	0	100	100
3	d	177/257 (69%)	170 (96%)	6 (3%)	1 (1%)	22	59
4	p	141/222 (64%)	130 (92%)	11 (8%)	0	100	100
5	e	129/134 (96%)	122 (95%)	7 (5%)	0	100	100
6	g	319/364 (88%)	306 (96%)	13 (4%)	0	100	100
7	G	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	H	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	I	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	J	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	K	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	L	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	M	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	N	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	O	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	P	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	Q	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	R	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
7	S	77/81 (95%)	74 (96%)	3 (4%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
7	T	77/81 (95%)	74 (96%)	3 (4%)	0	100	100
8	A	499/507 (98%)	485 (97%)	14 (3%)	0	100	100
8	C	499/507 (98%)	478 (96%)	21 (4%)	0	100	100
8	E	496/507 (98%)	483 (97%)	13 (3%)	0	100	100
9	B	476/498 (96%)	444 (93%)	31 (6%)	1 (0%)	44	78
9	D	478/498 (96%)	463 (97%)	14 (3%)	1 (0%)	44	78
9	F	477/498 (96%)	463 (97%)	13 (3%)	1 (0%)	44	78
All	All	5147/5557 (93%)	4942 (96%)	201 (4%)	4 (0%)	50	83

All (4) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
3	d	217	LEU
9	B	296	VAL
9	F	296	VAL
9	D	296	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	187/207 (90%)	187 (100%)	0	100	100
2	b	142/161 (88%)	142 (100%)	0	100	100
3	d	156/220 (71%)	156 (100%)	0	100	100
4	p	124/197 (63%)	124 (100%)	0	100	100
5	e	109/112 (97%)	109 (100%)	0	100	100
6	g	276/315 (88%)	275 (100%)	1 (0%)	89	91
7	G	51/53 (96%)	51 (100%)	0	100	100
7	H	51/53 (96%)	51 (100%)	0	100	100
7	I	51/53 (96%)	51 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
7	J	51/53 (96%)	51 (100%)	0	100	100
7	K	51/53 (96%)	51 (100%)	0	100	100
7	L	51/53 (96%)	51 (100%)	0	100	100
7	M	51/53 (96%)	51 (100%)	0	100	100
7	N	51/53 (96%)	51 (100%)	0	100	100
7	O	51/53 (96%)	51 (100%)	0	100	100
7	P	51/53 (96%)	51 (100%)	0	100	100
7	Q	51/53 (96%)	51 (100%)	0	100	100
7	R	51/53 (96%)	51 (100%)	0	100	100
7	S	51/53 (96%)	51 (100%)	0	100	100
7	T	51/53 (96%)	51 (100%)	0	100	100
8	A	410/414 (99%)	410 (100%)	0	100	100
8	C	410/414 (99%)	410 (100%)	0	100	100
8	E	408/414 (99%)	407 (100%)	1 (0%)	92	93
9	B	389/408 (95%)	389 (100%)	0	100	100
9	D	391/408 (96%)	390 (100%)	1 (0%)	91	92
9	F	390/408 (96%)	387 (99%)	3 (1%)	79	85
All	All	4106/4420 (93%)	4100 (100%)	6 (0%)	92	95

All (6) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
6	g	292	LEU
8	E	298	LEU
9	F	222	ILE
9	F	374	MET
9	F	433	GLN
9	D	279	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (72) such sidechains are listed below:

Mol	Chain	Res	Type
1	a	34	GLN
1	a	70	ASN
1	a	86	GLN

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Mol	Chain	Res	Type
1	a	193	ASN
2	b	33	ASN
2	b	133	GLN
2	b	161	ASN
3	d	141	HIS
3	d	188	ASN
3	d	190	HIS
3	d	199	GLN
3	d	240	GLN
4	p	167	GLN
4	p	195	GLN
5	e	27	ASN
6	g	56	ASN
6	g	81	ASN
6	g	228	HIS
6	g	301	ASN
6	g	303	GLN
7	S	42	GLN
7	N	42	GLN
7	O	42	GLN
7	P	42	GLN
7	Q	42	GLN
7	R	42	GLN
7	M	42	GLN
7	T	42	GLN
7	G	42	GLN
7	H	42	GLN
7	L	42	GLN
7	K	42	GLN
7	J	42	GLN
7	I	42	GLN
8	E	34	GLN
8	E	43	HIS
8	E	148	GLN
8	E	208	GLN
9	F	25	GLN
9	F	120	ASN
9	F	264	ASN
9	F	280	GLN
9	F	310	GLN
9	F	384	HIS
8	A	28	ASN

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Mol	Chain	Res	Type
8	A	34	GLN
8	A	66	ASN
8	A	148	GLN
8	A	186	ASN
8	A	187	GLN
8	A	188	GLN
8	A	201	GLN
8	A	214	GLN
8	A	256	HIS
8	A	409	ASN
8	A	423	GLN
8	A	497	GLN
9	B	71	ASN
9	B	112	ASN
9	B	210	ASN
9	B	238	GLN
9	B	345	HIS
9	B	389	GLN
8	C	187	GLN
8	C	267	GLN
8	C	408	GLN
8	C	409	ASN
8	C	423	GLN
9	D	60	ASN
9	D	67	GLN
9	D	72	ASN
9	D	389	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 ⓘ

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry

Of 10 ligands modelled in this entry, 5 are monoatomic - leaving 5 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. 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  $|Z| > 2$  is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z  > 2	Counts	RMSZ	# Z  > 2
10	ATP	C	601	11	26,33,33	0.95	1 (3%)	31,52,52	1.52	6 (19%)
12	ADP	D	600	11	24,29,29	1.32	3 (12%)	29,45,45	1.48	4 (13%)
12	ADP	B	501	11	24,29,29	1.08	1 (4%)	29,45,45	1.48	4 (13%)
10	ATP	A	600	11	26,33,33	1.07	1 (3%)	31,52,52	1.36	4 (12%)
10	ATP	E	600	11	26,33,33	1.19	2 (7%)	31,52,52	1.56	8 (25%)

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
10	ATP	C	601	11	-	0/18/38/38	0/3/3/3
12	ADP	D	600	11	-	5/12/32/32	0/3/3/3
12	ADP	B	501	11	-	5/12/32/32	0/3/3/3
10	ATP	A	600	11	-	3/18/38/38	0/3/3/3
10	ATP	E	600	11	-	4/18/38/38	0/3/3/3

All (8) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
12	D	600	ADP	C2'-C1'	-2.72	1.49	1.53
10	E	600	ATP	C2'-C1'	-2.61	1.49	1.53
12	B	501	ADP	C2'-C1'	-2.35	1.50	1.53
12	D	600	ADP	PB-O3B	-2.23	1.46	1.54
10	C	601	ATP	C2'-C1'	-2.13	1.50	1.53
12	D	600	ADP	O4'-C4'	-2.06	1.40	1.45
10	A	600	ATP	C5-C4	2.06	1.46	1.40
10	E	600	ATP	PG-O2G	-2.04	1.47	1.54

All (26) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	E	600	ATP	N3-C2-N1	-3.93	122.53	128.68
12	D	600	ADP	N3-C2-N1	-3.87	122.63	128.68
10	A	600	ATP	N3-C2-N1	-3.49	123.22	128.68
10	C	601	ATP	N3-C2-N1	-3.44	123.30	128.68
12	B	501	ADP	C3'-C2'-C1'	3.30	105.94	100.98
12	B	501	ADP	N3-C2-N1	-3.25	123.60	128.68
12	B	501	ADP	PA-O3A-PB	-3.24	121.69	132.83
10	E	600	ATP	C4-C5-N7	-3.18	106.08	109.40
10	C	601	ATP	C4-C5-N7	-2.66	106.62	109.40
10	A	600	ATP	O2B-PB-O1B	2.59	125.05	112.24
10	E	600	ATP	C2-N1-C6	2.53	123.09	118.75
10	C	601	ATP	C2-N1-C6	2.53	123.09	118.75
12	D	600	ADP	C4-C5-N7	-2.51	106.79	109.40
10	C	601	ATP	C3'-C2'-C1'	2.42	104.63	100.98
10	E	600	ATP	C3'-C2'-C1'	2.42	104.62	100.98
12	B	501	ADP	C4-C5-N7	-2.39	106.91	109.40
12	D	600	ADP	C2-N1-C6	2.33	122.75	118.75
12	D	600	ADP	O3B-PB-O2B	2.26	116.28	107.64
10	C	601	ATP	PB-O3B-PG	-2.25	125.12	132.83
10	A	600	ATP	C2-N1-C6	2.24	122.58	118.75
10	C	601	ATP	C1'-N9-C4	-2.19	122.79	126.64
10	E	600	ATP	O2B-PB-O1B	2.17	122.98	112.24
10	E	600	ATP	PB-O3B-PG	-2.11	125.59	132.83
10	E	600	ATP	C5'-C4'-C3'	-2.07	107.43	115.18
10	A	600	ATP	O4'-C1'-C2'	-2.04	103.95	106.93
10	E	600	ATP	O3B-PG-O1G	-2.00	100.09	111.19

There are no chirality outliers.

