Full wwPDB EM Validation Report ⓘ

Mar 8, 2026 – 01:30 AM UTC

PDB ID	:	5MVY / pdb_00005mvy
EMDB ID	:	EMD-3578
Title	:	Thin Filament at low calcium concentration
Authors	:	Paul, D.M.; Squire, J.M.; Morris, E.P.
Deposited on	:	2017-01-17
Resolution	:	28.40 Å(reported)

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

We welcome your comments at *validation@mail.wwpdb.org*
A user guide is available at
https://www.wwpdb.org/validation/2017/EMValidationReportHelp
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at
http://www.wwpdb.org/validation/2017/FAQs#types.

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

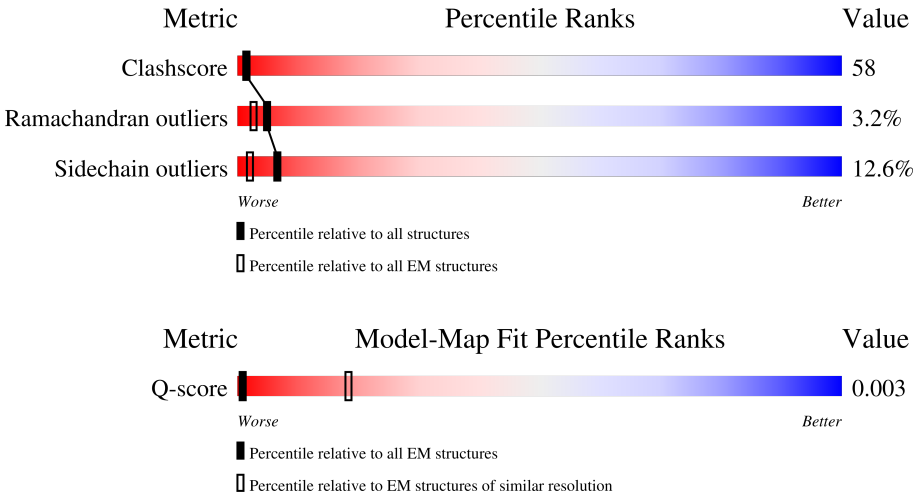
EMDB validation analysis	:	0.0.1.dev132
Mogul	:	2022.3.0, CSD as543be (2022)
MolProbity	:	4-5-2 with Phenix2.0
Buster-report	:	wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics	:	202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ	:	1.9.13
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.49

1 Overall quality at a glance (i)

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

The reported resolution of this entry is 28.40 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	11 (28.00 - 28.70)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for >=3, 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions <=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	375	80% 53% 33% 13%
1	B	375	70% 54% 32% 14%
1	C	375	9% 53% 34% 13%
1	D	375	56% 53% 34% 13%

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Mol	Chain	Length	Quality of chain
1	E	375	6% 53% 33% 14%
1	F	375	48% 54% 33% 13%
1	G	375	20% 53% 34% 13%
1	H	375	36% 52% 34% 13%
1	I	375	34% 53% 33% 14%
1	J	375	36% 52% 34% 13%
1	K	375	48% 54% 33% 13%
1	L	375	36% 52% 34% 13%
1	M	375	48% 53% 33% 14%
1	N	375	32% 52% 34% 13%
1	O	375	52% 53% 33% 14%
1	P	375	17% 53% 34% 13%
1	Q	375	43% 53% 33% 14%
1	R	375	9% 53% 34% 13%
1	S	375	38% 53% 33% 14%
1	T	375	18% 53% 33% 14%
1	U	375	33% 53% 33% 14%
1	V	375	33% 53% 33% 14%
1	W	375	28% 53% 34% 13%

2 Entry composition (i)

There are 2 unique types of molecules in this entry. The entry contains 68103 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 Actin, alpha skeletal muscle.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	B	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	C	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	D	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	E	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	F	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	G	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	H	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	I	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	J	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	K	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	L	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	M	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	N	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	O	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	P	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	Q	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		

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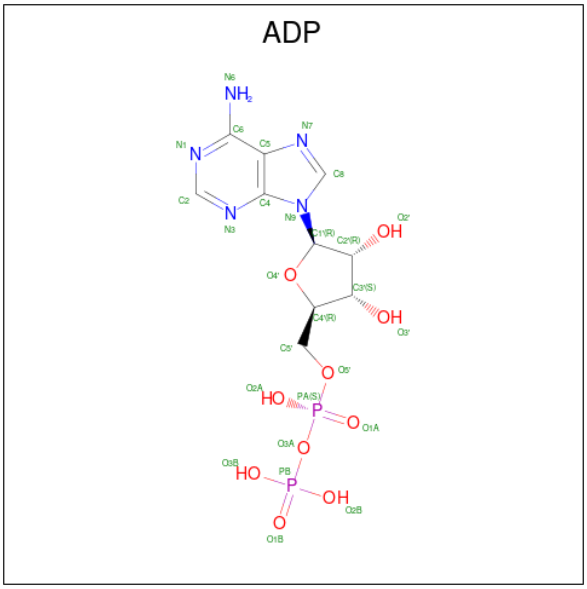
Mol	Chain	Residues	Atoms					AltConf	Trace
1	R	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	S	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	T	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	U	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	V	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		
1	W	375	Total	C	N	O	S	0	0
			2934	1855	493	565	21		

- Molecule 2 is ADENOSINE-5'-DIPHOSPHATE (CCD ID: ADP) (formula: $C_{10}H_{15}N_5O_{10}P_2$).



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

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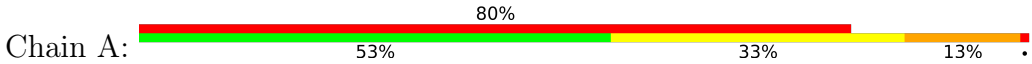
Mol	Chain	Residues	Atoms					AltConf
2	F	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	G	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	H	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	I	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	J	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	K	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	L	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	M	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	N	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	O	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	P	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	Q	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	R	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	S	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	T	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	U	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	V	1	Total	C	N	O	P	0
			27	10	5	10	2	
2	W	1	Total	C	N	O	P	0
			27	10	5	10	2	

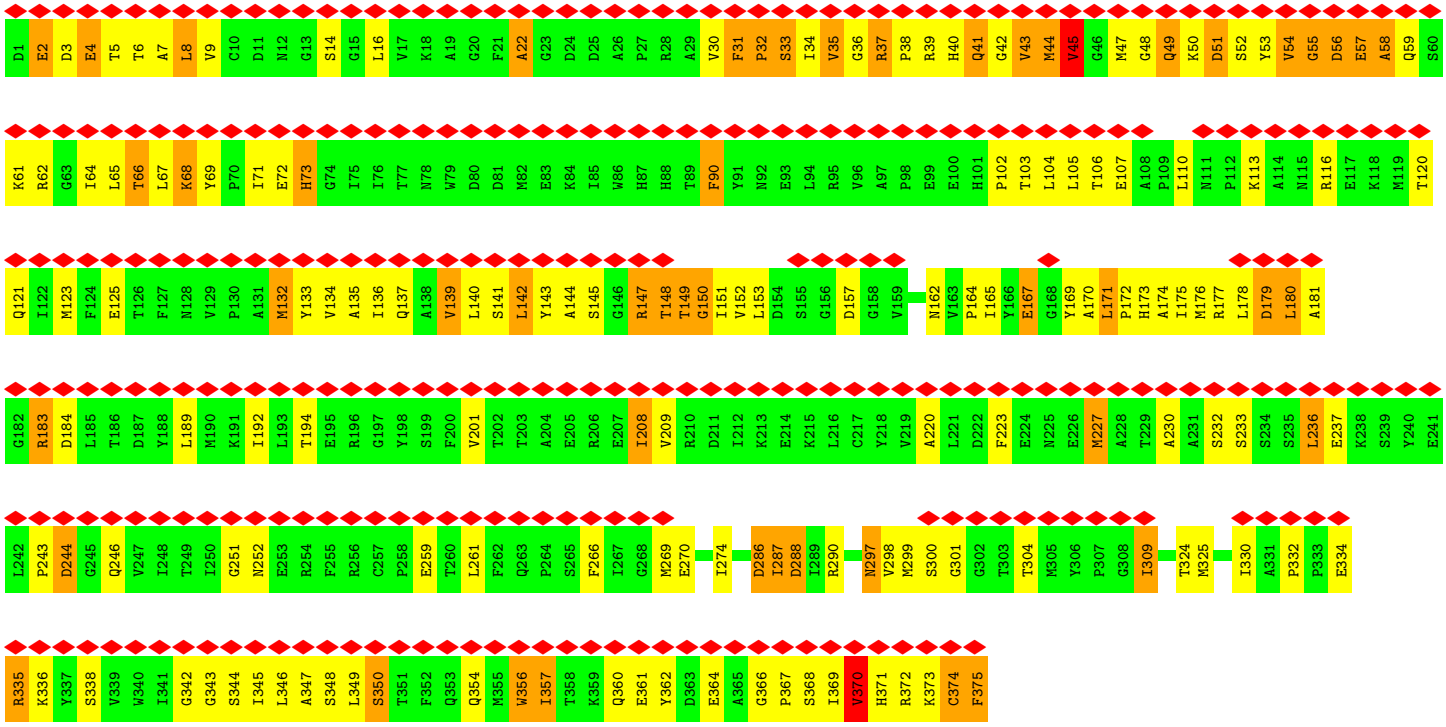
3 Residue-property plots (i)

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

- Molecule 1: Actin, alpha skeletal muscle

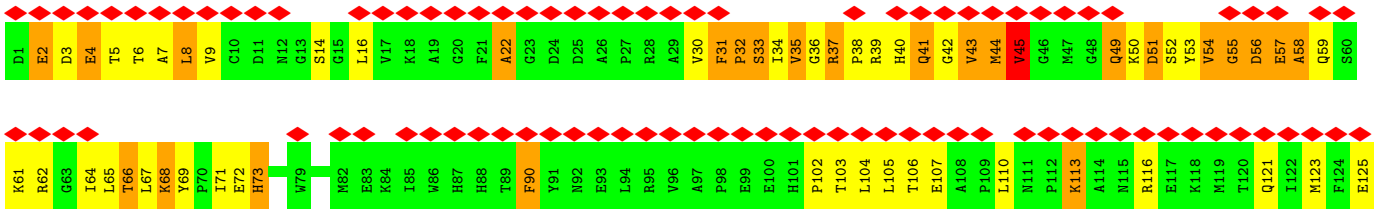
Chain A:

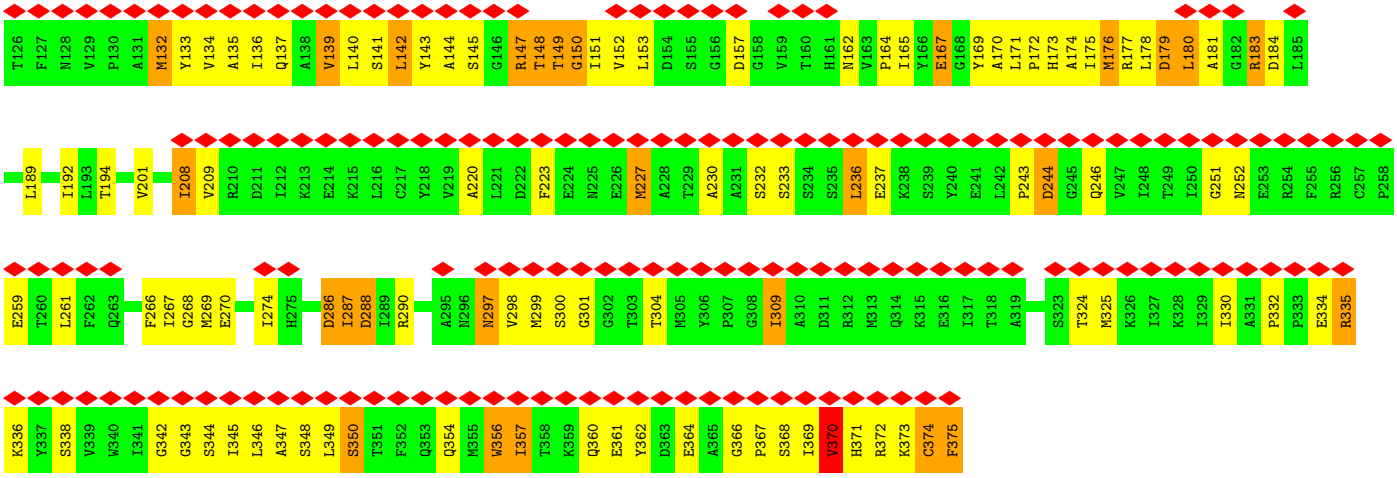




- Molecule 1: Actin, alpha skeletal muscle

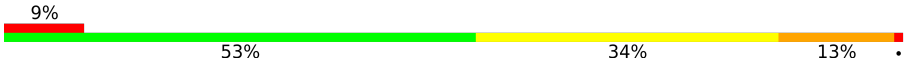
Chain B:

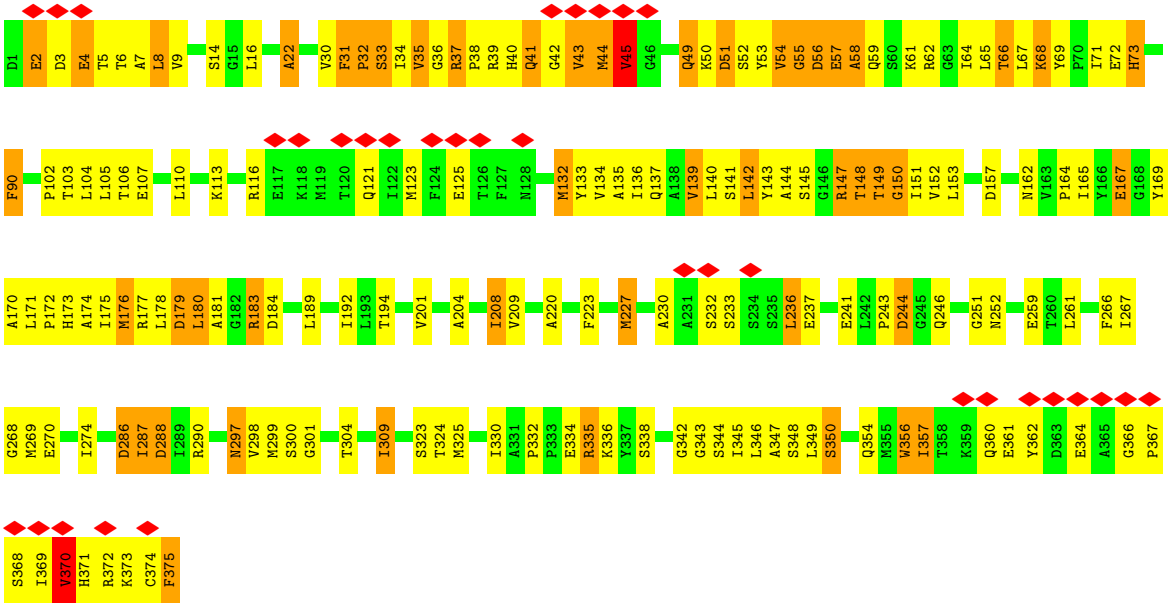




- Molecule 1: Actin, alpha skeletal muscle

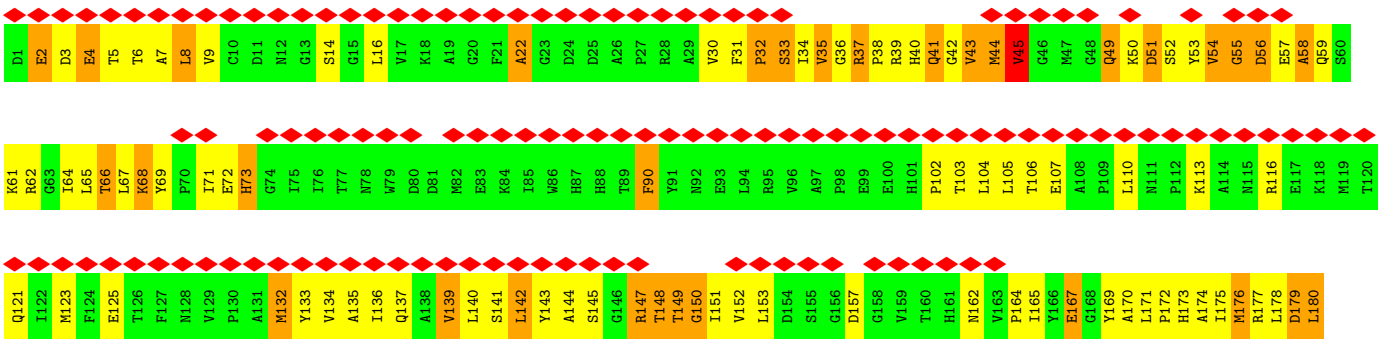
Chain C:

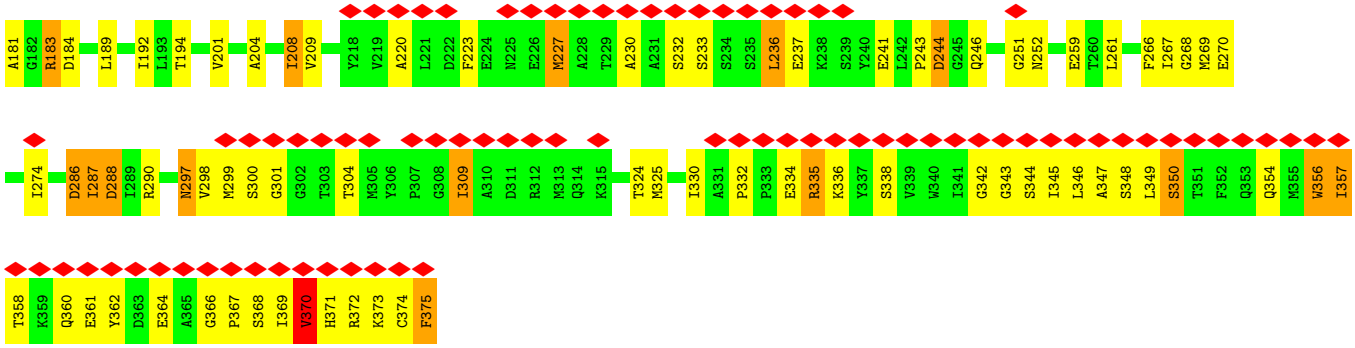




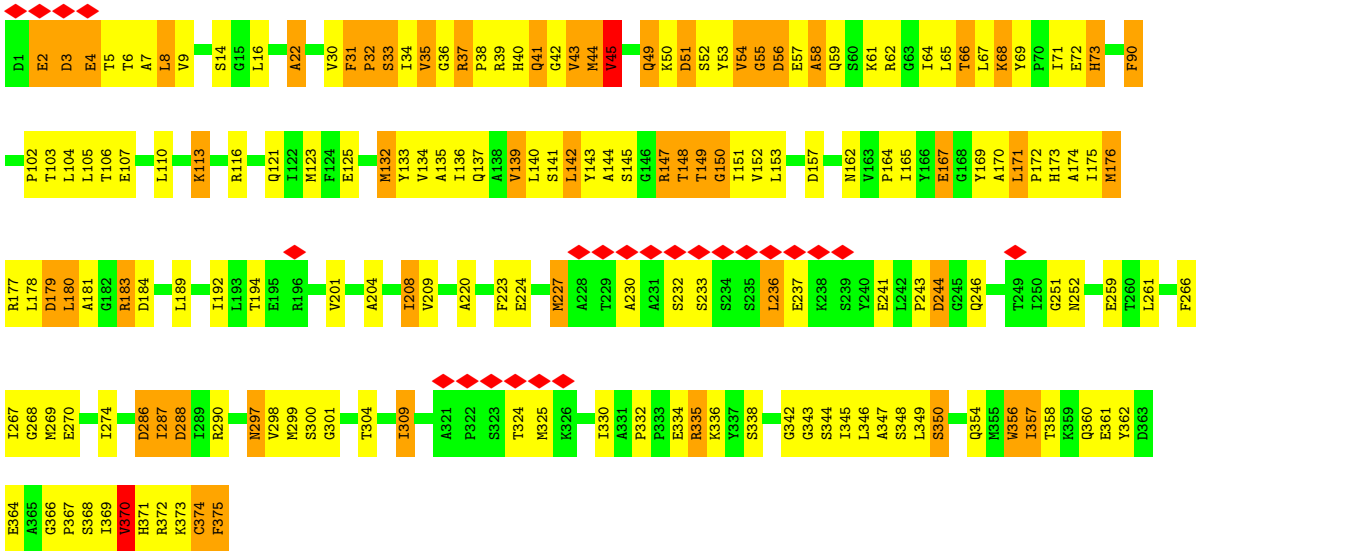
- Molecule 1: Actin, alpha skeletal muscle

Chain D:

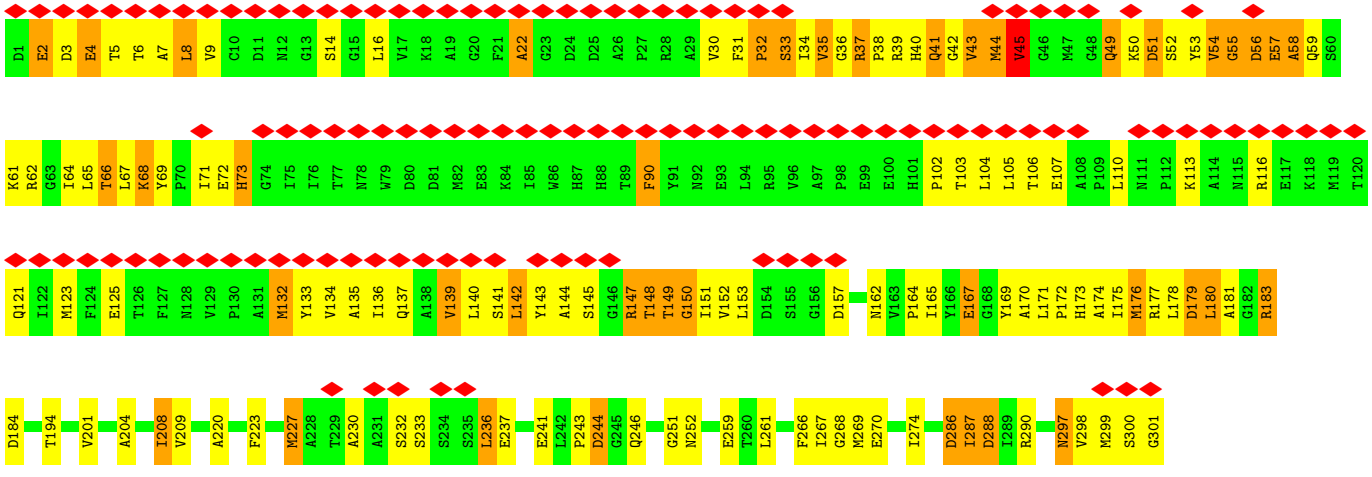


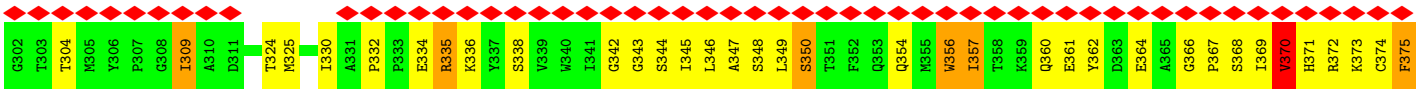


• Molecule 1: Actin, alpha skeletal muscle



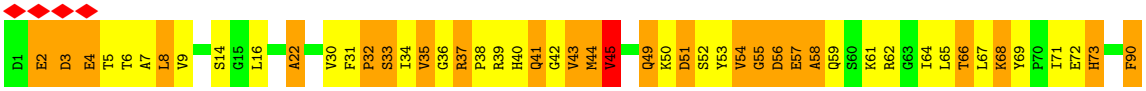
• Molecule 1: Actin, alpha skeletal muscle

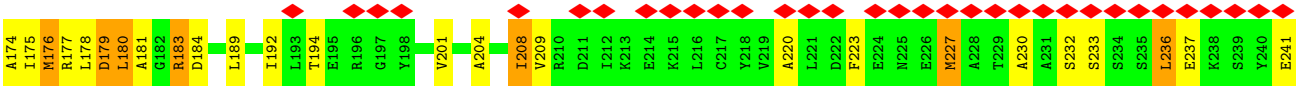


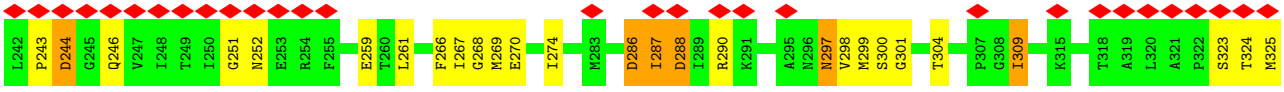


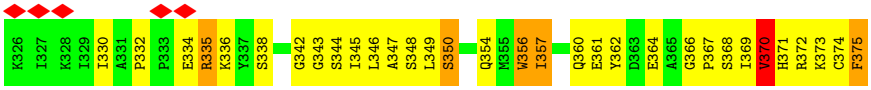
• Molecule 1: Actin, alpha skeletal muscle

Chain G:



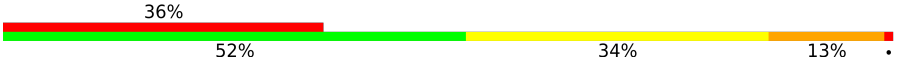


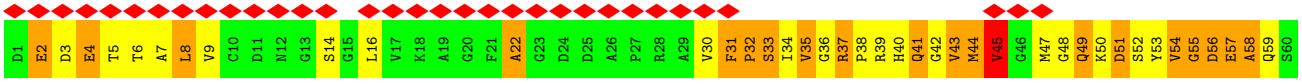


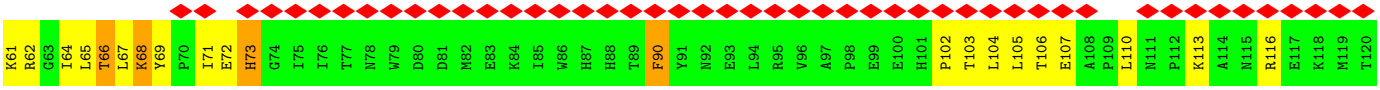


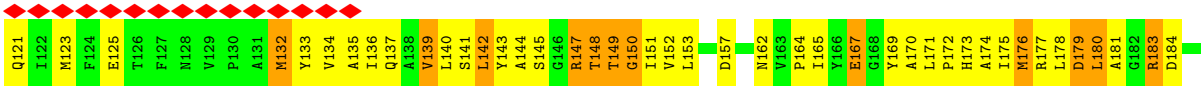
• Molecule 1: Actin, alpha skeletal muscle

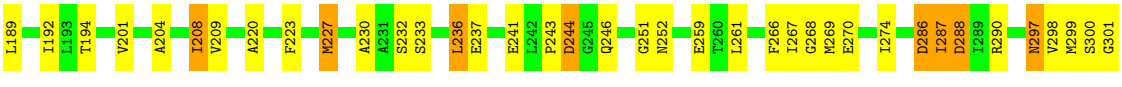
Chain H:

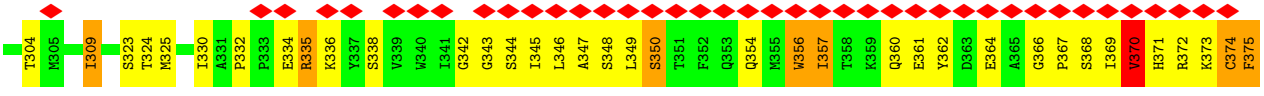






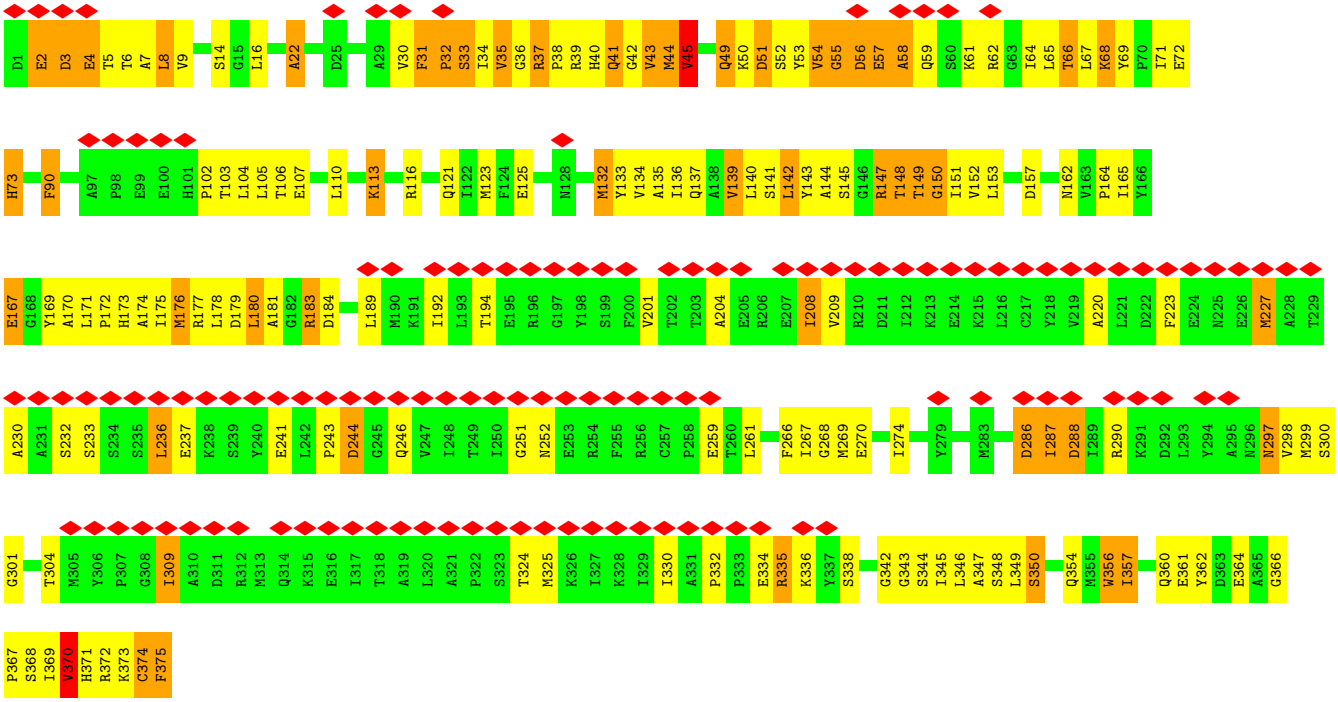






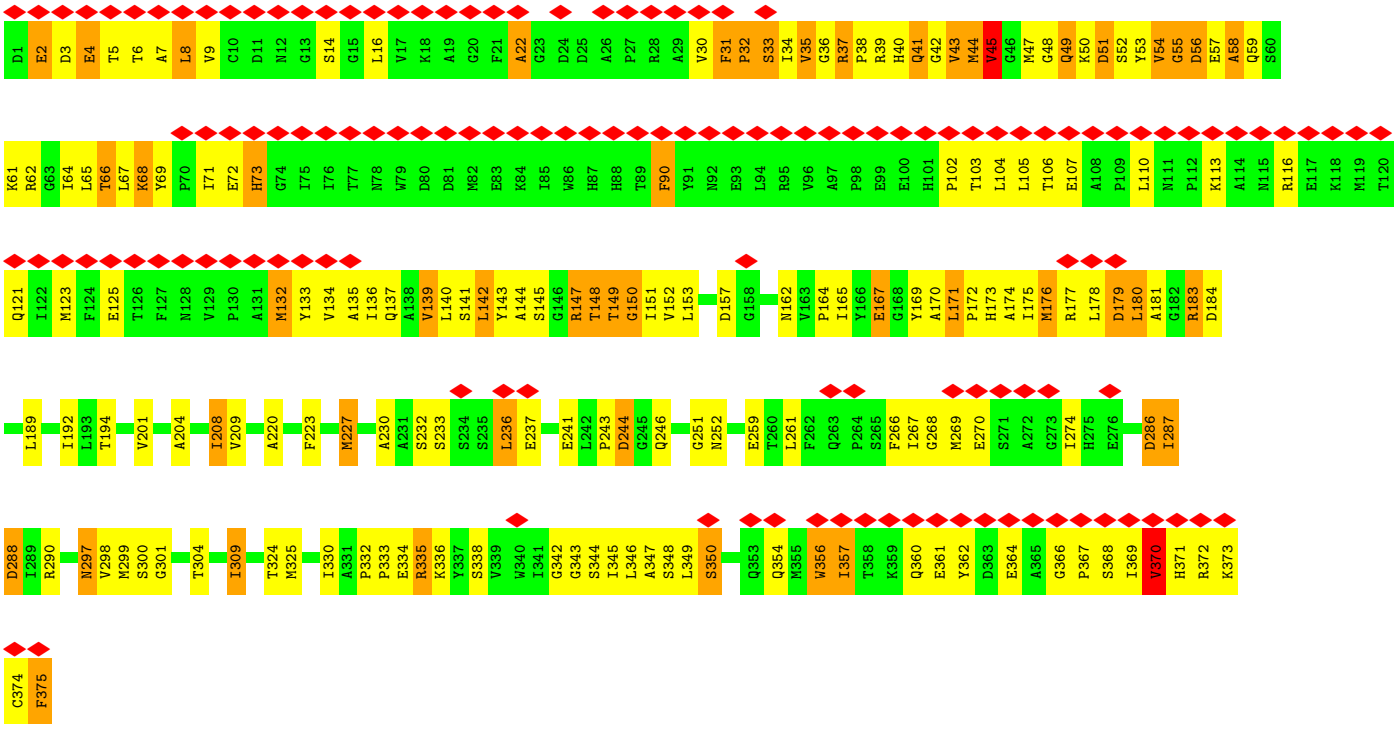
• Molecule 1: Actin, alpha skeletal muscle

Chain I:



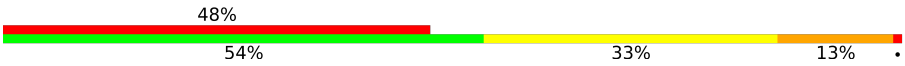
• Molecule 1: Actin, alpha skeletal muscle

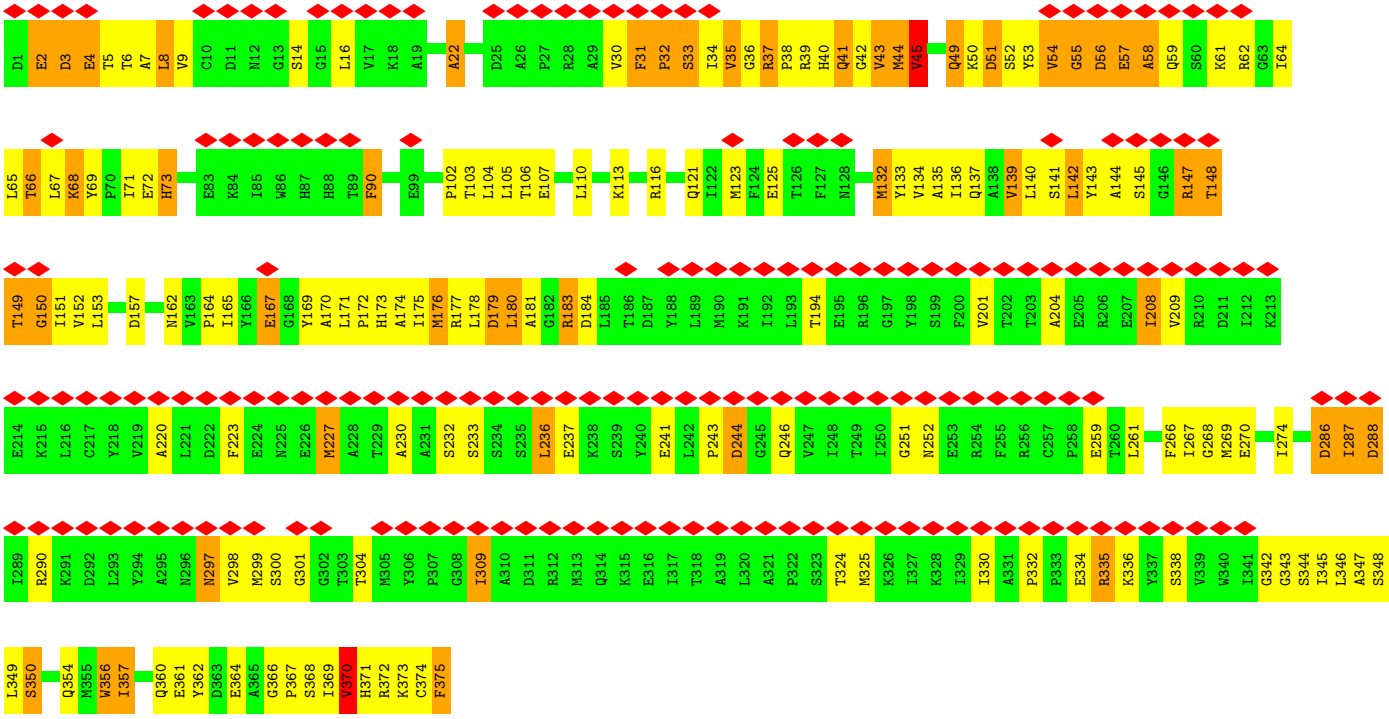
Chain J:



• Molecule 1: Actin, alpha skeletal muscle

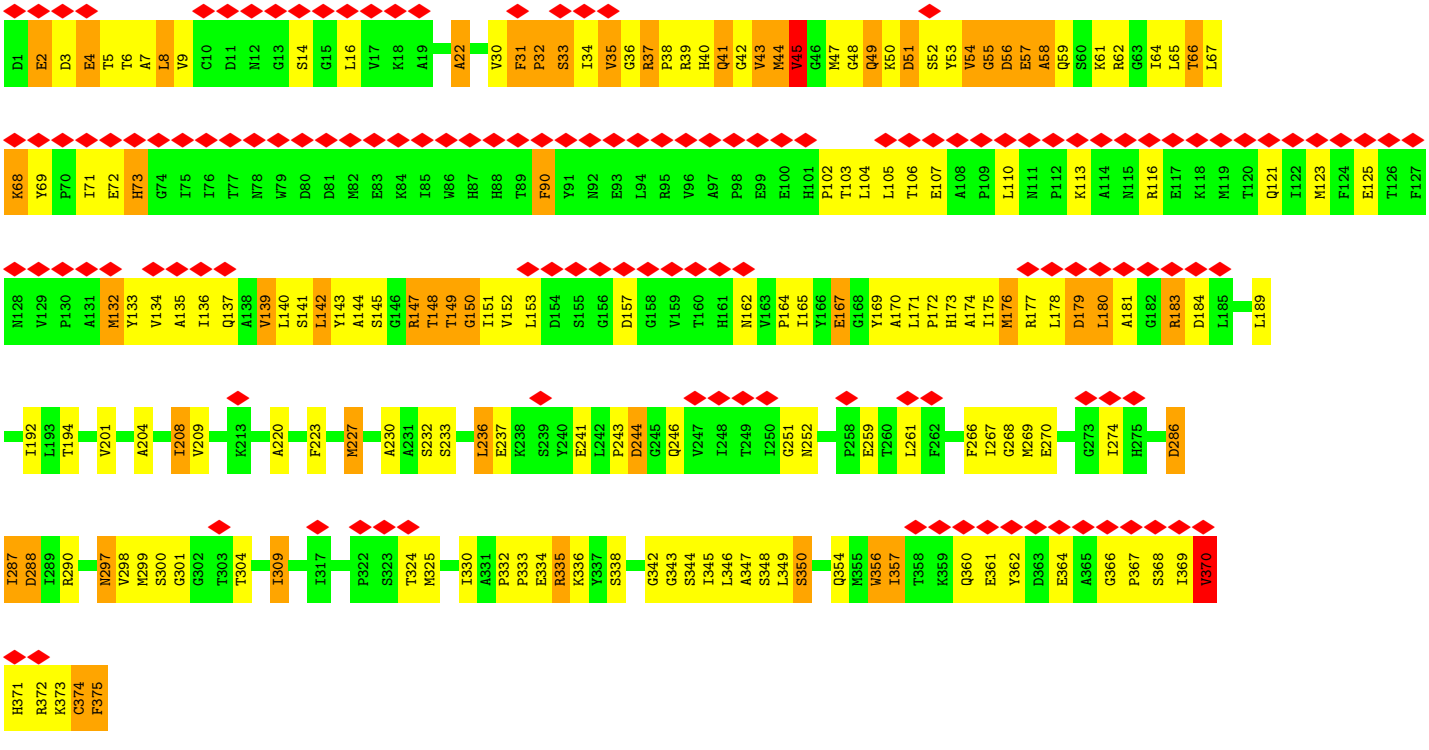
Chain K:





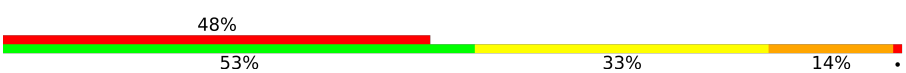
• Molecule 1: Actin, alpha skeletal muscle

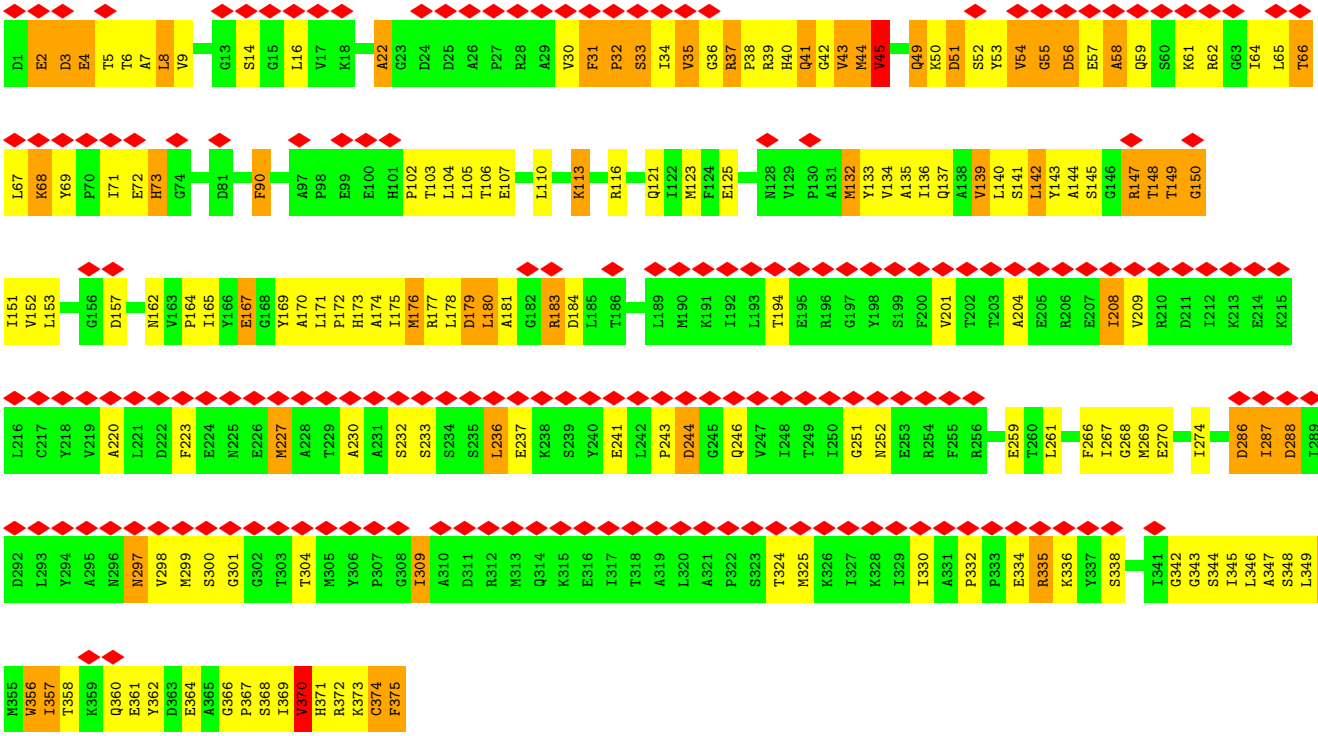
Chain L:



• Molecule 1: Actin, alpha skeletal muscle

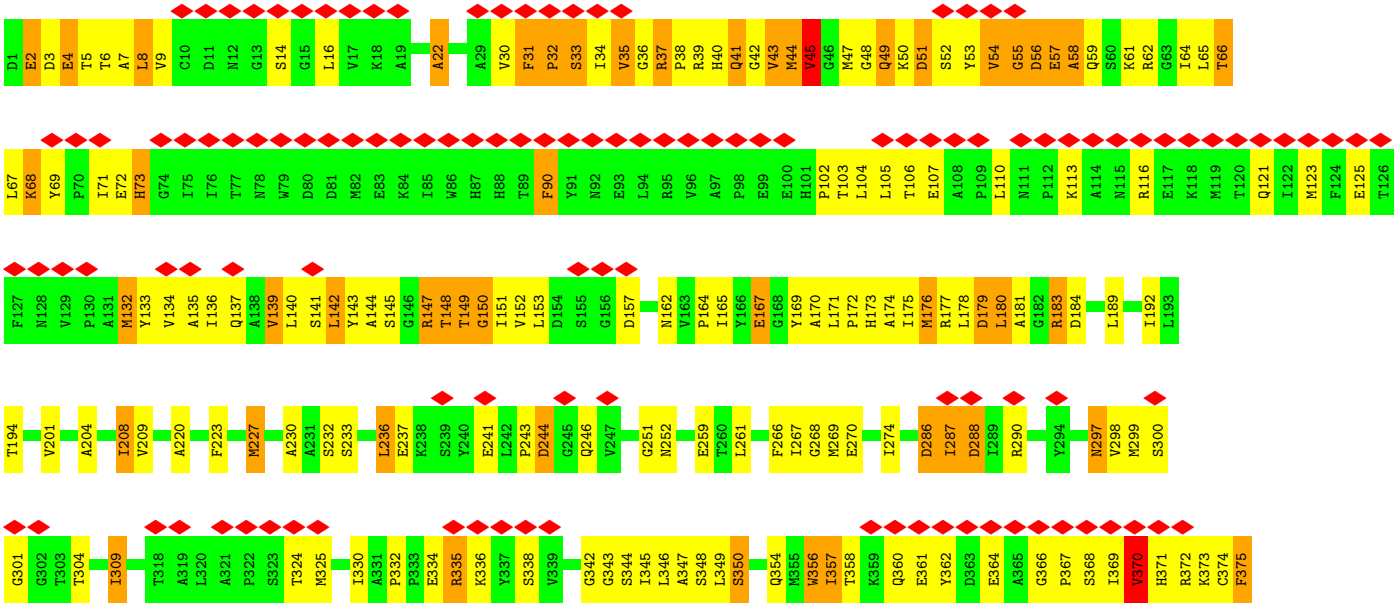
Chain M:





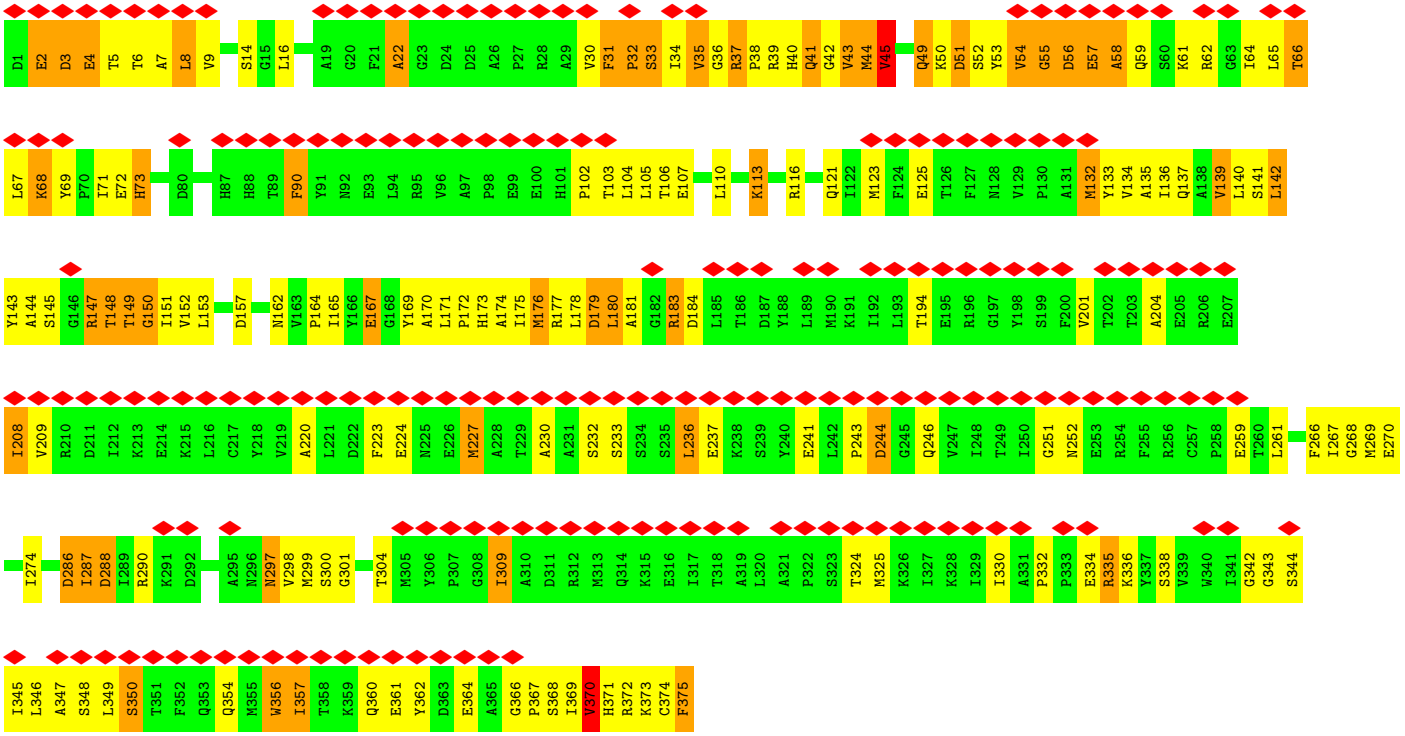
• Molecule 1: Actin, alpha skeletal muscle

Chain N:

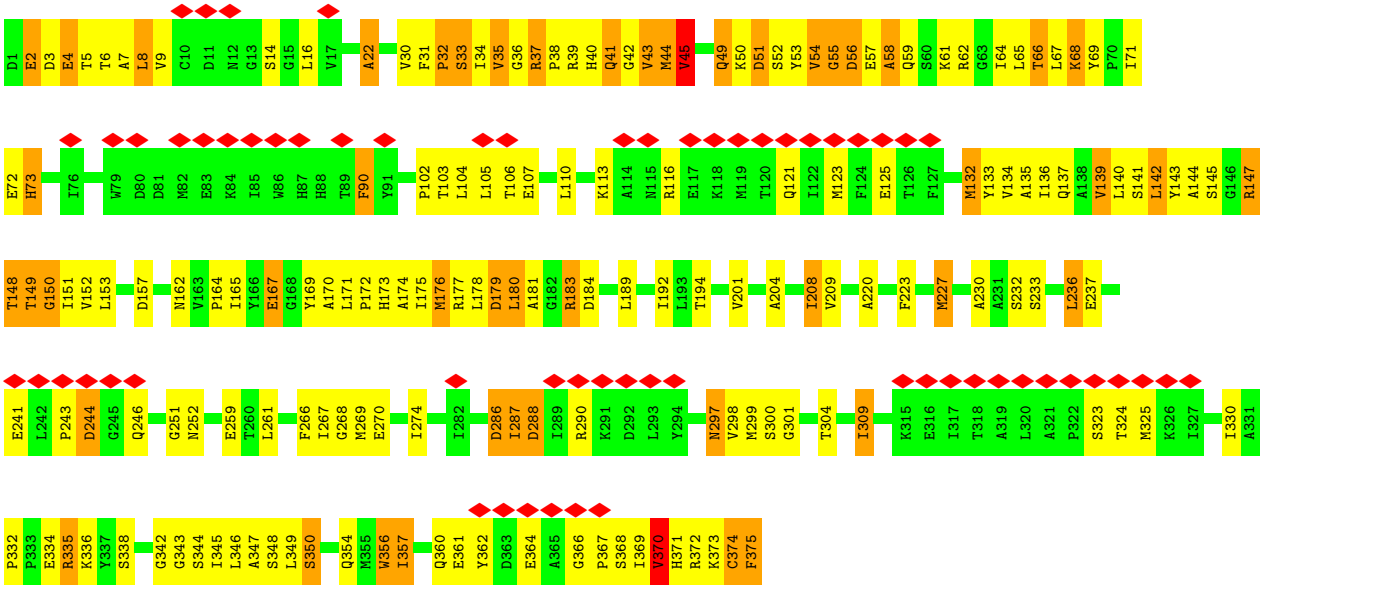


• Molecule 1: Actin, alpha skeletal muscle

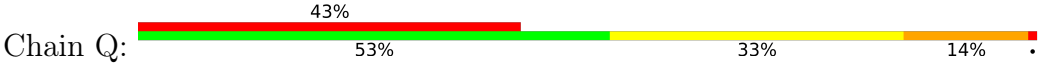
Chain O:

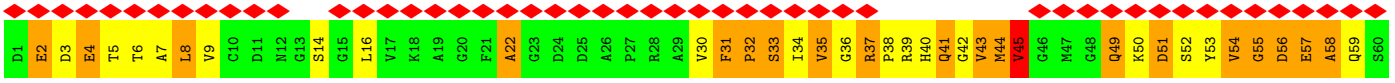


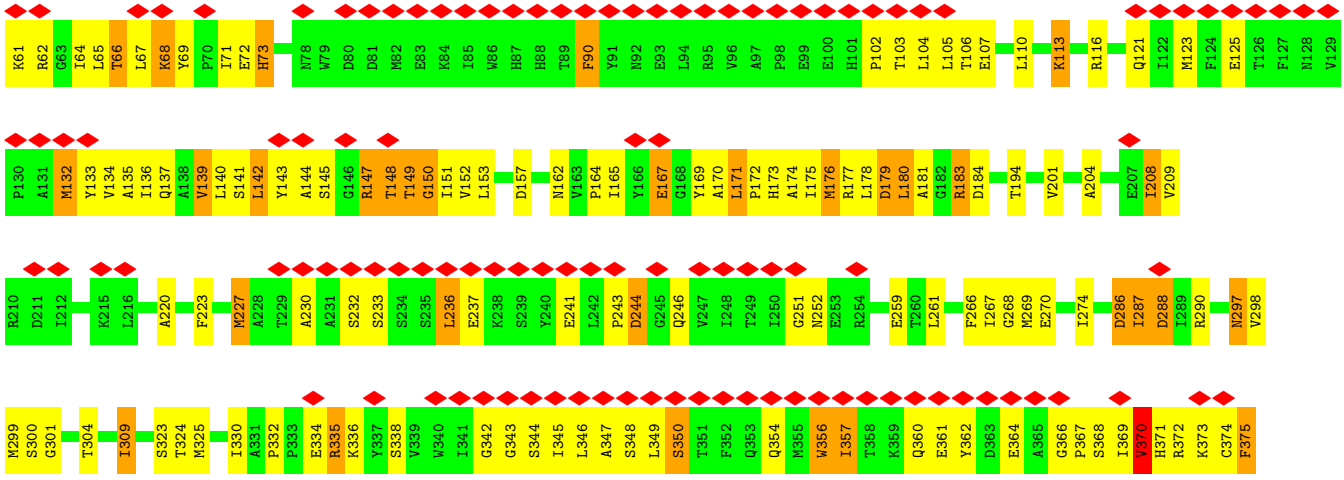
• Molecule 1: Actin, alpha skeletal muscle



• Molecule 1: Actin, alpha skeletal muscle

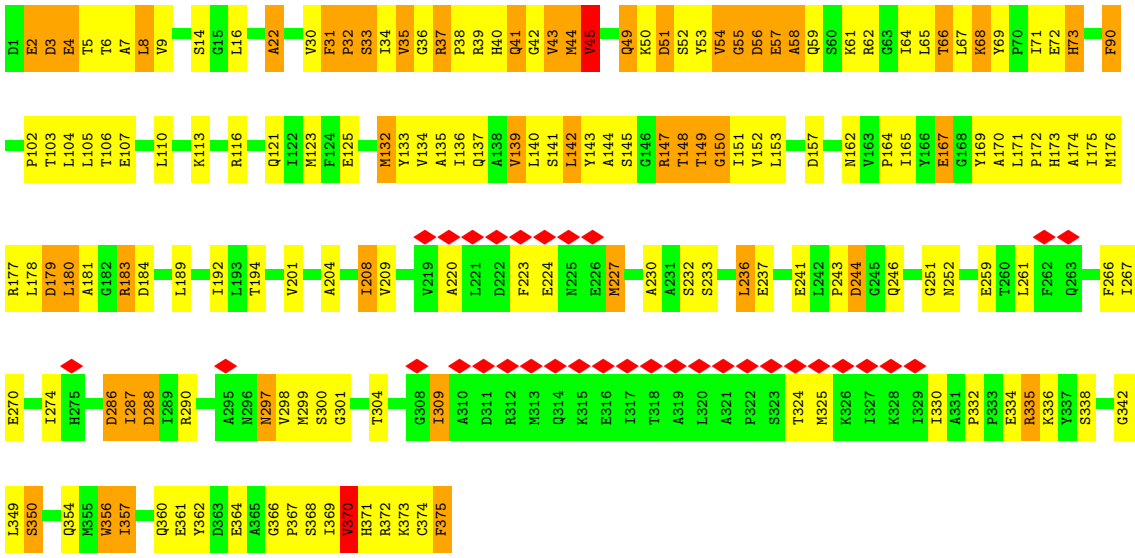






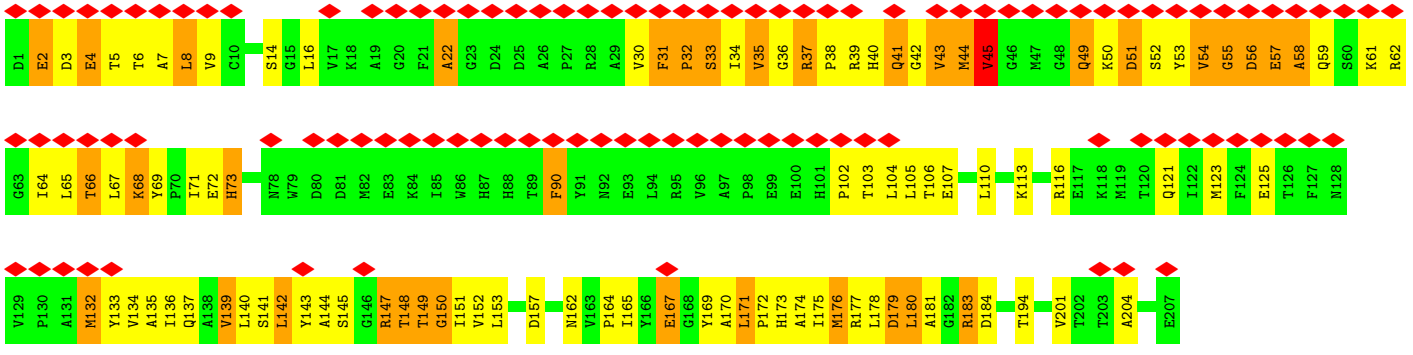
• Molecule 1: Actin, alpha skeletal muscle

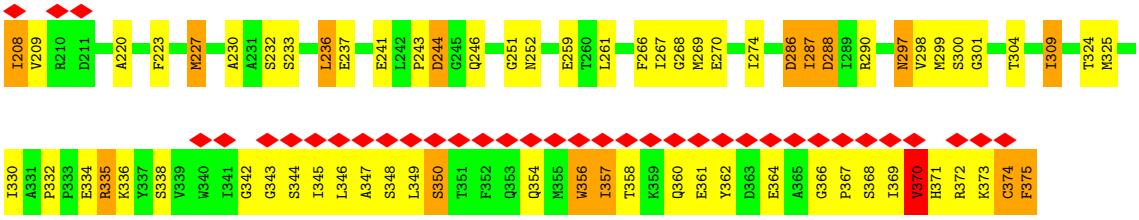
Chain R:



• Molecule 1: Actin, alpha skeletal muscle

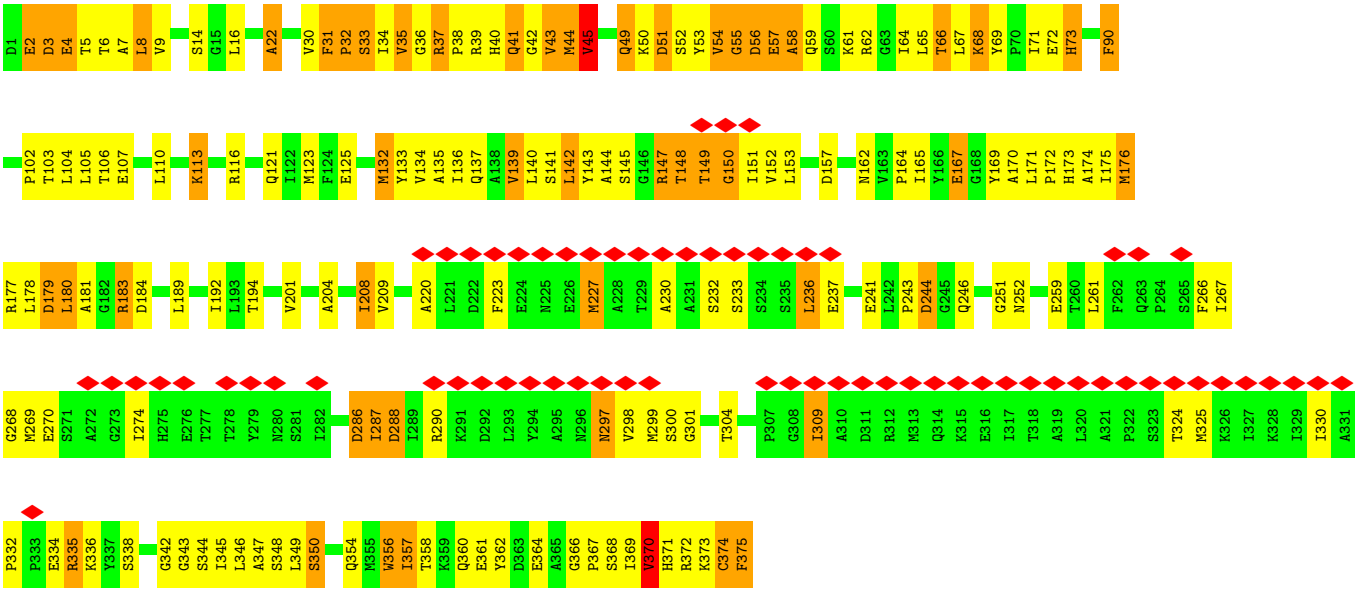
Chain S:





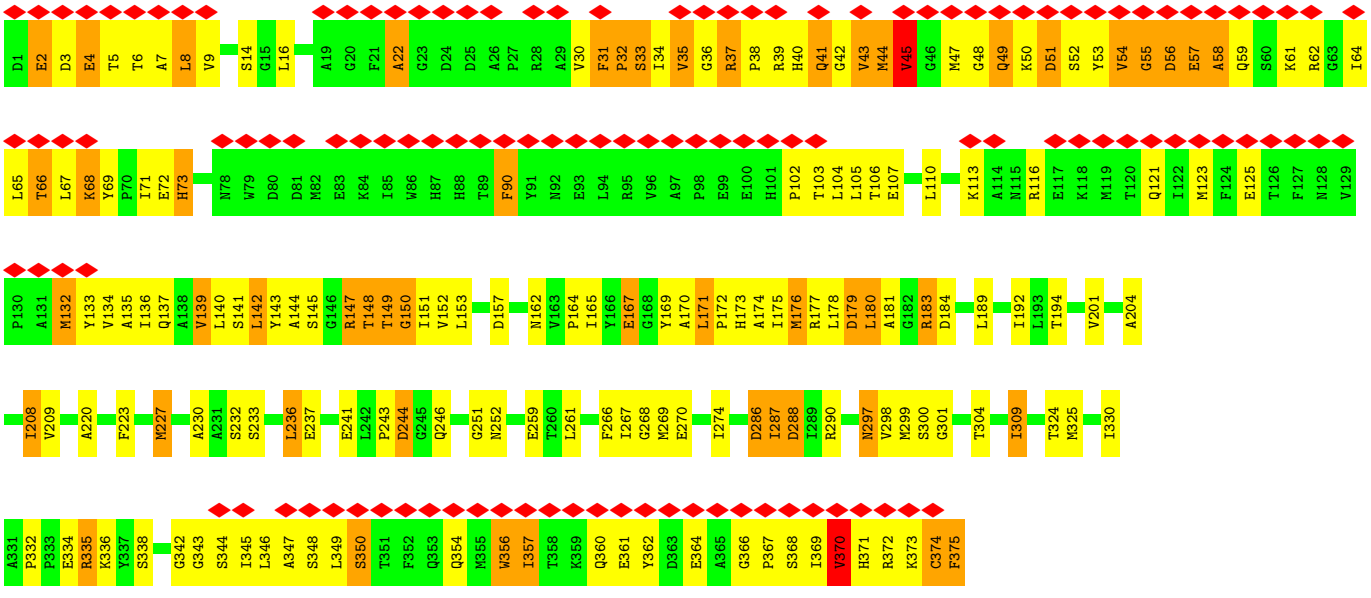
• Molecule 1: Actin, alpha skeletal muscle

Chain T:



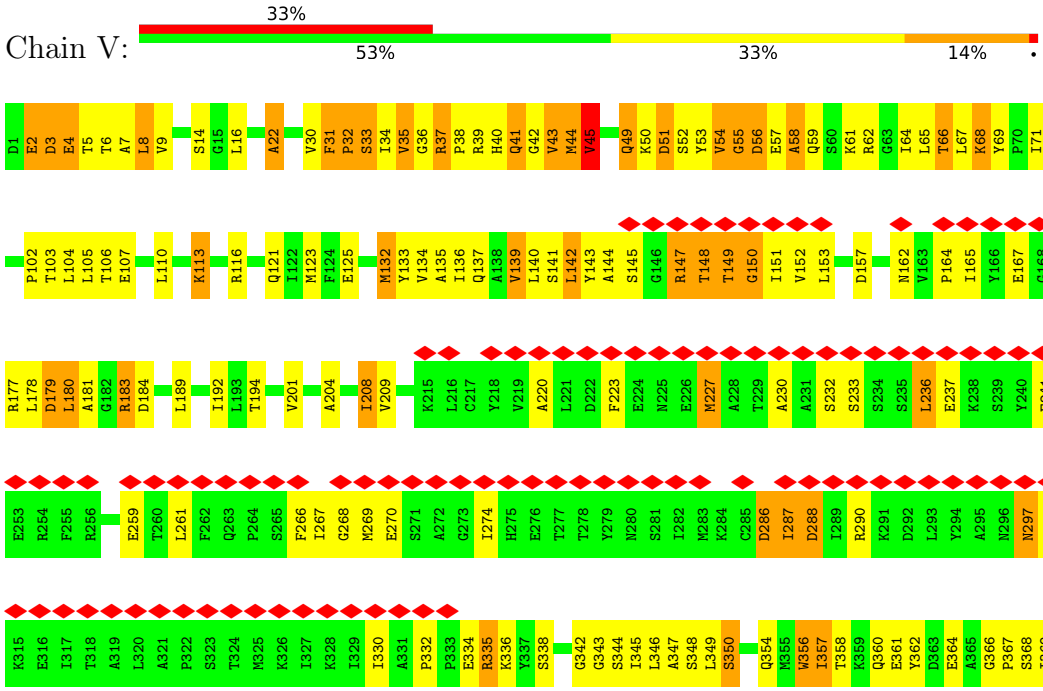
• Molecule 1: Actin, alpha skeletal muscle

Chain U:



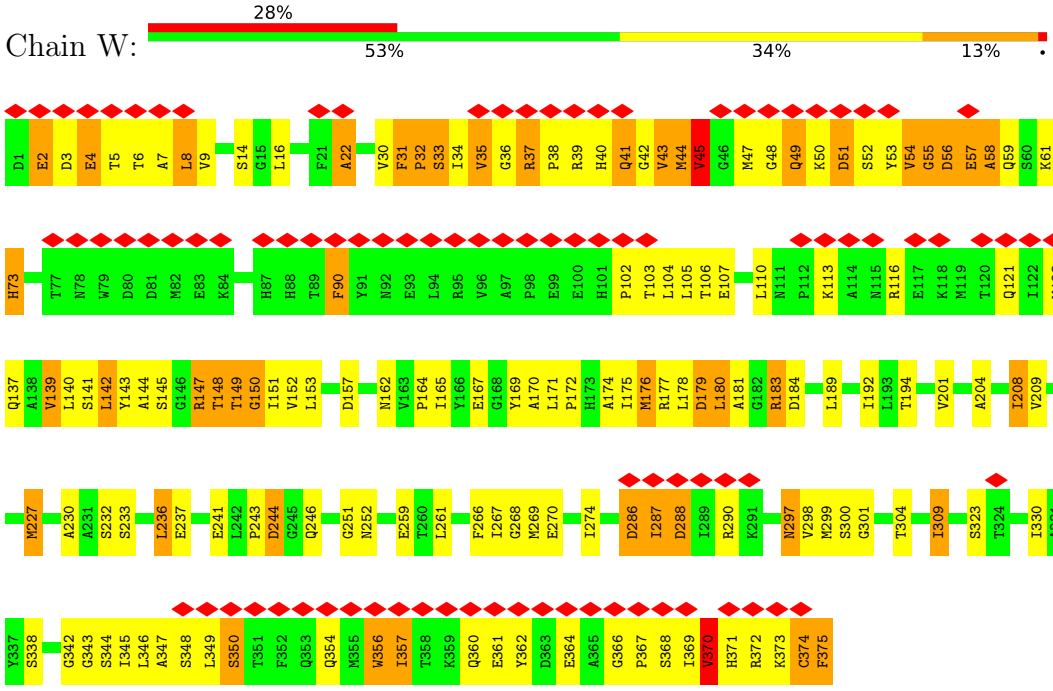
• Molecule 1: Actin, alpha skeletal muscle

Chain V:



• Molecule 1: Actin, alpha skeletal muscle

Chain W:



4 Experimental information (i)

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	1360	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	NONE	Depositor
Microscope	FEI/PHILIPS CM12	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/Å^2$)	12	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	AGFA SCIENTA FILM	Depositor
Maximum map value	20.259	Depositor
Minimum map value	-11.152	Depositor
Average map value	0.011	Depositor
Map value standard deviation	0.947	Depositor
Recommended contour level	1.8	Depositor
Map size (Å)	633.6, 633.6, 633.6	wwPDB
Map dimensions	96, 96, 96	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	6.6, 6.6, 6.6	Depositor

5 Model quality (i)

5.1 Standard geometry (i)

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

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	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	B	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	C	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
1	D	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	E	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	F	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	G	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
1	H	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
1	I	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	J	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	K	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	L	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	M	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	N	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	O	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	P	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
1	Q	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
1	R	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	S	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	T	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	U	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	V	0.88	1/2984 (0.0%)	1.42	29/4040 (0.7%)
1	W	0.88	1/2984 (0.0%)	1.42	30/4040 (0.7%)
All	All	0.88	23/68632 (0.0%)	1.42	673/92920 (0.7%)

All (23) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	P	31	PHE	C-N	-6.48	1.25	1.33
1	F	31	PHE	C-N	-6.47	1.25	1.33
1	G	31	PHE	C-N	-6.47	1.25	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	R	31	PHE	C-N	-6.47	1.25	1.33
1	E	31	PHE	C-N	-6.46	1.25	1.33
1	A	31	PHE	C-N	-6.46	1.25	1.33
1	K	31	PHE	C-N	-6.46	1.25	1.33
1	B	31	PHE	C-N	-6.44	1.25	1.33
1	J	31	PHE	C-N	-6.44	1.25	1.33
1	W	31	PHE	C-N	-6.44	1.25	1.33
1	S	31	PHE	C-N	-6.44	1.25	1.33
1	I	31	PHE	C-N	-6.43	1.25	1.33
1	O	31	PHE	C-N	-6.43	1.25	1.33
1	T	31	PHE	C-N	-6.42	1.25	1.33
1	Q	31	PHE	C-N	-6.42	1.25	1.33
1	L	31	PHE	C-N	-6.42	1.25	1.33
1	N	31	PHE	C-N	-6.42	1.25	1.33
1	V	31	PHE	C-N	-6.42	1.25	1.33
1	U	31	PHE	C-N	-6.42	1.25	1.33
1	C	31	PHE	C-N	-6.41	1.25	1.33
1	M	31	PHE	C-N	-6.40	1.25	1.33
1	D	31	PHE	C-N	-6.38	1.25	1.33
1	H	31	PHE	C-N	-6.38	1.25	1.33

All (673) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	H	286	ASP	CA-CB-CG	-8.00	104.60	112.60
1	S	286	ASP	CA-CB-CG	-8.00	104.60	112.60
1	F	286	ASP	CA-CB-CG	-8.00	104.60	112.60
1	B	286	ASP	CA-CB-CG	-7.99	104.61	112.60
1	L	286	ASP	CA-CB-CG	-7.99	104.61	112.60
1	N	286	ASP	CA-CB-CG	-7.98	104.62	112.60
1	U	286	ASP	CA-CB-CG	-7.98	104.62	112.60
1	G	286	ASP	CA-CB-CG	-7.98	104.62	112.60
1	I	286	ASP	CA-CB-CG	-7.98	104.62	112.60
1	J	286	ASP	CA-CB-CG	-7.97	104.63	112.60
1	W	286	ASP	CA-CB-CG	-7.97	104.63	112.60
1	T	286	ASP	CA-CB-CG	-7.97	104.63	112.60
1	Q	286	ASP	CA-CB-CG	-7.97	104.63	112.60
1	A	286	ASP	CA-CB-CG	-7.97	104.63	112.60
1	O	286	ASP	CA-CB-CG	-7.96	104.64	112.60
1	M	286	ASP	CA-CB-CG	-7.96	104.64	112.60
1	D	286	ASP	CA-CB-CG	-7.95	104.65	112.60
1	K	286	ASP	CA-CB-CG	-7.95	104.66	112.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	V	286	ASP	CA-CB-CG	-7.94	104.66	112.60
1	C	286	ASP	CA-CB-CG	-7.94	104.66	112.60
1	E	286	ASP	CA-CB-CG	-7.94	104.66	112.60
1	R	286	ASP	CA-CB-CG	-7.93	104.67	112.60
1	P	286	ASP	CA-CB-CG	-7.93	104.67	112.60
1	T	220	ALA	CA-C-N	7.83	132.21	120.31
1	T	220	ALA	C-N-CA	7.83	132.21	120.31
1	F	220	ALA	CA-C-N	7.82	132.20	120.31
1	F	220	ALA	C-N-CA	7.82	132.20	120.31
1	A	220	ALA	CA-C-N	7.82	132.19	120.31
1	A	220	ALA	C-N-CA	7.82	132.19	120.31
1	G	220	ALA	CA-C-N	7.81	132.18	120.31
1	G	220	ALA	C-N-CA	7.81	132.18	120.31
1	I	220	ALA	CA-C-N	7.80	132.17	120.31
1	I	220	ALA	C-N-CA	7.80	132.17	120.31
1	J	220	ALA	CA-C-N	7.80	132.17	120.31
1	J	220	ALA	C-N-CA	7.80	132.17	120.31
1	M	220	ALA	CA-C-N	7.80	132.17	120.31
1	M	220	ALA	C-N-CA	7.80	132.17	120.31
1	S	220	ALA	CA-C-N	7.80	132.17	120.31
1	S	220	ALA	C-N-CA	7.80	132.17	120.31
1	E	220	ALA	CA-C-N	7.79	132.16	120.31
1	E	220	ALA	C-N-CA	7.79	132.16	120.31
1	N	220	ALA	CA-C-N	7.79	132.16	120.31
1	N	220	ALA	C-N-CA	7.79	132.16	120.31
1	R	220	ALA	CA-C-N	7.79	132.15	120.31
1	R	220	ALA	C-N-CA	7.79	132.15	120.31
1	O	220	ALA	CA-C-N	7.79	132.15	120.31
1	O	220	ALA	C-N-CA	7.79	132.15	120.31
1	U	220	ALA	CA-C-N	7.79	132.14	120.31
1	U	220	ALA	C-N-CA	7.79	132.14	120.31
1	L	220	ALA	CA-C-N	7.78	132.14	120.31
1	L	220	ALA	C-N-CA	7.78	132.14	120.31
1	P	220	ALA	CA-C-N	7.78	132.14	120.31
1	P	220	ALA	C-N-CA	7.78	132.14	120.31
1	W	220	ALA	CA-C-N	7.78	132.13	120.31
1	W	220	ALA	C-N-CA	7.78	132.13	120.31
1	H	220	ALA	CA-C-N	7.78	132.13	120.31
1	H	220	ALA	C-N-CA	7.78	132.13	120.31
1	D	220	ALA	CA-C-N	7.77	132.12	120.31
1	D	220	ALA	C-N-CA	7.77	132.12	120.31
1	B	220	ALA	CA-C-N	7.77	132.12	120.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	B	220	ALA	C-N-CA	7.77	132.12	120.31
1	K	220	ALA	CA-C-N	7.77	132.12	120.31
1	K	220	ALA	C-N-CA	7.77	132.12	120.31
1	C	220	ALA	CA-C-N	7.77	132.12	120.31
1	C	220	ALA	C-N-CA	7.77	132.12	120.31
1	Q	220	ALA	CA-C-N	7.76	132.11	120.31
1	Q	220	ALA	C-N-CA	7.76	132.11	120.31
1	V	220	ALA	CA-C-N	7.75	132.09	120.31
1	V	220	ALA	C-N-CA	7.75	132.09	120.31
1	I	22	ALA	CA-C-N	-7.54	111.47	123.31
1	I	22	ALA	C-N-CA	-7.54	111.47	123.31
1	B	22	ALA	CA-C-N	-7.54	111.47	123.31
1	B	22	ALA	C-N-CA	-7.54	111.47	123.31
1	I	375	PHE	CA-CB-CG	-7.54	106.26	113.80
1	U	22	ALA	CA-C-N	-7.54	111.48	123.31
1	U	22	ALA	C-N-CA	-7.54	111.48	123.31
1	E	22	ALA	CA-C-N	-7.54	111.48	123.31
1	E	22	ALA	C-N-CA	-7.54	111.48	123.31
1	T	22	ALA	CA-C-N	-7.54	111.48	123.31
1	T	22	ALA	C-N-CA	-7.54	111.48	123.31
1	J	22	ALA	CA-C-N	-7.53	111.49	123.31
1	J	22	ALA	C-N-CA	-7.53	111.49	123.31
1	M	375	PHE	CA-CB-CG	-7.53	106.27	113.80
1	V	22	ALA	CA-C-N	-7.53	111.49	123.31
1	V	22	ALA	C-N-CA	-7.53	111.49	123.31
1	K	22	ALA	CA-C-N	-7.52	111.50	123.31
1	K	22	ALA	C-N-CA	-7.52	111.50	123.31
1	M	22	ALA	CA-C-N	-7.52	111.50	123.31
1	M	22	ALA	C-N-CA	-7.52	111.50	123.31
1	H	22	ALA	CA-C-N	-7.52	111.50	123.31
1	H	22	ALA	C-N-CA	-7.52	111.50	123.31
1	C	22	ALA	CA-C-N	-7.52	111.51	123.31
1	C	22	ALA	C-N-CA	-7.52	111.51	123.31
1	F	22	ALA	CA-C-N	-7.52	111.51	123.31
1	F	22	ALA	C-N-CA	-7.52	111.51	123.31
1	L	22	ALA	CA-C-N	-7.52	111.51	123.31
1	L	22	ALA	C-N-CA	-7.52	111.51	123.31
1	Q	22	ALA	CA-C-N	-7.51	111.51	123.31
1	Q	22	ALA	C-N-CA	-7.51	111.51	123.31
1	A	22	ALA	CA-C-N	-7.51	111.52	123.31
1	A	22	ALA	C-N-CA	-7.51	111.52	123.31
1	P	22	ALA	CA-C-N	-7.51	111.52	123.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	P	22	ALA	C-N-CA	-7.51	111.52	123.31
1	R	22	ALA	CA-C-N	-7.51	111.52	123.31
1	R	22	ALA	C-N-CA	-7.51	111.52	123.31
1	G	375	PHE	CA-CB-CG	-7.50	106.30	113.80
1	W	22	ALA	CA-C-N	-7.50	111.53	123.31
1	W	22	ALA	C-N-CA	-7.50	111.53	123.31
1	W	375	PHE	CA-CB-CG	-7.50	106.30	113.80
1	U	375	PHE	CA-CB-CG	-7.50	106.30	113.80
1	G	22	ALA	CA-C-N	-7.50	111.54	123.31
1	G	22	ALA	C-N-CA	-7.50	111.54	123.31
1	S	22	ALA	CA-C-N	-7.50	111.54	123.31
1	S	22	ALA	C-N-CA	-7.50	111.54	123.31
1	D	22	ALA	CA-C-N	-7.50	111.54	123.31
1	D	22	ALA	C-N-CA	-7.50	111.54	123.31
1	R	375	PHE	CA-CB-CG	-7.50	106.30	113.80
1	J	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	B	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	F	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	O	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	L	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	P	375	PHE	CA-CB-CG	-7.49	106.31	113.80
1	N	22	ALA	CA-C-N	-7.48	111.56	123.31
1	N	22	ALA	C-N-CA	-7.48	111.56	123.31
1	O	22	ALA	CA-C-N	-7.48	111.56	123.31
1	O	22	ALA	C-N-CA	-7.48	111.56	123.31
1	S	375	PHE	CA-CB-CG	-7.48	106.32	113.80
1	D	375	PHE	CA-CB-CG	-7.47	106.33	113.80
1	K	375	PHE	CA-CB-CG	-7.47	106.33	113.80
1	C	375	PHE	CA-CB-CG	-7.46	106.34	113.80
1	N	375	PHE	CA-CB-CG	-7.46	106.33	113.80
1	Q	375	PHE	CA-CB-CG	-7.46	106.34	113.80
1	T	375	PHE	CA-CB-CG	-7.45	106.36	113.80
1	H	375	PHE	CA-CB-CG	-7.44	106.36	113.80
1	V	375	PHE	CA-CB-CG	-7.43	106.37	113.80
1	E	375	PHE	CA-CB-CG	-7.43	106.37	113.80
1	A	375	PHE	CA-CB-CG	-7.43	106.37	113.80
1	B	201	VAL	O-C-N	-7.34	113.39	122.57
1	O	201	VAL	O-C-N	-7.33	113.40	122.57
1	G	201	VAL	O-C-N	-7.32	113.42	122.57
1	M	201	VAL	O-C-N	-7.32	113.42	122.57
1	K	201	VAL	O-C-N	-7.32	113.42	122.57
1	F	201	VAL	O-C-N	-7.32	113.42	122.57

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	H	201	VAL	O-C-N	-7.31	113.43	122.57
1	V	201	VAL	O-C-N	-7.31	113.43	122.57
1	A	201	VAL	O-C-N	-7.31	113.44	122.57
1	Q	201	VAL	O-C-N	-7.30	113.44	122.57
1	U	201	VAL	O-C-N	-7.30	113.44	122.57
1	N	201	VAL	O-C-N	-7.30	113.45	122.57
1	C	201	VAL	O-C-N	-7.30	113.45	122.57
1	T	201	VAL	O-C-N	-7.29	113.46	122.57
1	E	201	VAL	O-C-N	-7.28	113.47	122.57
1	R	201	VAL	O-C-N	-7.28	113.46	122.57
1	D	201	VAL	O-C-N	-7.28	113.47	122.57
1	L	201	VAL	O-C-N	-7.28	113.47	122.57
1	W	201	VAL	O-C-N	-7.28	113.47	122.57
1	I	201	VAL	O-C-N	-7.28	113.48	122.57
1	P	201	VAL	O-C-N	-7.27	113.48	122.57
1	S	201	VAL	O-C-N	-7.26	113.50	122.57
1	J	201	VAL	O-C-N	-7.25	113.50	122.57
1	M	297	ASN	CA-CB-CG	6.80	119.40	112.60
1	U	297	ASN	CA-CB-CG	6.79	119.39	112.60
1	H	297	ASN	CA-CB-CG	6.78	119.38	112.60
1	P	297	ASN	CA-CB-CG	6.76	119.36	112.60
1	R	297	ASN	CA-CB-CG	6.76	119.36	112.60
1	S	356	TRP	CA-C-O	-6.76	114.38	121.55
1	E	297	ASN	CA-CB-CG	6.76	119.36	112.60
1	J	297	ASN	CA-CB-CG	6.75	119.35	112.60
1	I	297	ASN	CA-CB-CG	6.75	119.35	112.60
1	K	297	ASN	CA-CB-CG	6.75	119.35	112.60
1	Q	297	ASN	CA-CB-CG	6.74	119.34	112.60
1	L	297	ASN	CA-CB-CG	6.74	119.34	112.60
1	O	297	ASN	CA-CB-CG	6.74	119.34	112.60
1	T	297	ASN	CA-CB-CG	6.73	119.33	112.60
1	T	356	TRP	CA-C-O	-6.73	114.42	121.55
1	A	297	ASN	CA-CB-CG	6.73	119.33	112.60
1	D	297	ASN	CA-CB-CG	6.73	119.33	112.60
1	F	356	TRP	CA-C-O	-6.73	114.42	121.55
1	O	356	TRP	CA-C-O	-6.73	114.42	121.55
1	G	356	TRP	CA-C-O	-6.73	114.42	121.55
1	N	297	ASN	CA-CB-CG	6.73	119.33	112.60
1	V	297	ASN	CA-CB-CG	6.72	119.32	112.60
1	S	297	ASN	CA-CB-CG	6.72	119.32	112.60
1	E	356	TRP	CA-C-O	-6.72	114.43	121.55
1	M	356	TRP	CA-C-O	-6.71	114.43	121.55

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	I	356	TRP	CA-C-O	-6.71	114.44	121.55
1	W	297	ASN	CA-CB-CG	6.71	119.31	112.60
1	C	297	ASN	CA-CB-CG	6.71	119.31	112.60
1	B	297	ASN	CA-CB-CG	6.71	119.31	112.60
1	R	356	TRP	CA-C-O	-6.71	114.44	121.55
1	B	356	TRP	CA-C-O	-6.71	114.44	121.55
1	N	356	TRP	CA-C-O	-6.71	114.44	121.55
1	P	356	TRP	CA-C-O	-6.70	114.44	121.55
1	F	297	ASN	CA-CB-CG	6.70	119.30	112.60
1	V	356	TRP	CA-C-O	-6.70	114.45	121.55
1	H	356	TRP	CA-C-O	-6.70	114.45	121.55
1	G	297	ASN	CA-CB-CG	6.69	119.29	112.60
1	A	356	TRP	CA-C-O	-6.69	114.46	121.55
1	W	356	TRP	CA-C-O	-6.68	114.47	121.55
1	L	356	TRP	CA-C-O	-6.68	114.47	121.55
1	J	356	TRP	CA-C-O	-6.67	114.47	121.55
1	K	356	TRP	CA-C-O	-6.67	114.48	121.55
1	C	356	TRP	CA-C-O	-6.67	114.48	121.55
1	D	356	TRP	CA-C-O	-6.66	114.49	121.55
1	Q	356	TRP	CA-C-O	-6.66	114.49	121.55
1	U	356	TRP	CA-C-O	-6.65	114.50	121.55
1	U	179	ASP	CA-CB-CG	-6.40	106.20	112.60
1	W	179	ASP	CA-CB-CG	-6.39	106.21	112.60
1	R	179	ASP	CA-CB-CG	-6.38	106.22	112.60
1	L	179	ASP	CA-CB-CG	-6.37	106.23	112.60
1	B	179	ASP	CA-CB-CG	-6.37	106.23	112.60
1	I	179	ASP	CA-CB-CG	-6.37	106.23	112.60
1	C	179	ASP	CA-CB-CG	-6.37	106.23	112.60
1	S	179	ASP	CA-CB-CG	-6.37	106.23	112.60
1	H	179	ASP	CA-CB-CG	-6.36	106.24	112.60
1	P	179	ASP	CA-CB-CG	-6.36	106.24	112.60
1	V	179	ASP	CA-CB-CG	-6.36	106.25	112.60
1	K	179	ASP	CA-CB-CG	-6.35	106.25	112.60
1	O	179	ASP	CA-CB-CG	-6.35	106.25	112.60
1	F	179	ASP	CA-CB-CG	-6.35	106.25	112.60
1	A	179	ASP	CA-CB-CG	-6.34	106.25	112.60
1	E	179	ASP	CA-CB-CG	-6.34	106.25	112.60
1	N	179	ASP	CA-CB-CG	-6.34	106.25	112.60
1	G	179	ASP	CA-CB-CG	-6.34	106.26	112.60
1	Q	179	ASP	CA-CB-CG	-6.34	106.26	112.60
1	D	179	ASP	CA-CB-CG	-6.33	106.27	112.60
1	T	179	ASP	CA-CB-CG	-6.33	106.27	112.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	M	179	ASP	CA-CB-CG	-6.33	106.27	112.60
1	J	179	ASP	CA-CB-CG	-6.31	106.29	112.60
1	S	125	GLU	CB-CG-CD	6.28	123.27	112.60
1	C	90	PHE	CA-CB-CG	6.27	120.07	113.80
1	W	125	GLU	CB-CG-CD	6.27	123.27	112.60
1	U	90	PHE	CA-CB-CG	6.27	120.07	113.80
1	E	90	PHE	CA-CB-CG	6.26	120.06	113.80
1	R	125	GLU	CB-CG-CD	6.26	123.24	112.60
1	D	125	GLU	CB-CG-CD	6.25	123.23	112.60
1	H	90	PHE	CA-CB-CG	6.25	120.06	113.80
1	V	125	GLU	CB-CG-CD	6.25	123.23	112.60
1	Q	125	GLU	CB-CG-CD	6.25	123.22	112.60
1	G	125	GLU	CB-CG-CD	6.25	123.22	112.60
1	N	125	GLU	CB-CG-CD	6.25	123.22	112.60
1	I	125	GLU	CB-CG-CD	6.25	123.22	112.60
1	M	90	PHE	CA-CB-CG	6.25	120.05	113.80
1	J	125	GLU	CB-CG-CD	6.25	123.22	112.60
1	D	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	K	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	T	125	GLU	CB-CG-CD	6.24	123.21	112.60
1	I	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	F	125	GLU	CB-CG-CD	6.24	123.21	112.60
1	P	125	GLU	CB-CG-CD	6.24	123.21	112.60
1	W	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	M	125	GLU	CB-CG-CD	6.24	123.20	112.60
1	S	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	V	90	PHE	CA-CB-CG	6.24	120.04	113.80
1	O	90	PHE	CA-CB-CG	6.23	120.03	113.80
1	J	90	PHE	CA-CB-CG	6.23	120.03	113.80
1	K	125	GLU	CB-CG-CD	6.23	123.19	112.60
1	P	90	PHE	CA-CB-CG	6.23	120.03	113.80
1	B	90	PHE	CA-CB-CG	6.23	120.03	113.80
1	E	125	GLU	CB-CG-CD	6.23	123.19	112.60
1	T	90	PHE	CA-CB-CG	6.23	120.03	113.80
1	H	125	GLU	CB-CG-CD	6.23	123.18	112.60
1	F	90	PHE	CA-CB-CG	6.22	120.03	113.80
1	L	125	GLU	CB-CG-CD	6.22	123.18	112.60
1	A	125	GLU	CB-CG-CD	6.22	123.18	112.60
1	C	125	GLU	CB-CG-CD	6.22	123.18	112.60
1	B	125	GLU	CB-CG-CD	6.22	123.17	112.60
1	Q	90	PHE	CA-CB-CG	6.22	120.02	113.80
1	U	125	GLU	CB-CG-CD	6.22	123.17	112.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	O	125	GLU	CB-CG-CD	6.22	123.17	112.60
1	R	90	PHE	CA-CB-CG	6.21	120.02	113.80
1	A	90	PHE	CA-CB-CG	6.21	120.01	113.80
1	L	90	PHE	CA-CB-CG	6.21	120.00	113.80
1	N	90	PHE	CA-CB-CG	6.20	120.00	113.80
1	G	90	PHE	CA-CB-CG	6.20	120.00	113.80
1	A	309	ILE	CA-C-O	-5.84	114.88	120.95
1	F	309	ILE	CA-C-O	-5.84	114.88	120.95
1	I	309	ILE	CA-C-O	-5.83	114.88	120.95
1	C	309	ILE	CA-C-O	-5.83	114.89	120.95
1	N	309	ILE	CA-C-O	-5.82	114.90	120.95
1	J	309	ILE	CA-C-O	-5.82	114.90	120.95
1	R	309	ILE	CA-C-O	-5.81	114.91	120.95
1	E	309	ILE	CA-C-O	-5.81	114.91	120.95
1	K	309	ILE	CA-C-O	-5.81	114.91	120.95
1	B	309	ILE	CA-C-O	-5.80	114.91	120.95
1	Q	309	ILE	CA-C-O	-5.80	114.91	120.95
1	S	309	ILE	CA-C-O	-5.80	114.92	120.95
1	G	309	ILE	CA-C-O	-5.80	114.92	120.95
1	T	309	ILE	CA-C-O	-5.80	114.92	120.95
1	H	309	ILE	CA-C-O	-5.79	114.93	120.95
1	W	309	ILE	CA-C-O	-5.79	114.93	120.95
1	O	309	ILE	CA-C-O	-5.78	114.94	120.95
1	D	184	ASP	CA-CB-CG	-5.78	106.82	112.60
1	V	309	ILE	CA-C-O	-5.78	114.94	120.95
1	M	309	ILE	CA-C-O	-5.78	114.94	120.95
1	P	309	ILE	CA-C-O	-5.78	114.94	120.95
1	P	184	ASP	CA-CB-CG	-5.77	106.83	112.60
1	B	184	ASP	CA-CB-CG	-5.76	106.84	112.60
1	D	309	ILE	CA-C-O	-5.76	114.95	120.95
1	U	309	ILE	CA-C-O	-5.75	114.97	120.95
1	K	184	ASP	CA-CB-CG	-5.75	106.85	112.60
1	U	184	ASP	CA-CB-CG	-5.75	106.85	112.60
1	F	184	ASP	CA-CB-CG	-5.75	106.85	112.60
1	A	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	H	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	R	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	I	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	G	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	L	309	ILE	CA-C-O	-5.74	114.98	120.95
1	N	184	ASP	CA-CB-CG	-5.74	106.86	112.60
1	C	184	ASP	CA-CB-CG	-5.73	106.87	112.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	L	184	ASP	CA-CB-CG	-5.73	106.87	112.60
1	V	184	ASP	CA-CB-CG	-5.73	106.87	112.60
1	T	184	ASP	CA-CB-CG	-5.73	106.87	112.60
1	W	184	ASP	CA-CB-CG	-5.73	106.87	112.60
1	O	184	ASP	CA-CB-CG	-5.73	106.87	112.60
1	E	184	ASP	CA-CB-CG	-5.72	106.88	112.60
1	J	184	ASP	CA-CB-CG	-5.72	106.88	112.60
1	M	184	ASP	CA-CB-CG	-5.71	106.89	112.60
1	Q	184	ASP	CA-CB-CG	-5.70	106.90	112.60
1	S	184	ASP	CA-CB-CG	-5.69	106.91	112.60
1	N	30	VAL	CA-C-O	-5.63	114.79	120.31
1	J	30	VAL	CA-C-O	-5.63	114.79	120.31
1	R	30	VAL	CA-C-O	-5.61	114.81	120.31
1	T	30	VAL	CA-C-O	-5.61	114.81	120.31
1	B	288	ASP	CA-CB-CG	-5.59	107.01	112.60
1	I	30	VAL	CA-C-O	-5.59	114.83	120.31
1	M	288	ASP	CA-CB-CG	-5.59	107.01	112.60
1	F	30	VAL	CA-C-O	-5.59	114.83	120.31
1	H	30	VAL	CA-C-O	-5.59	114.83	120.31
1	U	288	ASP	CA-CB-CG	-5.59	107.01	112.60
1	G	30	VAL	CA-C-O	-5.59	114.83	120.31
1	D	288	ASP	CA-CB-CG	-5.58	107.02	112.60
1	M	30	VAL	CA-C-O	-5.58	114.84	120.31
1	Q	30	VAL	CA-C-O	-5.58	114.84	120.31
1	D	30	VAL	CA-C-O	-5.58	114.85	120.31
1	E	30	VAL	CA-C-O	-5.57	114.85	120.31
1	W	288	ASP	CA-CB-CG	-5.57	107.03	112.60
1	N	288	ASP	CA-CB-CG	-5.56	107.04	112.60
1	S	288	ASP	CA-CB-CG	-5.56	107.04	112.60
1	V	30	VAL	CA-C-O	-5.56	114.86	120.31
1	L	30	VAL	CA-C-O	-5.55	114.87	120.31
1	S	30	VAL	CA-C-O	-5.55	114.87	120.31
1	B	30	VAL	CA-C-O	-5.55	114.87	120.31
1	C	30	VAL	CA-C-O	-5.55	114.87	120.31
1	H	288	ASP	CA-CB-CG	-5.55	107.05	112.60
1	W	30	VAL	CA-C-O	-5.55	114.87	120.31
1	G	32	PRO	N-CA-C	-5.55	102.56	111.38
1	U	30	VAL	CA-C-O	-5.55	114.88	120.31
1	A	30	VAL	CA-C-O	-5.54	114.88	120.31
1	A	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	O	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	E	288	ASP	CA-CB-CG	-5.54	107.06	112.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	F	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	G	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	R	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	V	288	ASP	CA-CB-CG	-5.54	107.06	112.60
1	J	288	ASP	CA-CB-CG	-5.53	107.07	112.60
1	O	30	VAL	CA-C-O	-5.53	114.89	120.31
1	P	30	VAL	CA-C-O	-5.53	114.89	120.31
1	R	32	PRO	N-CA-C	-5.53	102.58	111.38
1	B	32	PRO	N-CA-C	-5.53	102.59	111.38
1	K	288	ASP	CA-CB-CG	-5.53	107.07	112.60
1	Q	288	ASP	CA-CB-CG	-5.53	107.07	112.60
1	I	288	ASP	CA-CB-CG	-5.53	107.07	112.60
1	N	32	PRO	N-CA-C	-5.53	102.59	111.38
1	J	32	PRO	N-CA-C	-5.53	102.59	111.38
1	K	30	VAL	CA-C-O	-5.53	114.89	120.31
1	P	288	ASP	CA-CB-CG	-5.53	107.08	112.60
1	K	32	PRO	N-CA-C	-5.52	102.60	111.38
1	Q	32	PRO	N-CA-C	-5.52	102.60	111.38
1	D	32	PRO	N-CA-C	-5.52	102.61	111.38
1	W	32	PRO	N-CA-C	-5.52	102.60	111.38
1	C	288	ASP	CA-CB-CG	-5.52	107.08	112.60
1	F	32	PRO	N-CA-C	-5.52	102.61	111.38
1	V	32	PRO	N-CA-C	-5.51	102.61	111.38
1	U	32	PRO	N-CA-C	-5.51	102.61	111.38
1	E	32	PRO	N-CA-C	-5.51	102.62	111.38
1	L	288	ASP	CA-CB-CG	-5.51	107.09	112.60
1	O	32	PRO	N-CA-C	-5.50	102.63	111.38
1	T	288	ASP	CA-CB-CG	-5.50	107.10	112.60
1	A	32	PRO	N-CA-C	-5.50	102.64	111.38
1	I	32	PRO	N-CA-C	-5.50	102.64	111.38
1	P	32	PRO	N-CA-C	-5.49	102.65	111.38
1	M	32	PRO	N-CA-C	-5.49	102.65	111.38
1	S	32	PRO	N-CA-C	-5.49	102.65	111.38
1	H	32	PRO	N-CA-C	-5.49	102.65	111.38
1	T	32	PRO	N-CA-C	-5.49	102.65	111.38
1	C	32	PRO	N-CA-C	-5.49	102.66	111.38
1	L	32	PRO	N-CA-C	-5.47	102.67	111.38
1	W	194	THR	CA-C-N	5.46	128.14	120.28
1	W	194	THR	C-N-CA	5.46	128.14	120.28
1	B	194	THR	CA-C-N	5.45	128.13	120.28
1	B	194	THR	C-N-CA	5.45	128.13	120.28
1	J	194	THR	CA-C-N	5.44	128.12	120.28