All (17) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
12	B	501	ADP	C5'-O5'-PA-O2A
12	D	600	ADP	C5'-O5'-PA-O1A
12	D	600	ADP	C5'-O5'-PA-O2A
12	B	501	ADP	O4'-C4'-C5'-O5'
12	B	501	ADP	C3'-C4'-C5'-O5'
10	A	600	ATP	PB-O3B-PG-O1G
10	A	600	ATP	PB-O3B-PG-O2G
12	B	501	ADP	C5'-O5'-PA-O3A
10	E	600	ATP	PG-O3B-PB-O2B
10	E	600	ATP	PA-O3A-PB-O2B

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Mol	Chain	Res	Type	Atoms
12	D	600	ADP	PB-O3A-PA-O2A
12	B	501	ADP	C5'-O5'-PA-O1A
12	D	600	ADP	O4'-C4'-C5'-O5'
10	E	600	ATP	PA-O3A-PB-O1B
10	A	600	ATP	PA-O3A-PB-O2B
12	D	600	ADP	C5'-O5'-PA-O3A
10	E	600	ATP	PG-O3B-PB-O1B

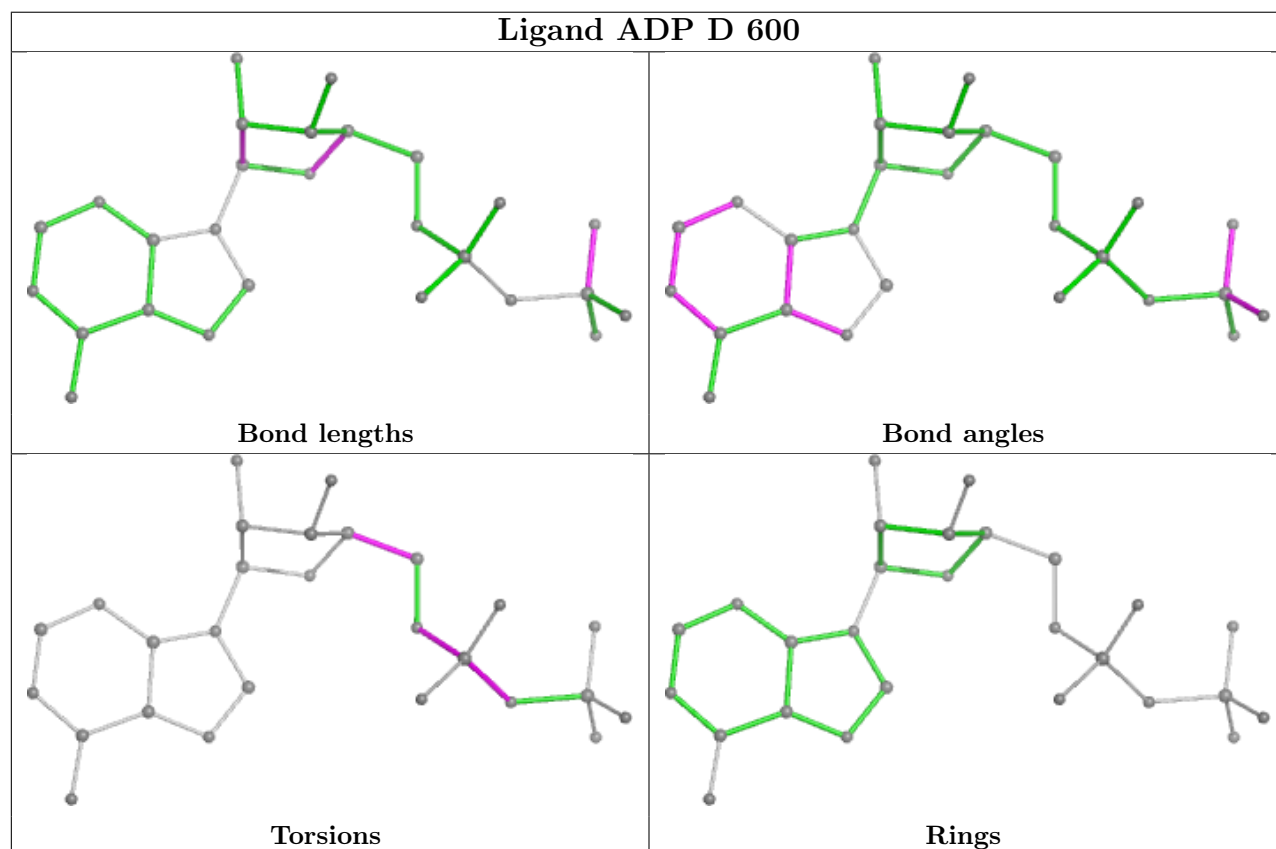
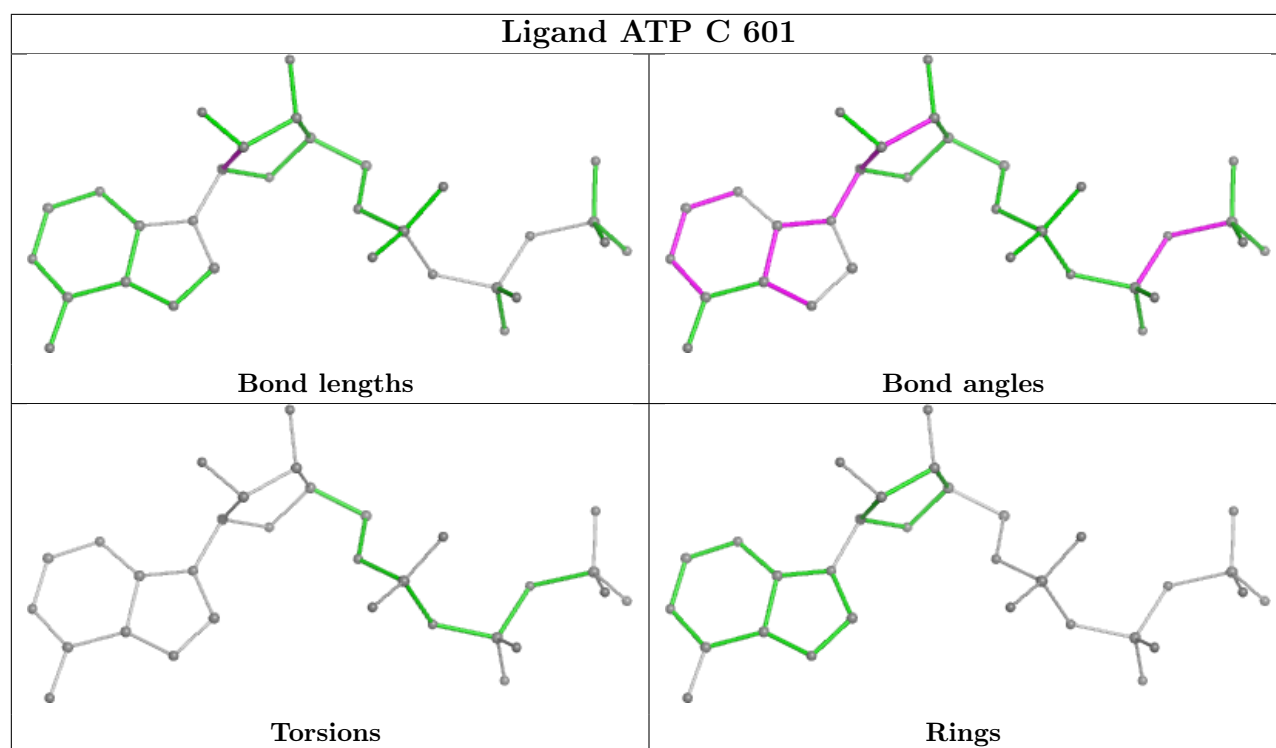
There are no ring outliers.