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	J	194	THR	C-N-CA	5.44	128.12	120.28
1	H	194	THR	CA-C-N	5.44	128.11	120.28
1	H	194	THR	C-N-CA	5.44	128.11	120.28
1	P	194	THR	CA-C-N	5.44	128.11	120.28
1	P	194	THR	C-N-CA	5.44	128.11	120.28
1	V	194	THR	CA-C-N	5.43	128.10	120.28
1	V	194	THR	C-N-CA	5.43	128.10	120.28
1	A	350	SER	CA-C-N	-5.43	112.06	120.31
1	A	350	SER	C-N-CA	-5.43	112.06	120.31
1	N	194	THR	CA-C-N	5.43	128.10	120.28
1	N	194	THR	C-N-CA	5.43	128.10	120.28
1	O	194	THR	CA-C-N	5.43	128.09	120.28
1	O	194	THR	C-N-CA	5.43	128.09	120.28
1	H	350	SER	CA-C-N	-5.42	112.06	120.31
1	H	350	SER	C-N-CA	-5.42	112.06	120.31
1	D	194	THR	CA-C-N	5.42	128.09	120.28
1	D	194	THR	C-N-CA	5.42	128.09	120.28
1	A	194	THR	CA-C-N	5.42	128.08	120.28
1	A	194	THR	C-N-CA	5.42	128.08	120.28
1	R	194	THR	CA-C-N	5.42	128.08	120.28
1	R	194	THR	C-N-CA	5.42	128.08	120.28
1	M	194	THR	CA-C-N	5.42	128.08	120.28
1	M	194	THR	C-N-CA	5.42	128.08	120.28
1	C	194	THR	CA-C-N	5.41	128.07	120.28
1	C	194	THR	C-N-CA	5.41	128.07	120.28
1	I	194	THR	CA-C-N	5.41	128.07	120.28
1	I	194	THR	C-N-CA	5.41	128.07	120.28
1	U	194	THR	CA-C-N	5.41	128.08	120.28
1	U	194	THR	C-N-CA	5.41	128.08	120.28
1	Q	350	SER	CA-C-N	-5.41	112.08	120.31
1	Q	350	SER	C-N-CA	-5.41	112.08	120.31
1	F	194	THR	CA-C-N	5.41	128.07	120.28
1	F	194	THR	C-N-CA	5.41	128.07	120.28
1	G	350	SER	CA-C-N	-5.41	112.09	120.31
1	G	350	SER	C-N-CA	-5.41	112.09	120.31
1	C	350	SER	CA-C-N	-5.41	112.09	120.31
1	C	350	SER	C-N-CA	-5.41	112.09	120.31
1	L	194	THR	CA-C-N	5.40	128.06	120.28
1	L	194	THR	C-N-CA	5.40	128.06	120.28
1	G	194	THR	CA-C-N	5.40	128.06	120.28
1	G	194	THR	C-N-CA	5.40	128.06	120.28
1	P	350	SER	CA-C-N	-5.40	112.10	120.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	P	350	SER	C-N-CA	-5.40	112.10	120.31
1	Q	194	THR	CA-C-N	5.40	128.06	120.28
1	Q	194	THR	C-N-CA	5.40	128.06	120.28
1	K	194	THR	CA-C-N	5.40	128.05	120.28
1	K	194	THR	C-N-CA	5.40	128.05	120.28
1	K	350	SER	CA-C-N	-5.40	112.11	120.31
1	K	350	SER	C-N-CA	-5.40	112.11	120.31
1	W	287	ILE	N-CA-C	-5.40	105.11	110.62
1	S	194	THR	CA-C-N	5.40	128.05	120.28
1	S	194	THR	C-N-CA	5.40	128.05	120.28
1	D	350	SER	CA-C-N	-5.39	112.11	120.31
1	D	350	SER	C-N-CA	-5.39	112.11	120.31
1	O	350	SER	CA-C-N	-5.39	112.11	120.31
1	O	350	SER	C-N-CA	-5.39	112.11	120.31
1	W	350	SER	CA-C-N	-5.39	112.11	120.31
1	W	350	SER	C-N-CA	-5.39	112.11	120.31
1	A	330	ILE	CA-C-N	-5.39	113.70	122.65
1	A	330	ILE	C-N-CA	-5.39	113.70	122.65
1	E	350	SER	CA-C-N	-5.39	112.12	120.31
1	E	350	SER	C-N-CA	-5.39	112.12	120.31
1	P	287	ILE	N-CA-C	-5.39	105.12	110.62
1	R	330	ILE	CA-C-N	-5.39	113.70	122.65
1	R	330	ILE	C-N-CA	-5.39	113.70	122.65
1	T	194	THR	CA-C-N	5.39	128.04	120.28
1	T	194	THR	C-N-CA	5.39	128.04	120.28
1	U	350	SER	CA-C-N	-5.39	112.12	120.31
1	U	350	SER	C-N-CA	-5.39	112.12	120.31
1	B	162	ASN	CA-CB-CG	5.39	117.99	112.60
1	E	194	THR	CA-C-N	5.39	128.04	120.28
1	E	194	THR	C-N-CA	5.39	128.04	120.28
1	I	350	SER	CA-C-N	-5.39	112.12	120.31
1	I	350	SER	C-N-CA	-5.39	112.12	120.31
1	F	350	SER	CA-C-N	-5.38	112.13	120.31
1	F	350	SER	C-N-CA	-5.38	112.13	120.31
1	L	330	ILE	CA-C-N	-5.38	113.71	122.65
1	L	330	ILE	C-N-CA	-5.38	113.71	122.65
1	B	350	SER	CA-C-N	-5.38	112.13	120.31
1	B	350	SER	C-N-CA	-5.38	112.13	120.31
1	J	330	ILE	CA-C-N	-5.38	113.72	122.65
1	J	330	ILE	C-N-CA	-5.38	113.72	122.65
1	M	350	SER	CA-C-N	-5.38	112.13	120.31
1	M	350	SER	C-N-CA	-5.38	112.13	120.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	V	350	SER	CA-C-N	-5.38	112.13	120.31
1	V	350	SER	C-N-CA	-5.38	112.13	120.31
1	M	162	ASN	CA-CB-CG	5.38	117.98	112.60
1	E	162	ASN	CA-CB-CG	5.38	117.98	112.60
1	R	350	SER	CA-C-N	-5.38	112.13	120.31
1	R	350	SER	C-N-CA	-5.38	112.13	120.31
1	S	350	SER	CA-C-N	-5.38	112.13	120.31
1	S	350	SER	C-N-CA	-5.38	112.13	120.31
1	C	330	ILE	CA-C-N	-5.38	113.72	122.65
1	C	330	ILE	C-N-CA	-5.38	113.72	122.65
1	J	350	SER	CA-C-N	-5.37	112.14	120.31
1	J	350	SER	C-N-CA	-5.37	112.14	120.31
1	V	330	ILE	CA-C-N	-5.37	113.73	122.65
1	V	330	ILE	C-N-CA	-5.37	113.73	122.65
1	V	274	ILE	CA-C-O	-5.37	114.82	120.57
1	L	350	SER	CA-C-N	-5.37	112.15	120.31
1	L	350	SER	C-N-CA	-5.37	112.15	120.31
1	A	287	ILE	N-CA-C	-5.37	105.14	110.62
1	J	287	ILE	N-CA-C	-5.37	105.14	110.62
1	L	287	ILE	N-CA-C	-5.37	105.14	110.62
1	T	287	ILE	N-CA-C	-5.37	105.14	110.62
1	D	330	ILE	CA-C-N	-5.37	113.74	122.65
1	D	330	ILE	C-N-CA	-5.37	113.74	122.65
1	C	287	ILE	N-CA-C	-5.37	105.15	110.62
1	T	350	SER	CA-C-N	-5.37	112.16	120.31
1	T	350	SER	C-N-CA	-5.37	112.16	120.31
1	G	287	ILE	N-CA-C	-5.36	105.15	110.62
1	M	330	ILE	CA-C-N	-5.36	113.75	122.65
1	M	330	ILE	C-N-CA	-5.36	113.75	122.65
1	H	162	ASN	CA-CB-CG	5.36	117.96	112.60
1	K	162	ASN	CA-CB-CG	5.36	117.96	112.60
1	N	350	SER	CA-C-N	-5.36	112.17	120.31
1	N	350	SER	C-N-CA	-5.36	112.17	120.31
1	O	287	ILE	N-CA-C	-5.36	105.15	110.62
1	S	330	ILE	CA-C-N	-5.36	113.76	122.65
1	S	330	ILE	C-N-CA	-5.36	113.76	122.65
1	V	287	ILE	N-CA-C	-5.36	105.16	110.62
1	W	330	ILE	CA-C-N	-5.36	113.76	122.65
1	W	330	ILE	C-N-CA	-5.36	113.76	122.65
1	G	330	ILE	CA-C-N	-5.36	113.76	122.65
1	G	330	ILE	C-N-CA	-5.36	113.76	122.65
1	P	330	ILE	CA-C-N	-5.35	113.76	122.65

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	P	330	ILE	C-N-CA	-5.35	113.76	122.65
1	C	162	ASN	CA-CB-CG	5.35	117.95	112.60
1	K	330	ILE	CA-C-N	-5.35	113.77	122.65
1	K	330	ILE	C-N-CA	-5.35	113.77	122.65
1	F	162	ASN	CA-CB-CG	5.35	117.95	112.60
1	F	287	ILE	N-CA-C	-5.35	105.17	110.62
1	S	287	ILE	N-CA-C	-5.35	105.17	110.62
1	T	330	ILE	CA-C-N	-5.35	113.77	122.65
1	T	330	ILE	C-N-CA	-5.35	113.77	122.65
1	Q	287	ILE	N-CA-C	-5.35	105.17	110.62
1	B	330	ILE	CA-C-N	-5.34	113.78	122.65
1	B	330	ILE	C-N-CA	-5.34	113.78	122.65
1	H	330	ILE	CA-C-N	-5.34	113.78	122.65
1	H	330	ILE	C-N-CA	-5.34	113.78	122.65
1	I	330	ILE	CA-C-N	-5.34	113.78	122.65
1	I	330	ILE	C-N-CA	-5.34	113.78	122.65
1	U	330	ILE	CA-C-N	-5.34	113.78	122.65
1	U	330	ILE	C-N-CA	-5.34	113.78	122.65
1	Q	330	ILE	CA-C-N	-5.34	113.78	122.65
1	Q	330	ILE	C-N-CA	-5.34	113.78	122.65
1	A	162	ASN	CA-CB-CG	5.34	117.94	112.60
1	N	330	ILE	CA-C-N	-5.34	113.78	122.65
1	N	330	ILE	C-N-CA	-5.34	113.78	122.65
1	S	162	ASN	CA-CB-CG	5.34	117.94	112.60
1	Q	162	ASN	CA-CB-CG	5.34	117.94	112.60
1	D	287	ILE	N-CA-C	-5.33	105.18	110.62
1	E	287	ILE	N-CA-C	-5.33	105.18	110.62
1	O	330	ILE	CA-C-N	-5.33	113.80	122.65
1	O	330	ILE	C-N-CA	-5.33	113.80	122.65
1	E	330	ILE	CA-C-N	-5.33	113.80	122.65
1	E	330	ILE	C-N-CA	-5.33	113.80	122.65
1	U	287	ILE	N-CA-C	-5.33	105.18	110.62
1	O	162	ASN	CA-CB-CG	5.33	117.93	112.60
1	W	162	ASN	CA-CB-CG	5.33	117.93	112.60
1	F	330	ILE	CA-C-N	-5.33	113.81	122.65
1	F	330	ILE	C-N-CA	-5.33	113.81	122.65
1	H	287	ILE	N-CA-C	-5.33	105.19	110.62
1	J	162	ASN	CA-CB-CG	5.33	117.93	112.60
1	N	162	ASN	CA-CB-CG	5.33	117.92	112.60
1	V	162	ASN	CA-CB-CG	5.33	117.93	112.60
1	L	274	ILE	CA-C-O	-5.32	114.88	120.57
1	C	150	GLY	CA-C-N	-5.32	116.20	123.12

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	C	150	GLY	C-N-CA	-5.32	116.20	123.12
1	S	274	ILE	CA-C-O	-5.32	114.88	120.57
1	A	274	ILE	CA-C-O	-5.32	114.88	120.57
1	G	162	ASN	CA-CB-CG	5.32	117.92	112.60
1	B	287	ILE	N-CA-C	-5.32	105.20	110.62
1	K	287	ILE	N-CA-C	-5.32	105.20	110.62
1	Q	274	ILE	CA-C-O	-5.32	114.88	120.57
1	I	274	ILE	CA-C-O	-5.32	114.88	120.57
1	P	162	ASN	CA-CB-CG	5.32	117.92	112.60
1	W	150	GLY	CA-C-N	-5.32	116.21	123.12
1	W	150	GLY	C-N-CA	-5.32	116.21	123.12
1	W	274	ILE	CA-C-O	-5.32	114.88	120.57
1	M	150	GLY	CA-C-N	-5.31	116.21	123.12
1	M	150	GLY	C-N-CA	-5.31	116.21	123.12
1	M	287	ILE	N-CA-C	-5.31	105.20	110.62
1	O	150	GLY	CA-C-N	-5.31	116.21	123.12
1	O	150	GLY	C-N-CA	-5.31	116.21	123.12
1	D	162	ASN	CA-CB-CG	5.31	117.91	112.60
1	I	150	GLY	CA-C-N	-5.31	116.22	123.12
1	I	150	GLY	C-N-CA	-5.31	116.22	123.12
1	C	274	ILE	CA-C-O	-5.31	114.89	120.57
1	I	162	ASN	CA-CB-CG	5.31	117.91	112.60
1	K	274	ILE	CA-C-O	-5.31	114.89	120.57
1	M	274	ILE	CA-C-O	-5.31	114.89	120.57
1	T	162	ASN	CA-CB-CG	5.31	117.91	112.60
1	O	274	ILE	CA-C-O	-5.31	114.89	120.57
1	R	162	ASN	CA-CB-CG	5.31	117.91	112.60
1	I	287	ILE	N-CA-C	-5.31	105.21	110.62
1	F	150	GLY	CA-C-N	-5.30	116.22	123.12
1	F	150	GLY	C-N-CA	-5.30	116.22	123.12
1	T	150	GLY	CA-C-N	-5.30	116.22	123.12
1	T	150	GLY	C-N-CA	-5.30	116.22	123.12
1	U	150	GLY	CA-C-N	-5.30	116.22	123.12
1	U	150	GLY	C-N-CA	-5.30	116.22	123.12
1	K	150	GLY	CA-C-N	-5.30	116.23	123.12
1	K	150	GLY	C-N-CA	-5.30	116.23	123.12
1	E	150	GLY	CA-C-N	-5.30	116.23	123.12
1	E	150	GLY	C-N-CA	-5.30	116.23	123.12
1	N	287	ILE	N-CA-C	-5.30	105.21	110.62
1	V	150	GLY	CA-C-N	-5.30	116.23	123.12
1	V	150	GLY	C-N-CA	-5.30	116.23	123.12
1	R	287	ILE	N-CA-C	-5.30	105.21	110.62

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	H	274	ILE	CA-C-O	-5.30	114.90	120.57
1	J	150	GLY	CA-C-N	-5.30	116.23	123.12
1	J	150	GLY	C-N-CA	-5.30	116.23	123.12
1	R	274	ILE	CA-C-O	-5.30	114.90	120.57
1	S	150	GLY	CA-C-N	-5.30	116.23	123.12
1	S	150	GLY	C-N-CA	-5.30	116.23	123.12
1	A	150	GLY	CA-C-N	-5.30	116.23	123.12
1	A	150	GLY	C-N-CA	-5.30	116.23	123.12
1	H	150	GLY	CA-C-N	-5.29	116.24	123.12
1	H	150	GLY	C-N-CA	-5.29	116.24	123.12
1	D	150	GLY	CA-C-N	-5.29	116.24	123.12
1	D	150	GLY	C-N-CA	-5.29	116.24	123.12
1	L	162	ASN	CA-CB-CG	5.29	117.89	112.60
1	L	150	GLY	CA-C-N	-5.29	116.25	123.12
1	L	150	GLY	C-N-CA	-5.29	116.25	123.12
1	P	150	GLY	CA-C-N	-5.29	116.24	123.12
1	P	150	GLY	C-N-CA	-5.29	116.24	123.12
1	U	162	ASN	CA-CB-CG	5.29	117.89	112.60
1	B	274	ILE	CA-C-O	-5.29	114.91	120.57
1	N	150	GLY	CA-C-N	-5.29	116.25	123.12
1	N	150	GLY	C-N-CA	-5.29	116.25	123.12
1	J	274	ILE	CA-C-O	-5.28	114.92	120.57
1	R	150	GLY	CA-C-N	-5.28	116.25	123.12
1	R	150	GLY	C-N-CA	-5.28	116.25	123.12
1	B	150	GLY	CA-C-N	-5.28	116.26	123.12
1	B	150	GLY	C-N-CA	-5.28	116.26	123.12
1	G	150	GLY	CA-C-N	-5.28	116.25	123.12
1	G	150	GLY	C-N-CA	-5.28	116.25	123.12
1	P	274	ILE	CA-C-O	-5.28	114.92	120.57
1	N	274	ILE	CA-C-O	-5.28	114.92	120.57
1	T	274	ILE	CA-C-O	-5.28	114.92	120.57
1	U	274	ILE	CA-C-O	-5.27	114.93	120.57
1	E	274	ILE	CA-C-O	-5.27	114.93	120.57
1	D	274	ILE	CA-C-O	-5.27	114.93	120.57
1	F	274	ILE	CA-C-O	-5.27	114.93	120.57
1	Q	150	GLY	CA-C-N	-5.27	116.27	123.12
1	Q	150	GLY	C-N-CA	-5.27	116.27	123.12
1	G	274	ILE	CA-C-O	-5.24	114.96	120.57
1	K	354	GLN	CA-C-O	-5.14	113.06	119.38
1	J	354	GLN	CA-C-O	-5.13	113.07	119.38
1	R	354	GLN	CA-C-O	-5.13	113.07	119.38
1	A	354	GLN	CA-C-O	-5.12	113.08	119.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	H	354	GLN	CA-C-O	-5.12	113.08	119.38
1	D	354	GLN	CA-C-O	-5.12	113.08	119.38
1	M	354	GLN	CA-C-O	-5.12	113.08	119.38
1	C	354	GLN	CA-C-O	-5.12	113.08	119.38
1	S	354	GLN	CA-C-O	-5.12	113.09	119.38
1	T	354	GLN	CA-C-O	-5.11	113.09	119.38
1	U	354	GLN	CA-C-O	-5.11	113.09	119.38
1	F	354	GLN	CA-C-O	-5.11	113.10	119.38
1	O	354	GLN	CA-C-O	-5.10	113.11	119.38
1	E	354	GLN	CA-C-O	-5.10	113.11	119.38
1	I	354	GLN	CA-C-O	-5.10	113.11	119.38
1	V	354	GLN	CA-C-O	-5.10	113.11	119.38
1	L	354	GLN	CA-C-O	-5.09	113.12	119.38
1	P	354	GLN	CA-C-O	-5.09	113.12	119.38
1	B	354	GLN	CA-C-O	-5.08	113.12	119.38
1	N	354	GLN	CA-C-O	-5.08	113.12	119.38
1	G	354	GLN	CA-C-O	-5.08	113.13	119.38
1	W	354	GLN	CA-C-O	-5.08	113.13	119.38
1	Q	354	GLN	CA-C-O	-5.08	113.14	119.38
1	C	323	SER	N-CA-C	5.01	117.47	111.71
1	W	323	SER	N-CA-C	5.01	117.47	111.71
1	H	323	SER	N-CA-C	5.01	117.47	111.71
1	Q	323	SER	N-CA-C	5.01	117.47	111.71
1	G	323	SER	N-CA-C	5.00	117.47	111.71
1	P	323	SER	N-CA-C	5.00	117.46	111.71

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts (i)

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	2934	0	2895	344	0
1	B	2934	0	2895	350	0
1	C	2934	0	2895	386	0
1	D	2934	0	2895	383	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	E	2934	0	2895	391	0
1	F	2934	0	2895	385	0
1	G	2934	0	2895	384	0
1	H	2934	0	2895	381	0
1	I	2934	0	2895	385	0
1	J	2934	0	2895	385	0
1	K	2934	0	2895	381	0
1	L	2934	0	2895	390	0
1	M	2934	0	2895	383	0
1	N	2934	0	2895	387	0
1	O	2934	0	2895	384	0
1	P	2934	0	2895	380	0
1	Q	2934	0	2895	387	0
1	R	2934	0	2895	379	0
1	S	2934	0	2895	387	0
1	T	2934	0	2895	385	0
1	U	2934	0	2895	386	0
1	V	2934	0	2895	349	0
1	W	2934	0	2895	343	0
2	A	27	0	12	4	0
2	B	27	0	12	4	0
2	C	27	0	12	4	0
2	D	27	0	12	4	0
2	E	27	0	12	4	0
2	F	27	0	12	5	0
2	G	27	0	12	4	0
2	H	27	0	12	4	0
2	I	27	0	12	4	0
2	J	27	0	12	4	0
2	K	27	0	12	4	0
2	L	27	0	12	4	0
2	M	27	0	12	5	0
2	N	27	0	12	4	0
2	O	27	0	12	5	0
2	P	27	0	12	4	0
2	Q	27	0	12	4	0
2	R	27	0	12	4	0
2	S	27	0	12	4	0
2	T	27	0	12	4	0
2	U	27	0	12	4	0
2	V	27	0	12	4	0
2	W	27	0	12	4	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
All	All	68103	0	66861	7800	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 58.

All (7800) 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:F:290:ARG:HD2	1:H:244:ASP:HB2	1.22	1.20
1:D:290:ARG:HD2	1:F:244:ASP:HB2	1.21	1.20
1:H:290:ARG:HD2	1:J:244:ASP:HB2	1.21	1.20
1:G:59:GLN:O	1:G:62:ARG:HG3	1.43	1.19
1:N:59:GLN:O	1:N:62:ARG:HG3	1.43	1.19
1:U:59:GLN:O	1:U:62:ARG:HG3	1.43	1.19
1:P:59:GLN:O	1:P:62:ARG:HG3	1.43	1.19
1:I:59:GLN:O	1:I:62:ARG:HG3	1.43	1.18
1:W:59:GLN:O	1:W:62:ARG:HG3	1.43	1.18
1:E:59:GLN:O	1:E:62:ARG:HG3	1.43	1.18
1:L:59:GLN:O	1:L:62:ARG:HG3	1.43	1.18
1:S:59:GLN:O	1:S:62:ARG:HG3	1.43	1.18
1:B:290:ARG:HD2	1:D:244:ASP:HB2	1.22	1.18
1:B:59:GLN:O	1:B:62:ARG:HG3	1.43	1.17
1:F:59:GLN:O	1:F:62:ARG:HG3	1.43	1.17
1:J:59:GLN:O	1:J:62:ARG:HG3	1.43	1.17
1:J:290:ARG:HD2	1:L:244:ASP:HB2	1.21	1.17
1:D:59:GLN:O	1:D:62:ARG:HG3	1.43	1.17
1:R:59:GLN:O	1:R:62:ARG:HG3	1.43	1.17
1:K:59:GLN:O	1:K:62:ARG:HG3	1.43	1.17
1:Q:59:GLN:O	1:Q:62:ARG:HG3	1.43	1.16
1:V:59:GLN:O	1:V:62:ARG:HG3	1.43	1.16
1:C:59:GLN:O	1:C:62:ARG:HG3	1.43	1.16
1:T:59:GLN:O	1:T:62:ARG:HG3	1.43	1.16
1:K:361:GLU:HB3	1:K:369:ILE:HD13	1.19	1.16
1:A:59:GLN:O	1:A:62:ARG:HG3	1.43	1.15
1:H:59:GLN:O	1:H:62:ARG:HG3	1.43	1.15
1:M:59:GLN:O	1:M:62:ARG:HG3	1.43	1.15
1:L:290:ARG:HD2	1:N:244:ASP:HB2	1.21	1.14
1:O:59:GLN:O	1:O:62:ARG:HG3	1.43	1.14
1:O:361:GLU:HB3	1:O:369:ILE:HD13	1.19	1.14
1:N:34:ILE:HB	1:N:54:VAL:HG11	1.30	1.14
1:A:34:ILE:HB	1:A:54:VAL:HG11	1.30	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:361:GLU:HB3	1:D:369:ILE:HD13	1.19	1.13
1:V:361:GLU:HB3	1:V:369:ILE:HD13	1.19	1.13
1:C:34:ILE:HB	1:C:54:VAL:HG11	1.30	1.13
1:J:43:VAL:HG13	1:J:44:MET:N	1.54	1.13
1:L:43:VAL:HG13	1:L:44:MET:N	1.54	1.13
1:Q:361:GLU:HB3	1:Q:369:ILE:HD13	1.19	1.13
1:S:357:ILE:HG12	1:S:370:VAL:HG23	1.13	1.13
1:D:357:ILE:HG12	1:D:370:VAL:HG23	1.13	1.13
1:I:357:ILE:HG12	1:I:370:VAL:HG23	1.13	1.13
1:M:357:ILE:HG12	1:M:370:VAL:HG23	1.13	1.13
1:F:357:ILE:HG12	1:F:370:VAL:HG23	1.13	1.12
1:H:43:VAL:HG13	1:H:44:MET:N	1.54	1.13
1:O:357:ILE:HG12	1:O:370:VAL:HG23	1.13	1.12
1:P:34:ILE:HB	1:P:54:VAL:HG11	1.30	1.13
1:W:43:VAL:HG13	1:W:44:MET:N	1.54	1.13
1:U:357:ILE:HG12	1:U:370:VAL:HG23	1.13	1.12
1:G:361:GLU:HB3	1:G:369:ILE:HD13	1.19	1.12
1:H:357:ILE:HG12	1:H:370:VAL:HG23	1.13	1.12
1:V:357:ILE:HG12	1:V:370:VAL:HG23	1.13	1.12
1:K:290:ARG:HD2	1:M:244:ASP:HB2	1.21	1.12
1:M:290:ARG:HD2	1:O:244:ASP:HB2	1.21	1.12
1:N:43:VAL:HG13	1:N:44:MET:N	1.54	1.12
1:L:34:ILE:HB	1:L:54:VAL:HG11	1.30	1.12
1:M:361:GLU:HB3	1:M:369:ILE:HD13	1.19	1.12
1:T:34:ILE:HB	1:T:54:VAL:HG11	1.30	1.12
1:T:357:ILE:HG12	1:T:370:VAL:HG23	1.13	1.12
1:W:357:ILE:HG12	1:W:370:VAL:HG23	1.13	1.12
1:N:361:GLU:HB3	1:N:369:ILE:HD13	1.19	1.12
1:A:43:VAL:HG13	1:A:44:MET:N	1.54	1.11
1:G:34:ILE:HB	1:G:54:VAL:HG11	1.30	1.11
1:G:357:ILE:HG12	1:G:370:VAL:HG23	1.13	1.11
1:U:43:VAL:HG13	1:U:44:MET:N	1.54	1.11
1:B:357:ILE:HG12	1:B:370:VAL:HG23	1.13	1.11
1:E:357:ILE:HG12	1:E:370:VAL:HG23	1.13	1.11
1:F:43:VAL:HG13	1:F:44:MET:N	1.54	1.11
1:I:290:ARG:HD2	1:K:244:ASP:HB2	1.21	1.11
1:J:357:ILE:HG12	1:J:370:VAL:HG23	1.13	1.11
1:K:357:ILE:HG12	1:K:370:VAL:HG23	1.13	1.11
1:L:357:ILE:HG12	1:L:370:VAL:HG23	1.13	1.11
1:N:290:ARG:HD2	1:P:244:ASP:HB2	1.21	1.11
1:O:290:ARG:HD2	1:Q:244:ASP:HB2	1.21	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:34:ILE:HB	1:E:54:VAL:HG11	1.30	1.11
1:R:34:ILE:HB	1:R:54:VAL:HG11	1.30	1.11
1:U:290:ARG:HD2	1:W:244:ASP:HB2	1.21	1.11
1:Q:357:ILE:HG12	1:Q:370:VAL:HG23	1.13	1.11
1:A:361:GLU:HB3	1:A:369:ILE:HD13	1.19	1.11
1:P:357:ILE:HG12	1:P:370:VAL:HG23	1.13	1.11
1:A:357:ILE:HG12	1:A:370:VAL:HG23	1.13	1.10
1:F:361:GLU:HB3	1:F:369:ILE:HD13	1.20	1.10
1:G:290:ARG:HD2	1:I:244:ASP:HB2	1.22	1.10
1:Q:180:LEU:HD21	1:Q:261:LEU:HD23	1.34	1.10
1:C:43:VAL:HG13	1:C:44:MET:N	1.54	1.10
1:E:180:LEU:HD21	1:E:261:LEU:HD23	1.34	1.10
1:J:34:ILE:HB	1:J:54:VAL:HG11	1.30	1.10
1:J:361:GLU:HB3	1:J:369:ILE:HD13	1.19	1.10
1:P:43:VAL:HG13	1:P:44:MET:N	1.54	1.10
1:R:180:LEU:HD21	1:R:261:LEU:HD23	1.34	1.10
1:A:290:ARG:HD2	1:C:244:ASP:HB2	1.21	1.10
1:D:180:LEU:HD21	1:D:261:LEU:HD23	1.34	1.10
1:E:290:ARG:HD2	1:G:244:ASP:HB2	1.21	1.10
1:P:361:GLU:HB3	1:P:369:ILE:HD13	1.19	1.10
1:Q:290:ARG:HD2	1:S:244:ASP:HB2	1.21	1.10
1:U:361:GLU:HB3	1:U:369:ILE:HD13	1.19	1.10
1:V:34:ILE:HB	1:V:54:VAL:HG11	1.30	1.10
1:W:361:GLU:HB3	1:W:369:ILE:HD13	1.19	1.10
1:C:290:ARG:HD2	1:E:244:ASP:HB2	1.21	1.10
1:F:180:LEU:HD21	1:F:261:LEU:HD23	1.34	1.10
1:I:34:ILE:HB	1:I:54:VAL:HG11	1.30	1.10
1:L:361:GLU:HB3	1:L:369:ILE:HD13	1.19	1.10
1:R:357:ILE:HG12	1:R:370:VAL:HG23	1.13	1.10
1:S:290:ARG:HD2	1:U:244:ASP:HB2	1.21	1.10
1:T:180:LEU:HD21	1:T:261:LEU:HD23	1.33	1.10
1:B:361:GLU:HB3	1:B:369:ILE:HD13	1.19	1.10
1:G:180:LEU:HD21	1:G:261:LEU:HD23	1.34	1.10
1:H:34:ILE:HB	1:H:54:VAL:HG11	1.30	1.10
1:N:357:ILE:HG12	1:N:370:VAL:HG23	1.13	1.10
1:S:43:VAL:HG13	1:S:44:MET:N	1.54	1.10
1:S:180:LEU:HD21	1:S:261:LEU:HD23	1.34	1.10
1:C:361:GLU:HB3	1:C:369:ILE:HD13	1.19	1.09
1:D:43:VAL:HG13	1:D:44:MET:N	1.54	1.09
1:J:208:ILE:HD11	1:J:243:PRO:HD2	1.34	1.09
1:K:180:LEU:HD21	1:K:261:LEU:HD23	1.34	1.09

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:361:GLU:HB3	1:R:369:ILE:HD13	1.19	1.09
1:B:208:ILE:HD11	1:B:243:PRO:HD2	1.34	1.09
1:M:180:LEU:HD21	1:M:261:LEU:HD23	1.34	1.09
1:O:208:ILE:HD11	1:O:243:PRO:HD2	1.34	1.09
1:P:208:ILE:HD11	1:P:243:PRO:HD2	1.34	1.09
1:V:180:LEU:HD21	1:V:261:LEU:HD23	1.34	1.09
1:W:34:ILE:HB	1:W:54:VAL:HG11	1.30	1.09
1:D:208:ILE:HD11	1:D:243:PRO:HD2	1.34	1.09
1:L:43:VAL:HG13	1:L:44:MET:H	0.92	1.09
1:M:208:ILE:HD11	1:M:243:PRO:HD2	1.34	1.09
1:O:180:LEU:HD21	1:O:261:LEU:HD23	1.34	1.09
1:U:34:ILE:HB	1:U:54:VAL:HG11	1.30	1.09
1:U:208:ILE:HD11	1:U:243:PRO:HD2	1.34	1.09
1:C:180:LEU:HD21	1:C:261:LEU:HD23	1.34	1.09
1:C:357:ILE:HG12	1:C:370:VAL:HG23	1.13	1.09
1:E:361:GLU:HB3	1:E:369:ILE:HD13	1.19	1.09
1:P:180:LEU:HD21	1:P:261:LEU:HD23	1.34	1.09
1:S:208:ILE:HD11	1:S:243:PRO:HD2	1.34	1.09
1:A:43:VAL:HG13	1:A:44:MET:H	0.92	1.09
1:A:180:LEU:HD21	1:A:261:LEU:HD23	1.34	1.09
1:C:43:VAL:HG13	1:C:44:MET:H	0.92	1.08
1:E:43:VAL:HG13	1:E:44:MET:N	1.54	1.08
1:H:361:GLU:HB3	1:H:369:ILE:HD13	1.19	1.08
1:I:208:ILE:HD11	1:I:243:PRO:HD2	1.34	1.08
1:L:180:LEU:HD21	1:L:261:LEU:HD23	1.34	1.08
1:S:361:GLU:HB3	1:S:369:ILE:HD13	1.19	1.08
1:U:43:VAL:HG13	1:U:44:MET:H	0.92	1.08
1:W:180:LEU:HD21	1:W:261:LEU:HD23	1.34	1.08
1:I:180:LEU:HD21	1:I:261:LEU:HD23	1.34	1.08
1:J:180:LEU:HD21	1:J:261:LEU:HD23	1.34	1.08
1:N:208:ILE:HD11	1:N:243:PRO:HD2	1.34	1.08
1:R:43:VAL:HG13	1:R:44:MET:N	1.54	1.08
1:T:43:VAL:HG13	1:T:44:MET:H	0.92	1.08
1:T:208:ILE:HD11	1:T:243:PRO:HD2	1.34	1.08
1:W:43:VAL:HG13	1:W:44:MET:H	0.92	1.08
1:B:180:LEU:HD21	1:B:261:LEU:HD23	1.34	1.08
1:C:208:ILE:HD11	1:C:243:PRO:HD2	1.34	1.08
1:E:208:ILE:HD11	1:E:243:PRO:HD2	1.34	1.08
1:H:180:LEU:HD21	1:H:261:LEU:HD23	1.34	1.08
1:H:208:ILE:HD11	1:H:243:PRO:HD2	1.34	1.08
1:J:43:VAL:HG13	1:J:44:MET:H	0.92	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:290:ARG:HD2	1:R:244:ASP:HB2	1.21	1.08
1:I:361:GLU:HB3	1:I:369:ILE:HD13	1.19	1.08
1:N:180:LEU:HD21	1:N:261:LEU:HD23	1.34	1.08
1:O:35:VAL:HG12	1:O:68:LYS:HB2	1.36	1.08
1:H:35:VAL:HG12	1:H:68:LYS:HB2	1.36	1.08
1:R:43:VAL:HG13	1:R:44:MET:H	0.92	1.08
1:S:35:VAL:HG12	1:S:68:LYS:HB2	1.36	1.08
1:U:180:LEU:HD21	1:U:261:LEU:HD23	1.34	1.08
1:B:43:VAL:HG13	1:B:44:MET:N	1.54	1.07
1:D:35:VAL:HG12	1:D:68:LYS:HB2	1.36	1.07
1:N:43:VAL:HG13	1:N:44:MET:H	0.92	1.07
1:Q:34:ILE:HB	1:Q:54:VAL:HG11	1.30	1.07
1:D:43:VAL:HG13	1:D:44:MET:H	0.92	1.07
1:F:34:ILE:HB	1:F:54:VAL:HG11	1.30	1.07
1:Q:43:VAL:HG13	1:Q:44:MET:N	1.54	1.07
1:T:290:ARG:HD2	1:V:244:ASP:HB2	1.21	1.07
1:T:361:GLU:HB3	1:T:369:ILE:HD13	1.20	1.07
1:G:43:VAL:HG13	1:G:44:MET:N	1.54	1.07
1:K:43:VAL:HG13	1:K:44:MET:H	0.92	1.07
1:R:290:ARG:HD2	1:T:244:ASP:HB2	1.21	1.07
1:S:34:ILE:HB	1:S:54:VAL:HG11	1.30	1.07
1:V:43:VAL:HG13	1:V:44:MET:H	0.92	1.07
1:B:43:VAL:HG13	1:B:44:MET:H	0.92	1.07
1:I:35:VAL:HG12	1:I:68:LYS:HB2	1.36	1.07
1:K:35:VAL:HG12	1:K:68:LYS:HB2	1.36	1.07
1:O:34:ILE:HB	1:O:54:VAL:HG11	1.30	1.07
1:S:43:VAL:HG13	1:S:44:MET:H	0.92	1.07
1:I:43:VAL:HG13	1:I:44:MET:H	0.92	1.07
1:V:35:VAL:HG12	1:V:68:LYS:HB2	1.36	1.07
1:K:34:ILE:HB	1:K:54:VAL:HG11	1.30	1.06
1:A:208:ILE:HD11	1:A:243:PRO:HD2	1.34	1.06
1:B:34:ILE:HB	1:B:54:VAL:HG11	1.30	1.06
1:L:208:ILE:HD11	1:L:243:PRO:HD2	1.34	1.06
1:M:43:VAL:HG13	1:M:44:MET:H	0.92	1.06
1:E:43:VAL:HG13	1:E:44:MET:H	0.92	1.06
1:F:43:VAL:HG13	1:F:44:MET:H	0.92	1.06
1:M:34:ILE:HB	1:M:54:VAL:HG11	1.30	1.06
1:T:35:VAL:HG12	1:T:68:LYS:HB2	1.36	1.06
1:U:35:VAL:HG12	1:U:68:LYS:HB2	1.36	1.06
1:T:43:VAL:HG13	1:T:44:MET:N	1.54	1.06
1:B:35:VAL:HG12	1:B:68:LYS:HB2	1.36	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:43:VAL:HG13	1:I:44:MET:N	1.54	1.06
1:O:43:VAL:HG13	1:O:44:MET:N	1.54	1.06
1:J:35:VAL:HG12	1:J:68:LYS:HB2	1.36	1.05
1:N:35:VAL:HG12	1:N:68:LYS:HB2	1.36	1.05
1:G:43:VAL:HG13	1:G:44:MET:H	0.92	1.05
1:H:43:VAL:HG13	1:H:44:MET:H	0.92	1.05
1:K:37:ARG:HG3	1:K:38:PRO:HD2	1.37	1.05
1:P:34:ILE:HB	1:P:54:VAL:CG1	1.87	1.05
1:V:208:ILE:HD11	1:V:243:PRO:HD2	1.34	1.05
1:W:208:ILE:HD11	1:W:243:PRO:HD2	1.34	1.05
1:D:34:ILE:HB	1:D:54:VAL:HG11	1.30	1.05
1:E:35:VAL:HG12	1:E:68:LYS:HB2	1.36	1.05
1:N:34:ILE:HB	1:N:54:VAL:CG1	1.87	1.05
1:R:34:ILE:HB	1:R:54:VAL:CG1	1.87	1.05
1:C:35:VAL:HG12	1:C:68:LYS:HB2	1.36	1.05
1:I:37:ARG:HG3	1:I:38:PRO:HD2	1.37	1.05
1:K:43:VAL:HG13	1:K:44:MET:N	1.54	1.05
1:K:208:ILE:HD11	1:K:243:PRO:HD2	1.34	1.05
1:L:34:ILE:HB	1:L:54:VAL:CG1	1.87	1.05
1:L:37:ARG:HG3	1:L:38:PRO:HD2	1.37	1.05
1:M:43:VAL:HG13	1:M:44:MET:N	1.54	1.05
1:O:43:VAL:HG13	1:O:44:MET:H	0.92	1.05
1:Q:208:ILE:HD11	1:Q:243:PRO:HD2	1.34	1.05
1:F:208:ILE:HD11	1:F:243:PRO:HD2	1.34	1.05
1:G:208:ILE:HD11	1:G:243:PRO:HD2	1.34	1.05
1:M:35:VAL:HG12	1:M:68:LYS:HB2	1.36	1.05
1:M:37:ARG:HG3	1:M:38:PRO:HD2	1.37	1.05
1:P:43:VAL:HG13	1:P:44:MET:H	0.92	1.05
1:T:34:ILE:HB	1:T:54:VAL:CG1	1.87	1.05
1:U:34:ILE:HB	1:U:54:VAL:CG1	1.87	1.05
1:W:35:VAL:HG12	1:W:68:LYS:HB2	1.36	1.05
1:J:34:ILE:HB	1:J:54:VAL:CG1	1.87	1.04
1:J:37:ARG:HG3	1:J:38:PRO:HD2	1.37	1.04
1:S:34:ILE:HB	1:S:54:VAL:CG1	1.87	1.04
1:V:43:VAL:HG13	1:V:44:MET:N	1.54	1.04
1:W:34:ILE:HB	1:W:54:VAL:CG1	1.87	1.04
1:G:34:ILE:HB	1:G:54:VAL:CG1	1.87	1.04
1:M:34:ILE:HG21	1:M:67:LEU:HD22	1.40	1.04
1:N:37:ARG:HG3	1:N:38:PRO:HD2	1.37	1.04
1:P:34:ILE:HG21	1:P:67:LEU:HD22	1.40	1.04
1:U:37:ARG:HG3	1:U:38:PRO:HD2	1.37	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:34:ILE:HG21	1:V:67:LEU:HD22	1.40	1.04
1:A:34:ILE:HB	1:A:54:VAL:CG1	1.87	1.04
1:C:34:ILE:HG21	1:C:67:LEU:HD22	1.40	1.04
1:G:34:ILE:HG21	1:G:67:LEU:HD22	1.40	1.04
1:G:37:ARG:HG3	1:G:38:PRO:HD2	1.37	1.04
1:H:37:ARG:HG3	1:H:38:PRO:HD2	1.37	1.04
1:I:34:ILE:HB	1:I:54:VAL:CG1	1.87	1.04
1:I:34:ILE:HG21	1:I:67:LEU:HD22	1.40	1.04
1:P:35:VAL:HG12	1:P:68:LYS:HB2	1.36	1.04
1:Q:34:ILE:HB	1:Q:54:VAL:CG1	1.87	1.04
1:R:34:ILE:HG21	1:R:67:LEU:HD22	1.40	1.04
1:U:45:VAL:O	1:U:45:VAL:HG22	1.57	1.04
1:W:37:ARG:HG3	1:W:38:PRO:HD2	1.37	1.04
1:A:45:VAL:HG22	1:A:45:VAL:O	1.57	1.04
1:E:34:ILE:HB	1:E:54:VAL:CG1	1.87	1.04
1:F:37:ARG:HG3	1:F:38:PRO:HD2	1.37	1.04
1:J:45:VAL:O	1:J:45:VAL:HG22	1.57	1.04
1:K:34:ILE:HB	1:K:54:VAL:CG1	1.87	1.04
1:L:35:VAL:HG12	1:L:68:LYS:HB2	1.36	1.04
1:A:34:ILE:HG21	1:A:67:LEU:HD22	1.40	1.04
1:A:37:ARG:HG3	1:A:38:PRO:HD2	1.37	1.04
1:H:34:ILE:HB	1:H:54:VAL:CG1	1.87	1.04
1:M:34:ILE:HB	1:M:54:VAL:CG1	1.87	1.04
1:O:34:ILE:HB	1:O:54:VAL:CG1	1.87	1.04
1:O:34:ILE:HG21	1:O:67:LEU:HD22	1.40	1.04
1:R:208:ILE:HD11	1:R:243:PRO:HD2	1.34	1.04
1:S:37:ARG:HG3	1:S:38:PRO:HD2	1.37	1.04
1:V:34:ILE:HB	1:V:54:VAL:CG1	1.87	1.04
1:K:34:ILE:HG21	1:K:67:LEU:HD22	1.40	1.03
1:N:45:VAL:HG22	1:N:45:VAL:O	1.57	1.03
1:O:37:ARG:HG3	1:O:38:PRO:HD2	1.37	1.03
1:Q:43:VAL:HG13	1:Q:44:MET:H	0.92	1.03
1:Q:45:VAL:HG22	1:Q:45:VAL:O	1.57	1.03
1:T:34:ILE:HG21	1:T:67:LEU:HD22	1.40	1.03
1:B:34:ILE:HG21	1:B:67:LEU:HD22	1.40	1.03
1:C:34:ILE:HB	1:C:54:VAL:CG1	1.87	1.03
1:F:35:VAL:HG12	1:F:68:LYS:HB2	1.36	1.03
1:H:45:VAL:O	1:H:45:VAL:HG22	1.57	1.03
1:Q:35:VAL:HG12	1:Q:68:LYS:HB2	1.36	1.03
1:E:34:ILE:HG21	1:E:67:LEU:HD22	1.40	1.03
1:N:34:ILE:HG21	1:N:67:LEU:HD22	1.40	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:34:ILE:HG21	1:D:67:LEU:HD22	1.40	1.03
1:F:34:ILE:HB	1:F:54:VAL:CG1	1.87	1.03
1:D:45:VAL:HG22	1:D:45:VAL:O	1.57	1.03
1:E:37:ARG:HG3	1:E:38:PRO:HD2	1.37	1.03
1:F:45:VAL:HG22	1:F:45:VAL:O	1.57	1.03
1:W:45:VAL:O	1:W:45:VAL:HG22	1.57	1.03
1:B:34:ILE:HB	1:B:54:VAL:CG1	1.87	1.02
1:F:34:ILE:HG21	1:F:67:LEU:HD22	1.40	1.02
1:A:35:VAL:HG12	1:A:68:LYS:HB2	1.36	1.02
1:D:34:ILE:HB	1:D:54:VAL:CG1	1.87	1.02
1:D:37:ARG:HG3	1:D:38:PRO:HD2	1.37	1.02
1:G:35:VAL:HG12	1:G:68:LYS:HB2	1.36	1.02
1:H:34:ILE:HG21	1:H:67:LEU:HD22	1.40	1.02
1:J:34:ILE:HG21	1:J:67:LEU:HD22	1.40	1.02
1:P:37:ARG:HG3	1:P:38:PRO:HD2	1.37	1.02
1:S:34:ILE:HG21	1:S:67:LEU:HD22	1.40	1.02
1:W:34:ILE:HG21	1:W:67:LEU:HD22	1.40	1.02
1:B:45:VAL:O	1:B:45:VAL:HG22	1.57	1.02
1:L:45:VAL:HG22	1:L:45:VAL:O	1.57	1.02
1:Q:34:ILE:HG21	1:Q:67:LEU:HD22	1.40	1.02
1:U:34:ILE:HG21	1:U:67:LEU:HD22	1.40	1.02
1:L:34:ILE:HG21	1:L:67:LEU:HD22	1.40	1.02
1:Q:37:ARG:HG3	1:Q:38:PRO:HD2	1.37	1.02
1:P:45:VAL:O	1:P:45:VAL:HG22	1.57	1.01
1:R:35:VAL:HG12	1:R:68:LYS:HB2	1.36	1.01
1:C:37:ARG:HG3	1:C:38:PRO:HD2	1.37	1.01
1:S:45:VAL:O	1:S:45:VAL:HG22	1.57	1.01
1:T:37:ARG:HG3	1:T:38:PRO:HD2	1.37	1.01
1:T:45:VAL:HG22	1:T:45:VAL:O	1.57	1.01
1:V:37:ARG:HG3	1:V:38:PRO:HD2	1.37	1.01
1:D:43:VAL:CG1	1:D:44:MET:H	1.74	1.01
1:K:45:VAL:O	1:K:45:VAL:HG22	1.57	1.01
1:O:45:VAL:O	1:O:45:VAL:HG22	1.57	1.01
1:R:37:ARG:HG3	1:R:38:PRO:HD2	1.37	1.01
1:B:43:VAL:CG1	1:B:44:MET:H	1.74	1.00
1:D:208:ILE:HD11	1:D:243:PRO:CD	1.92	1.00
1:E:45:VAL:O	1:E:45:VAL:HG22	1.57	1.00
1:F:43:VAL:CG1	1:F:44:MET:H	1.74	1.00
1:I:8:LEU:HD12	1:I:90:PHE:HE1	1.26	1.00
1:I:43:VAL:CG1	1:I:44:MET:H	1.74	1.00
1:A:208:ILE:HD11	1:A:243:PRO:CD	1.92	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:208:ILE:HD11	1:B:243:PRO:CD	1.92	1.00
1:F:208:ILE:HD11	1:F:243:PRO:CD	1.92	1.00
1:G:45:VAL:O	1:G:45:VAL:HG22	1.57	1.00
1:M:208:ILE:HD11	1:M:243:PRO:CD	1.92	1.00
1:S:8:LEU:HD12	1:S:90:PHE:HE1	1.26	1.00
1:C:208:ILE:HD11	1:C:243:PRO:CD	1.92	1.00
1:H:43:VAL:CG1	1:H:44:MET:H	1.74	1.00
1:K:208:ILE:HD11	1:K:243:PRO:CD	1.92	1.00
1:O:8:LEU:HD12	1:O:90:PHE:HE1	1.27	1.00
1:R:43:VAL:CG1	1:R:44:MET:H	1.74	1.00
1:T:208:ILE:HD11	1:T:243:PRO:CD	1.92	1.00
1:H:208:ILE:HD11	1:H:243:PRO:CD	1.92	1.00
1:O:208:ILE:HD11	1:O:243:PRO:CD	1.92	1.00
1:R:208:ILE:HD11	1:R:243:PRO:CD	1.92	1.00
1:B:37:ARG:HG3	1:B:38:PRO:HD2	1.37	1.00
1:D:8:LEU:HD12	1:D:90:PHE:HE1	1.27	1.00
1:E:8:LEU:HD12	1:E:90:PHE:HE1	1.27	1.00
1:T:8:LEU:HD12	1:T:90:PHE:HE1	1.27	1.00
1:V:208:ILE:HD11	1:V:243:PRO:CD	1.92	1.00
1:A:43:VAL:CG1	1:A:44:MET:H	1.74	1.00
1:H:8:LEU:HD12	1:H:90:PHE:HE1	1.27	1.00
1:C:45:VAL:O	1:C:45:VAL:HG22	1.57	1.00
1:M:45:VAL:O	1:M:45:VAL:HG22	1.57	0.99
1:W:208:ILE:HD11	1:W:243:PRO:CD	1.92	0.99
1:B:8:LEU:HD12	1:B:90:PHE:HE1	1.27	0.99
1:G:43:VAL:CG1	1:G:44:MET:H	1.74	0.99
1:J:208:ILE:HD11	1:J:243:PRO:CD	1.92	0.99
1:P:8:LEU:HD12	1:P:90:PHE:HE1	1.27	0.99
1:I:208:ILE:HD11	1:I:243:PRO:CD	1.92	0.99
1:R:45:VAL:O	1:R:45:VAL:HG22	1.57	0.99
1:V:45:VAL:HG22	1:V:45:VAL:O	1.57	0.99
1:E:208:ILE:HD11	1:E:243:PRO:CD	1.92	0.99
1:I:45:VAL:HG22	1:I:45:VAL:O	1.57	0.99
1:J:43:VAL:CG1	1:J:44:MET:H	1.74	0.99
1:P:208:ILE:HD11	1:P:243:PRO:CD	1.92	0.99
1:Q:208:ILE:HD11	1:Q:243:PRO:CD	1.92	0.99
1:P:43:VAL:CG1	1:P:44:MET:H	1.74	0.99
1:U:208:ILE:HD11	1:U:243:PRO:CD	1.92	0.99
1:L:8:LEU:HD12	1:L:90:PHE:HE1	1.27	0.99
1:L:208:ILE:HD11	1:L:243:PRO:CD	1.92	0.99
1:M:8:LEU:HD12	1:M:90:PHE:HE1	1.27	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:8:LEU:HD12	1:V:90:PHE:HE1	1.27	0.99
1:W:8:LEU:HD12	1:W:90:PHE:HE1	1.27	0.99
1:W:43:VAL:CG1	1:W:44:MET:H	1.74	0.99
1:G:208:ILE:HD11	1:G:243:PRO:CD	1.92	0.98
1:N:208:ILE:HD11	1:N:243:PRO:CD	1.92	0.98
1:S:208:ILE:HD11	1:S:243:PRO:CD	1.92	0.98
1:A:8:LEU:HD12	1:A:90:PHE:HE1	1.27	0.98
1:U:43:VAL:CG1	1:U:44:MET:H	1.74	0.98
1:K:8:LEU:HD12	1:K:90:PHE:HE1	1.27	0.98
1:L:43:VAL:CG1	1:L:44:MET:H	1.74	0.98
1:N:43:VAL:CG1	1:N:44:MET:H	1.74	0.98
1:E:43:VAL:CG1	1:E:44:MET:H	1.74	0.98
1:F:8:LEU:HD12	1:F:90:PHE:HE1	1.27	0.98
1:R:8:LEU:HD12	1:R:90:PHE:HE1	1.27	0.97
1:S:43:VAL:CG1	1:S:44:MET:H	1.74	0.97
1:C:8:LEU:HD12	1:C:90:PHE:HE1	1.27	0.97
1:N:8:LEU:HD12	1:N:90:PHE:HE1	1.27	0.97
1:U:8:LEU:HD12	1:U:90:PHE:HE1	1.27	0.97
1:Q:8:LEU:HD12	1:Q:90:PHE:HE1	1.27	0.97
1:G:8:LEU:HD12	1:G:90:PHE:HE1	1.27	0.96
1:Q:43:VAL:CG1	1:Q:44:MET:H	1.74	0.96
1:C:43:VAL:CG1	1:C:44:MET:H	1.74	0.96
1:H:300:SER:HA	1:H:335:ARG:CG	1.96	0.96
1:W:300:SER:HA	1:W:335:ARG:CG	1.96	0.96
1:A:300:SER:HA	1:A:335:ARG:CG	1.96	0.96
1:L:300:SER:HA	1:L:335:ARG:CG	1.96	0.96
1:N:300:SER:HA	1:N:335:ARG:CG	1.96	0.96
1:S:300:SER:HA	1:S:335:ARG:CG	1.96	0.96
1:J:8:LEU:HD12	1:J:90:PHE:HE1	1.27	0.95
1:R:300:SER:HA	1:R:335:ARG:CG	1.96	0.95
1:P:300:SER:HA	1:P:335:ARG:CG	1.96	0.95
1:C:300:SER:HA	1:C:335:ARG:CG	1.96	0.95
1:J:300:SER:HA	1:J:335:ARG:CG	1.96	0.95
1:I:300:SER:HA	1:I:335:ARG:CG	1.96	0.95
1:K:300:SER:HA	1:K:335:ARG:CG	1.96	0.95
1:E:300:SER:HA	1:E:335:ARG:CG	1.96	0.95
1:G:300:SER:HA	1:G:335:ARG:CG	1.96	0.95
1:B:300:SER:HA	1:B:335:ARG:CG	1.96	0.95
1:O:43:VAL:CG1	1:O:44:MET:H	1.74	0.95
1:U:300:SER:HA	1:U:335:ARG:CG	1.96	0.95
1:V:43:VAL:CG1	1:V:44:MET:H	1.74	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:39:ARG:HE	1:C:66:THR:HB	1.32	0.94
1:T:300:SER:HA	1:T:335:ARG:CG	1.96	0.94
1:D:300:SER:HA	1:D:335:ARG:CG	1.96	0.94
1:E:39:ARG:HE	1:E:66:THR:HB	1.32	0.94
1:Q:300:SER:HA	1:Q:335:ARG:CG	1.96	0.94
1:V:39:ARG:HE	1:V:66:THR:HB	1.32	0.94
1:O:300:SER:HA	1:O:335:ARG:CG	1.96	0.94
1:F:43:VAL:CG1	1:F:44:MET:N	2.30	0.94
1:L:133:TYR:CZ	1:L:375:PHE:HB2	2.03	0.94
1:N:133:TYR:CZ	1:N:375:PHE:HB2	2.03	0.94
1:M:300:SER:HA	1:M:335:ARG:CG	1.96	0.94
1:V:300:SER:HA	1:V:335:ARG:CG	1.96	0.94
1:A:39:ARG:HE	1:A:66:THR:HB	1.32	0.94
1:C:133:TYR:CZ	1:C:375:PHE:HB2	2.03	0.94
1:F:300:SER:HA	1:F:335:ARG:CG	1.96	0.94
1:J:133:TYR:CZ	1:J:375:PHE:HB2	2.03	0.94
1:A:133:TYR:CZ	1:A:375:PHE:HB2	2.03	0.94
1:L:43:VAL:CG1	1:L:44:MET:N	2.30	0.94
1:O:133:TYR:CZ	1:O:375:PHE:HB2	2.03	0.94
1:P:133:TYR:CZ	1:P:375:PHE:HB2	2.03	0.93
1:T:39:ARG:HE	1:T:66:THR:HB	1.32	0.93
1:W:133:TYR:CZ	1:W:375:PHE:HB2	2.03	0.93
1:E:133:TYR:CZ	1:E:375:PHE:HB2	2.03	0.93
1:M:133:TYR:CZ	1:M:375:PHE:HB2	2.03	0.93
1:U:43:VAL:CG1	1:U:44:MET:N	2.30	0.93
1:H:133:TYR:CZ	1:H:375:PHE:HB2	2.03	0.93
1:G:39:ARG:HE	1:G:66:THR:HB	1.32	0.93
1:L:39:ARG:HE	1:L:66:THR:HB	1.32	0.93
1:Q:133:TYR:CZ	1:Q:375:PHE:HB2	2.03	0.93
1:N:39:ARG:HE	1:N:66:THR:HB	1.32	0.93
1:V:133:TYR:CZ	1:V:375:PHE:HB2	2.03	0.93
1:M:43:VAL:CG1	1:M:44:MET:H	1.74	0.93
1:N:300:SER:HA	1:N:335:ARG:HG3	1.51	0.93
1:P:300:SER:HA	1:P:335:ARG:HG3	1.51	0.93
1:T:133:TYR:CZ	1:T:375:PHE:HB2	2.03	0.93
1:R:133:TYR:CZ	1:R:375:PHE:HB2	2.03	0.93
1:U:133:TYR:CZ	1:U:375:PHE:HB2	2.03	0.93
1:A:43:VAL:CG1	1:A:44:MET:N	2.30	0.93
1:G:133:TYR:CZ	1:G:375:PHE:HB2	2.02	0.93
1:L:300:SER:HA	1:L:335:ARG:HG3	1.51	0.93
1:M:34:ILE:CB	1:M:54:VAL:HG11	1.99	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:43:VAL:CG1	1:T:44:MET:H	1.74	0.93
1:B:133:TYR:CZ	1:B:375:PHE:HB2	2.03	0.92
1:K:133:TYR:CZ	1:K:375:PHE:HB2	2.03	0.92
1:R:300:SER:HA	1:R:335:ARG:HG3	1.51	0.92
1:S:133:TYR:CZ	1:S:375:PHE:HB2	2.03	0.92
1:K:34:ILE:CB	1:K:54:VAL:HG11	1.99	0.92
1:O:34:ILE:CB	1:O:54:VAL:HG11	1.99	0.92
1:O:357:ILE:CG1	1:O:370:VAL:HG23	2.00	0.92
1:S:39:ARG:HE	1:S:66:THR:HB	1.32	0.92
1:W:300:SER:HA	1:W:335:ARG:HG3	1.51	0.92
1:F:34:ILE:CB	1:F:54:VAL:HG11	1.99	0.92
1:J:39:ARG:HE	1:J:66:THR:HB	1.32	0.92
1:D:34:ILE:CB	1:D:54:VAL:HG11	1.99	0.92
1:F:133:TYR:CZ	1:F:375:PHE:HB2	2.03	0.92
1:M:39:ARG:HE	1:M:66:THR:HB	1.32	0.92
1:Q:39:ARG:HE	1:Q:66:THR:HB	1.32	0.92
1:U:39:ARG:HE	1:U:66:THR:HB	1.32	0.92
1:D:43:VAL:CG1	1:D:44:MET:N	2.30	0.92
1:D:357:ILE:CG1	1:D:370:VAL:HG23	2.00	0.92
1:J:300:SER:HA	1:J:335:ARG:HG3	1.51	0.92
1:P:39:ARG:HE	1:P:66:THR:HB	1.32	0.92
1:P:43:VAL:CG1	1:P:44:MET:N	2.30	0.92
1:C:300:SER:HA	1:C:335:ARG:HG3	1.51	0.92
1:E:300:SER:HA	1:E:335:ARG:HG3	1.51	0.92
1:I:133:TYR:CZ	1:I:375:PHE:HB2	2.03	0.92
1:K:39:ARG:HE	1:K:66:THR:HB	1.32	0.92
1:O:39:ARG:HE	1:O:66:THR:HB	1.32	0.92
1:Q:34:ILE:CB	1:Q:54:VAL:HG11	1.99	0.92
1:R:39:ARG:HE	1:R:66:THR:HB	1.32	0.92
1:D:133:TYR:CZ	1:D:375:PHE:HB2	2.03	0.92
1:H:34:ILE:CB	1:H:54:VAL:HG11	1.99	0.92
1:A:300:SER:HA	1:A:335:ARG:HG3	1.51	0.92
1:I:34:ILE:CB	1:I:54:VAL:HG11	1.99	0.92
1:T:34:ILE:CB	1:T:54:VAL:HG11	1.99	0.92
1:T:300:SER:HA	1:T:335:ARG:HG3	1.51	0.92
1:V:34:ILE:CB	1:V:54:VAL:HG11	1.99	0.92
1:B:34:ILE:CB	1:B:54:VAL:HG11	1.99	0.92
1:B:39:ARG:HE	1:B:66:THR:HB	1.32	0.92
1:U:300:SER:HA	1:U:335:ARG:HG3	1.51	0.92
1:G:300:SER:HA	1:G:335:ARG:HG3	1.51	0.91
1:J:43:VAL:CG1	1:J:44:MET:N	2.30	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:34:ILE:CB	1:P:54:VAL:HG11	1.99	0.91
1:C:357:ILE:CG1	1:C:370:VAL:HG23	2.00	0.91
1:I:39:ARG:HE	1:I:66:THR:HB	1.32	0.91
1:M:357:ILE:CG1	1:M:370:VAL:HG23	2.00	0.91
1:N:34:ILE:CB	1:N:54:VAL:HG11	1.99	0.91
1:B:357:ILE:CG1	1:B:370:VAL:HG23	2.00	0.91
1:D:39:ARG:HE	1:D:66:THR:HB	1.32	0.91
1:H:39:ARG:HE	1:H:66:THR:HB	1.32	0.91
1:R:34:ILE:CB	1:R:54:VAL:HG11	1.99	0.91
1:E:357:ILE:CG1	1:E:370:VAL:HG23	2.00	0.91
1:J:34:ILE:CB	1:J:54:VAL:HG11	1.99	0.91
1:N:357:ILE:CG1	1:N:370:VAL:HG23	2.00	0.91
1:S:34:ILE:CB	1:S:54:VAL:HG11	1.99	0.91
1:S:43:VAL:CG1	1:S:44:MET:N	2.30	0.91
1:W:34:ILE:CB	1:W:54:VAL:HG11	1.99	0.91
1:A:34:ILE:CB	1:A:54:VAL:HG11	1.99	0.91
1:H:300:SER:HA	1:H:335:ARG:HG3	1.51	0.91
1:L:357:ILE:CG1	1:L:370:VAL:HG23	2.00	0.91
1:P:357:ILE:CG1	1:P:370:VAL:HG23	2.00	0.91
1:S:357:ILE:CG1	1:S:370:VAL:HG23	2.00	0.91
1:U:34:ILE:CB	1:U:54:VAL:HG11	1.99	0.91
1:A:357:ILE:CG1	1:A:370:VAL:HG23	2.00	0.91
1:I:300:SER:HA	1:I:335:ARG:HG3	1.51	0.91
1:O:300:SER:HA	1:O:335:ARG:HG3	1.51	0.91
1:C:34:ILE:CB	1:C:54:VAL:HG11	1.99	0.91
1:W:39:ARG:HE	1:W:66:THR:HB	1.32	0.91
1:B:300:SER:HA	1:B:335:ARG:HG3	1.51	0.90
1:G:34:ILE:CB	1:G:54:VAL:HG11	1.99	0.90
1:G:290:ARG:HD2	1:I:244:ASP:CB	2.01	0.90
1:L:34:ILE:CB	1:L:54:VAL:HG11	1.99	0.90
1:P:290:ARG:HD2	1:R:244:ASP:CB	2.01	0.90
1:A:290:ARG:HD2	1:C:244:ASP:CB	2.01	0.90
1:E:290:ARG:HD2	1:G:244:ASP:CB	2.01	0.90
1:M:290:ARG:HD2	1:O:244:ASP:CB	2.01	0.90
1:H:357:ILE:CG1	1:H:370:VAL:HG23	2.00	0.90
1:K:290:ARG:HD2	1:M:244:ASP:CB	2.01	0.90
1:K:300:SER:HA	1:K:335:ARG:HG3	1.51	0.90
1:L:290:ARG:HD2	1:N:244:ASP:CB	2.01	0.90
1:M:300:SER:HA	1:M:335:ARG:HG3	1.51	0.90
1:R:290:ARG:HD2	1:T:244:ASP:CB	2.01	0.90
1:V:300:SER:HA	1:V:335:ARG:HG3	1.51	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:34:ILE:CB	1:E:54:VAL:HG11	1.99	0.90
1:F:39:ARG:HE	1:F:66:THR:HB	1.32	0.90
1:J:290:ARG:HD2	1:L:244:ASP:CB	2.02	0.90
1:S:300:SER:HA	1:S:335:ARG:HG3	1.51	0.90
1:G:357:ILE:CG1	1:G:370:VAL:HG23	2.00	0.90
1:Q:290:ARG:HD2	1:S:244:ASP:CB	2.01	0.90
1:Q:300:SER:HA	1:Q:335:ARG:HG3	1.51	0.90
1:T:357:ILE:CG1	1:T:370:VAL:HG23	2.00	0.90
1:B:290:ARG:HD2	1:D:244:ASP:CB	2.01	0.90
1:F:290:ARG:HD2	1:H:244:ASP:CB	2.02	0.90
1:L:104:LEU:HB2	1:L:356:TRP:CH2	2.07	0.90
1:S:73:HIC:HA	1:S:183:ARG:HH12	1.36	0.90
1:W:357:ILE:CG1	1:W:370:VAL:HG23	2.00	0.90
1:J:104:LEU:HB2	1:J:356:TRP:CH2	2.07	0.90
1:J:357:ILE:CG1	1:J:370:VAL:HG23	2.00	0.90
1:N:104:LEU:HB2	1:N:356:TRP:CH2	2.07	0.90
1:Q:73:HIC:HA	1:Q:183:ARG:HH12	1.36	0.90
1:D:300:SER:HA	1:D:335:ARG:HG3	1.51	0.90
1:E:43:VAL:CG1	1:E:44:MET:N	2.30	0.90
1:F:104:LEU:HB2	1:F:356:TRP:CH2	2.07	0.90
1:H:34:ILE:HG21	1:H:67:LEU:CD2	2.02	0.90
1:V:357:ILE:CG1	1:V:370:VAL:HG23	2.00	0.90
1:H:104:LEU:HB2	1:H:356:TRP:CH2	2.07	0.90
1:P:104:LEU:HB2	1:P:356:TRP:CH2	2.07	0.90
1:R:104:LEU:HB2	1:R:356:TRP:CH2	2.07	0.90
1:U:290:ARG:HD2	1:W:244:ASP:CB	2.02	0.90
1:W:34:ILE:HG21	1:W:67:LEU:CD2	2.02	0.90
1:B:34:ILE:HG21	1:B:67:LEU:CD2	2.02	0.90
1:D:104:LEU:HB2	1:D:356:TRP:CH2	2.07	0.90
1:Q:34:ILE:HG21	1:Q:67:LEU:CD2	2.02	0.90
1:R:357:ILE:CG1	1:R:370:VAL:HG23	2.00	0.90
1:N:34:ILE:HG21	1:N:67:LEU:CD2	2.02	0.89
1:O:34:ILE:HG21	1:O:67:LEU:CD2	2.02	0.89
1:U:34:ILE:HG21	1:U:67:LEU:CD2	2.02	0.89
1:B:104:LEU:HB2	1:B:356:TRP:CH2	2.07	0.89
1:F:34:ILE:HG21	1:F:67:LEU:CD2	2.02	0.89
1:F:73:HIC:HA	1:F:183:ARG:HH12	1.36	0.89
1:G:104:LEU:HB2	1:G:356:TRP:CH2	2.07	0.89
1:H:73:HIC:HA	1:H:183:ARG:HH12	1.36	0.89
1:I:104:LEU:HB2	1:I:356:TRP:CH2	2.07	0.89
1:K:43:VAL:CG1	1:K:44:MET:N	2.30	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:104:LEU:HB2	1:K:356:TRP:CH2	2.07	0.89
1:N:290:ARG:HD2	1:P:244:ASP:CB	2.01	0.89
1:I:290:ARG:HD2	1:K:244:ASP:CB	2.01	0.89
1:T:104:LEU:HB2	1:T:356:TRP:CH2	2.07	0.89
1:U:357:ILE:CG1	1:U:370:VAL:HG23	2.00	0.89
1:E:73:HIC:HA	1:E:183:ARG:HH12	1.36	0.89
1:E:104:LEU:HB2	1:E:356:TRP:CH2	2.07	0.89
1:G:236:LEU:HD13	1:G:251:GLY:CA	2.02	0.89
1:H:8:LEU:HD12	1:H:90:PHE:CE1	2.08	0.89
1:J:73:HIC:HA	1:J:183:ARG:HH12	1.36	0.89
1:M:104:LEU:HB2	1:M:356:TRP:CH2	2.07	0.89
1:U:73:HIC:HA	1:U:183:ARG:HH12	1.36	0.89
1:A:236:LEU:HD13	1:A:251:GLY:CA	2.02	0.89
1:C:73:HIC:HA	1:C:183:ARG:HH12	1.36	0.89
1:C:290:ARG:HD2	1:E:244:ASP:CB	2.01	0.89
1:I:236:LEU:HD13	1:I:251:GLY:CA	2.02	0.89
1:L:34:ILE:HG21	1:L:67:LEU:CD2	2.02	0.89
1:O:236:LEU:HD13	1:O:251:GLY:CA	2.02	0.89
1:S:8:LEU:HD12	1:S:90:PHE:CE1	2.08	0.89
1:S:290:ARG:HD2	1:U:244:ASP:CB	2.01	0.89
1:W:236:LEU:HD13	1:W:251:GLY:CA	2.02	0.89
1:D:290:ARG:HD2	1:F:244:ASP:CB	2.01	0.89
1:F:300:SER:HA	1:F:335:ARG:HG3	1.51	0.89
1:O:104:LEU:HB2	1:O:356:TRP:CH2	2.07	0.89
1:Q:236:LEU:HD13	1:Q:251:GLY:CA	2.02	0.89
1:S:104:LEU:HB2	1:S:356:TRP:CH2	2.07	0.89
1:U:104:LEU:HB2	1:U:356:TRP:CH2	2.07	0.89
1:A:8:LEU:HD12	1:A:90:PHE:CE1	2.08	0.89
1:A:34:ILE:HG21	1:A:67:LEU:CD2	2.02	0.89
1:C:104:LEU:HB2	1:C:356:TRP:CH2	2.07	0.89
1:E:236:LEU:HD13	1:E:251:GLY:CA	2.02	0.89
1:H:290:ARG:HD2	1:J:244:ASP:CB	2.01	0.89
1:L:8:LEU:HD12	1:L:90:PHE:CE1	2.08	0.89
1:O:73:HIC:HA	1:O:183:ARG:HH12	1.36	0.89
1:R:8:LEU:HD12	1:R:90:PHE:CE1	2.08	0.89
1:T:290:ARG:HD2	1:V:244:ASP:CB	2.01	0.89
1:U:236:LEU:HD13	1:U:251:GLY:CA	2.02	0.89
1:B:43:VAL:CG1	1:B:44:MET:N	2.30	0.89
1:M:236:LEU:HD13	1:M:251:GLY:CA	2.02	0.89
1:O:290:ARG:HD2	1:Q:244:ASP:CB	2.01	0.89
1:P:34:ILE:HG21	1:P:67:LEU:CD2	2.02	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:104:LEU:HB2	1:Q:356:TRP:CH2	2.07	0.89
1:S:236:LEU:HD13	1:S:251:GLY:CA	2.03	0.89
1:V:104:LEU:HB2	1:V:356:TRP:CH2	2.07	0.89
1:A:104:LEU:HB2	1:A:356:TRP:CH2	2.07	0.89
1:D:8:LEU:HD12	1:D:90:PHE:CE1	2.08	0.89
1:G:34:ILE:HG21	1:G:67:LEU:CD2	2.02	0.89
1:I:34:ILE:HG21	1:I:67:LEU:CD2	2.02	0.89
1:I:357:ILE:CG1	1:I:370:VAL:HG23	2.00	0.89
1:P:236:LEU:HD13	1:P:251:GLY:CA	2.02	0.89
1:B:8:LEU:HD12	1:B:90:PHE:CE1	2.08	0.89
1:D:34:ILE:HG21	1:D:67:LEU:CD2	2.02	0.89
1:G:8:LEU:HD12	1:G:90:PHE:CE1	2.08	0.89
1:G:73:HIC:HA	1:G:183:ARG:HH12	1.36	0.89
1:J:34:ILE:HG21	1:J:67:LEU:CD2	2.02	0.89
1:N:8:LEU:HD12	1:N:90:PHE:CE1	2.08	0.89
1:T:43:VAL:CG1	1:T:44:MET:N	2.30	0.89
1:V:34:ILE:HG21	1:V:67:LEU:CD2	2.02	0.89
1:W:104:LEU:HB2	1:W:356:TRP:CH2	2.07	0.89
1:K:236:LEU:HD13	1:K:251:GLY:CA	2.02	0.88
1:K:357:ILE:CG1	1:K:370:VAL:HG23	2.00	0.88
1:N:236:LEU:HD13	1:N:251:GLY:CA	2.02	0.88
1:W:8:LEU:HD12	1:W:90:PHE:CE1	2.08	0.88
1:C:8:LEU:HD12	1:C:90:PHE:CE1	2.08	0.88
1:L:73:HIC:HA	1:L:183:ARG:HH12	1.36	0.88
1:M:8:LEU:HD12	1:M:90:PHE:CE1	2.08	0.88
1:Q:357:ILE:CG1	1:Q:370:VAL:HG23	2.00	0.88
1:R:236:LEU:HD13	1:R:251:GLY:CA	2.02	0.88
1:U:8:LEU:HD12	1:U:90:PHE:CE1	2.08	0.88
1:C:236:LEU:HD13	1:C:251:GLY:CA	2.02	0.88
1:E:8:LEU:HD12	1:E:90:PHE:CE1	2.08	0.88
1:H:236:LEU:HD13	1:H:251:GLY:CA	2.02	0.88
1:J:236:LEU:HD13	1:J:251:GLY:CA	2.02	0.88
1:K:34:ILE:HG21	1:K:67:LEU:CD2	2.02	0.88
1:O:8:LEU:HD12	1:O:90:PHE:CE1	2.08	0.88
1:P:8:LEU:HD12	1:P:90:PHE:CE1	2.08	0.88
1:S:34:ILE:HG21	1:S:67:LEU:CD2	2.02	0.88
1:N:43:VAL:CG1	1:N:44:MET:N	2.30	0.88
1:V:236:LEU:HD13	1:V:251:GLY:CA	2.02	0.88
1:E:34:ILE:HG21	1:E:67:LEU:CD2	2.02	0.88
1:L:236:LEU:HD13	1:L:251:GLY:CA	2.02	0.88
1:H:43:VAL:CG1	1:H:44:MET:N	2.30	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:34:ILE:HG21	1:M:67:LEU:CD2	2.02	0.88
1:R:73:HIC:HA	1:R:183:ARG:HH12	1.36	0.88
1:T:73:HIC:HA	1:T:183:ARG:HH12	1.36	0.88
1:V:8:LEU:HD12	1:V:90:PHE:CE1	2.08	0.88
1:V:73:HIC:HA	1:V:183:ARG:HH12	1.36	0.88
1:A:73:HIC:HA	1:A:183:ARG:HH12	1.36	0.88
1:D:73:HIC:HA	1:D:183:ARG:HH12	1.36	0.88
1:F:8:LEU:HD12	1:F:90:PHE:CE1	2.08	0.88
1:F:236:LEU:HD13	1:F:251:GLY:CA	2.02	0.88
1:J:8:LEU:HD12	1:J:90:PHE:CE1	2.08	0.88
1:I:8:LEU:HD12	1:I:90:PHE:CE1	2.08	0.88
1:T:8:LEU:HD12	1:T:90:PHE:CE1	2.08	0.88
1:T:34:ILE:HG21	1:T:67:LEU:CD2	2.02	0.88
1:W:73:HIC:HA	1:W:183:ARG:HH12	1.36	0.88
1:Q:8:LEU:HD12	1:Q:90:PHE:CE1	2.08	0.88
1:B:236:LEU:HD13	1:B:251:GLY:CA	2.02	0.88
1:F:357:ILE:CG1	1:F:370:VAL:HG23	2.00	0.88
1:K:8:LEU:HD12	1:K:90:PHE:CE1	2.08	0.88
1:N:73:HIC:HA	1:N:183:ARG:HH12	1.36	0.88
1:T:236:LEU:HD13	1:T:251:GLY:CA	2.02	0.88
1:B:73:HIC:HA	1:B:183:ARG:HH12	1.36	0.87
1:D:236:LEU:HD13	1:D:251:GLY:CA	2.02	0.87
1:P:73:HIC:HA	1:P:183:ARG:HH12	1.36	0.87
1:M:73:HIC:HA	1:M:183:ARG:HH12	1.36	0.87
1:Q:43:VAL:CG1	1:Q:44:MET:N	2.30	0.87
1:R:34:ILE:HG21	1:R:67:LEU:CD2	2.02	0.87
1:C:34:ILE:HG21	1:C:67:LEU:CD2	2.02	0.87
1:I:73:HIC:HA	1:I:183:ARG:HH12	1.36	0.87
1:W:43:VAL:CG1	1:W:44:MET:N	2.30	0.87
1:J:134:VAL:O	1:J:375:PHE:HB3	1.76	0.86
1:W:134:VAL:O	1:W:375:PHE:HB3	1.76	0.86
1:L:134:VAL:O	1:L:375:PHE:HB3	1.76	0.86
1:A:134:VAL:O	1:A:375:PHE:HB3	1.76	0.86
1:H:134:VAL:O	1:H:375:PHE:HB3	1.76	0.86
1:N:134:VAL:O	1:N:375:PHE:HB3	1.76	0.86
1:U:134:VAL:O	1:U:375:PHE:HB3	1.76	0.86
1:M:361:GLU:CB	1:M:369:ILE:HD13	2.06	0.85
1:P:134:VAL:O	1:P:375:PHE:HB3	1.76	0.85
1:C:43:VAL:CG1	1:C:44:MET:N	2.30	0.85
1:C:134:VAL:O	1:C:375:PHE:HB3	1.76	0.85
1:D:361:GLU:CB	1:D:369:ILE:HD13	2.06	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:134:VAL:O	1:F:375:PHE:HB3	1.76	0.85
1:F:361:GLU:CB	1:F:369:ILE:HD13	2.06	0.85
1:V:361:GLU:CB	1:V:369:ILE:HD13	2.06	0.85
1:K:73:HIC:HA	1:K:183:ARG:HH12	1.36	0.85
1:S:134:VAL:O	1:S:375:PHE:HB3	1.76	0.85
1:I:43:VAL:CG1	1:I:44:MET:N	2.30	0.85
1:H:361:GLU:CB	1:H:369:ILE:HD13	2.06	0.85
1:D:134:VAL:O	1:D:375:PHE:HB3	1.76	0.85
1:H:236:LEU:HD13	1:H:251:GLY:HA3	1.58	0.85
1:R:134:VAL:O	1:R:375:PHE:HB3	1.76	0.85
1:U:236:LEU:HD13	1:U:251:GLY:HA3	1.58	0.85
1:J:236:LEU:HD13	1:J:251:GLY:HA3	1.58	0.85
1:K:361:GLU:CB	1:K:369:ILE:HD13	2.06	0.85
1:Q:134:VAL:O	1:Q:375:PHE:HB3	1.76	0.85
1:B:361:GLU:CB	1:B:369:ILE:HD13	2.06	0.85
1:E:134:VAL:O	1:E:375:PHE:HB3	1.76	0.85
1:I:236:LEU:HD13	1:I:251:GLY:HA3	1.58	0.85
1:W:236:LEU:HD13	1:W:251:GLY:HA3	1.58	0.85
1:B:134:VAL:O	1:B:375:PHE:HB3	1.76	0.84
1:E:361:GLU:CB	1:E:369:ILE:HD13	2.06	0.84
1:G:134:VAL:O	1:G:375:PHE:HB3	1.76	0.84
1:O:134:VAL:O	1:O:375:PHE:HB3	1.76	0.84
1:T:134:VAL:O	1:T:375:PHE:HB3	1.76	0.84
1:T:361:GLU:CB	1:T:369:ILE:HD13	2.06	0.84
1:C:361:GLU:CB	1:C:369:ILE:HD13	2.06	0.84
1:J:361:GLU:CB	1:J:369:ILE:HD13	2.06	0.84
1:M:134:VAL:O	1:M:375:PHE:HB3	1.76	0.84
1:A:361:GLU:CB	1:A:369:ILE:HD13	2.06	0.84
1:F:236:LEU:HD13	1:F:251:GLY:HA3	1.58	0.84
1:I:134:VAL:O	1:I:375:PHE:HB3	1.76	0.84
1:M:8:LEU:CD1	1:M:90:PHE:HE1	1.90	0.84
1:T:236:LEU:HD13	1:T:251:GLY:HA3	1.58	0.84
1:V:134:VAL:O	1:V:375:PHE:HB3	1.76	0.84
1:G:236:LEU:HD13	1:G:251:GLY:HA3	1.58	0.84
1:K:43:VAL:CG1	1:K:44:MET:H	1.74	0.84
1:K:134:VAL:O	1:K:375:PHE:HB3	1.76	0.84
1:S:236:LEU:HD13	1:S:251:GLY:HA3	1.58	0.84
1:V:8:LEU:CD1	1:V:90:PHE:HE1	1.90	0.84
1:K:236:LEU:HD13	1:K:251:GLY:HA3	1.58	0.84
1:L:236:LEU:HD13	1:L:251:GLY:HA3	1.58	0.84
1:R:43:VAL:CG1	1:R:44:MET:N	2.30	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:8:LEU:CD1	1:W:90:PHE:HE1	1.90	0.84
1:D:236:LEU:HD13	1:D:251:GLY:HA3	1.58	0.84
1:D:287:ILE:HD11	1:F:208:ILE:HD13	1.60	0.84
1:H:287:ILE:HD11	1:J:208:ILE:HD13	1.60	0.84
1:K:8:LEU:CD1	1:K:90:PHE:HE1	1.90	0.84
1:Q:236:LEU:HD13	1:Q:251:GLY:HA3	1.58	0.84
1:Q:287:ILE:HD11	1:S:208:ILE:HD13	1.60	0.84
1:W:361:GLU:CB	1:W:369:ILE:HD13	2.06	0.84
1:M:43:VAL:CG1	1:M:44:MET:N	2.30	0.84
1:N:236:LEU:HD13	1:N:251:GLY:HA3	1.58	0.84
1:O:8:LEU:CD1	1:O:90:PHE:HE1	1.90	0.84
1:B:236:LEU:HD13	1:B:251:GLY:HA3	1.58	0.84
1:B:287:ILE:HD11	1:D:208:ILE:HD13	1.60	0.84
1:J:287:ILE:HD11	1:L:208:ILE:HD13	1.60	0.84
1:L:361:GLU:CB	1:L:369:ILE:HD13	2.06	0.84
1:N:8:LEU:CD1	1:N:90:PHE:HE1	1.90	0.84
1:N:287:ILE:HD11	1:P:208:ILE:HD13	1.60	0.84
1:P:8:LEU:CD1	1:P:90:PHE:HE1	1.90	0.84
1:R:361:GLU:CB	1:R:369:ILE:HD13	2.06	0.84
1:A:236:LEU:HD13	1:A:251:GLY:HA3	1.58	0.83
1:A:287:ILE:HD11	1:C:208:ILE:HD13	1.60	0.83
1:D:8:LEU:CD1	1:D:90:PHE:HE1	1.90	0.83
1:F:8:LEU:CD1	1:F:90:PHE:HE1	1.90	0.83
1:H:8:LEU:CD1	1:H:90:PHE:HE1	1.90	0.83
1:I:361:GLU:CB	1:I:369:ILE:HD13	2.06	0.83
1:L:287:ILE:HD11	1:N:208:ILE:HD13	1.60	0.83
1:S:287:ILE:HD11	1:U:208:ILE:HD13	1.60	0.83
1:U:287:ILE:HD11	1:W:208:ILE:HD13	1.60	0.83
1:U:361:GLU:CB	1:U:369:ILE:HD13	2.06	0.83
1:E:8:LEU:CD1	1:E:90:PHE:HE1	1.90	0.83
1:F:287:ILE:HD11	1:H:208:ILE:HD13	1.60	0.83
1:R:236:LEU:HD13	1:R:251:GLY:HA3	1.58	0.83
1:T:8:LEU:CD1	1:T:90:PHE:HE1	1.90	0.83
1:B:8:LEU:CD1	1:B:90:PHE:HE1	1.90	0.83
1:M:236:LEU:HD13	1:M:251:GLY:HA3	1.58	0.83
1:O:236:LEU:HD13	1:O:251:GLY:HA3	1.58	0.83
1:O:287:ILE:HD11	1:Q:208:ILE:HD13	1.60	0.83
1:U:8:LEU:CD1	1:U:90:PHE:HE1	1.90	0.83
1:C:8:LEU:CD1	1:C:90:PHE:HE1	1.90	0.83
1:L:8:LEU:CD1	1:L:90:PHE:HE1	1.90	0.83
1:N:361:GLU:CB	1:N:369:ILE:HD13	2.06	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:287:ILE:HD11	1:R:208:ILE:HD13	1.60	0.83
1:R:8:LEU:CD1	1:R:90:PHE:HE1	1.90	0.83
1:A:8:LEU:CD1	1:A:90:PHE:HE1	1.90	0.83
1:G:8:LEU:CD1	1:G:90:PHE:HE1	1.90	0.83
1:P:236:LEU:HD13	1:P:251:GLY:HA3	1.58	0.83
1:M:287:ILE:HD11	1:O:208:ILE:HD13	1.60	0.83
1:E:236:LEU:HD13	1:E:251:GLY:HA3	1.58	0.83
1:I:8:LEU:CD1	1:I:90:PHE:HE1	1.90	0.83
1:J:8:LEU:CD1	1:J:90:PHE:HE1	1.90	0.83
1:P:361:GLU:CB	1:P:369:ILE:HD13	2.06	0.83
1:K:287:ILE:HD11	1:M:208:ILE:HD13	1.60	0.83
1:O:43:VAL:CG1	1:O:44:MET:N	2.30	0.83
1:S:361:GLU:CB	1:S:369:ILE:HD13	2.06	0.83
1:C:236:LEU:HD13	1:C:251:GLY:HA3	1.58	0.83
1:Q:8:LEU:CD1	1:Q:90:PHE:HE1	1.90	0.83
1:C:287:ILE:HD11	1:E:208:ILE:HD13	1.60	0.82
1:I:287:ILE:HD11	1:K:208:ILE:HD13	1.60	0.82
1:Q:35:VAL:N	1:Q:54:VAL:HG21	1.95	0.82
1:R:35:VAL:N	1:R:54:VAL:HG21	1.95	0.82
1:R:287:ILE:HD11	1:T:208:ILE:HD13	1.60	0.82
1:G:287:ILE:HD11	1:I:208:ILE:HD13	1.60	0.82
1:O:35:VAL:N	1:O:54:VAL:HG21	1.95	0.82
1:E:287:ILE:HD11	1:G:208:ILE:HD13	1.60	0.82
1:G:35:VAL:N	1:G:54:VAL:HG21	1.95	0.82
1:G:361:GLU:CB	1:G:369:ILE:HD13	2.06	0.82
1:M:35:VAL:N	1:M:54:VAL:HG21	1.95	0.82
1:S:8:LEU:CD1	1:S:90:PHE:HE1	1.90	0.82
1:S:35:VAL:N	1:S:54:VAL:HG21	1.95	0.82
1:F:35:VAL:N	1:F:54:VAL:HG21	1.95	0.82
1:V:236:LEU:HD13	1:V:251:GLY:HA3	1.58	0.82
1:C:35:VAL:N	1:C:54:VAL:HG21	1.95	0.82
1:T:287:ILE:HD11	1:V:208:ILE:HD13	1.60	0.82
1:A:35:VAL:N	1:A:54:VAL:HG21	1.95	0.82
1:K:35:VAL:N	1:K:54:VAL:HG21	1.95	0.82
1:P:35:VAL:N	1:P:54:VAL:HG21	1.95	0.82
1:G:140:LEU:O	1:G:342:GLY:HA3	1.80	0.82
1:B:35:VAL:N	1:B:54:VAL:HG21	1.95	0.82
1:I:140:LEU:O	1:I:342:GLY:HA3	1.80	0.82
1:L:35:VAL:N	1:L:54:VAL:HG21	1.95	0.82
1:N:35:VAL:N	1:N:54:VAL:HG21	1.95	0.82
1:T:35:VAL:N	1:T:54:VAL:HG21	1.95	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:35:VAL:N	1:U:54:VAL:HG21	1.95	0.82
1:V:35:VAL:N	1:V:54:VAL:HG21	1.95	0.82
1:W:35:VAL:N	1:W:54:VAL:HG21	1.95	0.82
1:E:169:TYR:CZ	1:G:40:HIS:HB3	2.15	0.81
1:I:35:VAL:N	1:I:54:VAL:HG21	1.95	0.81
1:K:140:LEU:O	1:K:342:GLY:HA3	1.80	0.81
1:P:169:TYR:CZ	1:R:40:HIS:HB3	2.15	0.81
1:S:34:ILE:C	1:S:54:VAL:HG21	2.05	0.81
1:D:35:VAL:N	1:D:54:VAL:HG21	1.95	0.81
1:E:35:VAL:N	1:E:54:VAL:HG21	1.95	0.81
1:G:169:TYR:CZ	1:I:40:HIS:HB3	2.15	0.81
1:O:34:ILE:C	1:O:54:VAL:HG21	2.06	0.81
1:R:34:ILE:C	1:R:54:VAL:HG21	2.05	0.81
1:R:169:TYR:CZ	1:T:40:HIS:HB3	2.16	0.81
1:T:169:TYR:CZ	1:V:40:HIS:HB3	2.16	0.81
1:B:34:ILE:C	1:B:54:VAL:HG21	2.06	0.81
1:B:362:TYR:HE1	1:B:367:PRO:HB3	1.46	0.81
1:E:34:ILE:C	1:E:54:VAL:HG21	2.06	0.81
1:E:140:LEU:O	1:E:342:GLY:HA3	1.80	0.81
1:F:34:ILE:C	1:F:54:VAL:HG21	2.06	0.81
1:H:140:LEU:O	1:H:342:GLY:HA3	1.80	0.81
1:I:34:ILE:C	1:I:54:VAL:HG21	2.05	0.81
1:M:140:LEU:O	1:M:342:GLY:HA3	1.80	0.81
1:P:140:LEU:O	1:P:342:GLY:HA3	1.80	0.81
1:Q:361:GLU:CB	1:Q:369:ILE:HD13	2.06	0.81
1:U:140:LEU:O	1:U:342:GLY:HA3	1.80	0.81
1:W:34:ILE:C	1:W:54:VAL:HG21	2.06	0.81
1:C:169:TYR:CZ	1:E:40:HIS:HB3	2.15	0.81
1:C:362:TYR:HE1	1:C:367:PRO:HB3	1.46	0.81
1:E:362:TYR:HE1	1:E:367:PRO:HB3	1.46	0.81
1:F:140:LEU:O	1:F:342:GLY:HA3	1.80	0.81
1:H:35:VAL:N	1:H:54:VAL:HG21	1.95	0.81
1:N:34:ILE:C	1:N:54:VAL:HG21	2.05	0.81
1:N:140:LEU:O	1:N:342:GLY:HA3	1.80	0.81
1:N:169:TYR:CZ	1:P:40:HIS:HB3	2.16	0.81
1:N:362:TYR:HE1	1:N:367:PRO:HB3	1.46	0.81
1:Q:140:LEU:O	1:Q:342:GLY:HA3	1.80	0.81
1:S:140:LEU:O	1:S:342:GLY:HA3	1.80	0.81
1:W:140:LEU:O	1:W:342:GLY:HA3	1.80	0.81
1:C:140:LEU:O	1:C:342:GLY:HA3	1.80	0.81
1:D:362:TYR:HE1	1:D:367:PRO:HB3	1.46	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:34:ILE:C	1:K:54:VAL:HG21	2.06	0.81
1:L:34:ILE:C	1:L:54:VAL:HG21	2.06	0.81
1:L:362:TYR:HE1	1:L:367:PRO:HB3	1.46	0.81
1:O:140:LEU:O	1:O:342:GLY:HA3	1.80	0.81
1:O:169:TYR:CZ	1:Q:40:HIS:HB3	2.15	0.81
1:Q:169:TYR:CZ	1:S:40:HIS:HB3	2.16	0.81
1:V:34:ILE:C	1:V:54:VAL:HG21	2.05	0.81
1:V:362:TYR:HE1	1:V:367:PRO:HB3	1.46	0.81
1:W:362:TYR:HE1	1:W:367:PRO:HB3	1.46	0.81
1:C:34:ILE:C	1:C:54:VAL:HG21	2.06	0.81
1:F:362:TYR:HE1	1:F:367:PRO:HB3	1.46	0.81
1:H:362:TYR:HE1	1:H:367:PRO:HB3	1.46	0.81
1:J:140:LEU:O	1:J:342:GLY:HA3	1.80	0.81
1:T:143:TYR:O	1:T:143:TYR:HD1	1.64	0.81
1:U:143:TYR:HD1	1:U:143:TYR:O	1.64	0.81
1:C:143:TYR:O	1:C:143:TYR:HD1	1.64	0.81
1:D:140:LEU:O	1:D:342:GLY:HA3	1.80	0.81
1:D:143:TYR:O	1:D:143:TYR:HD1	1.64	0.81
1:I:169:TYR:CZ	1:K:40:HIS:HB3	2.16	0.81
1:M:169:TYR:CZ	1:O:40:HIS:HB3	2.15	0.81
1:S:169:TYR:CZ	1:U:40:HIS:HB3	2.15	0.81
1:A:34:ILE:C	1:A:54:VAL:HG21	2.06	0.81
1:H:34:ILE:C	1:H:54:VAL:HG21	2.06	0.81
1:J:34:ILE:C	1:J:54:VAL:HG21	2.06	0.81
1:L:169:TYR:CZ	1:N:40:HIS:HB3	2.16	0.81
1:U:169:TYR:CZ	1:W:40:HIS:HB3	2.16	0.81
1:A:140:LEU:O	1:A:342:GLY:HA3	1.80	0.81
1:J:35:VAL:N	1:J:54:VAL:HG21	1.95	0.81
1:J:143:TYR:HD1	1:J:143:TYR:O	1.64	0.81
1:K:169:TYR:CZ	1:M:40:HIS:HB3	2.16	0.81
1:M:34:ILE:C	1:M:54:VAL:HG21	2.05	0.81
1:Q:143:TYR:O	1:Q:143:TYR:HD1	1.64	0.81
1:A:169:TYR:CZ	1:C:40:HIS:HB3	2.15	0.81
1:D:34:ILE:C	1:D:54:VAL:HG21	2.06	0.81
1:F:143:TYR:O	1:F:143:TYR:HD1	1.64	0.81
1:G:362:TYR:HE1	1:G:367:PRO:HB3	1.46	0.81
1:M:362:TYR:HE1	1:M:367:PRO:HB3	1.46	0.81
1:P:34:ILE:C	1:P:54:VAL:HG21	2.06	0.81
1:R:143:TYR:HD1	1:R:143:TYR:O	1.64	0.81
1:D:169:TYR:CZ	1:F:40:HIS:HB3	2.15	0.80
1:E:32:PRO:HG2	1:E:55:GLY:O	1.82	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:169:TYR:CZ	1:L:40:HIS:HB3	2.15	0.80
1:N:357:ILE:HG12	1:N:370:VAL:CG2	2.07	0.80
1:P:362:TYR:HE1	1:P:367:PRO:HB3	1.46	0.80
1:B:140:LEU:O	1:B:342:GLY:HA3	1.80	0.80
1:I:143:TYR:O	1:I:143:TYR:HD1	1.64	0.80
1:J:362:TYR:HE1	1:J:367:PRO:HB3	1.46	0.80
1:O:362:TYR:HE1	1:O:367:PRO:HB3	1.46	0.80
1:P:357:ILE:HG12	1:P:370:VAL:CG2	2.07	0.80
1:T:140:LEU:O	1:T:342:GLY:HA3	1.80	0.80
1:U:34:ILE:C	1:U:54:VAL:HG21	2.05	0.80
1:U:362:TYR:HE1	1:U:367:PRO:HB3	1.46	0.80
1:A:362:TYR:HE1	1:A:367:PRO:HB3	1.46	0.80
1:B:169:TYR:CZ	1:D:40:HIS:HB3	2.15	0.80
1:F:169:TYR:CZ	1:H:40:HIS:HB3	2.15	0.80
1:H:169:TYR:CZ	1:J:40:HIS:HB3	2.15	0.80
1:L:140:LEU:O	1:L:342:GLY:HA3	1.80	0.80
1:O:143:TYR:O	1:O:143:TYR:HD1	1.64	0.80
1:T:32:PRO:HG2	1:T:55:GLY:O	1.82	0.80
1:P:32:PRO:HG2	1:P:55:GLY:O	1.82	0.80
1:V:140:LEU:O	1:V:342:GLY:HA3	1.80	0.80
1:C:357:ILE:HG12	1:C:370:VAL:CG2	2.07	0.80
1:K:143:TYR:O	1:K:143:TYR:HD1	1.64	0.80
1:Q:34:ILE:C	1:Q:54:VAL:HG21	2.06	0.80
1:T:34:ILE:C	1:T:54:VAL:HG21	2.06	0.80
1:E:357:ILE:HG12	1:E:370:VAL:CG2	2.07	0.80
1:G:39:ARG:HE	1:G:66:THR:CB	1.95	0.80
1:N:32:PRO:HG2	1:N:55:GLY:O	1.82	0.80
1:H:58:ALA:HB1	1:H:65:LEU:HD22	1.64	0.80
1:K:32:PRO:HG2	1:K:55:GLY:O	1.82	0.80
1:N:143:TYR:O	1:N:143:TYR:HD1	1.64	0.80
1:P:39:ARG:HE	1:P:66:THR:CB	1.95	0.80
1:P:45:VAL:O	1:P:45:VAL:CG2	2.30	0.80
1:R:39:ARG:HE	1:R:66:THR:CB	1.95	0.80
1:T:362:TYR:HE1	1:T:367:PRO:HB3	1.46	0.80
1:C:45:VAL:O	1:C:45:VAL:CG2	2.30	0.80
1:F:45:VAL:O	1:F:45:VAL:CG2	2.30	0.80
1:G:43:VAL:CG1	1:G:44:MET:N	2.30	0.80
1:J:45:VAL:O	1:J:45:VAL:CG2	2.30	0.80
1:K:362:TYR:HE1	1:K:367:PRO:HB3	1.46	0.80
1:V:32:PRO:HG2	1:V:55:GLY:O	1.82	0.80
1:V:45:VAL:O	1:V:45:VAL:CG2	2.30	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:58:ALA:HB1	1:W:65:LEU:HD22	1.64	0.80
1:A:39:ARG:HE	1:A:66:THR:CB	1.95	0.80
1:A:143:TYR:O	1:A:143:TYR:HD1	1.64	0.80
1:G:45:VAL:O	1:G:45:VAL:CG2	2.30	0.80
1:H:143:TYR:O	1:H:143:TYR:HD1	1.64	0.80
1:L:357:ILE:HG12	1:L:370:VAL:CG2	2.07	0.80
1:R:140:LEU:O	1:R:342:GLY:HA3	1.80	0.80
1:S:45:VAL:O	1:S:45:VAL:CG2	2.30	0.80
1:E:39:ARG:HE	1:E:66:THR:CB	1.95	0.80
1:E:143:TYR:HD1	1:E:143:TYR:O	1.64	0.80
1:I:39:ARG:HE	1:I:66:THR:CB	1.95	0.80
1:N:58:ALA:HB1	1:N:65:LEU:HD22	1.64	0.80
1:W:45:VAL:O	1:W:45:VAL:CG2	2.30	0.80
1:C:32:PRO:HG2	1:C:55:GLY:O	1.82	0.79
1:G:34:ILE:C	1:G:54:VAL:HG21	2.06	0.79
1:J:39:ARG:HE	1:J:66:THR:CB	1.95	0.79
1:J:58:ALA:HB1	1:J:65:LEU:HD22	1.64	0.79
1:L:45:VAL:O	1:L:45:VAL:CG2	2.30	0.79
1:L:58:ALA:HB1	1:L:65:LEU:HD22	1.64	0.79
1:Q:32:PRO:HG2	1:Q:55:GLY:O	1.82	0.79
1:S:58:ALA:HB1	1:S:65:LEU:HD22	1.64	0.79
1:W:143:TYR:O	1:W:143:TYR:HD1	1.64	0.79
1:A:357:ILE:HG12	1:A:370:VAL:CG2	2.07	0.79
1:I:45:VAL:O	1:I:45:VAL:CG2	2.30	0.79
1:K:8:LEU:CD1	1:K:90:PHE:CE1	2.66	0.79
1:M:143:TYR:HD1	1:M:143:TYR:O	1.64	0.79
1:R:357:ILE:HG12	1:R:370:VAL:CG2	2.07	0.79
1:S:143:TYR:O	1:S:143:TYR:HD1	1.64	0.79
1:T:45:VAL:O	1:T:45:VAL:CG2	2.30	0.79
1:V:336:LYS:HE2	2:V:401:ADP:H5'2	1.65	0.79
1:A:58:ALA:HB1	1:A:65:LEU:HD22	1.64	0.79
1:F:32:PRO:HG2	1:F:55:GLY:O	1.82	0.79
1:I:8:LEU:CD1	1:I:90:PHE:CE1	2.66	0.79
1:I:32:PRO:HG2	1:I:55:GLY:O	1.82	0.79
1:O:32:PRO:HG2	1:O:55:GLY:O	1.82	0.79
1:S:32:PRO:HG2	1:S:55:GLY:O	1.82	0.79
1:U:58:ALA:HB1	1:U:65:LEU:HD22	1.64	0.79
1:A:32:PRO:HG2	1:A:55:GLY:O	1.82	0.79
1:C:58:ALA:HB1	1:C:65:LEU:HD22	1.64	0.79
1:D:58:ALA:HB1	1:D:65:LEU:HD22	1.64	0.79
1:F:58:ALA:HB1	1:F:65:LEU:HD22	1.64	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:143:TYR:HD1	1:G:143:TYR:O	1.64	0.79
1:H:32:PRO:HG2	1:H:55:GLY:O	1.82	0.79
1:H:39:ARG:HE	1:H:66:THR:CB	1.95	0.79
1:H:54:VAL:HA	1:H:58:ALA:HB2	1.65	0.79
1:I:362:TYR:HE1	1:I:367:PRO:HB3	1.46	0.79
1:L:143:TYR:O	1:L:143:TYR:HD1	1.64	0.79
1:M:32:PRO:HG2	1:M:55:GLY:O	1.82	0.79
1:O:361:GLU:CB	1:O:369:ILE:HD13	2.06	0.79
1:R:45:VAL:O	1:R:45:VAL:CG2	2.30	0.79
1:S:336:LYS:HE2	2:S:401:ADP:H5'2	1.65	0.79
1:S:362:TYR:HE1	1:S:367:PRO:HB3	1.46	0.79
1:B:8:LEU:CD1	1:B:90:PHE:CE1	2.66	0.79
1:B:143:TYR:HD1	1:B:143:TYR:O	1.64	0.79
1:B:336:LYS:HE2	2:B:401:ADP:H5'2	1.65	0.79
1:F:54:VAL:HA	1:F:58:ALA:HB2	1.65	0.79
1:G:32:PRO:HG2	1:G:55:GLY:O	1.82	0.79
1:N:39:ARG:HE	1:N:66:THR:CB	1.95	0.79
1:Q:336:LYS:HE2	2:Q:401:ADP:H5'2	1.65	0.79
1:Q:362:TYR:HE1	1:Q:367:PRO:HB3	1.46	0.79
1:R:362:TYR:HE1	1:R:367:PRO:HB3	1.46	0.79
1:T:8:LEU:CD1	1:T:90:PHE:CE1	2.66	0.79
1:T:39:ARG:HE	1:T:66:THR:CB	1.95	0.79
1:T:336:LYS:HE2	2:T:401:ADP:H5'2	1.65	0.79
1:U:54:VAL:HA	1:U:58:ALA:HB2	1.65	0.79
1:V:143:TYR:O	1:V:143:TYR:HD1	1.64	0.79
1:W:54:VAL:HA	1:W:58:ALA:HB2	1.65	0.79
1:D:32:PRO:HG2	1:D:55:GLY:O	1.82	0.79
1:E:45:VAL:O	1:E:45:VAL:CG2	2.30	0.79
1:L:39:ARG:HE	1:L:66:THR:CB	1.95	0.79
1:S:54:VAL:HA	1:S:58:ALA:HB2	1.65	0.79
1:U:45:VAL:O	1:U:45:VAL:CG2	2.30	0.79
1:W:357:ILE:HG12	1:W:370:VAL:CG2	2.07	0.79
1:A:45:VAL:O	1:A:45:VAL:CG2	2.30	0.79
1:D:54:VAL:HA	1:D:58:ALA:HB2	1.65	0.79
1:J:32:PRO:HG2	1:J:55:GLY:O	1.82	0.79
1:Q:39:ARG:HE	1:Q:66:THR:CB	1.95	0.79
1:S:39:ARG:HE	1:S:66:THR:CB	1.95	0.79
1:U:336:LYS:HE2	2:U:401:ADP:H5'2	1.65	0.79
1:V:43:VAL:CG1	1:V:44:MET:N	2.30	0.79
1:D:45:VAL:O	1:D:45:VAL:CG2	2.30	0.79
1:G:357:ILE:HG12	1:G:370:VAL:CG2	2.07	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:54:VAL:HA	1:J:58:ALA:HB2	1.65	0.79
1:O:39:ARG:HE	1:O:66:THR:CB	1.95	0.79
1:O:54:VAL:HA	1:O:58:ALA:HB2	1.65	0.79
1:P:58:ALA:HB1	1:P:65:LEU:HD22	1.64	0.79
1:R:8:LEU:CD1	1:R:90:PHE:CE1	2.66	0.79
1:V:39:ARG:HE	1:V:66:THR:CB	1.95	0.79
1:E:336:LYS:HE2	2:E:401:ADP:H5'2	1.65	0.79
1:F:39:ARG:HE	1:F:66:THR:CB	1.95	0.79
1:H:332:PRO:HG2	1:H:335:ARG:CZ	2.13	0.79
1:J:332:PRO:HG2	1:J:335:ARG:CZ	2.13	0.79
1:M:54:VAL:HA	1:M:58:ALA:HB2	1.65	0.79
1:Q:8:LEU:CD1	1:Q:90:PHE:CE1	2.66	0.79
1:U:32:PRO:HG2	1:U:55:GLY:O	1.82	0.79
1:U:332:PRO:HG2	1:U:335:ARG:CZ	2.13	0.79
1:W:8:LEU:CD1	1:W:90:PHE:CE1	2.66	0.79
1:B:32:PRO:HG2	1:B:55:GLY:O	1.82	0.79
1:C:8:LEU:CD1	1:C:90:PHE:CE1	2.66	0.79
1:C:336:LYS:HE2	2:C:401:ADP:H5'2	1.65	0.79
1:D:336:LYS:HE2	2:D:401:ADP:H5'2	1.65	0.79
1:E:8:LEU:CD1	1:E:90:PHE:CE1	2.66	0.79
1:O:336:LYS:HE2	2:O:401:ADP:H5'2	1.65	0.79
1:R:58:ALA:HB1	1:R:65:LEU:HD22	1.64	0.79
1:U:8:LEU:CD1	1:U:90:PHE:CE1	2.66	0.79
1:H:45:VAL:O	1:H:45:VAL:CG2	2.30	0.78
1:K:39:ARG:HE	1:K:66:THR:CB	1.95	0.78
1:L:332:PRO:HG2	1:L:335:ARG:CZ	2.13	0.78
1:N:45:VAL:O	1:N:45:VAL:CG2	2.30	0.78
1:O:8:LEU:CD1	1:O:90:PHE:CE1	2.66	0.78
1:Q:54:VAL:HA	1:Q:58:ALA:HB2	1.65	0.78
1:R:336:LYS:HE2	2:R:401:ADP:H5'2	1.65	0.78
1:B:54:VAL:HA	1:B:58:ALA:HB2	1.65	0.78
1:C:39:ARG:HE	1:C:66:THR:CB	1.95	0.78
1:F:332:PRO:HG2	1:F:335:ARG:CZ	2.13	0.78
1:G:336:LYS:HE2	2:G:401:ADP:H5'2	1.65	0.78
1:I:332:PRO:HG2	1:I:335:ARG:CZ	2.13	0.78
1:L:32:PRO:HG2	1:L:55:GLY:O	1.82	0.78
1:M:39:ARG:HE	1:M:66:THR:CB	1.95	0.78
1:Q:58:ALA:HB1	1:Q:65:LEU:HD22	1.64	0.78
1:S:332:PRO:HG2	1:S:335:ARG:CZ	2.13	0.78
1:U:39:ARG:HE	1:U:66:THR:CB	1.95	0.78
1:W:332:PRO:HG2	1:W:335:ARG:CZ	2.13	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:336:LYS:HE2	2:W:401:ADP:H5'2	1.65	0.78
1:A:8:LEU:CD1	1:A:90:PHE:CE1	2.66	0.78
1:G:332:PRO:HG2	1:G:335:ARG:CZ	2.13	0.78
1:O:58:ALA:HB1	1:O:65:LEU:HD22	1.64	0.78
1:W:39:ARG:HE	1:W:66:THR:CB	1.95	0.78
1:A:54:VAL:HA	1:A:58:ALA:HB2	1.65	0.78
1:A:236:LEU:HD12	1:A:237:GLU:N	1.99	0.78
1:A:336:LYS:HE2	2:A:401:ADP:H5'2	1.65	0.78
1:G:8:LEU:CD1	1:G:90:PHE:CE1	2.66	0.78
1:J:357:ILE:HG12	1:J:370:VAL:CG2	2.06	0.78
1:L:54:VAL:HA	1:L:58:ALA:HB2	1.65	0.78
1:P:143:TYR:O	1:P:143:TYR:HD1	1.64	0.78
1:Q:45:VAL:O	1:Q:45:VAL:CG2	2.30	0.78
1:D:332:PRO:HG2	1:D:335:ARG:CZ	2.13	0.78
1:E:58:ALA:HB1	1:E:65:LEU:HD22	1.64	0.78
1:I:336:LYS:HE2	2:I:401:ADP:H5'2	1.65	0.78
1:N:54:VAL:HA	1:N:58:ALA:HB2	1.65	0.78
1:W:32:PRO:HG2	1:W:55:GLY:O	1.82	0.78
1:D:8:LEU:CD1	1:D:90:PHE:CE1	2.66	0.78
1:K:54:VAL:HA	1:K:58:ALA:HB2	1.65	0.78
1:K:332:PRO:HG2	1:K:335:ARG:CZ	2.13	0.78
1:M:236:LEU:HD12	1:M:237:GLU:N	1.99	0.78
1:N:332:PRO:HG2	1:N:335:ARG:CZ	2.13	0.78
1:R:32:PRO:HG2	1:R:55:GLY:O	1.82	0.78
1:S:8:LEU:CD1	1:S:90:PHE:CE1	2.66	0.78
1:T:332:PRO:HG2	1:T:335:ARG:CZ	2.13	0.78
1:T:357:ILE:HG12	1:T:370:VAL:CG2	2.07	0.78
1:B:39:ARG:HE	1:B:66:THR:CB	1.95	0.78
1:B:58:ALA:HB1	1:B:65:LEU:HD22	1.64	0.78
1:D:39:ARG:HE	1:D:66:THR:CB	1.95	0.78
1:F:336:LYS:HE2	2:F:401:ADP:H5'2	1.65	0.78
1:G:167:GLU:CD	1:I:61:LYS:HE2	2.09	0.78
1:H:8:LEU:CD1	1:H:90:PHE:CE1	2.66	0.78
1:J:236:LEU:HD12	1:J:237:GLU:N	1.99	0.78
1:M:167:GLU:CD	1:O:61:LYS:HE2	2.09	0.78
1:M:336:LYS:HE2	2:M:401:ADP:H5'2	1.65	0.78
1:O:236:LEU:HD12	1:O:237:GLU:N	1.99	0.78
1:Q:332:PRO:HG2	1:Q:335:ARG:CZ	2.13	0.78
1:B:332:PRO:HG2	1:B:335:ARG:CZ	2.13	0.78
1:K:236:LEU:HD12	1:K:237:GLU:N	1.99	0.78
1:L:8:LEU:CD1	1:L:90:PHE:CE1	2.66	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:8:LEU:CD1	1:M:90:PHE:CE1	2.66	0.78
1:P:236:LEU:HD12	1:P:237:GLU:N	1.99	0.78
1:V:332:PRO:HG2	1:V:335:ARG:CZ	2.13	0.78
1:I:167:GLU:CD	1:K:61:LYS:HE2	2.09	0.78
1:N:8:LEU:CD1	1:N:90:PHE:CE1	2.66	0.78
1:O:167:GLU:CD	1:Q:61:LYS:HE2	2.09	0.78
1:P:336:LYS:HE2	2:P:401:ADP:H5'2	1.65	0.78
1:R:236:LEU:HD12	1:R:237:GLU:N	1.99	0.78
1:E:332:PRO:HG2	1:E:335:ARG:CZ	2.13	0.78
1:F:8:LEU:CD1	1:F:90:PHE:CE1	2.66	0.78
1:G:58:ALA:HB1	1:G:65:LEU:HD22	1.64	0.78
1:J:8:LEU:CD1	1:J:90:PHE:CE1	2.66	0.78
1:R:332:PRO:HG2	1:R:335:ARG:CZ	2.13	0.78
1:U:357:ILE:HG12	1:U:370:VAL:CG2	2.07	0.78
1:F:236:LEU:HD12	1:F:237:GLU:N	1.99	0.77
1:L:236:LEU:HD12	1:L:237:GLU:N	1.99	0.77
1:N:35:VAL:HG22	1:N:52:SER:HB2	1.66	0.77
1:P:332:PRO:HG2	1:P:335:ARG:CZ	2.13	0.77
1:T:58:ALA:HB1	1:T:65:LEU:HD22	1.64	0.77
1:U:167:GLU:CD	1:W:61:LYS:HE2	2.09	0.77
1:V:8:LEU:CD1	1:V:90:PHE:CE1	2.66	0.77
1:C:236:LEU:HD12	1:C:237:GLU:N	1.99	0.77
1:D:236:LEU:HD12	1:D:237:GLU:N	1.99	0.77
1:H:357:ILE:HG12	1:H:370:VAL:CG2	2.07	0.77
1:I:58:ALA:HB1	1:I:65:LEU:HD22	1.64	0.77
1:J:35:VAL:HG22	1:J:52:SER:HB2	1.66	0.77
1:M:58:ALA:HB1	1:M:65:LEU:HD22	1.64	0.77
1:M:332:PRO:HG2	1:M:335:ARG:CZ	2.13	0.77
1:P:54:VAL:HA	1:P:58:ALA:HB2	1.65	0.77
1:V:54:VAL:HA	1:V:58:ALA:HB2	1.65	0.77
1:V:58:ALA:HB1	1:V:65:LEU:HD22	1.64	0.77
1:V:357:ILE:HG12	1:V:370:VAL:CG2	2.07	0.77
1:I:357:ILE:HG12	1:I:370:VAL:CG2	2.07	0.77
1:M:45:VAL:O	1:M:45:VAL:CG2	2.30	0.77
1:P:8:LEU:CD1	1:P:90:PHE:CE1	2.66	0.77
1:A:35:VAL:HG22	1:A:52:SER:HB2	1.66	0.77
1:A:167:GLU:CD	1:C:61:LYS:HE2	2.09	0.77
1:G:54:VAL:HA	1:G:58:ALA:HB2	1.65	0.77
1:I:236:LEU:HD12	1:I:237:GLU:N	1.99	0.77
1:N:39:ARG:NE	1:N:66:THR:HA	2.00	0.77
1:O:332:PRO:HG2	1:O:335:ARG:CZ	2.13	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:287:ILE:CD1	1:R:208:ILE:HD13	2.15	0.77
1:Q:173:HIS:CE1	1:R:268:GLY:HA3	2.19	0.77
1:W:35:VAL:HG22	1:W:52:SER:HB2	1.66	0.77
1:A:287:ILE:CD1	1:C:208:ILE:HD13	2.15	0.77
1:C:54:VAL:HA	1:C:58:ALA:HB2	1.65	0.77
1:D:173:HIS:CE1	1:E:268:GLY:HA3	2.19	0.77
1:E:54:VAL:HA	1:E:58:ALA:HB2	1.65	0.77
1:F:173:HIS:CE1	1:G:268:GLY:HA3	2.19	0.77
1:J:167:GLU:CD	1:L:61:LYS:HE2	2.09	0.77
1:M:287:ILE:CD1	1:O:208:ILE:HD13	2.15	0.77
1:Q:236:LEU:HD12	1:Q:237:GLU:N	1.99	0.77
1:T:54:VAL:HA	1:T:58:ALA:HB2	1.65	0.77
1:W:39:ARG:NE	1:W:66:THR:HA	2.00	0.77
1:B:236:LEU:HD12	1:B:237:GLU:N	1.99	0.77
1:G:287:ILE:CD1	1:I:208:ILE:HD13	2.15	0.77
1:H:336:LYS:HE2	2:H:401:ADP:H5'2	1.65	0.77
1:O:173:HIS:CE1	1:P:268:GLY:HA3	2.19	0.77
1:O:287:ILE:CD1	1:Q:208:ILE:HD13	2.15	0.77
1:P:167:GLU:CD	1:R:61:LYS:HE2	2.09	0.77
1:V:236:LEU:HD12	1:V:237:GLU:N	1.99	0.77
1:F:287:ILE:CD1	1:H:208:ILE:HD13	2.15	0.77
1:G:236:LEU:HD12	1:G:237:GLU:N	1.99	0.77
1:I:54:VAL:HA	1:I:58:ALA:HB2	1.65	0.77
1:K:336:LYS:HE2	2:K:401:ADP:H5'2	1.65	0.77
1:N:336:LYS:HE2	2:N:401:ADP:H5'2	1.65	0.77
1:P:39:ARG:NE	1:P:66:THR:HA	2.00	0.77
1:T:236:LEU:HD12	1:T:237:GLU:N	1.99	0.77
1:A:332:PRO:HG2	1:A:335:ARG:CZ	2.13	0.77
1:D:167:GLU:CD	1:F:61:LYS:HE2	2.09	0.77
1:E:39:ARG:NE	1:E:66:THR:HA	2.00	0.77
1:E:173:HIS:CE1	1:F:268:GLY:HA3	2.19	0.77
1:F:167:GLU:CD	1:H:61:LYS:HE2	2.09	0.77
1:H:236:LEU:HD12	1:H:237:GLU:N	1.99	0.77
1:K:58:ALA:HB1	1:K:65:LEU:HD22	1.64	0.77
1:P:173:HIS:CE1	1:Q:268:GLY:HA3	2.19	0.77
1:R:167:GLU:CD	1:T:61:LYS:HE2	2.09	0.77
1:R:287:ILE:CD1	1:T:208:ILE:HD13	2.15	0.77
1:U:236:LEU:HD12	1:U:237:GLU:N	1.99	0.77
1:A:39:ARG:NE	1:A:66:THR:HA	2.00	0.77
1:C:287:ILE:CD1	1:E:208:ILE:HD13	2.15	0.77
1:D:36:GLY:O	1:D:52:SER:HA	1.85	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:39:ARG:NE	1:H:66:THR:HA	2.00	0.77
1:H:287:ILE:CD1	1:J:208:ILE:HD13	2.15	0.77
1:I:287:ILE:CD1	1:K:208:ILE:HD13	2.15	0.77
1:N:167:GLU:CD	1:P:61:LYS:HE2	2.09	0.77
1:R:35:VAL:HG22	1:R:52:SER:HB2	1.66	0.77
1:S:167:GLU:CD	1:U:61:LYS:HE2	2.09	0.77
1:S:173:HIS:CE1	1:T:268:GLY:HA3	2.20	0.77
1:T:167:GLU:CD	1:V:61:LYS:HE2	2.09	0.77
1:T:173:HIS:CE1	1:U:268:GLY:HA3	2.19	0.77
1:U:287:ILE:CD1	1:W:208:ILE:HD13	2.15	0.77
1:B:173:HIS:CE1	1:C:268:GLY:HA3	2.19	0.77
1:K:167:GLU:CD	1:M:61:LYS:HE2	2.09	0.77
1:L:39:ARG:NE	1:L:66:THR:HA	2.00	0.77
1:N:54:VAL:HG12	1:N:55:GLY:N	2.00	0.77
1:N:287:ILE:CD1	1:P:208:ILE:HD13	2.15	0.77
1:O:36:GLY:O	1:O:52:SER:HA	1.85	0.77
1:C:167:GLU:CD	1:E:61:LYS:HE2	2.09	0.76
1:C:173:HIS:CE1	1:D:268:GLY:HA3	2.19	0.76
1:C:332:PRO:HG2	1:C:335:ARG:CZ	2.13	0.76
1:D:287:ILE:CD1	1:F:208:ILE:HD13	2.15	0.76
1:E:36:GLY:O	1:E:52:SER:HA	1.85	0.76
1:F:35:VAL:HG22	1:F:52:SER:HB2	1.66	0.76
1:G:36:GLY:O	1:G:52:SER:HA	1.85	0.76
1:H:173:HIS:CE1	1:I:268:GLY:HA3	2.19	0.76
1:I:36:GLY:O	1:I:52:SER:HA	1.85	0.76
1:J:39:ARG:NE	1:J:66:THR:HA	2.00	0.76
1:L:167:GLU:CD	1:N:61:LYS:HE2	2.09	0.76
1:N:36:GLY:O	1:N:52:SER:HA	1.85	0.76
1:P:36:GLY:O	1:P:52:SER:HA	1.85	0.76
1:R:37:ARG:HG3	1:R:38:PRO:CD	2.15	0.76
1:R:54:VAL:HA	1:R:58:ALA:HB2	1.65	0.76
1:A:173:HIS:CE1	1:B:268:GLY:HA3	2.19	0.76
1:E:236:LEU:HD12	1:E:237:GLU:N	1.99	0.76
1:F:357:ILE:HG12	1:F:370:VAL:CG2	2.07	0.76
1:J:54:VAL:HG12	1:J:55:GLY:N	2.00	0.76
1:L:336:LYS:HE2	2:L:401:ADP:H5'2	1.65	0.76
1:N:173:HIS:CE1	1:O:268:GLY:HA3	2.19	0.76
1:P:37:ARG:HG3	1:P:38:PRO:CD	2.15	0.76
1:S:35:VAL:HG22	1:S:52:SER:HB2	1.66	0.76
1:U:36:GLY:O	1:U:52:SER:HA	1.85	0.76
1:U:37:ARG:HG3	1:U:38:PRO:CD	2.15	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:37:ARG:HG3	1:W:38:PRO:CD	2.15	0.76
1:B:167:GLU:CD	1:D:61:LYS:HE2	2.09	0.76
1:C:39:ARG:NE	1:C:66:THR:HA	2.00	0.76
1:E:35:VAL:HG22	1:E:52:SER:HB2	1.66	0.76
1:E:167:GLU:CD	1:G:61:LYS:HE2	2.09	0.76
1:F:36:GLY:O	1:F:52:SER:HA	1.85	0.76
1:K:36:GLY:O	1:K:52:SER:HA	1.85	0.76
1:K:39:ARG:NE	1:K:66:THR:HA	2.00	0.76
1:L:36:GLY:O	1:L:52:SER:HA	1.85	0.76
1:M:173:HIS:CE1	1:N:268:GLY:HA3	2.19	0.76
1:R:36:GLY:O	1:R:52:SER:HA	1.85	0.76
1:S:37:ARG:HG3	1:S:38:PRO:CD	2.15	0.76
1:V:173:HIS:CE1	1:W:268:GLY:HA3	2.20	0.76
1:C:36:GLY:O	1:C:52:SER:HA	1.85	0.76
1:C:54:VAL:HG12	1:C:55:GLY:N	2.00	0.76
1:F:39:ARG:NE	1:F:66:THR:HA	2.00	0.76
1:J:143:TYR:CZ	1:L:45:VAL:HG21	2.21	0.76
1:Q:37:ARG:HG3	1:Q:38:PRO:CD	2.15	0.76
1:Q:167:GLU:CD	1:S:61:LYS:HE2	2.09	0.76
1:R:173:HIS:CE1	1:S:268:GLY:HA3	2.20	0.76
1:S:287:ILE:CD1	1:U:208:ILE:HD13	2.15	0.76
1:W:236:LEU:HD12	1:W:237:GLU:N	1.99	0.76
1:F:143:TYR:CZ	1:H:45:VAL:HG21	2.21	0.76
1:G:173:HIS:CE1	1:H:268:GLY:HA3	2.19	0.76
1:H:167:GLU:CD	1:J:61:LYS:HE2	2.09	0.76
1:J:287:ILE:CD1	1:L:208:ILE:HD13	2.15	0.76
1:J:336:LYS:HE2	2:J:401:ADP:H5'2	1.65	0.76
1:N:37:ARG:HG3	1:N:38:PRO:CD	2.15	0.76
1:N:236:LEU:HD12	1:N:237:GLU:N	1.99	0.76
1:O:39:ARG:NE	1:O:66:THR:HA	2.00	0.76
1:P:143:TYR:CZ	1:R:45:VAL:HG21	2.21	0.76
1:Q:39:ARG:NE	1:Q:66:THR:HA	2.00	0.76
1:Q:143:TYR:CZ	1:S:45:VAL:HG21	2.21	0.76
1:R:54:VAL:HG12	1:R:55:GLY:N	2.00	0.76
1:S:236:LEU:HD12	1:S:237:GLU:N	1.99	0.76
1:S:357:ILE:HG12	1:S:370:VAL:CG2	2.07	0.76
1:T:36:GLY:O	1:T:52:SER:HA	1.85	0.76
1:U:143:TYR:CZ	1:W:45:VAL:HG21	2.21	0.76
1:E:54:VAL:HG12	1:E:55:GLY:N	2.00	0.76
1:G:39:ARG:NE	1:G:66:THR:HA	2.00	0.76
1:G:143:TYR:CZ	1:I:45:VAL:HG21	2.21	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:39:ARG:NE	1:I:66:THR:HA	2.00	0.76
1:I:173:HIS:CE1	1:J:268:GLY:HA3	2.20	0.76
1:J:36:GLY:O	1:J:52:SER:HA	1.85	0.76
1:L:173:HIS:CE1	1:M:268:GLY:HA3	2.19	0.76
1:S:36:GLY:O	1:S:52:SER:HA	1.85	0.76
1:V:39:ARG:NE	1:V:66:THR:HA	2.00	0.76
1:B:142:LEU:HD21	1:B:165:ILE:HD13	1.68	0.76
1:E:143:TYR:CZ	1:G:45:VAL:HG21	2.21	0.76
1:E:287:ILE:CD1	1:G:208:ILE:HD13	2.15	0.76
1:F:142:LEU:HD21	1:F:165:ILE:HD13	1.68	0.76
1:K:143:TYR:CZ	1:M:45:VAL:HG21	2.21	0.76
1:S:142:LEU:HD21	1:S:165:ILE:HD13	1.68	0.76
1:T:287:ILE:CD1	1:V:208:ILE:HD13	2.15	0.76
1:U:39:ARG:NE	1:U:66:THR:HA	2.00	0.76
1:A:36:GLY:O	1:A:52:SER:HA	1.85	0.76
1:A:143:TYR:CZ	1:C:45:VAL:HG21	2.21	0.76
1:B:287:ILE:CD1	1:D:208:ILE:HD13	2.15	0.76
1:K:287:ILE:CD1	1:M:208:ILE:HD13	2.15	0.76
1:M:36:GLY:O	1:M:52:SER:HA	1.85	0.76
1:O:37:ARG:HG3	1:O:38:PRO:CD	2.15	0.76
1:Q:287:ILE:CD1	1:S:208:ILE:HD13	2.15	0.76
1:U:142:LEU:HD21	1:U:165:ILE:HD13	1.68	0.76
1:U:173:HIS:CE1	1:V:268:GLY:HA3	2.19	0.76
1:B:143:TYR:CZ	1:D:45:VAL:HG21	2.21	0.76
1:D:143:TYR:CZ	1:F:45:VAL:HG21	2.21	0.76
1:I:44:MET:HG3	1:I:45:VAL:H	1.51	0.76
1:K:148:THR:O	1:K:165:ILE:HG22	1.86	0.76
1:K:357:ILE:HG12	1:K:370:VAL:CG2	2.07	0.76
1:M:142:LEU:HD21	1:M:165:ILE:HD13	1.68	0.76
1:O:142:LEU:HD21	1:O:165:ILE:HD13	1.68	0.76
1:R:143:TYR:CZ	1:T:45:VAL:HG21	2.21	0.76
1:T:39:ARG:NE	1:T:66:THR:HA	2.00	0.76
1:T:54:VAL:HG12	1:T:55:GLY:N	2.00	0.76
1:U:54:VAL:HG12	1:U:55:GLY:N	2.00	0.76
1:V:148:THR:O	1:V:165:ILE:HG22	1.86	0.76
1:C:35:VAL:HG22	1:C:52:SER:HB2	1.66	0.76
1:H:142:LEU:HD21	1:H:165:ILE:HD13	1.68	0.76
1:L:143:TYR:CZ	1:N:45:VAL:HG21	2.21	0.76
1:O:44:MET:HG3	1:O:45:VAL:H	1.51	0.76
1:V:44:MET:HG3	1:V:45:VAL:H	1.51	0.76
1:B:44:MET:HG3	1:B:45:VAL:H	1.51	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:45:VAL:O	1:B:45:VAL:CG2	2.30	0.75
1:D:39:ARG:NE	1:D:66:THR:HA	2.00	0.75
1:D:357:ILE:HG12	1:D:370:VAL:CG2	2.07	0.75
1:J:173:HIS:CE1	1:K:268:GLY:HA3	2.19	0.75
1:L:37:ARG:HG3	1:L:38:PRO:CD	2.15	0.75
1:L:287:ILE:CD1	1:N:208:ILE:HD13	2.15	0.75
1:Q:36:GLY:O	1:Q:52:SER:HA	1.85	0.75
1:S:39:ARG:NE	1:S:66:THR:HA	2.00	0.75
1:V:36:GLY:O	1:V:52:SER:HA	1.85	0.75
1:B:39:ARG:NE	1:B:66:THR:HA	2.00	0.75
1:D:142:LEU:HD21	1:D:165:ILE:HD13	1.68	0.75
1:H:36:GLY:O	1:H:52:SER:HA	1.85	0.75
1:I:54:VAL:HG12	1:I:55:GLY:N	2.00	0.75
1:K:173:HIS:CE1	1:L:268:GLY:HA3	2.19	0.75
1:M:37:ARG:HG3	1:M:38:PRO:CD	2.15	0.75
1:O:35:VAL:HG22	1:O:52:SER:HB2	1.66	0.75
1:R:39:ARG:NE	1:R:66:THR:HA	2.00	0.75
1:A:54:VAL:HG12	1:A:55:GLY:N	2.00	0.75
1:B:36:GLY:O	1:B:52:SER:HA	1.85	0.75
1:B:148:THR:O	1:B:165:ILE:HG22	1.86	0.75
1:B:357:ILE:HG12	1:B:370:VAL:CG2	2.07	0.75
1:H:143:TYR:CZ	1:J:45:VAL:HG21	2.21	0.75
1:I:35:VAL:HG22	1:I:52:SER:HB2	1.66	0.75
1:J:148:THR:O	1:J:165:ILE:HG22	1.86	0.75
1:K:142:LEU:HD11	1:K:165:ILE:HD11	1.69	0.75
1:L:148:THR:O	1:L:165:ILE:HG22	1.86	0.75
1:P:54:VAL:HG12	1:P:55:GLY:N	2.01	0.75
1:F:54:VAL:HG12	1:F:55:GLY:N	2.00	0.75
1:N:143:TYR:CZ	1:P:45:VAL:HG21	2.21	0.75
1:O:143:TYR:CZ	1:Q:45:VAL:HG21	2.21	0.75
1:F:44:MET:HG3	1:F:45:VAL:H	1.51	0.75
1:G:54:VAL:HG12	1:G:55:GLY:N	2.01	0.75
1:H:148:THR:O	1:H:165:ILE:HG22	1.86	0.75
1:M:39:ARG:NE	1:M:66:THR:HA	2.00	0.75
1:Q:142:LEU:HD21	1:Q:165:ILE:HD13	1.68	0.75
1:S:143:TYR:CZ	1:U:45:VAL:HG21	2.21	0.75
1:U:35:VAL:HG22	1:U:52:SER:HB2	1.66	0.75
1:E:148:THR:O	1:E:165:ILE:HG22	1.86	0.75
1:G:148:THR:O	1:G:165:ILE:HG22	1.86	0.75
1:K:35:VAL:HG22	1:K:52:SER:HB2	1.66	0.75
1:L:35:VAL:HG22	1:L:52:SER:HB2	1.66	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:143:TYR:CZ	1:O:45:VAL:HG21	2.21	0.75
1:M:148:THR:O	1:M:165:ILE:HG22	1.86	0.75
1:N:290:ARG:CD	1:P:244:ASP:HB2	2.12	0.75
1:Q:35:VAL:HG22	1:Q:52:SER:HB2	1.66	0.75
1:Q:142:LEU:HD11	1:Q:165:ILE:HD11	1.69	0.75
1:Q:148:THR:O	1:Q:165:ILE:HG22	1.86	0.75
1:R:44:MET:HG3	1:R:45:VAL:H	1.51	0.75
1:V:42:GLY:O	1:V:43:VAL:HG12	1.87	0.75
1:W:54:VAL:HG12	1:W:55:GLY:N	2.00	0.75
1:C:42:GLY:O	1:C:43:VAL:HG12	1.87	0.75
1:G:142:LEU:HD11	1:G:165:ILE:HD11	1.69	0.75
1:K:142:LEU:HD21	1:K:165:ILE:HD13	1.68	0.75
1:L:142:LEU:HD21	1:L:165:ILE:HD13	1.68	0.75
1:M:44:MET:HG3	1:M:45:VAL:H	1.51	0.75
1:N:148:THR:O	1:N:165:ILE:HG22	1.86	0.75
1:T:143:TYR:CZ	1:V:45:VAL:HG21	2.21	0.75
1:V:35:VAL:HG22	1:V:52:SER:HB2	1.66	0.75
1:V:142:LEU:HD11	1:V:165:ILE:HD11	1.69	0.75
1:W:142:LEU:HD21	1:W:165:ILE:HD13	1.68	0.75
1:D:42:GLY:O	1:D:43:VAL:HG12	1.87	0.75
1:E:42:GLY:O	1:E:43:VAL:HG12	1.87	0.75
1:F:148:THR:O	1:F:165:ILE:HG22	1.86	0.75
1:G:35:VAL:HG22	1:G:52:SER:HB2	1.66	0.75
1:J:37:ARG:HG3	1:J:38:PRO:CD	2.15	0.75
1:J:39:ARG:HE	1:J:66:THR:HA	1.52	0.75
1:J:142:LEU:HD21	1:J:165:ILE:HD13	1.68	0.75
1:K:37:ARG:HG3	1:K:38:PRO:CD	2.15	0.75
1:T:42:GLY:O	1:T:43:VAL:HG12	1.87	0.75
1:U:42:GLY:O	1:U:43:VAL:HG12	1.87	0.75
1:V:54:VAL:HG12	1:V:55:GLY:N	2.00	0.75
1:A:42:GLY:O	1:A:43:VAL:HG12	1.87	0.75
1:B:42:GLY:O	1:B:43:VAL:HG12	1.87	0.75
1:C:143:TYR:CZ	1:E:45:VAL:HG21	2.21	0.75
1:D:142:LEU:HD11	1:D:165:ILE:HD11	1.69	0.75
1:I:148:THR:O	1:I:165:ILE:HG22	1.86	0.75
1:Q:357:ILE:HG12	1:Q:370:VAL:CG2	2.07	0.75
1:S:44:MET:HG3	1:S:45:VAL:H	1.51	0.75
1:T:148:THR:O	1:T:165:ILE:HG22	1.86	0.75
1:W:42:GLY:O	1:W:43:VAL:HG12	1.87	0.75
1:A:39:ARG:HE	1:A:66:THR:HA	1.52	0.74
1:B:35:VAL:HG22	1:B:52:SER:HB2	1.66	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:142:LEU:HD21	1:I:165:ILE:HD13	1.68	0.74
1:I:143:TYR:CZ	1:K:45:VAL:HG21	2.21	0.74
1:M:35:VAL:HG22	1:M:52:SER:HB2	1.66	0.74
1:O:45:VAL:O	1:O:45:VAL:CG2	2.30	0.74
1:O:54:VAL:HG12	1:O:55:GLY:N	2.00	0.74
1:O:142:LEU:HD11	1:O:165:ILE:HD11	1.69	0.74
1:P:35:VAL:HG22	1:P:52:SER:HB2	1.66	0.74
1:W:36:GLY:O	1:W:52:SER:HA	1.85	0.74
1:E:44:MET:HG3	1:E:45:VAL:H	1.51	0.74
1:F:142:LEU:HD11	1:F:165:ILE:HD11	1.69	0.74
1:G:42:GLY:O	1:G:43:VAL:HG12	1.87	0.74
1:H:54:VAL:HG12	1:H:55:GLY:N	2.00	0.74
1:J:290:ARG:CD	1:L:244:ASP:HB2	2.12	0.74
1:M:142:LEU:HD11	1:M:165:ILE:HD11	1.69	0.74
1:T:35:VAL:HG22	1:T:52:SER:HB2	1.66	0.74
1:U:39:ARG:HE	1:U:66:THR:HA	1.52	0.74
1:D:35:VAL:HG22	1:D:52:SER:HB2	1.66	0.74
1:M:357:ILE:HG12	1:M:370:VAL:CG2	2.07	0.74
1:P:148:THR:O	1:P:165:ILE:HG22	1.86	0.74
1:I:37:ARG:HG3	1:I:38:PRO:CD	2.15	0.74
1:R:42:GLY:O	1:R:43:VAL:HG12	1.87	0.74
1:W:148:THR:O	1:W:165:ILE:HG22	1.86	0.74
1:A:142:LEU:HD21	1:A:165:ILE:HD13	1.68	0.74
1:G:44:MET:HG3	1:G:45:VAL:H	1.51	0.74
1:H:35:VAL:HG22	1:H:52:SER:HB2	1.66	0.74
1:K:44:MET:HG3	1:K:45:VAL:H	1.51	0.74
1:L:54:VAL:HG12	1:L:55:GLY:N	2.00	0.74
1:N:142:LEU:HD21	1:N:165:ILE:HD13	1.68	0.74
1:O:357:ILE:HG12	1:O:370:VAL:CG2	2.07	0.74
1:R:142:LEU:HD11	1:R:165:ILE:HD11	1.69	0.74
1:S:39:ARG:HE	1:S:66:THR:HA	1.52	0.74
1:S:42:GLY:O	1:S:43:VAL:HG12	1.87	0.74
1:U:142:LEU:HD11	1:U:165:ILE:HD11	1.69	0.74
1:V:142:LEU:HD21	1:V:165:ILE:HD13	1.68	0.74
1:A:148:THR:O	1:A:165:ILE:HG22	1.86	0.74
1:D:54:VAL:HG12	1:D:55:GLY:N	2.00	0.74
1:D:148:THR:O	1:D:165:ILE:HG22	1.86	0.74
1:I:42:GLY:O	1:I:43:VAL:HG12	1.87	0.74
1:L:42:GLY:O	1:L:43:VAL:HG12	1.87	0.74
1:M:42:GLY:O	1:M:43:VAL:HG12	1.87	0.74
1:Q:54:VAL:HG12	1:Q:55:GLY:N	2.01	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:180:LEU:HD12	1:Q:181:ALA:N	2.03	0.74
1:S:148:THR:O	1:S:165:ILE:HG22	1.86	0.74
1:B:54:VAL:HG12	1:B:55:GLY:N	2.00	0.74
1:B:142:LEU:HD11	1:B:165:ILE:HD11	1.69	0.74
1:F:42:GLY:O	1:F:43:VAL:HG12	1.87	0.74
1:G:142:LEU:HD21	1:G:165:ILE:HD13	1.68	0.74
1:H:37:ARG:HG3	1:H:38:PRO:CD	2.15	0.74
1:H:39:ARG:HE	1:H:66:THR:HA	1.52	0.74
1:M:39:ARG:HE	1:M:66:THR:HA	1.52	0.74
1:M:54:VAL:HG12	1:M:55:GLY:N	2.01	0.74
1:O:148:THR:O	1:O:165:ILE:HG22	1.86	0.74
1:Q:290:ARG:CD	1:S:244:ASP:HB2	2.12	0.74
1:S:54:VAL:HG12	1:S:55:GLY:N	2.01	0.74
1:S:180:LEU:HD12	1:S:181:ALA:N	2.03	0.74
1:C:44:MET:HG3	1:C:45:VAL:H	1.51	0.74
1:C:148:THR:O	1:C:165:ILE:HG22	1.86	0.74
1:G:37:ARG:HG3	1:G:38:PRO:CD	2.15	0.74
1:K:42:GLY:O	1:K:43:VAL:HG12	1.87	0.74
1:L:43:VAL:O	1:L:44:MET:CB	2.36	0.74
1:O:42:GLY:O	1:O:43:VAL:HG12	1.87	0.74
1:P:42:GLY:O	1:P:43:VAL:HG12	1.87	0.74
1:R:39:ARG:HE	1:R:66:THR:HA	1.52	0.74
1:T:142:LEU:HD21	1:T:165:ILE:HD13	1.68	0.74
1:U:180:LEU:HD12	1:U:181:ALA:N	2.03	0.74
1:J:42:GLY:O	1:J:43:VAL:HG12	1.87	0.74
1:K:39:ARG:HE	1:K:66:THR:HA	1.52	0.74
1:N:42:GLY:O	1:N:43:VAL:HG12	1.87	0.74
1:O:180:LEU:HD12	1:O:181:ALA:N	2.03	0.74
1:U:44:MET:HG3	1:U:45:VAL:H	1.51	0.74
1:U:148:THR:O	1:U:165:ILE:HG22	1.86	0.74
1:J:180:LEU:HD12	1:J:181:ALA:N	2.03	0.74
1:A:43:VAL:O	1:A:44:MET:CB	2.36	0.73
1:C:180:LEU:HD12	1:C:181:ALA:N	2.03	0.73
1:F:290:ARG:CD	1:H:244:ASP:HB2	2.12	0.73
1:H:180:LEU:HD12	1:H:181:ALA:N	2.03	0.73
1:J:44:MET:HG3	1:J:45:VAL:H	1.51	0.73
1:K:54:VAL:HG12	1:K:55:GLY:N	2.00	0.73
1:L:180:LEU:HD12	1:L:181:ALA:N	2.03	0.73
1:R:148:THR:O	1:R:165:ILE:HG22	1.86	0.73
1:T:44:MET:HG3	1:T:45:VAL:H	1.51	0.73
1:W:43:VAL:O	1:W:44:MET:CB	2.36	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:44:MET:HG3	1:W:45:VAL:H	1.51	0.73
1:C:43:VAL:O	1:C:44:MET:CB	2.36	0.73
1:D:44:MET:HG3	1:D:45:VAL:H	1.51	0.73
1:D:287:ILE:HD11	1:F:208:ILE:CD1	2.19	0.73
1:E:37:ARG:HG3	1:E:38:PRO:CD	2.16	0.73
1:E:142:LEU:HD11	1:E:165:ILE:HD11	1.69	0.73
1:J:43:VAL:O	1:J:44:MET:CB	2.36	0.73
1:N:43:VAL:O	1:N:44:MET:CB	2.36	0.73
1:N:44:MET:HG3	1:N:45:VAL:H	1.51	0.73
1:P:39:ARG:HE	1:P:66:THR:HA	1.52	0.73
1:P:142:LEU:HD21	1:P:165:ILE:HD13	1.68	0.73
1:Q:44:MET:HG3	1:Q:45:VAL:H	1.51	0.73
1:R:142:LEU:HD21	1:R:165:ILE:HD13	1.68	0.73
1:U:43:VAL:O	1:U:44:MET:CB	2.36	0.73
1:U:290:ARG:CD	1:W:244:ASP:HB2	2.12	0.73
1:A:44:MET:HG3	1:A:45:VAL:H	1.51	0.73
1:E:142:LEU:HD21	1:E:165:ILE:HD13	1.68	0.73
1:G:180:LEU:HD12	1:G:181:ALA:N	2.03	0.73
1:H:223:PHE:CD1	1:H:259:GLU:HG2	2.24	0.73
1:I:142:LEU:HD11	1:I:165:ILE:HD11	1.69	0.73
1:J:142:LEU:HD11	1:J:165:ILE:HD11	1.69	0.73
1:K:287:ILE:HD11	1:M:208:ILE:CD1	2.19	0.73
1:M:180:LEU:HD12	1:M:181:ALA:N	2.03	0.73
1:P:44:MET:HG3	1:P:45:VAL:H	1.51	0.73
1:P:287:ILE:HD11	1:R:208:ILE:CD1	2.19	0.73
1:R:180:LEU:HD12	1:R:181:ALA:N	2.03	0.73
1:R:287:ILE:HD11	1:T:208:ILE:CD1	2.19	0.73
1:T:142:LEU:HD11	1:T:165:ILE:HD11	1.69	0.73
1:T:142:LEU:HD11	1:T:165:ILE:CD1	2.19	0.73
1:T:180:LEU:HD12	1:T:181:ALA:N	2.03	0.73
1:A:142:LEU:HD11	1:A:165:ILE:CD1	2.19	0.73
1:B:34:ILE:HD13	1:B:67:LEU:HD13	1.71	0.73
1:B:287:ILE:HD11	1:D:208:ILE:CD1	2.19	0.73
1:C:142:LEU:HD21	1:C:165:ILE:HD13	1.68	0.73
1:E:142:LEU:HD11	1:E:165:ILE:CD1	2.19	0.73
1:G:287:ILE:HD11	1:I:208:ILE:CD1	2.19	0.73
1:I:287:ILE:HD11	1:K:208:ILE:CD1	2.19	0.73
1:J:223:PHE:CD1	1:J:259:GLU:HG2	2.23	0.73
1:L:44:MET:HG3	1:L:45:VAL:H	1.51	0.73
1:M:287:ILE:HD11	1:O:208:ILE:CD1	2.19	0.73
1:N:180:LEU:HD12	1:N:181:ALA:N	2.03	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:39:ARG:HE	1:O:66:THR:HA	1.52	0.73
1:W:180:LEU:HD12	1:W:181:ALA:N	2.03	0.73
1:A:142:LEU:HD11	1:A:165:ILE:HD11	1.69	0.73
1:D:34:ILE:HD13	1:D:67:LEU:HD13	1.71	0.73
1:F:287:ILE:HD11	1:H:208:ILE:CD1	2.19	0.73
1:I:39:ARG:HE	1:I:66:THR:HA	1.52	0.73
1:N:287:ILE:HD11	1:P:208:ILE:CD1	2.19	0.73
1:O:208:ILE:CD1	1:O:243:PRO:HD2	2.18	0.73
1:P:142:LEU:HD11	1:P:165:ILE:CD1	2.19	0.73
1:E:180:LEU:HD12	1:E:181:ALA:N	2.03	0.73
1:E:287:ILE:HD11	1:G:208:ILE:CD1	2.19	0.73
1:F:37:ARG:HG3	1:F:38:PRO:CD	2.15	0.73
1:F:180:LEU:HD12	1:F:181:ALA:N	2.03	0.73
1:F:223:PHE:CD1	1:F:259:GLU:HG2	2.24	0.73
1:H:44:MET:HG3	1:H:45:VAL:H	1.51	0.73
1:J:142:LEU:HD11	1:J:165:ILE:CD1	2.19	0.73
1:L:39:ARG:HE	1:L:66:THR:HA	1.52	0.73
1:O:287:ILE:HD11	1:Q:208:ILE:CD1	2.19	0.73
1:O:290:ARG:CD	1:Q:244:ASP:HB2	2.12	0.73
1:P:142:LEU:HD11	1:P:165:ILE:HD11	1.69	0.73
1:P:180:LEU:HD12	1:P:181:ALA:N	2.03	0.73
1:V:39:ARG:HE	1:V:66:THR:HA	1.52	0.73
1:V:180:LEU:HD12	1:V:181:ALA:N	2.03	0.73
1:V:223:PHE:CD1	1:V:259:GLU:HG2	2.24	0.73
1:W:142:LEU:HD11	1:W:165:ILE:HD11	1.69	0.73
1:C:37:ARG:HG3	1:C:38:PRO:CD	2.15	0.73
1:K:45:VAL:O	1:K:45:VAL:CG2	2.30	0.73
1:L:142:LEU:HD11	1:L:165:ILE:HD11	1.69	0.73
1:L:142:LEU:HD11	1:L:165:ILE:CD1	2.19	0.73
1:L:223:PHE:CD1	1:L:259:GLU:HG2	2.23	0.73
1:N:39:ARG:HE	1:N:66:THR:HA	1.52	0.73
1:N:142:LEU:HD11	1:N:165:ILE:HD11	1.69	0.73
1:P:223:PHE:CD1	1:P:259:GLU:HG2	2.24	0.73
1:R:290:ARG:CD	1:T:244:ASP:HB2	2.12	0.73
1:T:39:ARG:HE	1:T:66:THR:HA	1.52	0.73
1:T:287:ILE:HD11	1:V:208:ILE:CD1	2.19	0.73
1:D:208:ILE:CD1	1:D:243:PRO:HD2	2.18	0.73
1:E:208:ILE:CD1	1:E:243:PRO:HD2	2.18	0.73
1:F:34:ILE:HD13	1:F:67:LEU:HD13	1.71	0.73
1:G:39:ARG:HE	1:G:66:THR:HA	1.52	0.73
1:H:42:GLY:O	1:H:43:VAL:HG12	1.87	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:223:PHE:CD1	1:N:259:GLU:HG2	2.24	0.73
1:O:223:PHE:CD1	1:O:259:GLU:HG2	2.24	0.73
1:Q:39:ARG:HE	1:Q:66:THR:HA	1.52	0.73
1:Q:42:GLY:O	1:Q:43:VAL:HG12	1.87	0.73
1:Q:223:PHE:CD1	1:Q:259:GLU:HG2	2.24	0.73
1:R:223:PHE:CD1	1:R:259:GLU:HG2	2.24	0.73
1:T:223:PHE:CD1	1:T:259:GLU:HG2	2.24	0.73
1:A:180:LEU:HD12	1:A:181:ALA:N	2.03	0.73
1:I:180:LEU:HD12	1:I:181:ALA:N	2.03	0.73
1:S:142:LEU:HD11	1:S:165:ILE:HD11	1.69	0.73
1:U:34:ILE:HD13	1:U:67:LEU:HD13	1.71	0.73
1:V:43:VAL:O	1:V:44:MET:CB	2.36	0.73
1:B:43:VAL:O	1:B:44:MET:CB	2.36	0.73
1:B:180:LEU:HD12	1:B:181:ALA:N	2.03	0.73
1:D:223:PHE:CD1	1:D:259:GLU:HG2	2.24	0.73
1:F:43:VAL:O	1:F:44:MET:CB	2.36	0.73
1:G:142:LEU:HD11	1:G:165:ILE:CD1	2.19	0.73
1:S:223:PHE:CD1	1:S:259:GLU:HG2	2.24	0.73
1:V:142:LEU:HD11	1:V:165:ILE:CD1	2.19	0.73
1:W:34:ILE:HD13	1:W:67:LEU:HD13	1.71	0.73
1:W:223:PHE:CD1	1:W:259:GLU:HG2	2.24	0.73
1:C:61:LYS:O	1:C:64:ILE:HG22	1.90	0.72
1:D:180:LEU:HD12	1:D:181:ALA:N	2.03	0.72
1:K:142:LEU:HD11	1:K:165:ILE:CD1	2.19	0.72
1:K:180:LEU:HD12	1:K:181:ALA:N	2.03	0.72
1:M:223:PHE:CD1	1:M:259:GLU:HG2	2.24	0.72
1:S:34:ILE:HD13	1:S:67:LEU:HD13	1.71	0.72
1:U:142:LEU:HD11	1:U:165:ILE:CD1	2.19	0.72
1:U:208:ILE:CD1	1:U:243:PRO:HD2	2.18	0.72
1:U:287:ILE:HD11	1:W:208:ILE:CD1	2.19	0.72
1:A:39:ARG:HE	1:A:66:THR:CA	2.03	0.72
1:D:290:ARG:CD	1:F:244:ASP:HB2	2.12	0.72
1:L:287:ILE:HD11	1:N:208:ILE:CD1	2.19	0.72
1:P:208:ILE:CD1	1:P:243:PRO:HD2	2.18	0.72
1:R:142:LEU:HD11	1:R:165:ILE:CD1	2.19	0.72
1:A:61:LYS:O	1:A:64:ILE:HG22	1.90	0.72
1:C:142:LEU:HD11	1:C:165:ILE:CD1	2.19	0.72
1:C:287:ILE:HD11	1:E:208:ILE:CD1	2.19	0.72
1:D:244:ASP:OD1	1:D:246:GLN:HG3	1.89	0.72
1:E:43:VAL:O	1:E:44:MET:CB	2.36	0.72
1:E:61:LYS:O	1:E:64:ILE:HG22	1.90	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:142:LEU:HD11	1:F:165:ILE:CD1	2.19	0.72
1:H:244:ASP:OD1	1:H:246:GLN:HG3	1.90	0.72
1:H:287:ILE:HD11	1:J:208:ILE:CD1	2.19	0.72
1:I:34:ILE:HD13	1:I:67:LEU:HD13	1.71	0.72
1:J:39:ARG:HE	1:J:66:THR:CA	2.03	0.72
1:P:43:VAL:O	1:P:44:MET:CB	2.36	0.72
1:Q:287:ILE:HD11	1:S:208:ILE:CD1	2.19	0.72
1:S:142:LEU:HD11	1:S:165:ILE:CD1	2.19	0.72
1:S:244:ASP:OD1	1:S:246:GLN:HG3	1.90	0.72
1:V:34:ILE:HD13	1:V:67:LEU:HD13	1.71	0.72
1:W:244:ASP:OD1	1:W:246:GLN:HG3	1.89	0.72
1:B:43:VAL:O	1:B:44:MET:HB3	1.90	0.72
1:B:223:PHE:CD1	1:B:259:GLU:HG2	2.24	0.72
1:C:142:LEU:HD11	1:C:165:ILE:HD11	1.69	0.72
1:E:43:VAL:O	1:E:44:MET:HB3	1.90	0.72
1:F:301:GLY:H	1:F:335:ARG:HG3	1.55	0.72
1:G:61:LYS:O	1:G:64:ILE:HG22	1.90	0.72
1:G:290:ARG:CD	1:I:244:ASP:HB2	2.12	0.72
1:H:34:ILE:HD13	1:H:67:LEU:HD13	1.71	0.72
1:H:142:LEU:HD11	1:H:165:ILE:HD11	1.69	0.72
1:N:142:LEU:HD11	1:N:165:ILE:CD1	2.19	0.72
1:P:39:ARG:HE	1:P:66:THR:CA	2.03	0.72
1:Q:244:ASP:OD1	1:Q:246:GLN:HG3	1.90	0.72
1:V:37:ARG:HG3	1:V:38:PRO:CD	2.15	0.72
1:A:37:ARG:HG3	1:A:38:PRO:CD	2.15	0.72
1:A:223:PHE:CD1	1:A:259:GLU:HG2	2.24	0.72
1:D:37:ARG:HG3	1:D:38:PRO:CD	2.16	0.72
1:D:39:ARG:HE	1:D:66:THR:HA	1.52	0.72
1:D:142:LEU:HD11	1:D:165:ILE:CD1	2.19	0.72
1:D:301:GLY:H	1:D:335:ARG:HG3	1.55	0.72
1:G:34:ILE:HD13	1:G:67:LEU:HD13	1.71	0.72
1:G:43:VAL:O	1:G:44:MET:HB3	1.90	0.72
1:I:61:LYS:O	1:I:64:ILE:HG22	1.90	0.72
1:J:244:ASP:OD1	1:J:246:GLN:HG3	1.90	0.72
1:K:34:ILE:HD13	1:K:67:LEU:HD13	1.71	0.72
1:K:223:PHE:CD1	1:K:259:GLU:HG2	2.24	0.72
1:L:39:ARG:HE	1:L:66:THR:CA	2.03	0.72
1:M:142:LEU:HD11	1:M:165:ILE:CD1	2.19	0.72
1:N:301:GLY:H	1:N:335:ARG:HG3	1.55	0.72
1:O:301:GLY:H	1:O:335:ARG:HG3	1.55	0.72
1:R:39:ARG:HE	1:R:66:THR:CA	2.03	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:43:VAL:O	1:T:44:MET:CB	2.36	0.72
1:U:39:ARG:HE	1:U:66:THR:CA	2.03	0.72
1:U:223:PHE:CD1	1:U:259:GLU:HG2	2.24	0.72
1:W:142:LEU:HD11	1:W:165:ILE:CD1	2.19	0.72
1:W:301:GLY:H	1:W:335:ARG:HG3	1.55	0.72
1:B:39:ARG:HE	1:B:66:THR:HA	1.52	0.72
1:C:301:GLY:H	1:C:335:ARG:HG3	1.55	0.72
1:D:43:VAL:O	1:D:44:MET:CB	2.36	0.72
1:H:142:LEU:HD11	1:H:165:ILE:CD1	2.19	0.72
1:H:236:LEU:HD13	1:H:251:GLY:HA2	1.72	0.72
1:I:43:VAL:O	1:I:44:MET:CB	2.36	0.72
1:J:208:ILE:CD1	1:J:243:PRO:HD2	2.18	0.72
1:L:301:GLY:H	1:L:335:ARG:HG3	1.55	0.72
1:N:39:ARG:HE	1:N:66:THR:CA	2.03	0.72
1:O:142:LEU:HD11	1:O:165:ILE:CD1	2.19	0.72
1:O:244:ASP:OD1	1:O:246:GLN:HG3	1.89	0.72
1:Q:34:ILE:HD13	1:Q:67:LEU:HD13	1.71	0.72
1:T:39:ARG:HE	1:T:66:THR:CA	2.03	0.72
1:A:287:ILE:HD11	1:C:208:ILE:CD1	2.19	0.72
1:A:290:ARG:CD	1:C:244:ASP:HB2	2.12	0.72
1:B:142:LEU:HD11	1:B:165:ILE:CD1	2.19	0.72
1:C:43:VAL:O	1:C:44:MET:HB3	1.90	0.72
1:F:39:ARG:HE	1:F:66:THR:HA	1.52	0.72
1:F:236:LEU:HD13	1:F:251:GLY:HA2	1.72	0.72
1:K:43:VAL:O	1:K:44:MET:CB	2.36	0.72
1:K:61:LYS:O	1:K:64:ILE:HG22	1.90	0.72
1:M:43:VAL:O	1:M:44:MET:CB	2.36	0.72
1:M:301:GLY:H	1:M:335:ARG:HG3	1.55	0.72
1:O:43:VAL:O	1:O:44:MET:CB	2.36	0.72
1:U:301:GLY:H	1:U:335:ARG:HG3	1.55	0.72
1:D:43:VAL:O	1:D:44:MET:HB3	1.90	0.72
1:E:34:ILE:HD13	1:E:67:LEU:HD13	1.71	0.72
1:E:39:ARG:HE	1:E:66:THR:HA	1.52	0.72
1:E:301:GLY:H	1:E:335:ARG:HG3	1.55	0.72
1:I:39:ARG:HE	1:I:66:THR:CA	2.03	0.72
1:I:43:VAL:O	1:I:44:MET:HB3	1.90	0.72
1:I:142:LEU:HD11	1:I:165:ILE:CD1	2.19	0.72
1:I:223:PHE:CD1	1:I:259:GLU:HG2	2.24	0.72
1:L:244:ASP:OD1	1:L:246:GLN:HG3	1.90	0.72
1:M:61:LYS:O	1:M:64:ILE:HG22	1.90	0.72
1:M:236:LEU:HD13	1:M:251:GLY:HA2	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:43:VAL:O	1:Q:44:MET:CB	2.36	0.72
1:S:39:ARG:HE	1:S:66:THR:CA	2.03	0.72
1:S:287:ILE:HD11	1:U:208:ILE:CD1	2.19	0.72
1:S:301:GLY:H	1:S:335:ARG:HG3	1.55	0.72
1:W:61:LYS:O	1:W:64:ILE:HG22	1.90	0.72
1:C:39:ARG:HE	1:C:66:THR:CA	2.03	0.72
1:C:223:PHE:CD1	1:C:259:GLU:HG2	2.24	0.72
1:D:236:LEU:HD13	1:D:251:GLY:HA2	1.72	0.72
1:G:43:VAL:O	1:G:44:MET:CB	2.36	0.72
1:H:39:ARG:HE	1:H:66:THR:CA	2.03	0.72
1:H:43:VAL:O	1:H:44:MET:CB	2.36	0.72
1:J:236:LEU:HD13	1:J:251:GLY:HA2	1.72	0.72
1:O:236:LEU:HD13	1:O:251:GLY:HA2	1.72	0.72
1:Q:142:LEU:HD11	1:Q:165:ILE:CD1	2.19	0.72
1:A:34:ILE:HD13	1:A:67:LEU:HD13	1.71	0.72
1:C:34:ILE:HD13	1:C:67:LEU:HD13	1.71	0.72
1:F:244:ASP:OD1	1:F:246:GLN:HG3	1.90	0.72
1:G:39:ARG:HE	1:G:66:THR:CA	2.03	0.72
1:K:43:VAL:O	1:K:44:MET:HB3	1.90	0.72
1:K:236:LEU:HD13	1:K:251:GLY:HA2	1.72	0.72
1:L:34:ILE:CA	1:L:54:VAL:HG11	2.20	0.72
1:M:34:ILE:HD13	1:M:67:LEU:HD13	1.71	0.72
1:A:34:ILE:CA	1:A:54:VAL:HG11	2.20	0.71
1:A:301:GLY:H	1:A:335:ARG:HG3	1.55	0.71
1:B:39:ARG:HE	1:B:66:THR:CA	2.03	0.71
1:D:61:LYS:O	1:D:64:ILE:HG22	1.90	0.71
1:F:61:LYS:O	1:F:64:ILE:HG22	1.90	0.71
1:I:208:ILE:CD1	1:I:243:PRO:HD2	2.18	0.71
1:K:39:ARG:HE	1:K:66:THR:CA	2.03	0.71
1:M:43:VAL:O	1:M:44:MET:HB3	1.90	0.71
1:O:61:LYS:O	1:O:64:ILE:HG22	1.90	0.71
1:P:301:GLY:H	1:P:335:ARG:HG3	1.55	0.71
1:Q:301:GLY:H	1:Q:335:ARG:HG3	1.55	0.71
1:R:61:LYS:O	1:R:64:ILE:HG22	1.89	0.71
1:U:61:LYS:O	1:U:64:ILE:HG22	1.90	0.71
1:V:61:LYS:O	1:V:64:ILE:HG22	1.90	0.71
1:V:301:GLY:H	1:V:335:ARG:HG3	1.55	0.71
1:W:39:ARG:HE	1:W:66:THR:CA	2.03	0.71
1:A:43:VAL:O	1:A:44:MET:HB3	1.90	0.71
1:B:61:LYS:O	1:B:64:ILE:HG22	1.90	0.71
1:G:223:PHE:CD1	1:G:259:GLU:HG2	2.24	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:290:ARG:CD	1:J:244:ASP:HB2	2.12	0.71
1:J:287:ILE:HD11	1:L:208:ILE:CD1	2.19	0.71
1:R:43:VAL:O	1:R:44:MET:CB	2.36	0.71
1:T:34:ILE:HD13	1:T:67:LEU:HD13	1.71	0.71
1:W:34:ILE:CA	1:W:54:VAL:HG11	2.20	0.71
1:I:143:TYR:CD2	1:I:346:LEU:HD13	2.26	0.71
1:T:37:ARG:HG3	1:T:38:PRO:CD	2.15	0.71
1:B:37:ARG:HG3	1:B:38:PRO:CD	2.16	0.71
1:B:301:GLY:H	1:B:335:ARG:HG3	1.55	0.71
1:D:39:ARG:HE	1:D:66:THR:CA	2.03	0.71
1:E:223:PHE:CD1	1:E:259:GLU:HG2	2.24	0.71
1:J:34:ILE:CA	1:J:54:VAL:HG11	2.20	0.71
1:J:34:ILE:HD13	1:J:67:LEU:HD13	1.71	0.71
1:K:143:TYR:CD2	1:K:346:LEU:HD13	2.26	0.71
1:Q:61:LYS:O	1:Q:64:ILE:HG22	1.90	0.71
1:V:39:ARG:HE	1:V:66:THR:CA	2.03	0.71
1:F:43:VAL:O	1:F:44:MET:HB3	1.90	0.71
1:K:34:ILE:CA	1:K:54:VAL:HG11	2.20	0.71
1:L:236:LEU:HD13	1:L:251:GLY:HA2	1.72	0.71
1:M:244:ASP:OD1	1:M:246:GLN:HG3	1.90	0.71
1:O:34:ILE:HD13	1:O:67:LEU:HD13	1.71	0.71
1:O:43:VAL:O	1:O:44:MET:HB3	1.90	0.71
1:P:143:TYR:CD2	1:P:346:LEU:HD13	2.26	0.71
1:Q:39:ARG:HE	1:Q:66:THR:CA	2.03	0.71
1:R:143:TYR:CD2	1:R:346:LEU:HD13	2.26	0.71
1:U:34:ILE:CA	1:U:54:VAL:HG11	2.20	0.71
1:V:43:VAL:O	1:V:44:MET:HB3	1.90	0.71
1:B:236:LEU:HD13	1:B:251:GLY:HA2	1.72	0.71
1:B:244:ASP:OD1	1:B:246:GLN:HG3	1.89	0.71
1:E:244:ASP:OD1	1:E:246:GLN:HG3	1.89	0.71
1:F:34:ILE:CA	1:F:54:VAL:HG11	2.20	0.71
1:I:34:ILE:CA	1:I:54:VAL:HG11	2.20	0.71
1:I:236:LEU:HD13	1:I:251:GLY:HA2	1.72	0.71
1:L:61:LYS:O	1:L:64:ILE:HG22	1.90	0.71
1:Q:43:VAL:O	1:Q:44:MET:HB3	1.90	0.71
1:Q:236:LEU:HD13	1:Q:251:GLY:HA2	1.72	0.71
1:S:43:VAL:O	1:S:44:MET:CB	2.36	0.71
1:T:61:LYS:O	1:T:64:ILE:HG22	1.90	0.71
1:T:301:GLY:H	1:T:335:ARG:HG3	1.55	0.71
1:U:244:ASP:OD1	1:U:246:GLN:HG3	1.90	0.71
1:A:244:ASP:OD1	1:A:246:GLN:HG3	1.90	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:143:TYR:CD2	1:G:346:LEU:HD13	2.26	0.71
1:H:301:GLY:H	1:H:335:ARG:HG3	1.55	0.71
1:L:290:ARG:CD	1:N:244:ASP:HB2	2.12	0.71
1:M:39:ARG:HE	1:M:66:THR:CA	2.03	0.71
1:M:143:TYR:CD2	1:M:346:LEU:HD13	2.26	0.71
1:N:143:TYR:CD2	1:N:346:LEU:HD13	2.26	0.71
1:S:61:LYS:O	1:S:64:ILE:HG22	1.90	0.71
1:V:143:TYR:CD2	1:V:346:LEU:HD13	2.26	0.71
1:W:39:ARG:HE	1:W:66:THR:HA	1.52	0.71
1:B:34:ILE:CA	1:B:54:VAL:HG11	2.20	0.71
1:G:301:GLY:H	1:G:335:ARG:HG3	1.55	0.71
1:K:301:GLY:H	1:K:335:ARG:HG3	1.55	0.71
1:P:34:ILE:CA	1:P:54:VAL:HG11	2.20	0.71
1:R:34:ILE:CA	1:R:54:VAL:HG11	2.20	0.71
1:R:34:ILE:HD13	1:R:67:LEU:HD13	1.71	0.71
1:T:143:TYR:CD2	1:T:346:LEU:HD13	2.26	0.71
1:T:208:ILE:CD1	1:T:243:PRO:HD2	2.18	0.71
1:T:244:ASP:OD1	1:T:246:GLN:HG3	1.89	0.71
1:V:244:ASP:OD1	1:V:246:GLN:HG3	1.89	0.71
1:B:143:TYR:CD2	1:B:346:LEU:HD13	2.26	0.71
1:C:244:ASP:OD1	1:C:246:GLN:HG3	1.89	0.71
1:E:39:ARG:HE	1:E:66:THR:CA	2.03	0.71
1:H:61:LYS:O	1:H:64:ILE:HG22	1.90	0.71
1:J:301:GLY:H	1:J:335:ARG:HG3	1.55	0.71
1:N:61:LYS:O	1:N:64:ILE:HG22	1.90	0.71
1:T:43:VAL:O	1:T:44:MET:HB3	1.90	0.71
1:W:43:VAL:O	1:W:44:MET:HB3	1.90	0.71
1:W:236:LEU:HD13	1:W:251:GLY:HA2	1.72	0.71
1:H:43:VAL:O	1:H:44:MET:HB3	1.90	0.71
1:N:34:ILE:CA	1:N:54:VAL:HG11	2.20	0.71
1:R:43:VAL:O	1:R:44:MET:HB3	1.90	0.71
1:S:43:VAL:O	1:S:44:MET:HB3	1.90	0.71
1:S:208:ILE:CD1	1:S:243:PRO:HD2	2.18	0.71
1:U:43:VAL:O	1:U:44:MET:HB3	1.90	0.71
1:B:208:ILE:CD1	1:B:243:PRO:HD2	2.18	0.70
1:C:34:ILE:CA	1:C:54:VAL:HG11	2.20	0.70
1:D:143:TYR:CD2	1:D:346:LEU:HD13	2.26	0.70
1:G:236:LEU:HD13	1:G:251:GLY:HA2	1.72	0.70
1:G:244:ASP:OD1	1:G:246:GLN:HG3	1.90	0.70
1:M:290:ARG:CD	1:O:244:ASP:HB2	2.12	0.70
1:P:61:LYS:O	1:P:64:ILE:HG22	1.90	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:290:ARG:CD	1:U:244:ASP:HB2	2.12	0.70
1:A:143:TYR:CD2	1:A:346:LEU:HD13	2.26	0.70
1:E:143:TYR:CD2	1:E:346:LEU:HD13	2.26	0.70
1:F:39:ARG:HE	1:F:66:THR:CA	2.03	0.70
1:G:34:ILE:CA	1:G:54:VAL:HG11	2.20	0.70
1:M:34:ILE:CA	1:M:54:VAL:HG11	2.20	0.70
1:T:34:ILE:CA	1:T:54:VAL:HG11	2.20	0.70
1:W:143:TYR:CD2	1:W:346:LEU:HD13	2.26	0.70
1:C:236:LEU:CD1	1:C:237:GLU:HG2	2.22	0.70
1:E:169:TYR:CE2	1:G:40:HIS:HB3	2.27	0.70
1:I:236:LEU:CD1	1:I:237:GLU:HG2	2.22	0.70
1:J:61:LYS:O	1:J:64:ILE:HG22	1.90	0.70
1:K:169:TYR:CE2	1:M:40:HIS:HB3	2.27	0.70
1:K:244:ASP:OD1	1:K:246:GLN:HG3	1.90	0.70
1:L:34:ILE:HD13	1:L:67:LEU:HD13	1.71	0.70
1:N:236:LEU:HD13	1:N:251:GLY:HA2	1.72	0.70
1:N:244:ASP:OD1	1:N:246:GLN:HG3	1.90	0.70
1:P:244:ASP:OD1	1:P:246:GLN:HG3	1.89	0.70
1:Q:34:ILE:CA	1:Q:54:VAL:HG11	2.20	0.70
1:T:169:TYR:CE2	1:V:40:HIS:HB3	2.27	0.70
1:V:236:LEU:CD1	1:V:237:GLU:HG2	2.22	0.70
1:B:169:TYR:CE2	1:D:40:HIS:HB3	2.27	0.70
1:E:34:ILE:CA	1:E:54:VAL:HG11	2.20	0.70
1:F:143:TYR:CD2	1:F:346:LEU:HD13	2.26	0.70
1:H:236:LEU:CD1	1:H:237:GLU:HG2	2.22	0.70
1:L:143:TYR:CD2	1:L:346:LEU:HD13	2.26	0.70
1:L:236:LEU:CD1	1:L:237:GLU:HG2	2.22	0.70
1:M:208:ILE:CD1	1:M:243:PRO:HD2	2.18	0.70
1:P:236:LEU:CD1	1:P:237:GLU:HG2	2.22	0.70
1:R:169:TYR:CE2	1:T:40:HIS:HB3	2.27	0.70
1:S:236:LEU:HD13	1:S:251:GLY:HA2	1.72	0.70
1:U:143:TYR:CD2	1:U:346:LEU:HD13	2.26	0.70
1:V:34:ILE:CA	1:V:54:VAL:HG11	2.20	0.70
1:D:34:ILE:CA	1:D:54:VAL:HG11	2.20	0.70
1:F:37:ARG:CG	1:F:38:PRO:HD2	2.20	0.70
1:H:34:ILE:CA	1:H:54:VAL:HG11	2.20	0.70
1:I:37:ARG:CG	1:I:38:PRO:HD2	2.20	0.70
1:I:169:TYR:CE2	1:K:40:HIS:HB3	2.27	0.70
1:L:43:VAL:O	1:L:44:MET:HB3	1.90	0.70
1:M:169:TYR:CE2	1:O:40:HIS:HB3	2.27	0.70
1:P:34:ILE:HD13	1:P:67:LEU:HD13	1.71	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:43:VAL:O	1:P:44:MET:HB3	1.90	0.70
1:U:236:LEU:HD13	1:U:251:GLY:HA2	1.72	0.70
1:A:169:TYR:CE2	1:C:40:HIS:HB3	2.27	0.70
1:D:169:TYR:CE2	1:F:40:HIS:HB3	2.27	0.70
1:D:236:LEU:CD1	1:D:237:GLU:HG2	2.22	0.70
1:I:244:ASP:OD1	1:I:246:GLN:HG3	1.90	0.70
1:J:43:VAL:O	1:J:44:MET:HB3	1.90	0.70
1:M:37:ARG:CG	1:M:38:PRO:HD2	2.20	0.70
1:M:236:LEU:CD1	1:M:237:GLU:HG2	2.22	0.70
1:O:39:ARG:HE	1:O:66:THR:CA	2.03	0.70
1:O:143:TYR:CD2	1:O:346:LEU:HD13	2.26	0.70
1:U:236:LEU:CD1	1:U:237:GLU:HG2	2.22	0.70
1:C:39:ARG:HE	1:C:66:THR:HA	1.52	0.70
1:C:143:TYR:CD2	1:C:346:LEU:HD13	2.26	0.70
1:E:236:LEU:HD13	1:E:251:GLY:HA2	1.72	0.70
1:H:208:ILE:CD1	1:H:243:PRO:HD2	2.18	0.70
1:N:43:VAL:O	1:N:44:MET:HB3	1.90	0.70
1:R:236:LEU:CD1	1:R:237:GLU:HG2	2.22	0.70
1:R:244:ASP:OD1	1:R:246:GLN:HG3	1.90	0.70
1:S:143:TYR:CD2	1:S:346:LEU:HD13	2.26	0.70
1:A:236:LEU:HD13	1:A:251:GLY:HA2	1.72	0.70
1:C:169:TYR:CE2	1:E:40:HIS:HB3	2.27	0.70
1:J:143:TYR:CE1	1:L:45:VAL:HG21	2.27	0.70
1:O:236:LEU:CD1	1:O:237:GLU:HG2	2.22	0.70
1:P:143:TYR:CE1	1:R:45:VAL:HG21	2.27	0.70
1:P:169:TYR:CE2	1:R:40:HIS:HB3	2.27	0.70
1:R:301:GLY:H	1:R:335:ARG:HG3	1.55	0.70
1:S:34:ILE:CA	1:S:54:VAL:HG11	2.20	0.70
1:A:143:TYR:CE1	1:C:45:VAL:HG21	2.27	0.70
1:B:290:ARG:CD	1:D:244:ASP:HB2	2.12	0.70
1:C:236:LEU:HD13	1:C:251:GLY:HA2	1.72	0.70
1:K:34:ILE:HG22	1:K:35:VAL:N	2.07	0.70
1:Q:236:LEU:CD1	1:Q:237:GLU:HG2	2.22	0.70
1:U:143:TYR:CE1	1:W:45:VAL:HG21	2.27	0.70
1:C:208:ILE:CD1	1:C:243:PRO:HD2	2.18	0.70
1:E:143:TYR:CE1	1:G:45:VAL:HG21	2.27	0.70
1:F:169:TYR:CE2	1:H:40:HIS:HB3	2.27	0.70
1:G:34:ILE:HG22	1:G:35:VAL:N	2.07	0.70
1:G:169:TYR:CE2	1:I:40:HIS:HB3	2.27	0.70
1:G:236:LEU:CD1	1:G:237:GLU:HG2	2.22	0.70
1:H:143:TYR:CD2	1:H:346:LEU:HD13	2.26	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:301:GLY:H	1:I:335:ARG:HG3	1.55	0.70
1:N:34:ILE:HG22	1:N:35:VAL:N	2.07	0.70
1:N:34:ILE:HD13	1:N:67:LEU:HD13	1.71	0.70
1:O:34:ILE:CA	1:O:54:VAL:HG11	2.20	0.70
1:O:143:TYR:CE1	1:Q:45:VAL:HG21	2.27	0.70
1:Q:37:ARG:CG	1:Q:38:PRO:HD2	2.20	0.70
1:S:236:LEU:CD1	1:S:237:GLU:HG2	2.22	0.70
1:A:37:ARG:CG	1:A:38:PRO:HD2	2.20	0.69
1:C:34:ILE:HG22	1:C:35:VAL:N	2.07	0.69
1:D:143:TYR:CE1	1:F:45:VAL:HG21	2.27	0.69
1:E:236:LEU:CD1	1:E:237:GLU:HG2	2.22	0.69
1:P:236:LEU:HD13	1:P:251:GLY:HA2	1.72	0.69
1:S:143:TYR:CE1	1:U:45:VAL:HG21	2.27	0.69
1:V:34:ILE:HG22	1:V:35:VAL:N	2.07	0.69
1:H:143:TYR:CE1	1:J:45:VAL:HG21	2.27	0.69
1:N:208:ILE:CD1	1:N:243:PRO:HD2	2.18	0.69
1:O:169:TYR:CE2	1:Q:40:HIS:HB3	2.27	0.69
1:H:169:TYR:CE2	1:J:40:HIS:HB3	2.27	0.69
1:J:34:ILE:HG22	1:J:35:VAL:N	2.07	0.69
1:Q:143:TYR:CD2	1:Q:346:LEU:HD13	2.26	0.69
1:R:34:ILE:HG22	1:R:35:VAL:N	2.07	0.69
1:R:37:ARG:CG	1:R:38:PRO:HD2	2.20	0.69
1:U:50:LYS:HG3	1:U:53:TYR:CE2	2.28	0.69
1:A:236:LEU:CD1	1:A:237:GLU:HG2	2.22	0.69
1:B:236:LEU:CD1	1:B:237:GLU:HG2	2.22	0.69
1:E:34:ILE:HG22	1:E:35:VAL:N	2.07	0.69
1:J:143:TYR:CD2	1:J:346:LEU:HD13	2.26	0.69
1:K:143:TYR:CE1	1:M:45:VAL:HG21	2.27	0.69
1:K:236:LEU:CD1	1:K:237:GLU:HG2	2.22	0.69
1:L:143:TYR:CE1	1:N:45:VAL:HG21	2.27	0.69
1:N:169:TYR:CE2	1:P:40:HIS:HB3	2.27	0.69
1:O:34:ILE:HG22	1:O:35:VAL:N	2.07	0.69
1:T:236:LEU:CD1	1:T:237:GLU:HG2	2.22	0.69
1:F:50:LYS:HG3	1:F:53:TYR:CE2	2.28	0.69
1:H:50:LYS:HG3	1:H:53:TYR:CE2	2.28	0.69
1:H:58:ALA:CB	1:H:65:LEU:HD22	2.23	0.69
1:J:50:LYS:HG3	1:J:53:TYR:CE2	2.28	0.69
1:J:169:TYR:CE2	1:L:40:HIS:HB3	2.27	0.69
1:Q:50:LYS:HG3	1:Q:53:TYR:CE2	2.28	0.69
1:Q:143:TYR:CE1	1:S:45:VAL:HG21	2.27	0.69
1:R:58:ALA:CB	1:R:65:LEU:HD22	2.23	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:236:LEU:HD13	1:R:251:GLY:HA2	1.72	0.69
1:S:50:LYS:HG3	1:S:53:TYR:CE2	2.28	0.69
1:A:50:LYS:HG3	1:A:53:TYR:CE2	2.28	0.69
1:B:50:LYS:HG3	1:B:53:TYR:CE2	2.28	0.69
1:D:50:LYS:HG3	1:D:53:TYR:CE2	2.28	0.69
1:G:37:ARG:CG	1:G:38:PRO:HD2	2.20	0.69
1:G:58:ALA:CB	1:G:65:LEU:HD22	2.23	0.69
1:L:50:LYS:HG3	1:L:53:TYR:CE2	2.28	0.69
1:N:50:LYS:HG3	1:N:53:TYR:CE2	2.28	0.69
1:N:143:TYR:CE1	1:P:45:VAL:HG21	2.27	0.69
1:S:58:ALA:CB	1:S:65:LEU:HD22	2.23	0.69
1:T:143:TYR:CE1	1:V:45:VAL:HG21	2.27	0.69
1:U:169:TYR:CE2	1:W:40:HIS:HB3	2.27	0.69
1:W:50:LYS:HG3	1:W:53:TYR:CE2	2.28	0.69
1:A:58:ALA:CB	1:A:65:LEU:HD22	2.23	0.69
1:A:208:ILE:CD1	1:A:243:PRO:HD2	2.18	0.69
1:J:236:LEU:CD1	1:J:237:GLU:HG2	2.22	0.69
1:N:58:ALA:CB	1:N:65:LEU:HD22	2.23	0.69
1:N:236:LEU:CD1	1:N:237:GLU:HG2	2.22	0.69
1:P:50:LYS:HG3	1:P:53:TYR:CE2	2.28	0.69
1:U:34:ILE:HG22	1:U:35:VAL:N	2.07	0.69
1:B:58:ALA:CB	1:B:65:LEU:HD22	2.23	0.69
1:B:143:TYR:CE1	1:D:45:VAL:HG21	2.27	0.69
1:C:50:LYS:HG3	1:C:53:TYR:CE2	2.28	0.69
1:C:58:ALA:CB	1:C:65:LEU:HD22	2.23	0.69
1:C:143:TYR:CE1	1:E:45:VAL:HG21	2.27	0.69
1:D:34:ILE:HG22	1:D:35:VAL:N	2.07	0.69
1:D:58:ALA:CB	1:D:65:LEU:HD22	2.23	0.69
1:E:50:LYS:HG3	1:E:53:TYR:CE2	2.28	0.69
1:F:143:TYR:CE1	1:H:45:VAL:HG21	2.27	0.69
1:F:236:LEU:CD1	1:F:237:GLU:HG2	2.22	0.69
1:G:143:TYR:CE1	1:I:45:VAL:HG21	2.27	0.69
1:I:34:ILE:HG22	1:I:35:VAL:N	2.07	0.69
1:I:301:GLY:N	1:I:335:ARG:HG3	2.08	0.69
1:K:50:LYS:HG3	1:K:53:TYR:CE2	2.28	0.69
1:K:301:GLY:N	1:K:335:ARG:HG3	2.08	0.69
1:L:169:TYR:CE2	1:N:40:HIS:HB3	2.27	0.69
1:M:50:LYS:HG3	1:M:53:TYR:CE2	2.28	0.69
1:M:301:GLY:N	1:M:335:ARG:HG3	2.08	0.69
1:N:208:ILE:HD11	1:N:243:PRO:CG	2.23	0.69
1:O:50:LYS:HG3	1:O:53:TYR:CE2	2.28	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:208:ILE:HD11	1:P:243:PRO:CG	2.23	0.69
1:P:301:GLY:N	1:P:335:ARG:HG3	2.08	0.69
1:R:143:TYR:CE1	1:T:45:VAL:HG21	2.27	0.69
1:S:34:ILE:HG22	1:S:35:VAL:N	2.07	0.69
1:S:169:TYR:CE2	1:U:40:HIS:HB3	2.27	0.69
1:S:301:GLY:N	1:S:335:ARG:HG3	2.08	0.69
1:T:50:LYS:HG3	1:T:53:TYR:CE2	2.28	0.69
1:T:236:LEU:HD13	1:T:251:GLY:HA2	1.72	0.69
1:U:58:ALA:CB	1:U:65:LEU:HD22	2.23	0.69
1:V:236:LEU:HD13	1:V:251:GLY:HA2	1.72	0.69
1:W:236:LEU:CD1	1:W:237:GLU:HG2	2.22	0.69
1:D:22:ALA:HB1	1:D:348:SER:HB3	1.75	0.69
1:D:301:GLY:N	1:D:335:ARG:HG3	2.08	0.69
1:H:22:ALA:HB1	1:H:348:SER:HB3	1.75	0.69
1:L:58:ALA:CB	1:L:65:LEU:HD22	2.23	0.69
1:L:208:ILE:HD11	1:L:243:PRO:CG	2.23	0.69
1:M:58:ALA:CB	1:M:65:LEU:HD22	2.23	0.69
1:R:50:LYS:HG3	1:R:53:TYR:CE2	2.28	0.69
1:A:34:ILE:HG22	1:A:35:VAL:N	2.07	0.69
1:G:50:LYS:HG3	1:G:53:TYR:CE2	2.28	0.69
1:I:50:LYS:HG3	1:I:53:TYR:CE2	2.28	0.69
1:I:143:TYR:CE1	1:K:45:VAL:HG21	2.27	0.69
1:O:22:ALA:HB1	1:O:348:SER:HB3	1.75	0.69
1:O:58:ALA:CB	1:O:65:LEU:HD22	2.23	0.69
1:P:290:ARG:CD	1:R:244:ASP:HB2	2.12	0.69
1:Q:169:TYR:CE2	1:S:40:HIS:HB3	2.27	0.69
1:V:50:LYS:HG3	1:V:53:TYR:CE2	2.28	0.69
1:W:208:ILE:HD11	1:W:243:PRO:CG	2.23	0.69
1:A:301:GLY:N	1:A:335:ARG:HG3	2.08	0.68
1:E:58:ALA:CB	1:E:65:LEU:HD22	2.23	0.68
1:G:301:GLY:N	1:G:335:ARG:HG3	2.08	0.68
1:J:58:ALA:CB	1:J:65:LEU:HD22	2.23	0.68
1:J:173:HIS:CE1	1:K:268:GLY:CA	2.77	0.68
1:J:301:GLY:N	1:J:335:ARG:HG3	2.08	0.68
1:M:7:ALA:HB1	1:M:356:TRP:CH2	2.29	0.68
1:O:208:ILE:HD11	1:O:243:PRO:CG	2.23	0.68
1:R:208:ILE:HD11	1:R:243:PRO:CG	2.23	0.68
1:V:7:ALA:HB1	1:V:356:TRP:CH2	2.29	0.68
1:V:58:ALA:CB	1:V:65:LEU:HD22	2.23	0.68
1:W:34:ILE:HG22	1:W:35:VAL:N	2.07	0.68
1:W:58:ALA:CB	1:W:65:LEU:HD22	2.23	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:34:ILE:HG22	1:F:35:VAL:N	2.07	0.68
1:I:7:ALA:HB1	1:I:356:TRP:CH2	2.29	0.68
1:I:58:ALA:CB	1:I:65:LEU:HD22	2.23	0.68
1:M:208:ILE:HD11	1:M:243:PRO:CG	2.23	0.68
1:N:22:ALA:HB1	1:N:348:SER:HB3	1.76	0.68
1:T:58:ALA:CB	1:T:65:LEU:HD22	2.23	0.68
1:B:7:ALA:HB1	1:B:356:TRP:CH2	2.29	0.68
1:C:22:ALA:HB1	1:C:348:SER:HB3	1.75	0.68
1:C:362:TYR:CE1	1:C:367:PRO:HB3	2.28	0.68
1:D:7:ALA:HB1	1:D:356:TRP:CH2	2.29	0.68
1:E:290:ARG:CD	1:G:244:ASP:HB2	2.12	0.68
1:F:173:HIS:CE1	1:G:268:GLY:CA	2.77	0.68
1:H:34:ILE:HG22	1:H:35:VAL:N	2.07	0.68
1:K:7:ALA:HB1	1:K:356:TRP:CH2	2.29	0.68
1:K:208:ILE:HD11	1:K:243:PRO:CG	2.23	0.68
1:N:173:HIS:CE1	1:O:268:GLY:CA	2.77	0.68
1:O:7:ALA:HB1	1:O:356:TRP:CH2	2.29	0.68
1:O:301:GLY:N	1:O:335:ARG:HG3	2.08	0.68
1:R:301:GLY:N	1:R:335:ARG:HG3	2.08	0.68
1:T:7:ALA:HB1	1:T:356:TRP:CH2	2.29	0.68
1:U:208:ILE:HD11	1:U:243:PRO:CG	2.23	0.68
1:C:173:HIS:CE1	1:D:268:GLY:CA	2.77	0.68
1:J:208:ILE:HD11	1:J:243:PRO:CG	2.23	0.68
1:L:34:ILE:HG22	1:L:35:VAL:N	2.07	0.68
1:Q:34:ILE:HG22	1:Q:35:VAL:N	2.07	0.68
1:S:22:ALA:HB1	1:S:348:SER:HB3	1.75	0.68
1:U:22:ALA:HB1	1:U:348:SER:HB3	1.75	0.68
1:A:173:HIS:CE1	1:B:268:GLY:CA	2.77	0.68
1:A:362:TYR:CE1	1:A:367:PRO:HB3	2.28	0.68
1:F:7:ALA:HB1	1:F:356:TRP:CH2	2.29	0.68
1:L:22:ALA:HB1	1:L:348:SER:HB3	1.75	0.68
1:M:143:TYR:CE1	1:O:45:VAL:HG21	2.27	0.68
1:P:58:ALA:CB	1:P:65:LEU:HD22	2.23	0.68
1:Q:208:ILE:HD11	1:Q:243:PRO:CG	2.23	0.68
1:T:301:GLY:N	1:T:335:ARG:HG3	2.08	0.68
1:A:22:ALA:HB1	1:A:348:SER:HB3	1.75	0.68
1:B:301:GLY:N	1:B:335:ARG:HG3	2.08	0.68
1:E:208:ILE:HD11	1:E:243:PRO:CG	2.23	0.68
1:E:301:GLY:N	1:E:335:ARG:HG3	2.08	0.68
1:G:7:ALA:HB1	1:G:356:TRP:CH2	2.29	0.68
1:G:208:ILE:HD11	1:G:243:PRO:CG	2.23	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:362:TYR:O	1:G:366:GLY:HA2	1.94	0.68
1:N:38:PRO:CG	1:N:49:GLN:HE22	2.07	0.68
1:R:173:HIS:CE1	1:S:268:GLY:CA	2.77	0.68
1:S:173:HIS:CE1	1:T:268:GLY:CA	2.77	0.68
1:T:34:ILE:HG22	1:T:35:VAL:N	2.07	0.68
1:U:37:ARG:CG	1:U:38:PRO:HD2	2.20	0.68
1:V:301:GLY:N	1:V:335:ARG:HG3	2.08	0.68
1:B:34:ILE:HG22	1:B:35:VAL:N	2.07	0.68
1:C:7:ALA:HB1	1:C:356:TRP:CH2	2.29	0.68
1:C:208:ILE:HD11	1:C:243:PRO:CG	2.23	0.68
1:E:362:TYR:O	1:E:366:GLY:HA2	1.94	0.68
1:F:58:ALA:CB	1:F:65:LEU:HD22	2.23	0.68
1:F:301:GLY:N	1:F:335:ARG:HG3	2.08	0.68
1:H:362:TYR:O	1:H:366:GLY:HA2	1.94	0.68
1:I:22:ALA:HB1	1:I:348:SER:HB3	1.75	0.68
1:J:22:ALA:HB1	1:J:348:SER:HB3	1.75	0.68
1:K:58:ALA:CB	1:K:65:LEU:HD22	2.23	0.68
1:L:38:PRO:CG	1:L:49:GLN:HE22	2.07	0.68
1:O:362:TYR:CE1	1:O:367:PRO:HB3	2.28	0.68
1:P:173:HIS:CE1	1:Q:268:GLY:CA	2.77	0.68
1:T:362:TYR:O	1:T:366:GLY:HA2	1.94	0.68
1:A:7:ALA:HB1	1:A:356:TRP:CH2	2.29	0.68
1:E:7:ALA:HB1	1:E:356:TRP:CH2	2.29	0.68
1:E:173:HIS:CE1	1:F:268:GLY:CA	2.77	0.68
1:G:236:LEU:HD22	1:G:252:ASN:N	2.09	0.68
1:I:236:LEU:HD22	1:I:252:ASN:N	2.09	0.68
1:J:38:PRO:CG	1:J:49:GLN:HE22	2.07	0.68
1:N:301:GLY:N	1:N:335:ARG:HG3	2.08	0.68
1:P:34:ILE:HG22	1:P:35:VAL:N	2.07	0.68
1:P:38:PRO:CG	1:P:49:GLN:HE22	2.07	0.68
1:Q:58:ALA:CB	1:Q:65:LEU:HD22	2.23	0.68
1:Q:301:GLY:N	1:Q:335:ARG:HG3	2.08	0.68
1:Q:362:TYR:O	1:Q:366:GLY:HA2	1.94	0.68
1:R:7:ALA:HB1	1:R:356:TRP:CH2	2.29	0.68
1:S:208:ILE:HD11	1:S:243:PRO:CG	2.23	0.68
1:S:362:TYR:O	1:S:366:GLY:HA2	1.94	0.68
1:U:301:GLY:N	1:U:335:ARG:HG3	2.08	0.68
1:A:236:LEU:HD22	1:A:252:ASN:N	2.09	0.68
1:D:110:LEU:HD12	1:D:177:ARG:NH1	2.09	0.68
1:D:362:TYR:O	1:D:366:GLY:HA2	1.94	0.68
1:F:110:LEU:HD12	1:F:177:ARG:NH1	2.09	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:362:TYR:CE1	1:F:367:PRO:HB3	2.28	0.68
1:I:208:ILE:HD11	1:I:243:PRO:CG	2.23	0.68
1:K:110:LEU:HD12	1:K:177:ARG:NH1	2.09	0.68
1:M:34:ILE:HG22	1:M:35:VAL:N	2.07	0.68
1:M:362:TYR:CE1	1:M:367:PRO:HB3	2.28	0.68
1:P:236:LEU:HD22	1:P:252:ASN:N	2.09	0.68
1:Q:362:TYR:CE1	1:Q:367:PRO:HB3	2.28	0.68
1:R:22:ALA:HB1	1:R:348:SER:HB3	1.75	0.68
1:R:236:LEU:HD22	1:R:252:ASN:N	2.09	0.68
1:T:22:ALA:HB1	1:T:348:SER:HB3	1.75	0.68
1:T:208:ILE:HD11	1:T:243:PRO:CG	2.23	0.68
1:A:110:LEU:HD12	1:A:177:ARG:NH1	2.09	0.68
1:B:22:ALA:HB1	1:B:348:SER:HB3	1.75	0.68
1:B:362:TYR:O	1:B:366:GLY:HA2	1.94	0.68
1:D:362:TYR:CE1	1:D:367:PRO:HB3	2.28	0.68
1:E:110:LEU:HD12	1:E:177:ARG:NH1	2.09	0.68
1:H:110:LEU:HD12	1:H:177:ARG:NH1	2.09	0.68
1:H:301:GLY:N	1:H:335:ARG:HG3	2.08	0.68
1:J:37:ARG:CG	1:J:38:PRO:HD2	2.20	0.68
1:L:173:HIS:CE1	1:M:268:GLY:CA	2.77	0.68
1:L:301:GLY:N	1:L:335:ARG:HG3	2.08	0.68
1:M:110:LEU:HD12	1:M:177:ARG:NH1	2.09	0.68
1:P:110:LEU:HD12	1:P:177:ARG:NH1	2.09	0.68
1:Q:22:ALA:HB1	1:Q:348:SER:HB3	1.75	0.68
1:Q:208:ILE:CD1	1:Q:243:PRO:HD2	2.18	0.68
1:R:38:PRO:CG	1:R:49:GLN:HE22	2.07	0.68
1:U:173:HIS:CE1	1:V:268:GLY:CA	2.77	0.68
1:U:362:TYR:O	1:U:366:GLY:HA2	1.94	0.68
1:W:22:ALA:HB1	1:W:348:SER:HB3	1.75	0.68
1:W:362:TYR:O	1:W:366:GLY:HA2	1.94	0.68
1:B:58:ALA:HB1	1:B:65:LEU:CD2	2.24	0.67
1:B:110:LEU:HD12	1:B:177:ARG:NH1	2.09	0.67
1:B:208:ILE:HD11	1:B:243:PRO:CG	2.23	0.67
1:F:34:ILE:HD13	1:F:67:LEU:HD22	1.77	0.67
1:H:7:ALA:HB1	1:H:356:TRP:CH2	2.29	0.67
1:H:362:TYR:CE1	1:H:367:PRO:HB3	2.28	0.67
1:I:362:TYR:O	1:I:366:GLY:HA2	1.94	0.67
1:K:22:ALA:HB1	1:K:348:SER:HB3	1.75	0.67
1:K:236:LEU:HD22	1:K:252:ASN:N	2.09	0.67
1:M:38:PRO:CG	1:M:49:GLN:HE22	2.07	0.67
1:N:236:LEU:HD22	1:N:252:ASN:N	2.09	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:38:PRO:CG	1:O:49:GLN:HE22	2.07	0.67
1:Q:7:ALA:HB1	1:Q:356:TRP:CH2	2.29	0.67
1:T:362:TYR:CE1	1:T:367:PRO:HB3	2.28	0.67
1:V:362:TYR:CE1	1:V:367:PRO:HB3	2.28	0.67
1:C:236:LEU:HD22	1:C:252:ASN:N	2.09	0.67
1:C:301:GLY:N	1:C:335:ARG:HG3	2.08	0.67
1:D:38:PRO:CG	1:D:49:GLN:HE22	2.07	0.67
1:D:58:ALA:HB1	1:D:65:LEU:CD2	2.25	0.67
1:H:208:ILE:HD11	1:H:243:PRO:CG	2.23	0.67
1:K:37:ARG:CG	1:K:38:PRO:HD2	2.20	0.67
1:L:362:TYR:O	1:L:366:GLY:HA2	1.94	0.67
1:O:110:LEU:HD12	1:O:177:ARG:NH1	2.09	0.67
1:P:362:TYR:O	1:P:366:GLY:HA2	1.94	0.67
1:T:143:TYR:O	1:T:143:TYR:CD1	2.48	0.67
1:V:173:HIS:CE1	1:W:268:GLY:CA	2.77	0.67
1:V:362:TYR:O	1:V:366:GLY:HA2	1.94	0.67
1:A:208:ILE:HD11	1:A:243:PRO:CG	2.23	0.67
1:A:362:TYR:O	1:A:366:GLY:HA2	1.94	0.67
1:B:173:HIS:CE1	1:C:268:GLY:CA	2.77	0.67
1:B:236:LEU:HD22	1:B:252:ASN:N	2.09	0.67
1:B:370:VAL:O	1:B:370:VAL:HG13	1.95	0.67
1:E:143:TYR:O	1:E:143:TYR:CD1	2.48	0.67
1:G:58:ALA:HB1	1:G:65:LEU:CD2	2.25	0.67
1:G:208:ILE:CD1	1:G:243:PRO:HD2	2.18	0.67
1:H:38:PRO:CG	1:H:49:GLN:HE22	2.07	0.67
1:I:58:ALA:HB1	1:I:65:LEU:CD2	2.24	0.67
1:I:173:HIS:CE1	1:J:268:GLY:CA	2.77	0.67
1:J:362:TYR:O	1:J:366:GLY:HA2	1.94	0.67
1:L:104:LEU:HB2	1:L:356:TRP:CZ3	2.30	0.67
1:M:22:ALA:HB1	1:M:348:SER:HB3	1.76	0.67
1:N:104:LEU:HB2	1:N:356:TRP:CZ3	2.30	0.67
1:N:362:TYR:O	1:N:366:GLY:HA2	1.94	0.67
1:O:173:HIS:CE1	1:P:268:GLY:CA	2.77	0.67
1:O:362:TYR:O	1:O:366:GLY:HA2	1.94	0.67
1:Q:34:ILE:HD13	1:Q:67:LEU:HD22	1.77	0.67
1:T:110:LEU:HD12	1:T:177:ARG:NH1	2.09	0.67
1:U:34:ILE:HD13	1:U:67:LEU:HD22	1.77	0.67
1:B:38:PRO:CG	1:B:49:GLN:HE22	2.07	0.67
1:E:236:LEU:HD22	1:E:252:ASN:N	2.09	0.67
1:F:58:ALA:HB1	1:F:65:LEU:CD2	2.25	0.67
1:H:34:ILE:HD13	1:H:67:LEU:HD22	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:173:HIS:CE1	1:I:268:GLY:CA	2.77	0.67
1:I:143:TYR:O	1:I:143:TYR:CD1	2.48	0.67
1:K:104:LEU:HB2	1:K:356:TRP:CZ3	2.30	0.67
1:L:110:LEU:HD12	1:L:177:ARG:NH1	2.09	0.67
1:P:104:LEU:HB2	1:P:356:TRP:CZ3	2.30	0.67
1:P:143:TYR:O	1:P:143:TYR:CD1	2.48	0.67
1:R:362:TYR:CE1	1:R:367:PRO:HB3	2.28	0.67
1:U:370:VAL:O	1:U:370:VAL:HG13	1.95	0.67
1:W:370:VAL:HG13	1:W:370:VAL:O	1.95	0.67
1:B:34:ILE:HD13	1:B:67:LEU:HD22	1.77	0.67
1:C:362:TYR:O	1:C:366:GLY:HA2	1.94	0.67
1:D:34:ILE:HD13	1:D:67:LEU:HD22	1.77	0.67
1:E:58:ALA:HB1	1:E:65:LEU:CD2	2.25	0.67
1:F:208:ILE:CD1	1:F:243:PRO:HD2	2.18	0.67
1:G:22:ALA:HB1	1:G:348:SER:HB3	1.75	0.67
1:I:110:LEU:HD12	1:I:177:ARG:NH1	2.09	0.67
1:J:7:ALA:HB1	1:J:356:TRP:CH2	2.29	0.67
1:J:104:LEU:HB2	1:J:356:TRP:CZ3	2.30	0.67
1:K:58:ALA:HB1	1:K:65:LEU:CD2	2.25	0.67
1:K:153:LEU:HD23	1:K:299:MET:HE3	1.76	0.67
1:K:362:TYR:CE1	1:K:367:PRO:HB3	2.28	0.67
1:M:58:ALA:HB1	1:M:65:LEU:CD2	2.24	0.67
1:Q:110:LEU:HD12	1:Q:177:ARG:NH1	2.09	0.67
1:R:104:LEU:HB2	1:R:356:TRP:CZ3	2.30	0.67
1:T:38:PRO:CG	1:T:49:GLN:HE22	2.07	0.67
1:V:37:ARG:CG	1:V:38:PRO:HD2	2.20	0.67
1:V:38:PRO:CG	1:V:49:GLN:HE22	2.07	0.67
1:W:7:ALA:HB1	1:W:356:TRP:CH2	2.29	0.67
1:W:34:ILE:HD13	1:W:67:LEU:HD22	1.77	0.67
1:B:362:TYR:CE1	1:B:367:PRO:HB3	2.28	0.67
1:D:37:ARG:CG	1:D:38:PRO:HD2	2.20	0.67
1:F:38:PRO:CG	1:F:49:GLN:HE22	2.07	0.67
1:I:104:LEU:HB2	1:I:356:TRP:CZ3	2.30	0.67
1:I:153:LEU:HD23	1:I:299:MET:HE3	1.76	0.67
1:J:110:LEU:HD12	1:J:177:ARG:NH1	2.09	0.67
1:J:370:VAL:O	1:J:370:VAL:HG13	1.95	0.67
1:K:38:PRO:CG	1:K:49:GLN:HE22	2.07	0.67
1:K:362:TYR:O	1:K:366:GLY:HA2	1.94	0.67
1:M:104:LEU:HB2	1:M:356:TRP:CZ3	2.30	0.67
1:M:370:VAL:O	1:M:370:VAL:HG13	1.95	0.67
1:O:58:ALA:HB1	1:O:65:LEU:CD2	2.25	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:7:ALA:HB1	1:P:356:TRP:CH2	2.29	0.67
1:S:34:ILE:HD13	1:S:67:LEU:HD22	1.77	0.67
1:T:236:LEU:HD22	1:T:252:ASN:N	2.09	0.67
1:U:7:ALA:HB1	1:U:356:TRP:CH2	2.29	0.67
1:V:153:LEU:HD23	1:V:299:MET:HE3	1.76	0.67
1:V:208:ILE:HD11	1:V:243:PRO:CG	2.23	0.67
1:W:362:TYR:CE1	1:W:367:PRO:HB3	2.28	0.67
1:A:143:TYR:O	1:A:143:TYR:CD1	2.48	0.67
1:D:208:ILE:HD11	1:D:243:PRO:CG	2.23	0.67
1:D:236:LEU:HD22	1:D:252:ASN:N	2.09	0.67
1:F:22:ALA:HB1	1:F:348:SER:HB3	1.75	0.67
1:J:34:ILE:HD13	1:J:67:LEU:HD22	1.77	0.67
1:M:34:ILE:HD13	1:M:67:LEU:HD22	1.77	0.67
1:M:362:TYR:O	1:M:366:GLY:HA2	1.94	0.67
1:O:34:ILE:HD13	1:O:67:LEU:HD22	1.77	0.67
1:Q:58:ALA:HB1	1:Q:65:LEU:CD2	2.24	0.67
1:Q:236:LEU:HD22	1:Q:252:ASN:N	2.09	0.67
1:R:208:ILE:CD1	1:R:243:PRO:HD2	2.18	0.67
1:V:22:ALA:HB1	1:V:348:SER:HB3	1.75	0.67
1:C:38:PRO:CG	1:C:49:GLN:HE22	2.07	0.67
1:D:173:HIS:CE1	1:E:268:GLY:CA	2.77	0.67
1:D:370:VAL:HG13	1:D:370:VAL:O	1.95	0.67
1:E:22:ALA:HB1	1:E:348:SER:HB3	1.75	0.67
1:G:104:LEU:HB2	1:G:356:TRP:CZ3	2.30	0.67
1:G:173:HIS:CE1	1:H:268:GLY:CA	2.77	0.67
1:H:104:LEU:HB2	1:H:356:TRP:CZ3	2.30	0.67
1:J:362:TYR:CE1	1:J:367:PRO:HB3	2.29	0.67
1:K:173:HIS:CE1	1:L:268:GLY:CA	2.77	0.67
1:K:290:ARG:CD	1:M:244:ASP:HB2	2.12	0.67
1:M:236:LEU:HD22	1:M:252:ASN:N	2.09	0.67
1:N:143:TYR:O	1:N:143:TYR:CD1	2.48	0.67
1:O:104:LEU:HB2	1:O:356:TRP:CZ3	2.30	0.67
1:O:370:VAL:HG13	1:O:370:VAL:O	1.95	0.67
1:P:22:ALA:HB1	1:P:348:SER:HB3	1.75	0.67
1:Q:38:PRO:CG	1:Q:49:GLN:HE22	2.07	0.67
1:R:58:ALA:HB1	1:R:65:LEU:CD2	2.25	0.67
1:R:110:LEU:HD12	1:R:177:ARG:NH1	2.09	0.67
1:T:104:LEU:HB2	1:T:356:TRP:CZ3	2.30	0.67
1:A:38:PRO:CG	1:A:49:GLN:HE22	2.07	0.67
1:C:143:TYR:O	1:C:143:TYR:CD1	2.48	0.67
1:E:104:LEU:HB2	1:E:356:TRP:CZ3	2.30	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:153:LEU:HD23	1:E:299:MET:HE3	1.76	0.67
1:J:236:LEU:HD22	1:J:252:ASN:N	2.09	0.67
1:L:7:ALA:HB1	1:L:356:TRP:CH2	2.29	0.67
1:L:236:LEU:HD22	1:L:252:ASN:N	2.09	0.67
1:M:173:HIS:CE1	1:N:268:GLY:CA	2.77	0.67
1:P:37:ARG:CG	1:P:38:PRO:HD2	2.20	0.67
1:P:58:ALA:HB1	1:P:65:LEU:CD2	2.25	0.67
1:P:362:TYR:CE1	1:P:367:PRO:HB3	2.28	0.67
1:S:7:ALA:HB1	1:S:356:TRP:CH2	2.29	0.67
1:S:39:ARG:NE	1:S:66:THR:HB	2.09	0.67
1:U:153:LEU:HD23	1:U:299:MET:HE3	1.76	0.67
1:V:104:LEU:HB2	1:V:356:TRP:CZ3	2.30	0.67
1:V:110:LEU:HD12	1:V:177:ARG:NH1	2.09	0.67
1:W:236:LEU:HD22	1:W:252:ASN:N	2.09	0.67
1:W:301:GLY:N	1:W:335:ARG:HG3	2.08	0.67
1:C:58:ALA:HB1	1:C:65:LEU:CD2	2.25	0.67
1:F:208:ILE:HD11	1:F:243:PRO:CG	2.23	0.67
1:F:362:TYR:O	1:F:366:GLY:HA2	1.94	0.67
1:G:153:LEU:HD23	1:G:299:MET:HE3	1.76	0.67
1:M:143:TYR:O	1:M:143:TYR:CD1	2.48	0.67
1:M:153:LEU:HD23	1:M:299:MET:HE3	1.76	0.67
1:O:236:LEU:HD22	1:O:252:ASN:N	2.09	0.67
1:Q:39:ARG:NE	1:Q:66:THR:HB	2.09	0.67
1:S:110:LEU:HD12	1:S:177:ARG:NH1	2.09	0.67
1:S:236:LEU:HD22	1:S:252:ASN:N	2.09	0.67
1:T:58:ALA:HB1	1:T:65:LEU:CD2	2.25	0.67
1:U:236:LEU:HD22	1:U:252:ASN:N	2.09	0.67
1:W:110:LEU:HD12	1:W:177:ARG:NH1	2.09	0.67
1:G:110:LEU:HD12	1:G:177:ARG:NH1	2.09	0.66
1:H:58:ALA:HB1	1:H:65:LEU:CD2	2.25	0.66
1:H:370:VAL:O	1:H:370:VAL:HG13	1.95	0.66
1:L:34:ILE:HD13	1:L:67:LEU:HD22	1.77	0.66
1:L:143:TYR:O	1:L:143:TYR:CD1	2.48	0.66
1:L:370:VAL:HG13	1:L:370:VAL:O	1.95	0.66
1:N:110:LEU:HD12	1:N:177:ARG:NH1	2.09	0.66
1:O:34:ILE:HA	1:O:68:LYS:O	1.96	0.66
1:Q:104:LEU:HB2	1:Q:356:TRP:CZ3	2.30	0.66
1:R:362:TYR:O	1:R:366:GLY:HA2	1.94	0.66
1:S:153:LEU:HD23	1:S:299:MET:HE3	1.76	0.66
1:T:173:HIS:CE1	1:U:268:GLY:CA	2.77	0.66
1:U:38:PRO:CG	1:U:49:GLN:HE22	2.07	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:58:ALA:HB1	1:V:65:LEU:CD2	2.24	0.66
1:V:236:LEU:HD22	1:V:252:ASN:N	2.09	0.66
1:W:38:PRO:CG	1:W:49:GLN:HE22	2.07	0.66
1:W:153:LEU:HD23	1:W:299:MET:HE3	1.76	0.66
1:B:300:SER:CA	1:B:335:ARG:HG3	2.25	0.66
1:C:104:LEU:HB2	1:C:356:TRP:CZ3	2.30	0.66
1:K:34:ILE:HD13	1:K:67:LEU:HD22	1.77	0.66
1:K:370:VAL:O	1:K:370:VAL:HG13	1.95	0.66
1:L:153:LEU:HD23	1:L:299:MET:HE3	1.76	0.66
1:M:300:SER:CA	1:M:335:ARG:HG3	2.25	0.66
1:N:7:ALA:HB1	1:N:356:TRP:CH2	2.29	0.66
1:N:58:ALA:HB1	1:N:65:LEU:CD2	2.25	0.66
1:O:39:ARG:NE	1:O:66:THR:HB	2.09	0.66
1:R:143:TYR:O	1:R:143:TYR:CD1	2.48	0.66
1:R:153:LEU:HD23	1:R:299:MET:HE3	1.76	0.66
1:S:58:ALA:HB1	1:S:65:LEU:CD2	2.25	0.66
1:T:153:LEU:HD23	1:T:299:MET:HE3	1.76	0.66
1:U:362:TYR:CE1	1:U:367:PRO:HB3	2.28	0.66
1:V:34:ILE:HA	1:V:68:LYS:O	1.96	0.66
1:W:208:ILE:CD1	1:W:243:PRO:HD2	2.18	0.66
1:C:110:LEU:HD12	1:C:177:ARG:NH1	2.09	0.66
1:E:37:ARG:CG	1:E:38:PRO:HD2	2.20	0.66
1:E:38:PRO:CG	1:E:49:GLN:HE22	2.07	0.66
1:F:104:LEU:HB2	1:F:356:TRP:CZ3	2.30	0.66
1:Q:173:HIS:CE1	1:R:268:GLY:CA	2.77	0.66
1:S:370:VAL:O	1:S:370:VAL:HG13	1.95	0.66
1:V:143:TYR:O	1:V:143:TYR:CD1	2.48	0.66
1:D:34:ILE:HA	1:D:68:LYS:O	1.96	0.66
1:E:34:ILE:HA	1:E:68:LYS:O	1.96	0.66
1:F:236:LEU:HD22	1:F:252:ASN:N	2.09	0.66
1:H:236:LEU:HD22	1:H:252:ASN:N	2.09	0.66
1:J:143:TYR:O	1:J:143:TYR:CD1	2.48	0.66
1:J:153:LEU:HD23	1:J:299:MET:HE3	1.76	0.66
1:K:143:TYR:O	1:K:143:TYR:CD1	2.48	0.66
1:N:153:LEU:HD23	1:N:299:MET:HE3	1.76	0.66
1:O:37:ARG:CG	1:O:38:PRO:HD2	2.20	0.66
1:O:370:VAL:HG22	1:O:375:PHE:O	1.96	0.66
1:Q:34:ILE:HA	1:Q:68:LYS:O	1.96	0.66
1:S:104:LEU:HB2	1:S:356:TRP:CZ3	2.30	0.66
1:U:39:ARG:NE	1:U:66:THR:HB	2.09	0.66
1:U:110:LEU:HD12	1:U:177:ARG:NH1	2.09	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:58:ALA:HB1	1:W:65:LEU:CD2	2.25	0.66
1:A:104:LEU:HB2	1:A:356:TRP:CZ3	2.30	0.66
1:G:143:TYR:O	1:G:143:TYR:CD1	2.48	0.66
1:I:34:ILE:HD13	1:I:67:LEU:HD22	1.76	0.66
1:I:300:SER:CA	1:I:335:ARG:HG3	2.26	0.66
1:J:39:ARG:NE	1:J:66:THR:HB	2.09	0.66
1:K:34:ILE:HA	1:K:68:LYS:O	1.96	0.66
1:N:34:ILE:HD13	1:N:67:LEU:HD22	1.77	0.66
1:B:35:VAL:CG1	1:B:68:LYS:HB2	2.22	0.66
1:B:143:TYR:O	1:B:143:TYR:CD1	2.48	0.66
1:C:65:LEU:C	1:C:65:LEU:HD12	2.21	0.66
1:E:370:VAL:HG22	1:E:375:PHE:O	1.96	0.66
1:G:34:ILE:HA	1:G:68:LYS:O	1.96	0.66
1:G:38:PRO:CG	1:G:49:GLN:HE22	2.07	0.66
1:H:370:VAL:HG22	1:H:375:PHE:O	1.96	0.66
1:I:65:LEU:C	1:I:65:LEU:HD12	2.21	0.66
1:I:362:TYR:CE1	1:I:367:PRO:HB3	2.29	0.66
1:I:370:VAL:HG22	1:I:375:PHE:O	1.96	0.66
1:J:370:VAL:HG22	1:J:375:PHE:O	1.96	0.66
1:K:65:LEU:C	1:K:65:LEU:HD12	2.21	0.66
1:M:34:ILE:HA	1:M:68:LYS:O	1.96	0.66
1:O:153:LEU:HD23	1:O:299:MET:HE3	1.76	0.66
1:R:65:LEU:C	1:R:65:LEU:HD12	2.21	0.66
1:S:34:ILE:HA	1:S:68:LYS:O	1.96	0.66
1:U:35:VAL:CA	1:U:54:VAL:HG21	2.26	0.66
1:A:370:VAL:HG22	1:A:375:PHE:O	1.96	0.66
1:C:34:ILE:HA	1:C:68:LYS:O	1.96	0.66
1:C:153:LEU:HD23	1:C:299:MET:HE3	1.76	0.66
1:D:35:VAL:CG1	1:D:68:LYS:HB2	2.22	0.66
1:G:65:LEU:C	1:G:65:LEU:HD12	2.21	0.66
1:N:370:VAL:HG13	1:N:370:VAL:O	1.95	0.66
1:P:153:LEU:HD23	1:P:299:MET:HE3	1.76	0.66
1:P:370:VAL:HG22	1:P:375:PHE:O	1.96	0.66
1:S:35:VAL:CA	1:S:54:VAL:HG21	2.26	0.66
1:T:34:ILE:HA	1:T:68:LYS:O	1.96	0.66
1:U:104:LEU:HB2	1:U:356:TRP:CZ3	2.30	0.66
1:W:35:VAL:CA	1:W:54:VAL:HG21	2.26	0.66
1:W:143:TYR:O	1:W:143:TYR:CD1	2.48	0.66
1:A:58:ALA:HB1	1:A:65:LEU:CD2	2.25	0.66
1:B:153:LEU:HD23	1:B:299:MET:HE3	1.76	0.66
1:C:370:VAL:O	1:C:370:VAL:HG13	1.95	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:370:VAL:HG22	1:C:375:PHE:O	1.96	0.66
1:D:104:LEU:HB2	1:D:356:TRP:CZ3	2.30	0.66
1:G:35:VAL:CA	1:G:54:VAL:HG21	2.26	0.66
1:I:38:PRO:CG	1:I:49:GLN:HE22	2.07	0.66
1:L:208:ILE:CD1	1:L:243:PRO:HD2	2.18	0.66
1:L:362:TYR:CE1	1:L:367:PRO:HB3	2.28	0.66
1:N:362:TYR:CE1	1:N:367:PRO:HB3	2.28	0.66
1:P:65:LEU:HD12	1:P:65:LEU:C	2.21	0.66
1:Q:370:VAL:HG13	1:Q:370:VAL:O	1.95	0.66
1:S:362:TYR:CE1	1:S:367:PRO:HB3	2.28	0.66
1:T:300:SER:CA	1:T:335:ARG:HG3	2.26	0.66
1:W:370:VAL:HG22	1:W:375:PHE:O	1.96	0.66
1:E:35:VAL:CA	1:E:54:VAL:HG21	2.26	0.66
1:F:143:TYR:O	1:F:143:TYR:CD1	2.48	0.66
1:I:35:VAL:CA	1:I:54:VAL:HG21	2.26	0.66
1:L:58:ALA:HB1	1:L:65:LEU:CD2	2.25	0.66
1:M:370:VAL:HG22	1:M:375:PHE:O	1.96	0.66
1:O:300:SER:CA	1:O:335:ARG:HG3	2.25	0.66
1:P:34:ILE:HA	1:P:68:LYS:O	1.96	0.66
1:Q:35:VAL:CA	1:Q:54:VAL:HG21	2.26	0.66
1:Q:143:TYR:O	1:Q:143:TYR:CD1	2.48	0.66
1:Q:153:LEU:HD23	1:Q:299:MET:HE3	1.76	0.66
1:Q:300:SER:CA	1:Q:335:ARG:HG3	2.26	0.66
1:Q:324:THR:HG21	1:S:241:GLU:CD	2.21	0.66
1:Q:370:VAL:HG22	1:Q:375:PHE:O	1.96	0.66
1:R:34:ILE:HA	1:R:68:LYS:O	1.96	0.66
1:S:38:PRO:CG	1:S:49:GLN:HE22	2.07	0.66
1:T:370:VAL:HG13	1:T:370:VAL:O	1.95	0.66
1:U:58:ALA:HB1	1:U:65:LEU:CD2	2.25	0.66
1:U:143:TYR:O	1:U:143:TYR:CD1	2.48	0.66
1:U:324:THR:HG21	1:W:241:GLU:CD	2.21	0.66
1:A:34:ILE:HD13	1:A:67:LEU:HD22	1.77	0.66
1:B:65:LEU:C	1:B:65:LEU:HD12	2.21	0.66
1:D:324:THR:HG21	1:F:241:GLU:CD	2.21	0.66
1:G:370:VAL:HG13	1:G:370:VAL:O	1.95	0.66
1:H:324:THR:HG21	1:J:241:GLU:CD	2.21	0.66
1:J:324:THR:HG21	1:L:241:GLU:CD	2.21	0.66
1:M:65:LEU:HD12	1:M:65:LEU:C	2.21	0.66
1:O:65:LEU:C	1:O:65:LEU:HD12	2.21	0.66
1:O:143:TYR:O	1:O:143:TYR:CD1	2.48	0.66
1:P:370:VAL:O	1:P:370:VAL:HG13	1.95	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:300:SER:CA	1:S:335:ARG:HG3	2.25	0.66
1:T:370:VAL:HG22	1:T:375:PHE:O	1.96	0.66
1:A:370:VAL:HG13	1:A:370:VAL:O	1.95	0.65
1:B:324:THR:HG21	1:D:241:GLU:CD	2.21	0.65
1:B:370:VAL:HG22	1:B:375:PHE:O	1.96	0.65
1:C:34:ILE:HD13	1:C:67:LEU:HD22	1.77	0.65
1:I:370:VAL:HG13	1:I:370:VAL:O	1.95	0.65
1:J:58:ALA:HB1	1:J:65:LEU:CD2	2.25	0.65
1:L:39:ARG:NE	1:L:66:THR:HB	2.09	0.65
1:L:370:VAL:HG22	1:L:375:PHE:O	1.96	0.65
1:M:39:ARG:NE	1:M:66:THR:HB	2.09	0.65
1:Q:65:LEU:C	1:Q:65:LEU:HD12	2.21	0.65
1:R:370:VAL:HG22	1:R:375:PHE:O	1.96	0.65
1:S:324:THR:HG21	1:U:241:GLU:CD	2.21	0.65
1:T:34:ILE:HD13	1:T:67:LEU:HD22	1.77	0.65
1:V:370:VAL:O	1:V:370:VAL:HG13	1.95	0.65
1:V:370:VAL:HG22	1:V:375:PHE:O	1.96	0.65
1:W:104:LEU:HB2	1:W:356:TRP:CZ3	2.30	0.65
1:B:34:ILE:HA	1:B:68:LYS:O	1.96	0.65
1:D:35:VAL:CA	1:D:54:VAL:HG21	2.26	0.65
1:E:34:ILE:HD13	1:E:67:LEU:HD22	1.77	0.65
1:F:35:VAL:CG1	1:F:68:LYS:HB2	2.22	0.65
1:F:370:VAL:HG13	1:F:370:VAL:O	1.95	0.65
1:H:143:TYR:O	1:H:143:TYR:CD1	2.48	0.65
1:I:34:ILE:HA	1:I:68:LYS:O	1.96	0.65
1:J:34:ILE:HA	1:J:68:LYS:O	1.96	0.65
1:L:35:VAL:CA	1:L:54:VAL:HG21	2.26	0.65
1:R:34:ILE:HD13	1:R:67:LEU:HD22	1.77	0.65
1:T:65:LEU:C	1:T:65:LEU:HD12	2.21	0.65
1:V:35:VAL:CA	1:V:54:VAL:HG21	2.26	0.65
1:V:65:LEU:C	1:V:65:LEU:HD12	2.21	0.65
1:B:104:LEU:HB2	1:B:356:TRP:CZ3	2.30	0.65
1:D:153:LEU:HD23	1:D:299:MET:HE3	1.76	0.65
1:E:65:LEU:HD12	1:E:65:LEU:C	2.21	0.65
1:F:300:SER:CA	1:F:335:ARG:HG3	2.26	0.65
1:G:300:SER:CA	1:G:335:ARG:HG3	2.25	0.65
1:G:370:VAL:HG22	1:G:375:PHE:O	1.96	0.65
1:H:34:ILE:HA	1:H:68:LYS:O	1.96	0.65
1:H:153:LEU:HD23	1:H:299:MET:HE3	1.76	0.65
1:J:65:LEU:C	1:J:65:LEU:HD12	2.21	0.65
1:K:208:ILE:CD1	1:K:243:PRO:HD2	2.18	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:35:VAL:CA	1:O:54:VAL:HG21	2.26	0.65
1:V:34:ILE:HD13	1:V:67:LEU:HD22	1.77	0.65
1:V:300:SER:CA	1:V:335:ARG:HG3	2.25	0.65
1:A:34:ILE:HA	1:A:68:LYS:O	1.96	0.65
1:A:65:LEU:C	1:A:65:LEU:HD12	2.21	0.65
1:A:153:LEU:HD23	1:A:299:MET:HE3	1.76	0.65
1:D:143:TYR:O	1:D:143:TYR:CD1	2.48	0.65
1:E:369:ILE:CG2	1:E:370:VAL:N	2.60	0.65
1:F:370:VAL:HG22	1:F:375:PHE:O	1.96	0.65
1:G:34:ILE:HD13	1:G:67:LEU:HD22	1.77	0.65
1:K:35:VAL:CA	1:K:54:VAL:HG21	2.26	0.65
1:L:34:ILE:HA	1:L:68:LYS:O	1.96	0.65
1:L:65:LEU:C	1:L:65:LEU:HD12	2.21	0.65
1:N:370:VAL:HG22	1:N:375:PHE:O	1.96	0.65
1:P:34:ILE:HD13	1:P:67:LEU:HD22	1.77	0.65
1:S:143:TYR:O	1:S:143:TYR:CD1	2.48	0.65
1:T:290:ARG:CD	1:V:244:ASP:HB2	2.12	0.65
1:U:35:VAL:CG1	1:U:68:LYS:HB2	2.22	0.65
1:W:34:ILE:HA	1:W:68:LYS:O	1.96	0.65
1:W:39:ARG:NE	1:W:66:THR:HB	2.09	0.65
1:A:324:THR:HG21	1:C:241:GLU:CD	2.21	0.65
1:B:35:VAL:CA	1:B:54:VAL:HG21	2.26	0.65
1:C:35:VAL:CA	1:C:54:VAL:HG21	2.26	0.65
1:C:300:SER:CA	1:C:335:ARG:HG3	2.25	0.65
1:E:300:SER:CA	1:E:335:ARG:HG3	2.25	0.65
1:F:34:ILE:HA	1:F:68:LYS:O	1.96	0.65
1:P:35:VAL:CA	1:P:54:VAL:HG21	2.26	0.65
1:D:300:SER:CA	1:D:335:ARG:HG3	2.26	0.65
1:H:65:LEU:C	1:H:65:LEU:HD12	2.21	0.65
1:K:300:SER:CA	1:K:335:ARG:HG3	2.25	0.65
1:N:35:VAL:CA	1:N:54:VAL:HG21	2.26	0.65
1:N:369:ILE:CG2	1:N:370:VAL:N	2.60	0.65
1:T:369:ILE:CG2	1:T:370:VAL:N	2.60	0.65
1:E:362:TYR:CE1	1:E:367:PRO:HB3	2.29	0.65
1:H:300:SER:CA	1:H:335:ARG:HG3	2.26	0.65
1:K:324:THR:HG21	1:M:241:GLU:CD	2.21	0.65
1:L:369:ILE:CG2	1:L:370:VAL:N	2.60	0.65
1:N:65:LEU:C	1:N:65:LEU:HD12	2.21	0.65
1:S:143:TYR:CD1	1:S:143:TYR:C	2.75	0.65
1:T:324:THR:HG21	1:V:241:GLU:CD	2.21	0.65
1:U:370:VAL:HG22	1:U:375:PHE:O	1.96	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:369:ILE:CG2	1:C:370:VAL:N	2.60	0.65
1:G:369:ILE:CG2	1:G:370:VAL:N	2.60	0.65
1:J:35:VAL:CA	1:J:54:VAL:HG21	2.26	0.65
1:K:370:VAL:HG22	1:K:375:PHE:O	1.96	0.65
1:N:34:ILE:HA	1:N:68:LYS:O	1.96	0.65
1:O:143:TYR:CD1	1:O:143:TYR:C	2.75	0.65
1:P:324:THR:HG21	1:R:241:GLU:CD	2.21	0.65
1:R:324:THR:HG21	1:T:241:GLU:CD	2.21	0.65
1:V:369:ILE:CG2	1:V:370:VAL:N	2.60	0.65
1:D:58:ALA:O	1:D:61:LYS:N	2.30	0.65
1:F:153:LEU:HD23	1:F:299:MET:HE3	1.76	0.65
1:F:324:THR:HG21	1:H:241:GLU:CD	2.21	0.65
1:H:35:VAL:CA	1:H:54:VAL:HG21	2.26	0.65
1:H:143:TYR:CD1	1:H:143:TYR:C	2.75	0.65
1:L:324:THR:HG21	1:N:241:GLU:CD	2.21	0.65
1:M:324:THR:HG21	1:O:241:GLU:CD	2.21	0.65
1:O:58:ALA:O	1:O:61:LYS:N	2.30	0.65
1:R:35:VAL:CA	1:R:54:VAL:HG21	2.26	0.65
1:R:300:SER:CA	1:R:335:ARG:HG3	2.26	0.65
1:R:370:VAL:HG13	1:R:370:VAL:O	1.95	0.65
1:S:65:LEU:C	1:S:65:LEU:HD12	2.21	0.65
1:T:35:VAL:CA	1:T:54:VAL:HG21	2.26	0.65
1:W:37:ARG:CG	1:W:38:PRO:HD2	2.20	0.65
1:W:65:LEU:C	1:W:65:LEU:HD12	2.21	0.65
1:B:39:ARG:NE	1:B:66:THR:HB	2.09	0.65
1:C:58:ALA:O	1:C:61:LYS:N	2.30	0.65
1:E:58:ALA:O	1:E:61:LYS:N	2.30	0.65
1:E:370:VAL:O	1:E:370:VAL:HG13	1.95	0.65
1:G:362:TYR:CE1	1:G:367:PRO:HB3	2.29	0.65
1:N:58:ALA:O	1:N:61:LYS:N	2.30	0.65
1:N:324:THR:HG21	1:P:241:GLU:CD	2.21	0.65
1:U:34:ILE:HA	1:U:68:LYS:O	1.96	0.65
1:D:65:LEU:C	1:D:65:LEU:HD12	2.21	0.64
1:G:324:THR:HG21	1:I:241:GLU:CD	2.21	0.64
1:M:35:VAL:CA	1:M:54:VAL:HG21	2.26	0.64
1:N:300:SER:CA	1:N:335:ARG:HG3	2.26	0.64
1:P:369:ILE:CG2	1:P:370:VAL:N	2.60	0.64
1:S:370:VAL:HG22	1:S:375:PHE:O	1.96	0.64
1:T:58:ALA:O	1:T:61:LYS:N	2.30	0.64
1:W:300:SER:CA	1:W:335:ARG:HG3	2.25	0.64
1:W:369:ILE:CG2	1:W:370:VAL:N	2.60	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:35:VAL:CA	1:A:54:VAL:HG21	2.26	0.64
1:C:324:THR:HG21	1:E:241:GLU:CD	2.21	0.64
1:F:58:ALA:O	1:F:61:LYS:N	2.30	0.64
1:I:58:ALA:O	1:I:61:LYS:N	2.30	0.64
1:I:324:THR:HG21	1:K:241:GLU:CD	2.21	0.64
1:P:300:SER:CA	1:P:335:ARG:HG3	2.25	0.64
1:A:39:ARG:NE	1:A:66:THR:HB	2.09	0.64
1:F:7:ALA:HB1	1:F:356:TRP:CZ2	2.33	0.64
1:H:35:VAL:CG1	1:H:68:LYS:HB2	2.22	0.64
1:K:39:ARG:NE	1:K:66:THR:HB	2.09	0.64
1:N:39:ARG:NE	1:N:66:THR:HB	2.09	0.64
1:O:324:THR:HG21	1:Q:241:GLU:CD	2.21	0.64
1:P:58:ALA:O	1:P:61:LYS:N	2.30	0.64
1:Q:58:ALA:O	1:Q:61:LYS:N	2.30	0.64
1:V:58:ALA:O	1:V:61:LYS:N	2.30	0.64
1:A:7:ALA:HB1	1:A:356:TRP:CZ2	2.33	0.64
1:D:7:ALA:HB1	1:D:356:TRP:CZ2	2.33	0.64
1:F:35:VAL:CA	1:F:54:VAL:HG21	2.26	0.64
1:F:65:LEU:C	1:F:65:LEU:HD12	2.21	0.64
1:K:58:ALA:O	1:K:61:LYS:N	2.30	0.64
1:K:143:TYR:CD1	1:K:143:TYR:C	2.75	0.64
1:K:369:ILE:CG2	1:K:370:VAL:N	2.60	0.64
1:R:369:ILE:CG2	1:R:370:VAL:N	2.60	0.64
1:V:39:ARG:NE	1:V:66:THR:HB	2.09	0.64
1:W:35:VAL:CG1	1:W:68:LYS:HB2	2.22	0.64
1:C:7:ALA:HB1	1:C:356:TRP:CZ2	2.33	0.64
1:D:370:VAL:HG22	1:D:375:PHE:O	1.96	0.64
1:E:7:ALA:HB1	1:E:356:TRP:CZ2	2.33	0.64
1:H:7:ALA:HB1	1:H:356:TRP:CZ2	2.33	0.64
1:K:7:ALA:HB1	1:K:356:TRP:CZ2	2.33	0.64
1:K:34:ILE:HB	1:K:54:VAL:HG13	1.80	0.64
1:L:7:ALA:HB1	1:L:356:TRP:CZ2	2.33	0.64
1:O:369:ILE:CG2	1:O:370:VAL:N	2.60	0.64
1:P:7:ALA:HB1	1:P:356:TRP:CZ2	2.33	0.64
1:Q:7:ALA:HB1	1:Q:356:TRP:CZ2	2.33	0.64
1:R:7:ALA:HB1	1:R:356:TRP:CZ2	2.33	0.64
1:S:7:ALA:HB1	1:S:356:TRP:CZ2	2.33	0.64
1:S:37:ARG:CG	1:S:38:PRO:HD2	2.20	0.64
1:U:369:ILE:CG2	1:U:370:VAL:N	2.60	0.64
1:W:143:TYR:CD1	1:W:143:TYR:C	2.75	0.64
1:A:369:ILE:CG2	1:A:370:VAL:N	2.60	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:7:ALA:HB1	1:B:356:TRP:CZ2	2.33	0.64
1:C:290:ARG:CD	1:E:244:ASP:HB2	2.12	0.64
1:D:39:ARG:NE	1:D:66:THR:HB	2.09	0.64
1:H:37:ARG:CG	1:H:38:PRO:HD2	2.20	0.64
1:I:7:ALA:HB1	1:I:356:TRP:CZ2	2.33	0.64
1:J:7:ALA:HB1	1:J:356:TRP:CZ2	2.33	0.64
1:L:300:SER:CA	1:L:335:ARG:HG3	2.26	0.64
1:M:369:ILE:CG2	1:M:370:VAL:N	2.60	0.64
1:N:7:ALA:HB1	1:N:356:TRP:CZ2	2.33	0.64
1:N:37:ARG:CG	1:N:38:PRO:HD2	2.20	0.64
1:O:7:ALA:HB1	1:O:356:TRP:CZ2	2.33	0.64
1:V:208:ILE:CD1	1:V:243:PRO:HD2	2.18	0.64
1:A:300:SER:CA	1:A:335:ARG:HG3	2.26	0.64
1:C:39:ARG:NE	1:C:66:THR:HB	2.09	0.64
1:F:369:ILE:CG2	1:F:370:VAL:N	2.60	0.64
1:J:369:ILE:CG2	1:J:370:VAL:N	2.60	0.64
1:Q:369:ILE:CG2	1:Q:370:VAL:N	2.60	0.64
1:U:7:ALA:HB1	1:U:356:TRP:CZ2	2.33	0.64
1:U:300:SER:CA	1:U:335:ARG:HG3	2.26	0.64
1:B:58:ALA:O	1:B:61:LYS:N	2.30	0.64
1:E:324:THR:HG21	1:G:241:GLU:CD	2.21	0.64
1:H:369:ILE:CG2	1:H:370:VAL:N	2.60	0.64
1:M:7:ALA:HB1	1:M:356:TRP:CZ2	2.33	0.64
1:U:65:LEU:C	1:U:65:LEU:HD12	2.21	0.64
1:W:7:ALA:HB1	1:W:356:TRP:CZ2	2.33	0.64
1:D:369:ILE:CG2	1:D:370:VAL:N	2.60	0.64
1:G:7:ALA:HB1	1:G:356:TRP:CZ2	2.33	0.64
1:H:58:ALA:O	1:H:61:LYS:N	2.30	0.64
1:J:300:SER:CA	1:J:335:ARG:HG3	2.26	0.64
1:M:58:ALA:O	1:M:61:LYS:N	2.30	0.64
1:B:369:ILE:CG2	1:B:370:VAL:N	2.60	0.63
1:E:39:ARG:NE	1:E:66:THR:HB	2.09	0.63
1:I:298:VAL:HG12	1:I:335:ARG:HH11	1.63	0.63
1:I:334:GLU:OE2	1:I:334:GLU:HA	1.98	0.63
1:I:369:ILE:CG2	1:I:370:VAL:N	2.60	0.63
1:J:298:VAL:HG12	1:J:335:ARG:HH11	1.64	0.63
1:N:299:MET:HE2	1:N:304:THR:HB	1.81	0.63
1:T:143:TYR:CD1	1:T:143:TYR:C	2.75	0.63
1:V:35:VAL:CG1	1:V:68:LYS:HB2	2.22	0.63
1:E:299:MET:HE2	1:E:304:THR:HB	1.81	0.63
1:J:334:GLU:HA	1:J:334:GLU:OE2	1.98	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:334:GLU:HA	1:L:334:GLU:OE2	1.98	0.63
1:S:334:GLU:HA	1:S:334:GLU:OE2	1.98	0.63
1:T:299:MET:HE2	1:T:304:THR:HB	1.81	0.63
1:V:143:TYR:CD1	1:V:143:TYR:C	2.75	0.63
1:V:334:GLU:HA	1:V:334:GLU:OE2	1.98	0.63
1:C:299:MET:HE2	1:C:304:THR:HB	1.81	0.63
1:E:143:TYR:CD1	1:E:143:TYR:C	2.75	0.63
1:E:147:ARG:HH21	1:E:147:ARG:CG	2.12	0.63
1:F:334:GLU:HA	1:F:334:GLU:OE2	1.98	0.63
1:G:58:ALA:O	1:G:61:LYS:N	2.30	0.63
1:G:334:GLU:HA	1:G:334:GLU:OE2	1.98	0.63
1:L:143:TYR:CD1	1:L:143:TYR:C	2.75	0.63
1:P:299:MET:HE2	1:P:304:THR:HB	1.81	0.63
1:T:334:GLU:OE2	1:T:334:GLU:HA	1.98	0.63
1:A:298:VAL:HG12	1:A:335:ARG:HH11	1.64	0.63
1:A:299:MET:HE2	1:A:304:THR:HB	1.81	0.63
1:C:147:ARG:HH21	1:C:147:ARG:CG	2.12	0.63
1:G:39:ARG:NE	1:G:66:THR:HB	2.09	0.63
1:G:147:ARG:HH21	1:G:147:ARG:CG	2.12	0.63
1:H:299:MET:HE2	1:H:304:THR:HB	1.80	0.63
1:I:299:MET:HE2	1:I:304:THR:HB	1.81	0.63
1:J:299:MET:HE2	1:J:304:THR:HB	1.81	0.63
1:L:37:ARG:CG	1:L:38:PRO:HD2	2.20	0.63
1:P:298:VAL:HG12	1:P:335:ARG:HH11	1.64	0.63
1:R:299:MET:HE2	1:R:304:THR:HB	1.81	0.63
1:S:369:ILE:CG2	1:S:370:VAL:N	2.60	0.63
1:T:39:ARG:NE	1:T:66:THR:HB	2.09	0.63
1:V:298:VAL:HG12	1:V:335:ARG:HH11	1.64	0.63
1:W:147:ARG:HH21	1:W:147:ARG:CG	2.12	0.63
1:W:299:MET:HE2	1:W:304:THR:HB	1.81	0.63
1:C:298:VAL:HG12	1:C:335:ARG:HH11	1.64	0.63
1:D:34:ILE:HB	1:D:54:VAL:HG13	1.79	0.63
1:F:39:ARG:NE	1:F:66:THR:HB	2.09	0.63
1:G:143:TYR:CD1	1:G:143:TYR:C	2.75	0.63
1:G:298:VAL:HG12	1:G:335:ARG:HH11	1.64	0.63
1:J:35:VAL:CG1	1:J:68:LYS:HB2	2.22	0.63
1:J:37:ARG:O	1:J:66:THR:HG22	1.99	0.63
1:L:299:MET:HE2	1:L:304:THR:HB	1.81	0.63
1:O:37:ARG:HB2	1:O:51:ASP:O	1.99	0.63
1:Q:147:ARG:HH21	1:Q:147:ARG:CG	2.12	0.63
1:T:7:ALA:HB1	1:T:356:TRP:CZ2	2.33	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:37:ARG:HB2	1:T:51:ASP:O	1.99	0.63
1:W:334:GLU:HA	1:W:334:GLU:OE2	1.98	0.63
1:B:298:VAL:HG12	1:B:335:ARG:HH11	1.64	0.63
1:D:334:GLU:HA	1:D:334:GLU:OE2	1.98	0.63
1:E:169:TYR:CE2	1:G:40:HIS:CB	2.82	0.63
1:G:37:ARG:O	1:G:66:THR:HG22	1.99	0.63
1:I:37:ARG:HB2	1:I:51:ASP:O	1.99	0.63
1:K:334:GLU:HA	1:K:334:GLU:OE2	1.98	0.63
1:M:34:ILE:HB	1:M:54:VAL:HG13	1.80	0.63
1:O:147:ARG:HH21	1:O:147:ARG:CG	2.12	0.63
1:O:298:VAL:HG12	1:O:335:ARG:HH11	1.64	0.63
1:P:39:ARG:NE	1:P:66:THR:HB	2.09	0.63
1:Q:298:VAL:HG12	1:Q:335:ARG:HH11	1.64	0.63
1:U:147:ARG:HH21	1:U:147:ARG:CG	2.12	0.63
1:B:169:TYR:CE2	1:D:40:HIS:CB	2.82	0.63
1:C:169:TYR:CE2	1:E:40:HIS:CB	2.82	0.63
1:D:37:ARG:O	1:D:66:THR:HG22	1.99	0.63
1:E:37:ARG:O	1:E:66:THR:HG22	1.99	0.63