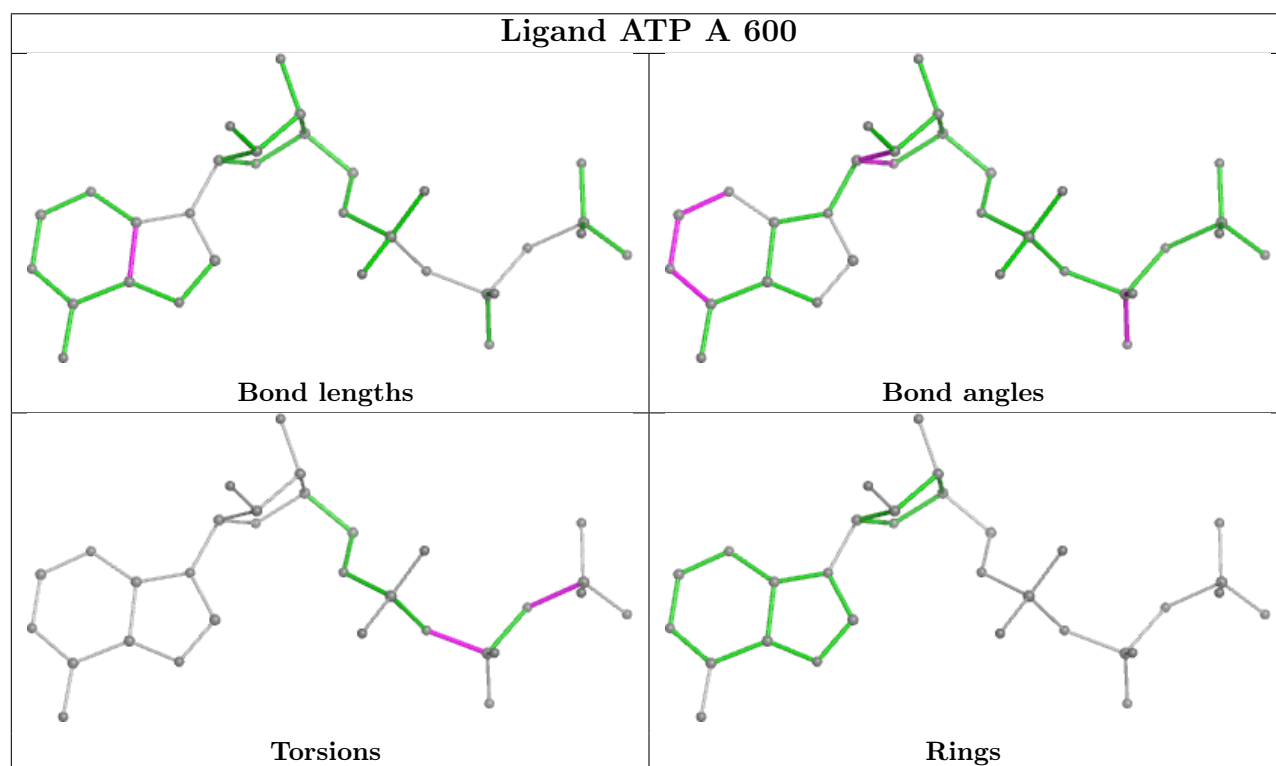
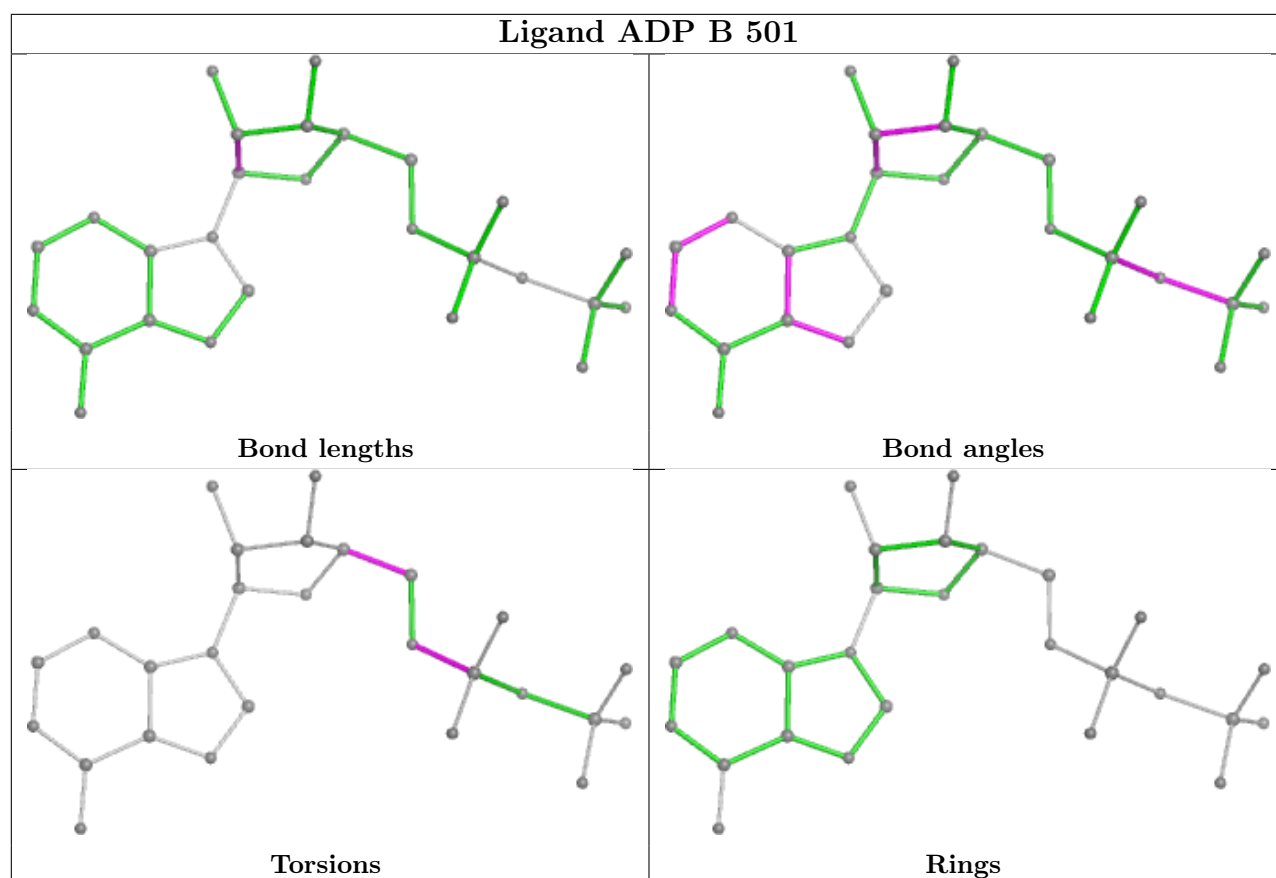
4 monomers are involved in 5 short contacts:

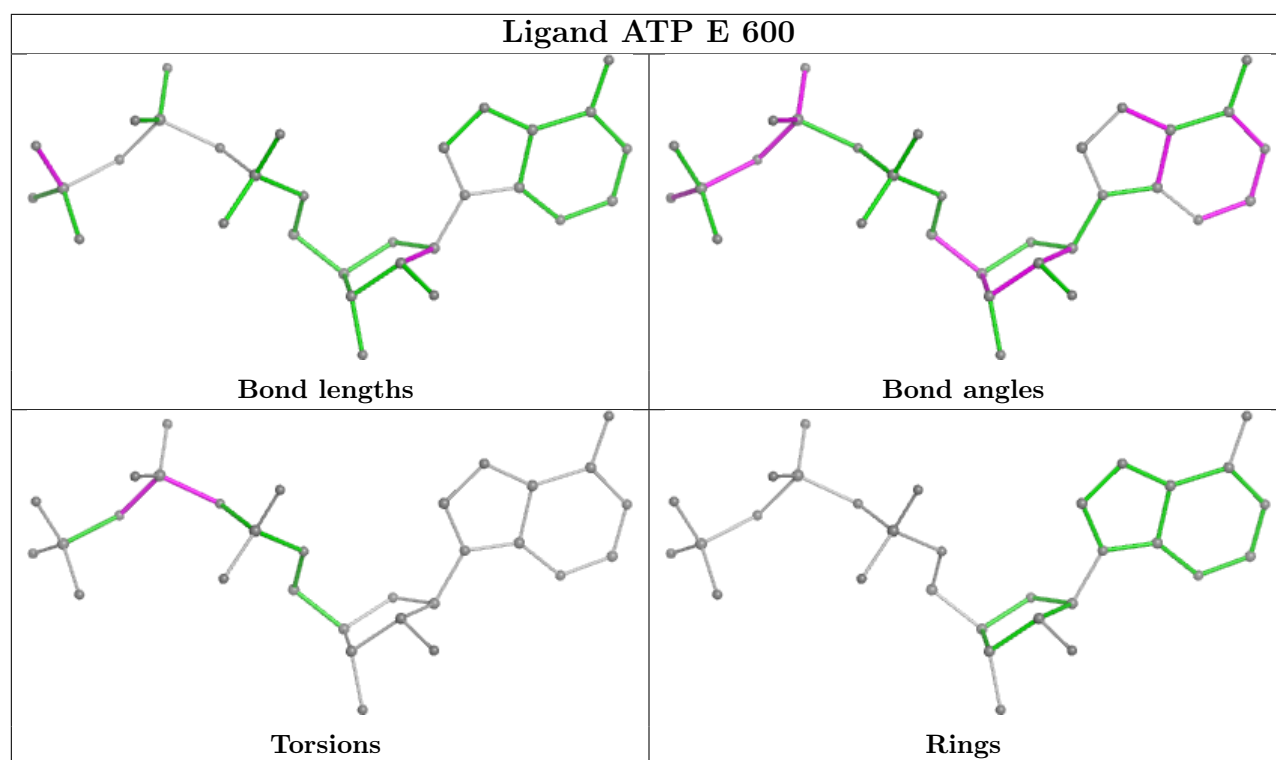
Mol	Chain	Res	Type	Clashes	Symm-Clashes
10	C	601	ATP	1	0
12	D	600	ADP	2	0
10	A	600	ATP	1	0
10	E	600	ATP	1	0

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.









## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

The following chains have linkage breaks:

Mol	Chain	Number of breaks
2	b	1
4	p	1
9	F	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	b	77:GLN	C	78:LEU	N	2.25
1	p	105:ASP	C	106:LYS	N	1.17
1	F	276:PHE	C	277:ARG	N	0.97

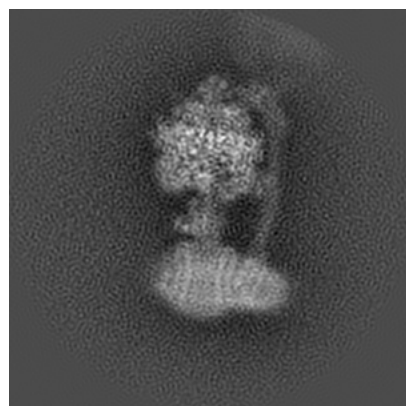
## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-4272. These allow visual inspection of the internal detail of the map and identification of artifacts.

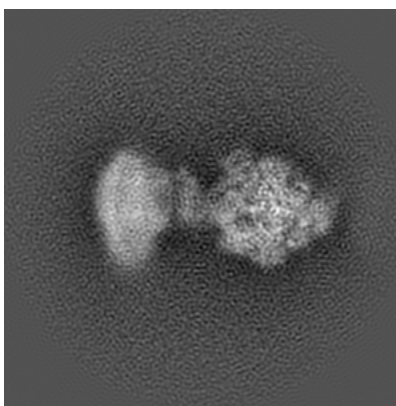
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

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

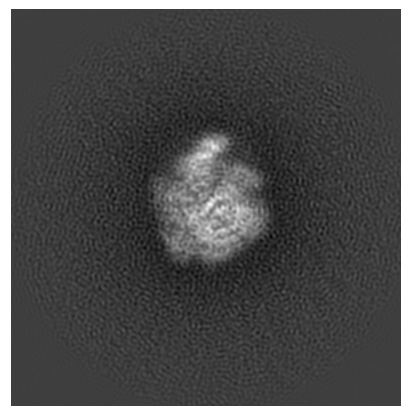
#### 6.1.1 Primary map



X

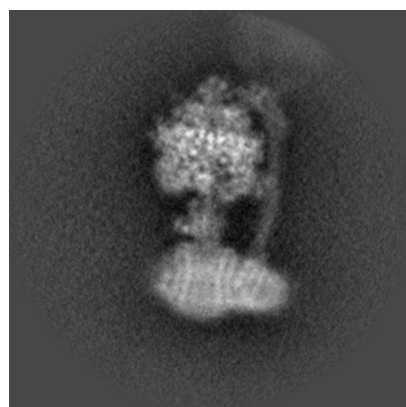


Y

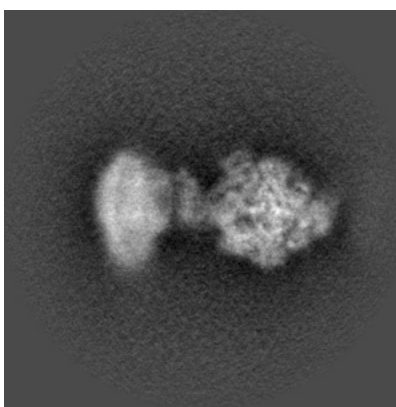


Z

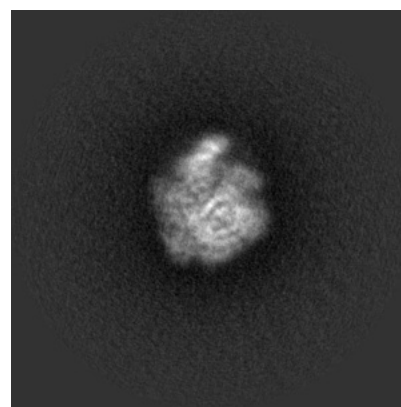
#### 6.1.2 Raw map



X



Y

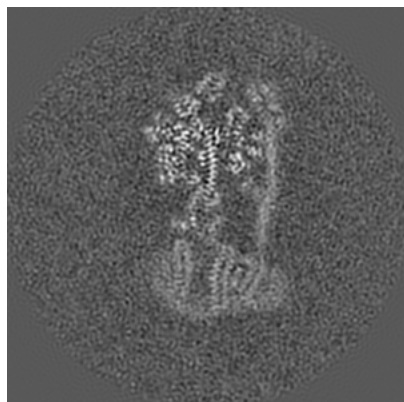


Z

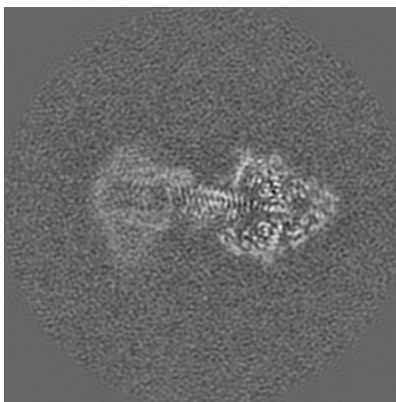
The images above show the map projected in three orthogonal directions.

## 6.2 Central slices [i](#)

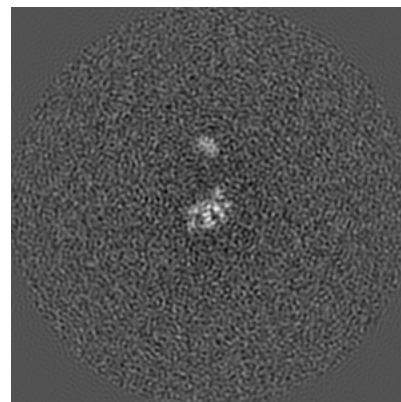
### 6.2.1 Primary map



X Index: 175

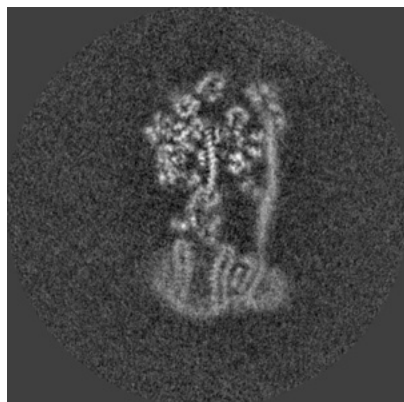


Y Index: 175

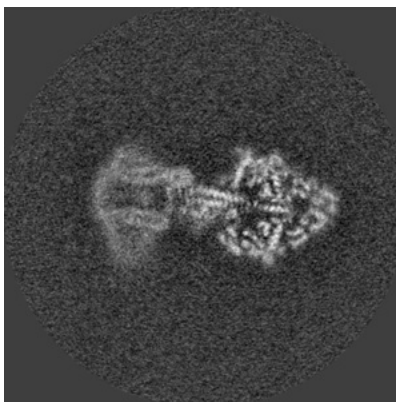


Z Index: 175

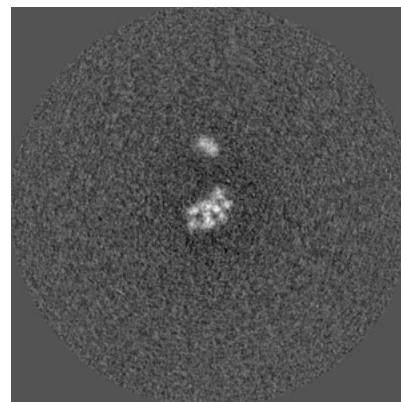
### 6.2.2 Raw map



X Index: 175



Y Index: 175

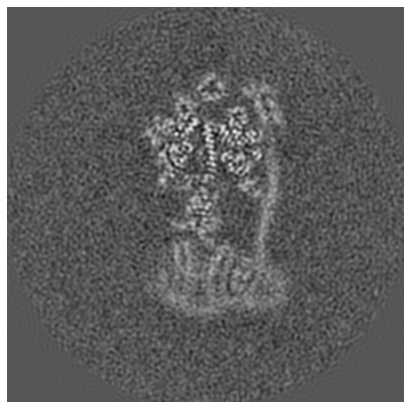


Z Index: 175

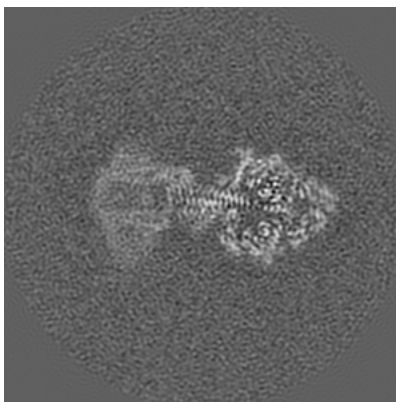
The images above show central slices of the map in three orthogonal directions.