1:G:299:MET:HE2	1:G:304:THR:HB	1.81	0.63
1:H:37:ARG:O	1:H:66:THR:HG22	1.99	0.63
1:K:37:ARG:HB2	1:K:51:ASP:O	1.99	0.63
1:L:37:ARG:O	1:L:66:THR:HG22	1.99	0.63
1:L:298:VAL:HG12	1:L:335:ARG:HH11	1.64	0.63
1:M:169:TYR:CE2	1:O:40:HIS:CB	2.82	0.63
1:O:35:VAL:CG1	1:O:68:LYS:HB2	2.22	0.63
1:O:37:ARG:O	1:O:66:THR:HG22	1.99	0.63
1:O:43:VAL:O	1:O:44:MET:HE2	1.99	0.63
1:R:58:ALA:O	1:R:61:LYS:N	2.30	0.63
1:R:298:VAL:HG12	1:R:335:ARG:HH11	1.64	0.63
1:S:58:ALA:O	1:S:61:LYS:N	2.30	0.63
1:S:147:ARG:HH21	1:S:147:ARG:CG	2.12	0.63
1:U:299:MET:HE2	1:U:304:THR:HB	1.81	0.63
1:V:7:ALA:HB1	1:V:356:TRP:CZ2	2.33	0.63
1:V:147:ARG:HH21	1:V:147:ARG:CG	2.12	0.63
1:A:37:ARG:O	1:A:66:THR:HG22	1.99	0.63
1:C:37:ARG:O	1:C:66:THR:HG22	1.99	0.63
1:E:43:VAL:O	1:E:44:MET:HE2	1.99	0.63
1:E:334:GLU:HA	1:E:334:GLU:OE2	1.98	0.63
1:F:37:ARG:O	1:F:66:THR:HG22	1.99	0.63
1:F:143:TYR:CD1	1:F:143:TYR:C	2.75	0.63
1:G:169:TYR:CE2	1:I:40:HIS:CB	2.82	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:298:VAL:HG12	1:H:335:ARG:HH11	1.64	0.63
1:I:290:ARG:CD	1:K:244:ASP:HB2	2.12	0.63
1:K:169:TYR:CE2	1:M:40:HIS:CB	2.82	0.63
1:M:35:VAL:CG1	1:M:68:LYS:HB2	2.22	0.63
1:M:37:ARG:O	1:M:66:THR:HG22	1.99	0.63
1:N:147:ARG:HH21	1:N:147:ARG:CG	2.12	0.63
1:O:169:TYR:CE2	1:Q:40:HIS:CB	2.82	0.63
1:Q:334:GLU:OE2	1:Q:334:GLU:HA	1.98	0.63
1:S:43:VAL:O	1:S:44:MET:HE2	1.99	0.63
1:S:299:MET:HE2	1:S:304:THR:HB	1.81	0.63
1:T:37:ARG:CG	1:T:38:PRO:HD2	2.20	0.63
1:T:43:VAL:O	1:T:44:MET:HE2	1.99	0.63
1:V:34:ILE:HB	1:V:54:VAL:HG13	1.80	0.63
1:W:298:VAL:HG12	1:W:335:ARG:HH11	1.64	0.63
1:D:169:TYR:CE2	1:F:40:HIS:CB	2.82	0.63
1:F:299:MET:HE2	1:F:304:THR:HB	1.81	0.63
1:I:39:ARG:NE	1:I:66:THR:HB	2.09	0.63
1:I:169:TYR:CE2	1:K:40:HIS:CB	2.82	0.63
1:J:147:ARG:HH21	1:J:147:ARG:CG	2.12	0.63
1:L:43:VAL:O	1:L:44:MET:HE2	1.99	0.63
1:L:147:ARG:HH21	1:L:147:ARG:CG	2.12	0.63
1:N:43:VAL:O	1:N:44:MET:HE2	1.99	0.63
1:P:147:ARG:HH21	1:P:147:ARG:CG	2.12	0.63
1:Q:35:VAL:CG1	1:Q:68:LYS:HB2	2.22	0.63
1:Q:37:ARG:HB2	1:Q:51:ASP:O	1.99	0.63
1:Q:37:ARG:O	1:Q:66:THR:HG22	1.99	0.63
1:Q:169:TYR:CE2	1:S:40:HIS:CB	2.82	0.63
1:R:143:TYR:CD1	1:R:143:TYR:C	2.75	0.63
1:S:37:ARG:O	1:S:66:THR:HG22	1.99	0.63
1:U:37:ARG:O	1:U:66:THR:HG22	1.99	0.63
1:V:299:MET:HE2	1:V:304:THR:HB	1.81	0.63
1:W:37:ARG:HB2	1:W:51:ASP:O	1.99	0.63
1:A:37:ARG:HB2	1:A:51:ASP:O	1.99	0.62
1:A:169:TYR:CE2	1:C:40:HIS:CB	2.82	0.62
1:B:143:TYR:CD1	1:B:143:TYR:C	2.75	0.62
1:B:362:TYR:HA	1:B:366:GLY:O	1.99	0.62
1:C:143:TYR:CD1	1:C:143:TYR:C	2.75	0.62
1:I:147:ARG:HH21	1:I:147:ARG:CG	2.12	0.62
1:Q:143:TYR:CD1	1:Q:143:TYR:C	2.75	0.62
1:T:35:VAL:CG1	1:T:68:LYS:HB2	2.22	0.62
1:U:334:GLU:HA	1:U:334:GLU:OE2	1.98	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:147:ARG:HH21	1:A:147:ARG:CG	2.12	0.62
1:D:362:TYR:HA	1:D:366:GLY:O	2.00	0.62
1:E:37:ARG:HB2	1:E:51:ASP:O	1.99	0.62
1:E:298:VAL:HG12	1:E:335:ARG:HH11	1.64	0.62
1:F:362:TYR:HA	1:F:366:GLY:O	2.00	0.62
1:I:37:ARG:O	1:I:66:THR:HG22	1.99	0.62
1:K:35:VAL:CG1	1:K:68:LYS:HB2	2.22	0.62
1:K:43:VAL:O	1:K:44:MET:HE2	1.99	0.62
1:L:169:TYR:CE2	1:N:40:HIS:CB	2.82	0.62
1:M:147:ARG:HH21	1:M:147:ARG:CG	2.12	0.62
1:M:334:GLU:HA	1:M:334:GLU:OE2	1.98	0.62
1:N:37:ARG:O	1:N:66:THR:HG22	1.99	0.62
1:N:169:TYR:CE2	1:P:40:HIS:CB	2.82	0.62
1:R:334:GLU:HA	1:R:334:GLU:OE2	1.98	0.62
1:B:147:ARG:HH21	1:B:147:ARG:CG	2.12	0.62
1:B:299:MET:HE2	1:B:304:THR:HB	1.81	0.62
1:D:43:VAL:O	1:D:44:MET:HE2	1.99	0.62
1:D:299:MET:HE2	1:D:304:THR:HB	1.81	0.62
1:F:37:ARG:HB2	1:F:51:ASP:O	1.99	0.62
1:F:43:VAL:O	1:F:44:MET:HE2	1.99	0.62
1:I:43:VAL:O	1:I:44:MET:HE2	1.99	0.62
1:K:299:MET:HE2	1:K:304:THR:HB	1.81	0.62
1:M:299:MET:HE2	1:M:304:THR:HB	1.81	0.62
1:P:169:TYR:CE2	1:R:40:HIS:CB	2.82	0.62
1:Q:299:MET:HE2	1:Q:304:THR:HB	1.81	0.62
1:S:169:TYR:CE2	1:U:40:HIS:CB	2.82	0.62
1:S:298:VAL:HG12	1:S:335:ARG:HH11	1.64	0.62
1:U:37:ARG:HB2	1:U:51:ASP:O	1.99	0.62
1:V:37:ARG:HB2	1:V:51:ASP:O	1.99	0.62
1:V:362:TYR:HA	1:V:366:GLY:O	2.00	0.62
1:A:143:TYR:CD1	1:A:143:TYR:C	2.75	0.62
1:C:37:ARG:CG	1:C:38:PRO:HD2	2.20	0.62
1:C:37:ARG:HB2	1:C:51:ASP:O	1.99	0.62
1:H:334:GLU:HA	1:H:334:GLU:OE2	1.98	0.62
1:J:43:VAL:O	1:J:44:MET:HE2	1.99	0.62
1:J:169:TYR:CE2	1:L:40:HIS:CB	2.82	0.62
1:M:37:ARG:HB2	1:M:51:ASP:O	1.99	0.62
1:M:362:TYR:HA	1:M:366:GLY:O	2.00	0.62
1:R:39:ARG:NE	1:R:66:THR:HB	2.09	0.62
1:T:147:ARG:HH21	1:T:147:ARG:CG	2.12	0.62
1:U:143:TYR:CD1	1:U:143:TYR:C	2.75	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:58:ALA:O	1:A:61:LYS:N	2.30	0.62
1:D:37:ARG:HB2	1:D:51:ASP:O	1.99	0.62
1:D:133:TYR:HB2	1:D:356:TRP:CE3	2.35	0.62
1:F:133:TYR:HB2	1:F:356:TRP:CE3	2.35	0.62
1:F:298:VAL:HG12	1:F:335:ARG:HH11	1.64	0.62
1:G:362:TYR:HA	1:G:366:GLY:O	1.99	0.62
1:J:58:ALA:O	1:J:61:LYS:N	2.30	0.62
1:P:334:GLU:HA	1:P:334:GLU:OE2	1.98	0.62
1:Q:43:VAL:O	1:Q:44:MET:HE2	1.99	0.62
1:Q:362:TYR:HA	1:Q:366:GLY:O	2.00	0.62
1:T:169:TYR:CE2	1:V:40:HIS:CB	2.82	0.62
1:T:298:VAL:HG12	1:T:335:ARG:HH11	1.64	0.62
1:U:43:VAL:O	1:U:44:MET:HE2	1.99	0.62
1:V:37:ARG:O	1:V:66:THR:HG22	1.99	0.62
1:V:133:TYR:HB2	1:V:356:TRP:CE3	2.35	0.62
1:A:40:HIS:O	1:A:41:GLN:C	2.43	0.62
1:B:43:VAL:O	1:B:44:MET:HE2	1.99	0.62
1:B:133:TYR:HB2	1:B:356:TRP:CE3	2.35	0.62
1:C:334:GLU:HA	1:C:334:GLU:OE2	1.98	0.62
1:D:298:VAL:HG12	1:D:335:ARG:HH11	1.64	0.62
1:E:40:HIS:O	1:E:41:GLN:C	2.43	0.62
1:E:362:TYR:HA	1:E:366:GLY:O	2.00	0.62
1:H:362:TYR:HA	1:H:366:GLY:O	2.00	0.62
1:K:147:ARG:HH21	1:K:147:ARG:CG	2.12	0.62
1:K:298:VAL:HG12	1:K:335:ARG:HH11	1.64	0.62
1:L:40:HIS:O	1:L:41:GLN:C	2.43	0.62
1:M:298:VAL:HG12	1:M:335:ARG:HH11	1.64	0.62
1:N:298:VAL:HG12	1:N:335:ARG:HH11	1.64	0.62
1:N:334:GLU:HA	1:N:334:GLU:OE2	1.98	0.62
1:P:40:HIS:O	1:P:41:GLN:C	2.43	0.62
1:P:104:LEU:HB2	1:P:356:TRP:HH2	1.63	0.62
1:P:143:TYR:CD1	1:P:143:TYR:C	2.75	0.62
1:Q:104:LEU:HB2	1:Q:356:TRP:HH2	1.63	0.62
1:S:35:VAL:CG1	1:S:68:LYS:HB2	2.22	0.62
1:T:362:TYR:HA	1:T:366:GLY:O	1.99	0.62
1:U:169:TYR:CE2	1:W:40:HIS:CB	2.82	0.62
1:W:133:TYR:HB2	1:W:356:TRP:CE3	2.35	0.62
1:A:35:VAL:CG1	1:A:68:LYS:HB2	2.22	0.62
1:B:37:ARG:HB2	1:B:51:ASP:O	1.99	0.62
1:H:169:TYR:CE2	1:J:40:HIS:CB	2.82	0.62
1:I:35:VAL:CG1	1:I:68:LYS:HB2	2.22	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:34:ILE:HB	1:J:54:VAL:HG13	1.79	0.62
1:M:44:MET:HG3	1:M:45:VAL:N	2.15	0.62
1:M:143:TYR:CD1	1:M:143:TYR:C	2.75	0.62
1:N:104:LEU:HB2	1:N:356:TRP:HH2	1.63	0.62
1:P:37:ARG:O	1:P:66:THR:HG22	1.99	0.62
1:R:43:VAL:O	1:R:44:MET:HE2	1.99	0.62
1:R:44:MET:HG3	1:R:45:VAL:N	2.15	0.62
1:R:147:ARG:HH21	1:R:147:ARG:CG	2.12	0.62
1:T:40:HIS:O	1:T:41:GLN:C	2.43	0.62
1:T:133:TYR:HB2	1:T:356:TRP:CE3	2.35	0.62
1:W:37:ARG:O	1:W:66:THR:HG22	1.99	0.62
1:W:40:HIS:O	1:W:41:GLN:C	2.43	0.62
1:A:44:MET:HG3	1:A:45:VAL:N	2.15	0.62
1:C:362:TYR:HA	1:C:366:GLY:O	2.00	0.62
1:F:169:TYR:CE2	1:H:40:HIS:CB	2.82	0.62
1:H:133:TYR:HB2	1:H:356:TRP:CE3	2.35	0.62
1:I:362:TYR:HA	1:I:366:GLY:O	2.00	0.62
1:O:299:MET:HE2	1:O:304:THR:HB	1.81	0.62
1:R:169:TYR:CE2	1:T:40:HIS:CB	2.82	0.62
1:S:37:ARG:HB2	1:S:51:ASP:O	1.99	0.62
1:V:44:MET:HG3	1:V:45:VAL:N	2.15	0.62
1:B:37:ARG:CG	1:B:38:PRO:HD2	2.20	0.62
1:B:37:ARG:O	1:B:66:THR:HG22	1.99	0.62
1:C:133:TYR:HB2	1:C:356:TRP:CE3	2.35	0.62
1:E:44:MET:HG3	1:E:45:VAL:N	2.15	0.62
1:E:133:TYR:HB2	1:E:356:TRP:CE3	2.35	0.62
1:H:40:HIS:O	1:H:41:GLN:C	2.43	0.62
1:I:44:MET:HG3	1:I:45:VAL:N	2.15	0.62
1:J:143:TYR:CD1	1:J:143:TYR:C	2.75	0.62
1:M:43:VAL:O	1:M:44:MET:HE2	1.99	0.62
1:N:44:MET:HG3	1:N:45:VAL:N	2.15	0.62
1:P:34:ILE:HB	1:P:54:VAL:HG13	1.79	0.62
1:P:43:VAL:O	1:P:44:MET:HE2	1.99	0.62
1:R:37:ARG:HB2	1:R:51:ASP:O	1.99	0.62
1:A:43:VAL:O	1:A:44:MET:HE2	1.99	0.62
1:A:133:TYR:HB2	1:A:356:TRP:CE3	2.35	0.62
1:A:334:GLU:OE2	1:A:334:GLU:HA	1.98	0.62
1:B:334:GLU:HA	1:B:334:GLU:OE2	1.98	0.62
1:F:34:ILE:HB	1:F:54:VAL:HG13	1.79	0.62
1:G:43:VAL:O	1:G:44:MET:HE2	1.99	0.62
1:G:133:TYR:HB2	1:G:356:TRP:CE3	2.35	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:39:ARG:NE	1:H:66:THR:HB	2.09	0.62
1:I:133:TYR:OH	1:I:375:PHE:HB2	2.00	0.62
1:J:362:TYR:HA	1:J:366:GLY:O	2.00	0.62
1:K:362:TYR:HA	1:K:366:GLY:O	2.00	0.62
1:L:35:VAL:CG1	1:L:68:LYS:HB2	2.22	0.62
1:O:334:GLU:HA	1:O:334:GLU:OE2	1.98	0.62
1:P:133:TYR:HB2	1:P:356:TRP:CE3	2.35	0.62
1:R:35:VAL:CG1	1:R:68:LYS:HB2	2.22	0.62
1:U:133:TYR:HB2	1:U:356:TRP:CE3	2.35	0.62
1:U:362:TYR:HA	1:U:366:GLY:O	2.00	0.62
1:W:362:TYR:HA	1:W:366:GLY:O	2.00	0.62
1:A:104:LEU:HB2	1:A:356:TRP:HH2	1.63	0.61
1:A:362:TYR:HA	1:A:366:GLY:O	2.00	0.61
1:D:44:MET:HG3	1:D:45:VAL:N	2.15	0.61
1:G:35:VAL:CG1	1:G:68:LYS:HB2	2.22	0.61
1:H:34:ILE:HB	1:H:54:VAL:HG13	1.80	0.61
1:I:40:HIS:O	1:I:41:GLN:C	2.43	0.61
1:K:37:ARG:O	1:K:66:THR:HG22	1.99	0.61
1:N:133:TYR:HB2	1:N:356:TRP:CE3	2.35	0.61
1:O:133:TYR:OH	1:O:375:PHE:HB2	2.00	0.61
1:R:37:ARG:O	1:R:66:THR:HG22	1.99	0.61
1:R:133:TYR:HB2	1:R:356:TRP:CE3	2.35	0.61
1:S:133:TYR:HB2	1:S:356:TRP:CE3	2.35	0.61
1:W:44:MET:HG3	1:W:45:VAL:N	2.15	0.61
1:C:43:VAL:O	1:C:44:MET:HE2	1.99	0.61
1:J:44:MET:HG3	1:J:45:VAL:N	2.15	0.61
1:K:157:ASP:HB2	2:K:401:ADP:H5'1	1.83	0.61
1:L:133:TYR:HB2	1:L:356:TRP:CE3	2.35	0.61
1:N:37:ARG:HB2	1:N:51:ASP:O	1.99	0.61
1:Q:44:MET:HG3	1:Q:45:VAL:N	2.15	0.61
1:S:362:TYR:HA	1:S:366:GLY:O	1.99	0.61
1:T:37:ARG:O	1:T:66:THR:HG22	1.99	0.61
1:W:58:ALA:O	1:W:61:LYS:N	2.30	0.61
1:W:104:LEU:HB2	1:W:356:TRP:HH2	1.63	0.61
1:C:35:VAL:CG1	1:C:68:LYS:HB2	2.22	0.61
1:D:147:ARG:HH21	1:D:147:ARG:CG	2.12	0.61
1:H:37:ARG:HB2	1:H:51:ASP:O	1.99	0.61
1:H:43:VAL:O	1:H:44:MET:HE2	1.99	0.61
1:I:133:TYR:HB2	1:I:356:TRP:CE3	2.35	0.61
1:J:133:TYR:HB2	1:J:356:TRP:CE3	2.35	0.61
1:N:35:VAL:CG1	1:N:68:LYS:HB2	2.22	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:362:TYR:HA	1:R:366:GLY:O	2.00	0.61
1:S:40:HIS:O	1:S:41:GLN:C	2.43	0.61
1:T:133:TYR:OH	1:T:375:PHE:HB2	2.00	0.61
1:V:157:ASP:HB2	2:V:401:ADP:H5'1	1.83	0.61
1:W:43:VAL:O	1:W:44:MET:HE2	1.99	0.61
1:D:157:ASP:HB2	2:D:401:ADP:H5'1	1.82	0.61
1:I:157:ASP:HB2	2:I:401:ADP:H5'1	1.83	0.61
1:O:34:ILE:HB	1:O:54:VAL:HG13	1.80	0.61
1:P:35:VAL:CG1	1:P:68:LYS:HB2	2.22	0.61
1:P:37:ARG:HB2	1:P:51:ASP:O	1.99	0.61
1:Q:133:TYR:HB2	1:Q:356:TRP:CE3	2.35	0.61
1:R:104:LEU:HB2	1:R:356:TRP:HH2	1.63	0.61
1:D:133:TYR:OH	1:D:375:PHE:HB2	2.00	0.61
1:E:35:VAL:CG1	1:E:68:LYS:HB2	2.22	0.61
1:E:157:ASP:HB2	2:E:401:ADP:H5'1	1.83	0.61
1:I:143:TYR:CD1	1:I:143:TYR:C	2.75	0.61
1:J:37:ARG:HB2	1:J:51:ASP:O	1.99	0.61
1:L:37:ARG:HB2	1:L:51:ASP:O	1.99	0.61
1:O:157:ASP:HB2	2:O:401:ADP:H5'1	1.83	0.61
1:Q:157:ASP:HB2	2:Q:401:ADP:H5'1	1.83	0.61
1:R:157:ASP:HB2	2:R:401:ADP:H5'1	1.83	0.61
1:V:43:VAL:O	1:V:44:MET:HE2	1.99	0.61
1:B:157:ASP:HB2	2:B:401:ADP:H5'1	1.83	0.61
1:G:37:ARG:HB2	1:G:51:ASP:O	1.99	0.61
1:H:147:ARG:HH21	1:H:147:ARG:CG	2.12	0.61
1:J:40:HIS:O	1:J:41:GLN:C	2.43	0.61
1:K:133:TYR:OH	1:K:375:PHE:HB2	2.00	0.61
1:L:362:TYR:HA	1:L:366:GLY:O	2.00	0.61
1:M:157:ASP:HB2	2:M:401:ADP:H5'1	1.83	0.61
1:U:298:VAL:HG12	1:U:335:ARG:HH11	1.64	0.61
1:C:40:HIS:O	1:C:41:GLN:C	2.43	0.61
1:C:157:ASP:HB2	2:C:401:ADP:H5'1	1.83	0.61
1:G:157:ASP:HB2	2:G:401:ADP:H5'1	1.83	0.61
1:I:36:GLY:N	1:I:54:VAL:HG23	2.16	0.61
1:K:44:MET:HG3	1:K:45:VAL:N	2.15	0.61
1:K:133:TYR:HB2	1:K:356:TRP:CE3	2.35	0.61
1:N:36:GLY:N	1:N:54:VAL:HG23	2.16	0.61
1:O:133:TYR:HB2	1:O:356:TRP:CE3	2.35	0.61
1:Q:133:TYR:OH	1:Q:375:PHE:HB2	2.00	0.61
1:T:157:ASP:HB2	2:T:401:ADP:H5'1	1.83	0.61
1:U:34:ILE:HB	1:U:54:VAL:HG13	1.80	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:34:ILE:HB	1:A:54:VAL:HG13	1.80	0.61
1:F:40:HIS:O	1:F:41:GLN:C	2.43	0.61
1:L:104:LEU:HB2	1:L:356:TRP:HH2	1.63	0.61
1:M:36:GLY:N	1:M:54:VAL:HG23	2.16	0.61
1:N:40:HIS:O	1:N:41:GLN:C	2.43	0.61
1:O:362:TYR:HA	1:O:366:GLY:O	2.00	0.61
1:P:44:MET:HG3	1:P:45:VAL:N	2.15	0.61
1:P:157:ASP:HB2	2:P:401:ADP:H5'1	1.83	0.61
1:U:40:HIS:O	1:U:41:GLN:C	2.43	0.61
1:U:58:ALA:O	1:U:61:LYS:N	2.30	0.61
1:D:40:HIS:O	1:D:41:GLN:C	2.43	0.61
1:F:147:ARG:HH21	1:F:147:ARG:CG	2.12	0.61
1:G:44:MET:HG3	1:G:45:VAL:N	2.15	0.61
1:J:36:GLY:N	1:J:54:VAL:HG23	2.16	0.61
1:L:36:GLY:N	1:L:54:VAL:HG23	2.16	0.61
1:M:40:HIS:O	1:M:41:GLN:C	2.43	0.61
1:P:36:GLY:N	1:P:54:VAL:HG23	2.16	0.61
1:P:325:MET:CE	1:R:244:ASP:O	2.49	0.61
1:A:61:LYS:HG2	1:A:64:ILE:HG21	1.83	0.61
1:B:40:HIS:O	1:B:41:GLN:C	2.43	0.61
1:F:61:LYS:HG2	1:F:64:ILE:HG21	1.83	0.61
1:I:34:ILE:CD1	1:I:67:LEU:HD22	2.31	0.61
1:L:58:ALA:O	1:L:61:LYS:N	2.30	0.61
1:L:61:LYS:HG2	1:L:64:ILE:HG21	1.83	0.61
1:M:133:TYR:HB2	1:M:356:TRP:CE3	2.35	0.61
1:N:325:MET:CE	1:P:244:ASP:O	2.49	0.61
1:O:104:LEU:HB2	1:O:356:TRP:HH2	1.62	0.61
1:R:40:HIS:O	1:R:41:GLN:C	2.43	0.61
1:S:36:GLY:N	1:S:54:VAL:HG23	2.16	0.61
1:S:44:MET:HG3	1:S:45:VAL:N	2.15	0.61
1:T:44:MET:HG3	1:T:45:VAL:N	2.15	0.61
1:V:133:TYR:OH	1:V:375:PHE:HB2	2.00	0.61
1:A:325:MET:CE	1:C:244:ASP:O	2.49	0.60
1:B:132:MET:O	1:B:357:ILE:HB	2.01	0.60
1:E:34:ILE:HB	1:E:54:VAL:HG13	1.80	0.60
1:H:61:LYS:HG2	1:H:64:ILE:HG21	1.83	0.60
1:K:325:MET:CE	1:M:244:ASP:O	2.49	0.60
1:N:362:TYR:HA	1:N:366:GLY:O	2.00	0.60
1:Q:36:GLY:N	1:Q:54:VAL:HG23	2.16	0.60
1:Q:40:HIS:O	1:Q:41:GLN:C	2.43	0.60
1:Q:61:LYS:HG2	1:Q:64:ILE:HG21	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:34:ILE:HB	1:S:54:VAL:HG13	1.79	0.60
1:S:157:ASP:HB2	2:S:401:ADP:H5'1	1.83	0.60
1:T:36:GLY:N	1:T:54:VAL:HG23	2.16	0.60
1:V:40:HIS:O	1:V:41:GLN:C	2.43	0.60
1:V:104:LEU:HB2	1:V:356:TRP:HH2	1.63	0.60
1:W:61:LYS:HG2	1:W:64:ILE:HG21	1.83	0.60
1:B:346:LEU:O	1:B:349:LEU:HB2	2.02	0.60
1:C:34:ILE:HB	1:C:54:VAL:HG13	1.79	0.60
1:C:133:TYR:OH	1:C:375:PHE:HB2	2.00	0.60
1:D:132:MET:O	1:D:357:ILE:HB	2.01	0.60
1:E:61:LYS:HG2	1:E:64:ILE:HG21	1.83	0.60
1:F:133:TYR:OH	1:F:375:PHE:HB2	2.00	0.60
1:F:157:ASP:HB2	2:F:401:ADP:H5'1	1.83	0.60
1:G:16:LEU:HD23	1:G:32:PRO:HA	1.83	0.60
1:G:40:HIS:O	1:G:41:GLN:C	2.43	0.60
1:G:133:TYR:OH	1:G:375:PHE:HB2	2.00	0.60
1:H:36:GLY:N	1:H:54:VAL:HG23	2.16	0.60
1:J:61:LYS:HG2	1:J:64:ILE:HG21	1.83	0.60
1:J:132:MET:O	1:J:357:ILE:HB	2.01	0.60
1:L:325:MET:CE	1:N:244:ASP:O	2.49	0.60
1:M:133:TYR:OH	1:M:375:PHE:HB2	2.00	0.60
1:N:61:LYS:HG2	1:N:64:ILE:HG21	1.83	0.60
1:O:40:HIS:O	1:O:41:GLN:C	2.43	0.60
1:O:325:MET:CE	1:Q:244:ASP:O	2.49	0.60
1:P:362:TYR:HA	1:P:366:GLY:O	2.00	0.60
1:R:36:GLY:N	1:R:54:VAL:HG23	2.16	0.60
1:R:61:LYS:HG2	1:R:64:ILE:HG21	1.83	0.60
1:R:325:MET:CE	1:T:244:ASP:O	2.49	0.60
1:S:61:LYS:HG2	1:S:64:ILE:HG21	1.83	0.60
1:T:132:MET:O	1:T:357:ILE:HB	2.01	0.60
1:U:132:MET:O	1:U:357:ILE:HB	2.01	0.60
1:U:133:TYR:OH	1:U:375:PHE:HB2	2.00	0.60
1:V:132:MET:O	1:V:357:ILE:HB	2.01	0.60
1:C:44:MET:HG3	1:C:45:VAL:N	2.15	0.60
1:D:61:LYS:HG2	1:D:64:ILE:HG21	1.83	0.60
1:E:133:TYR:OH	1:E:375:PHE:HB2	2.00	0.60
1:E:288:ASP:HA	1:G:243:PRO:HB2	1.83	0.60
1:I:325:MET:CE	1:K:244:ASP:O	2.49	0.60
1:K:40:HIS:O	1:K:41:GLN:C	2.43	0.60
1:O:36:GLY:N	1:O:54:VAL:HG23	2.16	0.60
1:P:132:MET:O	1:P:357:ILE:HB	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:325:MET:CE	1:S:244:ASP:O	2.49	0.60
1:Q:346:LEU:O	1:Q:349:LEU:HB2	2.02	0.60
1:R:132:MET:O	1:R:357:ILE:HB	2.01	0.60
1:S:325:MET:CE	1:U:244:ASP:O	2.49	0.60
1:U:36:GLY:N	1:U:54:VAL:HG23	2.16	0.60
1:U:104:LEU:HB2	1:U:356:TRP:HH2	1.62	0.60
1:W:132:MET:O	1:W:357:ILE:HB	2.01	0.60
1:A:157:ASP:HB2	2:A:401:ADP:H5'1	1.83	0.60
1:B:36:GLY:N	1:B:54:VAL:HG23	2.16	0.60
1:C:61:LYS:HG2	1:C:64:ILE:HG21	1.83	0.60
1:C:325:MET:CE	1:E:244:ASP:O	2.49	0.60
1:E:36:GLY:N	1:E:54:VAL:HG23	2.16	0.60
1:E:346:LEU:O	1:E:349:LEU:HB2	2.02	0.60
1:F:44:MET:HG3	1:F:45:VAL:N	2.15	0.60
1:G:288:ASP:HA	1:I:243:PRO:HB2	1.84	0.60
1:J:325:MET:CE	1:L:244:ASP:O	2.49	0.60
1:L:132:MET:O	1:L:357:ILE:HB	2.01	0.60
1:N:157:ASP:HB2	2:N:401:ADP:H5'1	1.83	0.60
1:P:61:LYS:HG2	1:P:64:ILE:HG21	1.83	0.60
1:R:288:ASP:HA	1:T:243:PRO:HB2	1.84	0.60
1:R:346:LEU:O	1:R:349:LEU:HB2	2.02	0.60
1:T:288:ASP:HA	1:V:243:PRO:HB2	1.84	0.60
1:T:346:LEU:O	1:T:349:LEU:HB2	2.02	0.60
1:U:61:LYS:HG2	1:U:64:ILE:HG21	1.83	0.60
1:V:34:ILE:CD1	1:V:67:LEU:HD22	2.31	0.60
1:C:346:LEU:O	1:C:349:LEU:HB2	2.02	0.60
1:E:325:MET:CE	1:G:244:ASP:O	2.49	0.60
1:F:34:ILE:CD1	1:F:67:LEU:HD22	2.31	0.60
1:F:132:MET:O	1:F:357:ILE:HB	2.01	0.60
1:G:61:LYS:HG2	1:G:64:ILE:HG21	1.83	0.60
1:H:34:ILE:CD1	1:H:67:LEU:HD22	2.31	0.60
1:K:346:LEU:O	1:K:349:LEU:HB2	2.02	0.60
1:L:34:ILE:HB	1:L:54:VAL:HG13	1.79	0.60
1:N:132:MET:O	1:N:357:ILE:HB	2.01	0.60
1:R:16:LEU:HD23	1:R:32:PRO:HA	1.83	0.60
1:R:34:ILE:CD1	1:R:67:LEU:HD22	2.31	0.60
1:T:34:ILE:CD1	1:T:67:LEU:HD22	2.31	0.60
1:T:61:LYS:HG2	1:T:64:ILE:HG21	1.83	0.60
1:T:325:MET:CE	1:V:244:ASP:O	2.49	0.60
1:U:34:ILE:CD1	1:U:67:LEU:HD22	2.31	0.60
1:U:157:ASP:HB2	2:U:401:ADP:H5'1	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:325:MET:CE	1:W:244:ASP:O	2.49	0.60
1:W:34:ILE:CD1	1:W:67:LEU:HD22	2.31	0.60
1:W:36:GLY:N	1:W:54:VAL:HG23	2.16	0.60
1:W:157:ASP:HB2	2:W:401:ADP:H5'1	1.83	0.60
1:A:36:GLY:N	1:A:54:VAL:HG23	2.16	0.60
1:B:61:LYS:HG2	1:B:64:ILE:HG21	1.83	0.60
1:B:133:TYR:OH	1:B:375:PHE:HB2	2.00	0.60
1:D:325:MET:CE	1:F:244:ASP:O	2.49	0.60
1:F:36:GLY:N	1:F:54:VAL:HG23	2.16	0.60
1:G:325:MET:CE	1:I:244:ASP:O	2.49	0.60
1:H:325:MET:CE	1:J:244:ASP:O	2.49	0.60
1:J:157:ASP:HB2	2:J:401:ADP:H5'1	1.83	0.60
1:L:54:VAL:O	1:L:55:GLY:C	2.45	0.60
1:N:133:TYR:OH	1:N:375:PHE:HB2	2.00	0.60
1:V:16:LEU:HD23	1:V:32:PRO:HA	1.83	0.60
1:W:54:VAL:O	1:W:55:GLY:C	2.45	0.60
1:A:54:VAL:O	1:A:55:GLY:C	2.45	0.60
1:A:132:MET:O	1:A:357:ILE:HB	2.01	0.60
1:E:16:LEU:HD23	1:E:32:PRO:HA	1.83	0.60
1:F:325:MET:CE	1:H:244:ASP:O	2.49	0.60
1:G:324:THR:CG2	1:I:241:GLU:OE2	2.50	0.60
1:H:132:MET:O	1:H:357:ILE:HB	2.01	0.60
1:J:133:TYR:OH	1:J:375:PHE:HB2	2.00	0.60
1:J:346:LEU:O	1:J:349:LEU:HB2	2.02	0.60
1:L:157:ASP:HB2	2:L:401:ADP:H5'1	1.83	0.60
1:M:34:ILE:CD1	1:M:67:LEU:HD22	2.31	0.60
1:P:133:TYR:OH	1:P:375:PHE:HB2	2.00	0.60
1:P:324:THR:CG2	1:R:241:GLU:OE2	2.50	0.60
1:A:288:ASP:HA	1:C:243:PRO:HB2	1.84	0.60
1:F:54:VAL:O	1:F:55:GLY:C	2.45	0.60
1:H:54:VAL:O	1:H:55:GLY:C	2.45	0.60
1:H:157:ASP:HB2	2:H:401:ADP:H5'1	1.83	0.60
1:H:346:LEU:O	1:H:349:LEU:HB2	2.02	0.60
1:I:346:LEU:O	1:I:349:LEU:HB2	2.02	0.60
1:K:288:ASP:HA	1:M:243:PRO:HB2	1.83	0.60
1:M:61:LYS:HG2	1:M:64:ILE:HG21	1.83	0.60
1:O:61:LYS:HG2	1:O:64:ILE:HG21	1.83	0.60
1:C:288:ASP:HA	1:E:243:PRO:HB2	1.84	0.60
1:D:34:ILE:CD1	1:D:67:LEU:HD22	2.31	0.60
1:K:36:GLY:N	1:K:54:VAL:HG23	2.16	0.60
1:L:346:LEU:O	1:L:349:LEU:HB2	2.02	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:16:LEU:HD23	1:M:32:PRO:HA	1.83	0.60
1:N:288:ASP:HA	1:P:243:PRO:HB2	1.83	0.60
1:O:44:MET:HG3	1:O:45:VAL:N	2.15	0.60
1:O:346:LEU:O	1:O:349:LEU:HB2	2.02	0.60
1:P:288:ASP:HA	1:R:243:PRO:HB2	1.84	0.60
1:R:133:TYR:OH	1:R:375:PHE:HB2	2.00	0.60
1:S:133:TYR:OH	1:S:375:PHE:HB2	2.00	0.60
1:S:346:LEU:O	1:S:349:LEU:HB2	2.02	0.60
1:W:133:TYR:OH	1:W:375:PHE:HB2	2.00	0.60
1:C:36:GLY:N	1:C:54:VAL:HG23	2.16	0.60
1:C:132:MET:O	1:C:357:ILE:HB	2.01	0.60
1:D:36:GLY:N	1:D:54:VAL:HG23	2.16	0.60
1:D:346:LEU:O	1:D:349:LEU:HB2	2.02	0.60
1:E:324:THR:CG2	1:G:241:GLU:OE2	2.50	0.60
1:G:36:GLY:N	1:G:54:VAL:HG23	2.16	0.60
1:H:16:LEU:HD23	1:H:32:PRO:HA	1.83	0.60
1:I:288:ASP:HA	1:K:243:PRO:HB2	1.84	0.60
1:J:34:ILE:CD1	1:J:67:LEU:HD22	2.31	0.60
1:K:34:ILE:CD1	1:K:67:LEU:HD22	2.31	0.60
1:K:61:LYS:HG2	1:K:64:ILE:HG21	1.83	0.60
1:L:44:MET:HG3	1:L:45:VAL:N	2.15	0.60
1:M:325:MET:CE	1:O:244:ASP:O	2.49	0.60
1:S:54:VAL:O	1:S:55:GLY:C	2.45	0.60
1:V:36:GLY:N	1:V:54:VAL:HG23	2.16	0.60
1:W:16:LEU:HD23	1:W:32:PRO:HA	1.83	0.60
1:A:324:THR:CG2	1:C:241:GLU:OE2	2.50	0.59
1:C:34:ILE:CD1	1:C:67:LEU:HD22	2.31	0.59
1:C:324:THR:CG2	1:E:241:GLU:OE2	2.50	0.59
1:H:104:LEU:HB2	1:H:356:TRP:HH2	1.63	0.59
1:H:287:ILE:CD1	1:J:208:ILE:CD1	2.80	0.59
1:I:324:THR:CG2	1:K:241:GLU:OE2	2.50	0.59
1:K:16:LEU:HD23	1:K:32:PRO:HA	1.83	0.59
1:L:133:TYR:OH	1:L:375:PHE:HB2	2.00	0.59
1:N:54:VAL:O	1:N:55:GLY:C	2.45	0.59
1:Q:54:VAL:O	1:Q:55:GLY:C	2.45	0.59
1:Q:132:MET:O	1:Q:357:ILE:HB	2.01	0.59
1:S:34:ILE:CD1	1:S:67:LEU:HD22	2.31	0.59
1:T:104:LEU:HB2	1:T:356:TRP:HH2	1.63	0.59
1:U:16:LEU:HD23	1:U:32:PRO:HA	1.83	0.59
1:V:61:LYS:HG2	1:V:64:ILE:HG21	1.83	0.59
1:C:43:VAL:O	1:C:44:MET:CG	2.51	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:54:VAL:O	1:C:55:GLY:C	2.45	0.59
1:D:143:TYR:HD1	1:D:143:TYR:C	2.09	0.59
1:E:132:MET:O	1:E:357:ILE:HB	2.01	0.59
1:G:34:ILE:CD1	1:G:67:LEU:HD22	2.31	0.59
1:K:8:LEU:HB3	1:K:103:THR:HG23	1.84	0.59
1:M:288:ASP:HA	1:O:243:PRO:HB2	1.83	0.59
1:M:346:LEU:O	1:M:349:LEU:HB2	2.02	0.59
1:N:346:LEU:O	1:N:349:LEU:HB2	2.02	0.59
1:O:132:MET:O	1:O:357:ILE:HB	2.01	0.59
1:P:16:LEU:HD23	1:P:32:PRO:HA	1.83	0.59
1:Q:34:ILE:HB	1:Q:54:VAL:HG13	1.79	0.59
1:R:54:VAL:O	1:R:55:GLY:C	2.45	0.59
1:A:34:ILE:CD1	1:A:67:LEU:HD22	2.31	0.59
1:F:43:VAL:O	1:F:44:MET:CG	2.51	0.59
1:F:324:THR:CG2	1:H:241:GLU:OE2	2.50	0.59
1:H:43:VAL:O	1:H:44:MET:CG	2.51	0.59
1:I:8:LEU:HB3	1:I:103:THR:HG23	1.84	0.59
1:J:43:VAL:O	1:J:44:MET:CG	2.51	0.59
1:J:143:TYR:HD1	1:J:143:TYR:C	2.09	0.59
1:L:288:ASP:HA	1:N:243:PRO:HB2	1.84	0.59
1:N:43:VAL:O	1:N:44:MET:CG	2.51	0.59
1:N:287:ILE:CD1	1:P:208:ILE:CD1	2.80	0.59
1:O:34:ILE:CD1	1:O:67:LEU:HD22	2.31	0.59
1:R:287:ILE:HG13	1:R:288:ASP:OD1	2.03	0.59
1:R:324:THR:CG2	1:T:241:GLU:OE2	2.50	0.59
1:S:324:THR:CG2	1:U:241:GLU:OE2	2.50	0.59
1:T:43:VAL:O	1:T:44:MET:CG	2.51	0.59
1:U:324:THR:CG2	1:W:241:GLU:OE2	2.50	0.59
1:W:34:ILE:HB	1:W:54:VAL:HG13	1.79	0.59
1:W:346:LEU:O	1:W:349:LEU:HB2	2.02	0.59
1:A:16:LEU:HD23	1:A:32:PRO:HA	1.83	0.59
1:B:325:MET:CE	1:D:244:ASP:O	2.49	0.59
1:E:104:LEU:HB2	1:E:356:TRP:HH2	1.63	0.59
1:F:104:LEU:HB2	1:F:356:TRP:HH2	1.63	0.59
1:G:104:LEU:HB2	1:G:356:TRP:HH2	1.63	0.59
1:G:346:LEU:O	1:G:349:LEU:HB2	2.02	0.59
1:H:133:TYR:OH	1:H:375:PHE:HB2	2.00	0.59
1:I:61:LYS:HG2	1:I:64:ILE:HG21	1.83	0.59
1:K:290:ARG:HH11	1:M:244:ASP:CB	2.16	0.59
1:K:324:THR:CG2	1:M:241:GLU:OE2	2.50	0.59
1:M:8:LEU:HB3	1:M:103:THR:HG23	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:43:VAL:O	1:R:44:MET:CG	2.51	0.59
1:S:16:LEU:HD23	1:S:32:PRO:HA	1.83	0.59
1:S:132:MET:O	1:S:357:ILE:HB	2.01	0.59
1:V:43:VAL:O	1:V:44:MET:CG	2.51	0.59
1:A:346:LEU:O	1:A:349:LEU:HB2	2.02	0.59
1:B:324:THR:CG2	1:D:241:GLU:OE2	2.50	0.59
1:D:43:VAL:O	1:D:44:MET:CG	2.51	0.59
1:F:16:LEU:HD23	1:F:32:PRO:HA	1.83	0.59
1:G:8:LEU:HB3	1:G:103:THR:HG23	1.84	0.59
1:G:132:MET:O	1:G:357:ILE:HB	2.01	0.59
1:I:16:LEU:HD23	1:I:32:PRO:HA	1.83	0.59
1:J:16:LEU:HD23	1:J:32:PRO:HA	1.83	0.59
1:N:34:ILE:CD1	1:N:67:LEU:HD22	2.31	0.59
1:N:324:THR:CG2	1:P:241:GLU:OE2	2.50	0.59
1:O:208:ILE:HD11	1:O:243:PRO:HG2	1.85	0.59
1:O:287:ILE:HG13	1:O:288:ASP:OD1	2.03	0.59
1:P:54:VAL:O	1:P:55:GLY:C	2.45	0.59
1:Q:16:LEU:HD23	1:Q:32:PRO:HA	1.83	0.59
1:S:208:ILE:HD11	1:S:243:PRO:HG2	1.85	0.59
1:S:288:ASP:HA	1:U:243:PRO:HB2	1.83	0.59
1:T:290:ARG:HH11	1:V:244:ASP:CB	2.16	0.59
1:A:64:ILE:HG23	1:A:65:LEU:N	2.18	0.59
1:B:16:LEU:HD23	1:B:32:PRO:HA	1.83	0.59
1:B:44:MET:HG3	1:B:45:VAL:N	2.15	0.59
1:D:324:THR:CG2	1:F:241:GLU:OE2	2.50	0.59
1:E:64:ILE:HG23	1:E:65:LEU:N	2.18	0.59
1:F:288:ASP:HA	1:H:243:PRO:HB2	1.84	0.59
1:H:324:THR:CG2	1:J:241:GLU:OE2	2.50	0.59
1:I:287:ILE:HG13	1:I:288:ASP:OD1	2.03	0.59
1:J:54:VAL:O	1:J:55:GLY:C	2.45	0.59
1:J:362:TYR:O	1:J:366:GLY:CA	2.51	0.59
1:K:361:GLU:HB3	1:K:369:ILE:CD1	2.14	0.59
1:M:132:MET:O	1:M:357:ILE:HB	2.01	0.59
1:P:34:ILE:CD1	1:P:67:LEU:HD22	2.31	0.59
1:P:43:VAL:O	1:P:44:MET:CG	2.51	0.59
1:P:64:ILE:HG23	1:P:65:LEU:N	2.18	0.59
1:P:287:ILE:HG13	1:P:288:ASP:OD1	2.03	0.59
1:P:346:LEU:O	1:P:349:LEU:HB2	2.02	0.59
1:Q:208:ILE:HD11	1:Q:243:PRO:HG2	1.85	0.59
1:R:362:TYR:O	1:R:366:GLY:CA	2.51	0.59
1:S:143:TYR:HD1	1:S:143:TYR:C	2.09	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:43:VAL:O	1:U:44:MET:CG	2.51	0.59
1:V:346:LEU:O	1:V:349:LEU:HB2	2.02	0.59
1:B:43:VAL:O	1:B:44:MET:CG	2.51	0.59
1:B:54:VAL:O	1:B:55:GLY:C	2.45	0.59
1:D:16:LEU:HD23	1:D:32:PRO:HA	1.83	0.59
1:G:43:VAL:O	1:G:44:MET:CG	2.51	0.59
1:J:324:THR:CG2	1:L:241:GLU:OE2	2.50	0.59
1:L:16:LEU:HD23	1:L:32:PRO:HA	1.83	0.59
1:L:287:ILE:HG13	1:L:288:ASP:OD1	2.03	0.59
1:L:362:TYR:O	1:L:366:GLY:CA	2.51	0.59
1:M:143:TYR:HD1	1:M:143:TYR:C	2.09	0.59
1:M:208:ILE:HD11	1:M:243:PRO:HG2	1.85	0.59
1:N:287:ILE:HG13	1:N:288:ASP:OD1	2.03	0.59
1:O:8:LEU:HB3	1:O:103:THR:HG23	1.84	0.59
1:Q:34:ILE:CD1	1:Q:67:LEU:HD22	2.31	0.59
1:S:43:VAL:O	1:S:44:MET:CG	2.51	0.59
1:T:324:THR:CG2	1:V:241:GLU:OE2	2.50	0.59
1:B:290:ARG:HH11	1:D:244:ASP:CB	2.16	0.59
1:C:362:TYR:O	1:C:366:GLY:CA	2.51	0.59
1:E:8:LEU:HB3	1:E:103:THR:HG23	1.84	0.59
1:E:34:ILE:CD1	1:E:67:LEU:HD22	2.31	0.59
1:E:223:PHE:CE1	1:E:259:GLU:HG2	2.38	0.59
1:J:104:LEU:HB2	1:J:356:TRP:HH2	1.63	0.59
1:K:54:VAL:O	1:K:55:GLY:C	2.45	0.59
1:L:43:VAL:O	1:L:44:MET:CG	2.51	0.59
1:L:223:PHE:CE1	1:L:259:GLU:HG2	2.38	0.59
1:L:287:ILE:CD1	1:N:208:ILE:CD1	2.80	0.59
1:N:362:TYR:O	1:N:366:GLY:CA	2.51	0.59
1:P:223:PHE:CE1	1:P:259:GLU:HG2	2.38	0.59
1:P:362:TYR:O	1:P:366:GLY:CA	2.51	0.59
1:S:287:ILE:CD1	1:U:208:ILE:CD1	2.80	0.59
1:T:54:VAL:O	1:T:55:GLY:C	2.45	0.59
1:T:287:ILE:HG13	1:T:288:ASP:OD1	2.03	0.59
1:U:54:VAL:O	1:U:55:GLY:C	2.45	0.59
1:U:346:LEU:O	1:U:349:LEU:HB2	2.02	0.59
1:W:43:VAL:O	1:W:44:MET:CG	2.51	0.59
1:W:223:PHE:CE1	1:W:259:GLU:HG2	2.38	0.59
1:W:287:ILE:HG13	1:W:288:ASP:OD1	2.03	0.59
1:D:143:TYR:CD1	1:D:143:TYR:C	2.75	0.59
1:D:287:ILE:HG13	1:D:288:ASP:OD1	2.03	0.59
1:D:288:ASP:HA	1:F:243:PRO:HB2	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:346:LEU:O	1:F:349:LEU:HB2	2.02	0.59
1:G:54:VAL:O	1:G:55:GLY:C	2.45	0.59
1:I:104:LEU:HB2	1:I:356:TRP:HH2	1.62	0.59
1:J:287:ILE:HG13	1:J:288:ASP:OD1	2.03	0.59
1:K:132:MET:O	1:K:357:ILE:HB	2.01	0.59
1:L:64:ILE:HG23	1:L:65:LEU:N	2.18	0.59
1:O:324:THR:CG2	1:Q:241:GLU:OE2	2.50	0.59
1:P:287:ILE:CD1	1:R:208:ILE:CD1	2.80	0.59
1:R:361:GLU:HB3	1:R:369:ILE:CD1	2.14	0.59
1:U:208:ILE:HD11	1:U:243:PRO:HG2	1.85	0.59
1:U:287:ILE:HG13	1:U:288:ASP:OD1	2.03	0.59
1:U:288:ASP:HA	1:W:243:PRO:HB2	1.84	0.59
1:A:133:TYR:OH	1:A:375:PHE:HB2	2.00	0.59
1:E:290:ARG:HH11	1:G:244:ASP:CB	2.16	0.59
1:E:362:TYR:O	1:E:366:GLY:CA	2.51	0.59
1:F:223:PHE:CE1	1:F:259:GLU:HG2	2.38	0.59
1:G:324:THR:HG21	1:I:241:GLU:OE2	2.03	0.59
1:I:54:VAL:O	1:I:55:GLY:C	2.45	0.59
1:I:132:MET:O	1:I:357:ILE:HB	2.01	0.59
1:I:324:THR:HG21	1:K:241:GLU:OE2	2.03	0.59
1:J:8:LEU:HB3	1:J:103:THR:HG23	1.84	0.59
1:J:223:PHE:CE1	1:J:259:GLU:HG2	2.38	0.59
1:J:288:ASP:HA	1:L:243:PRO:HB2	1.83	0.59
1:K:208:ILE:HD11	1:K:243:PRO:HG2	1.85	0.59
1:L:34:ILE:CD1	1:L:67:LEU:HD22	2.31	0.59
1:L:324:THR:CG2	1:N:241:GLU:OE2	2.50	0.59
1:M:324:THR:CG2	1:O:241:GLU:OE2	2.50	0.59
1:O:223:PHE:CE1	1:O:259:GLU:HG2	2.38	0.59
1:Q:287:ILE:HG13	1:Q:288:ASP:OD1	2.03	0.59
1:Q:288:ASP:HA	1:S:243:PRO:HB2	1.84	0.59
1:Q:324:THR:CG2	1:S:241:GLU:OE2	2.50	0.59
1:R:34:ILE:HB	1:R:54:VAL:HG13	1.80	0.59
1:S:287:ILE:HG13	1:S:288:ASP:OD1	2.03	0.59
1:T:16:LEU:HD23	1:T:32:PRO:HA	1.83	0.59
1:T:64:ILE:HG23	1:T:65:LEU:N	2.18	0.59
1:V:8:LEU:HB3	1:V:103:THR:HG23	1.84	0.59
1:V:287:ILE:HG13	1:V:288:ASP:OD1	2.03	0.59
1:W:8:LEU:HB3	1:W:103:THR:HG23	1.84	0.59
1:A:362:TYR:O	1:A:366:GLY:CA	2.51	0.58
1:B:208:ILE:HD11	1:B:243:PRO:HG2	1.85	0.58
1:D:54:VAL:O	1:D:55:GLY:C	2.45	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:324:THR:HG21	1:G:241:GLU:OE2	2.03	0.58
1:H:287:ILE:HG13	1:H:288:ASP:OD1	2.03	0.58
1:H:288:ASP:HA	1:J:243:PRO:HB2	1.84	0.58
1:I:34:ILE:HB	1:I:54:VAL:HG13	1.80	0.58
1:K:223:PHE:CE1	1:K:259:GLU:HG2	2.38	0.58
1:L:8:LEU:HB3	1:L:103:THR:HG23	1.84	0.58
1:M:290:ARG:HH11	1:O:244:ASP:CB	2.16	0.58
1:O:54:VAL:O	1:O:55:GLY:C	2.45	0.58
1:Q:43:VAL:O	1:Q:44:MET:CG	2.51	0.58
1:Q:223:PHE:CE1	1:Q:259:GLU:HG2	2.38	0.58
1:Q:290:ARG:HH11	1:S:244:ASP:CB	2.16	0.58
1:T:143:TYR:HD1	1:T:143:TYR:C	2.09	0.58
1:T:362:TYR:O	1:T:366:GLY:CA	2.51	0.58
1:V:223:PHE:CE1	1:V:259:GLU:HG2	2.38	0.58
1:A:223:PHE:CE1	1:A:259:GLU:HG2	2.38	0.58
1:C:324:THR:HG21	1:E:241:GLU:OE2	2.03	0.58
1:D:223:PHE:CE1	1:D:259:GLU:HG2	2.38	0.58
1:F:287:ILE:CD1	1:H:208:ILE:CD1	2.80	0.58
1:F:287:ILE:HG13	1:F:288:ASP:OD1	2.03	0.58
1:I:223:PHE:CE1	1:I:259:GLU:HG2	2.38	0.58
1:I:290:ARG:HH11	1:K:244:ASP:CB	2.16	0.58
1:J:64:ILE:HG23	1:J:65:LEU:N	2.18	0.58
1:K:287:ILE:HG13	1:K:288:ASP:OD1	2.03	0.58
1:K:324:THR:HG21	1:M:241:GLU:OE2	2.03	0.58
1:N:16:LEU:HD23	1:N:32:PRO:HA	1.83	0.58
1:O:288:ASP:HA	1:Q:243:PRO:HB2	1.83	0.58
1:R:287:ILE:CD1	1:T:208:ILE:CD1	2.80	0.58
1:R:290:ARG:HH11	1:T:244:ASP:CB	2.16	0.58
1:T:56:ASP:C	1:T:56:ASP:OD1	2.47	0.58
1:U:223:PHE:CE1	1:U:259:GLU:HG2	2.38	0.58
1:U:362:TYR:O	1:U:366:GLY:CA	2.51	0.58
1:W:362:TYR:O	1:W:366:GLY:CA	2.51	0.58
1:B:8:LEU:HB3	1:B:103:THR:HG23	1.84	0.58
1:B:34:ILE:CD1	1:B:67:LEU:HD22	2.31	0.58
1:B:38:PRO:CD	1:B:49:GLN:HE22	2.17	0.58
1:B:288:ASP:HA	1:D:243:PRO:HB2	1.83	0.58
1:C:104:LEU:HB2	1:C:356:TRP:HH2	1.63	0.58
1:C:287:ILE:CD1	1:E:208:ILE:CD1	2.80	0.58
1:C:290:ARG:HH11	1:E:244:ASP:CB	2.16	0.58
1:E:54:VAL:O	1:E:55:GLY:C	2.45	0.58
1:E:56:ASP:C	1:E:56:ASP:OD1	2.47	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:290:ARG:HH11	1:I:244:ASP:CB	2.16	0.58
1:G:362:TYR:O	1:G:366:GLY:CA	2.51	0.58
1:H:44:MET:HG3	1:H:45:VAL:N	2.15	0.58
1:H:290:ARG:HH11	1:J:244:ASP:CB	2.16	0.58
1:K:38:PRO:CD	1:K:49:GLN:HE22	2.17	0.58
1:K:287:ILE:CD1	1:M:208:ILE:CD1	2.80	0.58
1:M:38:PRO:CD	1:M:49:GLN:HE22	2.17	0.58
1:Q:8:LEU:HB3	1:Q:103:THR:HG23	1.84	0.58
1:R:324:THR:HG21	1:T:241:GLU:OE2	2.03	0.58
1:U:44:MET:HG3	1:U:45:VAL:N	2.15	0.58
1:U:64:ILE:HG23	1:U:65:LEU:N	2.18	0.58
1:A:324:THR:HG21	1:C:241:GLU:OE2	2.03	0.58
1:B:34:ILE:HB	1:B:54:VAL:HG13	1.79	0.58
1:B:56:ASP:OD1	1:B:56:ASP:C	2.47	0.58
1:C:16:LEU:HD23	1:C:32:PRO:HA	1.83	0.58
1:D:104:LEU:HB2	1:D:356:TRP:HH2	1.63	0.58
1:F:362:TYR:O	1:F:366:GLY:CA	2.51	0.58
1:H:8:LEU:HB3	1:H:103:THR:HG23	1.84	0.58
1:I:56:ASP:C	1:I:56:ASP:OD1	2.47	0.58
1:L:336:LYS:CE	2:L:401:ADP:H5'2	2.34	0.58
1:M:43:VAL:O	1:M:44:MET:CG	2.51	0.58
1:M:104:LEU:HB2	1:M:356:TRP:HH2	1.63	0.58
1:N:64:ILE:HG23	1:N:65:LEU:N	2.18	0.58
1:N:336:LYS:CE	2:N:401:ADP:H5'2	2.34	0.58
1:S:56:ASP:C	1:S:56:ASP:OD1	2.47	0.58
1:S:104:LEU:HB2	1:S:356:TRP:HH2	1.63	0.58
1:U:8:LEU:HB3	1:U:103:THR:HG23	1.84	0.58
1:V:38:PRO:CD	1:V:49:GLN:HE22	2.17	0.58
1:W:208:ILE:HD11	1:W:243:PRO:HG2	1.85	0.58
1:A:8:LEU:HB3	1:A:103:THR:HG23	1.84	0.58
1:A:43:VAL:O	1:A:44:MET:CG	2.51	0.58
1:B:361:GLU:HB3	1:B:369:ILE:CD1	2.14	0.58
1:D:8:LEU:HB3	1:D:103:THR:HG23	1.84	0.58
1:E:43:VAL:O	1:E:44:MET:CG	2.51	0.58
1:G:34:ILE:HB	1:G:54:VAL:HG13	1.79	0.58
1:J:290:ARG:HH11	1:L:244:ASP:CB	2.16	0.58
1:M:56:ASP:C	1:M:56:ASP:OD1	2.47	0.58
1:O:16:LEU:HD23	1:O:32:PRO:HA	1.83	0.58
1:O:43:VAL:O	1:O:44:MET:CG	2.51	0.58
1:P:56:ASP:C	1:P:56:ASP:OD1	2.47	0.58
1:T:223:PHE:CE1	1:T:259:GLU:HG2	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:287:ILE:CD1	1:W:208:ILE:CD1	2.80	0.58
1:U:336:LYS:CE	2:U:401:ADP:H5'2	2.34	0.58
1:V:208:ILE:HD11	1:V:243:PRO:HG2	1.85	0.58
1:B:287:ILE:HG13	1:B:288:ASP:OD1	2.03	0.58
1:C:8:LEU:HB3	1:C:103:THR:HG23	1.84	0.58
1:D:56:ASP:C	1:D:56:ASP:OD1	2.47	0.58
1:D:208:ILE:HD11	1:D:243:PRO:HG2	1.85	0.58
1:D:287:ILE:CD1	1:F:208:ILE:CD1	2.80	0.58
1:E:143:TYR:HD1	1:E:143:TYR:C	2.09	0.58
1:E:287:ILE:CD1	1:G:208:ILE:CD1	2.80	0.58
1:E:336:LYS:CE	2:E:401:ADP:H5'2	2.34	0.58
1:G:223:PHE:CE1	1:G:259:GLU:HG2	2.38	0.58
1:H:56:ASP:C	1:H:56:ASP:OD1	2.47	0.58
1:H:362:TYR:O	1:H:366:GLY:CA	2.51	0.58
1:I:362:TYR:O	1:I:366:GLY:CA	2.51	0.58
1:J:324:THR:HG21	1:L:241:GLU:OE2	2.03	0.58
1:M:54:VAL:O	1:M:55:GLY:C	2.45	0.58
1:M:324:THR:HG21	1:O:241:GLU:OE2	2.04	0.58
1:N:8:LEU:HB3	1:N:103:THR:HG23	1.84	0.58
1:N:34:ILE:CG2	1:N:67:LEU:HB3	2.34	0.58
1:N:173:HIS:CE1	1:O:267:ILE:C	2.82	0.58
1:O:38:PRO:CD	1:O:49:GLN:HE22	2.17	0.58
1:Q:56:ASP:C	1:Q:56:ASP:OD1	2.47	0.58
1:Q:173:HIS:CE1	1:R:267:ILE:C	2.82	0.58
1:W:56:ASP:C	1:W:56:ASP:OD1	2.47	0.58
1:W:64:ILE:HG23	1:W:65:LEU:N	2.18	0.58
1:B:173:HIS:CE1	1:C:267:ILE:C	2.82	0.58
1:C:287:ILE:HG13	1:C:288:ASP:OD1	2.03	0.58
1:D:290:ARG:HH11	1:F:244:ASP:CB	2.16	0.58
1:E:287:ILE:HG13	1:E:288:ASP:OD1	2.03	0.58
1:F:8:LEU:HB3	1:F:103:THR:HG23	1.84	0.58
1:F:56:ASP:C	1:F:56:ASP:OD1	2.47	0.58
1:I:43:VAL:O	1:I:44:MET:CG	2.51	0.58
1:I:208:ILE:HD11	1:I:243:PRO:HG2	1.85	0.58
1:K:43:VAL:O	1:K:44:MET:CG	2.51	0.58
1:K:173:HIS:CE1	1:L:267:ILE:C	2.82	0.58
1:L:324:THR:HG21	1:N:241:GLU:OE2	2.03	0.58
1:M:287:ILE:HG13	1:M:288:ASP:OD1	2.03	0.58
1:N:143:TYR:CD1	1:N:143:TYR:C	2.75	0.58
1:Q:287:ILE:CD1	1:S:208:ILE:CD1	2.80	0.58
1:S:38:PRO:CD	1:S:49:GLN:HE22	2.17	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:290:ARG:HH11	1:U:244:ASP:CB	2.16	0.58
1:S:336:LYS:CE	2:S:401:ADP:H5'2	2.34	0.58
1:T:208:ILE:HD11	1:T:243:PRO:HG2	1.85	0.58
1:U:173:HIS:CE1	1:V:267:ILE:C	2.82	0.58
1:V:54:VAL:O	1:V:55:GLY:C	2.45	0.58
1:V:173:HIS:CE1	1:W:267:ILE:C	2.82	0.58
1:A:56:ASP:C	1:A:56:ASP:OD1	2.47	0.58
1:D:38:PRO:CD	1:D:49:GLN:HE22	2.17	0.58
1:E:173:HIS:CE1	1:F:267:ILE:C	2.82	0.58
1:G:173:HIS:CE1	1:H:267:ILE:C	2.82	0.58
1:G:287:ILE:HG13	1:G:288:ASP:OD1	2.03	0.58
1:G:336:LYS:CE	2:G:401:ADP:H5'2	2.34	0.58
1:H:324:THR:HG21	1:J:241:GLU:OE2	2.03	0.58
1:I:64:ILE:HG23	1:I:65:LEU:N	2.18	0.58
1:J:336:LYS:CE	2:J:401:ADP:H5'2	2.34	0.58
1:L:4:GLU:O	1:L:5:THR:HB	2.04	0.58
1:L:56:ASP:C	1:L:56:ASP:OD1	2.47	0.58
1:M:362:TYR:O	1:M:366:GLY:CA	2.51	0.58
1:N:324:THR:HG21	1:P:241:GLU:OE2	2.03	0.58
1:O:56:ASP:C	1:O:56:ASP:OD1	2.47	0.58
1:P:34:ILE:CG2	1:P:67:LEU:HB3	2.34	0.58
1:P:173:HIS:CE1	1:Q:267:ILE:C	2.82	0.58
1:P:290:ARG:HH11	1:R:244:ASP:CB	2.16	0.58
1:P:336:LYS:CE	2:P:401:ADP:H5'2	2.34	0.58
1:T:8:LEU:HB3	1:T:103:THR:HG23	1.84	0.58
1:T:38:PRO:CD	1:T:49:GLN:HE22	2.17	0.58
1:W:336:LYS:CE	2:W:401:ADP:H5'2	2.34	0.58
1:C:56:ASP:C	1:C:56:ASP:OD1	2.47	0.58
1:C:336:LYS:CE	2:C:401:ADP:H5'2	2.34	0.58
1:D:368:SER:O	1:D:371:HIS:N	2.37	0.58
1:F:64:ILE:HG23	1:F:65:LEU:N	2.18	0.58
1:F:324:THR:HG21	1:H:241:GLU:OE2	2.03	0.58
1:H:64:ILE:HG23	1:H:65:LEU:N	2.18	0.58
1:H:223:PHE:CE1	1:H:259:GLU:HG2	2.38	0.58
1:L:143:TYR:HD1	1:L:143:TYR:C	2.09	0.58
1:L:173:HIS:CE1	1:M:267:ILE:C	2.82	0.58
1:O:34:ILE:HG21	1:O:67:LEU:CB	2.34	0.58
1:O:362:TYR:O	1:O:366:GLY:CA	2.51	0.58
1:P:4:GLU:O	1:P:5:THR:HB	2.04	0.58
1:P:324:THR:HG21	1:R:241:GLU:OE2	2.03	0.58
1:Q:362:TYR:O	1:Q:366:GLY:CA	2.51	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:34:ILE:CG2	1:R:67:LEU:HB3	2.34	0.58
1:R:208:ILE:HD11	1:R:243:PRO:HG2	1.85	0.58
1:R:223:PHE:CE1	1:R:259:GLU:HG2	2.38	0.58
1:S:368:SER:O	1:S:371:HIS:N	2.37	0.58
1:T:173:HIS:CE1	1:U:267:ILE:C	2.82	0.58
1:U:38:PRO:CD	1:U:49:GLN:HE22	2.17	0.58
1:U:56:ASP:C	1:U:56:ASP:OD1	2.47	0.58
1:W:4:GLU:O	1:W:5:THR:HB	2.04	0.58
1:A:4:GLU:O	1:A:5:THR:HB	2.04	0.58
1:B:34:ILE:CG2	1:B:67:LEU:HB3	2.34	0.58
1:B:362:TYR:O	1:B:366:GLY:CA	2.51	0.58
1:D:34:ILE:CG2	1:D:67:LEU:HB3	2.34	0.58
1:F:173:HIS:CE1	1:G:267:ILE:C	2.82	0.58
1:F:290:ARG:HH11	1:H:244:ASP:CB	2.16	0.58
1:G:4:GLU:O	1:G:5:THR:HB	2.04	0.58
1:H:173:HIS:CE1	1:I:267:ILE:C	2.82	0.58
1:I:38:PRO:CD	1:I:49:GLN:HE22	2.17	0.58
1:L:34:ILE:CG2	1:L:67:LEU:HB3	2.34	0.58
1:M:34:ILE:HG21	1:M:67:LEU:CB	2.34	0.58
1:M:223:PHE:CE1	1:M:259:GLU:HG2	2.38	0.58
1:N:290:ARG:HH11	1:P:244:ASP:CB	2.16	0.58
1:O:64:ILE:HG23	1:O:65:LEU:N	2.18	0.58
1:P:8:LEU:HB3	1:P:103:THR:HG23	1.84	0.58
1:Q:34:ILE:HG21	1:Q:67:LEU:CB	2.34	0.58
1:S:173:HIS:CE1	1:T:267:ILE:C	2.82	0.58
1:U:290:ARG:HH11	1:W:244:ASP:CB	2.16	0.58
1:V:362:TYR:O	1:V:366:GLY:CA	2.51	0.58
1:A:290:ARG:HH11	1:C:244:ASP:CB	2.16	0.57
1:B:64:ILE:HG23	1:B:65:LEU:N	2.18	0.57
1:B:223:PHE:CE1	1:B:259:GLU:HG2	2.38	0.57
1:B:324:THR:HG21	1:D:241:GLU:OE2	2.03	0.57
1:D:324:THR:HG21	1:F:241:GLU:OE2	2.03	0.57
1:D:362:TYR:O	1:D:366:GLY:CA	2.51	0.57
1:F:34:ILE:CG2	1:F:67:LEU:HB3	2.34	0.57
1:F:208:ILE:HD11	1:F:243:PRO:HG2	1.85	0.57
1:G:56:ASP:C	1:G:56:ASP:OD1	2.47	0.57
1:I:336:LYS:CE	2:I:401:ADP:H5'2	2.34	0.57
1:J:56:ASP:OD1	1:J:56:ASP:C	2.47	0.57
1:K:104:LEU:HB2	1:K:356:TRP:HH2	1.63	0.57
1:N:4:GLU:O	1:N:5:THR:HB	2.04	0.57
1:N:56:ASP:C	1:N:56:ASP:OD1	2.47	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:324:THR:HG21	1:Q:241:GLU:OE2	2.03	0.57
1:Q:38:PRO:CD	1:Q:49:GLN:HE22	2.17	0.57
1:Q:64:ILE:HG23	1:Q:65:LEU:N	2.18	0.57
1:Q:324:THR:HG21	1:S:241:GLU:OE2	2.03	0.57
1:R:34:ILE:HG21	1:R:67:LEU:CB	2.34	0.57
1:R:56:ASP:OD1	1:R:56:ASP:C	2.47	0.57
1:R:143:TYR:HD1	1:R:143:TYR:C	2.09	0.57
1:S:34:ILE:HG21	1:S:67:LEU:CB	2.34	0.57
1:S:223:PHE:CE1	1:S:259:GLU:HG2	2.38	0.57
1:T:34:ILE:HG21	1:T:67:LEU:CB	2.34	0.57
1:T:361:GLU:HB3	1:T:369:ILE:CD1	2.14	0.57
1:T:368:SER:O	1:T:371:HIS:N	2.37	0.57
1:A:287:ILE:CD1	1:C:208:ILE:CD1	2.80	0.57
1:A:287:ILE:HG13	1:A:288:ASP:OD1	2.03	0.57
1:B:287:ILE:CD1	1:D:208:ILE:CD1	2.80	0.57
1:B:336:LYS:CE	2:B:401:ADP:H5'2	2.34	0.57
1:C:38:PRO:CD	1:C:49:GLN:HE22	2.17	0.57
1:C:223:PHE:CE1	1:C:259:GLU:HG2	2.38	0.57
1:D:173:HIS:CE1	1:E:267:ILE:C	2.82	0.57
1:E:4:GLU:O	1:E:5:THR:HB	2.04	0.57
1:E:38:PRO:CD	1:E:49:GLN:HE22	2.17	0.57
1:G:34:ILE:CG2	1:G:67:LEU:HB3	2.34	0.57
1:G:64:ILE:HG23	1:G:65:LEU:N	2.18	0.57
1:H:4:GLU:O	1:H:5:THR:HB	2.04	0.57
1:H:34:ILE:HG21	1:H:67:LEU:CB	2.34	0.57
1:J:173:HIS:CE1	1:K:267:ILE:C	2.82	0.57
1:K:34:ILE:HG21	1:K:67:LEU:CB	2.34	0.57
1:K:64:ILE:HG23	1:K:65:LEU:N	2.18	0.57
1:K:362:TYR:O	1:K:366:GLY:CA	2.51	0.57
1:M:173:HIS:CE1	1:N:267:ILE:C	2.82	0.57
1:N:34:ILE:HG21	1:N:67:LEU:CB	2.34	0.57
1:N:143:TYR:HD1	1:N:143:TYR:C	2.09	0.57
1:O:173:HIS:CE1	1:P:267:ILE:C	2.82	0.57
1:O:368:SER:O	1:O:371:HIS:N	2.37	0.57
1:P:208:ILE:HD11	1:P:243:PRO:HG2	1.85	0.57
1:S:362:TYR:O	1:S:366:GLY:CA	2.51	0.57
1:V:34:ILE:HG21	1:V:67:LEU:CB	2.34	0.57
1:C:64:ILE:HG23	1:C:65:LEU:N	2.18	0.57
1:D:336:LYS:CE	2:D:401:ADP:H5'2	2.34	0.57
1:I:34:ILE:CG2	1:I:67:LEU:HB3	2.34	0.57
1:J:34:ILE:HG21	1:J:67:LEU:CB	2.34	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:38:PRO:CD	1:J:49:GLN:HE22	2.17	0.57
1:L:34:ILE:HG21	1:L:67:LEU:CB	2.34	0.57
1:P:34:ILE:HG21	1:P:67:LEU:CB	2.34	0.57
1:Q:336:LYS:CE	2:Q:401:ADP:H5'2	2.34	0.57
1:S:4:GLU:O	1:S:5:THR:HB	2.04	0.57
1:W:34:ILE:CG2	1:W:67:LEU:HB3	2.34	0.57
1:A:173:HIS:CE1	1:B:267:ILE:C	2.82	0.57
1:E:34:ILE:CG2	1:E:67:LEU:HB3	2.34	0.57
1:F:34:ILE:HG21	1:F:67:LEU:CB	2.34	0.57
1:F:38:PRO:CD	1:F:49:GLN:HE22	2.16	0.57
1:H:38:PRO:CD	1:H:49:GLN:HE22	2.17	0.57
1:I:34:ILE:HG21	1:I:67:LEU:CB	2.34	0.57
1:L:290:ARG:HH11	1:N:244:ASP:CB	2.16	0.57
1:N:208:ILE:HD11	1:N:243:PRO:HG2	1.85	0.57
1:N:223:PHE:CE1	1:N:259:GLU:HG2	2.38	0.57
1:R:336:LYS:CE	2:R:401:ADP:H5'2	2.34	0.57
1:T:287:ILE:CD1	1:V:208:ILE:CD1	2.80	0.57
1:U:34:ILE:HG21	1:U:67:LEU:CB	2.34	0.57
1:U:34:ILE:CG2	1:U:67:LEU:HB3	2.34	0.57
1:A:34:ILE:CG2	1:A:67:LEU:HB3	2.34	0.57
1:C:173:HIS:CE1	1:D:267:ILE:C	2.82	0.57
1:H:34:ILE:CG2	1:H:67:LEU:HB3	2.34	0.57
1:H:336:LYS:CE	2:H:401:ADP:H5'2	2.34	0.57
1:J:287:ILE:CD1	1:L:208:ILE:CD1	2.80	0.57
1:K:56:ASP:C	1:K:56:ASP:OD1	2.47	0.57
1:L:38:PRO:CD	1:L:49:GLN:HE22	2.17	0.57
1:L:208:ILE:HD11	1:L:243:PRO:HG2	1.85	0.57
1:R:64:ILE:HG23	1:R:65:LEU:N	2.18	0.57
1:S:8:LEU:HB3	1:S:103:THR:HG23	1.84	0.57
1:V:34:ILE:CG2	1:V:67:LEU:HB3	2.34	0.57
1:V:336:LYS:CE	2:V:401:ADP:H5'2	2.34	0.57
1:A:38:PRO:CD	1:A:49:GLN:HE22	2.17	0.57
1:A:208:ILE:HD11	1:A:243:PRO:HG2	1.85	0.57
1:B:34:ILE:HG21	1:B:67:LEU:CB	2.34	0.57
1:B:104:LEU:HB2	1:B:356:TRP:HH2	1.63	0.57
1:C:208:ILE:HD11	1:C:243:PRO:HG2	1.85	0.57
1:F:143:TYR:HD1	1:F:143:TYR:C	2.09	0.57
1:F:336:LYS:CE	2:F:401:ADP:H5'2	2.34	0.57
1:G:208:ILE:HD11	1:G:243:PRO:HG2	1.85	0.57
1:I:287:ILE:CD1	1:K:208:ILE:CD1	2.80	0.57
1:O:290:ARG:HH11	1:Q:244:ASP:CB	2.16	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:4:GLU:O	1:Q:5:THR:HB	2.04	0.57
1:Q:143:TYR:HD2	1:Q:346:LEU:HD13	1.70	0.57
1:T:4:GLU:O	1:T:5:THR:HB	2.04	0.57
1:T:34:ILE:CG2	1:T:67:LEU:HB3	2.34	0.57
1:T:324:THR:HG21	1:V:241:GLU:OE2	2.03	0.57
1:U:4:GLU:O	1:U:5:THR:HB	2.04	0.57
1:V:56:ASP:C	1:V:56:ASP:OD1	2.47	0.57
1:W:38:PRO:CD	1:W:49:GLN:HE22	2.17	0.57
1:A:336:LYS:CE	2:A:401:ADP:H5'2	2.34	0.57
1:D:34:ILE:HG21	1:D:67:LEU:CB	2.34	0.57
1:H:208:ILE:HD11	1:H:243:PRO:HG2	1.85	0.57
1:I:173:HIS:CE1	1:J:267:ILE:C	2.82	0.57
1:J:208:ILE:HD11	1:J:243:PRO:HG2	1.85	0.57
1:R:8:LEU:HB3	1:R:103:THR:HG23	1.84	0.57
1:R:173:HIS:CE1	1:S:267:ILE:C	2.82	0.57
1:V:143:TYR:HD1	1:V:143:TYR:C	2.09	0.57
1:G:34:ILE:HG21	1:G:67:LEU:CB	2.34	0.57
1:G:38:PRO:CD	1:G:49:GLN:HE22	2.17	0.57
1:H:143:TYR:HD2	1:H:346:LEU:HD13	1.70	0.57
1:I:4:GLU:O	1:I:5:THR:HB	2.04	0.57
1:J:143:TYR:HD2	1:J:346:LEU:HD13	1.70	0.57
1:K:34:ILE:CG2	1:K:67:LEU:HB3	2.34	0.57
1:K:336:LYS:CE	2:K:401:ADP:H5'2	2.34	0.57
1:K:368:SER:O	1:K:371:HIS:N	2.37	0.57
1:N:38:PRO:CD	1:N:49:GLN:HE22	2.17	0.57
1:O:34:ILE:CG2	1:O:67:LEU:HB3	2.34	0.57
1:R:38:PRO:CD	1:R:49:GLN:HE22	2.17	0.57
1:S:143:TYR:HD2	1:S:346:LEU:HD13	1.70	0.57
1:S:324:THR:HG21	1:U:241:GLU:OE2	2.03	0.57
1:U:324:THR:HG21	1:W:241:GLU:OE2	2.03	0.57
1:C:4:GLU:O	1:C:5:THR:HB	2.04	0.57
1:J:34:ILE:CG2	1:J:67:LEU:HB3	2.34	0.57
1:M:34:ILE:CG2	1:M:67:LEU:HB3	2.34	0.57
1:O:143:TYR:HD2	1:O:346:LEU:HD13	1.70	0.57
1:O:287:ILE:CD1	1:Q:208:ILE:CD1	2.80	0.57
1:O:336:LYS:CE	2:O:401:ADP:H5'2	2.34	0.57
1:Q:34:ILE:CG2	1:Q:67:LEU:HB3	2.34	0.57
1:S:34:ILE:CG2	1:S:67:LEU:HB3	2.34	0.57
1:T:336:LYS:CE	2:T:401:ADP:H5'2	2.34	0.57
1:C:34:ILE:CG2	1:C:67:LEU:HB3	2.34	0.57
1:M:64:ILE:HG23	1:M:65:LEU:N	2.18	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:38:PRO:CD	1:P:49:GLN:HE22	2.16	0.57
1:S:64:ILE:HG23	1:S:65:LEU:N	2.18	0.57
1:U:143:TYR:HD1	1:U:143:TYR:C	2.09	0.57
1:V:64:ILE:HG23	1:V:65:LEU:N	2.18	0.57
1:W:34:ILE:HG21	1:W:67:LEU:CB	2.34	0.57
1:B:143:TYR:CZ	1:D:45:VAL:CG2	2.88	0.56
1:B:290:ARG:HH11	1:D:244:ASP:HB2	1.71	0.56
1:G:143:TYR:HD1	1:G:143:TYR:C	2.09	0.56
1:G:290:ARG:HH11	1:I:244:ASP:HB2	1.70	0.56
1:I:290:ARG:HH11	1:K:244:ASP:HB2	1.70	0.56
1:M:287:ILE:CD1	1:O:208:ILE:CD1	2.80	0.56
1:R:4:GLU:O	1:R:5:THR:HB	2.04	0.56
1:U:143:TYR:HD2	1:U:346:LEU:HD13	1.70	0.56
1:A:34:ILE:HG21	1:A:67:LEU:CB	2.34	0.56
1:B:4:GLU:O	1:B:5:THR:HB	2.04	0.56
1:D:4:GLU:O	1:D:5:THR:HB	2.04	0.56
1:D:290:ARG:HH11	1:F:244:ASP:HB2	1.70	0.56
1:E:208:ILE:HD11	1:E:243:PRO:HG2	1.85	0.56
1:E:290:ARG:HH11	1:G:244:ASP:HB2	1.70	0.56
1:K:290:ARG:HH11	1:M:244:ASP:HB2	1.70	0.56
1:C:368:SER:O	1:C:371:HIS:N	2.37	0.56
1:D:64:ILE:HG23	1:D:65:LEU:N	2.18	0.56
1:E:34:ILE:HG21	1:E:67:LEU:CB	2.34	0.56
1:F:143:TYR:HD2	1:F:346:LEU:HD13	1.70	0.56
1:G:368:SER:O	1:G:371:HIS:N	2.37	0.56
1:J:4:GLU:O	1:J:5:THR:HB	2.04	0.56
1:L:143:TYR:HD2	1:L:346:LEU:HD13	1.70	0.56
1:M:4:GLU:O	1:M:5:THR:HB	2.04	0.56
1:N:34:ILE:HB	1:N:54:VAL:HG13	1.80	0.56
1:N:368:SER:O	1:N:371:HIS:N	2.37	0.56
1:R:290:ARG:HH11	1:T:244:ASP:HB2	1.70	0.56
1:T:143:TYR:CZ	1:V:45:VAL:CG2	2.88	0.56
1:C:34:ILE:HG21	1:C:67:LEU:CB	2.34	0.56
1:G:287:ILE:CD1	1:I:208:ILE:CD1	2.80	0.56
1:M:336:LYS:CE	2:M:401:ADP:H5'2	2.34	0.56
1:O:143:TYR:CZ	1:Q:45:VAL:CG2	2.88	0.56
1:P:290:ARG:HH11	1:R:244:ASP:HB2	1.70	0.56
1:Q:143:TYR:CZ	1:S:45:VAL:CG2	2.88	0.56
1:R:368:SER:O	1:R:371:HIS:N	2.37	0.56
1:V:368:SER:O	1:V:371:HIS:N	2.37	0.56
1:M:143:TYR:CZ	1:O:45:VAL:CG2	2.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:290:ARG:HH11	1:S:244:ASP:HB2	1.70	0.56
1:S:290:ARG:HH11	1:U:244:ASP:HB2	1.70	0.56
1:U:290:ARG:HH11	1:W:244:ASP:HB2	1.70	0.56
1:A:116:ARG:HD3	1:A:370:VAL:HG13	1.88	0.56
1:C:290:ARG:HH11	1:E:244:ASP:HB2	1.70	0.56
1:E:116:ARG:HD3	1:E:370:VAL:HG13	1.88	0.56
1:L:116:ARG:HD3	1:L:370:VAL:HG13	1.88	0.56
1:N:143:TYR:CZ	1:P:45:VAL:CG2	2.88	0.56
1:P:116:ARG:HD3	1:P:370:VAL:HG13	1.88	0.56
1:R:116:ARG:HD3	1:R:370:VAL:HG13	1.88	0.56
1:S:143:TYR:CZ	1:U:45:VAL:CG2	2.88	0.56
1:V:4:GLU:O	1:V:5:THR:HB	2.04	0.56
1:W:361:GLU:HB3	1:W:369:ILE:CD1	2.14	0.56
1:C:116:ARG:HD3	1:C:370:VAL:HG13	1.88	0.56
1:K:143:TYR:CZ	1:M:45:VAL:CG2	2.88	0.56
1:M:143:TYR:HD2	1:M:346:LEU:HD13	1.70	0.56
1:O:4:GLU:O	1:O:5:THR:HB	2.04	0.56
1:Q:362:TYR:HE1	1:Q:367:PRO:CB	2.19	0.56
1:J:368:SER:O	1:J:371:HIS:N	2.37	0.56
1:L:143:TYR:CZ	1:N:45:VAL:CG2	2.88	0.56
1:M:361:GLU:HB3	1:M:369:ILE:CD1	2.14	0.56
1:M:362:TYR:HE1	1:M:367:PRO:CB	2.19	0.56
1:N:116:ARG:HD3	1:N:370:VAL:HG13	1.88	0.56
1:R:143:TYR:CZ	1:T:45:VAL:CG2	2.88	0.56
1:U:34:ILE:CG2	1:U:67:LEU:HD22	2.28	0.56
1:W:143:TYR:HD2	1:W:346:LEU:HD13	1.70	0.56
1:D:143:TYR:HD2	1:D:346:LEU:HD13	1.70	0.56
1:K:4:GLU:O	1:K:5:THR:HB	2.04	0.56
1:N:290:ARG:HH11	1:P:244:ASP:HB2	1.70	0.56
1:P:143:TYR:CZ	1:R:45:VAL:CG2	2.88	0.56
1:V:361:GLU:HB3	1:V:369:ILE:CD1	2.14	0.56
1:I:143:TYR:CZ	1:K:45:VAL:CG2	2.88	0.56
1:L:290:ARG:HH11	1:N:244:ASP:HB2	1.70	0.56
1:M:290:ARG:HH11	1:O:244:ASP:HB2	1.70	0.56
1:T:116:ARG:HD3	1:T:370:VAL:HG13	1.88	0.56
1:U:143:TYR:CZ	1:W:45:VAL:CG2	2.88	0.56
1:F:290:ARG:HH11	1:H:244:ASP:HB2	1.70	0.55
1:F:362:TYR:HE1	1:F:367:PRO:CB	2.19	0.55
1:G:116:ARG:HD3	1:G:370:VAL:HG13	1.88	0.55
1:N:143:TYR:HD2	1:N:346:LEU:HD13	1.70	0.55
1:P:121:GLN:HA	1:P:362:TYR:OH	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:290:ARG:HH11	1:V:244:ASP:HB2	1.70	0.55
1:U:121:GLN:HA	1:U:362:TYR:OH	2.07	0.55
1:D:34:ILE:CG2	1:D:67:LEU:HD22	2.28	0.55
1:F:4:GLU:O	1:F:5:THR:HB	2.04	0.55
1:J:116:ARG:HD3	1:J:370:VAL:HG13	1.88	0.55
1:Q:116:ARG:HD3	1:Q:370:VAL:HG13	1.88	0.55
1:S:121:GLN:HA	1:S:362:TYR:OH	2.07	0.55
1:W:121:GLN:HA	1:W:362:TYR:OH	2.07	0.55
1:A:143:TYR:HD2	1:A:346:LEU:HD13	1.70	0.55
1:G:121:GLN:HA	1:G:362:TYR:OH	2.07	0.55
1:I:37:ARG:CG	1:I:38:PRO:CD	2.83	0.55
1:J:143:TYR:CZ	1:L:45:VAL:CG2	2.88	0.55
1:P:110:LEU:HD12	1:P:177:ARG:HH11	1.72	0.55
1:W:116:ARG:HD3	1:W:370:VAL:HG13	1.88	0.55
1:A:121:GLN:HA	1:A:362:TYR:OH	2.07	0.55
1:B:362:TYR:HE1	1:B:367:PRO:CB	2.19	0.55
1:C:143:TYR:HD2	1:C:346:LEU:HD13	1.70	0.55
1:D:116:ARG:HD3	1:D:370:VAL:HG13	1.88	0.55
1:N:110:LEU:HD12	1:N:177:ARG:HH11	1.72	0.55
1:W:37:ARG:CG	1:W:38:PRO:CD	2.83	0.55
1:A:143:TYR:CZ	1:C:45:VAL:CG2	2.88	0.55
1:E:369:ILE:C	1:E:371:HIS:H	2.15	0.55
1:F:368:SER:O	1:F:371:HIS:N	2.37	0.55
1:G:143:TYR:CZ	1:I:45:VAL:CG2	2.88	0.55
1:G:369:ILE:C	1:G:371:HIS:H	2.15	0.55
1:H:143:TYR:CZ	1:H:345:ILE:HG22	2.42	0.55
1:I:7:ALA:CB	1:I:347:ALA:HB1	2.37	0.55
1:I:110:LEU:HD12	1:I:177:ARG:HH11	1.72	0.55
1:I:121:GLN:HA	1:I:362:TYR:OH	2.07	0.55
1:J:290:ARG:HH11	1:L:244:ASP:HB2	1.70	0.55
1:K:110:LEU:HD12	1:K:177:ARG:HH11	1.72	0.55
1:M:116:ARG:HD3	1:M:370:VAL:HG13	1.88	0.55
1:N:121:GLN:HA	1:N:362:TYR:OH	2.07	0.55
1:O:32:PRO:HG3	1:O:59:GLN:NE2	2.22	0.55
1:O:116:ARG:HD3	1:O:370:VAL:HG13	1.87	0.55
1:O:121:GLN:HA	1:O:362:TYR:OH	2.07	0.55
1:O:290:ARG:HH11	1:Q:244:ASP:HB2	1.70	0.55
1:Q:121:GLN:HA	1:Q:362:TYR:OH	2.07	0.55
1:R:110:LEU:HD12	1:R:177:ARG:HH11	1.72	0.55
1:R:121:GLN:HA	1:R:362:TYR:OH	2.07	0.55
1:U:368:SER:O	1:U:371:HIS:N	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:116:ARG:HD3	1:V:370:VAL:HG13	1.88	0.55
1:W:369:ILE:C	1:W:371:HIS:H	2.15	0.55
1:B:116:ARG:HD3	1:B:370:VAL:HG13	1.88	0.55
1:C:369:ILE:C	1:C:371:HIS:H	2.15	0.55
1:D:32:PRO:HG3	1:D:59:GLN:NE2	2.22	0.55
1:E:143:TYR:CZ	1:E:345:ILE:HG22	2.42	0.55
1:F:116:ARG:HD3	1:F:370:VAL:HG13	1.87	0.55
1:G:143:TYR:CZ	1:G:345:ILE:HG22	2.42	0.55
1:I:369:ILE:C	1:I:371:HIS:H	2.15	0.55
1:J:7:ALA:CB	1:J:347:ALA:HB1	2.37	0.55
1:K:7:ALA:CB	1:K:347:ALA:HB1	2.37	0.55
1:L:7:ALA:CB	1:L:347:ALA:HB1	2.37	0.55
1:L:121:GLN:HA	1:L:362:TYR:OH	2.07	0.55
1:R:143:TYR:CZ	1:R:345:ILE:HG22	2.42	0.55
1:S:116:ARG:HD3	1:S:370:VAL:HG13	1.88	0.55
1:T:143:TYR:CZ	1:T:345:ILE:HG22	2.42	0.55
1:W:143:TYR:CZ	1:W:345:ILE:HG22	2.42	0.55
1:C:7:ALA:CB	1:C:347:ALA:HB1	2.37	0.55
1:E:32:PRO:HG3	1:E:59:GLN:NE2	2.22	0.55
1:G:7:ALA:CB	1:G:347:ALA:HB1	2.37	0.55
1:G:32:PRO:HG3	1:G:59:GLN:NE2	2.22	0.55
1:I:116:ARG:HD3	1:I:370:VAL:HG13	1.88	0.55
1:J:32:PRO:HG3	1:J:59:GLN:NE2	2.22	0.55
1:K:32:PRO:HG3	1:K:59:GLN:NE2	2.22	0.55
1:L:37:ARG:CG	1:L:38:PRO:CD	2.83	0.55
1:M:7:ALA:CB	1:M:347:ALA:HB1	2.37	0.55
1:N:7:ALA:CB	1:N:347:ALA:HB1	2.37	0.55
1:N:143:TYR:CZ	1:N:345:ILE:HG22	2.42	0.55
1:P:143:TYR:CZ	1:P:345:ILE:HG22	2.42	0.55
1:S:32:PRO:HG3	1:S:59:GLN:NE2	2.22	0.55
1:A:7:ALA:CB	1:A:347:ALA:HB1	2.37	0.55
1:A:290:ARG:HH11	1:C:244:ASP:HB2	1.70	0.55
1:A:369:ILE:C	1:A:371:HIS:H	2.15	0.55
1:B:121:GLN:HA	1:B:362:TYR:OH	2.07	0.55
1:C:37:ARG:CG	1:C:38:PRO:CD	2.83	0.55
1:F:143:TYR:CZ	1:H:45:VAL:CG2	2.88	0.55
1:F:236:LEU:HD12	1:F:237:GLU:CG	2.37	0.55
1:G:110:LEU:HD12	1:G:177:ARG:HH11	1.72	0.55
1:H:7:ALA:CB	1:H:347:ALA:HB1	2.37	0.55
1:H:37:ARG:CG	1:H:38:PRO:CD	2.83	0.55
1:H:236:LEU:HD12	1:H:237:GLU:CG	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:121:GLN:HA	1:M:362:TYR:OH	2.07	0.55
1:N:32:PRO:HG3	1:N:59:GLN:NE2	2.22	0.55
1:P:7:ALA:CB	1:P:347:ALA:HB1	2.37	0.55
1:V:121:GLN:HA	1:V:362:TYR:OH	2.07	0.55
1:V:143:TYR:CZ	1:V:345:ILE:HG22	2.42	0.55
1:V:143:TYR:HD2	1:V:346:LEU:HD13	1.70	0.55
1:W:32:PRO:HG3	1:W:59:GLN:NE2	2.22	0.55
1:W:110:LEU:HD12	1:W:177:ARG:HH11	1.72	0.55
1:A:143:TYR:CZ	1:A:345:ILE:HG22	2.42	0.55
1:B:236:LEU:HD12	1:B:237:GLU:CG	2.37	0.55
1:C:32:PRO:HG3	1:C:59:GLN:NE2	2.22	0.55
1:D:121:GLN:HA	1:D:362:TYR:OH	2.07	0.55
1:D:236:LEU:HD12	1:D:237:GLU:CG	2.37	0.55
1:D:361:GLU:HB3	1:D:369:ILE:CD1	2.14	0.55
1:E:7:ALA:CB	1:E:347:ALA:HB1	2.37	0.55
1:E:121:GLN:HA	1:E:362:TYR:OH	2.07	0.55
1:H:32:PRO:HG3	1:H:59:GLN:NE2	2.22	0.55
1:H:143:TYR:CZ	1:J:45:VAL:CG2	2.88	0.55
1:I:32:PRO:HG3	1:I:59:GLN:NE2	2.22	0.55
1:K:116:ARG:HD3	1:K:370:VAL:HG13	1.88	0.55
1:L:147:ARG:HG3	1:L:147:ARG:NH2	2.22	0.55
1:O:7:ALA:CB	1:O:347:ALA:HB1	2.37	0.55
1:O:143:TYR:CZ	1:O:345:ILE:HG22	2.42	0.55
1:T:236:LEU:HD12	1:T:237:GLU:CG	2.37	0.55
1:A:147:ARG:HG3	1:A:147:ARG:NH2	2.22	0.55
1:B:368:SER:O	1:B:371:HIS:N	2.37	0.55
1:C:143:TYR:CZ	1:E:45:VAL:CG2	2.88	0.55
1:E:143:TYR:CZ	1:G:45:VAL:CG2	2.88	0.55
1:E:362:TYR:HE1	1:E:367:PRO:CB	2.19	0.55
1:H:116:ARG:HD3	1:H:370:VAL:HG13	1.88	0.55
1:H:236:LEU:HD12	1:H:237:GLU:HG2	1.89	0.55
1:H:290:ARG:HH11	1:J:244:ASP:HB2	1.70	0.55
1:J:143:TYR:CZ	1:J:345:ILE:HG22	2.42	0.55
1:M:143:TYR:CZ	1:M:345:ILE:HG22	2.42	0.55
1:O:34:ILE:CG2	1:O:67:LEU:HD22	2.28	0.55
1:Q:143:TYR:CZ	1:Q:345:ILE:HG22	2.42	0.55
1:S:236:LEU:HD12	1:S:237:GLU:HG2	1.89	0.55
1:T:236:LEU:HD12	1:T:237:GLU:HG2	1.90	0.55
1:U:369:ILE:C	1:U:371:HIS:H	2.15	0.55
1:F:7:ALA:CB	1:F:347:ALA:HB1	2.37	0.54
1:F:143:TYR:CZ	1:F:345:ILE:HG22	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:236:LEU:HD12	1:F:237:GLU:HG2	1.89	0.54
1:H:147:ARG:NH2	1:H:147:ARG:HG3	2.22	0.54
1:J:121:GLN:HA	1:J:362:TYR:OH	2.07	0.54
1:Q:7:ALA:CB	1:Q:347:ALA:HB1	2.37	0.54
1:Q:147:ARG:HG3	1:Q:147:ARG:NH2	2.22	0.54
1:Q:236:LEU:HD12	1:Q:237:GLU:HG2	1.89	0.54
1:R:7:ALA:CB	1:R:347:ALA:HB1	2.37	0.54
1:R:236:LEU:HD12	1:R:237:GLU:CG	2.37	0.54
1:S:7:ALA:CB	1:S:347:ALA:HB1	2.37	0.54
1:T:110:LEU:HD12	1:T:177:ARG:HH11	1.72	0.54
1:T:369:ILE:C	1:T:371:HIS:H	2.15	0.54
1:V:32:PRO:HG3	1:V:59:GLN:NE2	2.22	0.54
1:W:147:ARG:HG3	1:W:147:ARG:NH2	2.22	0.54
1:A:32:PRO:HG3	1:A:59:GLN:NE2	2.22	0.54
1:A:110:LEU:HD12	1:A:177:ARG:HH11	1.72	0.54
1:B:143:TYR:HD2	1:B:346:LEU:HD13	1.70	0.54
1:D:34:ILE:HG21	1:D:67:LEU:HB3	1.90	0.54
1:D:37:ARG:CG	1:D:38:PRO:CD	2.83	0.54
1:D:143:TYR:CZ	1:F:45:VAL:CG2	2.88	0.54
1:E:143:TYR:HD2	1:E:346:LEU:HD13	1.70	0.54
1:F:34:ILE:HG21	1:F:67:LEU:HB3	1.90	0.54
1:F:147:ARG:HG3	1:F:147:ARG:NH2	2.22	0.54
1:G:236:LEU:HD12	1:G:237:GLU:CG	2.37	0.54
1:I:34:ILE:HG21	1:I:67:LEU:HB3	1.90	0.54
1:I:236:LEU:HD12	1:I:237:GLU:CG	2.37	0.54
1:I:236:LEU:HD12	1:I:237:GLU:HG2	1.89	0.54
1:J:236:LEU:HD12	1:J:237:GLU:CG	2.37	0.54
1:K:143:TYR:HD2	1:K:346:LEU:HD13	1.70	0.54
1:K:236:LEU:HD12	1:K:237:GLU:HG2	1.89	0.54
1:L:32:PRO:HG3	1:L:59:GLN:NE2	2.22	0.54
1:Q:368:SER:O	1:Q:371:HIS:N	2.37	0.54
1:R:236:LEU:HD12	1:R:237:GLU:HG2	1.89	0.54
1:S:34:ILE:HG21	1:S:67:LEU:HB3	1.90	0.54
1:U:7:ALA:CB	1:U:347:ALA:HB1	2.37	0.54
1:U:143:TYR:CZ	1:U:345:ILE:HG22	2.42	0.54
1:V:34:ILE:HG21	1:V:67:LEU:HB3	1.90	0.54
1:V:236:LEU:HD12	1:V:237:GLU:HG2	1.90	0.54
1:B:147:ARG:NH2	1:B:147:ARG:HG3	2.22	0.54
1:E:8:LEU:HB2	1:E:103:THR:OG1	2.08	0.54
1:E:34:ILE:HG21	1:E:67:LEU:HB3	1.90	0.54
1:J:369:ILE:C	1:J:371:HIS:H	2.15	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:34:ILE:HG21	1:K:67:LEU:HB3	1.90	0.54
1:K:369:ILE:C	1:K:371:HIS:H	2.15	0.54
1:L:143:TYR:CZ	1:L:345:ILE:HG22	2.42	0.54
1:O:236:LEU:HD12	1:O:237:GLU:HG2	1.89	0.54
1:Q:34:ILE:HG21	1:Q:67:LEU:HB3	1.90	0.54
1:Q:106:THR:HG22	1:Q:140:LEU:CD1	2.38	0.54
1:R:34:ILE:HG21	1:R:67:LEU:HB3	1.90	0.54
1:R:37:ARG:CG	1:R:38:PRO:CD	2.83	0.54
1:T:8:LEU:HB2	1:T:103:THR:OG1	2.08	0.54
1:U:34:ILE:HG21	1:U:67:LEU:HB3	1.90	0.54
1:U:110:LEU:HD12	1:U:177:ARG:HH11	1.72	0.54
1:U:116:ARG:HD3	1:U:370:VAL:HG13	1.88	0.54
1:U:147:ARG:NH2	1:U:147:ARG:HG3	2.22	0.54
1:V:8:LEU:HB2	1:V:103:THR:OG1	2.08	0.54
1:V:300:SER:HA	1:V:335:ARG:HG2	1.89	0.54
1:W:34:ILE:HG21	1:W:67:LEU:HB3	1.90	0.54
1:W:362:TYR:HE1	1:W:367:PRO:CB	2.19	0.54
1:A:37:ARG:CG	1:A:38:PRO:CD	2.83	0.54
1:B:34:ILE:HG21	1:B:67:LEU:HB3	1.89	0.54
1:B:300:SER:HA	1:B:335:ARG:HG2	1.89	0.54
1:C:8:LEU:HB2	1:C:103:THR:OG1	2.08	0.54
1:C:121:GLN:HA	1:C:362:TYR:OH	2.07	0.54
1:D:7:ALA:CB	1:D:347:ALA:HB1	2.37	0.54
1:E:223:PHE:CZ	1:E:266:PHE:HZ	2.26	0.54
1:F:32:PRO:HG3	1:F:59:GLN:NE2	2.22	0.54
1:G:236:LEU:HD12	1:G:237:GLU:HG2	1.90	0.54
1:I:8:LEU:HB2	1:I:103:THR:OG1	2.08	0.54
1:K:7:ALA:CB	1:K:356:TRP:CZ2	2.91	0.54
1:K:38:PRO:HB2	1:K:41:GLN:HB2	1.89	0.54
1:L:34:ILE:HG21	1:L:67:LEU:HB3	1.90	0.54
1:L:110:LEU:HD12	1:L:177:ARG:HH11	1.72	0.54
1:L:369:ILE:C	1:L:371:HIS:H	2.15	0.54
1:M:7:ALA:CB	1:M:356:TRP:CZ2	2.91	0.54
1:M:110:LEU:HD12	1:M:177:ARG:HH11	1.72	0.54
1:M:223:PHE:CZ	1:M:266:PHE:HZ	2.26	0.54
1:M:236:LEU:HD12	1:M:237:GLU:HG2	1.90	0.54
1:N:223:PHE:CZ	1:N:266:PHE:HZ	2.26	0.54
1:P:147:ARG:NH2	1:P:147:ARG:HG3	2.22	0.54
1:R:32:PRO:HG3	1:R:59:GLN:NE2	2.22	0.54
1:S:106:THR:HG22	1:S:140:LEU:CD1	2.38	0.54
1:T:38:PRO:HB2	1:T:41:GLN:HB2	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:32:PRO:HG3	1:U:59:GLN:NE2	2.22	0.54
1:V:236:LEU:HD12	1:V:237:GLU:CG	2.37	0.54
1:W:106:THR:HG22	1:W:140:LEU:CD1	2.38	0.54
1:A:236:LEU:HD12	1:A:237:GLU:CG	2.37	0.54
1:B:110:LEU:HD12	1:B:177:ARG:HH11	1.72	0.54
1:B:143:TYR:CZ	1:B:345:ILE:HG22	2.42	0.54
1:B:223:PHE:CZ	1:B:266:PHE:HZ	2.26	0.54
1:C:34:ILE:HG21	1:C:67:LEU:HB3	1.90	0.54
1:D:236:LEU:HD12	1:D:237:GLU:HG2	1.90	0.54
1:E:236:LEU:HD12	1:E:237:GLU:CG	2.37	0.54
1:H:34:ILE:HG21	1:H:67:LEU:HB3	1.90	0.54
1:H:121:GLN:HA	1:H:362:TYR:OH	2.07	0.54
1:I:7:ALA:CB	1:I:356:TRP:CZ2	2.91	0.54
1:I:38:PRO:HB2	1:I:41:GLN:HB2	1.89	0.54
1:J:34:ILE:HG21	1:J:67:LEU:HB3	1.90	0.54
1:J:236:LEU:HD12	1:J:237:GLU:HG2	1.90	0.54
1:M:8:LEU:HB2	1:M:103:THR:OG1	2.08	0.54
1:M:32:PRO:HG3	1:M:59:GLN:NE2	2.22	0.54
1:M:38:PRO:HB2	1:M:41:GLN:HB2	1.89	0.54
1:M:236:LEU:HD12	1:M:237:GLU:CG	2.37	0.54
1:O:7:ALA:CB	1:O:356:TRP:CZ2	2.91	0.54
1:O:34:ILE:HG21	1:O:67:LEU:HB3	1.90	0.54
1:P:8:LEU:HB2	1:P:103:THR:OG1	2.08	0.54
1:P:34:ILE:CG2	1:P:67:LEU:HD22	2.27	0.54
1:P:223:PHE:CZ	1:P:266:PHE:HZ	2.26	0.54
1:Q:236:LEU:HD12	1:Q:237:GLU:CG	2.37	0.54
1:R:369:ILE:C	1:R:371:HIS:H	2.15	0.54
1:S:37:ARG:CG	1:S:38:PRO:CD	2.83	0.54
1:S:236:LEU:HD12	1:S:237:GLU:CG	2.37	0.54
1:U:106:THR:HG22	1:U:140:LEU:CD1	2.38	0.54
1:U:236:LEU:HD12	1:U:237:GLU:CG	2.37	0.54
1:U:236:LEU:HD12	1:U:237:GLU:HG2	1.90	0.54
1:V:223:PHE:CZ	1:V:266:PHE:HZ	2.26	0.54
1:W:7:ALA:CB	1:W:347:ALA:HB1	2.37	0.54
1:C:223:PHE:CZ	1:C:266:PHE:HZ	2.26	0.54
1:D:110:LEU:HD12	1:D:177:ARG:HH11	1.72	0.54
1:D:223:PHE:CZ	1:D:266:PHE:HZ	2.26	0.54
1:E:38:PRO:HB2	1:E:41:GLN:HB2	1.89	0.54
1:F:121:GLN:HA	1:F:362:TYR:OH	2.07	0.54
1:G:34:ILE:HG21	1:G:67:LEU:HB3	1.90	0.54
1:H:369:ILE:C	1:H:371:HIS:H	2.15	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:143:TYR:CZ	1:I:345:ILE:HG22	2.42	0.54
1:K:143:TYR:CZ	1:K:345:ILE:HG22	2.42	0.54
1:K:236:LEU:HD12	1:K:237:GLU:CG	2.37	0.54
1:L:223:PHE:CZ	1:L:266:PHE:HZ	2.26	0.54
1:M:34:ILE:HG21	1:M:67:LEU:HB3	1.90	0.54
1:M:37:ARG:CG	1:M:38:PRO:CD	2.83	0.54
1:M:369:ILE:C	1:M:371:HIS:H	2.15	0.54
1:O:106:THR:HG22	1:O:140:LEU:CD1	2.38	0.54
1:O:223:PHE:CZ	1:O:266:PHE:HZ	2.26	0.54
1:O:236:LEU:HD12	1:O:237:GLU:CG	2.37	0.54
1:P:34:ILE:HG21	1:P:67:LEU:HB3	1.90	0.54
1:P:236:LEU:HD12	1:P:237:GLU:CG	2.37	0.54
1:S:369:ILE:C	1:S:371:HIS:H	2.15	0.54
1:T:7:ALA:CB	1:T:347:ALA:HB1	2.37	0.54
1:T:106:THR:HG22	1:T:140:LEU:CD1	2.38	0.54
1:T:121:GLN:HA	1:T:362:TYR:OH	2.07	0.54
1:W:223:PHE:CZ	1:W:266:PHE:HZ	2.26	0.54
1:A:140:LEU:HD22	1:A:343:GLY:HA2	1.90	0.54
1:B:7:ALA:CB	1:B:347:ALA:HB1	2.37	0.54
1:B:236:LEU:HD12	1:B:237:GLU:HG2	1.90	0.54
1:C:143:TYR:CZ	1:C:345:ILE:HG22	2.42	0.54
1:D:147:ARG:HG3	1:D:147:ARG:NH2	2.22	0.54
1:E:106:THR:HG22	1:E:140:LEU:CD1	2.38	0.54
1:E:110:LEU:HD12	1:E:177:ARG:HH11	1.72	0.54
1:G:7:ALA:CB	1:G:356:TRP:CZ2	2.91	0.54
1:G:38:PRO:HB2	1:G:41:GLN:HB2	1.89	0.54
1:K:106:THR:HG22	1:K:140:LEU:CD1	2.38	0.54
1:K:121:GLN:HA	1:K:362:TYR:OH	2.07	0.54
1:L:140:LEU:HD22	1:L:343:GLY:HA2	1.90	0.54
1:L:236:LEU:HD12	1:L:237:GLU:CG	2.37	0.54
1:N:369:ILE:C	1:N:371:HIS:H	2.15	0.54
1:O:8:LEU:HB2	1:O:103:THR:OG1	2.08	0.54
1:P:32:PRO:HG3	1:P:59:GLN:NE2	2.22	0.54
1:P:140:LEU:HD22	1:P:343:GLY:HA2	1.90	0.54
1:P:143:TYR:HD2	1:P:346:LEU:HD13	1.70	0.54
1:Q:7:ALA:CB	1:Q:356:TRP:CZ2	2.91	0.54
1:Q:32:PRO:HG3	1:Q:59:GLN:NE2	2.22	0.54
1:R:8:LEU:HB2	1:R:103:THR:OG1	2.08	0.54
1:R:38:PRO:HB2	1:R:41:GLN:HB2	1.90	0.54
1:R:143:TYR:HD2	1:R:346:LEU:HD13	1.70	0.54
1:R:223:PHE:CZ	1:R:266:PHE:HZ	2.26	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:147:ARG:NH2	1:S:147:ARG:HG3	2.22	0.54
1:T:34:ILE:HG21	1:T:67:LEU:HB3	1.90	0.54
1:T:300:SER:HA	1:T:335:ARG:HG2	1.89	0.54
1:V:7:ALA:CB	1:V:356:TRP:CZ2	2.91	0.54
1:A:8:LEU:HB2	1:A:103:THR:OG1	2.08	0.54
1:A:106:THR:HG22	1:A:140:LEU:CD1	2.38	0.54
1:A:143:TYR:HD1	1:A:143:TYR:C	2.09	0.54
1:D:300:SER:HA	1:D:335:ARG:HG2	1.89	0.54
1:F:223:PHE:CZ	1:F:266:PHE:HZ	2.26	0.54
1:G:143:TYR:HD2	1:G:346:LEU:HD13	1.70	0.54
1:J:147:ARG:NH2	1:J:147:ARG:HG3	2.22	0.54
1:K:223:PHE:CZ	1:K:266:PHE:HZ	2.26	0.54
1:M:147:ARG:NH2	1:M:147:ARG:HG3	2.22	0.54
1:N:34:ILE:HG21	1:N:67:LEU:HB3	1.89	0.54
1:P:236:LEU:HD12	1:P:237:GLU:HG2	1.89	0.54
1:R:7:ALA:CB	1:R:356:TRP:CZ2	2.91	0.54
1:S:143:TYR:CZ	1:S:345:ILE:HG22	2.42	0.54
1:V:38:PRO:HB2	1:V:41:GLN:HB2	1.90	0.54
1:A:34:ILE:HG21	1:A:67:LEU:HB3	1.89	0.54
1:B:106:THR:HG22	1:B:140:LEU:CD1	2.38	0.54
1:C:38:PRO:HB2	1:C:41:GLN:HB2	1.89	0.54
1:C:106:THR:HG22	1:C:140:LEU:CD1	2.38	0.54
1:C:143:TYR:HD1	1:C:143:TYR:C	2.09	0.54
1:D:208:ILE:HG22	1:D:209:VAL:N	2.23	0.54
1:E:236:LEU:HD12	1:E:237:GLU:HG2	1.90	0.54
1:G:223:PHE:CZ	1:G:266:PHE:HZ	2.26	0.54
1:I:360:GLN:O	1:I:364:GLU:HG3	2.08	0.54
1:J:106:THR:HG22	1:J:140:LEU:CD1	2.38	0.54
1:K:8:LEU:HB2	1:K:103:THR:OG1	2.08	0.54
1:L:106:THR:HG22	1:L:140:LEU:CD1	2.38	0.54
1:O:369:ILE:C	1:O:371:HIS:H	2.15	0.54
1:S:223:PHE:O	1:S:227:MET:HG2	2.08	0.54
1:T:7:ALA:CB	1:T:356:TRP:CZ2	2.91	0.54
1:T:32:PRO:CG	1:T:55:GLY:O	2.55	0.54
1:T:32:PRO:HG3	1:T:59:GLN:NE2	2.22	0.54
1:T:143:TYR:HD2	1:T:346:LEU:HD13	1.70	0.54
1:V:7:ALA:CB	1:V:347:ALA:HB1	2.37	0.54
1:V:362:TYR:HE1	1:V:367:PRO:CB	2.19	0.54
1:A:223:PHE:CZ	1:A:266:PHE:HZ	2.26	0.54
1:A:270:GLU:O	1:A:270:GLU:CD	2.51	0.54
1:B:32:PRO:HG3	1:B:59:GLN:NE2	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:208:ILE:HG22	1:B:209:VAL:N	2.23	0.54
1:C:140:LEU:HD22	1:C:343:GLY:HA2	1.90	0.54
1:C:236:LEU:HD12	1:C:237:GLU:CG	2.37	0.54
1:D:106:THR:HG22	1:D:140:LEU:CD1	2.38	0.54
1:E:147:ARG:NH2	1:E:147:ARG:HG3	2.22	0.54
1:F:106:THR:HG22	1:F:140:LEU:CD1	2.38	0.54
1:F:110:LEU:HD12	1:F:177:ARG:HH11	1.72	0.54
1:F:300:SER:HA	1:F:335:ARG:HG2	1.89	0.54
1:H:106:THR:HG22	1:H:140:LEU:CD1	2.38	0.54
1:J:140:LEU:HD22	1:J:343:GLY:HA2	1.90	0.54
1:J:223:PHE:CZ	1:J:266:PHE:HZ	2.26	0.54
1:M:368:SER:O	1:M:371:HIS:N	2.37	0.54
1:N:270:GLU:O	1:N:270:GLU:CD	2.52	0.54
1:P:37:ARG:CG	1:P:38:PRO:CD	2.83	0.54
1:Q:369:ILE:C	1:Q:371:HIS:H	2.15	0.54
1:T:223:PHE:CZ	1:T:266:PHE:HZ	2.26	0.54
1:W:236:LEU:HD12	1:W:237:GLU:CG	2.37	0.54
1:A:300:SER:HA	1:A:335:ARG:HG2	1.89	0.53
1:B:8:LEU:HB2	1:B:103:THR:OG1	2.08	0.53
1:C:32:PRO:CG	1:C:55:GLY:O	2.55	0.53
1:D:223:PHE:O	1:D:227:MET:HG2	2.08	0.53
1:F:270:GLU:O	1:F:270:GLU:CD	2.51	0.53
1:F:369:ILE:C	1:F:371:HIS:H	2.15	0.53
1:G:106:THR:HG22	1:G:140:LEU:CD1	2.38	0.53
1:H:135:ALA:HB1	1:H:140:LEU:HD21	1.91	0.53
1:H:147:ARG:CG	1:H:147:ARG:NH2	2.71	0.53
1:I:223:PHE:O	1:I:227:MET:HG2	2.09	0.53
1:L:362:TYR:HE1	1:L:367:PRO:CB	2.19	0.53
1:M:34:ILE:CG2	1:M:68:LYS:H	2.21	0.53
1:M:223:PHE:O	1:M:227:MET:HG2	2.09	0.53
1:N:223:PHE:O	1:N:227:MET:HG2	2.08	0.53
1:O:270:GLU:O	1:O:270:GLU:CD	2.51	0.53
1:P:7:ALA:CB	1:P:356:TRP:CZ2	2.91	0.53
1:P:38:PRO:HB2	1:P:41:GLN:HB2	1.89	0.53
1:P:360:GLN:O	1:P:364:GLU:HG3	2.08	0.53
1:R:32:PRO:CG	1:R:55:GLY:O	2.55	0.53
1:R:140:LEU:HD22	1:R:343:GLY:HA2	1.90	0.53
1:T:34:ILE:HB	1:T:54:VAL:HG13	1.80	0.53
1:T:34:ILE:CG2	1:T:68:LYS:H	2.21	0.53
1:U:223:PHE:CZ	1:U:266:PHE:HZ	2.26	0.53
1:U:270:GLU:O	1:U:270:GLU:CD	2.51	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:34:ILE:CG2	1:V:68:LYS:H	2.21	0.53
1:V:110:LEU:HD12	1:V:177:ARG:HH11	1.72	0.53
1:V:369:ILE:C	1:V:371:HIS:H	2.15	0.53
1:W:147:ARG:CG	1:W:147:ARG:NH2	2.71	0.53
1:W:236:LEU:HD12	1:W:237:GLU:HG2	1.90	0.53
1:W:270:GLU:O	1:W:270:GLU:CD	2.52	0.53
1:A:135:ALA:HB1	1:A:140:LEU:HD21	1.91	0.53
1:B:369:ILE:C	1:B:371:HIS:H	2.15	0.53
1:E:223:PHE:O	1:E:227:MET:HG2	2.08	0.53
1:G:360:GLN:O	1:G:364:GLU:HG3	2.08	0.53
1:H:270:GLU:O	1:H:270:GLU:CD	2.52	0.53
1:J:135:ALA:HB1	1:J:140:LEU:HD21	1.91	0.53
1:K:34:ILE:CG2	1:K:68:LYS:H	2.21	0.53
1:L:223:PHE:O	1:L:227:MET:HG2	2.09	0.53
1:L:236:LEU:HD12	1:L:237:GLU:HG2	1.89	0.53
1:L:270:GLU:O	1:L:270:GLU:CD	2.51	0.53
1:M:106:THR:HG22	1:M:140:LEU:CD1	2.38	0.53
1:N:106:THR:HG22	1:N:140:LEU:CD1	2.38	0.53
1:N:135:ALA:HB1	1:N:140:LEU:HD21	1.91	0.53
1:N:236:LEU:HD12	1:N:237:GLU:CG	2.37	0.53
1:P:369:ILE:C	1:P:371:HIS:H	2.15	0.53
1:Q:8:LEU:HB2	1:Q:103:THR:OG1	2.08	0.53
1:Q:270:GLU:O	1:Q:270:GLU:CD	2.52	0.53
1:R:270:GLU:O	1:R:270:GLU:CD	2.52	0.53
1:S:7:ALA:CB	1:S:356:TRP:CZ2	2.91	0.53
1:S:110:LEU:HD12	1:S:177:ARG:HH11	1.72	0.53
1:S:362:TYR:HE1	1:S:367:PRO:CB	2.19	0.53
1:U:38:PRO:HB2	1:U:41:GLN:HB2	1.89	0.53
1:V:173:HIS:NE2	1:W:267:ILE:C	2.67	0.53
1:W:38:PRO:HB2	1:W:41:GLN:HB2	1.90	0.53
1:A:360:GLN:O	1:A:364:GLU:HG3	2.08	0.53
1:B:7:ALA:CB	1:B:356:TRP:CZ2	2.91	0.53
1:B:116:ARG:HH12	1:B:375:PHE:HA	1.74	0.53
1:D:142:LEU:HD21	1:D:165:ILE:CD1	2.39	0.53
1:D:143:TYR:CZ	1:D:345:ILE:HG22	2.42	0.53
1:E:7:ALA:CB	1:E:356:TRP:CZ2	2.91	0.53
1:E:336:LYS:HE2	2:E:401:ADP:C5'	2.38	0.53
1:F:7:ALA:CB	1:F:356:TRP:CZ2	2.91	0.53
1:F:208:ILE:HG22	1:F:209:VAL:N	2.24	0.53
1:G:223:PHE:O	1:G:227:MET:HG2	2.09	0.53
1:H:223:PHE:CZ	1:H:266:PHE:HZ	2.26	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:106:THR:HG22	1:I:140:LEU:CD1	2.38	0.53
1:K:142:LEU:HD21	1:K:165:ILE:CD1	2.39	0.53
1:K:173:HIS:NE2	1:L:267:ILE:C	2.67	0.53
1:L:7:ALA:CB	1:L:356:TRP:CZ2	2.91	0.53
1:O:38:PRO:HB2	1:O:41:GLN:HB2	1.89	0.53
1:P:270:GLU:O	1:P:270:GLU:CD	2.52	0.53
1:R:106:THR:HG22	1:R:140:LEU:CD1	2.38	0.53
1:R:360:GLN:O	1:R:364:GLU:HG3	2.08	0.53
1:S:38:PRO:HB2	1:S:41:GLN:HB2	1.89	0.53
1:T:147:ARG:HG3	1:T:147:ARG:NH2	2.22	0.53
1:T:223:PHE:O	1:T:227:MET:HG2	2.08	0.53
1:U:135:ALA:HB1	1:U:140:LEU:HD21	1.91	0.53
1:U:140:LEU:HD22	1:U:343:GLY:HA2	1.90	0.53
1:V:116:ARG:HH12	1:V:375:PHE:HA	1.74	0.53
1:V:360:GLN:O	1:V:364:GLU:HG3	2.08	0.53
1:W:135:ALA:HB1	1:W:140:LEU:HD21	1.91	0.53
1:W:140:LEU:HD22	1:W:343:GLY:HA2	1.90	0.53
1:A:32:PRO:CG	1:A:55:GLY:O	2.55	0.53
1:A:236:LEU:HD12	1:A:237:GLU:HG2	1.89	0.53
1:B:173:HIS:NE2	1:C:267:ILE:C	2.67	0.53
1:C:135:ALA:HB1	1:C:140:LEU:HD21	1.91	0.53
1:C:270:GLU:O	1:C:270:GLU:CD	2.51	0.53
1:D:7:ALA:CB	1:D:356:TRP:CZ2	2.91	0.53
1:D:135:ALA:HB1	1:D:140:LEU:HD21	1.91	0.53
1:D:147:ARG:CG	1:D:147:ARG:NH2	2.71	0.53
1:D:270:GLU:O	1:D:270:GLU:CD	2.52	0.53
1:D:360:GLN:O	1:D:364:GLU:HG3	2.08	0.53
1:E:140:LEU:HD22	1:E:343:GLY:HA2	1.90	0.53
1:E:270:GLU:CD	1:E:270:GLU:O	2.51	0.53
1:F:135:ALA:HB1	1:F:140:LEU:HD21	1.90	0.53
1:F:223:PHE:O	1:F:227:MET:HG2	2.08	0.53
1:G:8:LEU:HB2	1:G:103:THR:OG1	2.08	0.53
1:H:110:LEU:HD12	1:H:177:ARG:HH11	1.72	0.53
1:J:270:GLU:O	1:J:270:GLU:CD	2.51	0.53
1:L:8:LEU:HB2	1:L:103:THR:OG1	2.08	0.53
1:N:140:LEU:HD22	1:N:343:GLY:HA2	1.90	0.53
1:O:173:HIS:NE2	1:P:267:ILE:C	2.67	0.53
1:P:135:ALA:HB1	1:P:140:LEU:HD21	1.91	0.53
1:Q:173:HIS:NE2	1:R:267:ILE:C	2.67	0.53
1:Q:223:PHE:CZ	1:Q:266:PHE:HZ	2.26	0.53
1:S:116:ARG:HH12	1:S:375:PHE:HA	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:147:ARG:CG	1:S:147:ARG:NH2	2.71	0.53
1:S:173:HIS:NE2	1:T:267:ILE:C	2.67	0.53
1:U:223:PHE:O	1:U:227:MET:HG2	2.09	0.53
1:V:106:THR:HG22	1:V:140:LEU:CD1	2.38	0.53
1:V:223:PHE:O	1:V:227:MET:HG2	2.08	0.53
1:W:7:ALA:CB	1:W:356:TRP:CZ2	2.91	0.53
1:A:34:ILE:CG2	1:A:67:LEU:HD22	2.28	0.53
1:A:362:TYR:HE1	1:A:367:PRO:CB	2.19	0.53
1:B:360:GLN:O	1:B:364:GLU:HG3	2.08	0.53
1:C:110:LEU:HD12	1:C:177:ARG:HH11	1.72	0.53
1:E:173:HIS:NE2	1:F:267:ILE:C	2.67	0.53
1:F:360:GLN:O	1:F:364:GLU:HG3	2.08	0.53
1:G:270:GLU:O	1:G:270:GLU:CD	2.52	0.53
1:H:7:ALA:CB	1:H:356:TRP:CZ2	2.91	0.53
1:I:32:PRO:CG	1:I:55:GLY:O	2.55	0.53
1:I:34:ILE:CG2	1:I:68:LYS:H	2.21	0.53
1:I:147:ARG:HG3	1:I:147:ARG:NH2	2.22	0.53
1:J:7:ALA:CB	1:J:356:TRP:CZ2	2.91	0.53
1:K:360:GLN:O	1:K:364:GLU:HG3	2.08	0.53
1:N:8:LEU:HB2	1:N:103:THR:OG1	2.08	0.53
1:O:208:ILE:HG22	1:O:209:VAL:N	2.23	0.53
1:Q:223:PHE:O	1:Q:227:MET:HG2	2.08	0.53
1:R:34:ILE:CG2	1:R:68:LYS:H	2.21	0.53
1:R:300:SER:HA	1:R:335:ARG:HG2	1.89	0.53
1:S:135:ALA:HB1	1:S:140:LEU:HD21	1.91	0.53
1:T:336:LYS:HE2	2:T:401:ADP:C5'	2.38	0.53
1:W:142:LEU:HD21	1:W:165:ILE:CD1	2.39	0.53
1:A:38:PRO:HB2	1:A:41:GLN:HB2	1.89	0.53
1:B:37:ARG:CG	1:B:38:PRO:CD	2.83	0.53
1:B:270:GLU:O	1:B:270:GLU:CD	2.52	0.53
1:C:147:ARG:CG	1:C:147:ARG:NH2	2.71	0.53
1:C:360:GLN:O	1:C:364:GLU:HG3	2.08	0.53
1:D:116:ARG:HH12	1:D:375:PHE:HA	1.74	0.53
1:D:173:HIS:NE2	1:E:267:ILE:C	2.67	0.53
1:D:369:ILE:C	1:D:371:HIS:H	2.15	0.53
1:F:336:LYS:HE2	2:F:401:ADP:C5'	2.38	0.53
1:H:8:LEU:HB2	1:H:103:THR:OG1	2.08	0.53
1:I:116:ARG:HH12	1:I:375:PHE:HA	1.74	0.53
1:I:173:HIS:NE2	1:J:267:ILE:C	2.67	0.53
1:I:270:GLU:O	1:I:270:GLU:CD	2.51	0.53
1:J:8:LEU:HB2	1:J:103:THR:OG1	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:110:LEU:HD12	1:J:177:ARG:HH11	1.72	0.53
1:K:208:ILE:HG22	1:K:209:VAL:N	2.23	0.53
1:M:208:ILE:HG22	1:M:209:VAL:N	2.23	0.53
1:N:7:ALA:CB	1:N:356:TRP:CZ2	2.91	0.53
1:O:147:ARG:HG3	1:O:147:ARG:NH2	2.22	0.53
1:P:223:PHE:O	1:P:227:MET:HG2	2.08	0.53
1:Q:116:ARG:HH12	1:Q:375:PHE:HA	1.74	0.53
1:Q:135:ALA:HB1	1:Q:140:LEU:HD21	1.91	0.53
1:Q:208:ILE:HG22	1:Q:209:VAL:N	2.24	0.53
1:R:223:PHE:O	1:R:227:MET:HG2	2.08	0.53
1:S:236:LEU:HD22	1:S:251:GLY:C	2.34	0.53
1:U:116:ARG:HH12	1:U:375:PHE:HA	1.74	0.53
1:U:236:LEU:HD22	1:U:251:GLY:C	2.34	0.53
1:V:227:MET:HA	1:V:227:MET:HE3	1.91	0.53
1:A:7:ALA:CB	1:A:356:TRP:CZ2	2.91	0.53
1:B:38:PRO:HB2	1:B:41:GLN:HB2	1.90	0.53
1:B:135:ALA:HB1	1:B:140:LEU:HD21	1.91	0.53
1:B:223:PHE:O	1:B:227:MET:HG2	2.08	0.53
1:G:140:LEU:HD22	1:G:343:GLY:HA2	1.90	0.53
1:H:360:GLN:O	1:H:364:GLU:HG3	2.08	0.53
1:L:135:ALA:HB1	1:L:140:LEU:HD21	1.91	0.53
1:M:173:HIS:NE2	1:N:267:ILE:C	2.67	0.53
1:N:147:ARG:NH2	1:N:147:ARG:HG3	2.22	0.53
1:O:135:ALA:HB1	1:O:140:LEU:HD21	1.91	0.53
1:O:143:TYR:CE2	1:Q:45:VAL:HG21	2.44	0.53
1:O:236:LEU:HD22	1:O:251:GLY:C	2.34	0.53
1:O:362:TYR:HE1	1:O:367:PRO:CB	2.19	0.53
1:Q:236:LEU:HD22	1:Q:251:GLY:C	2.34	0.53
1:S:208:ILE:HG22	1:S:209:VAL:N	2.23	0.53
1:S:223:PHE:CZ	1:S:266:PHE:HZ	2.26	0.53
1:T:116:ARG:HH12	1:T:375:PHE:HA	1.74	0.53
1:T:173:HIS:NE2	1:U:267:ILE:C	2.67	0.53
1:T:270:GLU:O	1:T:270:GLU:CD	2.51	0.53
1:U:7:ALA:CB	1:U:356:TRP:CZ2	2.91	0.53
1:V:336:LYS:HE2	2:V:401:ADP:C5'	2.38	0.53
1:A:223:PHE:O	1:A:227:MET:HG2	2.08	0.53
1:B:34:ILE:CG2	1:B:68:LYS:H	2.21	0.53
1:C:116:ARG:HH12	1:C:375:PHE:HA	1.74	0.53
1:C:147:ARG:NH2	1:C:147:ARG:HG3	2.22	0.53
1:C:173:HIS:NE2	1:D:267:ILE:C	2.67	0.53
1:D:8:LEU:HB2	1:D:103:THR:OG1	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:32:PRO:CG	1:E:55:GLY:O	2.55	0.53
1:E:116:ARG:HH12	1:E:375:PHE:HA	1.74	0.53
1:E:227:MET:HA	1:E:227:MET:HE3	1.91	0.53
1:F:8:LEU:HB2	1:F:103:THR:OG1	2.08	0.53
1:F:140:LEU:HD22	1:F:343:GLY:HA2	1.90	0.53
1:G:32:PRO:CG	1:G:55:GLY:O	2.55	0.53
1:G:37:ARG:CG	1:G:38:PRO:CD	2.83	0.53
1:G:116:ARG:HH12	1:G:375:PHE:HA	1.74	0.53
1:H:34:ILE:CG2	1:H:68:LYS:H	2.21	0.53
1:H:140:LEU:HD22	1:H:343:GLY:HA2	1.90	0.53
1:H:173:HIS:NE2	1:I:267:ILE:C	2.67	0.53
1:H:208:ILE:HG22	1:H:209:VAL:N	2.24	0.53
1:I:143:TYR:HD2	1:I:346:LEU:HD13	1.70	0.53
1:J:223:PHE:O	1:J:227:MET:HG2	2.09	0.53
1:K:223:PHE:O	1:K:227:MET:HG2	2.08	0.53
1:O:34:ILE:CG2	1:O:68:LYS:H	2.21	0.53
1:O:110:LEU:HD12	1:O:177:ARG:HH11	1.72	0.53
1:Q:143:TYR:CE2	1:S:45:VAL:HG21	2.44	0.53
1:Q:336:LYS:HE2	2:Q:401:ADP:C5'	2.38	0.53
1:S:336:LYS:HE2	2:S:401:ADP:C5'	2.38	0.53
1:T:140:LEU:HD22	1:T:343:GLY:HA2	1.90	0.53
1:U:142:LEU:HD21	1:U:165:ILE:CD1	2.38	0.53
1:V:32:PRO:CG	1:V:55:GLY:O	2.55	0.53
1:V:147:ARG:NH2	1:V:147:ARG:HG3	2.22	0.53
1:W:236:LEU:HD22	1:W:251:GLY:C	2.34	0.53
1:B:142:LEU:HD21	1:B:165:ILE:CD1	2.38	0.53
1:C:7:ALA:CB	1:C:356:TRP:CZ2	2.91	0.53
1:D:38:PRO:HB2	1:D:41:GLN:HB2	1.89	0.53
1:D:236:LEU:HD22	1:D:251:GLY:C	2.34	0.53
1:E:34:ILE:CG2	1:E:67:LEU:HD22	2.28	0.53
1:F:173:HIS:NE2	1:G:267:ILE:C	2.67	0.53
1:F:236:LEU:HD22	1:F:251:GLY:C	2.34	0.53
1:G:147:ARG:HG3	1:G:147:ARG:NH2	2.22	0.53
1:G:173:HIS:NE2	1:H:267:ILE:C	2.67	0.53
1:H:143:TYR:CE2	1:J:45:VAL:HG21	2.44	0.53
1:I:142:LEU:HD21	1:I:165:ILE:CD1	2.38	0.53
1:I:223:PHE:CZ	1:I:266:PHE:HZ	2.26	0.53
1:I:368:SER:O	1:I:371:HIS:N	2.37	0.53
1:M:143:TYR:CE2	1:O:45:VAL:HG21	2.44	0.53
1:M:236:LEU:HD22	1:M:251:GLY:C	2.34	0.53
1:M:270:GLU:O	1:M:270:GLU:CD	2.52	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:116:ARG:HH12	1:O:375:PHE:HA	1.74	0.53
1:O:223:PHE:O	1:O:227:MET:HG2	2.08	0.53
1:O:360:GLN:O	1:O:364:GLU:HG3	2.08	0.53
1:Q:38:PRO:HB2	1:Q:41:GLN:HB2	1.89	0.53
1:R:142:LEU:HD21	1:R:165:ILE:CD1	2.39	0.53
1:T:227:MET:HA	1:T:227:MET:HE3	1.91	0.53
1:W:116:ARG:HH12	1:W:375:PHE:HA	1.74	0.53
1:B:143:TYR:CE2	1:D:45:VAL:HG21	2.44	0.53
1:C:208:ILE:HG22	1:C:209:VAL:N	2.23	0.53
1:C:223:PHE:O	1:C:227:MET:HG2	2.08	0.53
1:F:34:ILE:CG2	1:F:68:LYS:H	2.21	0.53
1:G:227:MET:HA	1:G:227:MET:HE3	1.91	0.53
1:H:236:LEU:HD22	1:H:251:GLY:C	2.34	0.53
1:I:208:ILE:HG22	1:I:209:VAL:N	2.24	0.53
1:I:227:MET:HA	1:I:227:MET:HE3	1.91	0.53
1:J:143:TYR:CE2	1:L:45:VAL:HG21	2.44	0.53
1:J:360:GLN:O	1:J:364:GLU:HG3	2.08	0.53
1:K:270:GLU:O	1:K:270:GLU:CD	2.51	0.53
1:K:362:TYR:HE1	1:K:367:PRO:CB	2.19	0.53
1:N:32:PRO:CG	1:N:55:GLY:O	2.55	0.53
1:N:360:GLN:O	1:N:364:GLU:HG3	2.08	0.53
1:P:106:THR:HG22	1:P:140:LEU:CD1	2.38	0.53
1:S:270:GLU:O	1:S:270:GLU:CD	2.52	0.53
1:U:8:LEU:HB2	1:U:103:THR:OG1	2.08	0.53
1:U:54:VAL:CG1	1:U:55:GLY:N	2.71	0.53
1:U:173:HIS:NE2	1:V:267:ILE:C	2.67	0.53
1:A:116:ARG:HH12	1:A:375:PHE:HA	1.74	0.52
1:D:34:ILE:CG2	1:D:68:LYS:H	2.21	0.52
1:D:169:TYR:CZ	1:D:172:PRO:HD3	2.45	0.52
1:F:38:PRO:HB2	1:F:41:GLN:HB2	1.89	0.52
1:G:34:ILE:CG2	1:G:68:LYS:H	2.21	0.52
1:G:143:TYR:CE2	1:I:45:VAL:HG21	2.44	0.52
1:G:336:LYS:HE2	2:G:401:ADP:C5'	2.38	0.52
1:J:173:HIS:NE2	1:K:267:ILE:C	2.67	0.52
1:M:169:TYR:CZ	1:M:172:PRO:HD3	2.45	0.52
1:N:37:ARG:CG	1:N:38:PRO:CD	2.83	0.52
1:N:58:ALA:O	1:N:59:GLN:C	2.53	0.52
1:N:147:ARG:CG	1:N:147:ARG:NH2	2.71	0.52
1:O:140:LEU:HD22	1:O:343:GLY:HA2	1.90	0.52
1:O:336:LYS:HE2	2:O:401:ADP:C5'	2.38	0.52
1:P:32:PRO:CG	1:P:55:GLY:O	2.55	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:336:LYS:HE2	2:U:401:ADP:C5'	2.38	0.52
1:V:169:TYR:CZ	1:V:172:PRO:HD3	2.45	0.52
1:V:362:TYR:HD1	1:V:367:PRO:HA	1.74	0.52
1:W:8:LEU:HB2	1:W:103:THR:OG1	2.08	0.52
1:W:34:ILE:CG2	1:W:68:LYS:H	2.21	0.52
1:E:34:ILE:CG2	1:E:68:LYS:H	2.21	0.52
1:E:135:ALA:HB1	1:E:140:LEU:HD21	1.91	0.52
1:G:169:TYR:CZ	1:G:172:PRO:HD3	2.45	0.52
1:H:362:TYR:HD1	1:H:367:PRO:HA	1.75	0.52
1:H:368:SER:O	1:H:371:HIS:N	2.37	0.52
1:J:34:ILE:CG2	1:J:68:LYS:H	2.21	0.52
1:J:37:ARG:CG	1:J:38:PRO:CD	2.83	0.52
1:J:58:ALA:O	1:J:59:GLN:C	2.53	0.52
1:K:32:PRO:CG	1:K:55:GLY:O	2.55	0.52
1:K:37:ARG:CG	1:K:38:PRO:CD	2.83	0.52
1:K:116:ARG:HH12	1:K:375:PHE:HA	1.74	0.52
1:K:169:TYR:CZ	1:K:172:PRO:HD3	2.45	0.52
1:L:32:PRO:CG	1:L:55:GLY:O	2.55	0.52
1:N:38:PRO:HB2	1:N:41:GLN:HB2	1.90	0.52
1:P:142:LEU:HD21	1:P:165:ILE:CD1	2.39	0.52
1:R:116:ARG:HH12	1:R:375:PHE:HA	1.74	0.52
1:R:208:ILE:HG22	1:R:209:VAL:N	2.23	0.52
1:R:227:MET:HA	1:R:227:MET:HE3	1.91	0.52
1:S:169:TYR:CZ	1:S:172:PRO:HD3	2.45	0.52
1:T:135:ALA:HB1	1:T:140:LEU:HD21	1.91	0.52
1:V:270:GLU:O	1:V:270:GLU:CD	2.52	0.52
1:A:173:HIS:NE2	1:B:267:ILE:C	2.67	0.52
1:C:58:ALA:O	1:C:59:GLN:C	2.53	0.52
1:C:362:TYR:HE1	1:C:367:PRO:CB	2.19	0.52
1:E:360:GLN:O	1:E:364:GLU:HG3	2.08	0.52
1:E:368:SER:O	1:E:371:HIS:N	2.37	0.52
1:F:143:TYR:CE2	1:H:45:VAL:HG21	2.44	0.52
1:G:135:ALA:HB1	1:G:140:LEU:HD21	1.90	0.52
1:H:362:TYR:HE1	1:H:367:PRO:CB	2.19	0.52
1:J:361:GLU:HB3	1:J:369:ILE:CD1	2.14	0.52
1:K:227:MET:HA	1:K:227:MET:HE3	1.91	0.52
1:M:135:ALA:HB1	1:M:140:LEU:HD21	1.91	0.52
1:M:227:MET:HA	1:M:227:MET:HE3	1.91	0.52
1:P:34:ILE:CG2	1:P:68:LYS:H	2.21	0.52
1:P:173:HIS:NE2	1:Q:267:ILE:C	2.67	0.52
1:P:208:ILE:HG22	1:P:209:VAL:N	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:362:TYR:HD1	1:P:367:PRO:HA	1.75	0.52
1:Q:58:ALA:O	1:Q:59:GLN:C	2.53	0.52
1:Q:110:LEU:HD12	1:Q:177:ARG:HH11	1.72	0.52
1:R:135:ALA:HB1	1:R:140:LEU:HD21	1.91	0.52
1:S:8:LEU:HB2	1:S:103:THR:OG1	2.08	0.52
1:U:58:ALA:O	1:U:59:GLN:C	2.53	0.52
1:V:37:ARG:CG	1:V:38:PRO:CD	2.83	0.52
1:V:140:LEU:HD22	1:V:343:GLY:HA2	1.90	0.52
1:W:368:SER:O	1:W:371:HIS:N	2.37	0.52
1:A:34:ILE:CG2	1:A:68:LYS:H	2.21	0.52
1:A:208:ILE:HG22	1:A:209:VAL:N	2.24	0.52
1:A:368:SER:O	1:A:371:HIS:N	2.37	0.52
1:B:140:LEU:HD22	1:B:343:GLY:HA2	1.90	0.52
1:B:169:TYR:CZ	1:B:172:PRO:HD3	2.45	0.52
1:B:236:LEU:HD22	1:B:251:GLY:C	2.34	0.52
1:C:227:MET:HA	1:C:227:MET:HE3	1.91	0.52
1:C:236:LEU:HD22	1:C:251:GLY:C	2.34	0.52
1:E:37:ARG:CG	1:E:38:PRO:CD	2.83	0.52
1:E:54:VAL:CG1	1:E:55:GLY:N	2.71	0.52
1:E:169:TYR:CZ	1:E:172:PRO:HD3	2.45	0.52
1:F:58:ALA:O	1:F:59:GLN:C	2.53	0.52
1:F:116:ARG:HH12	1:F:375:PHE:HA	1.74	0.52
1:F:169:TYR:CZ	1:F:172:PRO:HD3	2.45	0.52
1:F:362:TYR:HD1	1:F:367:PRO:HA	1.75	0.52
1:G:208:ILE:HG22	1:G:209:VAL:N	2.23	0.52
1:H:38:PRO:HB2	1:H:41:GLN:HB2	1.90	0.52
1:I:135:ALA:HB1	1:I:140:LEU:HD21	1.91	0.52
1:K:140:LEU:HD22	1:K:343:GLY:HA2	1.90	0.52
1:K:147:ARG:NH2	1:K:147:ARG:HG3	2.22	0.52
1:L:169:TYR:CZ	1:L:172:PRO:HD3	2.44	0.52
1:L:173:HIS:NE2	1:M:267:ILE:C	2.67	0.52
1:M:140:LEU:HD22	1:M:343:GLY:HA2	1.90	0.52
1:M:336:LYS:HE2	2:M:401:ADP:C5'	2.38	0.52
1:M:360:GLN:O	1:M:364:GLU:HG3	2.08	0.52
1:N:173:HIS:NE2	1:O:267:ILE:C	2.67	0.52
1:O:58:ALA:O	1:O:59:GLN:C	2.53	0.52
1:O:169:TYR:CZ	1:O:172:PRO:HD3	2.45	0.52
1:O:227:MET:HA	1:O:227:MET:HE3	1.91	0.52
1:P:143:TYR:CE2	1:R:45:VAL:HG21	2.44	0.52
1:Q:140:LEU:HD22	1:Q:343:GLY:HA2	1.90	0.52
1:Q:360:GLN:O	1:Q:364:GLU:HG3	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:147:ARG:NH2	1:R:147:ARG:HG3	2.22	0.52
1:S:140:LEU:HD22	1:S:343:GLY:HA2	1.90	0.52
1:S:143:TYR:CE2	1:U:45:VAL:HG21	2.44	0.52
1:T:169:TYR:CZ	1:T:172:PRO:HD3	2.45	0.52
1:T:236:LEU:HD22	1:T:251:GLY:C	2.34	0.52
1:U:34:ILE:CG2	1:U:68:LYS:H	2.21	0.52
1:W:164:PRO:HG3	1:W:174:ALA:HB1	1.92	0.52
1:W:223:PHE:O	1:W:227:MET:HG2	2.09	0.52
1:A:362:TYR:HD1	1:A:367:PRO:HA	1.75	0.52
1:B:58:ALA:O	1:B:59:GLN:C	2.53	0.52
1:B:227:MET:HA	1:B:227:MET:HE3	1.91	0.52
1:C:34:ILE:CG2	1:C:68:LYS:H	2.21	0.52
1:C:236:LEU:HD12	1:C:237:GLU:HG2	1.89	0.52
1:C:362:TYR:HD1	1:C:367:PRO:HA	1.75	0.52
1:D:58:ALA:O	1:D:59:GLN:C	2.53	0.52
1:E:236:LEU:HD22	1:E:251:GLY:C	2.34	0.52
1:E:362:TYR:HD1	1:E:367:PRO:HA	1.75	0.52
1:H:116:ARG:HH12	1:H:375:PHE:HA	1.74	0.52
1:I:140:LEU:HD22	1:I:343:GLY:HA2	1.90	0.52
1:I:143:TYR:CE2	1:K:45:VAL:HG21	2.44	0.52
1:J:362:TYR:HD1	1:J:367:PRO:HA	1.75	0.52
1:L:38:PRO:HB2	1:L:41:GLN:HB2	1.90	0.52
1:O:362:TYR:HD1	1:O:367:PRO:HA	1.75	0.52
1:Q:34:ILE:CG2	1:Q:68:LYS:H	2.21	0.52
1:R:58:ALA:O	1:R:59:GLN:C	2.53	0.52
1:R:362:TYR:HE1	1:R:367:PRO:CB	2.19	0.52
1:T:360:GLN:O	1:T:364:GLU:HG3	2.08	0.52
1:U:37:ARG:CG	1:U:38:PRO:CD	2.83	0.52
1:U:164:PRO:HG3	1:U:174:ALA:HB1	1.92	0.52
1:U:169:TYR:CZ	1:U:172:PRO:HD3	2.45	0.52
1:U:208:ILE:HG22	1:U:209:VAL:N	2.24	0.52
1:V:135:ALA:HB1	1:V:140:LEU:HD21	1.91	0.52
1:W:300:SER:HA	1:W:335:ARG:HG2	1.89	0.52
1:A:143:TYR:CE2	1:C:45:VAL:HG21	2.44	0.52
1:A:236:LEU:HD22	1:A:251:GLY:C	2.34	0.52
1:D:336:LYS:HE2	2:D:401:ADP:C5'	2.38	0.52
1:E:208:ILE:HG22	1:E:209:VAL:N	2.23	0.52
1:J:38:PRO:HB2	1:J:41:GLN:HB2	1.89	0.52
1:J:169:TYR:CZ	1:J:172:PRO:HD3	2.45	0.52
1:K:58:ALA:O	1:K:59:GLN:C	2.53	0.52
1:K:236:LEU:HD22	1:K:251:GLY:C	2.34	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:34:ILE:CG2	1:M:67:LEU:HD22	2.28	0.52
1:M:58:ALA:O	1:M:59:GLN:C	2.53	0.52
1:P:300:SER:HA	1:P:335:ARG:HG2	1.89	0.52
1:Q:362:TYR:HD1	1:Q:367:PRO:HA	1.75	0.52
1:R:236:LEU:HD22	1:R:251:GLY:C	2.34	0.52
1:S:142:LEU:HD21	1:S:165:ILE:CD1	2.38	0.52
1:T:143:TYR:CE2	1:V:45:VAL:HG21	2.44	0.52
1:U:360:GLN:O	1:U:364:GLU:HG3	2.08	0.52
1:V:236:LEU:HD22	1:V:251:GLY:C	2.34	0.52
1:W:360:GLN:O	1:W:364:GLU:HG3	2.08	0.52
1:A:164:PRO:HG3	1:A:174:ALA:HB1	1.92	0.52
1:D:140:LEU:HD22	1:D:343:GLY:HA2	1.90	0.52
1:G:58:ALA:O	1:G:59:GLN:C	2.53	0.52
1:H:223:PHE:O	1:H:227:MET:HG2	2.08	0.52
1:I:169:TYR:CZ	1:I:172:PRO:HD3	2.45	0.52
1:J:208:ILE:HG22	1:J:209:VAL:N	2.23	0.52
1:J:236:LEU:HD22	1:J:251:GLY:C	2.34	0.52
1:K:135:ALA:HB1	1:K:140:LEU:HD21	1.91	0.52
1:K:164:PRO:HG2	1:K:174:ALA:HB3	1.92	0.52
1:L:360:GLN:O	1:L:364:GLU:HG3	2.08	0.52
1:N:142:LEU:HD21	1:N:165:ILE:CD1	2.39	0.52
1:N:236:LEU:HD12	1:N:237:GLU:HG2	1.90	0.52
1:O:361:GLU:HB3	1:O:369:ILE:CD1	2.14	0.52
1:R:173:HIS:NE2	1:S:267:ILE:C	2.67	0.52
1:S:360:GLN:O	1:S:364:GLU:HG3	2.08	0.52
1:T:362:TYR:HD1	1:T:367:PRO:HA	1.75	0.52
1:U:143:TYR:CE2	1:W:45:VAL:HG21	2.44	0.52
1:V:34:ILE:CG2	1:V:67:LEU:HD22	2.27	0.52
1:V:58:ALA:O	1:V:59:GLN:C	2.53	0.52
1:V:164:PRO:HG2	1:V:174:ALA:HB3	1.92	0.52
1:V:208:ILE:HG22	1:V:209:VAL:N	2.23	0.52
1:A:227:MET:HA	1:A:227:MET:HE3	1.91	0.52
1:A:361:GLU:HB3	1:A:369:ILE:CD1	2.14	0.52
1:D:143:TYR:CE2	1:F:45:VAL:HG21	2.44	0.52
1:K:336:LYS:HE2	2:K:401:ADP:C5'	2.38	0.52
1:L:147:ARG:CG	1:L:147:ARG:NH2	2.71	0.52
1:L:208:ILE:HG22	1:L:209:VAL:N	2.23	0.52
1:M:116:ARG:HH12	1:M:375:PHE:HA	1.74	0.52
1:M:362:TYR:HD1	1:M:367:PRO:HA	1.75	0.52
1:N:143:TYR:CE2	1:P:45:VAL:HG21	2.44	0.52
1:N:169:TYR:CZ	1:N:172:PRO:HD3	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:227:MET:HE3	1:N:227:MET:HA	1.91	0.52
1:P:58:ALA:O	1:P:59:GLN:C	2.53	0.52
1:P:368:SER:O	1:P:371:HIS:N	2.37	0.52
1:Q:169:TYR:CZ	1:Q:172:PRO:HD3	2.45	0.52
1:U:300:SER:HA	1:U:335:ARG:HG2	1.89	0.52
1:E:143:TYR:CE2	1:G:45:VAL:HG21	2.44	0.52
1:G:142:LEU:HD21	1:G:165:ILE:CD1	2.39	0.52
1:G:236:LEU:HD22	1:G:251:GLY:C	2.34	0.52
1:I:164:PRO:HG2	1:I:174:ALA:HB3	1.92	0.52
1:I:362:TYR:HD1	1:I:367:PRO:HA	1.74	0.52
1:K:143:TYR:CE2	1:M:45:VAL:HG21	2.44	0.52
1:L:34:ILE:CG2	1:L:68:LYS:H	2.21	0.52
1:L:143:TYR:CE2	1:N:45:VAL:HG21	2.44	0.52
1:M:167:GLU:OE1	1:O:61:LYS:CE	2.58	0.52
1:O:164:PRO:HG2	1:O:174:ALA:HB3	1.92	0.52
1:O:167:GLU:OE1	1:Q:61:LYS:CE	2.58	0.52
1:P:116:ARG:HH12	1:P:375:PHE:HA	1.74	0.52
1:P:227:MET:HA	1:P:227:MET:HE3	1.91	0.52
1:S:58:ALA:O	1:S:59:GLN:C	2.53	0.52
1:T:167:GLU:OE1	1:V:61:LYS:CE	2.58	0.52
1:T:208:ILE:HG22	1:T:209:VAL:N	2.24	0.52
1:A:58:ALA:O	1:A:59:GLN:C	2.53	0.52
1:B:336:LYS:HE2	2:B:401:ADP:C5'	2.38	0.52
1:D:362:TYR:HE1	1:D:367:PRO:CB	2.19	0.52
1:E:36:GLY:HA3	1:E:66:THR:O	2.10	0.52
1:I:167:GLU:OE1	1:K:61:LYS:CE	2.58	0.52
1:I:236:LEU:HD22	1:I:251:GLY:C	2.34	0.52
1:J:116:ARG:HH12	1:J:375:PHE:HA	1.74	0.52
1:J:230:ALA:HA	1:J:233:SER:HB2	1.92	0.52
1:P:36:GLY:HA3	1:P:66:THR:O	2.10	0.52
1:Q:167:GLU:OE1	1:S:61:LYS:CE	2.58	0.52
1:Q:227:MET:HE3	1:Q:227:MET:HA	1.91	0.52
1:R:167:GLU:OE1	1:T:61:LYS:CE	2.58	0.52
1:S:36:GLY:HA3	1:S:66:THR:O	2.10	0.52
1:T:58:ALA:O	1:T:59:GLN:C	2.53	0.52
1:A:36:GLY:HA3	1:A:66:THR:O	2.10	0.51
1:A:169:TYR:CZ	1:A:172:PRO:HD3	2.45	0.51
1:B:36:GLY:HA3	1:B:66:THR:O	2.10	0.51
1:C:143:TYR:CE2	1:E:45:VAL:HG21	2.44	0.51
1:D:362:TYR:HD1	1:D:367:PRO:HA	1.75	0.51
1:E:58:ALA:O	1:E:59:GLN:C	2.53	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:164:PRO:HG3	1:F:174:ALA:HB1	1.92	0.51
1:G:362:TYR:HD1	1:G:367:PRO:HA	1.75	0.51
1:I:58:ALA:O	1:I:59:GLN:C	2.53	0.51
1:J:147:ARG:CG	1:J:147:ARG:NH2	2.71	0.51
1:K:167:GLU:OE1	1:M:61:LYS:CE	2.58	0.51
1:L:167:GLU:OE1	1:N:61:LYS:CE	2.58	0.51
1:L:230:ALA:HA	1:L:233:SER:HB2	1.92	0.51
1:L:362:TYR:HD1	1:L:367:PRO:HA	1.75	0.51
1:L:368:SER:O	1:L:371:HIS:N	2.37	0.51
1:M:164:PRO:HG2	1:M:174:ALA:HB3	1.92	0.51
1:P:164:PRO:HG3	1:P:174:ALA:HB1	1.92	0.51
1:S:34:ILE:CG2	1:S:68:LYS:H	2.21	0.51
1:S:164:PRO:HG3	1:S:174:ALA:HB1	1.92	0.51
1:B:22:ALA:HB1	1:B:348:SER:CB	2.41	0.51
1:B:164:PRO:HG2	1:B:174:ALA:HB3	1.92	0.51
1:B:167:GLU:OE1	1:D:61:LYS:CE	2.58	0.51
1:C:164:PRO:HG3	1:C:174:ALA:HB1	1.92	0.51
1:D:164:PRO:HG3	1:D:174:ALA:HB1	1.92	0.51
1:F:361:GLU:HB3	1:F:369:ILE:CD1	2.14	0.51
1:H:58:ALA:O	1:H:59:GLN:C	2.53	0.51
1:I:58:ALA:CA	1:I:65:LEU:CD2	2.89	0.51
1:I:336:LYS:HE2	2:I:401:ADP:C5'	2.38	0.51
1:L:22:ALA:HB1	1:L:348:SER:CB	2.41	0.51
1:L:58:ALA:O	1:L:59:GLN:C	2.53	0.51
1:L:116:ARG:HH12	1:L:375:PHE:HA	1.74	0.51
1:N:34:ILE:CG2	1:N:68:LYS:H	2.21	0.51
1:N:362:TYR:HE1	1:N:367:PRO:CB	2.19	0.51
1:N:362:TYR:HD1	1:N:367:PRO:HA	1.75	0.51
1:O:58:ALA:CA	1:O:65:LEU:CD2	2.89	0.51
1:P:236:LEU:HD22	1:P:251:GLY:C	2.34	0.51
1:Q:22:ALA:HB1	1:Q:348:SER:CB	2.41	0.51
1:Q:36:GLY:HA3	1:Q:66:THR:O	2.10	0.51
1:Q:58:ALA:CA	1:Q:65:LEU:CD2	2.89	0.51
1:S:300:SER:HA	1:S:335:ARG:HG2	1.89	0.51
1:T:164:PRO:HG2	1:T:174:ALA:HB3	1.92	0.51
1:T:362:TYR:HE1	1:T:367:PRO:CB	2.19	0.51
1:W:32:PRO:CG	1:W:55:GLY:O	2.55	0.51
1:W:230:ALA:HA	1:W:233:SER:HB2	1.92	0.51
1:A:332:PRO:HG2	1:A:335:ARG:NE	2.26	0.51
1:B:58:ALA:CA	1:B:65:LEU:CD2	2.89	0.51
1:C:169:TYR:CZ	1:C:172:PRO:HD3	2.45	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:332:PRO:HG2	1:C:335:ARG:NE	2.26	0.51
1:D:39:ARG:NE	1:D:66:THR:CA	2.69	0.51
1:E:22:ALA:HB1	1:E:348:SER:CB	2.41	0.51
1:E:164:PRO:HG2	1:E:174:ALA:HB3	1.92	0.51
1:G:39:ARG:HG2	1:G:66:THR:HB	1.93	0.51
1:G:164:PRO:HG2	1:G:174:ALA:HB3	1.92	0.51
1:G:362:TYR:HE1	1:G:367:PRO:CB	2.19	0.51
1:H:164:PRO:HG3	1:H:174:ALA:HB1	1.92	0.51
1:H:230:ALA:HA	1:H:233:SER:HB2	1.92	0.51
1:I:34:ILE:CG2	1:I:35:VAL:N	2.74	0.51
1:J:32:PRO:CG	1:J:55:GLY:O	2.55	0.51
1:K:39:ARG:HG2	1:K:66:THR:HB	1.93	0.51
1:K:58:ALA:CA	1:K:65:LEU:CD2	2.89	0.51
1:L:227:MET:HA	1:L:227:MET:HE3	1.91	0.51
1:M:36:GLY:HA3	1:M:66:THR:O	2.10	0.51
1:M:58:ALA:CA	1:M:65:LEU:CD2	2.89	0.51
1:N:36:GLY:HA3	1:N:66:THR:O	2.10	0.51
1:N:164:PRO:HG3	1:N:174:ALA:HB1	1.92	0.51
1:P:167:GLU:OE1	1:R:61:LYS:CE	2.58	0.51
1:P:169:TYR:CZ	1:P:172:PRO:HD3	2.44	0.51
1:R:143:TYR:CE2	1:T:45:VAL:HG21	2.44	0.51
1:R:164:PRO:HG2	1:R:174:ALA:HB3	1.92	0.51
1:S:332:PRO:HG2	1:S:335:ARG:NE	2.26	0.51
1:S:362:TYR:HD1	1:S:367:PRO:HA	1.75	0.51
1:W:362:TYR:HD1	1:W:367:PRO:HA	1.75	0.51
1:A:34:ILE:CG2	1:A:35:VAL:N	2.74	0.51
1:A:167:GLU:OE1	1:C:61:LYS:CE	2.58	0.51
1:D:22:ALA:HB1	1:D:348:SER:CB	2.41	0.51
1:D:36:GLY:HA3	1:D:66:THR:O	2.10	0.51
1:G:36:GLY:HA3	1:G:66:THR:O	2.10	0.51
1:G:58:ALA:CA	1:G:65:LEU:CD2	2.89	0.51
1:H:169:TYR:CZ	1:H:172:PRO:HD3	2.45	0.51
1:J:332:PRO:HG2	1:J:335:ARG:NE	2.26	0.51
1:L:34:ILE:CG2	1:L:67:LEU:HD22	2.28	0.51
1:L:54:VAL:CG1	1:L:55:GLY:N	2.72	0.51
1:L:142:LEU:HD21	1:L:165:ILE:CD1	2.39	0.51
1:N:208:ILE:HG22	1:N:209:VAL:N	2.23	0.51
1:N:230:ALA:HA	1:N:233:SER:HB2	1.92	0.51
1:O:37:ARG:CG	1:O:38:PRO:CD	2.83	0.51
1:P:362:TYR:HE1	1:P:367:PRO:CB	2.19	0.51
1:Q:164:PRO:HG2	1:Q:174:ALA:HB3	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:332:PRO:HG2	1:Q:335:ARG:NE	2.26	0.51
1:R:34:ILE:CG2	1:R:35:VAL:N	2.74	0.51
1:R:169:TYR:CZ	1:R:172:PRO:HD3	2.45	0.51
1:S:22:ALA:HB1	1:S:348:SER:CB	2.41	0.51
1:S:58:ALA:CA	1:S:65:LEU:CD2	2.89	0.51
1:S:167:GLU:OE1	1:U:61:LYS:CE	2.58	0.51
1:T:36:GLY:HA3	1:T:66:THR:O	2.10	0.51
1:T:37:ARG:CG	1:T:38:PRO:CD	2.83	0.51
1:U:36:GLY:HA3	1:U:66:THR:O	2.10	0.51
1:U:167:GLU:OE1	1:W:61:LYS:CE	2.58	0.51
1:U:227:MET:HA	1:U:227:MET:HE3	1.91	0.51
1:U:230:ALA:HA	1:U:233:SER:HB2	1.92	0.51
1:U:361:GLU:HB3	1:U:369:ILE:CD1	2.14	0.51
1:W:34:ILE:CG2	1:W:67:LEU:HD22	2.27	0.51
1:W:227:MET:HA	1:W:227:MET:HE3	1.91	0.51
1:C:36:GLY:HA3	1:C:66:THR:O	2.10	0.51
1:C:361:GLU:HB3	1:C:369:ILE:CD1	2.14	0.51
1:D:58:ALA:CA	1:D:65:LEU:CD2	2.89	0.51
1:D:164:PRO:HG2	1:D:174:ALA:HB3	1.92	0.51
1:G:34:ILE:CG2	1:G:35:VAL:N	2.74	0.51
1:J:167:GLU:OE1	1:L:61:LYS:CE	2.58	0.51
1:K:22:ALA:HB1	1:K:348:SER:CB	2.41	0.51
1:N:22:ALA:HB1	1:N:348:SER:CB	2.41	0.51
1:N:169:TYR:OH	1:P:40:HIS:HB3	2.10	0.51
1:P:34:ILE:CG2	1:P:35:VAL:N	2.74	0.51
1:Q:37:ARG:CG	1:Q:38:PRO:CD	2.83	0.51
1:R:164:PRO:HG3	1:R:174:ALA:HB1	1.92	0.51
1:T:22:ALA:HB1	1:T:348:SER:CB	2.41	0.51
1:T:39:ARG:HG2	1:T:66:THR:HB	1.93	0.51
1:V:58:ALA:CA	1:V:65:LEU:CD2	2.89	0.51
1:W:58:ALA:O	1:W:59:GLN:C	2.53	0.51
1:W:169:TYR:CZ	1:W:172:PRO:HD3	2.45	0.51
1:B:164:PRO:HG3	1:B:174:ALA:HB1	1.92	0.51
1:C:34:ILE:CG2	1:C:67:LEU:HD22	2.28	0.51
1:D:167:GLU:OE1	1:F:61:LYS:CE	2.58	0.51
1:F:36:GLY:HA3	1:F:66:THR:O	2.10	0.51
1:F:37:ARG:CG	1:F:38:PRO:CD	2.83	0.51
1:F:58:ALA:CA	1:F:65:LEU:CD2	2.89	0.51
1:F:227:MET:HE3	1:F:227:MET:HA	1.91	0.51
1:G:34:ILE:HG23	1:G:68:LYS:N	2.26	0.51
1:G:167:GLU:OE1	1:I:61:LYS:CE	2.58	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:332:PRO:HG2	1:H:335:ARG:NE	2.26	0.51
1:J:22:ALA:HB1	1:J:348:SER:CB	2.41	0.51
1:L:36:GLY:HA3	1:L:66:THR:O	2.10	0.51
1:N:116:ARG:HH12	1:N:375:PHE:HA	1.74	0.51
1:N:167:GLU:OE1	1:P:61:LYS:CE	2.58	0.51
1:O:332:PRO:HG2	1:O:335:ARG:NE	2.26	0.51
1:P:164:PRO:HG2	1:P:174:ALA:HB3	1.92	0.51
1:P:169:TYR:OH	1:R:40:HIS:HB3	2.10	0.51
1:S:227:MET:HA	1:S:227:MET:HE3	1.91	0.51
1:S:230:ALA:HA	1:S:233:SER:HB2	1.92	0.51
1:T:332:PRO:HG2	1:T:335:ARG:NE	2.26	0.51
1:U:34:ILE:HG23	1:U:68:LYS:N	2.26	0.51
1:U:332:PRO:HG2	1:U:335:ARG:NE	2.26	0.51
1:W:208:ILE:HG22	1:W:209:VAL:N	2.23	0.51
1:A:169:TYR:OH	1:C:40:HIS:HB3	2.10	0.51
1:B:34:ILE:HG23	1:B:68:LYS:N	2.26	0.51
1:B:169:TYR:OH	1:D:40:HIS:HB3	2.10	0.51
1:C:34:ILE:CG2	1:C:35:VAL:N	2.74	0.51
1:C:34:ILE:HG12	1:C:69:TYR:CE1	2.46	0.51
1:C:164:PRO:HG2	1:C:174:ALA:HB3	1.92	0.51
1:D:227:MET:HA	1:D:227:MET:HE3	1.91	0.51
1:D:230:ALA:HA	1:D:233:SER:HB2	1.92	0.51
1:E:34:ILE:HG23	1:E:68:LYS:N	2.26	0.51
1:E:58:ALA:CA	1:E:65:LEU:CD2	2.89	0.51
1:E:332:PRO:HG2	1:E:335:ARG:NE	2.26	0.51
1:F:34:ILE:CG2	1:F:67:LEU:HD22	2.28	0.51
1:F:167:GLU:OE1	1:H:61:LYS:CE	2.58	0.51
1:I:34:ILE:HG23	1:I:68:LYS:N	2.26	0.51
1:K:362:TYR:HD1	1:K:367:PRO:HA	1.75	0.51
1:L:332:PRO:HG2	1:L:335:ARG:NE	2.26	0.51
1:O:36:GLY:HA3	1:O:66:THR:O	2.10	0.51
1:Q:230:ALA:HA	1:Q:233:SER:HB2	1.92	0.51
1:Q:300:SER:HA	1:Q:335:ARG:HG2	1.89	0.51
1:R:362:TYR:HD1	1:R:367:PRO:HA	1.75	0.51
1:T:34:ILE:HG12	1:T:69:TYR:CE1	2.46	0.51
1:T:58:ALA:CA	1:T:65:LEU:CD2	2.89	0.51
1:T:164:PRO:HG3	1:T:174:ALA:HB1	1.92	0.51
1:U:58:ALA:CA	1:U:65:LEU:CD2	2.89	0.51
1:V:332:PRO:HG2	1:V:335:ARG:NE	2.26	0.51
1:A:34:ILE:HG23	1:A:68:LYS:N	2.26	0.51
1:A:230:ALA:HA	1:A:233:SER:HB2	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:230:ALA:HA	1:C:233:SER:HB2	1.92	0.51
1:D:34:ILE:HG23	1:D:68:LYS:N	2.26	0.51
1:D:34:ILE:HG12	1:D:69:TYR:CE1	2.46	0.51
1:F:230:ALA:HA	1:F:233:SER:HB2	1.92	0.51
1:H:2:GLU:OE2	1:H:2:GLU:HA	2.10	0.51
1:H:34:ILE:CG2	1:H:67:LEU:HD22	2.28	0.51
1:H:36:GLY:HA3	1:H:66:THR:O	2.10	0.51
1:H:58:ALA:CA	1:H:65:LEU:CD2	2.89	0.51
1:H:227:MET:HA	1:H:227:MET:HE3	1.91	0.51
1:J:164:PRO:HG3	1:J:174:ALA:HB1	1.92	0.51
1:K:34:ILE:HG12	1:K:69:TYR:CE1	2.46	0.51
1:L:164:PRO:HG3	1:L:174:ALA:HB1	1.92	0.51
1:L:169:TYR:OH	1:N:40:HIS:HB3	2.10	0.51
1:L:236:LEU:HD22	1:L:251:GLY:C	2.34	0.51
1:M:2:GLU:OE2	1:M:2:GLU:HA	2.10	0.51
1:O:34:ILE:HG12	1:O:69:TYR:CE1	2.46	0.51
1:O:39:ARG:HG2	1:O:66:THR:HB	1.93	0.51
1:P:347:ALA:HA	1:P:356:TRP:CZ2	2.46	0.51
1:Q:164:PRO:HG3	1:Q:174:ALA:HB1	1.92	0.51
1:W:2:GLU:OE2	1:W:2:GLU:HA	2.10	0.51
1:B:39:ARG:HG2	1:B:66:THR:HB	1.93	0.51
1:B:332:PRO:HG2	1:B:335:ARG:NE	2.26	0.51
1:C:34:ILE:HG23	1:C:68:LYS:N	2.26	0.51
1:C:39:ARG:HG2	1:C:66:THR:HB	1.92	0.51
1:E:164:PRO:HG3	1:E:174:ALA:HB1	1.92	0.51
1:E:167:GLU:OE1	1:G:61:LYS:CE	2.58	0.51
1:F:34:ILE:HG23	1:F:68:LYS:N	2.26	0.51
1:H:167:GLU:OE1	1:J:61:LYS:CE	2.58	0.51
1:I:164:PRO:HG3	1:I:174:ALA:HB1	1.92	0.51
1:J:34:ILE:HG23	1:J:68:LYS:N	2.26	0.51
1:J:164:PRO:HG2	1:J:174:ALA:HB3	1.92	0.51
1:J:227:MET:HA	1:J:227:MET:HE3	1.91	0.51
1:L:34:ILE:HG23	1:L:68:LYS:N	2.26	0.51
1:L:34:ILE:HG12	1:L:69:TYR:CE1	2.46	0.51
1:M:39:ARG:NE	1:M:66:THR:CA	2.69	0.51
1:O:300:SER:HA	1:O:335:ARG:HG2	1.89	0.51
1:Q:142:LEU:HD21	1:Q:165:ILE:CD1	2.39	0.51
1:R:58:ALA:CA	1:R:65:LEU:CD2	2.89	0.51
1:R:347:ALA:HA	1:R:356:TRP:CZ2	2.46	0.51
1:T:347:ALA:HA	1:T:356:TRP:CZ2	2.46	0.51
1:U:34:ILE:HG12	1:U:69:TYR:CE1	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:164:PRO:HG2	1:W:174:ALA:HB3	1.92	0.51
1:B:39:ARG:NE	1:B:66:THR:CA	2.69	0.51
1:C:54:VAL:CG1	1:C:55:GLY:N	2.72	0.51
1:D:169:TYR:OH	1:F:40:HIS:HB3	2.10	0.51
1:E:34:ILE:HG12	1:E:69:TYR:CE1	2.46	0.51
1:F:332:PRO:HG2	1:F:335:ARG:NE	2.26	0.51
1:I:36:GLY:HA3	1:I:66:THR:O	2.10	0.51
1:J:347:ALA:HA	1:J:356:TRP:CZ2	2.46	0.51
1:K:34:ILE:CG2	1:K:35:VAL:N	2.74	0.51
1:K:34:ILE:HG23	1:K:68:LYS:N	2.26	0.51
1:K:164:PRO:HG3	1:K:174:ALA:HB1	1.92	0.51
1:L:347:ALA:HA	1:L:356:TRP:CZ2	2.46	0.51
1:M:347:ALA:HA	1:M:356:TRP:CZ2	2.46	0.51
1:N:236:LEU:HD22	1:N:251:GLY:C	2.34	0.51
1:N:300:SER:HA	1:N:335:ARG:HG2	1.89	0.51
1:R:332:PRO:HG2	1:R:335:ARG:NE	2.26	0.51
1:R:336:LYS:HE2	2:R:401:ADP:C5'	2.38	0.51
1:T:34:ILE:CG2	1:T:35:VAL:N	2.74	0.51
1:U:22:ALA:HB1	1:U:348:SER:CB	2.41	0.51
1:V:36:GLY:HA3	1:V:66:THR:O	2.10	0.51
1:V:347:ALA:HA	1:V:356:TRP:CZ2	2.47	0.51
1:A:58:ALA:CA	1:A:65:LEU:CD2	2.89	0.50
1:B:230:ALA:HA	1:B:233:SER:HB2	1.92	0.50
1:E:142:LEU:HD21	1:E:165:ILE:CD1	2.39	0.50
1:F:22:ALA:HB1	1:F:348:SER:CB	2.41	0.50
1:F:147:ARG:CG	1:F:147:ARG:NH2	2.71	0.50
1:F:299:MET:HE1	1:F:309:ILE:HD13	1.94	0.50
1:G:22:ALA:HB1	1:G:348:SER:CB	2.41	0.50
1:G:332:PRO:HG2	1:G:335:ARG:NE	2.26	0.50
1:H:34:ILE:HG23	1:H:68:LYS:N	2.26	0.50
1:H:34:ILE:HG12	1:H:69:TYR:CE1	2.46	0.50
1:H:347:ALA:HA	1:H:356:TRP:CZ2	2.46	0.50
1:J:34:ILE:HG12	1:J:69:TYR:CE1	2.46	0.50
1:K:169:TYR:OH	1:M:40:HIS:HB3	2.10	0.50
1:M:22:ALA:HB1	1:M:348:SER:CB	2.41	0.50
1:M:32:PRO:CG	1:M:55:GLY:O	2.55	0.50
1:M:164:PRO:HG3	1:M:174:ALA:HB1	1.92	0.50
1:M:169:TYR:OH	1:O:40:HIS:HB3	2.10	0.50
1:N:34:ILE:HG12	1:N:69:TYR:CE1	2.46	0.50
1:N:347:ALA:HA	1:N:356:TRP:CZ2	2.46	0.50
1:O:32:PRO:CG	1:O:55:GLY:O	2.55	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:34:ILE:HG23	1:P:68:LYS:N	2.26	0.50
1:P:230:ALA:HA	1:P:233:SER:HB2	1.92	0.50
1:Q:34:ILE:HG12	1:Q:69:TYR:CE1	2.46	0.50
1:U:32:PRO:CG	1:U:55:GLY:O	2.55	0.50
1:U:147:ARG:CG	1:U:147:ARG:NH2	2.71	0.50
1:U:362:TYR:HD1	1:U:367:PRO:HA	1.75	0.50
1:V:34:ILE:CG2	1:V:35:VAL:N	2.74	0.50
1:V:34:ILE:HG12	1:V:69:TYR:CE1	2.46	0.50
1:V:164:PRO:HG3	1:V:174:ALA:HB1	1.92	0.50
1:B:299:MET:HE1	1:B:309:ILE:HD13	1.94	0.50
1:C:43:VAL:C	1:C:44:MET:HG2	2.37	0.50
1:C:167:GLU:OE1	1:E:61:LYS:CE	2.58	0.50
1:E:230:ALA:HA	1:E:233:SER:HB2	1.92	0.50
1:F:32:PRO:CG	1:F:55:GLY:O	2.55	0.50
1:F:347:ALA:HA	1:F:356:TRP:CZ2	2.46	0.50
1:G:164:PRO:HG3	1:G:174:ALA:HB1	1.92	0.50
1:I:169:TYR:OH	1:K:40:HIS:HB3	2.10	0.50
1:I:347:ALA:HA	1:I:356:TRP:CZ2	2.46	0.50
1:J:34:ILE:CG2	1:J:35:VAL:N	2.74	0.50
1:J:58:ALA:CA	1:J:65:LEU:CD2	2.89	0.50
1:K:347:ALA:HA	1:K:356:TRP:CZ2	2.47	0.50
1:L:164:PRO:HG2	1:L:174:ALA:HB3	1.92	0.50
1:M:34:ILE:HG23	1:M:68:LYS:N	2.26	0.50
1:M:332:PRO:HG2	1:M:335:ARG:NE	2.26	0.50
1:N:332:PRO:HG2	1:N:335:ARG:NE	2.26	0.50
1:O:169:TYR:OH	1:Q:40:HIS:HB3	2.10	0.50
1:O:347:ALA:HA	1:O:356:TRP:CZ2	2.46	0.50
1:Q:32:PRO:CG	1:Q:55:GLY:O	2.55	0.50
1:R:34:ILE:HG12	1:R:69:TYR:CE1	2.46	0.50
1:R:230:ALA:HA	1:R:233:SER:HB2	1.92	0.50
1:V:230:ALA:HA	1:V:233:SER:HB2	1.92	0.50
1:A:34:ILE:HG12	1:A:69:TYR:CE1	2.46	0.50
1:F:164:PRO:HG2	1:F:174:ALA:HB3	1.92	0.50
1:I:362:TYR:HE1	1:I:367:PRO:CB	2.19	0.50
1:J:36:GLY:HA3	1:J:66:THR:O	2.10	0.50
1:L:34:ILE:CG2	1:L:35:VAL:N	2.74	0.50
1:M:300:SER:HA	1:M:335:ARG:HG2	1.89	0.50
1:O:299:MET:HE1	1:O:309:ILE:HD13	1.94	0.50
1:P:22:ALA:HB1	1:P:348:SER:CB	2.41	0.50
1:P:39:ARG:HG2	1:P:66:THR:HB	1.93	0.50
1:P:58:ALA:CA	1:P:65:LEU:CD2	2.89	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:34:ILE:HD13	1:Q:67:LEU:CD1	2.41	0.50
1:S:34:ILE:HG23	1:S:68:LYS:N	2.26	0.50
1:S:34:ILE:HG12	1:S:69:TYR:CE1	2.46	0.50
1:S:164:PRO:HG2	1:S:174:ALA:HB3	1.92	0.50
1:S:299:MET:HE1	1:S:309:ILE:HD13	1.94	0.50
1:U:164:PRO:HG2	1:U:174:ALA:HB3	1.92	0.50
1:U:169:TYR:OH	1:W:40:HIS:HB3	2.10	0.50
1:V:22:ALA:HB1	1:V:348:SER:CB	2.41	0.50
1:V:299:MET:HE1	1:V:309:ILE:HD13	1.94	0.50
1:W:36:GLY:HA3	1:W:66:THR:O	2.10	0.50
1:W:58:ALA:CA	1:W:65:LEU:CD2	2.89	0.50
1:A:43:VAL:C	1:A:44:MET:HG2	2.37	0.50
1:A:164:PRO:HG2	1:A:174:ALA:HB3	1.92	0.50
1:B:362:TYR:HD1	1:B:367:PRO:HA	1.75	0.50
1:C:58:ALA:CA	1:C:65:LEU:CD2	2.89	0.50
1:D:40:HIS:O	1:D:41:GLN:O	2.30	0.50
1:D:347:ALA:HA	1:D:356:TRP:CZ2	2.46	0.50
1:E:34:ILE:CG2	1:E:35:VAL:N	2.74	0.50
1:E:43:VAL:C	1:E:44:MET:HG2	2.37	0.50
1:F:34:ILE:HG12	1:F:69:TYR:CE1	2.46	0.50
1:G:43:VAL:C	1:G:44:MET:HG2	2.37	0.50
1:G:299:MET:HE1	1:G:309:ILE:HD13	1.94	0.50
1:H:37:ARG:CG	1:H:38:PRO:N	2.75	0.50
1:H:164:PRO:HG2	1:H:174:ALA:HB3	1.92	0.50
1:J:142:LEU:HD21	1:J:165:ILE:CD1	2.39	0.50
1:K:36:GLY:HA3	1:K:66:THR:O	2.10	0.50
1:N:34:ILE:CG2	1:N:35:VAL:N	2.74	0.50
1:N:34:ILE:HG23	1:N:68:LYS:N	2.26	0.50
1:O:40:HIS:O	1:O:41:GLN:O	2.30	0.50
1:O:230:ALA:HA	1:O:233:SER:HB2	1.92	0.50
1:R:36:GLY:HA3	1:R:66:THR:O	2.10	0.50
1:R:169:TYR:OH	1:T:40:HIS:HB3	2.10	0.50
1:S:2:GLU:HA	1:S:2:GLU:OE2	2.10	0.50
1:U:43:VAL:C	1:U:44:MET:HG2	2.37	0.50
1:V:34:ILE:HG23	1:V:68:LYS:N	2.26	0.50
1:V:39:ARG:HG2	1:V:66:THR:HB	1.93	0.50
1:W:37:ARG:CG	1:W:38:PRO:N	2.75	0.50
1:W:43:VAL:C	1:W:44:MET:HG2	2.37	0.50
1:A:22:ALA:HB1	1:A:348:SER:CB	2.41	0.50
1:B:34:ILE:CG2	1:B:35:VAL:N	2.74	0.50
1:B:40:HIS:O	1:B:41:GLN:O	2.30	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:347:ALA:HA	1:B:356:TRP:CZ2	2.46	0.50
1:C:169:TYR:OH	1:E:40:HIS:HB3	2.10	0.50
1:F:37:ARG:CG	1:F:38:PRO:N	2.75	0.50
1:F:40:HIS:O	1:F:41:GLN:O	2.30	0.50
1:G:169:TYR:OH	1:I:40:HIS:HB3	2.10	0.50
1:G:347:ALA:HA	1:G:356:TRP:CZ2	2.46	0.50
1:M:40:HIS:O	1:M:41:GLN:O	2.30	0.50
1:O:34:ILE:CG2	1:O:35:VAL:N	2.74	0.50
1:O:34:ILE:HG23	1:O:68:LYS:N	2.26	0.50
1:O:164:PRO:HG3	1:O:174:ALA:HB1	1.92	0.50
1:P:43:VAL:C	1:P:44:MET:HG2	2.37	0.50
1:Q:40:HIS:O	1:Q:41:GLN:O	2.30	0.50
1:R:34:ILE:HG23	1:R:68:LYS:N	2.26	0.50
1:S:40:HIS:O	1:S:41:GLN:O	2.30	0.50
1:U:37:ARG:CG	1:U:38:PRO:N	2.75	0.50
1:W:39:ARG:HG2	1:W:66:THR:HB	1.92	0.50
1:W:332:PRO:HG2	1:W:335:ARG:NE	2.26	0.50
1:A:290:ARG:NH1	1:C:244:ASP:OD1	2.45	0.50
1:C:299:MET:HE1	1:C:309:ILE:HD13	1.94	0.50
1:D:32:PRO:CG	1:D:55:GLY:O	2.55	0.50
1:E:290:ARG:NH1	1:G:244:ASP:OD1	2.45	0.50
1:E:361:GLU:HB3	1:E:369:ILE:CD1	2.14	0.50
1:F:34:ILE:HD13	1:F:67:LEU:CD1	2.41	0.50
1:H:34:ILE:CG2	1:H:35:VAL:N	2.74	0.50
1:H:40:HIS:O	1:H:41:GLN:O	2.30	0.50
1:J:362:TYR:HE1	1:J:367:PRO:CB	2.19	0.50
1:K:40:HIS:O	1:K:41:GLN:O	2.30	0.50
1:K:299:MET:HE1	1:K:309:ILE:HD13	1.94	0.50
1:M:34:ILE:HG12	1:M:69:TYR:CE1	2.46	0.50
1:N:164:PRO:HG2	1:N:174:ALA:HB3	1.92	0.50
1:N:290:ARG:NH1	1:P:244:ASP:OD1	2.45	0.50
1:O:34:ILE:HD13	1:O:67:LEU:CD1	2.41	0.50
1:P:290:ARG:NH1	1:R:244:ASP:OD1	2.45	0.50
1:P:299:MET:HE1	1:P:309:ILE:HD13	1.94	0.50
1:P:332:PRO:HG2	1:P:335:ARG:NE	2.26	0.50
1:Q:34:ILE:CG2	1:Q:35:VAL:N	2.74	0.50
1:Q:347:ALA:HA	1:Q:356:TRP:CZ2	2.47	0.50
1:S:43:VAL:C	1:S:44:MET:HG2	2.37	0.50
1:T:290:ARG:NH1	1:V:244:ASP:OD1	2.45	0.50
1:C:287:ILE:HG13	1:C:288:ASP:N	2.27	0.50
1:D:2:GLU:OE2	1:D:2:GLU:HA	2.10	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:332:PRO:HG2	1:D:335:ARG:NE	2.26	0.50
1:E:347:ALA:HA	1:E:356:TRP:CZ2	2.46	0.50
1:F:169:TYR:OH	1:H:40:HIS:HB3	2.10	0.50
1:G:290:ARG:NH1	1:I:244:ASP:OD1	2.45	0.50
1:I:34:ILE:HG12	1:I:69:TYR:CE1	2.46	0.50
1:I:290:ARG:NH1	1:K:244:ASP:OD1	2.45	0.50
1:I:299:MET:HE1	1:I:309:ILE:HD13	1.94	0.50
1:I:332:PRO:HG2	1:I:335:ARG:NE	2.26	0.50
1:J:34:ILE:CG2	1:J:67:LEU:HD22	2.27	0.50
1:K:332:PRO:HG2	1:K:335:ARG:NE	2.26	0.50
1:L:39:ARG:HG2	1:L:66:THR:HB	1.93	0.50
1:L:58:ALA:CA	1:L:65:LEU:CD2	2.89	0.50
1:M:35:VAL:CA	1:M:54:VAL:CG2	2.90	0.50
1:N:43:VAL:C	1:N:44:MET:HG2	2.37	0.50
1:N:58:ALA:CA	1:N:65:LEU:CD2	2.89	0.50
1:P:34:ILE:HG12	1:P:69:TYR:CE1	2.46	0.50
1:Q:169:TYR:OH	1:S:40:HIS:HB3	2.10	0.50
1:R:39:ARG:HG2	1:R:66:THR:HB	1.92	0.50
1:S:37:ARG:CG	1:S:38:PRO:N	2.75	0.50
1:S:347:ALA:HA	1:S:356:TRP:CZ2	2.46	0.50
1:T:230:ALA:HA	1:T:233:SER:HB2	1.92	0.50
1:U:40:HIS:O	1:U:41:GLN:O	2.30	0.50
1:W:34:ILE:CG2	1:W:35:VAL:N	2.74	0.50
1:W:34:ILE:HG23	1:W:68:LYS:N	2.26	0.50
1:W:40:HIS:O	1:W:41:GLN:O	2.30	0.50
1:A:287:ILE:HG13	1:A:288:ASP:N	2.27	0.50
1:E:169:TYR:OH	1:G:40:HIS:HB3	2.10	0.50
1:E:287:ILE:HG13	1:E:288:ASP:N	2.27	0.50
1:G:37:ARG:CG	1:G:38:PRO:N	2.75	0.50
1:H:32:PRO:CG	1:H:55:GLY:O	2.55	0.50
1:I:230:ALA:HA	1:I:233:SER:HB2	1.92	0.50
1:J:169:TYR:OH	1:L:40:HIS:HB3	2.10	0.50
1:J:299:MET:HE1	1:J:309:ILE:HD13	1.94	0.50
1:K:34:ILE:HD13	1:K:67:LEU:CD1	2.41	0.50
1:K:300:SER:HA	1:K:335:ARG:HG2	1.89	0.50
1:O:35:VAL:CA	1:O:54:VAL:CG2	2.90	0.50
1:Q:34:ILE:CG2	1:Q:67:LEU:HD22	2.28	0.50
1:R:35:VAL:CA	1:R:54:VAL:CG2	2.90	0.50
1:R:43:VAL:C	1:R:44:MET:HG2	2.37	0.50
1:T:35:VAL:CA	1:T:54:VAL:CG2	2.90	0.50
1:T:299:MET:HE1	1:T:309:ILE:HD13	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:32:PRO:CG	1:B:55:GLY:O	2.55	0.50
1:B:147:ARG:CG	1:B:147:ARG:NH2	2.71	0.50
1:D:34:ILE:HD13	1:D:67:LEU:CD1	2.41	0.50
1:D:37:ARG:CG	1:D:38:PRO:N	2.75	0.50
1:G:34:ILE:HG12	1:G:69:TYR:CE1	2.46	0.50
1:G:230:ALA:HA	1:G:233:SER:HB2	1.92	0.50
1:H:169:TYR:OH	1:J:40:HIS:HB3	2.10	0.50
1:I:40:HIS:O	1:I:41:GLN:O	2.30	0.50
1:I:43:VAL:C	1:I:44:MET:HG2	2.37	0.50
1:K:35:VAL:CA	1:K:54:VAL:CG2	2.90	0.50
1:L:290:ARG:NH1	1:N:244:ASP:OD1	2.45	0.50
1:M:299:MET:HE1	1:M:309:ILE:HD13	1.94	0.50
1:O:290:ARG:NH1	1:Q:244:ASP:OD1	2.45	0.50
1:P:35:VAL:CA	1:P:54:VAL:CG2	2.90	0.50
1:Q:147:ARG:CG	1:Q:147:ARG:NH2	2.71	0.50
1:U:347:ALA:HA	1:U:356:TRP:CZ2	2.46	0.50
1:V:37:ARG:CG	1:V:38:PRO:N	2.75	0.50
1:B:34:ILE:HD13	1:B:67:LEU:CD1	2.41	0.49
1:B:34:ILE:HG12	1:B:69:TYR:CE1	2.46	0.49
1:C:347:ALA:HA	1:C:356:TRP:CZ2	2.46	0.49
1:F:39:ARG:HG2	1:F:66:THR:HB	1.93	0.49
1:G:287:ILE:HG13	1:G:288:ASP:N	2.27	0.49
1:J:39:ARG:HG2	1:J:66:THR:HB	1.93	0.49
1:J:40:HIS:O	1:J:41:GLN:O	2.30	0.49
1:K:34:ILE:CG2	1:K:67:LEU:HD22	2.28	0.49
1:K:290:ARG:NH1	1:M:244:ASP:OD1	2.45	0.49
1:M:34:ILE:CG2	1:M:35:VAL:N	2.74	0.49
1:M:290:ARG:NH1	1:O:244:ASP:OD1	2.45	0.49
1:Q:34:ILE:HG23	1:Q:68:LYS:N	2.26	0.49
1:T:34:ILE:HG23	1:T:68:LYS:N	2.26	0.49
1:U:299:MET:HE1	1:U:309:ILE:HD13	1.94	0.49
1:V:40:HIS:O	1:V:41:GLN:O	2.30	0.49
1:V:142:LEU:HD21	1:V:165:ILE:CD1	2.39	0.49
1:W:22:ALA:HB1	1:W:348:SER:CB	2.41	0.49
1:D:39:ARG:HG2	1:D:66:THR:HB	1.92	0.49
1:E:147:ARG:CG	1:E:147:ARG:NH2	2.71	0.49
1:F:44:MET:CG	1:F:45:VAL:N	2.75	0.49
1:I:287:ILE:HG22	1:I:290:ARG:CZ	2.43	0.49
1:K:230:ALA:HA	1:K:233:SER:HB2	1.92	0.49
1:L:40:HIS:O	1:L:41:GLN:O	2.30	0.49
1:L:43:VAL:C	1:L:44:MET:HG2	2.37	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:287:ILE:HG22	1:L:290:ARG:CZ	2.43	0.49
1:L:299:MET:HE1	1:L:309:ILE:HD13	1.94	0.49
1:N:39:ARG:HG2	1:N:66:THR:HB	1.93	0.49
1:Q:39:ARG:NE	1:Q:66:THR:CA	2.69	0.49
1:S:34:ILE:CG2	1:S:35:VAL:N	2.74	0.49
1:S:169:TYR:OH	1:U:40:HIS:HB3	2.10	0.49
1:W:34:ILE:HG12	1:W:69:TYR:CE1	2.46	0.49
1:W:299:MET:HE1	1:W:309:ILE:HD13	1.94	0.49
1:A:347:ALA:HA	1:A:356:TRP:CZ2	2.46	0.49
1:C:35:VAL:CA	1:C:54:VAL:CG2	2.90	0.49
1:F:287:ILE:HG22	1:F:290:ARG:CZ	2.43	0.49
1:H:43:VAL:C	1:H:44:MET:HG2	2.37	0.49
1:I:39:ARG:HG2	1:I:66:THR:HB	1.93	0.49
1:M:230:ALA:HA	1:M:233:SER:HB2	1.92	0.49
1:O:287:ILE:HG22	1:O:290:ARG:CZ	2.43	0.49
1:Q:35:VAL:CA	1:Q:54:VAL:CG2	2.90	0.49
1:Q:43:VAL:C	1:Q:44:MET:HG2	2.37	0.49
1:Q:299:MET:HE1	1:Q:309:ILE:HD13	1.94	0.49
1:R:37:ARG:CG	1:R:38:PRO:N	2.75	0.49
1:T:40:HIS:O	1:T:41:GLN:O	2.30	0.49
1:V:35:VAL:CA	1:V:54:VAL:CG2	2.90	0.49
1:V:287:ILE:HG22	1:V:290:ARG:CZ	2.43	0.49
1:W:287:ILE:HG13	1:W:288:ASP:N	2.27	0.49
1:W:347:ALA:HA	1:W:356:TRP:CZ2	2.47	0.49
1:A:39:ARG:HG2	1:A:66:THR:HB	1.93	0.49
1:H:22:ALA:HB1	1:H:348:SER:CB	2.41	0.49
1:I:35:VAL:CA	1:I:54:VAL:CG2	2.90	0.49
1:M:43:VAL:C	1:M:44:MET:HG2	2.37	0.49
1:N:35:VAL:CA	1:N:54:VAL:CG2	2.90	0.49
1:O:142:LEU:HD21	1:O:165:ILE:CD1	2.39	0.49
1:S:39:ARG:HG2	1:S:66:THR:HB	1.93	0.49
1:S:287:ILE:HG22	1:S:290:ARG:CZ	2.43	0.49
1:T:43:VAL:C	1:T:44:MET:HG2	2.37	0.49
1:T:169:TYR:OH	1:V:40:HIS:HB3	2.10	0.49
1:A:147:ARG:CG	1:A:147:ARG:NH2	2.71	0.49
1:B:287:ILE:HG22	1:B:290:ARG:CZ	2.43	0.49
1:D:290:ARG:NH1	1:F:244:ASP:OD1	2.45	0.49
1:E:35:VAL:CA	1:E:54:VAL:CG2	2.90	0.49
1:G:40:HIS:O	1:G:41:GLN:O	2.30	0.49
1:H:290:ARG:NH1	1:J:244:ASP:OD1	2.45	0.49
1:I:43:VAL:O	1:I:44:MET:HG2	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:43:VAL:C	1:J:44:MET:HG2	2.37	0.49
1:K:43:VAL:C	1:K:44:MET:HG2	2.37	0.49
1:L:287:ILE:HG13	1:L:288:ASP:N	2.27	0.49
1:M:34:ILE:HD13	1:M:67:LEU:CD1	2.41	0.49
1:M:287:ILE:HG22	1:M:290:ARG:CZ	2.43	0.49
1:N:40:HIS:O	1:N:41:GLN:O	2.30	0.49
1:O:43:VAL:C	1:O:44:MET:HG2	2.37	0.49
1:R:299:MET:HE1	1:R:309:ILE:HD13	1.94	0.49
1:S:32:PRO:CG	1:S:55:GLY:O	2.55	0.49
1:S:34:ILE:HD13	1:S:67:LEU:CD1	2.41	0.49
1:T:43:VAL:O	1:T:44:MET:HG2	2.13	0.49
1:U:39:ARG:HG2	1:U:66:THR:HB	1.93	0.49
1:U:287:ILE:HG22	1:U:290:ARG:CZ	2.43	0.49
1:U:362:TYR:HE1	1:U:367:PRO:CB	2.19	0.49
1:A:40:HIS:O	1:A:41:GLN:O	2.30	0.49
1:C:142:LEU:HD21	1:C:165:ILE:CD1	2.39	0.49
1:D:34:ILE:CG2	1:D:35:VAL:N	2.74	0.49
1:E:37:ARG:CG	1:E:38:PRO:N	2.75	0.49
1:E:39:ARG:HG2	1:E:66:THR:HB	1.93	0.49
1:E:40:HIS:O	1:E:41:GLN:O	2.30	0.49
1:F:34:ILE:CG2	1:F:35:VAL:N	2.74	0.49
1:F:43:VAL:C	1:F:44:MET:HG2	2.37	0.49
1:F:290:ARG:NH1	1:H:244:ASP:OD1	2.45	0.49
1:H:39:ARG:HG2	1:H:66:THR:HB	1.93	0.49
1:H:287:ILE:HG22	1:H:290:ARG:CZ	2.43	0.49
1:H:299:MET:HE1	1:H:309:ILE:HD13	1.94	0.49
1:I:34:ILE:HG22	1:I:35:VAL:H	1.78	0.49
1:I:37:ARG:CG	1:I:38:PRO:N	2.75	0.49
1:I:147:ARG:CG	1:I:147:ARG:NH2	2.71	0.49
1:I:300:SER:HA	1:I:335:ARG:HG2	1.89	0.49
1:J:43:VAL:O	1:J:44:MET:HG2	2.13	0.49
1:K:34:ILE:HG22	1:K:35:VAL:H	1.78	0.49
1:O:2:GLU:OE2	1:O:2:GLU:HA	2.10	0.49
1:P:37:ARG:CG	1:P:38:PRO:N	2.75	0.49
1:P:40:HIS:O	1:P:41:GLN:O	2.30	0.49
1:P:147:ARG:CG	1:P:147:ARG:NH2	2.71	0.49
1:Q:290:ARG:NH1	1:S:244:ASP:OD1	2.45	0.49
1:R:34:ILE:HG22	1:R:35:VAL:H	1.78	0.49
1:R:40:HIS:O	1:R:41:GLN:O	2.30	0.49
1:R:290:ARG:NH1	1:T:244:ASP:OD1	2.45	0.49
1:U:43:VAL:O	1:U:44:MET:HG2	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:34:ILE:HD13	1:V:67:LEU:CD1	2.41	0.49
1:B:34:ILE:HG22	1:B:35:VAL:H	1.78	0.49
1:C:290:ARG:NH1	1:E:244:ASP:OD1	2.45	0.49
1:D:43:VAL:C	1:D:44:MET:HG2	2.37	0.49
1:E:8:LEU:HB2	1:E:103:THR:HA	1.95	0.49
1:F:43:VAL:O	1:F:44:MET:HG2	2.13	0.49
1:H:336:LYS:HE2	2:H:401:ADP:C5'	2.38	0.49
1:I:287:ILE:HG13	1:I:288:ASP:N	2.27	0.49
1:J:287:ILE:HG13	1:J:288:ASP:N	2.27	0.49
1:K:37:ARG:CG	1:K:38:PRO:N	2.75	0.49
1:K:43:VAL:O	1:K:44:MET:HG2	2.13	0.49
1:L:300:SER:HA	1:L:335:ARG:HG2	1.89	0.49
1:M:39:ARG:HG2	1:M:66:THR:HB	1.93	0.49
1:M:147:ARG:CG	1:M:147:ARG:NH2	2.71	0.49
1:O:22:ALA:HB1	1:O:348:SER:CB	2.41	0.49
1:O:37:ARG:CG	1:O:38:PRO:N	2.75	0.49
1:P:43:VAL:O	1:P:44:MET:HG2	2.13	0.49
1:Q:298:VAL:HG12	1:Q:335:ARG:NH1	2.28	0.49
1:S:290:ARG:NH1	1:U:244:ASP:OD1	2.45	0.49
1:T:147:ARG:CG	1:T:147:ARG:NH2	2.71	0.49
1:U:34:ILE:HD13	1:U:67:LEU:CD1	2.41	0.49
1:V:43:VAL:C	1:V:44:MET:HG2	2.37	0.49
1:W:43:VAL:O	1:W:44:MET:HG2	2.13	0.49
1:B:290:ARG:NH1	1:D:244:ASP:OD1	2.45	0.49
1:C:37:ARG:CG	1:C:38:PRO:N	2.75	0.49
1:C:40:HIS:O	1:C:41:GLN:O	2.30	0.49
1:D:34:ILE:HG22	1:D:35:VAL:H	1.78	0.49
1:E:43:VAL:O	1:E:44:MET:HG2	2.13	0.49
1:E:287:ILE:HG22	1:E:290:ARG:CZ	2.43	0.49
1:H:142:LEU:HD21	1:H:165:ILE:CD1	2.39	0.49
1:H:298:VAL:HG12	1:H:335:ARG:NH1	2.28	0.49
1:H:361:GLU:HB3	1:H:369:ILE:CD1	2.14	0.49
1:M:37:ARG:CG	1:M:38:PRO:N	2.75	0.49
1:M:43:VAL:O	1:M:44:MET:HG2	2.13	0.49
1:N:287:ILE:HG13	1:N:288:ASP:N	2.27	0.49
1:O:287:ILE:HG13	1:O:288:ASP:N	2.27	0.49
1:Q:8:LEU:HB2	1:Q:103:THR:HA	1.95	0.49
1:Q:39:ARG:HG2	1:Q:66:THR:HB	1.93	0.49
1:T:34:ILE:HG22	1:T:35:VAL:H	1.78	0.49
1:T:37:ARG:CG	1:T:38:PRO:N	2.75	0.49
1:U:34:ILE:CG2	1:U:35:VAL:N	2.74	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:34:ILE:HD13	1:W:67:LEU:CD1	2.41	0.49
1:B:43:VAL:O	1:B:44:MET:HG2	2.13	0.49
1:D:8:LEU:HB2	1:D:103:THR:HA	1.95	0.49
1:D:299:MET:HE1	1:D:309:ILE:HD13	1.94	0.49
1:F:8:LEU:HB2	1:F:103:THR:HA	1.95	0.49
1:F:298:VAL:HG12	1:F:335:ARG:NH1	2.28	0.49
1:G:8:LEU:HB2	1:G:103:THR:HA	1.95	0.49
1:I:34:ILE:HD13	1:I:67:LEU:CD1	2.41	0.49
1:J:287:ILE:HG22	1:J:290:ARG:CZ	2.43	0.49
1:O:298:VAL:HG12	1:O:335:ARG:NH1	2.28	0.49
1:P:287:ILE:HG22	1:P:290:ARG:CZ	2.43	0.49
1:Q:43:VAL:O	1:Q:44:MET:HG2	2.13	0.49
1:U:287:ILE:HG13	1:U:288:ASP:N	2.27	0.49
1:U:290:ARG:NH1	1:W:244:ASP:OD1	2.45	0.49
1:A:37:ARG:CG	1:A:38:PRO:N	2.75	0.49
1:A:43:VAL:O	1:A:44:MET:HG2	2.13	0.49
1:A:287:ILE:HG22	1:A:290:ARG:CZ	2.43	0.49
1:A:299:MET:HE1	1:A:309:ILE:HD13	1.94	0.49
1:B:58:ALA:HA	1:B:65:LEU:CD2	2.43	0.49
1:D:43:VAL:O	1:D:44:MET:HG2	2.13	0.49
1:E:299:MET:HE1	1:E:309:ILE:HD13	1.94	0.49
1:G:2:GLU:OE2	1:G:2:GLU:HA	2.10	0.49
1:G:35:VAL:CA	1:G:54:VAL:CG2	2.90	0.49
1:H:34:ILE:HD13	1:H:67:LEU:CD1	2.41	0.49
1:H:35:VAL:CA	1:H:54:VAL:CG2	2.90	0.49
1:H:287:ILE:HG13	1:H:288:ASP:N	2.27	0.49
1:I:22:ALA:HB1	1:I:348:SER:CB	2.41	0.49
1:I:58:ALA:HA	1:I:65:LEU:CD2	2.43	0.49
1:J:180:LEU:HD12	1:J:181:ALA:O	2.13	0.49
1:K:2:GLU:OE2	1:K:2:GLU:HA	2.11	0.49
1:K:335:ARG:O	1:K:338:SER:HB3	2.13	0.49
1:L:35:VAL:CA	1:L:54:VAL:CG2	2.90	0.49
1:O:8:LEU:HB2	1:O:103:THR:HA	1.95	0.49
1:S:298:VAL:HG12	1:S:335:ARG:NH1	2.28	0.49
1:U:180:LEU:HD12	1:U:181:ALA:O	2.13	0.49
1:B:43:VAL:C	1:B:44:MET:HG2	2.37	0.48
1:C:8:LEU:HB2	1:C:103:THR:HA	1.95	0.48
1:D:58:ALA:HA	1:D:65:LEU:CD2	2.43	0.48
1:F:335:ARG:O	1:F:338:SER:HB3	2.13	0.48
1:H:335:ARG:O	1:H:338:SER:HB3	2.13	0.48
1:I:335:ARG:O	1:I:338:SER:HB3	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:35:VAL:CA	1:J:54:VAL:CG2	2.90	0.48
1:J:290:ARG:NH1	1:L:244:ASP:OD1	2.45	0.48
1:J:298:VAL:HG12	1:J:335:ARG:NH1	2.28	0.48
1:M:335:ARG:O	1:M:338:SER:HB3	2.13	0.48
1:O:335:ARG:O	1:O:338:SER:HB3	2.13	0.48
1:P:34:ILE:HG22	1:P:35:VAL:H	1.78	0.48
1:Q:287:ILE:HG13	1:Q:288:ASP:N	2.27	0.48
1:Q:335:ARG:O	1:Q:338:SER:HB3	2.13	0.48
1:R:287:ILE:HG22	1:R:290:ARG:CZ	2.43	0.48
1:S:8:LEU:HB2	1:S:103:THR:HA	1.95	0.48
1:V:180:LEU:HD12	1:V:181:ALA:O	2.13	0.48
1:B:143:TYR:HD1	1:B:143:TYR:C	2.09	0.48
1:B:180:LEU:HD12	1:B:181:ALA:O	2.13	0.48
1:C:22:ALA:HB1	1:C:348:SER:CB	2.41	0.48
1:D:35:VAL:CA	1:D:54:VAL:CG2	2.90	0.48
1:D:287:ILE:HG22	1:D:290:ARG:CZ	2.43	0.48
1:D:335:ARG:O	1:D:338:SER:HB3	2.13	0.48
1:E:180:LEU:HD12	1:E:181:ALA:O	2.13	0.48
1:G:34:ILE:HG22	1:G:35:VAL:H	1.78	0.48
1:G:43:VAL:O	1:G:44:MET:HG2	2.13	0.48
1:H:8:LEU:HB2	1:H:103:THR:HA	1.95	0.48
1:K:58:ALA:HA	1:K:65:LEU:CD2	2.43	0.48
1:K:287:ILE:HG22	1:K:290:ARG:CZ	2.43	0.48
1:M:132:MET:HE2	1:M:132:MET:HB2	1.75	0.48
1:R:8:LEU:HB2	1:R:103:THR:HA	1.95	0.48
1:R:22:ALA:HB1	1:R:348:SER:CB	2.41	0.48
1:R:287:ILE:HG13	1:R:288:ASP:N	2.27	0.48
1:S:335:ARG:O	1:S:338:SER:HB3	2.13	0.48
1:T:287:ILE:HG13	1:T:288:ASP:N	2.27	0.48
1:U:35:VAL:CA	1:U:54:VAL:CG2	2.90	0.48
1:U:58:ALA:HA	1:U:65:LEU:CD2	2.43	0.48
1:B:58:ALA:CB	1:B:65:LEU:CD2	2.90	0.48
1:C:287:ILE:HG22	1:C:290:ARG:CZ	2.43	0.48
1:G:58:ALA:HA	1:G:65:LEU:CD2	2.43	0.48
1:H:43:VAL:O	1:H:44:MET:HG2	2.13	0.48
1:I:167:GLU:CD	1:K:61:LYS:CE	2.85	0.48
1:K:180:LEU:HD12	1:K:181:ALA:O	2.13	0.48
1:M:58:ALA:CB	1:M:65:LEU:CD2	2.90	0.48
1:N:43:VAL:O	1:N:44:MET:HG2	2.13	0.48
1:P:8:LEU:HB2	1:P:103:THR:HA	1.95	0.48
1:P:180:LEU:HD12	1:P:181:ALA:O	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:287:ILE:HG13	1:P:288:ASP:N	2.27	0.48
1:R:34:ILE:HG22	1:R:54:VAL:HG21	1.96	0.48
1:S:58:ALA:HA	1:S:65:LEU:CD2	2.44	0.48
1:T:287:ILE:HG22	1:T:290:ARG:CZ	2.43	0.48
1:V:2:GLU:OE2	1:V:2:GLU:HA	2.10	0.48
1:V:335:ARG:O	1:V:338:SER:HB3	2.13	0.48
1:W:35:VAL:CA	1:W:54:VAL:CG2	2.90	0.48
1:W:287:ILE:HG22	1:W:290:ARG:CZ	2.43	0.48
1:A:8:LEU:HB2	1:A:103:THR:HA	1.95	0.48
1:B:8:LEU:HB2	1:B:103:THR:HA	1.95	0.48
1:B:35:VAL:CA	1:B:54:VAL:CG2	2.90	0.48
1:B:335:ARG:O	1:B:338:SER:HB3	2.14	0.48
1:C:2:GLU:OE2	1:C:2:GLU:HA	2.10	0.48
1:D:180:LEU:HD12	1:D:181:ALA:O	2.13	0.48
1:F:35:VAL:CA	1:F:54:VAL:CG2	2.90	0.48
1:G:180:LEU:HD12	1:G:181:ALA:O	2.13	0.48
1:I:8:LEU:HB2	1:I:103:THR:HA	1.95	0.48
1:J:8:LEU:HB2	1:J:103:THR:HA	1.95	0.48
1:J:34:ILE:HD13	1:J:67:LEU:CD1	2.41	0.48
1:J:37:ARG:CG	1:J:38:PRO:N	2.75	0.48
1:K:287:ILE:HG13	1:K:288:ASP:N	2.27	0.48
1:L:8:LEU:HB2	1:L:103:THR:HA	1.95	0.48
1:L:43:VAL:O	1:L:44:MET:HG2	2.13	0.48
1:N:8:LEU:HB2	1:N:103:THR:HA	1.95	0.48
1:N:287:ILE:HG22	1:N:290:ARG:CZ	2.43	0.48
1:P:34:ILE:HG22	1:P:54:VAL:HG21	1.96	0.48
1:Q:287:ILE:HG22	1:Q:290:ARG:CZ	2.43	0.48
1:T:335:ARG:O	1:T:338:SER:HB3	2.13	0.48
1:U:8:LEU:HB2	1:U:103:THR:HA	1.95	0.48
1:U:335:ARG:O	1:U:338:SER:HB3	2.13	0.48
1:A:223:PHE:CD1	1:A:259:GLU:CG	2.96	0.48
1:A:298:VAL:HG12	1:A:335:ARG:NH1	2.28	0.48
1:C:43:VAL:O	1:C:44:MET:HG2	2.13	0.48
1:G:300:SER:HA	1:G:335:ARG:HG2	1.89	0.48
1:H:167:GLU:CD	1:J:61:LYS:CE	2.85	0.48
1:J:335:ARG:O	1:J:338:SER:HB3	2.14	0.48
1:M:8:LEU:HB2	1:M:103:THR:HA	1.95	0.48
1:N:299:MET:HE1	1:N:309:ILE:HD13	1.94	0.48
1:Q:58:ALA:CB	1:Q:65:LEU:CD2	2.90	0.48
1:Q:58:ALA:HA	1:Q:65:LEU:CD2	2.43	0.48
1:R:2:GLU:OE2	1:R:2:GLU:HA	2.10	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:43:VAL:O	1:R:44:MET:HG2	2.13	0.48
1:R:180:LEU:HD12	1:R:181:ALA:O	2.13	0.48
1:S:287:ILE:HG13	1:S:288:ASP:N	2.27	0.48
1:T:8:LEU:HB2	1:T:103:THR:HA	1.95	0.48
1:V:43:VAL:O	1:V:44:MET:HG2	2.13	0.48
1:V:298:VAL:HG12	1:V:335:ARG:NH1	2.28	0.48
1:W:34:ILE:HG22	1:W:35:VAL:H	1.78	0.48
1:A:180:LEU:HD12	1:A:181:ALA:O	2.13	0.48
1:E:34:ILE:HG23	1:E:68:LYS:H	1.79	0.48
1:E:298:VAL:HG12	1:E:335:ARG:NH1	2.28	0.48
1:F:58:ALA:HA	1:F:65:LEU:CD2	2.43	0.48
1:F:287:ILE:HG13	1:F:288:ASP:N	2.27	0.48
1:K:147:ARG:CG	1:K:147:ARG:NH2	2.71	0.48
1:L:34:ILE:HD13	1:L:67:LEU:CD1	2.41	0.48
1:L:34:ILE:HG23	1:L:68:LYS:H	1.79	0.48
1:L:298:VAL:HG12	1:L:335:ARG:NH1	2.28	0.48
1:M:142:LEU:HD21	1:M:165:ILE:CD1	2.38	0.48
1:M:287:ILE:HG13	1:M:288:ASP:N	2.27	0.48
1:M:298:VAL:HG12	1:M:335:ARG:NH1	2.28	0.48
1:N:34:ILE:HG22	1:N:54:VAL:HG21	1.96	0.48
1:N:34:ILE:HG23	1:N:68:LYS:H	1.79	0.48
1:Q:34:ILE:HG23	1:Q:68:LYS:H	1.79	0.48
1:Q:361:GLU:HB3	1:Q:369:ILE:CD1	2.14	0.48
1:T:34:ILE:HG22	1:T:54:VAL:HG21	1.96	0.48
1:U:223:PHE:CD1	1:U:259:GLU:CG	2.96	0.48
1:V:58:ALA:HA	1:V:65:LEU:CD2	2.44	0.48
1:V:223:PHE:CD1	1:V:259:GLU:CG	2.96	0.48
1:V:287:ILE:HG13	1:V:288:ASP:N	2.27	0.48
1:W:8:LEU:HB2	1:W:103:THR:HA	1.95	0.48
1:W:58:ALA:HA	1:W:65:LEU:CD2	2.43	0.48
1:W:223:PHE:CD1	1:W:259:GLU:CG	2.96	0.48
1:C:298:VAL:HG12	1:C:335:ARG:NH1	2.28	0.48
1:E:34:ILE:HD13	1:E:67:LEU:CD1	2.41	0.48
1:E:58:ALA:HA	1:E:65:LEU:CD2	2.43	0.48
1:F:180:LEU:HD12	1:F:181:ALA:O	2.13	0.48
1:G:287:ILE:HG22	1:G:290:ARG:CZ	2.43	0.48
1:H:180:LEU:HD12	1:H:181:ALA:O	2.13	0.48
1:I:298:VAL:HG12	1:I:335:ARG:NH1	2.28	0.48
1:N:58:ALA:HA	1:N:65:LEU:CD2	2.43	0.48
1:N:167:GLU:CD	1:P:61:LYS:CE	2.85	0.48
1:O:34:ILE:HG23	1:O:68:LYS:H	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:58:ALA:CB	1:O:65:LEU:CD2	2.90	0.48
1:O:180:LEU:HD12	1:O:181:ALA:O	2.13	0.48
1:P:34:ILE:HG23	1:P:68:LYS:H	1.79	0.48
1:Q:223:PHE:CD1	1:Q:259:GLU:CG	2.96	0.48
1:S:223:PHE:CD1	1:S:259:GLU:CG	2.96	0.48
1:T:34:ILE:HD13	1:T:67:LEU:CD1	2.41	0.48
1:U:298:VAL:HG12	1:U:335:ARG:NH1	2.28	0.48
1:A:34:ILE:HG22	1:A:54:VAL:HG21	1.96	0.48
1:A:335:ARG:O	1:A:338:SER:HB3	2.13	0.48
1:B:287:ILE:HG13	1:B:288:ASP:N	2.27	0.48
1:D:58:ALA:CB	1:D:65:LEU:CD2	2.90	0.48
1:D:167:GLU:CD	1:F:61:LYS:CE	2.85	0.48
1:F:34:ILE:HG23	1:F:68:LYS:H	1.79	0.48
1:F:39:ARG:NE	1:F:66:THR:CA	2.69	0.48
1:G:298:VAL:HG12	1:G:335:ARG:NH1	2.28	0.48
1:G:335:ARG:O	1:G:338:SER:HB3	2.14	0.48
1:G:361:GLU:HB3	1:G:369:ILE:CD1	2.14	0.48
1:I:8:LEU:O	1:I:104:LEU:N	2.47	0.48
1:K:58:ALA:CB	1:K:65:LEU:CD2	2.90	0.48
1:M:58:ALA:HA	1:M:65:LEU:CD2	2.43	0.48
1:O:223:PHE:CD1	1:O:259:GLU:CG	2.96	0.48
1:P:58:ALA:HA	1:P:65:LEU:CD2	2.43	0.48
1:Q:180:LEU:HD12	1:Q:181:ALA:O	2.14	0.48
1:R:58:ALA:HA	1:R:65:LEU:CD2	2.43	0.48
1:T:58:ALA:HA	1:T:65:LEU:CD2	2.43	0.48
1:T:223:PHE:CD1	1:T:259:GLU:CG	2.96	0.48
1:W:34:ILE:HG22	1:W:54:VAL:HG21	1.96	0.48
1:C:223:PHE:CD1	1:C:259:GLU:CG	2.96	0.48
1:D:287:ILE:HG13	1:D:288:ASP:N	2.27	0.48
1:E:369:ILE:HG22	1:E:370:VAL:N	2.29	0.48
1:F:288:ASP:OD2	1:H:204:ALA:HB1	2.14	0.48
1:G:34:ILE:HD13	1:G:67:LEU:CD1	2.41	0.48
1:G:34:ILE:HG23	1:G:68:LYS:H	1.79	0.48
1:K:369:ILE:HG22	1:K:370:VAL:N	2.29	0.48
1:L:180:LEU:HD12	1:L:181:ALA:O	2.13	0.48
1:N:34:ILE:HG22	1:N:35:VAL:H	1.78	0.48
1:N:369:ILE:HG22	1:N:370:VAL:N	2.29	0.48
1:P:335:ARG:O	1:P:338:SER:HB3	2.13	0.48
1:S:43:VAL:O	1:S:44:MET:HG2	2.13	0.48
1:T:8:LEU:O	1:T:104:LEU:N	2.47	0.48
1:T:142:LEU:HD21	1:T:165:ILE:CD1	2.39	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:369:ILE:HG22	1:V:370:VAL:N	2.29	0.48
1:W:180:LEU:HD12	1:W:181:ALA:O	2.13	0.48
1:A:142:LEU:HD21	1:A:165:ILE:CD1	2.38	0.48
1:B:37:ARG:CG	1:B:38:PRO:N	2.75	0.48
1:B:288:ASP:OD2	1:D:204:ALA:HB1	2.14	0.48
1:B:298:VAL:HG12	1:B:335:ARG:NH1	2.28	0.48
1:C:34:ILE:HG23	1:C:68:LYS:H	1.79	0.48
1:C:44:MET:CG	1:C:45:VAL:N	2.75	0.48
1:C:58:ALA:HA	1:C:65:LEU:CD2	2.43	0.48
1:G:369:ILE:HG22	1:G:370:VAL:N	2.29	0.48
1:J:288:ASP:OD2	1:L:204:ALA:HB1	2.14	0.48
1:L:34:ILE:HG22	1:L:54:VAL:HG21	1.96	0.48
1:L:58:ALA:HA	1:L:65:LEU:CD2	2.43	0.48
1:M:223:PHE:CD1	1:M:259:GLU:CG	2.96	0.48
1:N:180:LEU:HD12	1:N:181:ALA:O	2.13	0.48
1:P:34:ILE:HD13	1:P:67:LEU:CD1	2.41	0.48
1:Q:288:ASP:OD2	1:S:204:ALA:HB1	2.14	0.48
1:S:34:ILE:HG23	1:S:68:LYS:H	1.79	0.48
1:S:180:LEU:HD12	1:S:181:ALA:O	2.13	0.48
1:T:369:ILE:HG22	1:T:370:VAL:N	2.29	0.48
1:U:288:ASP:OD2	1:W:204:ALA:HB1	2.14	0.48
1:V:8:LEU:O	1:V:104:LEU:N	2.47	0.48
1:V:335:ARG:HA	1:V:335:ARG:HD3	1.39	0.48
1:W:34:ILE:HG23	1:W:68:LYS:H	1.79	0.48
1:W:335:ARG:O	1:W:338:SER:HB3	2.14	0.48
1:W:369:ILE:HG22	1:W:370:VAL:N	2.29	0.48
1:A:31:PHE:HA	1:A:32:PRO:HD3	1.66	0.47
1:A:58:ALA:HA	1:A:65:LEU:CD2	2.43	0.47
1:D:288:ASP:OD2	1:F:204:ALA:HB1	2.14	0.47
1:E:8:LEU:O	1:E:104:LEU:N	2.47	0.47
1:E:44:MET:CG	1:E:45:VAL:N	2.76	0.47
1:H:34:ILE:HG23	1:H:68:LYS:H	1.79	0.47
1:H:58:ALA:HA	1:H:65:LEU:CD2	2.43	0.47
1:I:44:MET:CG	1:I:45:VAL:N	2.75	0.47
1:I:180:LEU:HD12	1:I:181:ALA:O	2.13	0.47
1:K:31:PHE:HA	1:K:32:PRO:HD3	1.66	0.47
1:K:223:PHE:CD1	1:K:259:GLU:CG	2.96	0.47
1:L:37:ARG:CG	1:L:38:PRO:N	2.75	0.47
1:M:34:ILE:HG23	1:M:68:LYS:H	1.79	0.47
1:N:2:GLU:OE2	1:N:2:GLU:HA	2.10	0.47
1:N:37:ARG:CG	1:N:38:PRO:N	2.75	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:288:ASP:OD2	1:P:204:ALA:HB1	2.14	0.47
1:N:335:ARG:O	1:N:338:SER:HB3	2.14	0.47
1:O:58:ALA:HA	1:O:65:LEU:CD2	2.43	0.47
1:P:288:ASP:OD2	1:R:204:ALA:HB1	2.14	0.47
1:R:335:ARG:O	1:R:338:SER:HB3	2.14	0.47
1:U:34:ILE:HG22	1:U:35:VAL:H	1.78	0.47
1:V:147:ARG:CG	1:V:147:ARG:NH2	2.71	0.47
1:W:298:VAL:HG12	1:W:335:ARG:NH1	2.28	0.47
1:A:34:ILE:HD13	1:A:67:LEU:CD1	2.41	0.47
1:C:34:ILE:HG22	1:C:54:VAL:HG21	1.96	0.47
1:D:34:ILE:HG23	1:D:68:LYS:H	1.79	0.47
1:H:369:ILE:HG22	1:H:370:VAL:N	2.29	0.47
1:I:223:PHE:CD1	1:I:259:GLU:CG	2.96	0.47
1:J:58:ALA:HA	1:J:65:LEU:CD2	2.43	0.47
1:K:8:LEU:O	1:K:104:LEU:N	2.47	0.47
1:K:288:ASP:OD2	1:M:204:ALA:HB1	2.14	0.47
1:L:335:ARG:O	1:L:338:SER:HB3	2.13	0.47
1:M:180:LEU:HD12	1:M:181:ALA:O	2.13	0.47
1:O:335:ARG:HA	1:O:335:ARG:HD3	1.39	0.47
1:R:288:ASP:OD2	1:T:204:ALA:HB1	2.14	0.47
1:S:35:VAL:CA	1:S:54:VAL:CG2	2.90	0.47
1:S:270:GLU:CD	1:S:270:GLU:C	2.83	0.47
1:T:180:LEU:HD12	1:T:181:ALA:O	2.14	0.47
1:T:298:VAL:HG12	1:T:335:ARG:NH1	2.28	0.47
1:U:34:ILE:HG22	1:U:54:VAL:HG21	1.96	0.47
1:V:34:ILE:HG23	1:V:68:LYS:H	1.79	0.47
1:V:58:ALA:CB	1:V:65:LEU:CD2	2.90	0.47
1:W:39:ARG:NE	1:W:66:THR:CA	2.69	0.47
1:A:44:MET:CG	1:A:45:VAL:N	2.75	0.47
1:A:236:LEU:HD11	1:A:237:GLU:HG2	1.96	0.47
1:A:288:ASP:OD2	1:C:204:ALA:HB1	2.14	0.47
1:C:167:GLU:CD	1:E:61:LYS:CE	2.85	0.47
1:C:180:LEU:HD12	1:C:181:ALA:O	2.13	0.47
1:C:236:LEU:HD11	1:C:237:GLU:HG2	1.96	0.47
1:E:34:ILE:HG22	1:E:35:VAL:H	1.78	0.47
1:E:223:PHE:CD1	1:E:259:GLU:CG	2.96	0.47
1:F:142:LEU:HD21	1:F:165:ILE:CD1	2.38	0.47
1:G:270:GLU:CD	1:G:270:GLU:C	2.83	0.47
1:H:143:TYR:CZ	1:H:345:ILE:CG2	2.98	0.47
1:J:44:MET:CG	1:J:45:VAL:N	2.75	0.47
1:J:336:LYS:HE2	2:J:401:ADP:C5'	2.38	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:298:VAL:HG12	1:N:335:ARG:NH1	2.28	0.47
1:P:143:TYR:CZ	1:P:345:ILE:CG2	2.98	0.47
1:P:369:ILE:HG22	1:P:370:VAL:N	2.29	0.47
1:Q:270:GLU:CD	1:Q:270:GLU:C	2.83	0.47
1:R:34:ILE:HD13	1:R:67:LEU:CD1	2.41	0.47
1:R:34:ILE:HG23	1:R:68:LYS:H	1.79	0.47
1:U:34:ILE:HG23	1:U:68:LYS:H	1.79	0.47
1:V:8:LEU:HB2	1:V:103:THR:HA	1.95	0.47
1:V:34:ILE:HG22	1:V:54:VAL:HG21	1.96	0.47
1:W:143:TYR:CZ	1:W:345:ILE:CG2	2.98	0.47
1:A:35:VAL:CA	1:A:54:VAL:CG2	2.90	0.47
1:A:143:TYR:CZ	1:A:345:ILE:CG2	2.98	0.47
1:E:270:GLU:CD	1:E:270:GLU:C	2.83	0.47
1:G:223:PHE:CD1	1:G:259:GLU:CG	2.96	0.47
1:I:34:ILE:HG23	1:I:68:LYS:H	1.79	0.47
1:I:288:ASP:OD2	1:K:204:ALA:HB1	2.14	0.47
1:J:300:SER:HA	1:J:335:ARG:HG2	1.89	0.47
1:J:369:ILE:HG22	1:J:370:VAL:N	2.29	0.47
1:K:8:LEU:HB2	1:K:103:THR:HA	1.95	0.47
1:K:132:MET:HE2	1:K:132:MET:HB2	1.75	0.47
1:K:143:TYR:HD1	1:K:143:TYR:C	2.09	0.47
1:K:298:VAL:HG12	1:K:335:ARG:NH1	2.28	0.47
1:M:288:ASP:OD2	1:O:204:ALA:HB1	2.14	0.47
1:O:270:GLU:CD	1:O:270:GLU:C	2.83	0.47
1:R:34:ILE:CG2	1:R:67:LEU:HD22	2.28	0.47
1:S:34:ILE:HG22	1:S:35:VAL:H	1.78	0.47
1:U:270:GLU:CD	1:U:270:GLU:C	2.83	0.47
1:A:369:ILE:HG22	1:A:370:VAL:N	2.29	0.47
1:B:61:LYS:HG2	1:B:64:ILE:CG2	2.44	0.47
1:B:369:ILE:HG22	1:B:370:VAL:N	2.29	0.47
1:C:288:ASP:OD2	1:E:204:ALA:HB1	2.14	0.47
1:C:335:ARG:O	1:C:338:SER:HB3	2.14	0.47
1:E:300:SER:HA	1:E:335:ARG:HG2	1.89	0.47
1:F:288:ASP:OD1	1:H:208:ILE:HD12	2.15	0.47
1:H:176:MET:HE3	1:H:176:MET:HB3	1.81	0.47
1:H:236:LEU:HD11	1:H:237:GLU:HG2	1.96	0.47
1:H:288:ASP:OD1	1:J:208:ILE:HD12	2.15	0.47
1:I:270:GLU:CD	1:I:270:GLU:C	2.83	0.47
1:J:2:GLU:OE2	1:J:2:GLU:HA	2.10	0.47
1:J:143:TYR:CZ	1:J:345:ILE:CG2	2.98	0.47
1:K:270:GLU:CD	1:K:270:GLU:C	2.83	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:369:ILE:HG22	1:L:370:VAL:N	2.29	0.47
1:M:270:GLU:CD	1:M:270:GLU:C	2.83	0.47
1:M:369:ILE:HG22	1:M:370:VAL:N	2.29	0.47
1:N:270:GLU:CD	1:N:270:GLU:C	2.83	0.47
1:O:43:VAL:O	1:O:44:MET:HG2	2.13	0.47
1:R:143:TYR:CZ	1:R:345:ILE:CG2	2.98	0.47
1:R:287:ILE:HD12	1:T:208:ILE:HD13	1.96	0.47
1:T:169:TYR:CD1	1:V:42:GLY:HA3	2.50	0.47
1:U:143:TYR:CZ	1:U:345:ILE:CG2	2.98	0.47
1:A:37:ARG:O	1:A:66:THR:CG2	2.63	0.47
1:B:34:ILE:HG22	1:B:54:VAL:HG21	1.95	0.47
1:C:8:LEU:O	1:C:104:LEU:N	2.47	0.47
1:C:270:GLU:CD	1:C:270:GLU:C	2.83	0.47
1:C:369:ILE:HG22	1:C:370:VAL:N	2.29	0.47
1:E:288:ASP:OD1	1:G:208:ILE:HD12	2.15	0.47
1:J:14:SER:HB2	1:J:183:ARG:HH22	1.80	0.47
1:K:288:ASP:OD1	1:M:208:ILE:HD12	2.15	0.47
1:L:270:GLU:CD	1:L:270:GLU:C	2.83	0.47
1:N:143:TYR:CZ	1:N:345:ILE:CG2	2.98	0.47
1:N:176:MET:HE3	1:N:176:MET:HB3	1.81	0.47
1:O:8:LEU:O	1:O:104:LEU:N	2.47	0.47
1:O:167:GLU:CD	1:Q:61:LYS:CE	2.85	0.47
1:P:270:GLU:CD	1:P:270:GLU:C	2.83	0.47
1:Q:288:ASP:OD1	1:S:208:ILE:HD12	2.15	0.47
1:R:169:TYR:CD1	1:T:42:GLY:HA3	2.50	0.47
1:S:14:SER:HB2	1:S:183:ARG:HH22	1.80	0.47
1:S:143:TYR:CZ	1:S:345:ILE:CG2	2.98	0.47
1:S:167:GLU:CD	1:U:61:LYS:CE	2.85	0.47
1:U:14:SER:HB2	1:U:183:ARG:HH22	1.80	0.47
1:A:336:LYS:HE2	2:A:401:ADP:C5'	2.38	0.47
1:B:14:SER:HB2	1:B:183:ARG:HH22	1.80	0.47
1:B:134:VAL:O	1:B:375:PHE:OXT	2.33	0.47
1:B:288:ASP:OD1	1:D:208:ILE:HD12	2.15	0.47
1:C:288:ASP:OD1	1:E:208:ILE:HD12	2.15	0.47
1:C:369:ILE:HG23	1:C:370:VAL:N	2.30	0.47
1:D:134:VAL:O	1:D:375:PHE:OXT	2.33	0.47
1:E:14:SER:HB2	1:E:183:ARG:HH22	1.80	0.47
1:E:34:ILE:HG22	1:E:54:VAL:HG21	1.96	0.47
1:E:335:ARG:O	1:E:338:SER:HB3	2.13	0.47
1:F:134:VAL:O	1:F:375:PHE:OXT	2.33	0.47
1:F:143:TYR:CZ	1:F:345:ILE:CG2	2.98	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:369:ILE:HG22	1:F:370:VAL:N	2.29	0.47
1:G:61:LYS:HG2	1:G:64:ILE:CG2	2.45	0.47
1:G:288:ASP:OD1	1:I:208:ILE:HD12	2.15	0.47
1:H:14:SER:HB2	1:H:183:ARG:HH22	1.80	0.47
1:H:134:VAL:O	1:H:375:PHE:OXT	2.33	0.47
1:H:288:ASP:OD2	1:J:204:ALA:HB1	2.14	0.47
1:I:34:ILE:HG22	1:I:54:VAL:HG21	1.95	0.47
1:I:61:LYS:HG2	1:I:64:ILE:CG2	2.45	0.47
1:J:31:PHE:HA	1:J:32:PRO:HD3	1.66	0.47
1:J:34:ILE:HG22	1:J:54:VAL:HG21	1.96	0.47
1:J:37:ARG:O	1:J:66:THR:CG2	2.63	0.47
1:J:134:VAL:O	1:J:375:PHE:OXT	2.33	0.47
1:J:147:ARG:HH21	1:J:147:ARG:HG3	1.80	0.47
1:J:270:GLU:CD	1:J:270:GLU:C	2.83	0.47
1:K:34:ILE:HG23	1:K:68:LYS:H	1.79	0.47
1:L:14:SER:HB2	1:L:183:ARG:HH22	1.80	0.47
1:L:134:VAL:O	1:L:375:PHE:OXT	2.33	0.47
1:L:143:TYR:CZ	1:L:345:ILE:CG2	2.98	0.47
1:M:34:ILE:HG22	1:M:54:VAL:HG21	1.95	0.47
1:M:288:ASP:OD1	1:O:208:ILE:HD12	2.15	0.47
1:N:34:ILE:CG2	1:N:67:LEU:HD22	2.27	0.47
1:O:34:ILE:HG22	1:O:54:VAL:HG21	1.96	0.47
1:P:8:LEU:O	1:P:104:LEU:N	2.47	0.47
1:P:44:MET:CG	1:P:45:VAL:N	2.76	0.47
1:P:169:TYR:CD1	1:R:42:GLY:HA3	2.50	0.47
1:Q:14:SER:HB2	1:Q:183:ARG:HH22	1.80	0.47
1:Q:37:ARG:CG	1:Q:38:PRO:N	2.75	0.47
1:Q:134:VAL:O	1:Q:375:PHE:OXT	2.33	0.47
1:Q:369:ILE:HG22	1:Q:370:VAL:N	2.29	0.47
1:R:37:ARG:O	1:R:66:THR:CG2	2.63	0.47
1:R:44:MET:CG	1:R:45:VAL:N	2.75	0.47
1:R:167:GLU:CD	1:T:61:LYS:CE	2.85	0.47
1:R:270:GLU:CD	1:R:270:GLU:C	2.83	0.47
1:R:369:ILE:HG23	1:R:370:VAL:N	2.30	0.47
1:S:34:ILE:HG22	1:S:54:VAL:HG21	1.96	0.47
1:S:134:VAL:O	1:S:375:PHE:OXT	2.33	0.47
1:S:288:ASP:OD2	1:U:204:ALA:HB1	2.14	0.47
1:T:31:PHE:HA	1:T:32:PRO:HD3	1.66	0.47
1:T:37:ARG:O	1:T:66:THR:CG2	2.63	0.47
1:T:288:ASP:OD2	1:V:204:ALA:HB1	2.14	0.47
1:U:134:VAL:O	1:U:375:PHE:OXT	2.33	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:236:LEU:HD11	1:V:237:GLU:HG2	1.96	0.47
1:W:134:VAL:O	1:W:375:PHE:OXT	2.33	0.47
1:A:134:VAL:O	1:A:375:PHE:OXT	2.33	0.47
1:A:169:TYR:CD1	1:C:42:GLY:HA3	2.50	0.47
1:B:270:GLU:CD	1:B:270:GLU:C	2.83	0.47
1:C:14:SER:HB2	1:C:183:ARG:HH22	1.80	0.47
1:D:143:TYR:CZ	1:D:345:ILE:CG2	2.98	0.47
1:D:335:ARG:HA	1:D:335:ARG:HD3	1.39	0.47
1:D:369:ILE:HG22	1:D:370:VAL:N	2.29	0.47
1:G:14:SER:HB2	1:G:183:ARG:HH22	1.80	0.47
1:G:73:HIC:HA	1:G:183:ARG:NH1	2.18	0.47
1:G:288:ASP:OD2	1:I:204:ALA:HB1	2.14	0.47
1:G:290:ARG:HB2	1:I:244:ASP:HA	1.97	0.47
1:H:147:ARG:HH21	1:H:147:ARG:HG3	1.80	0.47
1:I:14:SER:HB2	1:I:183:ARG:HH22	1.80	0.47
1:I:37:ARG:O	1:I:66:THR:CG2	2.63	0.47
1:J:236:LEU:HD11	1:J:237:GLU:HG2	1.96	0.47
1:L:37:ARG:O	1:L:66:THR:CG2	2.63	0.47
1:L:61:LYS:HG2	1:L:64:ILE:CG2	2.45	0.47
1:L:132:MET:HE2	1:L:132:MET:HB2	1.75	0.47
1:L:369:ILE:HG23	1:L:370:VAL:N	2.30	0.47
1:M:8:LEU:O	1:M:104:LEU:N	2.47	0.47
1:M:134:VAL:O	1:M:375:PHE:OXT	2.33	0.47
1:N:134:VAL:O	1:N:375:PHE:OXT	2.33	0.47
1:N:288:ASP:OD1	1:P:208:ILE:HD12	2.15	0.47
1:O:134:VAL:O	1:O:375:PHE:OXT	2.33	0.47
1:O:288:ASP:OD2	1:Q:204:ALA:HB1	2.14	0.47
1:P:287:ILE:HD12	1:R:208:ILE:HD13	1.96	0.47
1:P:298:VAL:HG12	1:P:335:ARG:NH1	2.28	0.47
1:Q:34:ILE:HG22	1:Q:35:VAL:H	1.78	0.47
1:Q:143:TYR:CZ	1:Q:345:ILE:CG2	2.97	0.47
1:R:288:ASP:OD1	1:T:208:ILE:HD12	2.15	0.47
1:R:290:ARG:HB2	1:T:244:ASP:HA	1.97	0.47
1:R:298:VAL:HG12	1:R:335:ARG:NH1	2.28	0.47
1:S:58:ALA:CB	1:S:65:LEU:CD2	2.90	0.47
1:S:288:ASP:OD1	1:U:208:ILE:HD12	2.15	0.47
1:S:369:ILE:HG22	1:S:370:VAL:N	2.29	0.47
1:T:58:ALA:CB	1:T:65:LEU:CD2	2.90	0.47
1:U:37:ARG:O	1:U:66:THR:CG2	2.63	0.47
1:V:44:MET:CG	1:V:45:VAL:N	2.75	0.47
1:W:270:GLU:CD	1:W:270:GLU:C	2.83	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:34:ILE:HG23	1:A:68:LYS:H	1.79	0.47
1:A:290:ARG:HB2	1:C:244:ASP:HA	1.97	0.47
1:D:34:ILE:HG22	1:D:54:VAL:HG21	1.96	0.47
1:D:270:GLU:CD	1:D:270:GLU:C	2.83	0.47
1:E:143:TYR:CZ	1:E:345:ILE:CG2	2.98	0.47
1:F:73:HIC:HA	1:F:183:ARG:NH1	2.18	0.47
1:F:287:ILE:HD12	1:H:208:ILE:HD13	1.96	0.47
1:G:8:LEU:O	1:G:104:LEU:N	2.47	0.47
1:G:34:ILE:HG22	1:G:54:VAL:HG21	1.96	0.47
1:G:37:ARG:O	1:G:66:THR:CG2	2.63	0.47
1:G:143:TYR:CZ	1:G:345:ILE:CG2	2.98	0.47
1:I:169:TYR:CD1	1:K:42:GLY:HA3	2.50	0.47
1:K:34:ILE:HG22	1:K:54:VAL:HG21	1.95	0.47
1:K:287:ILE:HD12	1:M:208:ILE:HD13	1.96	0.47
1:L:288:ASP:OD1	1:N:208:ILE:HD12	2.15	0.47
1:M:143:TYR:CZ	1:M:345:ILE:CG2	2.97	0.47
1:N:236:LEU:HD11	1:N:237:GLU:HG2	1.96	0.47
1:O:14:SER:HB2	1:O:183:ARG:HH22	1.80	0.47
1:O:287:ILE:HD12	1:Q:208:ILE:HD13	1.96	0.47
1:O:369:ILE:HG22	1:O:370:VAL:N	2.29	0.47
1:Q:61:LYS:HG2	1:Q:64:ILE:CG2	2.45	0.47
1:Q:106:THR:HB	1:Q:137:GLN:HG2	1.97	0.47
1:T:34:ILE:HG23	1:T:68:LYS:H	1.79	0.47
1:B:34:ILE:HG23	1:B:68:LYS:H	1.79	0.47
1:B:335:ARG:HA	1:B:335:ARG:HD3	1.39	0.47
1:C:34:ILE:HD13	1:C:67:LEU:CD1	2.41	0.47
1:C:37:ARG:O	1:C:66:THR:CG2	2.63	0.47
1:D:288:ASP:OD1	1:F:208:ILE:HD12	2.15	0.47
1:D:298:VAL:HG12	1:D:335:ARG:NH1	2.28	0.47
1:E:288:ASP:OD2	1:G:204:ALA:HB1	2.14	0.47
1:H:37:ARG:O	1:H:66:THR:CG2	2.63	0.47
1:J:173:HIS:HE1	1:K:268:GLY:CA	2.28	0.47
1:K:61:LYS:HG2	1:K:64:ILE:CG2	2.45	0.47
1:K:143:TYR:CZ	1:K:345:ILE:CG2	2.98	0.47
1:L:34:ILE:HG22	1:L:35:VAL:H	1.78	0.47
1:L:236:LEU:HD11	1:L:237:GLU:HG2	1.96	0.47
1:N:39:ARG:NE	1:N:66:THR:CA	2.69	0.47
1:N:44:MET:CG	1:N:45:VAL:N	2.76	0.47
1:N:61:LYS:HG2	1:N:64:ILE:CG2	2.45	0.47
1:P:134:VAL:O	1:P:375:PHE:OXT	2.33	0.47
1:P:288:ASP:OD1	1:R:208:ILE:HD12	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:34:ILE:HG22	1:Q:54:VAL:HG21	1.95	0.47
1:S:106:THR:HB	1:S:137:GLN:HG2	1.97	0.47
1:T:335:ARG:HA	1:T:335:ARG:HD3	1.39	0.47
1:V:14:SER:HB2	1:V:183:ARG:HH22	1.80	0.47
1:W:44:MET:CG	1:W:45:VAL:N	2.75	0.47
1:W:349:LEU:HD23	1:W:349:LEU:HA	1.80	0.47
1:A:14:SER:HB2	1:A:183:ARG:HH22	1.80	0.46
1:A:44:MET:CG	1:A:45:VAL:H	2.18	0.46
1:A:143:TYR:CE2	1:A:346:LEU:HD13	2.51	0.46
1:A:152:VAL:HG22	1:A:298:VAL:HB	1.98	0.46
1:A:270:GLU:CD	1:A:270:GLU:C	2.83	0.46
1:B:152:VAL:HG22	1:B:298:VAL:HB	1.98	0.46
1:C:134:VAL:O	1:C:375:PHE:OXT	2.33	0.46
1:C:169:TYR:CD1	1:E:42:GLY:HA3	2.50	0.46
1:D:14:SER:HB2	1:D:183:ARG:HH22	1.80	0.46
1:D:73:HIC:HA	1:D:183:ARG:NH1	2.18	0.46
1:E:287:ILE:HD12	1:G:208:ILE:HD13	1.96	0.46
1:F:2:GLU:OE2	1:F:2:GLU:HA	2.10	0.46
1:F:58:ALA:CB	1:F:65:LEU:CD2	2.90	0.46
1:G:287:ILE:HD12	1:I:208:ILE:HD13	1.96	0.46
1:H:35:VAL:C	1:H:54:VAL:CG2	2.88	0.46
1:H:270:GLU:CD	1:H:270:GLU:C	2.83	0.46
1:I:143:TYR:CZ	1:I:345:ILE:CG2	2.98	0.46
1:I:369:ILE:HG22	1:I:370:VAL:N	2.29	0.46
1:J:35:VAL:C	1:J:54:VAL:CG2	2.88	0.46
1:J:58:ALA:CB	1:J:65:LEU:CD2	2.90	0.46
1:J:132:MET:HE2	1:J:132:MET:HB2	1.74	0.46
1:K:134:VAL:O	1:K:375:PHE:OXT	2.33	0.46
1:K:169:TYR:CD1	1:M:42:GLY:HA3	2.50	0.46
1:L:288:ASP:OD2	1:N:204:ALA:HB1	2.14	0.46
1:L:361:GLU:HB3	1:L:369:ILE:CD1	2.14	0.46
1:N:8:LEU:O	1:N:104:LEU:N	2.47	0.46
1:N:14:SER:HB2	1:N:183:ARG:HH22	1.80	0.46
1:O:152:VAL:HG22	1:O:298:VAL:HB	1.98	0.46
1:Q:35:VAL:C	1:Q:54:VAL:CG2	2.88	0.46
1:Q:147:ARG:HH21	1:Q:147:ARG:HG3	1.80	0.46
1:R:8:LEU:O	1:R:104:LEU:N	2.47	0.46
1:T:35:VAL:C	1:T:54:VAL:CG2	2.88	0.46
1:T:270:GLU:CD	1:T:270:GLU:C	2.83	0.46
1:U:106:THR:HB	1:U:137:GLN:HG2	1.97	0.46
1:W:14:SER:HB2	1:W:183:ARG:HH22	1.80	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:336:LYS:HE2	2:W:401:ADP:C5'	2.38	0.46
1:A:147:ARG:HH21	1:A:147:ARG:HG3	1.80	0.46
1:B:106:THR:HB	1:B:137:GLN:HG2	1.97	0.46
1:C:143:TYR:CE2	1:C:346:LEU:HD13	2.51	0.46
1:C:336:LYS:HE2	2:C:401:ADP:C5'	2.38	0.46
1:E:61:LYS:HG2	1:E:64:ILE:CG2	2.45	0.46
1:F:143:TYR:CE2	1:F:346:LEU:HD13	2.51	0.46
1:F:270:GLU:CD	1:F:270:GLU:C	2.83	0.46
1:H:143:TYR:CE2	1:H:346:LEU:HD13	2.51	0.46
1:I:58:ALA:CB	1:I:65:LEU:CD2	2.90	0.46
1:J:61:LYS:HG2	1:J:64:ILE:CG2	2.45	0.46
1:J:152:VAL:HG22	1:J:298:VAL:HB	1.98	0.46
1:J:223:PHE:CD1	1:J:259:GLU:CG	2.96	0.46
1:K:44:MET:CG	1:K:45:VAL:N	2.75	0.46
1:L:35:VAL:C	1:L:54:VAL:CG2	2.88	0.46
1:L:290:ARG:HB2	1:N:244:ASP:HA	1.97	0.46
1:N:152:VAL:HG22	1:N:298:VAL:HB	1.98	0.46
1:N:369:ILE:HG23	1:N:370:VAL:N	2.30	0.46
1:O:35:VAL:C	1:O:54:VAL:CG2	2.88	0.46
1:O:147:ARG:HH21	1:O:147:ARG:HG3	1.80	0.46
1:P:37:ARG:O	1:P:66:THR:CG2	2.63	0.46
1:Q:152:VAL:HG22	1:Q:298:VAL:HB	1.98	0.46
1:R:35:VAL:C	1:R:54:VAL:CG2	2.88	0.46
1:R:58:ALA:CB	1:R:65:LEU:CD2	2.90	0.46
1:S:35:VAL:C	1:S:54:VAL:CG2	2.88	0.46
1:S:61:LYS:HG2	1:S:64:ILE:CG2	2.45	0.46
1:U:44:MET:CG	1:U:45:VAL:N	2.75	0.46
1:U:152:VAL:HG22	1:U:298:VAL:HB	1.98	0.46
1:V:35:VAL:C	1:V:54:VAL:CG2	2.89	0.46
1:V:143:TYR:CZ	1:V:345:ILE:CG2	2.98	0.46
1:B:143:TYR:CZ	1:B:345:ILE:CG2	2.98	0.46
1:B:287:ILE:HD12	1:D:208:ILE:HD13	1.96	0.46
1:C:143:TYR:CZ	1:C:345:ILE:CG2	2.98	0.46
1:C:152:VAL:HG22	1:C:298:VAL:HB	1.98	0.46
1:D:37:ARG:O	1:D:66:THR:CG2	2.63	0.46
1:D:61:LYS:HG2	1:D:64:ILE:CG2	2.45	0.46
1:E:134:VAL:O	1:E:375:PHE:OXT	2.33	0.46
1:F:34:ILE:HG22	1:F:54:VAL:HG21	1.96	0.46
1:F:35:VAL:C	1:F:54:VAL:CG2	2.89	0.46
1:F:167:GLU:CD	1:H:61:LYS:CE	2.85	0.46
1:F:169:TYR:CD1	1:H:42:GLY:HA3	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:169:TYR:CD1	1:I:42:GLY:HA3	2.50	0.46
1:H:223:PHE:CD1	1:H:259:GLU:CG	2.96	0.46
1:I:134:VAL:O	1:I:375:PHE:OXT	2.33	0.46
1:K:167:GLU:CD	1:M:61:LYS:CE	2.85	0.46
1:L:39:ARG:NE	1:L:66:THR:CA	2.69	0.46
1:L:147:ARG:HH21	1:L:147:ARG:HG3	1.80	0.46
1:L:152:VAL:HG22	1:L:298:VAL:HB	1.98	0.46
1:M:143:TYR:CE2	1:M:346:LEU:HD13	2.51	0.46
1:N:169:TYR:CD1	1:P:42:GLY:HA3	2.50	0.46
1:O:143:TYR:CZ	1:O:345:ILE:CG2	2.98	0.46
1:O:143:TYR:CE2	1:O:346:LEU:HD13	2.51	0.46
1:O:288:ASP:OD1	1:Q:208:ILE:HD12	2.15	0.46
1:P:35:VAL:C	1:P:54:VAL:CG2	2.88	0.46
1:P:290:ARG:HB2	1:R:244:ASP:HA	1.97	0.46
1:Q:8:LEU:O	1:Q:104:LEU:N	2.47	0.46
1:R:134:VAL:O	1:R:375:PHE:OXT	2.33	0.46
1:R:173:HIS:HE1	1:S:268:GLY:CA	2.28	0.46
1:R:369:ILE:HG22	1:R:370:VAL:N	2.29	0.46
1:S:37:ARG:O	1:S:66:THR:CG2	2.63	0.46
1:T:44:MET:CG	1:T:45:VAL:N	2.76	0.46
1:T:61:LYS:HG2	1:T:64:ILE:CG2	2.45	0.46
1:T:143:TYR:CZ	1:T:345:ILE:CG2	2.98	0.46
1:U:2:GLU:OE2	1:U:2:GLU:HA	2.10	0.46
1:U:61:LYS:HG2	1:U:64:ILE:CG2	2.45	0.46
1:V:270:GLU:CD	1:V:270:GLU:C	2.83	0.46
1:W:37:ARG:O	1:W:66:THR:CG2	2.63	0.46
1:A:35:VAL:C	1:A:54:VAL:CG2	2.89	0.46
1:B:369:ILE:HG23	1:B:370:VAL:N	2.30	0.46
1:C:39:ARG:NE	1:C:66:THR:CA	2.69	0.46
1:D:152:VAL:HG22	1:D:298:VAL:HB	1.98	0.46
1:D:169:TYR:CD1	1:F:42:GLY:HA3	2.50	0.46
1:F:14:SER:HB2	1:F:183:ARG:HH22	1.80	0.46
1:G:134:VAL:O	1:G:375:PHE:OXT	2.33	0.46
1:H:152:VAL:HG22	1:H:298:VAL:HB	1.98	0.46
1:H:300:SER:HA	1:H:335:ARG:HG2	1.89	0.46
1:I:9:VAL:HG21	1:I:344:SER:HA	1.98	0.46
1:I:34:ILE:HD13	1:I:67:LEU:CD2	2.46	0.46
1:I:132:MET:HE2	1:I:132:MET:HB2	1.75	0.46
1:J:143:TYR:CE2	1:J:346:LEU:HD13	2.51	0.46
1:M:152:VAL:HG22	1:M:298:VAL:HB	1.98	0.46
1:N:34:ILE:HD13	1:N:67:LEU:CD1	2.41	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:236:LEU:CD1	1:N:237:GLU:CG	2.93	0.46
1:O:37:ARG:O	1:O:66:THR:CG2	2.63	0.46
1:O:106:THR:HB	1:O:137:GLN:HG2	1.98	0.46
1:T:9:VAL:HG21	1:T:344:SER:HA	1.98	0.46
1:T:134:VAL:O	1:T:375:PHE:OXT	2.33	0.46
1:T:236:LEU:HD11	1:T:237:GLU:HG2	1.96	0.46
1:T:288:ASP:OD1	1:V:208:ILE:HD12	2.15	0.46
1:U:35:VAL:C	1:U:54:VAL:CG2	2.88	0.46
1:U:58:ALA:CB	1:U:65:LEU:CD2	2.90	0.46
1:U:369:ILE:HG22	1:U:370:VAL:N	2.29	0.46
1:V:106:THR:HB	1:V:137:GLN:HG2	1.97	0.46
1:V:134:VAL:O	1:V:375:PHE:OXT	2.33	0.46
1:W:106:THR:HB	1:W:137:GLN:HG2	1.97	0.46
1:W:236:LEU:HD11	1:W:237:GLU:HG2	1.96	0.46
1:W:369:ILE:HG23	1:W:370:VAL:N	2.30	0.46
1:B:8:LEU:O	1:B:104:LEU:N	2.47	0.46
1:C:9:VAL:HG21	1:C:344:SER:HA	1.98	0.46
1:C:173:HIS:HE1	1:D:268:GLY:CA	2.28	0.46
1:D:8:LEU:O	1:D:104:LEU:N	2.47	0.46
1:E:290:ARG:HB2	1:G:244:ASP:HA	1.97	0.46
1:F:152:VAL:HG22	1:F:298:VAL:HB	1.98	0.46
1:F:223:PHE:CD1	1:F:259:GLU:CG	2.96	0.46
1:H:58:ALA:CB	1:H:65:LEU:CD2	2.90	0.46
1:I:287:ILE:HD12	1:K:208:ILE:HD13	1.96	0.46
1:J:8:LEU:O	1:J:104:LEU:N	2.47	0.46
1:L:44:MET:CG	1:L:45:VAL:N	2.75	0.46
1:L:223:PHE:CD1	1:L:259:GLU:CG	2.96	0.46
1:M:35:VAL:C	1:M:54:VAL:CG2	2.88	0.46
1:M:369:ILE:HG23	1:M:370:VAL:N	2.30	0.46
1:N:35:VAL:C	1:N:54:VAL:CG2	2.89	0.46
1:N:173:HIS:HE1	1:O:268:GLY:CA	2.28	0.46
1:P:61:LYS:HG2	1:P:64:ILE:CG2	2.45	0.46
1:P:152:VAL:HG22	1:P:298:VAL:HB	1.98	0.46
1:S:152:VAL:HG22	1:S:298:VAL:HB	1.98	0.46
1:T:143:TYR:CE2	1:T:346:LEU:HD13	2.51	0.46
1:U:288:ASP:OD1	1:W:208:ILE:HD12	2.15	0.46
1:V:152:VAL:HG22	1:V:298:VAL:HB	1.98	0.46
1:W:61:LYS:HG2	1:W:64:ILE:CG2	2.45	0.46
1:W:152:VAL:HG22	1:W:298:VAL:HB	1.98	0.46
1:A:288:ASP:OD1	1:C:208:ILE:HD12	2.15	0.46
1:C:35:VAL:C	1:C:54:VAL:CG2	2.88	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:44:MET:CG	1:C:45:VAL:H	2.18	0.46
1:C:147:ARG:HH21	1:C:147:ARG:HG3	1.80	0.46
1:C:290:ARG:HB2	1:E:244:ASP:HA	1.97	0.46
1:D:35:VAL:C	1:D:54:VAL:CG2	2.88	0.46
1:D:143:TYR:CE2	1:D:346:LEU:HD13	2.51	0.46
1:E:173:HIS:HE1	1:F:268:GLY:CA	2.28	0.46
1:G:152:VAL:HG22	1:G:298:VAL:HB	1.98	0.46
1:H:34:ILE:HG22	1:H:54:VAL:HG21	1.95	0.46
1:H:73:HIC:HA	1:H:183:ARG:NH1	2.18	0.46
1:I:152:VAL:HG22	1:I:298:VAL:HB	1.98	0.46
1:I:290:ARG:HB2	1:K:244:ASP:HA	1.97	0.46
1:K:14:SER:HB2	1:K:183:ARG:HH22	1.80	0.46
1:L:336:LYS:HE2	2:L:401:ADP:C5'	2.38	0.46
1:M:106:THR:HB	1:M:137:GLN:HG2	1.97	0.46
1:M:135:ALA:HB3	1:M:140:LEU:HD11	1.98	0.46
1:M:169:TYR:CD1	1:O:42:GLY:HA3	2.50	0.46
1:N:37:ARG:O	1:N:66:THR:CG2	2.63	0.46
1:N:336:LYS:HE2	2:N:401:ADP:C5'	2.38	0.46
1:O:61:LYS:HG2	1:O:64:ILE:CG2	2.45	0.46
1:P:14:SER:HB2	1:P:183:ARG:HH22	1.80	0.46
1:Q:369:ILE:HG23	1:Q:370:VAL:N	2.30	0.46
1:R:143:TYR:CE2	1:R:346:LEU:HD13	2.51	0.46
1:S:176:MET:HE3	1:S:176:MET:HB3	1.81	0.46
1:T:152:VAL:HG22	1:T:298:VAL:HB	1.98	0.46
1:T:167:GLU:CD	1:V:61:LYS:CE	2.85	0.46
1:T:290:ARG:HB2	1:V:244:ASP:HA	1.97	0.46
1:V:37:ARG:O	1:V:66:THR:CG2	2.63	0.46
1:W:35:VAL:C	1:W:54:VAL:CG2	2.88	0.46
1:W:135:ALA:HB3	1:W:140:LEU:HD11	1.98	0.46
1:A:2:GLU:OE2	1:A:2:GLU:HA	2.10	0.46
1:A:33:SER:O	1:A:69:TYR:CD1	2.69	0.46
1:A:135:ALA:HB3	1:A:140:LEU:HD11	1.98	0.46
1:C:61:LYS:HG2	1:C:64:ILE:CG2	2.45	0.46
1:D:106:THR:HB	1:D:137:GLN:HG2	1.97	0.46
1:D:223:PHE:CD1	1:D:259:GLU:CG	2.96	0.46
1:E:143:TYR:CE2	1:E:346:LEU:HD13	2.51	0.46
1:E:169:TYR:CD1	1:G:42:GLY:HA3	2.50	0.46
1:F:135:ALA:HB3	1:F:140:LEU:HD11	1.98	0.46
1:G:34:ILE:CG2	1:G:67:LEU:HD22	2.27	0.46
1:G:39:ARG:NE	1:G:66:THR:CA	2.69	0.46
1:H:169:TYR:CD1	1:J:42:GLY:HA3	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:287:ILE:HB	1:K:244:ASP:HB3	1.97	0.46
1:I:369:ILE:HG23	1:I:370:VAL:N	2.30	0.46
1:J:135:ALA:HB3	1:J:140:LEU:HD11	1.98	0.46
1:J:167:GLU:CD	1:L:61:LYS:CE	2.85	0.46
1:J:288:ASP:OD1	1:L:208:ILE:HD12	2.15	0.46
1:K:35:VAL:C	1:K:54:VAL:CG2	2.88	0.46
1:L:143:TYR:CE2	1:L:346:LEU:HD13	2.51	0.46
1:M:14:SER:HB2	1:M:183:ARG:HH22	1.80	0.46
1:O:169:TYR:CD1	1:Q:42:GLY:HA3	2.50	0.46
1:Q:143:TYR:CE2	1:Q:346:LEU:HD13	2.51	0.46
1:Q:169:TYR:CD1	1:S:42:GLY:HA3	2.50	0.46
1:T:106:THR:HB	1:T:137:GLN:HG2	1.97	0.46
1:T:173:HIS:HE1	1:U:268:GLY:CA	2.28	0.46
1:T:349:LEU:HD23	1:T:349:LEU:HA	1.80	0.46
1:U:39:ARG:NE	1:U:66:THR:CA	2.69	0.46
1:U:236:LEU:HD11	1:U:237:GLU:HG2	1.96	0.46
1:V:33:SER:O	1:V:69:TYR:CD1	2.69	0.46
1:V:143:TYR:CE2	1:V:346:LEU:HD13	2.51	0.46
1:V:144:ALA:HB2	1:V:342:GLY:N	2.31	0.46
1:W:33:SER:O	1:W:69:TYR:CD1	2.69	0.46
1:W:143:TYR:CE2	1:W:346:LEU:HD13	2.51	0.46
1:B:169:TYR:CD1	1:D:42:GLY:HA3	2.50	0.46
1:B:223:PHE:CD1	1:B:259:GLU:CG	2.96	0.46
1:C:135:ALA:HB3	1:C:140:LEU:HD11	1.98	0.46
1:D:135:ALA:HB3	1:D:140:LEU:HD11	1.98	0.46
1:D:287:ILE:HD12	1:F:208:ILE:HD13	1.96	0.46
1:E:33:SER:O	1:E:69:TYR:CD1	2.69	0.46
1:F:8:LEU:O	1:F:104:LEU:N	2.47	0.46
1:F:37:ARG:O	1:F:66:THR:CG2	2.63	0.46
1:H:135:ALA:HB3	1:H:140:LEU:HD11	1.98	0.46
1:I:236:LEU:HD11	1:I:237:GLU:HG2	1.96	0.46
1:I:288:ASP:OD1	1:K:208:ILE:HD12	2.15	0.46
1:J:7:ALA:HB1	1:J:347:ALA:HB1	1.98	0.46
1:K:37:ARG:O	1:K:66:THR:CG2	2.63	0.46
1:K:152:VAL:HG22	1:K:298:VAL:HB	1.98	0.46
1:L:33:SER:O	1:L:69:TYR:CD1	2.69	0.46
1:M:37:ARG:O	1:M:66:THR:CG2	2.63	0.46
1:M:61:LYS:HG2	1:M:64:ILE:CG2	2.45	0.46
1:M:144:ALA:HB2	1:M:342:GLY:N	2.31	0.46
1:M:290:ARG:HB2	1:O:244:ASP:HA	1.97	0.46
1:N:135:ALA:HB3	1:N:140:LEU:HD11	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:135:ALA:HB3	1:O:140:LEU:HD11	1.98	0.46
1:P:33:SER:O	1:P:69:TYR:CD1	2.69	0.46
1:P:135:ALA:HB3	1:P:140:LEU:HD11	1.98	0.46
1:P:336:LYS:HE2	2:P:401:ADP:C5'	2.38	0.46
1:Q:44:MET:CG	1:Q:45:VAL:H	2.18	0.46
1:Q:73:HIC:HA	1:Q:183:ARG:NH1	2.18	0.46
1:R:9:VAL:HG21	1:R:344:SER:HA	1.98	0.46
1:R:152:VAL:HG22	1:R:298:VAL:HB	1.98	0.46
1:T:14:SER:HB2	1:T:183:ARG:HH22	1.80	0.46
1:T:33:SER:O	1:T:69:TYR:CD1	2.69	0.46
1:T:144:ALA:HB2	1:T:342:GLY:N	2.31	0.46
1:T:287:ILE:HB	1:V:244:ASP:HB3	1.97	0.46
1:T:369:ILE:HG23	1:T:370:VAL:N	2.30	0.46
1:U:33:SER:O	1:U:69:TYR:CD1	2.69	0.46
1:U:135:ALA:HB3	1:U:140:LEU:HD11	1.98	0.46
1:U:287:ILE:HB	1:W:244:ASP:HB3	1.97	0.46
1:W:374:CYS:HB2	1:W:375:PHE:H	1.61	0.46
1:B:35:VAL:C	1:B:54:VAL:CG2	2.88	0.46
1:B:37:ARG:O	1:B:66:THR:CG2	2.63	0.46
1:B:44:MET:CG	1:B:45:VAL:H	2.18	0.46
1:B:135:ALA:HB3	1:B:140:LEU:HD11	1.98	0.46
1:B:369:ILE:HG22	1:B:370:VAL:H	1.81	0.46
1:C:300:SER:HA	1:C:335:ARG:HG2	1.89	0.46
1:E:37:ARG:O	1:E:66:THR:CG2	2.63	0.46
1:E:73:HIC:HA	1:E:183:ARG:NH1	2.18	0.46
1:E:152:VAL:HG22	1:E:298:VAL:HB	1.98	0.46
1:E:167:GLU:CD	1:G:61:LYS:CE	2.85	0.46
1:H:61:LYS:HG2	1:H:64:ILE:CG2	2.45	0.46
1:I:7:ALA:HB1	1:I:347:ALA:HB1	1.98	0.46
1:I:35:VAL:C	1:I:54:VAL:CG2	2.88	0.46
1:I:38:PRO:HD3	1:I:49:GLN:HE22	1.81	0.46
1:K:7:ALA:HB1	1:K:347:ALA:HB1	1.98	0.46
1:K:143:TYR:CE2	1:K:346:LEU:HD13	2.51	0.46
1:M:31:PHE:HA	1:M:32:PRO:HD3	1.66	0.46
1:N:147:ARG:HH21	1:N:147:ARG:HG3	1.80	0.46
1:O:34:ILE:HG22	1:O:35:VAL:H	1.78	0.46
1:O:144:ALA:HB2	1:O:342:GLY:N	2.31	0.46
1:P:143:TYR:CE2	1:P:346:LEU:HD13	2.51	0.46
1:R:144:ALA:HB2	1:R:342:GLY:N	2.31	0.46
1:R:223:PHE:CD1	1:R:259:GLU:CG	2.96	0.46
1:R:236:LEU:HD11	1:R:237:GLU:HG2	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:135:ALA:HB3	1:S:140:LEU:HD11	1.98	0.46
1:T:374:CYS:HB2	1:T:375:PHE:H	1.61	0.46
1:V:135:ALA:HB3	1:V:140:LEU:HD11	1.98	0.46
1:A:61:LYS:HG2	1:A:64:ILE:CG2	2.45	0.46
1:A:144:ALA:HB2	1:A:342:GLY:N	2.31	0.46
1:C:34:ILE:HG22	1:C:35:VAL:H	1.78	0.46
1:C:176:MET:HE3	1:C:176:MET:HB3	1.81	0.46
1:E:287:ILE:HB	1:G:244:ASP:HB3	1.98	0.46
1:E:369:ILE:HG23	1:E:370:VAL:N	2.30	0.46
1:G:9:VAL:HG21	1:G:344:SER:HA	1.98	0.46
1:G:135:ALA:HB3	1:G:140:LEU:HD11	1.98	0.46
1:J:34:ILE:HG23	1:J:68:LYS:H	1.79	0.46
1:J:290:ARG:HB2	1:L:244:ASP:HA	1.97	0.46
1:J:369:ILE:HG22	1:J:370:VAL:H	1.81	0.46
1:K:369:ILE:HG22	1:K:370:VAL:H	1.81	0.46
1:L:7:ALA:HB1	1:L:347:ALA:HB1	1.98	0.46
1:L:34:ILE:N	1:L:54:VAL:HG11	2.31	0.46
1:L:167:GLU:CD	1:N:61:LYS:CE	2.85	0.46
1:M:7:ALA:HB1	1:M:347:ALA:HB1	1.98	0.46
1:M:34:ILE:HD13	1:M:67:LEU:CD2	2.46	0.46
1:N:9:VAL:HG21	1:N:344:SER:HA	1.98	0.46
1:N:287:ILE:HD12	1:P:208:ILE:HD13	1.96	0.46
1:N:290:ARG:HB2	1:P:244:ASP:HA	1.97	0.46
1:P:2:GLU:OE2	1:P:2:GLU:HA	2.10	0.46
1:P:58:ALA:CB	1:P:65:LEU:CD2	2.90	0.46
1:R:33:SER:O	1:R:69:TYR:CD1	2.69	0.46
1:R:39:ARG:NE	1:R:66:THR:CA	2.69	0.46
1:R:61:LYS:HG2	1:R:64:ILE:CG2	2.45	0.46
1:R:135:ALA:HB3	1:R:140:LEU:HD11	1.98	0.46
1:R:335:ARG:HA	1:R:335:ARG:HD3	1.39	0.46
1:S:33:SER:O	1:S:69:TYR:CD1	2.69	0.46
1:S:34:ILE:CG2	1:S:67:LEU:HD22	2.28	0.46
1:S:143:TYR:CE2	1:S:346:LEU:HD13	2.51	0.46
1:T:135:ALA:HB3	1:T:140:LEU:HD11	1.98	0.46
1:U:133:TYR:CE2	1:U:375:PHE:HB2	2.51	0.46
1:U:169:TYR:CD1	1:W:42:GLY:HA3	2.50	0.46
1:A:8:LEU:O	1:A:104:LEU:N	2.47	0.45
1:A:34:ILE:HD13	1:A:67:LEU:CD2	2.46	0.45
1:C:144:ALA:HB2	1:C:342:GLY:N	2.31	0.45
1:C:287:ILE:HD12	1:E:208:ILE:HD13	1.96	0.45
1:E:35:VAL:C	1:E:54:VAL:CG2	2.89	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:132:MET:HE2	1:E:132:MET:HB2	1.75	0.45
1:F:349:LEU:HD23	1:F:349:LEU:HA	1.80	0.45
1:G:35:VAL:C	1:G:54:VAL:CG2	2.88	0.45
1:H:7:ALA:HB1	1:H:347:ALA:HB1	1.98	0.45
1:H:369:ILE:HG22	1:H:370:VAL:H	1.81	0.45
1:H:369:ILE:HG23	1:H:370:VAL:N	2.30	0.45
1:I:135:ALA:HB3	1:I:140:LEU:HD11	1.98	0.45
1:J:169:TYR:CD1	1:L:42:GLY:HA3	2.50	0.45
1:K:135:ALA:HB3	1:K:140:LEU:HD11	1.98	0.45
1:L:135:ALA:HB3	1:L:140:LEU:HD11	1.98	0.45
1:L:169:TYR:CD1	1:N:42:GLY:HA3	2.50	0.45
1:L:287:ILE:HG22	1:L:290:ARG:NH1	2.32	0.45
1:N:7:ALA:HB1	1:N:347:ALA:HB1	1.98	0.45
1:N:106:THR:HB	1:N:137:GLN:HG2	1.97	0.45
1:N:223:PHE:CD1	1:N:259:GLU:CG	2.96	0.45
1:O:9:VAL:HG21	1:O:344:SER:HA	1.98	0.45
1:O:287:ILE:HG22	1:O:290:ARG:NH1	2.32	0.45
1:P:144:ALA:HB2	1:P:342:GLY:N	2.31	0.45
1:Q:135:ALA:HB3	1:Q:140:LEU:HD11	1.98	0.45
1:Q:335:ARG:HA	1:Q:335:ARG:HD3	1.39	0.45
1:Q:369:ILE:HG22	1:Q:370:VAL:H	1.81	0.45
1:R:106:THR:HB	1:R:137:GLN:HG2	1.97	0.45
1:R:287:ILE:HB	1:T:244:ASP:HB3	1.97	0.45
1:S:73:HIC:HA	1:S:183:ARG:NH1	2.18	0.45
1:S:169:TYR:CD1	1:U:42:GLY:HA3	2.50	0.45
1:S:236:LEU:CD1	1:S:237:GLU:CG	2.93	0.45
1:S:287:ILE:HD12	1:U:208:ILE:HD13	1.96	0.45
1:S:287:ILE:HB	1:U:244:ASP:HB3	1.97	0.45
1:W:287:ILE:HG22	1:W:290:ARG:NH1	2.32	0.45
1:A:106:THR:HB	1:A:137:GLN:HG2	1.97	0.45
1:B:38:PRO:HD3	1:B:49:GLN:HE22	1.82	0.45
1:B:73:HIC:HA	1:B:183:ARG:NH1	2.18	0.45
1:B:143:TYR:CE2	1:B:346:LEU:HD13	2.51	0.45
1:D:236:LEU:CD1	1:D:237:GLU:CG	2.93	0.45
1:D:369:ILE:HG23	1:D:370:VAL:N	2.30	0.45
1:E:147:ARG:HH21	1:E:147:ARG:HG3	1.80	0.45
1:F:144:ALA:HB2	1:F:342:GLY:N	2.31	0.45
1:G:7:ALA:HB1	1:G:347:ALA:HB1	1.98	0.45
1:G:144:ALA:HB2	1:G:342:GLY:N	2.31	0.45
1:H:34:ILE:N	1:H:54:VAL:HG11	2.31	0.45
1:H:144:ALA:HB2	1:H:342:GLY:N	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:33:SER:O	1:I:69:TYR:CD1	2.69	0.45
1:I:38:PRO:CG	1:I:49:GLN:NE2	2.79	0.45
1:J:38:PRO:HD3	1:J:49:GLN:HE22	1.82	0.45
1:J:106:THR:HB	1:J:137:GLN:HG2	1.97	0.45
1:K:144:ALA:HB2	1:K:342:GLY:N	2.31	0.45
1:K:290:ARG:HB2	1:M:244:ASP:HA	1.97	0.45
1:L:8:LEU:O	1:L:104:LEU:N	2.47	0.45
1:L:106:THR:HB	1:L:137:GLN:HG2	1.97	0.45
1:N:34:ILE:HD13	1:N:67:LEU:CD2	2.46	0.45
1:N:287:ILE:HG22	1:N:290:ARG:NH1	2.32	0.45
1:O:369:ILE:HG22	1:O:370:VAL:H	1.82	0.45
1:P:38:PRO:HD3	1:P:49:GLN:HE22	1.81	0.45
1:P:133:TYR:CE2	1:P:375:PHE:HB2	2.51	0.45
1:Q:37:ARG:O	1:Q:66:THR:CG2	2.63	0.45
1:Q:38:PRO:HD3	1:Q:49:GLN:HE22	1.82	0.45
1:Q:144:ALA:HB2	1:Q:342:GLY:N	2.31	0.45
1:Q:287:ILE:HG22	1:Q:290:ARG:NH1	2.32	0.45
1:S:38:PRO:HD3	1:S:49:GLN:HE22	1.82	0.45
1:S:133:TYR:CE2	1:S:375:PHE:HB2	2.51	0.45
1:S:287:ILE:HG22	1:S:290:ARG:NH1	2.32	0.45
1:V:164:PRO:CG	1:V:174:ALA:CB	2.95	0.45
1:A:9:VAL:HG21	1:A:344:SER:HA	1.98	0.45
1:A:34:ILE:N	1:A:54:VAL:HG11	2.31	0.45
1:C:287:ILE:HB	1:E:244:ASP:HB3	1.97	0.45
1:D:369:ILE:HG22	1:D:370:VAL:H	1.82	0.45
1:E:335:ARG:HA	1:E:335:ARG:HD3	1.39	0.45
1:F:7:ALA:HB1	1:F:347:ALA:HB1	1.98	0.45
1:F:34:ILE:N	1:F:54:VAL:HG11	2.31	0.45
1:F:61:LYS:HG2	1:F:64:ILE:CG2	2.45	0.45
1:F:106:THR:HB	1:F:137:GLN:HG2	1.97	0.45
1:F:324:THR:HG23	1:H:241:GLU:OE2	2.16	0.45
1:F:335:ARG:HA	1:F:335:ARG:HD3	1.39	0.45
1:G:287:ILE:HB	1:I:244:ASP:HB3	1.97	0.45
1:H:33:SER:O	1:H:69:TYR:CD1	2.69	0.45
1:I:144:ALA:HB2	1:I:342:GLY:N	2.31	0.45
1:I:369:ILE:HG22	1:I:370:VAL:H	1.82	0.45
1:J:34:ILE:HG22	1:J:35:VAL:H	1.78	0.45
1:J:287:ILE:HG22	1:J:290:ARG:NH1	2.32	0.45
1:K:9:VAL:HG21	1:K:344:SER:HA	1.98	0.45
1:K:33:SER:O	1:K:69:TYR:CD1	2.69	0.45
1:M:287:ILE:HG22	1:M:290:ARG:NH1	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:73:HIC:HA	1:O:183:ARG:NH1	2.18	0.45
1:P:106:THR:HB	1:P:137:GLN:HG2	1.97	0.45
1:P:223:PHE:CD1	1:P:259:GLU:CG	2.96	0.45
1:P:287:ILE:HB	1:R:244:ASP:HB3	1.97	0.45
1:Q:33:SER:O	1:Q:69:TYR:CD1	2.69	0.45
1:R:180:LEU:CD1	1:R:181:ALA:N	2.78	0.45
1:T:164:PRO:CG	1:T:174:ALA:CB	2.95	0.45
1:T:176:MET:HE3	1:T:176:MET:HB3	1.81	0.45
1:U:287:ILE:HG22	1:U:290:ARG:NH1	2.32	0.45
1:V:38:PRO:HD3	1:V:49:GLN:HE22	1.81	0.45
1:W:133:TYR:CE2	1:W:375:PHE:HB2	2.51	0.45
1:W:236:LEU:CD1	1:W:237:GLU:CG	2.93	0.45
1:W:369:ILE:HG22	1:W:370:VAL:H	1.82	0.45
1:A:38:PRO:HD3	1:A:49:GLN:HE22	1.81	0.45
1:A:167:GLU:CD	1:C:61:LYS:CE	2.85	0.45
1:A:287:ILE:HG22	1:A:290:ARG:NH1	2.32	0.45
1:B:290:ARG:HB2	1:D:244:ASP:HA	1.97	0.45
1:C:38:PRO:HD3	1:C:49:GLN:HE22	1.81	0.45
1:C:287:ILE:HG22	1:C:290:ARG:NH1	2.32	0.45
1:D:38:PRO:HD3	1:D:49:GLN:HE22	1.82	0.45
1:D:144:ALA:HB2	1:D:342:GLY:N	2.31	0.45
1:E:116:ARG:NH1	1:E:375:PHE:HA	2.32	0.45
1:E:135:ALA:HB3	1:E:140:LEU:HD11	1.98	0.45
1:E:287:ILE:HG22	1:E:290:ARG:NH1	2.32	0.45
1:F:38:PRO:CG	1:F:49:GLN:NE2	2.79	0.45
1:F:290:ARG:HB2	1:H:244:ASP:HA	1.97	0.45
1:F:369:ILE:HG22	1:F:370:VAL:H	1.81	0.45
1:F:369:ILE:HG23	1:F:370:VAL:N	2.30	0.45
1:G:38:PRO:HD3	1:G:49:GLN:HE22	1.82	0.45
1:G:335:ARG:HA	1:G:335:ARG:HD3	1.39	0.45
1:H:287:ILE:HG22	1:H:290:ARG:NH1	2.32	0.45
1:K:106:THR:HB	1:K:137:GLN:HG2	1.97	0.45
1:K:287:ILE:HB	1:M:244:ASP:HB3	1.98	0.45
1:M:33:SER:O	1:M:69:TYR:CD1	2.69	0.45
1:M:34:ILE:N	1:M:54:VAL:HG11	2.31	0.45
1:N:34:ILE:N	1:N:54:VAL:HG11	2.31	0.45
1:N:116:ARG:NH1	1:N:375:PHE:HA	2.32	0.45
1:N:143:TYR:CE2	1:N:346:LEU:HD13	2.51	0.45
1:O:7:ALA:HB1	1:O:347:ALA:HB1	1.98	0.45
1:O:33:SER:O	1:O:69:TYR:CD1	2.69	0.45
1:P:236:LEU:HD11	1:P:237:GLU:HG2	1.96	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:14:SER:HB2	1:R:183:ARG:HH22	1.80	0.45
1:R:38:PRO:HD3	1:R:49:GLN:HE22	1.81	0.45
1:R:287:ILE:HG22	1:R:290:ARG:NH1	2.32	0.45
1:S:324:THR:HG23	1:U:241:GLU:OE2	2.16	0.45
1:T:287:ILE:HG22	1:T:290:ARG:NH1	2.32	0.45
1:U:34:ILE:N	1:U:54:VAL:HG11	2.31	0.45
1:U:143:TYR:CE2	1:U:346:LEU:HD13	2.51	0.45
1:U:369:ILE:HG22	1:U:370:VAL:H	1.82	0.45
1:V:9:VAL:HG21	1:V:344:SER:HA	1.98	0.45
1:V:369:ILE:HG22	1:V:370:VAL:H	1.82	0.45
1:W:8:LEU:O	1:W:104:LEU:N	2.47	0.45
1:A:173:HIS:HE1	1:B:268:GLY:CA	2.28	0.45
1:B:116:ARG:NH1	1:B:375:PHE:HA	2.32	0.45
1:B:164:PRO:CG	1:B:174:ALA:CB	2.95	0.45
1:C:33:SER:O	1:C:69:TYR:CD1	2.69	0.45
1:C:73:HIC:HA	1:C:183:ARG:NH1	2.18	0.45
1:C:116:ARG:NH1	1:C:375:PHE:HA	2.32	0.45
1:C:233:SER:HB3	1:C:236:LEU:HG	1.99	0.45
1:D:7:ALA:HB1	1:D:347:ALA:HB1	1.98	0.45
1:E:58:ALA:CB	1:E:65:LEU:CD2	2.90	0.45
1:E:324:THR:HG23	1:G:241:GLU:OE2	2.16	0.45
1:F:287:ILE:HG22	1:F:290:ARG:NH1	2.32	0.45
1:G:33:SER:O	1:G:69:TYR:CD1	2.69	0.45
1:G:116:ARG:NH1	1:G:375:PHE:HA	2.32	0.45
1:G:143:TYR:CE2	1:G:346:LEU:HD13	2.51	0.45
1:G:369:ILE:HG22	1:G:370:VAL:H	1.81	0.45
1:H:8:LEU:O	1:H:104:LEU:N	2.47	0.45
1:H:106:THR:HB	1:H:137:GLN:HG2	1.97	0.45
1:H:324:THR:HG23	1:J:241:GLU:OE2	2.16	0.45
1:I:176:MET:HE3	1:I:176:MET:HB3	1.81	0.45
1:J:144:ALA:HB2	1:J:342:GLY:N	2.31	0.45
1:L:2:GLU:OE2	1:L:2:GLU:HA	2.11	0.45
1:L:236:LEU:CD1	1:L:237:GLU:CG	2.93	0.45
1:M:9:VAL:HG21	1:M:344:SER:HA	1.98	0.45
1:N:33:SER:O	1:N:69:TYR:CD1	2.69	0.45
1:P:7:ALA:HB1	1:P:347:ALA:HB1	1.98	0.45
1:Q:287:ILE:HB	1:S:244:ASP:HB3	1.97	0.45
1:T:34:ILE:HD13	1:T:67:LEU:CD2	2.46	0.45
1:T:324:THR:HG23	1:V:241:GLU:OE2	2.16	0.45
1:U:73:HIC:HA	1:U:183:ARG:NH1	2.18	0.45
1:U:144:ALA:HB2	1:U:342:GLY:N	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:324:THR:HG23	1:W:241:GLU:OE2	2.16	0.45
1:U:349:LEU:HD23	1:U:349:LEU:HA	1.80	0.45
1:W:116:ARG:NH1	1:W:375:PHE:HA	2.32	0.45
1:W:144:ALA:HB2	1:W:342:GLY:N	2.31	0.45
1:B:2:GLU:OE2	1:B:2:GLU:HA	2.10	0.45
1:B:287:ILE:HG22	1:B:290:ARG:NH1	2.32	0.45
1:C:106:THR:HB	1:C:137:GLN:HG2	1.97	0.45
1:C:180:LEU:CD1	1:C:181:ALA:N	2.78	0.45
1:D:9:VAL:HG21	1:D:344:SER:HA	1.98	0.45
1:D:116:ARG:NH1	1:D:375:PHE:HA	2.32	0.45
1:D:287:ILE:HG22	1:D:290:ARG:NH1	2.32	0.45
1:D:324:THR:HG23	1:F:241:GLU:OE2	2.16	0.45
1:G:164:PRO:CG	1:G:174:ALA:CB	2.95	0.45
1:H:290:ARG:HB2	1:J:244:ASP:HA	1.97	0.45
1:K:38:PRO:HD3	1:K:49:GLN:HE22	1.81	0.45
1:K:287:ILE:HG22	1:K:290:ARG:NH1	2.32	0.45
1:L:38:PRO:HD3	1:L:49:GLN:HE22	1.81	0.45
1:L:116:ARG:NH1	1:L:375:PHE:HA	2.32	0.45
1:M:335:ARG:HA	1:M:335:ARG:HD3	1.39	0.45
1:M:369:ILE:HG22	1:M:370:VAL:H	1.82	0.45
1:N:144:ALA:HB2	1:N:342:GLY:N	2.31	0.45
1:O:31:PHE:HA	1:O:32:PRO:HD3	1.65	0.45
1:P:34:ILE:N	1:P:54:VAL:HG11	2.31	0.45
1:P:180:LEU:CD1	1:P:181:ALA:N	2.78	0.45
1:Q:2:GLU:OE2	1:Q:2:GLU:HA	2.11	0.45
1:Q:290:ARG:HB2	1:S:244:ASP:HA	1.97	0.45
1:Q:324:THR:HG23	1:S:241:GLU:OE2	2.17	0.45
1:S:8:LEU:O	1:S:104:LEU:N	2.47	0.45
1:S:369:ILE:HG22	1:S:370:VAL:H	1.82	0.45
1:U:116:ARG:NH1	1:U:375:PHE:HA	2.32	0.45
1:W:110:LEU:CD1	1:W:177:ARG:NH1	2.80	0.45
1:A:180:LEU:CD1	1:A:181:ALA:N	2.78	0.45
1:A:233:SER:HB3	1:A:236:LEU:HG	1.99	0.45
1:B:7:ALA:HB1	1:B:347:ALA:HB1	1.98	0.45
1:C:34:ILE:N	1:C:54:VAL:HG11	2.31	0.45
1:C:369:ILE:HG22	1:C:370:VAL:H	1.81	0.45
1:E:7:ALA:HB1	1:E:347:ALA:HB1	1.98	0.45
1:E:9:VAL:HG21	1:E:344:SER:HA	1.98	0.45
1:E:106:THR:HB	1:E:137:GLN:HG2	1.98	0.45
1:E:167:GLU:OE1	1:G:61:LYS:CD	2.65	0.45
1:E:233:SER:HB3	1:E:236:LEU:HG	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:106:THR:HB	1:G:137:GLN:HG2	1.97	0.45
1:G:132:MET:HE2	1:G:132:MET:HB2	1.75	0.45
1:G:167:GLU:OE1	1:I:61:LYS:CD	2.65	0.45
1:G:180:LEU:CD1	1:G:181:ALA:N	2.78	0.45
1:G:236:LEU:HD11	1:G:237:GLU:HG2	1.96	0.45
1:G:324:THR:HG23	1:I:241:GLU:OE2	2.16	0.45
1:I:164:PRO:CG	1:I:174:ALA:CB	2.95	0.45
1:I:361:GLU:HB3	1:I:369:ILE:CD1	2.14	0.45
1:J:33:SER:O	1:J:69:TYR:CD1	2.69	0.45
1:J:164:PRO:CG	1:J:174:ALA:CB	2.95	0.45
1:J:167:GLU:OE1	1:L:61:LYS:CD	2.65	0.45
1:J:324:THR:HG23	1:L:241:GLU:OE2	2.16	0.45
1:K:7:ALA:HB3	1:K:347:ALA:HB1	1.99	0.45
1:K:34:ILE:N	1:K:54:VAL:HG11	2.31	0.45
1:K:164:PRO:CG	1:K:174:ALA:CB	2.95	0.45
1:K:167:GLU:OE1	1:M:61:LYS:CD	2.65	0.45
1:L:164:PRO:CG	1:L:174:ALA:CB	2.95	0.45
1:L:369:ILE:HG22	1:L:370:VAL:H	1.81	0.45
1:M:7:ALA:HB3	1:M:347:ALA:HB1	1.99	0.45
1:M:287:ILE:HB	1:O:244:ASP:HB3	1.97	0.45
1:N:361:GLU:HB3	1:N:369:ILE:CD1	2.14	0.45
1:O:7:ALA:HB3	1:O:347:ALA:HB1	1.99	0.45
1:P:116:ARG:NH1	1:P:375:PHE:HA	2.32	0.45
1:P:287:ILE:HG22	1:P:290:ARG:NH1	2.32	0.45
1:Q:7:ALA:HB1	1:Q:347:ALA:HB1	1.98	0.45
1:S:290:ARG:HB2	1:U:244:ASP:HA	1.97	0.45
1:S:369:ILE:HG23	1:S:370:VAL:N	2.30	0.45
1:T:369:ILE:HG22	1:T:370:VAL:H	1.81	0.45
1:U:8:LEU:O	1:U:104:LEU:N	2.47	0.45
1:A:357:ILE:HD12	1:A:357:ILE:HA	1.79	0.45
1:A:369:ILE:HG22	1:A:370:VAL:H	1.81	0.45
1:B:33:SER:O	1:B:69:TYR:CD1	2.69	0.45
1:B:34:ILE:N	1:B:54:VAL:HG11	2.31	0.45
1:B:144:ALA:HB2	1:B:342:GLY:N	2.31	0.45
1:B:167:GLU:OE1	1:D:61:LYS:CD	2.65	0.45
1:B:300:SER:HA	1:B:335:ARG:NE	2.32	0.45
1:D:38:PRO:CG	1:D:49:GLN:NE2	2.79	0.45
1:D:164:PRO:CG	1:D:174:ALA:CB	2.95	0.45
1:D:290:ARG:HB2	1:F:244:ASP:HA	1.97	0.45
1:E:2:GLU:OE2	1:E:2:GLU:HA	2.10	0.45
1:E:144:ALA:HB2	1:E:342:GLY:N	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:164:PRO:CG	1:E:174:ALA:CB	2.95	0.45
1:G:300:SER:HA	1:G:335:ARG:NE	2.32	0.45
1:H:38:PRO:HD3	1:H:49:GLN:HE22	1.81	0.45
1:H:164:PRO:CG	1:H:174:ALA:CB	2.95	0.45
1:H:236:LEU:CD1	1:H:237:GLU:CG	2.93	0.45
1:I:7:ALA:HB3	1:I:347:ALA:HB1	1.99	0.45
1:I:106:THR:HB	1:I:137:GLN:HG2	1.97	0.45
1:I:116:ARG:NH1	1:I:375:PHE:HA	2.32	0.45
1:I:167:GLU:OE1	1:K:61:LYS:CD	2.65	0.45
1:K:38:PRO:CG	1:K:49:GLN:NE2	2.79	0.45
1:L:287:ILE:HB	1:N:244:ASP:HB3	1.97	0.45
1:L:362:TYR:CE1	1:L:367:PRO:CB	2.99	0.45
1:M:167:GLU:CD	1:O:61:LYS:CE	2.85	0.45
1:M:167:GLU:OE1	1:O:61:LYS:CD	2.65	0.45
1:O:35:VAL:H	1:O:68:LYS:H	1.65	0.45
1:O:290:ARG:HB2	1:Q:244:ASP:HA	1.97	0.45
1:O:369:ILE:HG23	1:O:370:VAL:N	2.30	0.45
1:Q:7:ALA:HB3	1:Q:347:ALA:HB1	1.99	0.45
1:R:164:PRO:CG	1:R:174:ALA:CB	2.95	0.45
1:S:116:ARG:NH1	1:S:375:PHE:HA	2.32	0.45
1:S:236:LEU:HD11	1:S:237:GLU:HG2	1.96	0.45
1:T:38:PRO:CG	1:T:49:GLN:NE2	2.79	0.45
1:U:6:THR:O	1:U:102:PRO:HD2	2.17	0.45
1:U:300:SER:HA	1:U:335:ARG:NE	2.32	0.45
1:V:132:MET:HB2	1:V:132:MET:HE2	1.75	0.45
1:V:287:ILE:HG22	1:V:290:ARG:NH1	2.32	0.45
1:A:287:ILE:HB	1:C:244:ASP:HB3	1.97	0.45
1:B:9:VAL:HG21	1:B:344:SER:HA	1.98	0.45
1:B:38:PRO:CG	1:B:49:GLN:NE2	2.79	0.45
1:B:110:LEU:CD1	1:B:177:ARG:NH1	2.80	0.45
1:B:287:ILE:HB	1:D:244:ASP:HB3	1.97	0.45
1:C:7:ALA:HB1	1:C:347:ALA:HB1	1.98	0.45
1:D:34:ILE:HD13	1:D:67:LEU:CD2	2.46	0.45
1:D:34:ILE:N	1:D:54:VAL:HG11	2.31	0.45
1:E:300:SER:HA	1:E:335:ARG:NE	2.32	0.45
1:E:369:ILE:HG22	1:E:370:VAL:H	1.82	0.45
1:F:287:ILE:HB	1:H:244:ASP:HB3	1.97	0.45
1:G:133:TYR:CE2	1:G:375:PHE:HB2	2.51	0.45
1:H:34:ILE:HD13	1:H:67:LEU:CD2	2.46	0.45
1:H:167:GLU:OE1	1:J:61:LYS:CD	2.65	0.45
1:H:287:ILE:HB	1:J:244:ASP:HB3	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:34:ILE:N	1:J:54:VAL:HG11	2.31	0.45
1:L:9:VAL:HG21	1:L:344:SER:HA	1.98	0.45
1:L:167:GLU:OE1	1:N:61:LYS:CD	2.65	0.45
1:M:44:MET:CG	1:M:45:VAL:N	2.75	0.45
1:O:34:ILE:N	1:O:54:VAL:HG11	2.31	0.45
1:O:38:PRO:HD3	1:O:49:GLN:HE22	1.82	0.45
1:O:132:MET:HE2	1:O:132:MET:HB2	1.74	0.45
1:O:324:THR:HG23	1:Q:241:GLU:OE2	2.16	0.45
1:P:34:ILE:HD13	1:P:67:LEU:CD2	2.46	0.45
1:Q:167:GLU:OE1	1:S:61:LYS:CD	2.65	0.45
1:R:324:THR:HG23	1:T:241:GLU:OE2	2.16	0.45
1:S:7:ALA:HB1	1:S:347:ALA:HB1	1.98	0.45
1:S:7:ALA:HB3	1:S:347:ALA:HB1	1.99	0.45
1:S:335:ARG:HA	1:S:335:ARG:HD3	1.39	0.45
1:T:180:LEU:CD1	1:T:181:ALA:N	2.78	0.45
1:U:290:ARG:HB2	1:W:244:ASP:HA	1.97	0.45
1:V:34:ILE:HG22	1:V:35:VAL:H	1.78	0.45
1:V:34:ILE:N	1:V:54:VAL:HG11	2.31	0.45
1:V:233:SER:HB3	1:V:236:LEU:HG	1.99	0.45
1:W:34:ILE:N	1:W:54:VAL:HG11	2.31	0.45
1:W:38:PRO:HD3	1:W:49:GLN:HE22	1.81	0.45
1:W:73:HIC:HA	1:W:183:ARG:NH1	2.18	0.45
1:W:164:PRO:CG	1:W:174:ALA:CB	2.95	0.45
1:A:132:MET:HB2	1:A:132:MET:HE2	1.74	0.45
1:A:164:PRO:CG	1:A:174:ALA:CB	2.95	0.45
1:A:300:SER:HA	1:A:335:ARG:NE	2.32	0.45
1:B:167:GLU:CD	1:D:61:LYS:CE	2.85	0.45
1:B:324:THR:HG23	1:D:241:GLU:OE2	2.16	0.45
1:C:167:GLU:OE1	1:E:61:LYS:CD	2.65	0.45
1:D:300:SER:HA	1:D:335:ARG:NE	2.32	0.45
1:E:34:ILE:N	1:E:54:VAL:HG11	2.31	0.45
1:F:33:SER:O	1:F:69:TYR:CD1	2.69	0.45
1:G:34:ILE:N	1:G:54:VAL:HG11	2.31	0.45
1:G:147:ARG:HH21	1:G:147:ARG:HG3	1.80	0.45
1:G:233:SER:HB3	1:G:236:LEU:HG	1.99	0.45
1:H:9:VAL:HG21	1:H:344:SER:HA	1.98	0.45
1:I:180:LEU:CD1	1:I:181:ALA:N	2.78	0.45
1:I:300:SER:HA	1:I:335:ARG:NE	2.32	0.45
1:J:34:ILE:C	1:J:54:VAL:HG11	2.42	0.45
1:J:34:ILE:HD13	1:J:67:LEU:CD2	2.46	0.45
1:J:287:ILE:HB	1:L:244:ASP:HB3	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:300:SER:HA	1:L:335:ARG:NE	2.32	0.45
1:L:357:ILE:HD12	1:L:357:ILE:HA	1.79	0.45
1:L:374:CYS:HB2	1:L:375:PHE:H	1.61	0.45
1:P:9:VAL:HG21	1:P:344:SER:HA	1.98	0.45
1:P:369:ILE:HG22	1:P:370:VAL:H	1.82	0.45
1:Q:34:ILE:N	1:Q:54:VAL:HG11	2.31	0.45
1:Q:236:LEU:CD1	1:Q:237:GLU:CG	2.93	0.45
1:R:7:ALA:HB1	1:R:347:ALA:HB1	1.98	0.45
1:R:369:ILE:HG22	1:R:370:VAL:H	1.82	0.45
1:S:9:VAL:HG21	1:S:344:SER:HA	1.98	0.45
1:S:144:ALA:HB2	1:S:342:GLY:N	2.31	0.45
1:S:300:SER:HA	1:S:335:ARG:NE	2.32	0.45
1:T:233:SER:HB3	1:T:236:LEU:HG	1.99	0.45
1:U:164:PRO:CG	1:U:174:ALA:CB	2.95	0.45
1:W:6:THR:O	1:W:102:PRO:HD2	2.17	0.45
1:W:34:ILE:HD13	1:W:67:LEU:CD2	2.46	0.45
1:W:58:ALA:CB	1:W:65:LEU:CD2	2.90	0.45
1:W:143:TYR:HD1	1:W:143:TYR:C	2.09	0.45
1:A:73:HIC:HA	1:A:183:ARG:NH1	2.18	0.44
1:A:167:GLU:OE1	1:C:61:LYS:CD	2.65	0.44
1:C:324:THR:HG23	1:E:241:GLU:OE2	2.16	0.44
1:D:6:THR:O	1:D:102:PRO:HD2	2.17	0.44
1:F:38:PRO:HD3	1:F:49:GLN:HE22	1.81	0.44
1:F:116:ARG:NH1	1:F:375:PHE:HA	2.32	0.44
1:G:7:ALA:HB3	1:G:347:ALA:HB1	1.99	0.44
1:G:287:ILE:HG22	1:G:290:ARG:NH1	2.32	0.44
1:H:65:LEU:C	1:H:65:LEU:CD1	2.90	0.44
1:J:65:LEU:C	1:J:65:LEU:CD1	2.90	0.44
1:K:300:SER:HA	1:K:335:ARG:NE	2.32	0.44
1:L:65:LEU:C	1:L:65:LEU:CD1	2.90	0.44
1:M:164:PRO:CG	1:M:174:ALA:CB	2.95	0.44
1:N:164:PRO:CG	1:N:174:ALA:CB	2.95	0.44
1:N:287:ILE:HB	1:P:244:ASP:HB3	1.98	0.44
1:N:369:ILE:HG22	1:N:370:VAL:H	1.82	0.44
1:O:116:ARG:NH1	1:O:375:PHE:HA	2.32	0.44
1:O:167:GLU:OE1	1:Q:61:LYS:CD	2.65	0.44
1:P:164:PRO:CG	1:P:174:ALA:CB	2.95	0.44
1:P:167:GLU:OE1	1:R:61:LYS:CD	2.65	0.44
1:Q:34:ILE:HD13	1:Q:67:LEU:CD2	2.46	0.44
1:Q:176:MET:HE3	1:Q:176:MET:HB3	1.81	0.44
1:R:116:ARG:NH1	1:R:375:PHE:HA	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:R:133:TYR:CE2	1:R:375:PHE:HB2	2.51	0.44
1:R:167:GLU:OE1	1:T:61:LYS:CD	2.65	0.44
1:S:6:THR:O	1:S:102:PRO:HD2	2.17	0.44
1:S:361:GLU:HB3	1:S:369:ILE:CD1	2.14	0.44
1:U:7:ALA:HB3	1:U:347:ALA:HB1	1.99	0.44
1:U:38:PRO:HD3	1:U:49:GLN:HE22	1.82	0.44
1:V:300:SER:HA	1:V:335:ARG:NE	2.32	0.44
1:W:300:SER:HA	1:W:335:ARG:NE	2.32	0.44
1:W:362:TYR:CE1	1:W:367:PRO:CB	2.99	0.44
1:A:7:ALA:HB1	1:A:347:ALA:HB1	1.98	0.44
1:A:110:LEU:CD1	1:A:177:ARG:NH1	2.80	0.44
1:A:116:ARG:NH1	1:A:375:PHE:HA	2.32	0.44
1:B:6:THR:O	1:B:102:PRO:HD2	2.17	0.44
1:B:31:PHE:HA	1:B:32:PRO:HD3	1.66	0.44
1:C:132:MET:HE2	1:C:132:MET:HB2	1.74	0.44
1:C:164:PRO:CG	1:C:174:ALA:CB	2.95	0.44
1:C:300:SER:HA	1:C:335:ARG:NE	2.32	0.44
1:D:33:SER:O	1:D:69:TYR:CD1	2.69	0.44
1:D:287:ILE:HB	1:F:244:ASP:HB3	1.98	0.44
1:F:65:LEU:C	1:F:65:LEU:CD1	2.90	0.44
1:G:34:ILE:HD13	1:G:67:LEU:CD2	2.46	0.44
1:G:58:ALA:CB	1:G:65:LEU:CD2	2.90	0.44
1:H:6:THR:O	1:H:102:PRO:HD2	2.17	0.44
1:H:34:ILE:C	1:H:54:VAL:HG11	2.42	0.44
1:I:287:ILE:HG22	1:I:290:ARG:NH1	2.32	0.44
1:I:362:TYR:CE1	1:I:367:PRO:CB	2.99	0.44
1:J:73:HIC:HA	1:J:183:ARG:NH1	2.18	0.44
1:J:116:ARG:NH1	1:J:375:PHE:HA	2.32	0.44
1:K:369:ILE:HG23	1:K:370:VAL:N	2.30	0.44
1:L:34:ILE:C	1:L:54:VAL:HG11	2.42	0.44
1:L:144:ALA:HB2	1:L:342:GLY:N	2.31	0.44
1:M:35:VAL:H	1:M:68:LYS:H	1.65	0.44
1:N:65:LEU:C	1:N:65:LEU:CD1	2.90	0.44
1:N:132:MET:HE2	1:N:132:MET:HB2	1.75	0.44
1:N:167:GLU:OE1	1:P:61:LYS:CD	2.65	0.44
1:N:300:SER:HA	1:N:335:ARG:NE	2.32	0.44
1:O:300:SER:HA	1:O:335:ARG:NE	2.32	0.44
1:P:300:SER:HA	1:P:335:ARG:NE	2.32	0.44
1:Q:35:VAL:H	1:Q:68:LYS:H	1.65	0.44
1:Q:132:MET:HE2	1:Q:132:MET:HB2	1.75	0.44
1:S:34:ILE:HD13	1:S:67:LEU:CD2	2.46	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:167:GLU:OE1	1:U:61:LYS:CD	2.65	0.44
1:T:167:GLU:OE1	1:V:61:LYS:CD	2.65	0.44
1:U:7:ALA:HB1	1:U:347:ALA:HB1	1.98	0.44
1:V:61:LYS:HG2	1:V:64:ILE:CG2	2.45	0.44
1:B:34:ILE:O	1:B:54:VAL:HB	2.18	0.44
1:C:170:ALA:O	1:C:172:PRO:HD3	2.18	0.44
1:D:35:VAL:H	1:D:68:LYS:H	1.65	0.44
1:D:110:LEU:CD1	1:D:177:ARG:NH1	2.80	0.44
1:D:167:GLU:OE1	1:F:61:LYS:CD	2.65	0.44
1:E:6:THR:O	1:E:102:PRO:HD2	2.17	0.44
1:F:6:THR:O	1:F:102:PRO:HD2	2.17	0.44
1:F:34:ILE:C	1:F:54:VAL:HG11	2.42	0.44
1:F:35:VAL:H	1:F:68:LYS:H	1.65	0.44
1:F:164:PRO:CG	1:F:174:ALA:CB	2.95	0.44
1:G:34:ILE:O	1:G:54:VAL:HB	2.18	0.44
1:G:110:LEU:CD1	1:G:177:ARG:NH1	2.80	0.44
1:J:34:ILE:O	1:J:54:VAL:HB	2.18	0.44
1:K:116:ARG:NH1	1:K:375:PHE:HA	2.32	0.44
1:L:170:ALA:O	1:L:172:PRO:HD3	2.18	0.44
1:L:324:THR:HG23	1:N:241:GLU:OE2	2.16	0.44
1:M:34:ILE:HG22	1:M:35:VAL:H	1.78	0.44
1:M:34:ILE:O	1:M:54:VAL:HB	2.18	0.44
1:M:38:PRO:CG	1:M:49:GLN:NE2	2.79	0.44
1:O:164:PRO:CG	1:O:174:ALA:CB	2.95	0.44
1:P:173:HIS:HE1	1:Q:268:GLY:CA	2.28	0.44
1:Q:34:ILE:O	1:Q:54:VAL:HB	2.18	0.44
1:Q:300:SER:HA	1:Q:335:ARG:NE	2.32	0.44
1:R:233:SER:HB3	1:R:236:LEU:HG	1.99	0.44
1:S:164:PRO:CG	1:S:174:ALA:CB	2.95	0.44
1:T:6:THR:O	1:T:102:PRO:HD2	2.17	0.44
1:V:34:ILE:O	1:V:54:VAL:HB	2.18	0.44
1:V:38:PRO:CG	1:V:49:GLN:NE2	2.79	0.44
1:V:362:TYR:CE1	1:V:367:PRO:CB	2.99	0.44
1:A:34:ILE:O	1:A:54:VAL:HB	2.18	0.44
1:C:110:LEU:CD1	1:C:177:ARG:NH1	2.80	0.44
1:H:34:ILE:HG22	1:H:35:VAL:H	1.78	0.44
1:I:6:THR:O	1:I:102:PRO:HD2	2.17	0.44
1:I:143:TYR:CE2	1:I:346:LEU:HD13	2.51	0.44
1:I:233:SER:HB3	1:I:236:LEU:HG	1.99	0.44
1:J:7:ALA:HB3	1:J:347:ALA:HB1	1.99	0.44
1:J:9:VAL:HG21	1:J:344:SER:HA	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:151:ILE:O	1:J:297:ASN:HA	2.17	0.44
1:J:369:ILE:HG23	1:J:370:VAL:N	2.30	0.44
1:K:34:ILE:O	1:K:54:VAL:HB	2.18	0.44
1:P:369:ILE:HG23	1:P:370:VAL:N	2.30	0.44
1:Q:148:THR:CG2	1:Q:149:THR:N	2.81	0.44
1:R:357:ILE:HD12	1:R:357:ILE:HA	1.79	0.44
1:U:9:VAL:HG21	1:U:344:SER:HA	1.98	0.44
1:V:7:ALA:HB1	1:V:347:ALA:HB1	1.98	0.44
1:A:287:ILE:HD12	1:C:208:ILE:HD13	1.96	0.44
1:B:34:ILE:CG2	1:B:67:LEU:HD22	2.28	0.44
1:C:34:ILE:HD13	1:C:67:LEU:CD2	2.46	0.44
1:E:34:ILE:O	1:E:54:VAL:HB	2.18	0.44
1:E:110:LEU:CD1	1:E:177:ARG:NH1	2.80	0.44
1:E:180:LEU:CD1	1:E:181:ALA:N	2.78	0.44
1:F:233:SER:HB3	1:F:236:LEU:HG	1.99	0.44
1:H:7:ALA:HB3	1:H:347:ALA:HB1	1.99	0.44
1:H:132:MET:HE2	1:H:132:MET:HB2	1.75	0.44
1:H:151:ILE:O	1:H:297:ASN:HA	2.17	0.44
1:M:73:HIC:HA	1:M:183:ARG:NH1	2.18	0.44
1:M:170:ALA:O	1:M:172:PRO:HD3	2.18	0.44
1:O:287:ILE:HB	1:Q:244:ASP:HB3	1.97	0.44
1:P:299:MET:HE2	1:P:299:MET:HB3	1.95	0.44
1:Q:116:ARG:NH1	1:Q:375:PHE:HA	2.32	0.44
1:Q:233:SER:HB3	1:Q:236:LEU:HG	1.99	0.44
1:R:34:ILE:N	1:R:54:VAL:HG11	2.31	0.44
1:R:34:ILE:O	1:R:54:VAL:HB	2.18	0.44
1:R:298:VAL:CG1	1:R:335:ARG:NH1	2.81	0.44
1:S:34:ILE:N	1:S:54:VAL:HG11	2.31	0.44
1:S:233:SER:HB3	1:S:236:LEU:HG	1.99	0.44
1:U:38:PRO:CG	1:U:49:GLN:NE2	2.79	0.44
1:U:170:ALA:O	1:U:172:PRO:HD3	2.18	0.44
1:V:116:ARG:NH1	1:V:375:PHE:HA	2.32	0.44
1:W:38:PRO:CG	1:W:49:GLN:NE2	2.79	0.44
1:A:298:VAL:CG1	1:A:335:ARG:NH1	2.81	0.44
1:D:233:SER:HB3	1:D:236:LEU:HG	1.99	0.44
1:F:7:ALA:HB3	1:F:347:ALA:HB1	1.99	0.44
1:F:9:VAL:HG21	1:F:344:SER:HA	1.98	0.44
1:F:298:VAL:CG1	1:F:335:ARG:NH1	2.81	0.44
1:F:300:SER:HA	1:F:335:ARG:NE	2.32	0.44
1:G:6:THR:O	1:G:102:PRO:HD2	2.17	0.44
1:G:357:ILE:HD12	1:G:357:ILE:HA	1.78	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:298:VAL:CG1	1:H:335:ARG:NH1	2.81	0.44
1:I:34:ILE:N	1:I:54:VAL:HG11	2.31	0.44
1:J:298:VAL:CG1	1:J:335:ARG:NH1	2.81	0.44
1:K:335:ARG:HA	1:K:335:ARG:HD3	1.39	0.44
1:L:7:ALA:HB3	1:L:347:ALA:HB1	1.99	0.44
1:L:173:HIS:HE1	1:M:268:GLY:CA	2.28	0.44
1:M:116:ARG:NH1	1:M:375:PHE:HA	2.32	0.44
1:M:324:THR:HG23	1:O:241:GLU:OE2	2.17	0.44
1:M:374:CYS:HB2	1:M:375:PHE:H	1.61	0.44
1:N:7:ALA:HB3	1:N:347:ALA:HB1	1.99	0.44
1:N:34:ILE:C	1:N:54:VAL:HG11	2.42	0.44
1:N:58:ALA:CB	1:N:65:LEU:CD2	2.90	0.44
1:N:148:THR:CG2	1:N:149:THR:N	2.81	0.44
1:N:170:ALA:O	1:N:172:PRO:HD3	2.18	0.44
1:O:38:PRO:CG	1:O:49:GLN:NE2	2.79	0.44
1:O:298:VAL:CG1	1:O:335:ARG:NH1	2.81	0.44
1:P:7:ALA:HB3	1:P:347:ALA:HB1	1.99	0.44
1:P:34:ILE:O	1:P:54:VAL:HB	2.18	0.44
1:P:298:VAL:CG1	1:P:335:ARG:NH1	2.81	0.44
1:P:361:GLU:HB3	1:P:369:ILE:CD1	2.14	0.44
1:Q:6:THR:O	1:Q:102:PRO:HD2	2.17	0.44
1:Q:151:ILE:O	1:Q:297:ASN:HA	2.18	0.44
1:Q:298:VAL:CG1	1:Q:335:ARG:NH1	2.81	0.44
1:R:7:ALA:HB3	1:R:347:ALA:HB1	1.99	0.44
1:T:38:PRO:HD3	1:T:49:GLN:HE22	1.81	0.44
1:T:151:ILE:O	1:T:297:ASN:HA	2.17	0.44
1:V:6:THR:O	1:V:102:PRO:HD2	2.17	0.44
1:V:180:LEU:CD1	1:V:181:ALA:N	2.78	0.44
1:W:7:ALA:HB1	1:W:347:ALA:HB1	1.98	0.44
1:W:9:VAL:HG21	1:W:344:SER:HA	1.98	0.44
1:W:170:ALA:O	1:W:172:PRO:HD3	2.18	0.44
1:A:170:ALA:O	1:A:172:PRO:HD3	2.18	0.44
1:B:170:ALA:O	1:B:172:PRO:HD3	2.18	0.44
1:D:298:VAL:CG1	1:D:335:ARG:NH1	2.81	0.44
1:F:110:LEU:CD1	1:F:177:ARG:NH1	2.80	0.44
1:F:167:GLU:OE1	1:H:61:LYS:CD	2.65	0.44
1:G:151:ILE:O	1:G:297:ASN:HA	2.18	0.44
1:I:133:TYR:CE2	1:I:375:PHE:HB2	2.51	0.44
1:I:170:ALA:O	1:I:172:PRO:HD3	2.18	0.44
1:I:298:VAL:CG1	1:I:335:ARG:NH1	2.81	0.44
1:J:170:ALA:O	1:J:172:PRO:HD3	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:287:ILE:HD12	1:L:208:ILE:HD13	1.96	0.44
1:J:300:SER:HA	1:J:335:ARG:NE	2.32	0.44
1:K:34:ILE:HD13	1:K:67:LEU:CD2	2.46	0.44
1:K:298:VAL:CG1	1:K:335:ARG:NH1	2.81	0.44
1:L:6:THR:O	1:L:102:PRO:HD2	2.17	0.44
1:L:287:ILE:HD12	1:N:208:ILE:HD13	1.96	0.44
1:M:298:VAL:CG1	1:M:335:ARG:NH1	2.81	0.44
1:N:6:THR:O	1:N:102:PRO:HD2	2.17	0.44
1:P:6:THR:O	1:P:102:PRO:HD2	2.17	0.44
1:Q:38:PRO:CG	1:Q:49:GLN:NE2	2.79	0.44
1:Q:164:PRO:CG	1:Q:174:ALA:CB	2.95	0.44
1:R:151:ILE:O	1:R:297:ASN:HA	2.17	0.44
1:S:34:ILE:O	1:S:54:VAL:HB	2.18	0.44
1:S:38:PRO:CG	1:S:49:GLN:NE2	2.79	0.44
1:T:7:ALA:HB1	1:T:347:ALA:HB1	1.98	0.44
1:T:34:ILE:N	1:T:54:VAL:HG11	2.31	0.44
1:T:170:ALA:O	1:T:172:PRO:HD3	2.18	0.44
1:T:298:VAL:CG1	1:T:335:ARG:NH1	2.81	0.44
1:V:110:LEU:CD1	1:V:177:ARG:NH1	2.80	0.44
1:W:34:ILE:O	1:W:54:VAL:HB	2.18	0.44
1:A:34:ILE:HG22	1:A:35:VAL:H	1.78	0.44
1:B:104:LEU:HD23	1:B:104:LEU:C	2.43	0.44
1:D:170:ALA:O	1:D:172:PRO:HD3	2.18	0.44
1:E:7:ALA:HB3	1:E:347:ALA:HB1	1.99	0.44
1:E:38:PRO:HD3	1:E:49:GLN:HE22	1.82	0.44
1:E:151:ILE:O	1:E:297:ASN:HA	2.17	0.44
1:E:183:ARG:HE	1:E:183:ARG:HB3	1.58	0.44
1:E:236:LEU:HD11	1:E:237:GLU:HG2	1.97	0.44
1:F:34:ILE:O	1:F:54:VAL:HB	2.18	0.44
1:G:34:ILE:C	1:G:54:VAL:HG11	2.42	0.44
1:H:35:VAL:H	1:H:68:LYS:H	1.65	0.44
1:H:104:LEU:HD23	1:H:104:LEU:C	2.43	0.44
1:H:300:SER:HA	1:H:335:ARG:NE	2.32	0.44
1:I:104:LEU:C	1:I:104:LEU:HD23	2.43	0.44
1:K:104:LEU:HD23	1:K:104:LEU:C	2.43	0.44
1:K:176:MET:HE3	1:K:176:MET:HB3	1.81	0.44
1:N:34:ILE:CG2	1:N:67:LEU:CB	2.96	0.44
1:N:104:LEU:HD23	1:N:104:LEU:C	2.43	0.44
1:N:180:LEU:CD1	1:N:181:ALA:N	2.78	0.44
1:P:148:THR:CG2	1:P:149:THR:N	2.81	0.44
1:R:300:SER:HA	1:R:335:ARG:NE	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:151:ILE:O	1:S:297:ASN:HA	2.18	0.44
1:S:298:VAL:CG1	1:S:335:ARG:NH1	2.81	0.44
1:T:7:ALA:HB3	1:T:347:ALA:HB1	1.99	0.44
1:V:7:ALA:HB3	1:V:347:ALA:HB1	1.99	0.44
1:V:151:ILE:O	1:V:297:ASN:HA	2.17	0.44
1:W:233:SER:HB3	1:W:236:LEU:HG	1.99	0.44
1:B:133:TYR:CE2	1:B:375:PHE:HB2	2.51	0.44
1:B:233:SER:HB3	1:B:236:LEU:HG	1.99	0.44
1:C:151:ILE:O	1:C:297:ASN:HA	2.18	0.44
1:C:298:VAL:CG1	1:C:335:ARG:NH1	2.81	0.44
1:E:39:ARG:NE	1:E:66:THR:CA	2.69	0.44
1:F:8:LEU:CB	1:F:103:THR:HG23	2.48	0.44
1:H:34:ILE:O	1:H:54:VAL:HB	2.18	0.44
1:H:116:ARG:NH1	1:H:375:PHE:HA	2.32	0.44
1:H:233:SER:HB3	1:H:236:LEU:HG	1.99	0.44
1:H:362:TYR:CE1	1:H:367:PRO:CB	2.99	0.44
1:I:147:ARG:HH21	1:I:147:ARG:HG3	1.80	0.44
1:I:151:ILE:O	1:I:297:ASN:HA	2.18	0.44
1:J:233:SER:HB3	1:J:236:LEU:HG	1.99	0.44
1:K:65:LEU:C	1:K:65:LEU:CD1	2.90	0.44
1:K:299:MET:O	1:K:332:PRO:HD2	2.18	0.44
1:K:325:MET:HE3	1:M:244:ASP:O	2.18	0.44
1:L:151:ILE:O	1:L:297:ASN:HA	2.18	0.44
1:L:298:VAL:CG1	1:L:335:ARG:NH1	2.81	0.44
1:M:8:LEU:CB	1:M:103:THR:HG23	2.48	0.44
1:N:357:ILE:HD12	1:N:357:ILE:HA	1.79	0.44
1:O:151:ILE:O	1:O:297:ASN:HA	2.18	0.44
1:O:233:SER:HB3	1:O:236:LEU:HG	1.99	0.44
1:P:34:ILE:CG2	1:P:67:LEU:CB	2.96	0.44
1:P:104:LEU:HD23	1:P:104:LEU:C	2.43	0.44
1:P:233:SER:HB3	1:P:236:LEU:HG	1.99	0.44
1:Q:9:VAL:HG21	1:Q:344:SER:HA	1.98	0.44
1:Q:104:LEU:C	1:Q:104:LEU:HD23	2.43	0.44
1:Q:170:ALA:O	1:Q:172:PRO:HD3	2.18	0.44
1:R:6:THR:O	1:R:102:PRO:HD2	2.17	0.44
1:S:34:ILE:C	1:S:54:VAL:HG11	2.42	0.44
1:T:2:GLU:OE2	1:T:2:GLU:HA	2.10	0.44
1:T:8:LEU:CB	1:T:103:THR:HG23	2.48	0.44
1:T:116:ARG:NH1	1:T:375:PHE:HA	2.32	0.44
1:T:300:SER:HA	1:T:335:ARG:NE	2.32	0.44
1:U:167:GLU:OE1	1:W:61:LYS:CD	2.65	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:233:SER:HB3	1:U:236:LEU:HG	1.99	0.44
1:V:34:ILE:C	1:V:54:VAL:HG11	2.42	0.44
1:W:7:ALA:HB3	1:W:347:ALA:HB1	1.99	0.44
1:W:34:ILE:C	1:W:54:VAL:HG11	2.42	0.44
1:W:104:LEU:HD23	1:W:104:LEU:C	2.43	0.44
1:A:34:ILE:C	1:A:54:VAL:HG11	2.42	0.43
1:A:349:LEU:HD23	1:A:349:LEU:HA	1.79	0.43
1:A:369:ILE:HG23	1:A:370:VAL:N	2.30	0.43
1:B:298:VAL:CG1	1:B:335:ARG:NH1	2.81	0.43
1:D:7:ALA:HB3	1:D:347:ALA:HB1	1.99	0.43
1:D:34:ILE:C	1:D:54:VAL:HG11	2.42	0.43
1:F:34:ILE:HD13	1:F:67:LEU:CD2	2.46	0.43
1:F:151:ILE:O	1:F:297:ASN:HA	2.18	0.43
1:F:367:PRO:O	1:F:368:SER:C	2.61	0.43
1:G:170:ALA:O	1:G:172:PRO:HD3	2.18	0.43
1:G:173:HIS:HE1	1:H:268:GLY:CA	2.28	0.43
1:H:110:LEU:CD1	1:H:177:ARG:NH1	2.80	0.43
1:I:34:ILE:C	1:I:54:VAL:HG11	2.42	0.43
1:I:324:THR:HG23	1:K:241:GLU:OE2	2.16	0.43
1:I:325:MET:HE3	1:K:244:ASP:O	2.18	0.43
1:L:34:ILE:O	1:L:54:VAL:HB	2.17	0.43
1:L:69:TYR:CD2	1:L:69:TYR:N	2.86	0.43
1:L:180:LEU:CD1	1:L:181:ALA:N	2.78	0.43
1:L:233:SER:HB3	1:L:236:LEU:HG	1.99	0.43
1:M:233:SER:HB3	1:M:236:LEU:HG	1.99	0.43
1:M:300:SER:HA	1:M:335:ARG:NE	2.32	0.43
1:M:325:MET:HE3	1:O:244:ASP:O	2.18	0.43
1:N:34:ILE:O	1:N:54:VAL:HB	2.18	0.43
1:N:64:ILE:CG2	1:N:65:LEU:N	2.81	0.43
1:N:233:SER:HB3	1:N:236:LEU:HG	1.99	0.43
1:N:298:VAL:CG1	1:N:335:ARG:NH1	2.81	0.43
1:O:8:LEU:CB	1:O:103:THR:HG23	2.48	0.43
1:O:34:ILE:HD13	1:O:67:LEU:CD2	2.46	0.43
1:Q:34:ILE:C	1:Q:54:VAL:HG11	2.42	0.43
1:R:8:LEU:CB	1:R:103:THR:HG23	2.48	0.43
1:R:349:LEU:HD23	1:R:349:LEU:HA	1.79	0.43
1:S:104:LEU:HD23	1:S:104:LEU:C	2.43	0.43
1:S:148:THR:CG2	1:S:149:THR:N	2.81	0.43
1:S:349:LEU:HD23	1:S:349:LEU:HA	1.80	0.43
1:U:104:LEU:HD23	1:U:104:LEU:C	2.43	0.43
1:U:151:ILE:O	1:U:297:ASN:HA	2.17	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:236:LEU:CD1	1:U:237:GLU:CG	2.93	0.43
1:U:298:VAL:CG1	1:U:335:ARG:NH1	2.81	0.43
1:V:35:VAL:H	1:V:68:LYS:H	1.65	0.43
1:V:298:VAL:CG1	1:V:335:ARG:NH1	2.81	0.43
1:W:298:VAL:CG1	1:W:335:ARG:NH1	2.81	0.43
1:A:6:THR:O	1:A:102:PRO:HD2	2.17	0.43
1:A:324:THR:HG23	1:C:241:GLU:OE2	2.16	0.43
1:B:35:VAL:H	1:B:68:LYS:H	1.65	0.43
1:C:34:ILE:C	1:C:54:VAL:HG11	2.42	0.43
1:C:58:ALA:CB	1:C:65:LEU:CD2	2.90	0.43
1:C:64:ILE:CG2	1:C:65:LEU:N	2.81	0.43
1:D:8:LEU:CB	1:D:103:THR:HG23	2.48	0.43
1:D:34:ILE:O	1:D:54:VAL:HB	2.17	0.43
1:E:34:ILE:C	1:E:54:VAL:HG11	2.42	0.43
1:E:64:ILE:CG2	1:E:65:LEU:N	2.81	0.43
1:G:104:LEU:HD23	1:G:104:LEU:C	2.43	0.43
1:G:298:VAL:CG1	1:G:335:ARG:NH1	2.81	0.43
1:G:367:PRO:O	1:G:368:SER:C	2.62	0.43
1:H:64:ILE:CG2	1:H:65:LEU:N	2.81	0.43
1:K:6:THR:O	1:K:102:PRO:HD2	2.17	0.43
1:K:151:ILE:O	1:K:297:ASN:HA	2.18	0.43
1:L:58:ALA:CB	1:L:65:LEU:CD2	2.90	0.43
1:L:104:LEU:C	1:L:104:LEU:HD23	2.43	0.43
1:M:104:LEU:HD23	1:M:104:LEU:C	2.43	0.43
1:M:180:LEU:O	1:M:180:LEU:HG	2.11	0.43
1:M:367:PRO:O	1:M:368:SER:C	2.61	0.43
1:P:39:ARG:NE	1:P:66:THR:CA	2.69	0.43
1:P:324:THR:HG23	1:R:241:GLU:OE2	2.16	0.43
1:P:357:ILE:HD12	1:P:357:ILE:HA	1.79	0.43
1:R:34:ILE:CG2	1:R:67:LEU:CB	2.96	0.43
1:R:104:LEU:HD23	1:R:104:LEU:C	2.43	0.43
1:R:148:THR:CG2	1:R:149:THR:N	2.81	0.43
1:R:299:MET:HE2	1:R:299:MET:HB3	1.95	0.43
1:S:31:PHE:HA	1:S:32:PRO:HD3	1.66	0.43
1:T:172:PRO:HA	1:T:175:ILE:CD1	2.49	0.43
1:T:367:PRO:O	1:T:368:SER:C	2.61	0.43
1:U:69:TYR:CD2	1:U:69:TYR:N	2.87	0.43
1:U:171:LEU:HA	1:U:172:PRO:HD3	1.79	0.43
1:U:362:TYR:CE1	1:U:367:PRO:CB	2.99	0.43
1:V:8:LEU:CB	1:V:103:THR:HG23	2.48	0.43
1:V:299:MET:O	1:V:332:PRO:HD2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:69:TYR:N	1:W:69:TYR:CD2	2.87	0.43
1:W:149:THR:HG23	1:W:150:GLY:N	2.34	0.43
1:C:34:ILE:O	1:C:54:VAL:HB	2.18	0.43
1:D:104:LEU:HD23	1:D:104:LEU:C	2.43	0.43
1:D:367:PRO:O	1:D:368:SER:C	2.61	0.43
1:E:172:PRO:HA	1:E:175:ILE:CD1	2.49	0.43
1:E:367:PRO:O	1:E:368:SER:C	2.61	0.43
1:G:35:VAL:H	1:G:68:LYS:H	1.65	0.43
1:G:149:THR:HG23	1:G:150:GLY:N	2.34	0.43
1:G:299:MET:O	1:G:332:PRO:HD2	2.18	0.43
1:H:8:LEU:CB	1:H:103:THR:HG23	2.48	0.43
1:J:6:THR:O	1:J:102:PRO:HD2	2.17	0.43
1:J:64:ILE:CG2	1:J:65:LEU:N	2.81	0.43
1:J:69:TYR:CD2	1:J:69:TYR:N	2.86	0.43
1:J:180:LEU:O	1:J:180:LEU:HG	2.11	0.43
1:K:35:VAL:H	1:K:68:LYS:H	1.65	0.43
1:K:180:LEU:CD1	1:K:181:ALA:N	2.78	0.43
1:L:31:PHE:HA	1:L:32:PRO:HD3	1.66	0.43
1:L:34:ILE:CG2	1:L:67:LEU:CB	2.96	0.43
1:L:149:THR:HG23	1:L:150:GLY:N	2.34	0.43
1:M:38:PRO:HD3	1:M:49:GLN:HE22	1.82	0.43
1:M:287:ILE:HD12	1:O:208:ILE:HD13	1.96	0.43
1:N:69:TYR:CD2	1:N:69:TYR:N	2.86	0.43
1:N:324:THR:HG23	1:P:241:GLU:OE2	2.16	0.43
1:O:6:THR:O	1:O:102:PRO:HD2	2.17	0.43
1:O:34:ILE:C	1:O:54:VAL:HG11	2.42	0.43
1:O:367:PRO:O	1:O:368:SER:C	2.61	0.43
1:P:34:ILE:C	1:P:54:VAL:HG11	2.42	0.43
1:P:172:PRO:HA	1:P:175:ILE:CD1	2.49	0.43
1:Q:31:PHE:HA	1:Q:32:PRO:HD3	1.66	0.43
1:R:170:ALA:O	1:R:172:PRO:HD3	2.18	0.43
1:S:64:ILE:CG2	1:S:65:LEU:N	2.81	0.43
1:S:69:TYR:CD2	1:S:69:TYR:N	2.87	0.43
1:S:170:ALA:O	1:S:172:PRO:HD3	2.18	0.43
1:S:367:PRO:O	1:S:368:SER:C	2.61	0.43
1:T:64:ILE:CG2	1:T:65:LEU:N	2.81	0.43
1:T:104:LEU:C	1:T:104:LEU:HD23	2.43	0.43
1:T:110:LEU:CD1	1:T:177:ARG:NH1	2.80	0.43
1:U:367:PRO:O	1:U:368:SER:C	2.61	0.43
1:V:367:PRO:O	1:V:368:SER:C	2.62	0.43
1:W:35:VAL:H	1:W:68:LYS:H	1.65	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:7:ALA:HB3	1:B:347:ALA:HB1	1.99	0.43
1:B:236:LEU:HD11	1:B:237:GLU:HG2	1.96	0.43
1:C:6:THR:O	1:C:102:PRO:HD2	2.17	0.43
1:C:31:PHE:HA	1:C:32:PRO:HD3	1.66	0.43
1:E:104:LEU:HD23	1:E:104:LEU:C	2.43	0.43
1:E:149:THR:HG23	1:E:150:GLY:N	2.34	0.43
1:F:170:ALA:O	1:F:172:PRO:HD3	2.18	0.43
1:G:325:MET:HE3	1:I:244:ASP:O	2.18	0.43
1:I:65:LEU:C	1:I:65:LEU:CD1	2.90	0.43
1:I:148:THR:CG2	1:I:149:THR:N	2.81	0.43
1:J:176:MET:HE3	1:J:176:MET:HB3	1.81	0.43
1:J:299:MET:HE2	1:J:299:MET:HB3	1.95	0.43
1:K:324:THR:HG23	1:M:241:GLU:OE2	2.16	0.43
1:M:151:ILE:O	1:M:297:ASN:HA	2.18	0.43
1:N:172:PRO:HA	1:N:175:ILE:CD1	2.49	0.43
1:O:170:ALA:O	1:O:172:PRO:HD3	2.18	0.43
1:P:136:ILE:O	1:P:139:VAL:HB	2.19	0.43
1:P:151:ILE:O	1:P:297:ASN:HA	2.18	0.43
1:Q:8:LEU:CB	1:Q:103:THR:HG23	2.48	0.43
1:Q:69:TYR:N	1:Q:69:TYR:CD2	2.87	0.43
1:R:31:PHE:HA	1:R:32:PRO:HD3	1.66	0.43
1:U:34:ILE:C	1:U:54:VAL:HG11	2.42	0.43
1:U:34:ILE:HD13	1:U:67:LEU:CD2	2.46	0.43
1:W:107:GLU:O	1:W:137:GLN:HG3	2.19	0.43
1:W:151:ILE:O	1:W:297:ASN:HA	2.17	0.43
1:A:8:LEU:CB	1:A:103:THR:HG23	2.48	0.43
1:A:58:ALA:CB	1:A:65:LEU:CD2	2.90	0.43
1:A:149:THR:HG23	1:A:150:GLY:N	2.34	0.43
1:B:148:THR:CG2	1:B:149:THR:N	2.81	0.43
1:C:8:LEU:CB	1:C:103:THR:HG23	2.48	0.43
1:C:136:ILE:O	1:C:139:VAL:HB	2.19	0.43
1:C:172:PRO:HA	1:C:175:ILE:CD1	2.49	0.43
1:D:133:TYR:CE2	1:D:375:PHE:HB2	2.51	0.43
1:E:104:LEU:HD13	1:E:347:ALA:HB2	2.01	0.43
1:E:170:ALA:O	1:E:172:PRO:HD3	2.18	0.43
1:E:176:MET:HE3	1:E:176:MET:HB3	1.81	0.43
1:F:34:ILE:HG22	1:F:35:VAL:H	1.78	0.43
1:G:172:PRO:HA	1:G:175:ILE:CD1	2.49	0.43
1:H:148:THR:CG2	1:H:149:THR:N	2.81	0.43
1:I:34:ILE:O	1:I:54:VAL:HB	2.18	0.43
1:I:149:THR:HG23	1:I:150:GLY:N	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:104:LEU:HD23	1:J:104:LEU:C	2.43	0.43
1:J:110:LEU:CD1	1:J:177:ARG:NH1	2.80	0.43
1:J:362:TYR:CE1	1:J:367:PRO:CB	2.99	0.43
1:K:149:THR:HG23	1:K:150:GLY:N	2.34	0.43
1:K:170:ALA:O	1:K:172:PRO:HD3	2.18	0.43
1:K:236:LEU:HD11	1:K:237:GLU:HG2	1.96	0.43
1:L:176:MET:HE3	1:L:176:MET:HB3	1.81	0.43
1:M:6:THR:O	1:M:102:PRO:HD2	2.17	0.43
1:M:299:MET:O	1:M:332:PRO:HD2	2.18	0.43
1:N:38:PRO:HD3	1:N:49:GLN:HE22	1.81	0.43
1:N:151:ILE:O	1:N:297:ASN:HA	2.18	0.43
1:O:172:PRO:HA	1:O:175:ILE:CD1	2.49	0.43
1:O:299:MET:O	1:O:332:PRO:HD2	2.18	0.43
1:O:369:ILE:C	1:O:371:HIS:N	2.77	0.43
1:P:8:LEU:CB	1:P:103:THR:HG23	2.48	0.43
1:P:367:PRO:O	1:P:368:SER:C	2.61	0.43
1:Q:369:ILE:C	1:Q:371:HIS:N	2.77	0.43
1:R:172:PRO:HA	1:R:175:ILE:CD1	2.49	0.43
1:S:35:VAL:H	1:S:68:LYS:H	1.65	0.43
1:T:299:MET:O	1:T:332:PRO:HD2	2.18	0.43
1:U:167:GLU:CD	1:W:61:LYS:CE	2.85	0.43
1:V:31:PHE:HA	1:V:32:PRO:HD3	1.66	0.43
1:V:173:HIS:HE1	1:W:268:GLY:CA	2.28	0.43
1:A:39:ARG:NE	1:A:66:THR:CA	2.69	0.43
1:A:136:ILE:O	1:A:139:VAL:HB	2.19	0.43
1:A:367:PRO:O	1:A:368:SER:C	2.62	0.43
1:B:34:ILE:HD13	1:B:67:LEU:CD2	2.46	0.43
1:C:7:ALA:HB3	1:C:347:ALA:HB1	1.99	0.43
1:C:104:LEU:HD13	1:C:347:ALA:HB2	2.01	0.43
1:C:104:LEU:HD23	1:C:104:LEU:C	2.43	0.43
1:D:151:ILE:O	1:D:297:ASN:HA	2.18	0.43
1:E:35:VAL:H	1:E:68:LYS:H	1.65	0.43
1:E:136:ILE:O	1:E:139:VAL:HB	2.19	0.43
1:E:298:VAL:CG1	1:E:335:ARG:NH1	2.81	0.43
1:E:374:CYS:HB2	1:E:375:PHE:H	1.61	0.43
1:F:104:LEU:C	1:F:104:LEU:HD23	2.43	0.43
1:F:362:TYR:CE1	1:F:367:PRO:CB	2.99	0.43
1:H:69:TYR:CD2	1:H:69:TYR:N	2.87	0.43
1:H:104:LEU:HD13	1:H:347:ALA:HB2	2.01	0.43
1:I:2:GLU:OE2	1:I:2:GLU:HA	2.10	0.43
1:I:2:GLU:HB3	1:I:3:ASP:H	1.66	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:104:LEU:HD13	1:J:347:ALA:HB2	2.01	0.43
1:K:148:THR:CG2	1:K:149:THR:N	2.81	0.43
1:K:233:SER:HB3	1:K:236:LEU:HG	1.99	0.43
1:K:362:TYR:CE1	1:K:367:PRO:CB	2.99	0.43
1:M:34:ILE:C	1:M:54:VAL:HG11	2.42	0.43
1:M:172:PRO:HA	1:M:175:ILE:CD1	2.49	0.43
1:N:136:ILE:O	1:N:139:VAL:HB	2.19	0.43
1:O:34:ILE:O	1:O:54:VAL:HB	2.18	0.43
1:O:69:TYR:N	1:O:69:TYR:CD2	2.86	0.43
1:P:104:LEU:HD13	1:P:347:ALA:HB2	2.01	0.43
1:Q:172:PRO:HA	1:Q:175:ILE:CD1	2.49	0.43
1:R:64:ILE:CG2	1:R:65:LEU:N	2.81	0.43
1:R:136:ILE:O	1:R:139:VAL:HB	2.19	0.43
1:R:369:ILE:C	1:R:371:HIS:N	2.77	0.43
1:S:132:MET:HE2	1:S:132:MET:HB2	1.75	0.43
1:S:369:ILE:C	1:S:371:HIS:N	2.77	0.43
1:T:34:ILE:C	1:T:54:VAL:HG11	2.42	0.43
1:T:34:ILE:O	1:T:54:VAL:HB	2.18	0.43
1:T:133:TYR:CE2	1:T:375:PHE:HB2	2.51	0.43
1:T:149:THR:HG23	1:T:150:GLY:N	2.34	0.43
1:V:104:LEU:HD23	1:V:104:LEU:C	2.43	0.43
1:V:149:THR:HG23	1:V:150:GLY:N	2.34	0.43
1:V:172:PRO:HA	1:V:175:ILE:CD1	2.49	0.43
1:V:369:ILE:HG23	1:V:370:VAL:N	2.30	0.43
1:W:64:ILE:CG2	1:W:65:LEU:N	2.82	0.43
1:W:104:LEU:HD13	1:W:347:ALA:HB2	2.01	0.43
1:B:34:ILE:C	1:B:54:VAL:HG11	2.42	0.43
1:B:69:TYR:CD2	1:B:69:TYR:N	2.86	0.43
1:B:369:ILE:C	1:B:371:HIS:N	2.77	0.43
1:C:149:THR:HG23	1:C:150:GLY:N	2.34	0.43
1:H:31:PHE:HA	1:H:32:PRO:HD3	1.66	0.43
1:H:107:GLU:O	1:H:137:GLN:HG3	2.19	0.43
1:J:8:LEU:CB	1:J:103:THR:HG23	2.48	0.43
1:J:149:THR:HG23	1:J:150:GLY:N	2.34	0.43
1:J:180:LEU:CD1	1:J:181:ALA:N	2.78	0.43
1:J:367:PRO:O	1:J:368:SER:C	2.61	0.43
1:K:34:ILE:C	1:K:54:VAL:HG11	2.42	0.43
1:K:367:PRO:O	1:K:368:SER:C	2.61	0.43
1:L:104:LEU:HD13	1:L:347:ALA:HB2	2.01	0.43
1:L:107:GLU:O	1:L:137:GLN:HG3	2.19	0.43
1:L:172:PRO:HA	1:L:175:ILE:CD1	2.49	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:335:ARG:HD3	1:N:335:ARG:HA	1.39	0.43
1:P:369:ILE:C	1:P:371:HIS:N	2.77	0.43
1:Q:65:LEU:HD13	1:Q:67:LEU:HD23	2.01	0.43
1:R:104:LEU:HD13	1:R:347:ALA:HB2	2.01	0.43
1:R:110:LEU:CD1	1:R:177:ARG:NH1	2.80	0.43
1:R:149:THR:HG23	1:R:150:GLY:N	2.34	0.43
1:S:172:PRO:HA	1:S:175:ILE:CD1	2.49	0.43
1:T:132:MET:HE2	1:T:132:MET:HB2	1.75	0.43
1:T:148:THR:CG2	1:T:149:THR:N	2.81	0.43
1:U:34:ILE:O	1:U:54:VAL:HB	2.18	0.43
1:U:35:VAL:H	1:U:68:LYS:H	1.65	0.43
1:U:64:ILE:CG2	1:U:65:LEU:N	2.81	0.43
1:U:149:THR:HG23	1:U:150:GLY:N	2.34	0.43
1:U:172:PRO:HA	1:U:175:ILE:CD1	2.49	0.43
1:W:8:LEU:CB	1:W:103:THR:HG23	2.48	0.43
1:W:172:PRO:HA	1:W:175:ILE:CD1	2.49	0.43
1:A:104:LEU:HD13	1:A:347:ALA:HB2	2.01	0.43
1:A:133:TYR:CE2	1:A:375:PHE:HB2	2.51	0.43
1:B:64:ILE:CG2	1:B:65:LEU:N	2.81	0.43
1:B:65:LEU:HD13	1:B:67:LEU:HD23	2.01	0.43
1:C:35:VAL:H	1:C:68:LYS:H	1.65	0.43
1:F:107:GLU:O	1:F:137:GLN:HG3	2.19	0.43
1:F:236:LEU:HD11	1:F:237:GLU:HG2	1.96	0.43
1:H:39:ARG:NE	1:H:66:THR:CA	2.69	0.43
1:H:170:ALA:O	1:H:172:PRO:HD3	2.18	0.43
1:H:299:MET:O	1:H:332:PRO:HD2	2.18	0.43
1:I:31:PHE:HA	1:I:32:PRO:HD3	1.66	0.43
1:I:64:ILE:CG2	1:I:65:LEU:N	2.81	0.43
1:J:39:ARG:NE	1:J:66:THR:CA	2.69	0.43
1:J:299:MET:O	1:J:332:PRO:HD2	2.18	0.43
1:K:64:ILE:CG2	1:K:65:LEU:N	2.81	0.43
1:K:133:TYR:CE2	1:K:375:PHE:HB2	2.51	0.43
1:N:369:ILE:C	1:N:371:HIS:N	2.77	0.43
1:O:65:LEU:HD13	1:O:67:LEU:HD23	2.01	0.43
1:P:69:TYR:CD2	1:P:69:TYR:N	2.86	0.43
1:P:149:THR:HG23	1:P:150:GLY:N	2.34	0.43
1:Q:34:ILE:CG2	1:Q:67:LEU:CB	2.96	0.43
1:Q:64:ILE:CG2	1:Q:65:LEU:N	2.81	0.43
1:T:39:ARG:NE	1:T:66:THR:CA	2.69	0.43
1:T:104:LEU:HD13	1:T:347:ALA:HB2	2.01	0.43
1:T:136:ILE:O	1:T:139:VAL:HB	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:U:104:LEU:HD13	1:U:347:ALA:HB2	2.01	0.43
1:V:64:ILE:CG2	1:V:65:LEU:N	2.81	0.43
1:V:72:GLU:O	1:V:183:ARG:NH2	2.52	0.43
1:A:151:ILE:O	1:A:297:ASN:HA	2.18	0.43
1:A:172:PRO:HA	1:A:175:ILE:CD1	2.49	0.43
1:B:151:ILE:O	1:B:297:ASN:HA	2.17	0.43
1:B:178:LEU:HG	1:B:180:LEU:H	1.84	0.43
1:B:325:MET:HE3	1:D:244:ASP:O	2.18	0.43
1:C:107:GLU:O	1:C:137:GLN:HG3	2.19	0.43
1:D:64:ILE:CG2	1:D:65:LEU:N	2.81	0.43
1:D:349:LEU:HD23	1:D:349:LEU:HA	1.80	0.43
1:E:8:LEU:CB	1:E:103:THR:HG23	2.48	0.43
1:E:34:ILE:CG2	1:E:67:LEU:CB	2.96	0.43
1:E:178:LEU:HG	1:E:180:LEU:H	1.84	0.43
1:G:2:GLU:HB3	1:G:3:ASP:H	1.66	0.43
1:G:34:ILE:CG2	1:G:67:LEU:CB	2.96	0.43
1:G:72:GLU:O	1:G:183:ARG:NH2	2.52	0.43
1:G:136:ILE:O	1:G:139:VAL:HB	2.19	0.43
1:I:172:PRO:HA	1:I:175:ILE:CD1	2.49	0.43
1:L:110:LEU:CD1	1:L:177:ARG:NH1	2.80	0.43
1:L:136:ILE:O	1:L:139:VAL:HB	2.19	0.43
1:M:236:LEU:HD11	1:M:237:GLU:HG2	1.96	0.43
1:M:357:ILE:HD12	1:M:357:ILE:HA	1.79	0.43
1:N:107:GLU:O	1:N:137:GLN:HG3	2.19	0.43
1:O:104:LEU:HD23	1:O:104:LEU:C	2.43	0.43
1:O:236:LEU:HD11	1:O:237:GLU:HG2	1.96	0.43
1:P:35:VAL:H	1:P:68:LYS:H	1.65	0.43
1:P:110:LEU:CD1	1:P:177:ARG:NH1	2.80	0.43
1:S:65:LEU:HD13	1:S:67:LEU:HD23	2.01	0.43
1:T:369:ILE:C	1:T:371:HIS:N	2.77	0.43
1:W:299:MET:O	1:W:332:PRO:HD2	2.18	0.43
1:A:35:VAL:H	1:A:68:LYS:H	1.65	0.43
1:B:374:CYS:HB2	1:B:375:PHE:H	1.62	0.43
1:C:72:GLU:O	1:C:183:ARG:NH2	2.52	0.43
1:D:65:LEU:HD13	1:D:67:LEU:HD23	2.01	0.43
1:D:172:PRO:HA	1:D:175:ILE:CD1	2.49	0.43
1:D:236:LEU:HD11	1:D:237:GLU:HG2	1.96	0.43
1:E:105:LEU:HD11	1:E:123:MET:HG3	2.01	0.43
1:F:65:LEU:HD13	1:F:67:LEU:HD23	2.01	0.43
1:F:236:LEU:CD1	1:F:237:GLU:CG	2.93	0.43
1:G:176:MET:HE3	1:G:176:MET:HB3	1.81	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:369:ILE:HG23	1:G:370:VAL:N	2.30	0.43
1:J:172:PRO:HA	1:J:175:ILE:CD1	2.49	0.43
1:K:72:GLU:O	1:K:183:ARG:NH2	2.52	0.43
1:K:107:GLU:O	1:K:137:GLN:HG3	2.19	0.43
1:K:172:PRO:HA	1:K:175:ILE:CD1	2.49	0.43
1:K:178:LEU:HG	1:K:180:LEU:H	1.84	0.43
1:L:34:ILE:HD13	1:L:67:LEU:CD2	2.46	0.43
1:L:35:VAL:H	1:L:68:LYS:H	1.65	0.43
1:L:64:ILE:CG2	1:L:65:LEU:N	2.81	0.43
1:L:106:THR:HG22	1:L:140:LEU:HD11	2.01	0.43
1:L:367:PRO:O	1:L:368:SER:C	2.62	0.43
1:M:107:GLU:O	1:M:137:GLN:HG3	2.19	0.43
1:M:368:SER:O	1:M:369:ILE:C	2.62	0.43
1:N:35:VAL:H	1:N:68:LYS:H	1.65	0.43
1:N:104:LEU:HD13	1:N:347:ALA:HB2	2.01	0.43
1:N:110:LEU:CD1	1:N:177:ARG:NH1	2.80	0.43
1:P:105:LEU:HD11	1:P:123:MET:HG3	2.01	0.43
1:P:170:ALA:O	1:P:172:PRO:HD3	2.18	0.43
1:Q:107:GLU:O	1:Q:137:GLN:HG3	2.19	0.43
1:Q:178:LEU:HG	1:Q:180:LEU:H	1.84	0.43
1:Q:236:LEU:HD11	1:Q:237:GLU:HG2	1.96	0.43
1:Q:299:MET:O	1:Q:332:PRO:HD2	2.18	0.43
1:R:35:VAL:H	1:R:68:LYS:H	1.65	0.43
1:R:72:GLU:O	1:R:183:ARG:NH2	2.52	0.43
1:R:299:MET:O	1:R:332:PRO:HD2	2.18	0.43
1:U:8:LEU:CB	1:U:103:THR:HG23	2.48	0.43
1:U:34:ILE:HG23	1:U:67:LEU:HB3	2.01	0.43
1:V:39:ARG:NE	1:V:66:THR:CA	2.69	0.43
1:W:106:THR:HG22	1:W:140:LEU:HD11	2.01	0.43
1:A:7:ALA:HB3	1:A:347:ALA:HB1	1.99	0.42
1:A:106:THR:HG22	1:A:140:LEU:HD11	2.01	0.42
1:A:107:GLU:O	1:A:137:GLN:HG3	2.19	0.42
1:A:299:MET:O	1:A:332:PRO:HD2	2.18	0.42
1:B:132:MET:HB2	1:B:132:MET:HE2	1.75	0.42
1:B:172:PRO:HA	1:B:175:ILE:CD1	2.49	0.42
1:B:368:SER:O	1:B:369:ILE:C	2.62	0.42
1:D:110:LEU:CD1	1:D:177:ARG:HH11	2.32	0.42
1:D:299:MET:O	1:D:332:PRO:HD2	2.18	0.42
1:E:299:MET:O	1:E:332:PRO:HD2	2.18	0.42
1:F:69:TYR:CD2	1:F:69:TYR:N	2.86	0.42
1:F:133:TYR:CE2	1:F:375:PHE:HB2	2.51	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:149:THR:HG23	1:F:150:GLY:N	2.34	0.42
1:F:172:PRO:HA	1:F:175:ILE:CD1	2.49	0.42
1:G:104:LEU:HD13	1:G:347:ALA:HB2	2.01	0.42
1:G:148:THR:CG2	1:G:149:THR:N	2.81	0.42
1:G:167:GLU:CD	1:I:61:LYS:CE	2.85	0.42
1:H:325:MET:HE3	1:J:244:ASP:O	2.18	0.42
1:I:35:VAL:H	1:I:68:LYS:H	1.65	0.42
1:I:105:LEU:HD11	1:I:123:MET:HG3	2.01	0.42
1:K:105:LEU:HD11	1:K:123:MET:HG3	2.01	0.42
1:L:105:LEU:HD11	1:L:123:MET:HG3	2.01	0.42
1:L:299:MET:O	1:L:332:PRO:HD2	2.18	0.42
1:M:149:THR:HG23	1:M:150:GLY:N	2.34	0.42
1:N:105:LEU:HD11	1:N:123:MET:HG3	2.01	0.42
1:N:106:THR:HG22	1:N:140:LEU:HD11	2.01	0.42
1:N:149:THR:HG23	1:N:150:GLY:N	2.34	0.42
1:N:325:MET:HE3	1:P:244:ASP:O	2.18	0.42
1:O:44:MET:CG	1:O:45:VAL:N	2.75	0.42
1:Q:167:GLU:OE1	1:S:61:LYS:HE2	2.19	0.42
1:R:34:ILE:C	1:R:54:VAL:HG11	2.42	0.42
1:S:136:ILE:O	1:S:139:VAL:HB	2.19	0.42
1:T:178:LEU:HG	1:T:180:LEU:H	1.84	0.42
1:V:2:GLU:HB3	1:V:3:ASP:H	1.66	0.42
1:V:105:LEU:HD11	1:V:123:MET:HG3	2.01	0.42
1:V:170:ALA:O	1:V:172:PRO:HD3	2.18	0.42
1:W:34:ILE:HG23	1:W:67:LEU:HB3	2.02	0.42
1:W:105:LEU:HD11	1:W:123:MET:HG3	2.01	0.42
1:A:104:LEU:C	1:A:104:LEU:HD23	2.43	0.42
1:B:173:HIS:HE1	1:C:268:GLY:CA	2.28	0.42
1:C:105:LEU:HD11	1:C:123:MET:HG3	2.01	0.42
1:C:299:MET:O	1:C:332:PRO:HD2	2.18	0.42
1:C:335:ARG:HD3	1:C:335:ARG:HA	1.39	0.42
1:D:69:TYR:CD2	1:D:69:TYR:N	2.86	0.42
1:D:325:MET:HE3	1:F:244:ASP:O	2.18	0.42
1:F:64:ILE:CG2	1:F:65:LEU:N	2.81	0.42
1:F:104:LEU:HD13	1:F:347:ALA:HB2	2.01	0.42
1:G:69:TYR:N	1:G:69:TYR:CD2	2.87	0.42
1:H:167:GLU:OE1	1:J:64:ILE:HD13	2.19	0.42
1:H:172:PRO:HA	1:H:175:ILE:CD1	2.49	0.42
1:I:107:GLU:O	1:I:137:GLN:HG3	2.19	0.42
1:I:136:ILE:O	1:I:139:VAL:HB	2.19	0.42
1:I:143:TYR:HD1	1:I:143:TYR:C	2.09	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:367:PRO:O	1:I:368:SER:C	2.62	0.42
1:J:34:ILE:CG2	1:J:67:LEU:CB	2.96	0.42
1:J:105:LEU:HD11	1:J:123:MET:HG3	2.02	0.42
1:J:180:LEU:HD12	1:J:181:ALA:CA	2.50	0.42
1:K:167:GLU:OE1	1:M:64:ILE:HD13	2.19	0.42
1:K:349:LEU:HD23	1:K:349:LEU:HA	1.79	0.42
1:L:8:LEU:N	1:L:102:PRO:O	2.48	0.42
1:L:178:LEU:HG	1:L:180:LEU:H	1.84	0.42
1:L:369:ILE:C	1:L:371:HIS:N	2.77	0.42
1:M:65:LEU:HD13	1:M:67:LEU:HD23	2.01	0.42
1:M:136:ILE:O	1:M:139:VAL:HB	2.19	0.42
1:N:8:LEU:CB	1:N:103:THR:HG23	2.48	0.42
1:N:299:MET:O	1:N:332:PRO:HD2	2.18	0.42
1:O:107:GLU:O	1:O:137:GLN:HG3	2.19	0.42
1:O:362:TYR:CE1	1:O:367:PRO:CB	2.99	0.42
1:P:325:MET:HE3	1:R:244:ASP:O	2.18	0.42
1:Q:287:ILE:HD12	1:S:208:ILE:HD13	1.96	0.42
1:Q:349:LEU:HD23	1:Q:349:LEU:HA	1.80	0.42
1:Q:368:SER:O	1:Q:369:ILE:C	2.62	0.42
1:R:367:PRO:O	1:R:368:SER:C	2.61	0.42
1:S:8:LEU:CB	1:S:103:THR:HG23	2.48	0.42
1:S:104:LEU:HD13	1:S:347:ALA:HB2	2.01	0.42
1:U:106:THR:HG22	1:U:140:LEU:HD11	2.01	0.42
1:U:107:GLU:O	1:U:137:GLN:HG3	2.19	0.42
1:U:110:LEU:CD1	1:U:177:ARG:HH11	2.32	0.42
1:U:180:LEU:O	1:U:180:LEU:HG	2.11	0.42
1:U:369:ILE:C	1:U:371:HIS:N	2.77	0.42
1:V:136:ILE:O	1:V:139:VAL:HB	2.19	0.42
1:A:167:GLU:OE1	1:C:61:LYS:HE2	2.19	0.42
1:B:65:LEU:C	1:B:65:LEU:CD1	2.90	0.42
1:B:367:PRO:O	1:B:368:SER:C	2.61	0.42
1:C:34:ILE:CG2	1:C:67:LEU:CB	2.96	0.42
1:C:367:PRO:O	1:C:368:SER:C	2.61	0.42
1:D:44:MET:CG	1:D:45:VAL:N	2.75	0.42
1:D:104:LEU:HD13	1:D:347:ALA:HB2	2.01	0.42
1:D:176:MET:HB3	1:D:176:MET:HE3	1.81	0.42
1:D:178:LEU:HG	1:D:180:LEU:H	1.84	0.42
1:F:135:ALA:CB	1:F:140:LEU:HD21	2.49	0.42
1:F:136:ILE:O	1:F:139:VAL:HB	2.19	0.42
1:G:44:MET:CG	1:G:45:VAL:H	2.18	0.42
1:G:65:LEU:C	1:G:65:LEU:CD1	2.90	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:167:GLU:OE1	1:I:64:ILE:HD13	2.19	0.42
1:H:72:GLU:O	1:H:183:ARG:NH2	2.52	0.42
1:H:135:ALA:CB	1:H:140:LEU:HD21	2.49	0.42
1:H:149:THR:HG23	1:H:150:GLY:N	2.34	0.42
1:H:178:LEU:HG	1:H:180:LEU:H	1.84	0.42
1:H:367:PRO:O	1:H:368:SER:C	2.62	0.42
1:I:34:ILE:CG2	1:I:67:LEU:CB	2.96	0.42
1:I:69:TYR:CD2	1:I:69:TYR:N	2.87	0.42
1:I:72:GLU:O	1:I:183:ARG:NH2	2.52	0.42
1:I:368:SER:O	1:I:369:ILE:C	2.62	0.42
1:I:374:CYS:HB2	1:I:375:PHE:H	1.62	0.42
1:J:106:THR:HG22	1:J:140:LEU:HD11	2.01	0.42
1:J:136:ILE:O	1:J:139:VAL:HB	2.19	0.42
1:K:69:TYR:CD2	1:K:69:TYR:N	2.86	0.42
1:L:180:LEU:HD12	1:L:181:ALA:CA	2.50	0.42
1:M:64:ILE:CG2	1:M:65:LEU:N	2.81	0.42
1:N:34:ILE:HG23	1:N:67:LEU:HB3	2.01	0.42
1:N:180:LEU:HD12	1:N:181:ALA:CA	2.50	0.42
1:N:368:SER:O	1:N:369:ILE:C	2.62	0.42
1:Q:104:LEU:HD13	1:Q:347:ALA:HB2	2.01	0.42
1:Q:299:MET:HE2	1:Q:299:MET:HB3	1.95	0.42
1:Q:367:PRO:O	1:Q:368:SER:C	2.62	0.42
1:R:105:LEU:HD11	1:R:123:MET:HG3	2.01	0.42
1:R:180:LEU:HD12	1:R:181:ALA:CA	2.50	0.42
1:T:69:TYR:N	1:T:69:TYR:CD2	2.86	0.42
1:T:167:GLU:OE1	1:V:64:ILE:HD13	2.19	0.42
1:U:105:LEU:HD11	1:U:123:MET:HG3	2.01	0.42
1:U:178:LEU:HG	1:U:180:LEU:H	1.84	0.42
1:V:148:THR:CG2	1:V:149:THR:N	2.81	0.42
1:V:299:MET:HE2	1:V:299:MET:HB3	1.95	0.42
1:W:31:PHE:HA	1:W:32:PRO:HD3	1.66	0.42
1:A:105:LEU:HD11	1:A:123:MET:HG3	2.01	0.42
1:B:180:LEU:O	1:B:180:LEU:HG	2.11	0.42
1:C:106:THR:HG22	1:C:140:LEU:HD11	2.01	0.42
1:F:178:LEU:HG	1:F:180:LEU:H	1.84	0.42
1:G:34:ILE:HG23	1:G:67:LEU:HB3	2.02	0.42
1:G:105:LEU:HD11	1:G:123:MET:HG3	2.01	0.42
1:G:107:GLU:O	1:G:137:GLN:HG3	2.19	0.42
1:H:180:LEU:HD12	1:H:181:ALA:CA	2.50	0.42
1:H:299:MET:HE2	1:H:299:MET:HB3	1.95	0.42
1:I:104:LEU:HD13	1:I:347:ALA:HB2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:299:MET:O	1:I:332:PRO:HD2	2.18	0.42
1:J:110:LEU:CD1	1:J:177:ARG:HH11	2.32	0.42
1:K:73:HIC:HA	1:K:183:ARG:NH1	2.18	0.42
1:K:368:SER:O	1:K:369:ILE:C	2.62	0.42
1:L:8:LEU:CB	1:L:103:THR:HG23	2.48	0.42
1:M:148:THR:CG2	1:M:149:THR:N	2.81	0.42
1:M:167:GLU:OE1	1:O:64:ILE:HD13	2.19	0.42
1:M:173:HIS:HE1	1:N:268:GLY:CA	2.28	0.42
1:M:362:TYR:CE1	1:M:367:PRO:CB	2.99	0.42
1:N:178:LEU:HG	1:N:180:LEU:H	1.84	0.42
1:O:110:LEU:CD1	1:O:177:ARG:HH11	2.32	0.42
1:P:180:LEU:HD12	1:P:181:ALA:CA	2.50	0.42
1:Q:362:TYR:CE1	1:Q:367:PRO:CB	2.99	0.42
1:R:107:GLU:O	1:R:137:GLN:HG3	2.19	0.42
1:S:34:ILE:HG23	1:S:67:LEU:HB3	2.01	0.42
1:S:72:GLU:O	1:S:183:ARG:NH2	2.52	0.42
1:S:178:LEU:HG	1:S:180:LEU:H	1.84	0.42
1:T:105:LEU:HD11	1:T:123:MET:HG3	2.01	0.42
1:T:299:MET:HE2	1:T:299:MET:HB3	1.95	0.42
1:U:65:LEU:HD13	1:U:67:LEU:HD23	2.01	0.42
1:V:44:MET:CG	1:V:45:VAL:H	2.18	0.42
1:W:72:GLU:O	1:W:183:ARG:NH2	2.52	0.42
1:W:136:ILE:O	1:W:139:VAL:HB	2.19	0.42
1:W:148:THR:CG2	1:W:149:THR:N	2.81	0.42
1:W:178:LEU:HG	1:W:180:LEU:H	1.85	0.42
1:A:374:CYS:HB2	1:A:375:PHE:H	1.61	0.42
1:B:34:ILE:CG2	1:B:67:LEU:CB	2.96	0.42
1:B:34:ILE:HG23	1:B:67:LEU:HB3	2.02	0.42
1:B:104:LEU:HD13	1:B:347:ALA:HB2	2.01	0.42
1:B:176:MET:HE3	1:B:176:MET:HB3	1.81	0.42
1:C:167:GLU:OE1	1:E:61:LYS:HE2	2.20	0.42
1:C:300:SER:HA	1:C:335:ARG:CD	2.50	0.42
1:D:136:ILE:O	1:D:139:VAL:HB	2.19	0.42
1:E:34:ILE:HG23	1:E:67:LEU:HB3	2.01	0.42
1:E:65:LEU:HD13	1:E:67:LEU:HD23	2.01	0.42
1:E:300:SER:HA	1:E:335:ARG:CD	2.50	0.42
1:F:167:GLU:OE1	1:H:64:ILE:HD13	2.19	0.42
1:H:105:LEU:HD11	1:H:123:MET:HG3	2.01	0.42
1:I:335:ARG:HA	1:I:335:ARG:HD3	1.39	0.42
1:J:325:MET:HE3	1:L:244:ASP:O	2.18	0.42
1:J:368:SER:O	1:J:369:ILE:C	2.62	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:104:LEU:HD13	1:K:347:ALA:HB2	2.01	0.42
1:L:34:ILE:HG23	1:L:67:LEU:HB3	2.01	0.42
1:L:73:HIC:HA	1:L:183:ARG:NH1	2.18	0.42
1:M:2:GLU:HB3	1:M:3:ASP:H	1.65	0.42
1:M:72:GLU:O	1:M:183:ARG:NH2	2.52	0.42
1:N:72:GLU:O	1:N:183:ARG:NH2	2.52	0.42
1:O:72:GLU:O	1:O:183:ARG:NH2	2.52	0.42
1:O:105:LEU:HD11	1:O:123:MET:HG3	2.01	0.42
1:O:368:SER:O	1:O:369:ILE:C	2.62	0.42
1:P:72:GLU:O	1:P:183:ARG:NH2	2.52	0.42
1:P:374:CYS:HB2	1:P:375:PHE:H	1.61	0.42
1:Q:110:LEU:CD1	1:Q:177:ARG:HH11	2.32	0.42
1:Q:136:ILE:O	1:Q:139:VAL:HB	2.19	0.42
1:Q:180:LEU:O	1:Q:180:LEU:HG	2.11	0.42
1:R:69:TYR:CD2	1:R:69:TYR:N	2.86	0.42
1:R:167:GLU:OE1	1:T:64:ILE:HD13	2.19	0.42
1:R:178:LEU:HG	1:R:180:LEU:H	1.84	0.42
1:S:107:GLU:O	1:S:137:GLN:HG3	2.19	0.42
1:S:149:THR:HG23	1:S:150:GLY:N	2.34	0.42
1:S:167:GLU:OE1	1:U:64:ILE:HD13	2.19	0.42
1:S:167:GLU:OE1	1:U:61:LYS:HE2	2.20	0.42
1:T:72:GLU:O	1:T:183:ARG:NH2	2.52	0.42
1:T:107:GLU:O	1:T:137:GLN:HG3	2.19	0.42
1:T:180:LEU:HD12	1:T:181:ALA:CA	2.50	0.42
1:U:183:ARG:HE	1:U:183:ARG:HB3	1.58	0.42
1:V:107:GLU:O	1:V:137:GLN:HG3	2.19	0.42
1:V:369:ILE:C	1:V:371:HIS:N	2.77	0.42
1:W:135:ALA:CB	1:W:140:LEU:HD21	2.50	0.42
1:W:180:LEU:CD1	1:W:181:ALA:N	2.78	0.42
1:A:64:ILE:CG2	1:A:65:LEU:N	2.82	0.42
1:A:121:GLN:HG3	1:A:362:TYR:OH	2.20	0.42
1:A:178:LEU:HG	1:A:180:LEU:H	1.84	0.42
1:B:167:GLU:OE1	1:D:61:LYS:HE2	2.19	0.42
1:B:299:MET:O	1:B:332:PRO:HD2	2.18	0.42
1:C:178:LEU:HG	1:C:180:LEU:H	1.84	0.42
1:D:105:LEU:HD11	1:D:123:MET:HG3	2.01	0.42
1:D:106:THR:HG22	1:D:140:LEU:HD11	2.01	0.42
1:D:148:THR:CG2	1:D:149:THR:N	2.81	0.42
1:E:357:ILE:HD12	1:E:357:ILE:HA	1.78	0.42
1:F:299:MET:O	1:F:332:PRO:HD2	2.18	0.42
1:F:325:MET:HE3	1:H:244:ASP:O	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:368:SER:O	1:F:369:ILE:C	2.62	0.42
1:G:8:LEU:CB	1:G:103:THR:HG23	2.48	0.42
1:G:64:ILE:CG2	1:G:65:LEU:N	2.81	0.42
1:H:65:LEU:HD13	1:H:67:LEU:HD23	2.01	0.42
1:H:110:LEU:CD1	1:H:177:ARG:HH11	2.32	0.42
1:I:34:ILE:HG23	1:I:67:LEU:HB3	2.02	0.42
1:I:180:LEU:HD12	1:I:181:ALA:CA	2.50	0.42
1:K:65:LEU:HD13	1:K:67:LEU:HD23	2.01	0.42
1:K:180:LEU:HD12	1:K:181:ALA:CA	2.50	0.42
1:L:300:SER:HA	1:L:335:ARG:CD	2.50	0.42
1:M:105:LEU:HD11	1:M:123:MET:HG3	2.01	0.42
1:M:180:LEU:HD12	1:M:181:ALA:CA	2.50	0.42
1:O:104:LEU:HD13	1:O:347:ALA:HB2	2.01	0.42
1:O:136:ILE:O	1:O:139:VAL:HB	2.19	0.42
1:P:34:ILE:HG23	1:P:67:LEU:HB3	2.01	0.42
1:Q:105:LEU:HD11	1:Q:123:MET:HG3	2.01	0.42
1:T:2:GLU:HB3	1:T:3:ASP:H	1.66	0.42
1:U:148:THR:CG2	1:U:149:THR:N	2.81	0.42
1:V:374:CYS:HB2	1:V:375:PHE:H	1.61	0.42
1:A:72:GLU:O	1:A:183:ARG:NH2	2.52	0.42
1:A:135:ALA:CB	1:A:140:LEU:HD21	2.49	0.42
1:A:368:SER:O	1:A:369:ILE:C	2.62	0.42
1:B:72:GLU:O	1:B:183:ARG:NH2	2.52	0.42
1:B:106:THR:HG22	1:B:140:LEU:HD11	2.01	0.42
1:B:107:GLU:O	1:B:137:GLN:HG3	2.19	0.42
1:B:149:THR:HG23	1:B:150:GLY:N	2.34	0.42
1:C:368:SER:O	1:C:369:ILE:C	2.62	0.42
1:D:135:ALA:CB	1:D:140:LEU:HD21	2.49	0.42
1:D:149:THR:HG23	1:D:150:GLY:N	2.34	0.42
1:D:362:TYR:CE1	1:D:367:PRO:CB	2.99	0.42
1:F:110:LEU:CD1	1:F:177:ARG:HH11	2.32	0.42
1:F:180:LEU:HD12	1:F:181:ALA:CA	2.50	0.42
1:G:236:LEU:CD1	1:G:237:GLU:CG	2.93	0.42
1:G:368:SER:O	1:G:369:ILE:C	2.62	0.42
1:H:180:LEU:CD1	1:H:181:ALA:N	2.78	0.42
1:J:35:VAL:H	1:J:68:LYS:H	1.65	0.42
1:J:135:ALA:CB	1:J:140:LEU:HD21	2.50	0.42
1:J:167:GLU:OE1	1:L:64:ILE:HD13	2.19	0.42
1:K:2:GLU:HB3	1:K:3:ASP:H	1.66	0.42
1:K:136:ILE:O	1:K:139:VAL:HB	2.19	0.42
1:K:149:THR:CG2	1:K:150:GLY:N	2.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:69:TYR:CD2	1:M:69:TYR:N	2.86	0.42
1:N:65:LEU:HD13	1:N:67:LEU:HD23	2.01	0.42
1:N:167:GLU:OE1	1:P:61:LYS:HE2	2.19	0.42
1:O:180:LEU:HD12	1:O:181:ALA:CA	2.50	0.42
1:Q:180:LEU:HD12	1:Q:181:ALA:CA	2.50	0.42
1:S:105:LEU:HD11	1:S:123:MET:HG3	2.01	0.42
1:S:110:LEU:CD1	1:S:177:ARG:HH11	2.32	0.42
1:S:299:MET:O	1:S:332:PRO:HD2	2.18	0.42
1:T:35:VAL:H	1:T:68:LYS:H	1.65	0.42
1:T:65:LEU:HD13	1:T:67:LEU:HD23	2.01	0.42
1:U:121:GLN:HG3	1:U:362:TYR:OH	2.20	0.42
1:U:136:ILE:O	1:U:139:VAL:HB	2.19	0.42
1:U:176:MET:HE3	1:U:176:MET:HB3	1.81	0.42
1:U:299:MET:O	1:U:332:PRO:HD2	2.18	0.42
1:V:104:LEU:HD13	1:V:347:ALA:HB2	2.01	0.42
1:V:176:MET:HB3	1:V:176:MET:HE3	1.81	0.42
1:V:180:LEU:HD12	1:V:181:ALA:CA	2.50	0.42
1:V:368:SER:O	1:V:369:ILE:C	2.62	0.42
1:W:121:GLN:HG3	1:W:362:TYR:OH	2.20	0.42
1:A:65:LEU:HD13	1:A:67:LEU:HD23	2.01	0.42
1:B:105:LEU:HD11	1:B:123:MET:HG3	2.02	0.42
1:B:136:ILE:O	1:B:139:VAL:HB	2.19	0.42
1:B:149:THR:CG2	1:B:150:GLY:N	2.82	0.42
1:B:362:TYR:CE1	1:B:367:PRO:CB	2.99	0.42
1:D:72:GLU:O	1:D:183:ARG:NH2	2.52	0.42
1:D:368:SER:O	1:D:369:ILE:C	2.62	0.42
1:E:72:GLU:O	1:E:183:ARG:NH2	2.52	0.42
1:F:105:LEU:HD11	1:F:123:MET:HG3	2.02	0.42
1:G:180:LEU:HD12	1:G:181:ALA:CA	2.50	0.42
1:G:300:SER:HA	1:G:335:ARG:CD	2.50	0.42
1:H:133:TYR:CE2	1:H:375:PHE:HB2	2.51	0.42
1:H:136:ILE:O	1:H:139:VAL:HB	2.19	0.42
1:I:39:ARG:NE	1:I:66:THR:CA	2.69	0.42
1:I:167:GLU:OE1	1:K:64:ILE:HD13	2.19	0.42
1:I:299:MET:HE2	1:I:299:MET:HB3	1.95	0.42
1:K:110:LEU:CD1	1:K:177:ARG:HH11	2.32	0.42
1:L:72:GLU:O	1:L:183:ARG:NH2	2.52	0.42
1:L:121:GLN:HG3	1:L:362:TYR:OH	2.20	0.42
1:L:167:GLU:OE1	1:N:64:ILE:HD13	2.19	0.42
1:L:325:MET:HE3	1:N:244:ASP:O	2.18	0.42
1:L:368:SER:O	1:L:369:ILE:C	2.62	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:8:LEU:N	1:O:102:PRO:O	2.48	0.42
1:P:106:THR:HG22	1:P:140:LEU:HD11	2.01	0.42
1:P:167:GLU:CD	1:R:61:LYS:CE	2.85	0.42
1:Q:171:LEU:HA	1:Q:172:PRO:HD3	1.79	0.42
1:Q:173:HIS:HE1	1:R:268:GLY:CA	2.28	0.42
1:R:65:LEU:HD13	1:R:67:LEU:HD23	2.01	0.42
1:S:121:GLN:HG3	1:S:362:TYR:OH	2.20	0.42
1:S:362:TYR:CE1	1:S:367:PRO:CB	2.99	0.42
1:T:368:SER:O	1:T:369:ILE:C	2.62	0.42
1:U:287:ILE:HD12	1:W:208:ILE:HD13	1.96	0.42
1:U:368:SER:O	1:U:369:ILE:C	2.62	0.42
1:V:135:ALA:CB	1:V:140:LEU:HD21	2.49	0.42
1:W:367:PRO:O	1:W:368:SER:C	2.62	0.42
1:A:38:PRO:CG	1:A:49:GLN:NE2	2.79	0.42
1:B:180:LEU:CD1	1:B:181:ALA:N	2.78	0.42
1:C:121:GLN:HG3	1:C:362:TYR:OH	2.20	0.42
1:D:167:GLU:OE1	1:F:64:ILE:HD13	2.19	0.42
1:E:167:GLU:OE1	1:G:61:LYS:HE2	2.20	0.42
1:E:180:LEU:HD12	1:E:181:ALA:CA	2.49	0.42
1:F:180:LEU:O	1:F:180:LEU:HG	2.11	0.42
1:J:107:GLU:O	1:J:137:GLN:HG3	2.19	0.42
1:J:121:GLN:HG3	1:J:362:TYR:OH	2.20	0.42
1:J:148:THR:CG2	1:J:149:THR:N	2.81	0.42
1:J:369:ILE:C	1:J:371:HIS:N	2.77	0.42
1:K:34:ILE:CG2	1:K:67:LEU:CB	2.96	0.42
1:K:110:LEU:CD1	1:K:177:ARG:NH1	2.80	0.42
1:L:65:LEU:HD13	1:L:67:LEU:HD23	2.01	0.42
1:M:104:LEU:HD13	1:M:347:ALA:HB2	2.01	0.42
1:M:133:TYR:CE2	1:M:375:PHE:HB2	2.51	0.42
1:N:103:THR:O	1:N:132:MET:HA	2.20	0.42
1:O:149:THR:HG23	1:O:150:GLY:N	2.34	0.42
1:O:167:GLU:OE1	1:Q:64:ILE:HD13	2.19	0.42
1:Q:34:ILE:HG23	1:Q:67:LEU:HB3	2.02	0.42
1:R:368:SER:O	1:R:369:ILE:C	2.62	0.42
1:S:180:LEU:HD12	1:S:181:ALA:CA	2.50	0.42
1:U:132:MET:HE2	1:U:132:MET:HB2	1.74	0.42
1:U:180:LEU:HD12	1:U:181:ALA:CA	2.49	0.42
1:V:121:GLN:HG3	1:V:362:TYR:OH	2.20	0.42
1:W:65:LEU:HD13	1:W:67:LEU:HD23	2.01	0.42
1:W:176:MET:HE3	1:W:176:MET:HB3	1.81	0.42
1:A:167:GLU:OE1	1:C:64:ILE:HD13	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:299:MET:HE2	1:A:299:MET:HB3	1.95	0.42
1:A:325:MET:HE3	1:C:244:ASP:O	2.18	0.42
1:C:34:ILE:HG23	1:C:67:LEU:HB3	2.01	0.42
1:C:148:THR:CG2	1:C:149:THR:N	2.81	0.42
1:E:103:THR:O	1:E:132:MET:HA	2.20	0.42
1:E:106:THR:HG22	1:E:140:LEU:HD11	2.01	0.42
1:F:106:THR:HG22	1:F:140:LEU:HD11	2.01	0.42
1:F:132:MET:HE2	1:F:132:MET:HB2	1.75	0.42
1:G:65:LEU:HD13	1:G:67:LEU:HD23	2.01	0.42
1:I:178:LEU:HG	1:I:180:LEU:H	1.84	0.42
1:J:178:LEU:HG	1:J:180:LEU:H	1.84	0.42
1:L:47:MET:HB3	1:L:48:GLY:H	1.61	0.42
1:L:103:THR:O	1:L:132:MET:HA	2.20	0.42
1:M:110:LEU:CD1	1:M:177:ARG:NH1	2.80	0.42
1:M:180:LEU:CD1	1:M:181:ALA:N	2.78	0.42
1:N:121:GLN:HG3	1:N:362:TYR:OH	2.20	0.42
1:O:64:ILE:CG2	1:O:65:LEU:N	2.81	0.42
1:O:178:LEU:HG	1:O:180:LEU:H	1.84	0.42
1:O:180:LEU:CD1	1:O:181:ALA:N	2.78	0.42
1:P:107:GLU:O	1:P:137:GLN:HG3	2.19	0.42
1:P:299:MET:O	1:P:332:PRO:HD2	2.18	0.42
1:P:368:SER:O	1:P:369:ILE:C	2.62	0.42
1:Q:72:GLU:O	1:Q:183:ARG:NH2	2.52	0.42
1:T:135:ALA:CB	1:T:140:LEU:HD21	2.49	0.42
1:U:135:ALA:CB	1:U:140:LEU:HD21	2.50	0.42
1:U:300:SER:HA	1:U:335:ARG:CD	2.50	0.42
1:A:34:ILE:CG2	1:A:67:LEU:CB	2.96	0.41
1:B:8:LEU:CB	1:B:103:THR:HG23	2.48	0.41
1:C:133:TYR:CE2	1:C:375:PHE:HB2	2.51	0.41
1:C:167:GLU:OE1	1:E:64:ILE:HD13	2.19	0.41
1:C:325:MET:HE3	1:E:244:ASP:O	2.18	0.41
1:E:69:TYR:CD2	1:E:69:TYR:N	2.87	0.41
1:G:103:THR:O	1:G:132:MET:HA	2.20	0.41
1:G:178:LEU:HG	1:G:180:LEU:H	1.84	0.41
1:H:106:THR:HG22	1:H:140:LEU:HD11	2.01	0.41
1:K:140:LEU:HB3	1:K:343:GLY:N	2.35	0.41
1:K:173:HIS:HE1	1:L:268:GLY:CA	2.28	0.41
1:M:178:LEU:HG	1:M:180:LEU:H	1.84	0.41
1:P:103:THR:O	1:P:132:MET:HA	2.20	0.41
1:P:178:LEU:HG	1:P:180:LEU:H	1.84	0.41
1:Q:121:GLN:HG3	1:Q:362:TYR:OH	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:149:THR:CG2	1:Q:150:GLY:N	2.82	0.41
1:R:140:LEU:HB3	1:R:343:GLY:N	2.35	0.41
1:S:106:THR:HG22	1:S:140:LEU:HD11	2.01	0.41
1:S:325:MET:HE3	1:U:244:ASP:O	2.18	0.41
1:V:103:THR:O	1:V:132:MET:HA	2.20	0.41
1:V:133:TYR:CE2	1:V:375:PHE:HB2	2.51	0.41
1:W:103:THR:O	1:W:132:MET:HA	2.20	0.41
1:W:140:LEU:HB3	1:W:343:GLY:N	2.35	0.41
1:W:369:ILE:C	1:W:371:HIS:N	2.77	0.41
1:A:171:LEU:HA	1:A:172:PRO:HD3	1.79	0.41
1:A:189:LEU:HA	1:A:192:ILE:HG12	2.02	0.41
1:C:38:PRO:CG	1:C:49:GLN:NE2	2.79	0.41
1:D:107:GLU:O	1:D:137:GLN:HG3	2.19	0.41
1:D:121:GLN:HA	1:D:362:TYR:CZ	2.55	0.41
1:D:167:GLU:OE1	1:F:61:LYS:HE2	2.20	0.41
1:E:107:GLU:O	1:E:137:GLN:HG3	2.19	0.41
1:E:121:GLN:HG3	1:E:362:TYR:OH	2.20	0.41
1:E:325:MET:HE3	1:G:244:ASP:O	2.18	0.41
1:E:368:SER:O	1:E:369:ILE:C	2.62	0.41
1:F:72:GLU:O	1:F:183:ARG:NH2	2.52	0.41
1:G:167:GLU:OE1	1:I:61:LYS:HE2	2.19	0.41
1:H:34:ILE:CG2	1:H:67:LEU:CB	2.96	0.41
1:H:121:GLN:HG3	1:H:362:TYR:OH	2.20	0.41
1:I:65:LEU:HD13	1:I:67:LEU:HD23	2.01	0.41
1:J:34:ILE:HG23	1:J:67:LEU:HB3	2.01	0.41
1:J:47:MET:HB3	1:J:48:GLY:H	1.62	0.41
1:J:65:LEU:HD13	1:J:67:LEU:HD23	2.01	0.41
1:J:72:GLU:O	1:J:183:ARG:NH2	2.52	0.41
1:L:167:GLU:OE1	1:N:61:LYS:HE2	2.19	0.41
1:N:110:LEU:CD1	1:N:177:ARG:HH11	2.32	0.41
1:N:189:LEU:HA	1:N:192:ILE:HG12	2.02	0.41
1:N:300:SER:HA	1:N:335:ARG:CD	2.50	0.41
1:O:106:THR:HG22	1:O:140:LEU:HD11	2.01	0.41
1:O:173:HIS:HE1	1:P:268:GLY:CA	2.28	0.41
1:Q:106:THR:HG22	1:Q:140:LEU:HD11	2.01	0.41
1:Q:121:GLN:HA	1:Q:362:TYR:CZ	2.55	0.41
1:Q:140:LEU:HB3	1:Q:343:GLY:N	2.35	0.41
1:Q:180:LEU:CD1	1:Q:181:ALA:N	2.78	0.41
1:R:34:ILE:HG23	1:R:67:LEU:HB3	2.01	0.41
1:S:121:GLN:HA	1:S:362:TYR:CZ	2.55	0.41
1:S:368:SER:O	1:S:369:ILE:C	2.62	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:34:ILE:CG2	1:T:67:LEU:CB	2.96	0.41
1:T:121:GLN:HG3	1:T:362:TYR:OH	2.20	0.41
1:T:140:LEU:HB3	1:T:343:GLY:N	2.36	0.41
1:U:103:THR:O	1:U:132:MET:HA	2.20	0.41
1:U:140:LEU:HB3	1:U:343:GLY:N	2.36	0.41
1:U:167:GLU:OE1	1:W:64:ILE:HD13	2.19	0.41
1:V:34:ILE:HG23	1:V:67:LEU:HB3	2.02	0.41
1:V:65:LEU:HD13	1:V:67:LEU:HD23	2.01	0.41
1:V:69:TYR:CD2	1:V:69:TYR:N	2.86	0.41
1:V:73:HIC:HA	1:V:183:ARG:NH1	2.18	0.41
1:A:148:THR:CG2	1:A:149:THR:N	2.81	0.41
1:B:121:GLN:HA	1:B:362:TYR:CZ	2.55	0.41
1:B:121:GLN:HG3	1:B:362:TYR:OH	2.20	0.41
1:C:65:LEU:HD13	1:C:67:LEU:HD23	2.01	0.41
1:C:103:THR:O	1:C:132:MET:HA	2.20	0.41
1:C:189:LEU:HA	1:C:192:ILE:HG12	2.02	0.41
1:D:132:MET:HB2	1:D:132:MET:HE2	1.75	0.41
1:D:180:LEU:HD12	1:D:181:ALA:CA	2.50	0.41
1:E:121:GLN:HA	1:E:362:TYR:CZ	2.55	0.41
1:E:167:GLU:OE1	1:G:64:ILE:HD13	2.19	0.41
1:F:140:LEU:HB3	1:F:343:GLY:N	2.35	0.41
1:H:121:GLN:HA	1:H:362:TYR:CZ	2.55	0.41
1:I:121:GLN:HA	1:I:362:TYR:CZ	2.55	0.41
1:I:369:ILE:C	1:I:371:HIS:N	2.77	0.41
1:J:103:THR:O	1:J:132:MET:HA	2.20	0.41
1:J:121:GLN:HA	1:J:362:TYR:CZ	2.55	0.41
1:J:335:ARG:O	1:J:338:SER:CB	2.69	0.41
1:N:33:SER:C	1:N:34:ILE:HG13	2.46	0.41
1:N:167:GLU:OE1	1:P:64:ILE:HD13	2.19	0.41
1:O:34:ILE:HG23	1:O:67:LEU:HB3	2.01	0.41
1:O:140:LEU:HB3	1:O:343:GLY:N	2.35	0.41
1:O:349:LEU:HD23	1:O:349:LEU:HA	1.80	0.41
1:P:132:MET:HE2	1:P:132:MET:HB2	1.75	0.41
1:P:140:LEU:HB3	1:P:343:GLY:N	2.35	0.41
1:P:149:THR:CG2	1:P:150:GLY:N	2.82	0.41
1:Q:335:ARG:O	1:Q:338:SER:CB	2.69	0.41
1:S:103:THR:O	1:S:132:MET:HA	2.20	0.41
1:S:140:LEU:HB3	1:S:343:GLY:N	2.35	0.41
1:S:374:CYS:HB2	1:S:375:PHE:H	1.61	0.41
1:T:103:THR:O	1:T:132:MET:HA	2.20	0.41
1:V:34:ILE:CG2	1:V:67:LEU:CB	2.96	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:140:LEU:HB3	1:V:343:GLY:N	2.35	0.41
1:W:8:LEU:N	1:W:102:PRO:O	2.48	0.41
1:W:368:SER:O	1:W:369:ILE:C	2.62	0.41
1:A:140:LEU:HB3	1:A:343:GLY:N	2.36	0.41
1:B:135:ALA:CB	1:B:140:LEU:HD21	2.49	0.41
1:C:33:SER:C	1:C:34:ILE:HG13	2.46	0.41
1:C:180:LEU:HD12	1:C:181:ALA:CA	2.50	0.41
1:D:33:SER:C	1:D:34:ILE:HG13	2.46	0.41
1:D:183:ARG:HE	1:D:183:ARG:HB3	1.59	0.41
1:E:31:PHE:HA	1:E:32:PRO:HD3	1.66	0.41
1:E:171:LEU:HA	1:E:172:PRO:HD3	1.79	0.41
1:F:33:SER:C	1:F:34:ILE:HG13	2.46	0.41
1:F:236:LEU:HD12	1:F:237:GLU:CA	2.51	0.41
1:G:33:SER:C	1:G:34:ILE:HG13	2.46	0.41
1:G:121:GLN:HA	1:G:362:TYR:CZ	2.56	0.41
1:G:369:ILE:C	1:G:371:HIS:N	2.77	0.41
1:H:335:ARG:O	1:H:338:SER:CB	2.69	0.41
1:H:368:SER:O	1:H:369:ILE:C	2.62	0.41
1:H:374:CYS:HB2	1:H:375:PHE:H	1.61	0.41
1:I:110:LEU:CD1	1:I:177:ARG:NH1	2.80	0.41
1:I:140:LEU:HB3	1:I:343:GLY:N	2.35	0.41
1:I:300:SER:HA	1:I:335:ARG:CD	2.50	0.41
1:M:33:SER:C	1:M:34:ILE:HG13	2.46	0.41
1:M:106:THR:HG22	1:M:140:LEU:HD11	2.01	0.41
1:M:135:ALA:CB	1:M:140:LEU:HD21	2.49	0.41
1:N:31:PHE:HA	1:N:32:PRO:HD3	1.66	0.41
1:O:110:LEU:CD1	1:O:177:ARG:NH1	2.80	0.41
1:O:121:GLN:HG3	1:O:362:TYR:OH	2.20	0.41
1:P:167:GLU:OE1	1:R:64:ILE:HD13	2.19	0.41
1:P:176:MET:HB3	1:P:176:MET:HE3	1.81	0.41
1:P:189:LEU:HA	1:P:192:ILE:HG12	2.02	0.41
1:R:2:GLU:HB3	1:R:3:ASP:H	1.66	0.41
1:R:33:SER:C	1:R:34:ILE:HG13	2.46	0.41
1:R:189:LEU:HA	1:R:192:ILE:HG12	2.02	0.41
1:S:171:LEU:HA	1:S:172:PRO:HD3	1.79	0.41
1:T:189:LEU:HA	1:T:192:ILE:HG12	2.02	0.41
1:U:369:ILE:HG23	1:U:370:VAL:N	2.30	0.41
1:V:110:LEU:CD1	1:V:177:ARG:HH11	2.32	0.41
1:W:33:SER:C	1:W:34:ILE:HG13	2.46	0.41
1:W:121:GLN:HA	1:W:362:TYR:CZ	2.55	0.41
1:W:180:LEU:HD12	1:W:181:ALA:CA	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:110:LEU:CD1	1:B:177:ARG:HH11	2.32	0.41
1:B:140:LEU:HB3	1:B:343:GLY:N	2.35	0.41
1:B:335:ARG:O	1:B:338:SER:CB	2.69	0.41
1:C:69:TYR:CD2	1:C:69:TYR:N	2.86	0.41
1:C:140:LEU:HB3	1:C:343:GLY:N	2.35	0.41
1:C:299:MET:HE2	1:C:299:MET:HB3	1.95	0.41
1:E:2:GLU:HB3	1:E:3:ASP:H	1.66	0.41
1:E:135:ALA:CB	1:E:140:LEU:HD21	2.49	0.41
1:E:369:ILE:C	1:E:371:HIS:N	2.77	0.41
1:I:103:THR:O	1:I:132:MET:HA	2.20	0.41
1:I:335:ARG:O	1:I:338:SER:CB	2.69	0.41
1:L:189:LEU:HA	1:L:192:ILE:HG12	2.02	0.41
1:L:335:ARG:HD3	1:L:335:ARG:HA	1.39	0.41
1:M:121:GLN:HG3	1:M:362:TYR:OH	2.20	0.41
1:M:236:LEU:HD12	1:M:237:GLU:CA	2.51	0.41
1:O:103:THR:O	1:O:132:MET:HA	2.20	0.41
1:P:64:ILE:CG2	1:P:65:LEU:N	2.82	0.41
1:P:65:LEU:HD13	1:P:67:LEU:HD23	2.01	0.41
1:P:135:ALA:CB	1:P:140:LEU:HD21	2.49	0.41
1:P:349:LEU:HD23	1:P:349:LEU:HA	1.80	0.41
1:Q:103:THR:O	1:Q:132:MET:HA	2.20	0.41
1:Q:167:GLU:OE1	1:S:64:ILE:HD13	2.19	0.41
1:Q:325:MET:HE3	1:S:244:ASP:O	2.18	0.41
1:R:34:ILE:HD13	1:R:67:LEU:CD2	2.46	0.41
1:R:135:ALA:CB	1:R:140:LEU:HD21	2.49	0.41
1:S:34:ILE:CG2	1:S:67:LEU:CB	2.96	0.41
1:S:180:LEU:CD1	1:S:181:ALA:N	2.78	0.41
1:S:357:ILE:CG2	1:S:358:THR:N	2.82	0.41
1:U:72:GLU:O	1:U:183:ARG:NH2	2.52	0.41
1:U:180:LEU:CD1	1:U:181:ALA:N	2.78	0.41
1:U:335:ARG:O	1:U:338:SER:CB	2.69	0.41
1:V:171:LEU:HA	1:V:172:PRO:HD3	1.79	0.41
1:A:180:LEU:HD12	1:A:181:ALA:CA	2.50	0.41
1:B:236:LEU:HD12	1:B:237:GLU:CA	2.51	0.41
1:C:121:GLN:HA	1:C:362:TYR:CZ	2.55	0.41
1:C:369:ILE:C	1:C:371:HIS:N	2.77	0.41
1:D:121:GLN:HG3	1:D:362:TYR:OH	2.20	0.41
1:E:189:LEU:HA	1:E:192:ILE:HG12	2.02	0.41
1:F:223:PHE:CZ	1:F:266:PHE:CZ	3.09	0.41
1:G:106:THR:HG22	1:G:140:LEU:HD11	2.01	0.41
1:G:135:ALA:CB	1:G:140:LEU:HD21	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:189:LEU:HA	1:G:192:ILE:HG12	2.02	0.41
1:H:33:SER:C	1:H:34:ILE:HG13	2.46	0.41
1:I:34:ILE:CG2	1:I:67:LEU:HD22	2.28	0.41
1:J:133:TYR:CE2	1:J:375:PHE:HB2	2.51	0.41
1:J:167:GLU:OE1	1:L:61:LYS:HE2	2.20	0.41
1:K:121:GLN:HA	1:K:362:TYR:CZ	2.55	0.41
1:K:335:ARG:O	1:K:338:SER:CB	2.69	0.41
1:M:34:ILE:CG2	1:M:67:LEU:CB	2.96	0.41
1:M:121:GLN:HA	1:M:362:TYR:CZ	2.55	0.41
1:M:335:ARG:O	1:M:338:SER:CB	2.69	0.41
1:N:135:ALA:CB	1:N:140:LEU:HD21	2.49	0.41
1:N:140:LEU:HB3	1:N:343:GLY:N	2.35	0.41
1:O:299:MET:HE2	1:O:299:MET:HB3	1.95	0.41
1:P:121:GLN:HG3	1:P:362:TYR:OH	2.20	0.41
1:S:135:ALA:CB	1:S:140:LEU:HD21	2.49	0.41
1:S:335:ARG:O	1:S:338:SER:CB	2.69	0.41
1:U:31:PHE:HA	1:U:32:PRO:HD3	1.66	0.41
1:V:178:LEU:HG	1:V:180:LEU:H	1.84	0.41
1:W:47:MET:HB3	1:W:48:GLY:H	1.61	0.41
1:W:236:LEU:HD12	1:W:237:GLU:CA	2.51	0.41
1:A:103:THR:O	1:A:132:MET:HA	2.20	0.41
1:A:149:THR:CG2	1:A:150:GLY:N	2.82	0.41
1:B:167:GLU:OE1	1:D:64:ILE:HD13	2.19	0.41
1:C:357:ILE:HD12	1:C:357:ILE:HA	1.79	0.41
1:D:8:LEU:N	1:D:102:PRO:O	2.48	0.41
1:D:140:LEU:HB3	1:D:343:GLY:N	2.35	0.41
1:D:335:ARG:O	1:D:338:SER:CB	2.69	0.41
1:F:57:GLU:O	1:F:58:ALA:C	2.64	0.41
1:F:121:GLN:HA	1:F:362:TYR:CZ	2.55	0.41
1:F:121:GLN:HG3	1:F:362:TYR:OH	2.20	0.41
1:F:299:MET:HE2	1:F:299:MET:HB3	1.95	0.41
1:F:335:ARG:O	1:F:338:SER:CB	2.69	0.41
1:G:110:LEU:CD1	1:G:177:ARG:HH11	2.32	0.41
1:G:121:GLN:HG3	1:G:362:TYR:OH	2.20	0.41
1:G:236:LEU:HD12	1:G:237:GLU:CA	2.51	0.41
1:H:8:LEU:N	1:H:102:PRO:O	2.48	0.41
1:H:369:ILE:C	1:H:371:HIS:N	2.77	0.41
1:I:8:LEU:CD1	1:I:90:PHE:CD1	3.04	0.41
1:I:57:GLU:O	1:I:58:ALA:C	2.64	0.41
1:I:106:THR:HG22	1:I:140:LEU:HD11	2.01	0.41
1:I:236:LEU:HD12	1:I:237:GLU:CA	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:33:SER:C	1:K:34:ILE:HG13	2.46	0.41
1:K:121:GLN:HG3	1:K:362:TYR:OH	2.20	0.41
1:L:236:LEU:HD12	1:L:237:GLU:CA	2.51	0.41
1:M:110:LEU:CD1	1:M:177:ARG:HH11	2.32	0.41
1:M:183:ARG:HE	1:M:183:ARG:HB3	1.58	0.41
1:N:38:PRO:CG	1:N:49:GLN:NE2	2.79	0.41
1:N:121:GLN:HA	1:N:362:TYR:CZ	2.55	0.41
1:N:367:PRO:O	1:N:368:SER:C	2.61	0.41
1:O:33:SER:C	1:O:34:ILE:HG13	2.46	0.41
1:O:176:MET:HE3	1:O:176:MET:HB3	1.81	0.41
1:P:65:LEU:C	1:P:65:LEU:CD1	2.90	0.41
1:P:121:GLN:HA	1:P:362:TYR:CZ	2.55	0.41
1:Q:110:LEU:CD1	1:Q:177:ARG:NH1	2.80	0.41
1:Q:149:THR:HG23	1:Q:150:GLY:N	2.34	0.41
1:Q:236:LEU:HD12	1:Q:237:GLU:CA	2.51	0.41
1:R:106:THR:HG22	1:R:140:LEU:HD11	2.01	0.41
1:R:110:LEU:CD1	1:R:177:ARG:HH11	2.32	0.41
1:R:325:MET:HE3	1:T:244:ASP:O	2.18	0.41
1:R:362:TYR:CE1	1:R:367:PRO:CB	2.99	0.41
1:T:33:SER:C	1:T:34:ILE:HG13	2.46	0.41
1:T:335:ARG:O	1:T:338:SER:CB	2.69	0.41
1:U:33:SER:C	1:U:34:ILE:HG13	2.46	0.41
1:W:110:LEU:CD1	1:W:177:ARG:HH11	2.32	0.41
1:W:335:ARG:O	1:W:338:SER:CB	2.69	0.41
1:A:33:SER:C	1:A:34:ILE:HG13	2.46	0.41
1:A:34:ILE:HG23	1:A:67:LEU:HB3	2.01	0.41
1:B:33:SER:C	1:B:34:ILE:HG13	2.46	0.41
1:B:180:LEU:HD12	1:B:181:ALA:CA	2.50	0.41
1:B:357:ILE:HD12	1:B:357:ILE:HA	1.79	0.41
1:C:110:LEU:CD1	1:C:177:ARG:HH11	2.32	0.41
1:C:135:ALA:CB	1:C:140:LEU:HD21	2.49	0.41
1:D:300:SER:HA	1:D:335:ARG:CD	2.50	0.41
1:H:34:ILE:HG23	1:H:67:LEU:HB3	2.01	0.41
1:I:8:LEU:CB	1:I:103:THR:HG23	2.49	0.41
1:I:33:SER:C	1:I:34:ILE:HG13	2.46	0.41
1:I:121:GLN:HG3	1:I:362:TYR:OH	2.20	0.41
1:J:149:THR:CG2	1:J:150:GLY:N	2.82	0.41
1:K:106:THR:HG22	1:K:140:LEU:HD11	2.01	0.41
1:O:39:ARG:NE	1:O:66:THR:CA	2.69	0.41
1:O:148:THR:CG2	1:O:149:THR:N	2.81	0.41
1:P:142:LEU:CD1	1:P:165:ILE:HD11	2.47	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:300:SER:HA	1:P:335:ARG:CD	2.50	0.41
1:P:335:ARG:O	1:P:338:SER:CB	2.69	0.41
1:Q:57:GLU:O	1:Q:58:ALA:C	2.64	0.41
1:R:236:LEU:HD12	1:R:237:GLU:CA	2.51	0.41
1:S:33:SER:C	1:S:34:ILE:HG13	2.46	0.41
1:S:39:ARG:NE	1:S:66:THR:CA	2.69	0.41
1:S:236:LEU:HD12	1:S:237:GLU:CA	2.51	0.41
1:T:236:LEU:HD12	1:T:237:GLU:CA	2.51	0.41
1:U:8:LEU:N	1:U:102:PRO:O	2.48	0.41
1:U:325:MET:HE3	1:W:244:ASP:O	2.18	0.41
1:V:33:SER:C	1:V:34:ILE:HG13	2.46	0.41
1:W:300:SER:HA	1:W:335:ARG:CD	2.50	0.41
1:A:110:LEU:CD1	1:A:177:ARG:HH11	2.32	0.41
1:A:121:GLN:HA	1:A:362:TYR:CZ	2.56	0.41
1:A:236:LEU:HD12	1:A:237:GLU:CA	2.51	0.41
1:B:103:THR:O	1:B:132:MET:HA	2.20	0.41
1:B:175:ILE:H	1:B:175:ILE:HG13	1.71	0.41
1:B:349:LEU:HD23	1:B:349:LEU:HA	1.80	0.41
1:C:164:PRO:HG2	1:C:174:ALA:CB	2.51	0.41
1:D:164:PRO:CG	1:D:174:ALA:HB3	2.51	0.41
1:D:180:LEU:CD1	1:D:181:ALA:N	2.78	0.41
1:D:357:ILE:CG2	1:D:358:THR:N	2.83	0.41
1:E:38:PRO:CG	1:E:49:GLN:NE2	2.79	0.41
1:E:65:LEU:C	1:E:65:LEU:CD1	2.90	0.41
1:E:140:LEU:HB3	1:E:343:GLY:N	2.35	0.41
1:E:148:THR:CG2	1:E:149:THR:N	2.81	0.41
1:E:335:ARG:O	1:E:338:SER:CB	2.69	0.41
1:F:8:LEU:N	1:F:102:PRO:O	2.48	0.41
1:F:173:HIS:HE1	1:G:268:GLY:CA	2.28	0.41
1:F:180:LEU:CD1	1:F:181:ALA:N	2.78	0.41
1:G:8:LEU:CD1	1:G:90:PHE:CD1	3.04	0.41
1:G:57:GLU:O	1:G:58:ALA:C	2.64	0.41
1:H:57:GLU:O	1:H:58:ALA:C	2.64	0.41
1:H:103:THR:O	1:H:132:MET:HA	2.20	0.41
1:H:140:LEU:HB3	1:H:343:GLY:N	2.35	0.41
1:H:167:GLU:OE1	1:J:61:LYS:HE2	2.20	0.41
1:H:236:LEU:HD12	1:H:237:GLU:CA	2.51	0.41
1:H:287:ILE:HD12	1:J:208:ILE:HD13	1.96	0.41
1:I:34:ILE:CG2	1:I:68:LYS:N	2.84	0.41
1:I:135:ALA:CB	1:I:140:LEU:HD21	2.49	0.41
1:I:189:LEU:HA	1:I:192:ILE:HG12	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:33:SER:C	1:J:34:ILE:HG13	2.46	0.41
1:J:164:PRO:HG2	1:J:174:ALA:CB	2.51	0.41
1:J:164:PRO:CG	1:J:174:ALA:HB3	2.51	0.41
1:K:8:LEU:CD1	1:K:90:PHE:CD1	3.04	0.41
1:K:34:ILE:CG2	1:K:68:LYS:N	2.84	0.41
1:K:103:THR:O	1:K:132:MET:HA	2.20	0.41
1:K:135:ALA:CB	1:K:140:LEU:HD21	2.50	0.41
1:K:299:MET:HE2	1:K:299:MET:HB3	1.95	0.41
1:L:33:SER:C	1:L:34:ILE:HG13	2.46	0.41
1:L:110:LEU:CD1	1:L:177:ARG:HH11	2.32	0.41
1:L:121:GLN:HA	1:L:362:TYR:CZ	2.55	0.41
1:L:164:PRO:HG2	1:L:174:ALA:CB	2.51	0.41
1:L:164:PRO:CG	1:L:174:ALA:HB3	2.51	0.41
1:L:335:ARG:O	1:L:338:SER:CB	2.69	0.41
1:M:34:ILE:CG2	1:M:68:LYS:N	2.84	0.41
1:M:34:ILE:HG23	1:M:67:LEU:HB3	2.02	0.41
1:M:103:THR:O	1:M:132:MET:HA	2.20	0.41
1:M:176:MET:HE3	1:M:176:MET:HB3	1.81	0.41
1:M:357:ILE:CG2	1:M:358:THR:N	2.83	0.41
1:N:164:PRO:CG	1:N:174:ALA:HB3	2.51	0.41
1:N:335:ARG:O	1:N:338:SER:CB	2.69	0.41
1:O:2:GLU:HB3	1:O:3:ASP:H	1.66	0.41
1:O:57:GLU:O	1:O:58:ALA:C	2.64	0.41
1:O:133:TYR:CE2	1:O:375:PHE:HB2	2.51	0.41
1:O:142:LEU:CD1	1:O:165:ILE:HD11	2.47	0.41
1:P:38:PRO:CG	1:P:49:GLN:NE2	2.79	0.41
1:P:110:LEU:CD1	1:P:177:ARG:HH11	2.32	0.41
1:P:362:TYR:CE1	1:P:367:PRO:CB	2.99	0.41
1:R:8:LEU:CD1	1:R:90:PHE:CD1	3.04	0.41
1:R:57:GLU:O	1:R:58:ALA:C	2.64	0.41
1:R:103:THR:O	1:R:132:MET:HA	2.20	0.41
1:R:121:GLN:HA	1:R:362:TYR:CZ	2.56	0.41
1:R:121:GLN:HG3	1:R:362:TYR:OH	2.20	0.41
1:S:173:HIS:HE1	1:T:268:GLY:CA	2.28	0.41
1:T:8:LEU:CD1	1:T:90:PHE:CD1	3.04	0.41
1:T:34:ILE:HG23	1:T:67:LEU:HB3	2.01	0.41
1:T:149:THR:CG2	1:T:150:GLY:N	2.82	0.41
1:T:236:LEU:CD1	1:T:237:GLU:CG	2.93	0.41
1:U:44:MET:CG	1:U:45:VAL:H	2.18	0.41
1:V:189:LEU:HA	1:V:192:ILE:HG12	2.03	0.41
1:V:335:ARG:O	1:V:338:SER:CB	2.69	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:34:ILE:CG2	1:W:67:LEU:CB	2.96	0.41
1:W:57:GLU:O	1:W:58:ALA:C	2.64	0.41
1:W:223:PHE:CZ	1:W:266:PHE:CZ	3.09	0.41
1:A:69:TYR:CD2	1:A:69:TYR:N	2.86	0.41
1:B:8:LEU:CD1	1:B:90:PHE:CD1	3.04	0.41
1:C:57:GLU:O	1:C:58:ALA:C	2.64	0.41
1:C:335:ARG:O	1:C:338:SER:CB	2.69	0.41
1:D:173:HIS:HE1	1:E:268:GLY:CA	2.28	0.41
1:E:164:PRO:HG2	1:E:174:ALA:CB	2.51	0.41
1:E:224:GLU:OE1	1:E:224:GLU:N	2.51	0.41
1:F:164:PRO:CG	1:F:174:ALA:HB3	2.51	0.41
1:F:169:TYR:CE2	1:H:40:HIS:HB2	2.56	0.41
1:F:176:MET:HE3	1:F:176:MET:HB3	1.81	0.41
1:G:34:ILE:CG2	1:G:68:LYS:N	2.84	0.41
1:H:164:PRO:CG	1:H:174:ALA:HB3	2.51	0.41
1:H:169:TYR:CE2	1:J:40:HIS:HB2	2.56	0.41
1:I:73:HIC:HA	1:I:183:ARG:NH1	2.18	0.41
1:I:167:GLU:OE1	1:K:61:LYS:HE2	2.20	0.41
1:J:171:LEU:HA	1:J:172:PRO:HD3	1.79	0.41
1:J:189:LEU:HA	1:J:192:ILE:HG12	2.02	0.41
1:L:38:PRO:CG	1:L:49:GLN:NE2	2.79	0.41
1:L:57:GLU:O	1:L:58:ALA:C	2.64	0.41
1:L:140:LEU:HB3	1:L:343:GLY:N	2.35	0.41
1:L:183:ARG:HE	1:L:183:ARG:HB3	1.58	0.41
1:L:332:PRO:HA	1:L:333:PRO:HD3	1.95	0.41
1:N:73:HIC:HA	1:N:183:ARG:NH1	2.18	0.41
1:O:135:ALA:CB	1:O:140:LEU:HD21	2.49	0.41
1:Q:8:LEU:CD1	1:Q:90:PHE:CD1	3.04	0.41
1:Q:33:SER:C	1:Q:34:ILE:HG13	2.46	0.41
1:Q:164:PRO:HG2	1:Q:174:ALA:CB	2.51	0.41
1:R:224:GLU:OE1	1:R:224:GLU:N	2.51	0.41
1:S:164:PRO:HG2	1:S:174:ALA:CB	2.51	0.41
1:T:34:ILE:CG2	1:T:67:LEU:HD22	2.28	0.41
1:T:57:GLU:O	1:T:58:ALA:C	2.64	0.41
1:T:121:GLN:HA	1:T:362:TYR:CZ	2.55	0.41
1:U:236:LEU:HD12	1:U:237:GLU:CA	2.51	0.41
1:V:34:ILE:HD13	1:V:67:LEU:CD2	2.46	0.41
1:V:357:ILE:HD12	1:V:357:ILE:HA	1.78	0.41
1:V:357:ILE:CG2	1:V:358:THR:N	2.83	0.41
1:W:189:LEU:HA	1:W:192:ILE:HG12	2.02	0.41
1:W:332:PRO:HA	1:W:333:PRO:HD3	1.95	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:369:ILE:C	1:A:371:HIS:N	2.77	0.40
1:B:143:TYR:CE2	1:D:45:VAL:CG2	3.04	0.40
1:C:143:TYR:CE2	1:E:45:VAL:CG2	3.04	0.40
1:D:43:VAL:O	1:D:44:MET:CE	2.69	0.40
1:E:142:LEU:CD1	1:E:165:ILE:HD11	2.47	0.40
1:F:34:ILE:CG2	1:F:67:LEU:CB	2.96	0.40
1:H:164:PRO:HG2	1:H:174:ALA:CB	2.51	0.40
1:J:169:TYR:CE2	1:L:40:HIS:HB2	2.56	0.40
1:J:236:LEU:HD12	1:J:237:GLU:CA	2.51	0.40
1:J:332:PRO:HA	1:J:333:PRO:HD3	1.95	0.40
1:M:8:LEU:N	1:M:102:PRO:O	2.48	0.40
1:N:8:LEU:N	1:N:102:PRO:O	2.48	0.40
1:N:183:ARG:HE	1:N:183:ARG:HB3	1.58	0.40
1:N:362:TYR:CE1	1:N:367:PRO:CB	2.99	0.40
1:O:34:ILE:CG2	1:O:68:LYS:N	2.84	0.40
1:O:335:ARG:O	1:O:338:SER:CB	2.69	0.40
1:Q:143:TYR:CE2	1:S:45:VAL:CG2	3.04	0.40
1:S:8:LEU:N	1:S:102:PRO:O	2.48	0.40
1:S:110:LEU:CD1	1:S:177:ARG:NH1	2.80	0.40
1:S:164:PRO:CG	1:S:174:ALA:HB3	2.51	0.40
1:T:110:LEU:CD1	1:T:177:ARG:HH11	2.32	0.40
1:U:47:MET:HB3	1:U:48:GLY:H	1.62	0.40
1:U:57:GLU:O	1:U:58:ALA:C	2.64	0.40
1:V:164:PRO:HG2	1:V:174:ALA:CB	2.51	0.40
1:V:164:PRO:CG	1:V:174:ALA:HB3	2.51	0.40
1:A:47:MET:HB3	1:A:48:GLY:H	1.61	0.40
1:A:143:TYR:CE2	1:C:45:VAL:CG2	3.04	0.40
1:A:164:PRO:HG2	1:A:174:ALA:CB	2.51	0.40
1:B:50:LYS:CG	1:B:53:TYR:CE2	3.03	0.40
1:B:113:LYS:HB3	1:B:371:HIS:CD2	2.57	0.40
1:H:47:MET:HB3	1:H:48:GLY:H	1.62	0.40
1:H:189:LEU:HA	1:H:192:ILE:HG12	2.02	0.40
1:K:57:GLU:O	1:K:58:ALA:C	2.64	0.40
1:K:236:LEU:HD12	1:K:237:GLU:CA	2.51	0.40
1:K:300:SER:HA	1:K:335:ARG:CD	2.50	0.40
1:L:135:ALA:CB	1:L:140:LEU:HD21	2.50	0.40
1:M:140:LEU:HB3	1:M:343:GLY:N	2.35	0.40
1:N:47:MET:HB3	1:N:48:GLY:H	1.62	0.40
1:O:164:PRO:HG2	1:O:174:ALA:CB	2.51	0.40
1:Q:44:MET:CG	1:Q:45:VAL:N	2.76	0.40
1:T:34:ILE:CG2	1:T:68:LYS:N	2.84	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:106:THR:HG22	1:T:140:LEU:HD11	2.01	0.40
1:T:164:PRO:CG	1:T:174:ALA:HB3	2.51	0.40
1:T:325:MET:HE3	1:V:244:ASP:O	2.18	0.40
1:T:357:ILE:CG2	1:T:358:THR:N	2.82	0.40
1:U:164:PRO:CG	1:U:174:ALA:HB3	2.51	0.40
1:U:189:LEU:HA	1:U:192:ILE:HG12	2.02	0.40
1:A:8:LEU:CD1	1:A:90:PHE:CD1	3.04	0.40
1:B:57:GLU:O	1:B:58:ALA:C	2.64	0.40
1:B:189:LEU:HA	1:B:192:ILE:HG12	2.02	0.40
1:C:50:LYS:CG	1:C:53:TYR:CE2	3.03	0.40
1:D:34:ILE:CG2	1:D:67:LEU:CB	2.96	0.40
1:D:103:THR:O	1:D:132:MET:HA	2.20	0.40
1:D:143:TYR:CE2	1:F:45:VAL:CG2	3.04	0.40
1:D:189:LEU:HA	1:D:192:ILE:HG12	2.02	0.40
1:E:299:MET:HE2	1:E:299:MET:HB3	1.95	0.40
1:F:173:HIS:CE1	1:G:268:GLY:N	2.90	0.40
1:G:140:LEU:HB3	1:G:343:GLY:N	2.35	0.40
1:G:164:PRO:HG2	1:G:174:ALA:CB	2.51	0.40
1:J:140:LEU:HB3	1:J:343:GLY:N	2.36	0.40
1:K:369:ILE:C	1:K:371:HIS:N	2.77	0.40
1:N:57:GLU:O	1:N:58:ALA:C	2.64	0.40
1:N:164:PRO:HG2	1:N:174:ALA:CB	2.51	0.40
1:N:173:HIS:CE1	1:O:268:GLY:N	2.90	0.40
1:N:357:ILE:CG2	1:N:358:THR:N	2.83	0.40
1:O:8:LEU:CD1	1:O:90:PHE:CD1	3.04	0.40
1:O:121:GLN:HA	1:O:362:TYR:CZ	2.55	0.40
1:O:143:TYR:CE2	1:Q:45:VAL:CG2	3.04	0.40
1:O:149:THR:CG2	1:O:150:GLY:N	2.82	0.40
1:O:236:LEU:HD12	1:O:237:GLU:CA	2.51	0.40
1:P:33:SER:C	1:P:34:ILE:HG13	2.46	0.40
1:Q:113:LYS:HB3	1:Q:371:HIS:CD2	2.57	0.40
1:Q:164:PRO:CG	1:Q:174:ALA:HB3	2.51	0.40
1:R:34:ILE:CG2	1:R:68:LYS:N	2.84	0.40
1:R:50:LYS:CG	1:R:53:TYR:CE2	3.03	0.40
1:S:183:ARG:HE	1:S:183:ARG:HB3	1.58	0.40
1:T:164:PRO:HG2	1:T:174:ALA:CB	2.51	0.40
1:U:121:GLN:HA	1:U:362:TYR:CZ	2.55	0.40
1:U:223:PHE:CZ	1:U:266:PHE:CZ	3.09	0.40
1:W:357:ILE:HD12	1:W:357:ILE:HA	1.78	0.40
1:A:57:GLU:O	1:A:58:ALA:C	2.64	0.40
1:A:335:ARG:O	1:A:338:SER:CB	2.69	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:34:ILE:CG2	1:E:68:LYS:N	2.84	0.40
1:E:143:TYR:CE2	1:G:45:VAL:CG2	3.04	0.40
1:E:173:HIS:CE1	1:F:268:GLY:N	2.90	0.40
1:E:357:ILE:CG2	1:E:358:THR:N	2.82	0.40
1:F:369:ILE:C	1:F:371:HIS:N	2.77	0.40
1:I:173:HIS:CE1	1:J:268:GLY:N	2.90	0.40
1:I:236:LEU:CD1	1:I:237:GLU:CG	2.93	0.40
1:K:173:HIS:CE1	1:L:268:GLY:N	2.90	0.40
1:M:8:LEU:CD1	1:M:90:PHE:CD1	3.04	0.40
1:M:157:ASP:H	2:M:401:ADP:PB	2.45	0.40
1:O:224:GLU:OE1	1:O:224:GLU:N	2.51	0.40
1:P:8:LEU:CD1	1:P:90:PHE:CD1	3.04	0.40
1:P:173:HIS:CE1	1:Q:268:GLY:N	2.90	0.40
1:Q:34:ILE:CG2	1:Q:68:LYS:N	2.84	0.40
1:Q:135:ALA:CB	1:Q:140:LEU:HD21	2.49	0.40
1:S:143:TYR:CE2	1:U:45:VAL:CG2	3.04	0.40
1:T:287:ILE:HD12	1:V:208:ILE:HD13	1.96	0.40
1:U:164:PRO:HG2	1:U:174:ALA:CB	2.51	0.40
1:U:374:CYS:HB2	1:U:375:PHE:H	1.61	0.40
1:V:43:VAL:O	1:V:44:MET:CE	2.69	0.40
1:V:113:LYS:HB3	1:V:371:HIS:CD2	2.57	0.40
1:V:236:LEU:HD12	1:V:237:GLU:CA	2.51	0.40
1:W:183:ARG:HE	1:W:183:ARG:HB3	1.58	0.40
1:A:120:THR:HG23	1:A:132:MET:SD	2.62	0.40
1:C:8:LEU:CD1	1:C:90:PHE:CD1	3.04	0.40
1:C:236:LEU:CD1	1:C:237:GLU:CG	2.93	0.40
1:D:169:TYR:CE2	1:F:40:HIS:HB2	2.56	0.40
1:D:173:HIS:CE1	1:E:268:GLY:N	2.90	0.40
1:E:8:LEU:CD1	1:E:90:PHE:CD1	3.04	0.40
1:E:33:SER:C	1:E:34:ILE:HG13	2.46	0.40
1:E:113:LYS:HB3	1:E:371:HIS:CD2	2.57	0.40
1:F:8:LEU:CD1	1:F:90:PHE:CD1	3.04	0.40
1:F:157:ASP:H	2:F:401:ADP:PB	2.45	0.40
1:G:44:MET:CG	1:G:45:VAL:N	2.75	0.40
1:G:335:ARG:O	1:G:338:SER:CB	2.69	0.40
1:I:110:LEU:CD1	1:I:177:ARG:HH11	2.32	0.40
1:I:113:LYS:HB3	1:I:371:HIS:CD2	2.57	0.40
1:K:39:ARG:NE	1:K:66:THR:CA	2.69	0.40
1:L:169:TYR:CE2	1:N:40:HIS:HB2	2.56	0.40
1:M:113:LYS:HB3	1:M:371:HIS:CD2	2.57	0.40
1:M:349:LEU:HD23	1:M:349:LEU:HA	1.80	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:113:LYS:HB3	1:O:371:HIS:CD2	2.57	0.40
1:O:157:ASP:H	2:O:401:ADP:PB	2.45	0.40
1:P:164:PRO:CG	1:P:174:ALA:HB3	2.51	0.40
1:Q:173:HIS:CE1	1:R:268:GLY:N	2.90	0.40
1:R:32:PRO:HD2	1:R:55:GLY:C	2.47	0.40
1:R:335:ARG:O	1:R:338:SER:CB	2.69	0.40
1:S:8:LEU:CD1	1:S:90:PHE:CD1	3.04	0.40
1:S:57:GLU:O	1:S:58:ALA:C	2.64	0.40
1:S:65:LEU:C	1:S:65:LEU:CD1	2.90	0.40
1:S:223:PHE:CZ	1:S:266:PHE:CZ	3.09	0.40
1:T:32:PRO:HD2	1:T:55:GLY:C	2.47	0.40
1:T:113:LYS:HB3	1:T:371:HIS:CD2	2.57	0.40
1:V:8:LEU:CD1	1:V:90:PHE:CD1	3.04	0.40
1:V:32:PRO:HD2	1:V:55:GLY:C	2.47	0.40
1:V:106:THR:HG22	1:V:140:LEU:HD11	2.01	0.40
1:V:121:GLN:HA	1:V:362:TYR:CZ	2.55	0.40
1:W:132:MET:HE2	1:W:132:MET:HB2	1.75	0.40
1:W:299:MET:HE2	1:W:299:MET:HB3	1.95	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	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	B	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	C	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	D	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	E	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	F	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	G	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	H	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	I	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	J	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	K	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	L	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	M	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	N	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	O	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	P	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	Q	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	R	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	S	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	T	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	U	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	V	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
1	W	372/375 (99%)	329 (88%)	31 (8%)	12 (3%)	3	21
All	All	8556/8625 (99%)	7567 (88%)	713 (8%)	276 (3%)	5	21