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

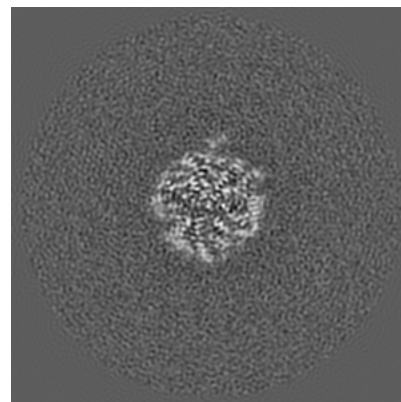
### 6.3.1 Primary map



X Index: 172

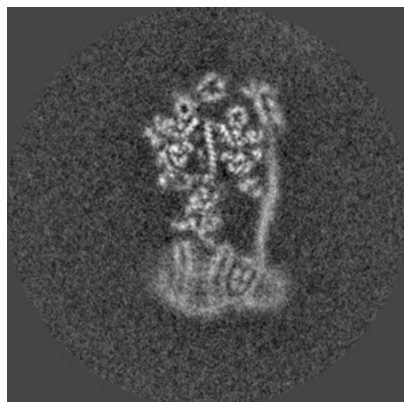


Y Index: 174

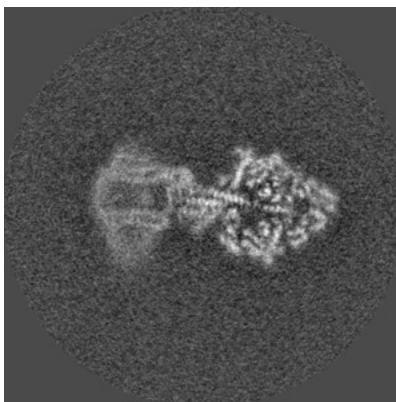


Z Index: 233

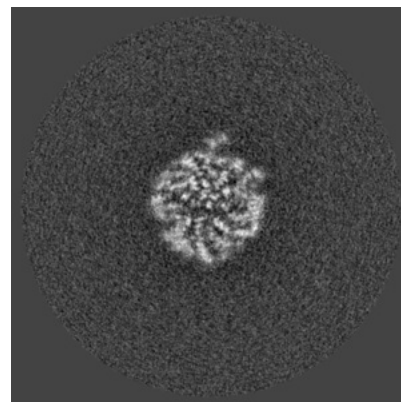
### 6.3.2 Raw map



X Index: 172



Y Index: 174



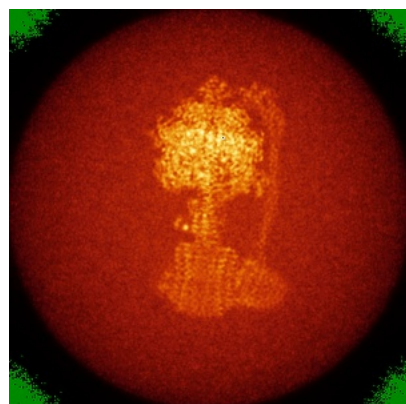
Z Index: 233

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

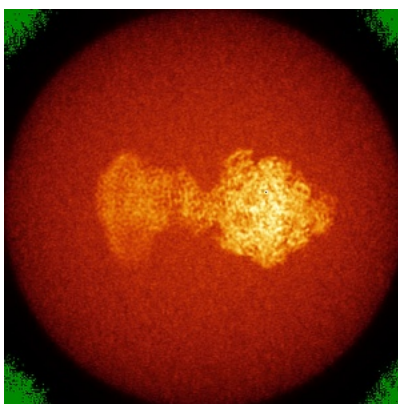


## 6.4 Orthogonal standard-deviation projections (False-color) [i](#)

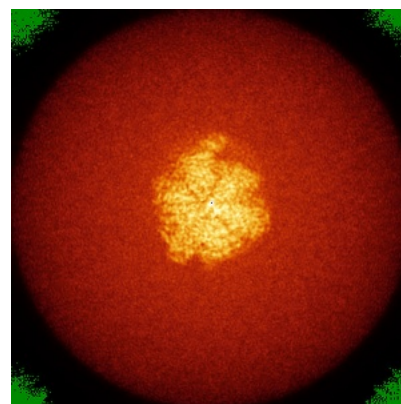
### 6.4.1 Primary map



X

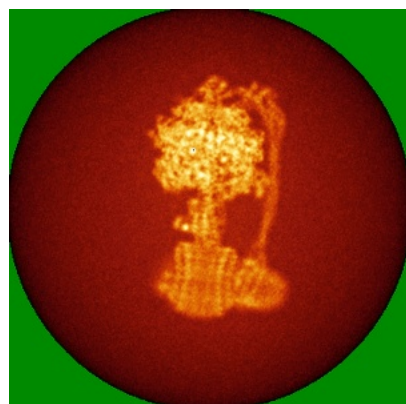


Y

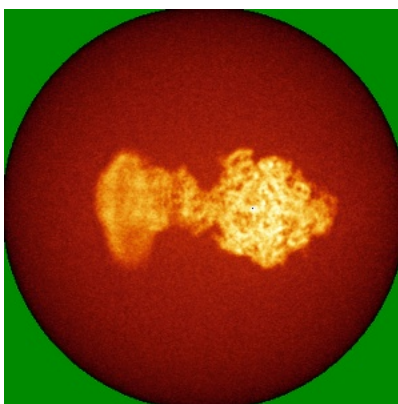


Z

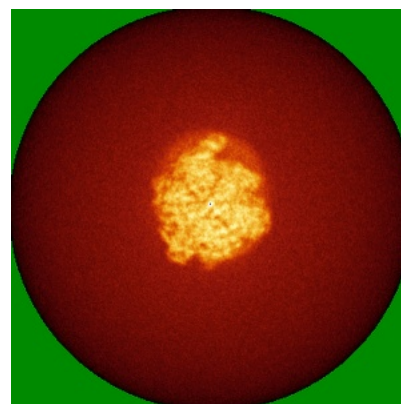
### 6.4.2 Raw map



X



Y



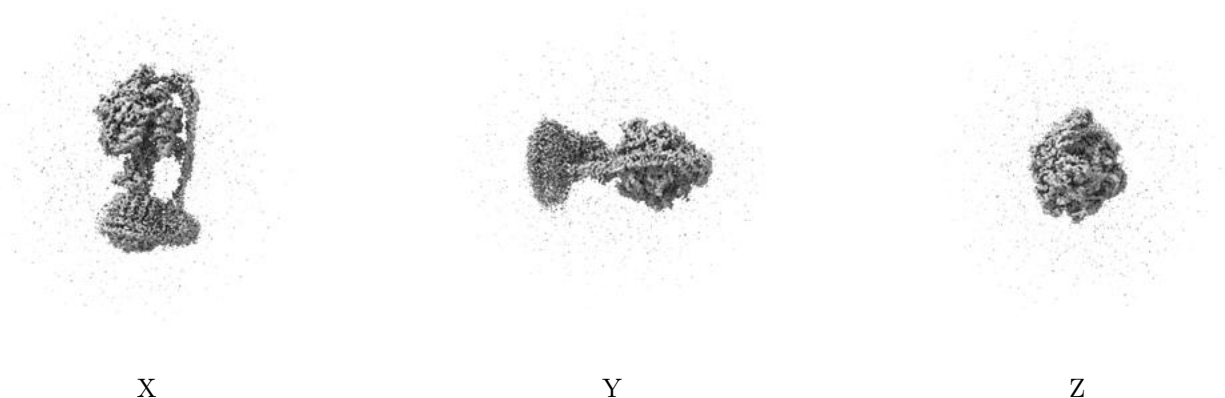
Z

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



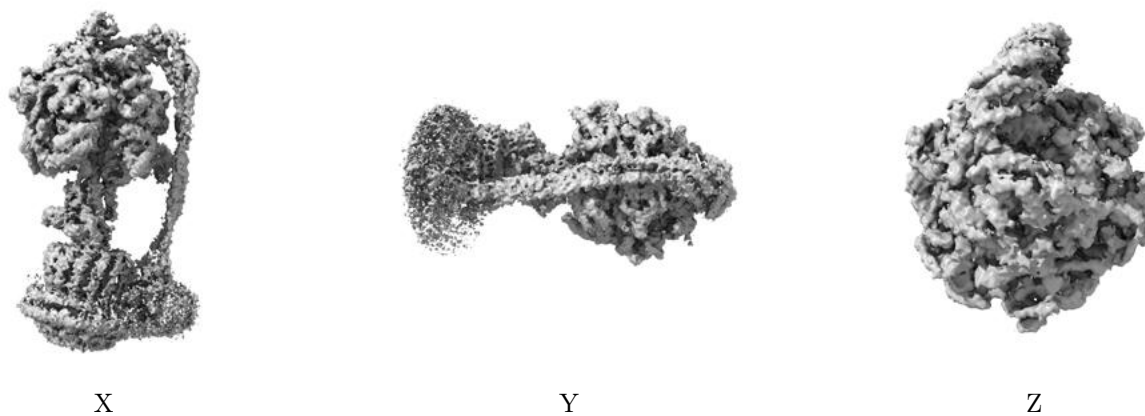
## 6.5 Orthogonal surface views [i](#)

### 6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.05. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

### 6.5.2 Raw map



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

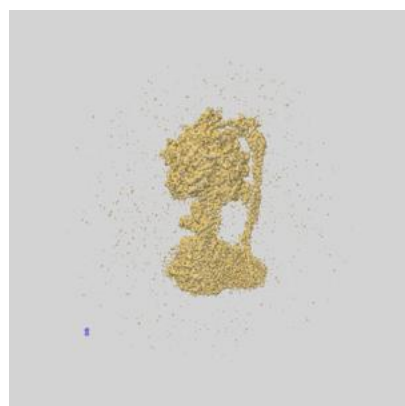
## 6.6 Mask visualisation [i](#)

This section shows the 3D surface view of the primary map at 50% transparency overlaid with the specified mask at 0% transparency