All (276) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	3	ASP
1	A	41	GLN
1	A	44	MET
1	B	3	ASP
1	B	41	GLN
1	B	44	MET
1	C	3	ASP
1	C	41	GLN
1	C	44	MET
1	D	3	ASP
1	D	41	GLN
1	D	44	MET
1	E	3	ASP
1	E	41	GLN

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Mol	Chain	Res	Type
1	E	44	MET
1	F	3	ASP
1	F	41	GLN
1	F	44	MET
1	G	3	ASP
1	G	41	GLN
1	G	44	MET
1	H	3	ASP
1	H	41	GLN
1	H	44	MET
1	I	3	ASP
1	I	41	GLN
1	I	44	MET
1	J	3	ASP
1	J	41	GLN
1	J	44	MET
1	K	3	ASP
1	K	41	GLN
1	K	44	MET
1	L	3	ASP
1	L	41	GLN
1	L	44	MET
1	M	3	ASP
1	M	41	GLN
1	M	44	MET
1	N	3	ASP
1	N	41	GLN
1	N	44	MET
1	O	3	ASP
1	O	41	GLN
1	O	44	MET
1	P	3	ASP
1	P	41	GLN
1	P	44	MET
1	Q	3	ASP
1	Q	41	GLN
1	Q	44	MET
1	R	3	ASP
1	R	41	GLN
1	R	44	MET
1	S	3	ASP
1	S	41	GLN