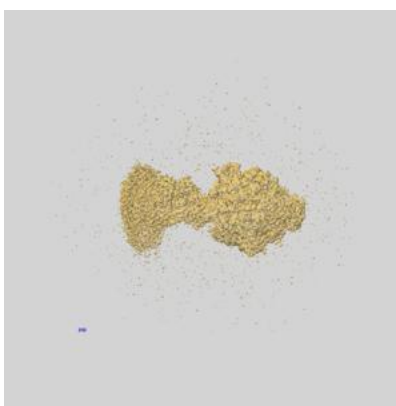
A mask typically either:

- Encompasses the whole structure
- Separates out a domain, a functional unit, a monomer or an area of interest from a larger structure

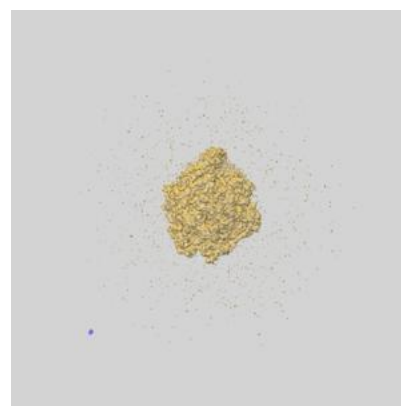
### 6.6.1 emd\_4272\_msk\_1.map [i](#)



X



Y

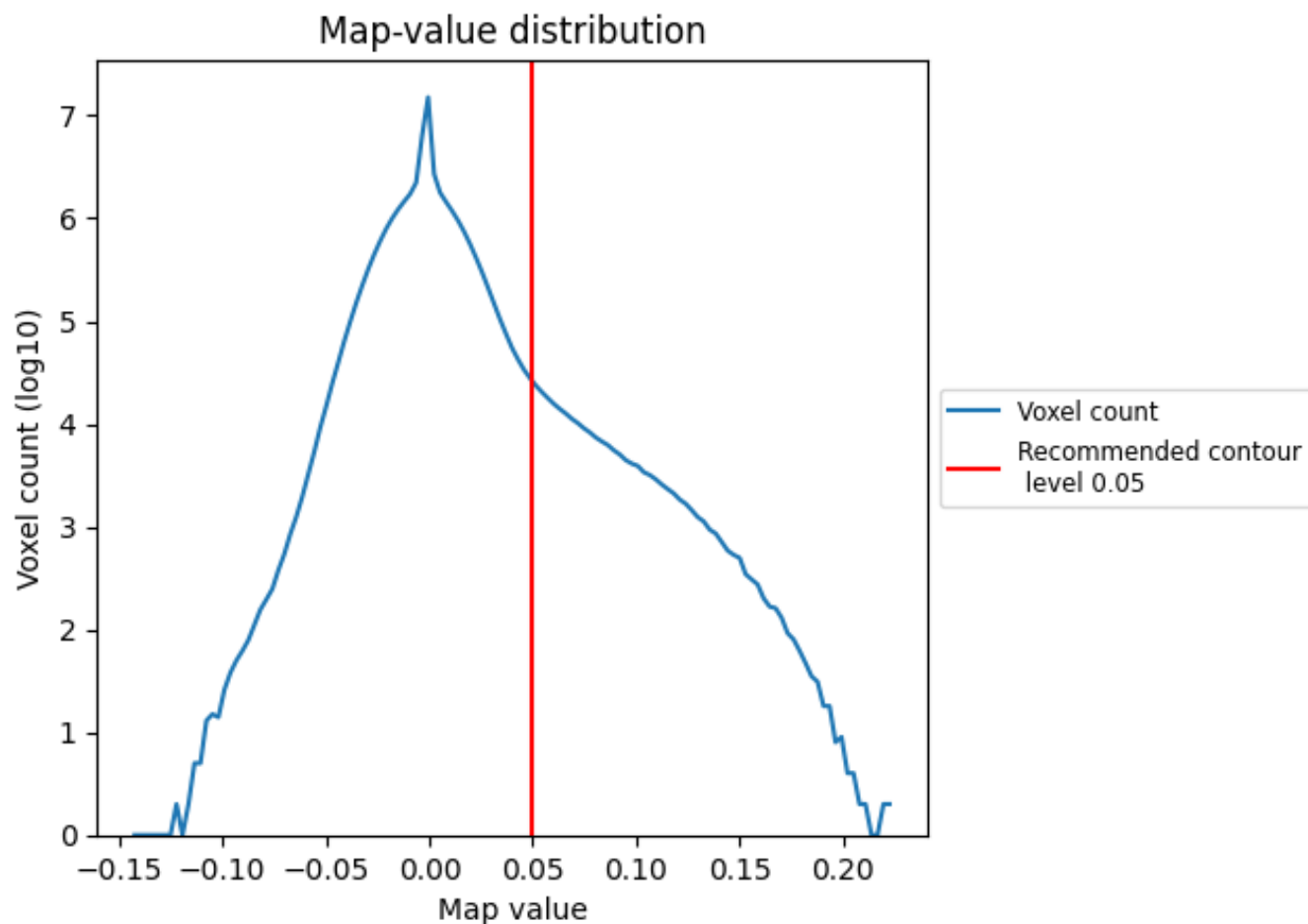


Z

## 7 Map analysis [i](#)

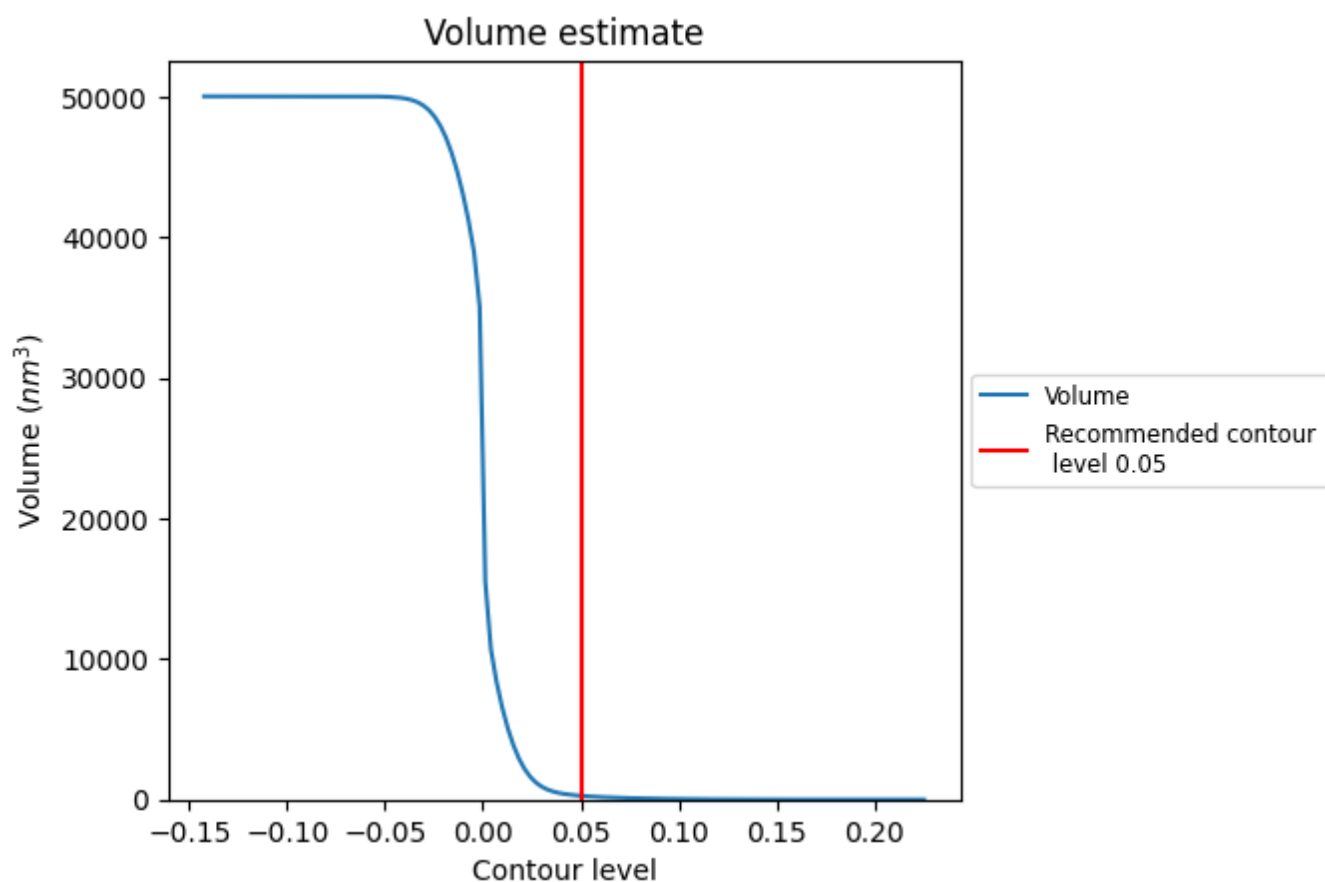
This section contains the results of statistical analysis of the map.