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Mol	Chain	Res	Type
1	S	44	MET
1	T	3	ASP
1	T	41	GLN
1	T	44	MET
1	U	3	ASP
1	U	41	GLN
1	U	44	MET
1	V	3	ASP
1	V	41	GLN
1	V	44	MET
1	W	3	ASP
1	W	41	GLN
1	W	44	MET
1	A	4	GLU
1	A	55	GLY
1	A	374	CYS
1	B	4	GLU
1	B	55	GLY
1	B	374	CYS
1	C	4	GLU
1	C	55	GLY
1	C	374	CYS
1	D	4	GLU
1	D	55	GLY
1	D	374	CYS
1	E	4	GLU
1	E	55	GLY
1	E	374	CYS
1	F	4	GLU
1	F	55	GLY
1	F	374	CYS
1	G	4	GLU
1	G	55	GLY
1	G	374	CYS
1	H	4	GLU
1	H	55	GLY
1	H	374	CYS
1	I	4	GLU
1	I	55	GLY
1	I	374	CYS
1	J	4	GLU
1	J	55	GLY

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Mol	Chain	Res	Type
1	J	374	CYS
1	K	4	GLU
1	K	55	GLY
1	K	374	CYS
1	L	4	GLU
1	L	55	GLY
1	L	374	CYS
1	M	4	GLU
1	M	55	GLY
1	M	374	CYS
1	N	4	GLU
1	N	55	GLY
1	N	374	CYS
1	O	4	GLU
1	O	55	GLY
1	O	374	CYS
1	P	4	GLU
1	P	55	GLY
1	P	374	CYS
1	Q	4	GLU
1	Q	55	GLY
1	Q	374	CYS
1	R	4	GLU
1	R	55	GLY
1	R	374	CYS
1	S	4	GLU
1	S	55	GLY
1	S	374	CYS
1	T	4	GLU
1	T	55	GLY
1	T	374	CYS
1	U	4	GLU
1	U	55	GLY
1	U	374	CYS
1	V	4	GLU
1	V	55	GLY
1	V	374	CYS
1	W	4	GLU
1	W	55	GLY
1	W	374	CYS
1	A	2	GLU
1	B	2	GLU