### 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.

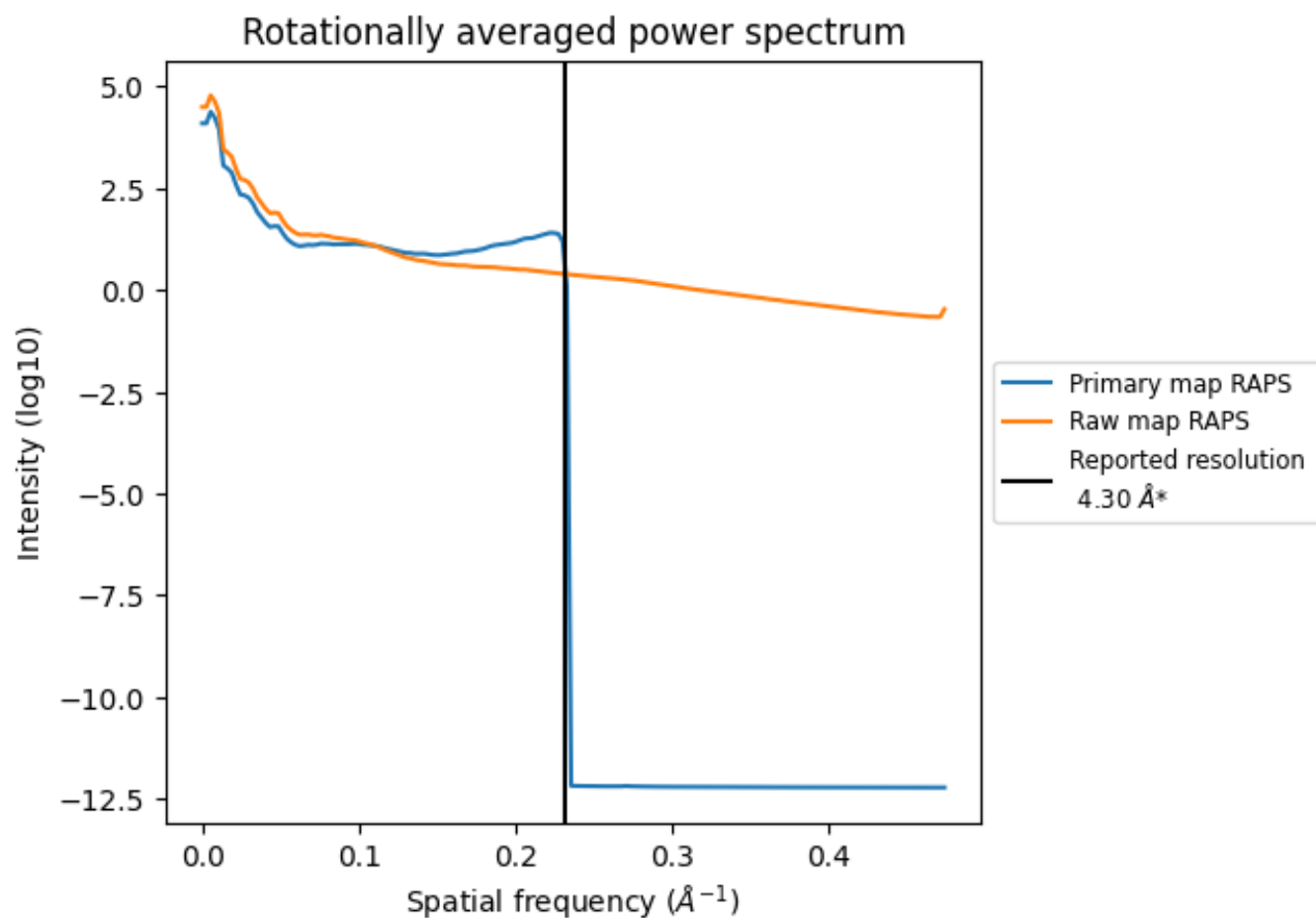
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 269 nm<sup>3</sup>; this corresponds to an approximate mass of 243 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 ⓘ

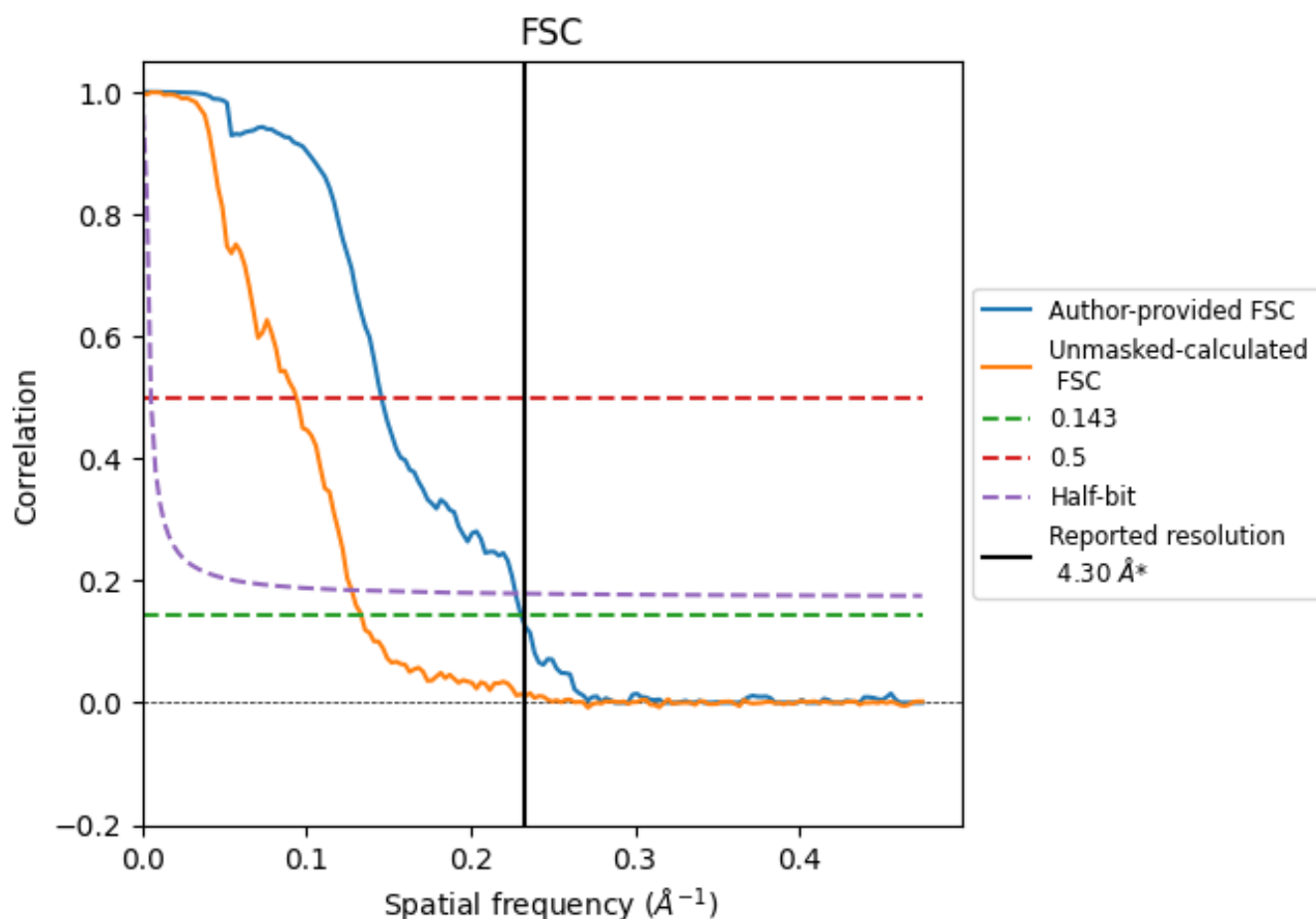


\*Reported resolution corresponds to spatial frequency of 0.233  $\text{\AA}^{-1}$

## 8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

### 8.1 FSC [i](#)



\*Reported resolution corresponds to spatial frequency of 0.233  $\text{\AA}^{-1}$

## 8.2 Resolution estimates [i](#)

Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	4.30	-	-
Author-provided FSC curve	4.33	6.87	4.40
Unmasked-calculated*	7.50	10.68	7.84

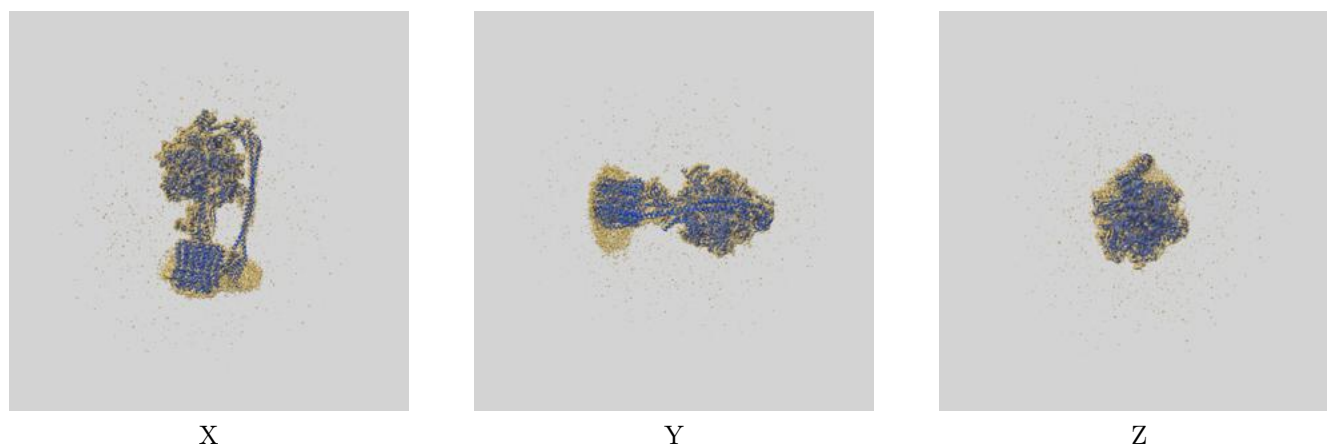
\*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 7.50 differs from the reported value 4.3 by more than 10 %



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

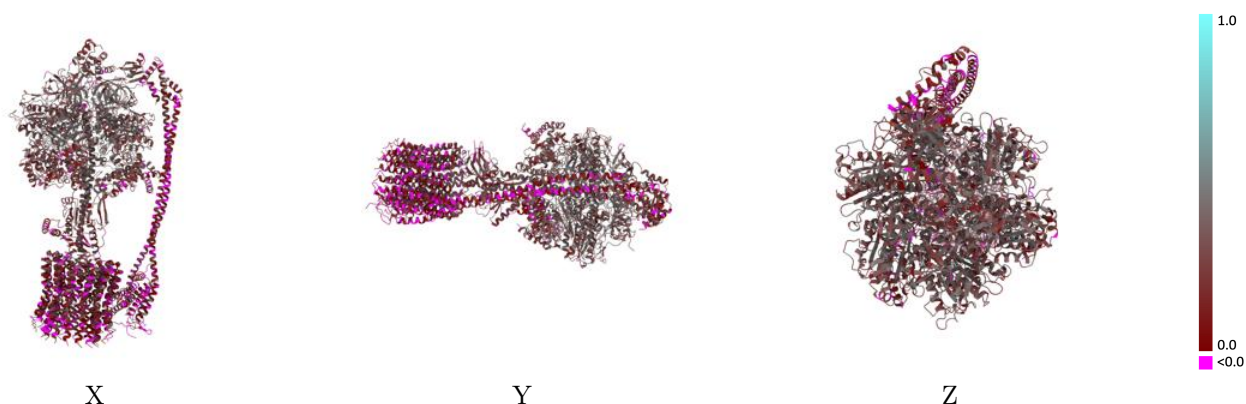
This section contains information regarding the fit between EMDB map EMD-4272 and PDB model 6FKI. Per-residue inclusion information can be found in section [3](#) on page [8](#).

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



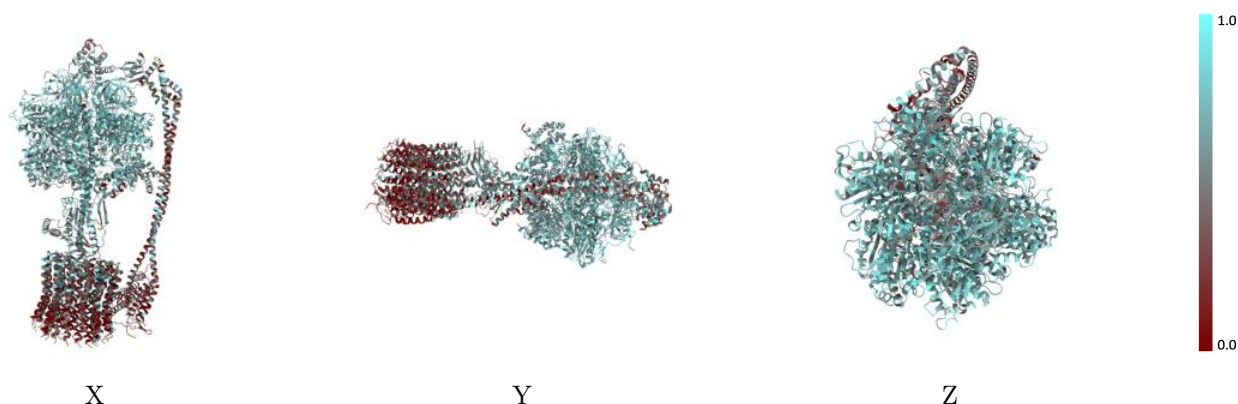
The images above show the 3D surface view of the map at the recommended contour level 0.05 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



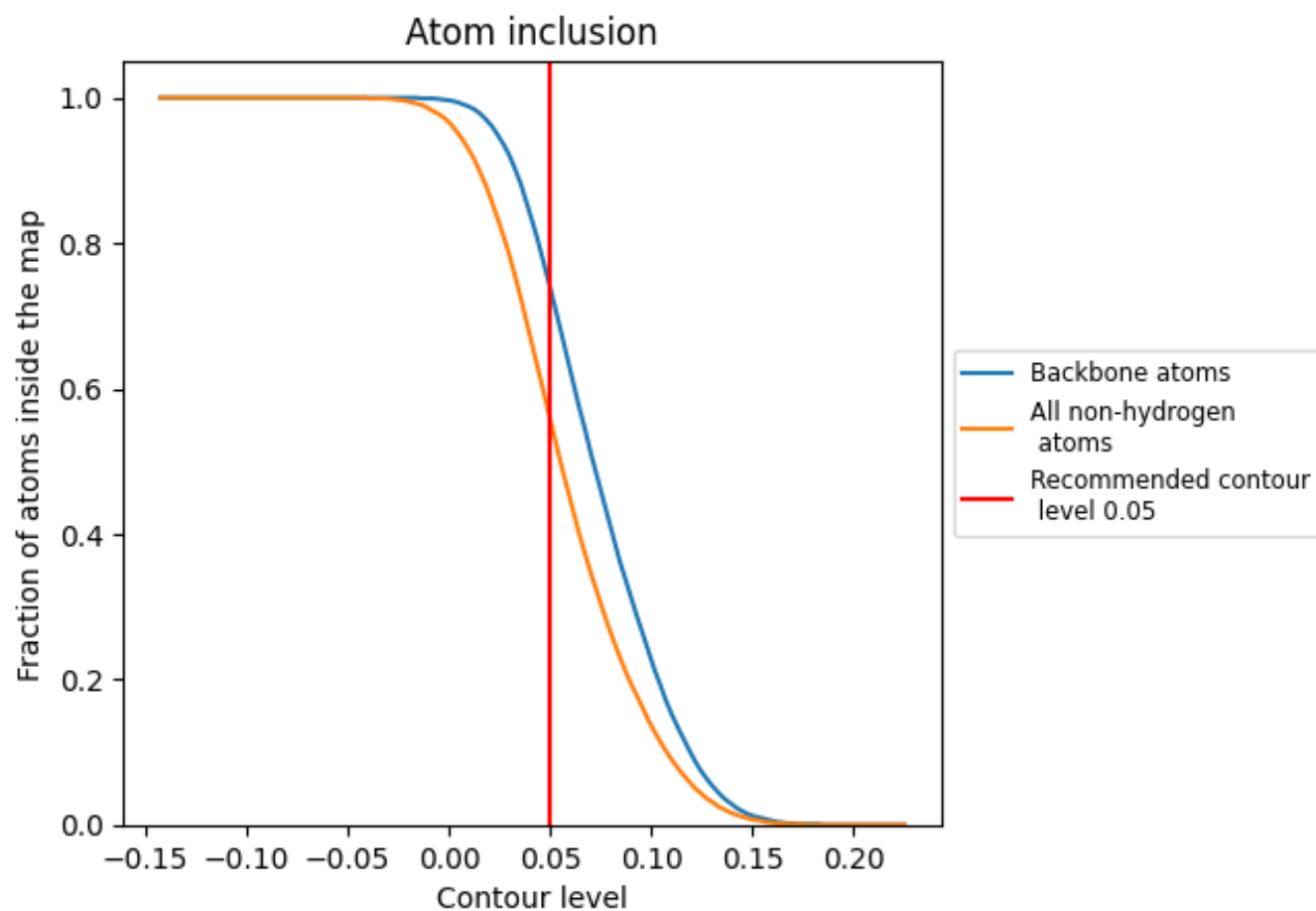
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.05).























































## 9.4 Atom inclusion [i](#)



At the recommended contour level, 74% of all backbone atoms, 56% 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 (0.05) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.5600	 0.2530
A	 0.6440	 0.3040
B	 0.6910	 0.3370
C	 0.6810	 0.3170
D	 0.7010	 0.3390
E	 0.6820	 0.3290
F	 0.6910	 0.3440
G	 0.3150	 0.1020
H	 0.3190	 0.1190
I	 0.3670	 0.1480
J	 0.3900	 0.1260
K	 0.3580	 0.1500
L	 0.3340	 0.1390
M	 0.3400	 0.1560
N	 0.3270	 0.1300
O	 0.3380	 0.1450
P	 0.3920	 0.1470
Q	 0.3340	 0.0960
R	 0.2410	 0.1100
S	 0.2650	 0.1010
T	 0.2670	 0.1140
a	 0.2850	 0.0850
b	 0.3610	 0.0810
d	 0.5470	 0.2520
e	 0.5370	 0.1940
g	 0.5880	 0.2630
p	 0.3650	 0.0900