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Mol	Chain	Res	Type
1	C	2	GLU
1	D	2	GLU
1	E	2	GLU
1	F	2	GLU
1	G	2	GLU
1	H	2	GLU
1	I	2	GLU
1	J	2	GLU
1	K	2	GLU
1	L	2	GLU
1	M	2	GLU
1	N	2	GLU
1	O	2	GLU
1	P	2	GLU
1	Q	2	GLU
1	R	2	GLU
1	S	2	GLU
1	T	2	GLU
1	U	2	GLU
1	V	2	GLU
1	W	2	GLU
1	A	43	VAL
1	A	54	VAL
1	B	43	VAL
1	B	54	VAL
1	C	43	VAL
1	C	54	VAL
1	D	43	VAL
1	D	54	VAL
1	E	43	VAL
1	E	54	VAL
1	F	43	VAL
1	F	54	VAL
1	G	43	VAL
1	G	54	VAL
1	H	43	VAL
1	H	54	VAL
1	I	43	VAL
1	I	54	VAL
1	J	43	VAL
1	J	54	VAL
1	K	43	VAL

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Mol	Chain	Res	Type
1	K	54	VAL
1	L	43	VAL
1	L	54	VAL
1	M	43	VAL
1	M	54	VAL
1	N	43	VAL
1	N	54	VAL
1	O	43	VAL
1	O	54	VAL
1	P	43	VAL
1	P	54	VAL
1	Q	43	VAL
1	Q	54	VAL
1	R	43	VAL
1	R	54	VAL
1	S	43	VAL
1	S	54	VAL
1	T	43	VAL
1	T	54	VAL
1	U	43	VAL
1	U	54	VAL
1	V	43	VAL
1	V	54	VAL
1	W	43	VAL
1	W	54	VAL
1	A	58	ALA
1	C	58	ALA
1	D	58	ALA
1	F	58	ALA
1	H	58	ALA
1	I	58	ALA
1	J	58	ALA
1	L	58	ALA
1	N	58	ALA
1	O	58	ALA
1	P	58	ALA
1	Q	58	ALA
1	R	58	ALA
1	T	58	ALA
1	U	58	ALA
1	W	58	ALA
1	B	58	ALA

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Mol	Chain	Res	Type
1	E	58	ALA
1	G	58	ALA
1	K	58	ALA
1	M	58	ALA
1	S	58	ALA
1	V	58	ALA
1	A	45	VAL
1	B	45	VAL
1	C	45	VAL
1	F	45	VAL
1	M	45	VAL
1	P	45	VAL
1	V	45	VAL
1	W	45	VAL
1	D	45	VAL
1	E	45	VAL
1	G	45	VAL
1	H	45	VAL
1	I	45	VAL
1	J	45	VAL
1	K	45	VAL
1	L	45	VAL
1	N	45	VAL
1	O	45	VAL
1	Q	45	VAL
1	R	45	VAL
1	S	45	VAL
1	T	45	VAL
1	U	45	VAL
1	A	370	VAL
1	B	370	VAL
1	C	370	VAL
1	D	370	VAL
1	E	370	VAL
1	F	370	VAL
1	G	370	VAL
1	H	370	VAL
1	I	370	VAL
1	J	370	VAL
1	K	370	VAL
1	L	370	VAL
1	M	370	VAL

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Mol	Chain	Res	Type
1	N	370	VAL
1	O	370	VAL
1	P	370	VAL
1	Q	370	VAL
1	R	370	VAL
1	S	370	VAL
1	T	370	VAL
1	U	370	VAL
1	V	370	VAL
1	W	370	VAL

5.3.2 Protein sidechains (i)

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	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	B	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	C	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	D	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	E	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	F	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	G	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	H	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	I	317/317 (100%)	278 (88%)	39 (12%)	4	17
1	J	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	K	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	L	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	M	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	N	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	O	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	P	317/317 (100%)	277 (87%)	40 (13%)	4	16

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	Q	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	R	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	S	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	T	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	U	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	V	317/317 (100%)	277 (87%)	40 (13%)	4	16
1	W	317/317 (100%)	277 (87%)	40 (13%)	4	16
All	All	7291/7291 (100%)	6372 (87%)	919 (13%)	6	16

All (919) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	8	LEU
1	A	33	SER
1	A	35	VAL
1	A	37	ARG
1	A	45	VAL
1	A	49	GLN
1	A	51	ASP
1	A	56	ASP
1	A	57	GLU
1	A	66	THR
1	A	68	LYS
1	A	71	ILE
1	A	113	LYS
1	A	132	MET
1	A	139	VAL
1	A	141	SER
1	A	142	LEU
1	A	145	SER
1	A	147	ARG
1	A	148	THR
1	A	149	THR
1	A	167	GLU
1	A	171	LEU
1	A	176	MET
1	A	179	ASP
1	A	180	LEU
1	A	183	ARG
1	A	208	ILE

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Mol	Chain	Res	Type
1	A	227	MET
1	A	232	SER
1	A	236	LEU
1	A	244	ASP
1	A	269	MET
1	A	286	ASP
1	A	335	ARG
1	A	350	SER
1	A	357	ILE
1	A	370	VAL
1	A	372	ARG
1	A	373	LYS
1	B	8	LEU
1	B	33	SER
1	B	35	VAL
1	B	37	ARG
1	B	45	VAL
1	B	49	GLN
1	B	51	ASP
1	B	56	ASP
1	B	57	GLU
1	B	66	THR
1	B	68	LYS
1	B	71	ILE
1	B	113	LYS
1	B	132	MET
1	B	139	VAL
1	B	141	SER
1	B	142	LEU
1	B	145	SER
1	B	147	ARG
1	B	148	THR
1	B	149	THR
1	B	167	GLU
1	B	171	LEU
1	B	176	MET
1	B	179	ASP
1	B	180	LEU
1	B	183	ARG
1	B	208	ILE
1	B	227	MET
1	B	232	SER

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Mol	Chain	Res	Type
1	B	236	LEU
1	B	244	ASP
1	B	269	MET
1	B	286	ASP
1	B	335	ARG
1	B	350	SER
1	B	357	ILE
1	B	370	VAL
1	B	372	ARG
1	B	373	LYS
1	C	8	LEU
1	C	33	SER
1	C	35	VAL
1	C	37	ARG
1	C	45	VAL
1	C	49	GLN
1	C	51	ASP
1	C	56	ASP
1	C	57	GLU
1	C	66	THR
1	C	68	LYS
1	C	71	ILE
1	C	113	LYS
1	C	132	MET
1	C	139	VAL
1	C	141	SER
1	C	142	LEU
1	C	145	SER
1	C	147	ARG
1	C	148	THR
1	C	149	THR
1	C	167	GLU
1	C	171	LEU
1	C	176	MET
1	C	179	ASP
1	C	180	LEU
1	C	183	ARG
1	C	208	ILE
1	C	227	MET
1	C	232	SER
1	C	236	LEU
1	C	244	ASP

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Mol	Chain	Res	Type
1	C	269	MET
1	C	286	ASP
1	C	335	ARG
1	C	350	SER
1	C	357	ILE
1	C	370	VAL
1	C	372	ARG
1	C	373	LYS
1	D	8	LEU
1	D	33	SER
1	D	35	VAL
1	D	37	ARG
1	D	45	VAL
1	D	49	GLN
1	D	51	ASP
1	D	56	ASP
1	D	57	GLU
1	D	66	THR
1	D	68	LYS
1	D	71	ILE
1	D	113	LYS
1	D	132	MET
1	D	139	VAL
1	D	141	SER
1	D	142	LEU
1	D	145	SER
1	D	147	ARG
1	D	148	THR
1	D	149	THR
1	D	167	GLU
1	D	171	LEU
1	D	176	MET
1	D	179	ASP
1	D	180	LEU
1	D	183	ARG
1	D	208	ILE
1	D	227	MET
1	D	232	SER
1	D	236	LEU
1	D	244	ASP
1	D	269	MET
1	D	286	ASP

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Mol	Chain	Res	Type
1	D	335	ARG
1	D	350	SER
1	D	357	ILE
1	D	370	VAL
1	D	372	ARG
1	D	373	LYS
1	E	8	LEU
1	E	33	SER
1	E	35	VAL
1	E	37	ARG
1	E	45	VAL
1	E	49	GLN
1	E	51	ASP
1	E	56	ASP
1	E	57	GLU
1	E	66	THR
1	E	68	LYS
1	E	71	ILE
1	E	113	LYS
1	E	132	MET
1	E	139	VAL
1	E	141	SER
1	E	142	LEU
1	E	145	SER
1	E	147	ARG
1	E	148	THR
1	E	149	THR
1	E	167	GLU
1	E	171	LEU
1	E	176	MET
1	E	179	ASP
1	E	180	LEU
1	E	183	ARG
1	E	208	ILE
1	E	227	MET
1	E	232	SER
1	E	236	LEU
1	E	244	ASP
1	E	269	MET
1	E	286	ASP
1	E	335	ARG
1	E	350	SER

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Mol	Chain	Res	Type
1	E	357	ILE
1	E	370	VAL
1	E	372	ARG
1	E	373	LYS
1	F	8	LEU
1	F	33	SER
1	F	35	VAL
1	F	37	ARG
1	F	45	VAL
1	F	49	GLN
1	F	51	ASP
1	F	56	ASP
1	F	57	GLU
1	F	66	THR
1	F	68	LYS
1	F	71	ILE
1	F	113	LYS
1	F	132	MET
1	F	139	VAL
1	F	141	SER
1	F	142	LEU
1	F	145	SER
1	F	147	ARG
1	F	148	THR
1	F	149	THR
1	F	167	GLU
1	F	171	LEU
1	F	176	MET
1	F	179	ASP
1	F	180	LEU
1	F	183	ARG
1	F	208	ILE
1	F	227	MET
1	F	232	SER
1	F	236	LEU
1	F	244	ASP
1	F	269	MET
1	F	286	ASP
1	F	335	ARG
1	F	350	SER
1	F	357	ILE
1	F	370	VAL

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Mol	Chain	Res	Type
1	F	372	ARG
1	F	373	LYS
1	G	8	LEU
1	G	33	SER
1	G	35	VAL
1	G	37	ARG
1	G	45	VAL
1	G	49	GLN
1	G	51	ASP
1	G	56	ASP
1	G	57	GLU
1	G	66	THR
1	G	68	LYS
1	G	71	ILE
1	G	113	LYS
1	G	132	MET
1	G	139	VAL
1	G	141	SER
1	G	142	LEU
1	G	145	SER
1	G	147	ARG
1	G	148	THR
1	G	149	THR
1	G	167	GLU
1	G	171	LEU
1	G	176	MET
1	G	179	ASP
1	G	180	LEU
1	G	183	ARG
1	G	208	ILE
1	G	227	MET
1	G	232	SER
1	G	236	LEU
1	G	244	ASP
1	G	269	MET
1	G	286	ASP
1	G	335	ARG
1	G	350	SER
1	G	357	ILE
1	G	370	VAL
1	G	372	ARG
1	G	373	LYS

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Mol	Chain	Res	Type
1	H	8	LEU
1	H	33	SER
1	H	35	VAL
1	H	37	ARG
1	H	45	VAL
1	H	49	GLN
1	H	51	ASP
1	H	56	ASP
1	H	57	GLU
1	H	66	THR
1	H	68	LYS
1	H	71	ILE
1	H	113	LYS
1	H	132	MET
1	H	139	VAL
1	H	141	SER
1	H	142	LEU
1	H	145	SER
1	H	147	ARG
1	H	148	THR
1	H	149	THR
1	H	167	GLU
1	H	171	LEU
1	H	176	MET
1	H	179	ASP
1	H	180	LEU
1	H	183	ARG
1	H	208	ILE
1	H	227	MET
1	H	232	SER
1	H	236	LEU
1	H	244	ASP
1	H	269	MET
1	H	286	ASP
1	H	335	ARG
1	H	350	SER
1	H	357	ILE
1	H	370	VAL
1	H	372	ARG
1	H	373	LYS
1	I	8	LEU
1	I	33	SER

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Mol	Chain	Res	Type
1	I	35	VAL
1	I	37	ARG
1	I	45	VAL
1	I	49	GLN
1	I	51	ASP
1	I	56	ASP
1	I	57	GLU
1	I	66	THR
1	I	68	LYS
1	I	71	ILE
1	I	113	LYS
1	I	132	MET
1	I	139	VAL
1	I	141	SER
1	I	142	LEU
1	I	145	SER
1	I	147	ARG
1	I	148	THR
1	I	149	THR
1	I	167	GLU
1	I	171	LEU
1	I	176	MET
1	I	180	LEU
1	I	183	ARG
1	I	208	ILE
1	I	227	MET
1	I	232	SER
1	I	236	LEU
1	I	244	ASP
1	I	269	MET
1	I	286	ASP
1	I	335	ARG
1	I	350	SER
1	I	357	ILE
1	I	370	VAL
1	I	372	ARG
1	I	373	LYS
1	J	8	LEU
1	J	33	SER
1	J	35	VAL
1	J	37	ARG
1	J	45	VAL

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Mol	Chain	Res	Type
1	J	49	GLN
1	J	51	ASP
1	J	56	ASP
1	J	57	GLU
1	J	66	THR
1	J	68	LYS
1	J	71	ILE
1	J	113	LYS
1	J	132	MET
1	J	139	VAL
1	J	141	SER
1	J	142	LEU
1	J	145	SER
1	J	147	ARG
1	J	148	THR
1	J	149	THR
1	J	167	GLU
1	J	171	LEU
1	J	176	MET
1	J	179	ASP
1	J	180	LEU
1	J	183	ARG
1	J	208	ILE
1	J	227	MET
1	J	232	SER
1	J	236	LEU
1	J	244	ASP
1	J	269	MET
1	J	286	ASP
1	J	335	ARG
1	J	350	SER
1	J	357	ILE
1	J	370	VAL
1	J	372	ARG
1	J	373	LYS
1	K	8	LEU
1	K	33	SER
1	K	35	VAL
1	K	37	ARG
1	K	45	VAL
1	K	49	GLN
1	K	51	ASP

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Mol	Chain	Res	Type
1	K	56	ASP
1	K	57	GLU
1	K	66	THR
1	K	68	LYS
1	K	71	ILE
1	K	113	LYS
1	K	132	MET
1	K	139	VAL
1	K	141	SER
1	K	142	LEU
1	K	145	SER
1	K	147	ARG
1	K	148	THR
1	K	149	THR
1	K	167	GLU
1	K	171	LEU
1	K	176	MET
1	K	179	ASP
1	K	180	LEU
1	K	183	ARG
1	K	208	ILE
1	K	227	MET
1	K	232	SER
1	K	236	LEU
1	K	244	ASP
1	K	269	MET
1	K	286	ASP
1	K	335	ARG
1	K	350	SER
1	K	357	ILE
1	K	370	VAL
1	K	372	ARG
1	K	373	LYS
1	L	8	LEU
1	L	33	SER
1	L	35	VAL
1	L	37	ARG
1	L	45	VAL
1	L	49	GLN
1	L	51	ASP
1	L	56	ASP
1	L	57	GLU

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Mol	Chain	Res	Type
1	L	66	THR
1	L	68	LYS
1	L	71	ILE
1	L	113	LYS
1	L	132	MET
1	L	139	VAL
1	L	141	SER
1	L	142	LEU
1	L	145	SER
1	L	147	ARG
1	L	148	THR
1	L	149	THR
1	L	167	GLU
1	L	171	LEU
1	L	176	MET
1	L	179	ASP
1	L	180	LEU
1	L	183	ARG
1	L	208	ILE
1	L	227	MET
1	L	232	SER
1	L	236	LEU
1	L	244	ASP
1	L	269	MET
1	L	286	ASP
1	L	335	ARG
1	L	350	SER
1	L	357	ILE
1	L	370	VAL
1	L	372	ARG
1	L	373	LYS
1	M	8	LEU
1	M	33	SER
1	M	35	VAL
1	M	37	ARG
1	M	45	VAL
1	M	49	GLN
1	M	51	ASP
1	M	56	ASP
1	M	57	GLU
1	M	66	THR
1	M	68	LYS

Continued on next page...

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Mol	Chain	Res	Type
1	M	71	ILE
1	M	113	LYS
1	M	132	MET
1	M	139	VAL
1	M	141	SER
1	M	142	LEU
1	M	145	SER
1	M	147	ARG
1	M	148	THR
1	M	149	THR
1	M	167	GLU
1	M	171	LEU
1	M	176	MET
1	M	179	ASP
1	M	180	LEU
1	M	183	ARG
1	M	208	ILE
1	M	227	MET
1	M	232	SER
1	M	236	LEU
1	M	244	ASP
1	M	269	MET
1	M	286	ASP
1	M	335	ARG
1	M	350	SER
1	M	357	ILE
1	M	370	VAL
1	M	372	ARG
1	M	373	LYS
1	N	8	LEU
1	N	33	SER
1	N	35	VAL
1	N	37	ARG
1	N	45	VAL
1	N	49	GLN
1	N	51	ASP
1	N	56	ASP
1	N	57	GLU
1	N	66	THR
1	N	68	LYS
1	N	71	ILE
1	N	113	LYS

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Mol	Chain	Res	Type
1	N	132	MET
1	N	139	VAL
1	N	141	SER
1	N	142	LEU
1	N	145	SER
1	N	147	ARG
1	N	148	THR
1	N	149	THR
1	N	167	GLU
1	N	171	LEU
1	N	176	MET
1	N	179	ASP
1	N	180	LEU
1	N	183	ARG
1	N	208	ILE
1	N	227	MET
1	N	232	SER
1	N	236	LEU
1	N	244	ASP
1	N	269	MET
1	N	286	ASP
1	N	335	ARG
1	N	350	SER
1	N	357	ILE
1	N	370	VAL
1	N	372	ARG
1	N	373	LYS
1	O	8	LEU
1	O	33	SER
1	O	35	VAL
1	O	37	ARG
1	O	45	VAL
1	O	49	GLN
1	O	51	ASP
1	O	56	ASP
1	O	57	GLU
1	O	66	THR
1	O	68	LYS
1	O	71	ILE
1	O	113	LYS
1	O	132	MET
1	O	139	VAL

Continued on next page...

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Mol	Chain	Res	Type
1	O	141	SER
1	O	142	LEU
1	O	145	SER
1	O	147	ARG
1	O	148	THR
1	O	149	THR
1	O	167	GLU
1	O	171	LEU
1	O	176	MET
1	O	179	ASP
1	O	180	LEU
1	O	183	ARG
1	O	208	ILE
1	O	227	MET
1	O	232	SER
1	O	236	LEU
1	O	244	ASP
1	O	269	MET
1	O	286	ASP
1	O	335	ARG
1	O	350	SER
1	O	357	ILE
1	O	370	VAL
1	O	372	ARG
1	O	373	LYS
1	P	8	LEU
1	P	33	SER
1	P	35	VAL
1	P	37	ARG
1	P	45	VAL
1	P	49	GLN
1	P	51	ASP
1	P	56	ASP
1	P	57	GLU
1	P	66	THR
1	P	68	LYS
1	P	71	ILE
1	P	113	LYS
1	P	132	MET
1	P	139	VAL
1	P	141	SER
1	P	142	LEU

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Mol	Chain	Res	Type
1	P	145	SER
1	P	147	ARG
1	P	148	THR
1	P	149	THR
1	P	167	GLU
1	P	171	LEU
1	P	176	MET
1	P	179	ASP
1	P	180	LEU
1	P	183	ARG
1	P	208	ILE
1	P	227	MET
1	P	232	SER
1	P	236	LEU
1	P	244	ASP
1	P	269	MET
1	P	286	ASP
1	P	335	ARG
1	P	350	SER
1	P	357	ILE
1	P	370	VAL
1	P	372	ARG
1	P	373	LYS
1	Q	8	LEU
1	Q	33	SER
1	Q	35	VAL
1	Q	37	ARG
1	Q	45	VAL
1	Q	49	GLN
1	Q	51	ASP
1	Q	56	ASP
1	Q	57	GLU
1	Q	66	THR
1	Q	68	LYS
1	Q	71	ILE
1	Q	113	LYS
1	Q	132	MET
1	Q	139	VAL
1	Q	141	SER
1	Q	142	LEU
1	Q	145	SER
1	Q	147	ARG

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Mol	Chain	Res	Type
1	Q	148	THR
1	Q	149	THR
1	Q	167	GLU
1	Q	171	LEU
1	Q	176	MET
1	Q	179	ASP
1	Q	180	LEU
1	Q	183	ARG
1	Q	208	ILE
1	Q	227	MET
1	Q	232	SER
1	Q	236	LEU
1	Q	244	ASP
1	Q	269	MET
1	Q	286	ASP
1	Q	335	ARG
1	Q	350	SER
1	Q	357	ILE
1	Q	370	VAL
1	Q	372	ARG
1	Q	373	LYS
1	R	8	LEU
1	R	33	SER
1	R	35	VAL
1	R	37	ARG
1	R	45	VAL
1	R	49	GLN
1	R	51	ASP
1	R	56	ASP
1	R	57	GLU
1	R	66	THR
1	R	68	LYS
1	R	71	ILE
1	R	113	LYS
1	R	132	MET
1	R	139	VAL
1	R	141	SER
1	R	142	LEU
1	R	145	SER
1	R	147	ARG
1	R	148	THR
1	R	149	THR

Continued on next page...

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Mol	Chain	Res	Type
1	R	167	GLU
1	R	171	LEU
1	R	176	MET
1	R	179	ASP
1	R	180	LEU
1	R	183	ARG
1	R	208	ILE
1	R	227	MET
1	R	232	SER
1	R	236	LEU
1	R	244	ASP
1	R	269	MET
1	R	286	ASP
1	R	335	ARG
1	R	350	SER
1	R	357	ILE
1	R	370	VAL
1	R	372	ARG
1	R	373	LYS
1	S	8	LEU
1	S	33	SER
1	S	35	VAL
1	S	37	ARG
1	S	45	VAL
1	S	49	GLN
1	S	51	ASP
1	S	56	ASP
1	S	57	GLU
1	S	66	THR
1	S	68	LYS
1	S	71	ILE
1	S	113	LYS
1	S	132	MET
1	S	139	VAL
1	S	141	SER
1	S	142	LEU
1	S	145	SER
1	S	147	ARG
1	S	148	THR
1	S	149	THR
1	S	167	GLU
1	S	171	LEU

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Mol	Chain	Res	Type
1	S	176	MET
1	S	179	ASP
1	S	180	LEU
1	S	183	ARG
1	S	208	ILE
1	S	227	MET
1	S	232	SER
1	S	236	LEU
1	S	244	ASP
1	S	269	MET
1	S	286	ASP
1	S	335	ARG
1	S	350	SER
1	S	357	ILE
1	S	370	VAL
1	S	372	ARG
1	S	373	LYS
1	T	8	LEU
1	T	33	SER
1	T	35	VAL
1	T	37	ARG
1	T	45	VAL
1	T	49	GLN
1	T	51	ASP
1	T	56	ASP
1	T	57	GLU
1	T	66	THR
1	T	68	LYS
1	T	71	ILE
1	T	113	LYS
1	T	132	MET
1	T	139	VAL
1	T	141	SER
1	T	142	LEU
1	T	145	SER
1	T	147	ARG
1	T	148	THR
1	T	149	THR
1	T	167	GLU
1	T	171	LEU
1	T	176	MET
1	T	179	ASP

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Mol	Chain	Res	Type
1	T	180	LEU
1	T	183	ARG
1	T	208	ILE
1	T	227	MET
1	T	232	SER
1	T	236	LEU
1	T	244	ASP
1	T	269	MET
1	T	286	ASP
1	T	335	ARG
1	T	350	SER
1	T	357	ILE
1	T	370	VAL
1	T	372	ARG
1	T	373	LYS
1	U	8	LEU
1	U	33	SER
1	U	35	VAL
1	U	37	ARG
1	U	45	VAL
1	U	49	GLN
1	U	51	ASP
1	U	56	ASP
1	U	57	GLU
1	U	66	THR
1	U	68	LYS
1	U	71	ILE
1	U	113	LYS
1	U	132	MET
1	U	139	VAL
1	U	141	SER
1	U	142	LEU
1	U	145	SER
1	U	147	ARG
1	U	148	THR
1	U	149	THR
1	U	167	GLU
1	U	171	LEU
1	U	176	MET
1	U	179	ASP
1	U	180	LEU
1	U	183	ARG

Continued on next page...

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Mol	Chain	Res	Type
1	U	208	ILE
1	U	227	MET
1	U	232	SER
1	U	236	LEU
1	U	244	ASP
1	U	269	MET
1	U	286	ASP
1	U	335	ARG
1	U	350	SER
1	U	357	ILE
1	U	370	VAL
1	U	372	ARG
1	U	373	LYS
1	V	8	LEU
1	V	33	SER
1	V	35	VAL
1	V	37	ARG
1	V	45	VAL
1	V	49	GLN
1	V	51	ASP
1	V	56	ASP
1	V	57	GLU
1	V	66	THR
1	V	68	LYS
1	V	71	ILE
1	V	113	LYS
1	V	132	MET
1	V	139	VAL
1	V	141	SER
1	V	142	LEU
1	V	145	SER
1	V	147	ARG
1	V	148	THR
1	V	149	THR
1	V	167	GLU
1	V	171	LEU
1	V	176	MET
1	V	179	ASP
1	V	180	LEU
1	V	183	ARG
1	V	208	ILE
1	V	227	MET

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Mol	Chain	Res	Type
1	V	232	SER
1	V	236	LEU
1	V	244	ASP
1	V	269	MET
1	V	286	ASP
1	V	335	ARG
1	V	350	SER
1	V	357	ILE
1	V	370	VAL
1	V	372	ARG
1	V	373	LYS
1	W	8	LEU
1	W	33	SER
1	W	35	VAL
1	W	37	ARG
1	W	45	VAL
1	W	49	GLN
1	W	51	ASP
1	W	56	ASP
1	W	57	GLU
1	W	66	THR
1	W	68	LYS
1	W	71	ILE
1	W	113	LYS
1	W	132	MET
1	W	139	VAL
1	W	141	SER
1	W	142	LEU
1	W	145	SER
1	W	147	ARG
1	W	148	THR
1	W	149	THR
1	W	167	GLU
1	W	171	LEU
1	W	176	MET
1	W	179	ASP
1	W	180	LEU
1	W	183	ARG
1	W	208	ILE
1	W	227	MET
1	W	232	SER
1	W	236	LEU

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Mol	Chain	Res	Type
1	W	244	ASP
1	W	269	MET
1	W	286	ASP
1	W	335	ARG
1	W	350	SER
1	W	357	ILE
1	W	370	VAL
1	W	372	ARG
1	W	373	LYS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (100) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	49	GLN
1	A	59	GLN
1	A	121	GLN
1	A	246	GLN
1	A	353	GLN
1	B	49	GLN
1	B	59	GLN
1	B	121	GLN
1	B	246	GLN
1	B	353	GLN
1	C	49	GLN
1	C	59	GLN
1	C	121	GLN
1	C	173	HIS
1	C	353	GLN
1	D	49	GLN
1	D	59	GLN
1	D	121	GLN
1	D	353	GLN
1	E	49	GLN
1	E	59	GLN
1	E	121	GLN
1	E	173	HIS
1	E	353	GLN
1	F	49	GLN
1	F	59	GLN
1	F	121	GLN
1	F	353	GLN
1	G	49	GLN

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Mol	Chain	Res	Type
1	G	59	GLN
1	G	121	GLN
1	G	173	HIS
1	G	353	GLN
1	H	49	GLN
1	H	59	GLN
1	H	121	GLN
1	H	353	GLN
1	I	49	GLN
1	I	59	GLN
1	I	121	GLN
1	I	353	GLN
1	J	49	GLN
1	J	59	GLN
1	J	121	GLN
1	J	353	GLN
1	K	49	GLN
1	K	59	GLN
1	K	121	GLN
1	K	353	GLN
1	L	49	GLN
1	L	59	GLN
1	L	121	GLN
1	L	353	GLN
1	M	49	GLN
1	M	59	GLN
1	M	121	GLN
1	M	353	GLN
1	N	49	GLN
1	N	59	GLN
1	N	121	GLN
1	N	173	HIS
1	N	353	GLN
1	O	49	GLN
1	O	59	GLN
1	O	121	GLN
1	O	353	GLN
1	P	49	GLN
1	P	59	GLN
1	P	121	GLN
1	P	353	GLN
1	Q	49	GLN

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Mol	Chain	Res	Type
1	Q	59	GLN
1	Q	121	GLN
1	Q	353	GLN
1	R	49	GLN
1	R	59	GLN
1	R	121	GLN
1	R	173	HIS
1	R	353	GLN
1	S	49	GLN
1	S	59	GLN
1	S	121	GLN
1	S	353	GLN
1	T	49	GLN
1	T	59	GLN
1	T	121	GLN
1	T	173	HIS
1	T	353	GLN
1	U	49	GLN
1	U	59	GLN
1	U	121	GLN
1	U	353	GLN
1	V	49	GLN
1	V	59	GLN
1	V	121	GLN
1	V	353	GLN
1	W	49	GLN
1	W	59	GLN
1	W	121	GLN
1	W	353	GLN

5.3.3 RNA

There are no RNA molecules in this entry.

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

23 non-standard protein/DNA/RNA residues are modelled in this entry.

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	$\#\|Z\| > 2$
1	HIC	L	73	1	10,11,12	0.67	0	9,14,16	3.27	4 (44%)
1	HIC	D	73	1	10,11,12	0.66	0	9,14,16	3.30	4 (44%)
1	HIC	F	73	1	10,11,12	0.66	0	9,14,16	3.30	4 (44%)
1	HIC	E	73	1	10,11,12	0.67	0	9,14,16	3.31	4 (44%)
1	HIC	P	73	1	10,11,12	0.67	0	9,14,16	3.26	4 (44%)
1	HIC	Q	73	1	10,11,12	0.67	0	9,14,16	3.30	4 (44%)
1	HIC	M	73	1	10,11,12	0.66	0	9,14,16	3.32	4 (44%)
1	HIC	W	73	1	10,11,12	0.66	0	9,14,16	3.32	4 (44%)
1	HIC	A	73	1	10,11,12	0.67	0	9,14,16	3.28	4 (44%)
1	HIC	K	73	1	10,11,12	0.66	0	9,14,16	3.29	4 (44%)
1	HIC	S	73	1	10,11,12	0.67	0	9,14,16	3.33	4 (44%)
1	HIC	U	73	1	10,11,12	0.66	0	9,14,16	3.27	4 (44%)
1	HIC	C	73	1	10,11,12	0.67	0	9,14,16	3.32	4 (44%)
1	HIC	V	73	1	10,11,12	0.66	0	9,14,16	3.27	4 (44%)
1	HIC	H	73	1	10,11,12	0.66	0	9,14,16	3.28	4 (44%)
1	HIC	O	73	1	10,11,12	0.66	0	9,14,16	3.29	4 (44%)
1	HIC	R	73	1	10,11,12	0.66	0	9,14,16	3.27	4 (44%)
1	HIC	J	73	1	10,11,12	0.66	0	9,14,16	3.27	4 (44%)
1	HIC	T	73	1	10,11,12	0.67	0	9,14,16	3.29	4 (44%)
1	HIC	N	73	1	10,11,12	0.66	0	9,14,16	3.28	4 (44%)
1	HIC	I	73	1	10,11,12	0.67	0	9,14,16	3.31	4 (44%)
1	HIC	G	73	1	10,11,12	0.67	0	9,14,16	3.29	4 (44%)
1	HIC	B	73	1	10,11,12	0.67	0	9,14,16	3.29	4 (44%)

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
1	HIC	L	73	1	-	1/5/6/8	0/1/1/1
1	HIC	D	73	1	-	1/5/6/8	0/1/1/1
1	HIC	F	73	1	-	1/5/6/8	0/1/1/1

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
1	HIC	E	73	1	-	1/5/6/8	0/1/1/1
1	HIC	P	73	1	-	1/5/6/8	0/1/1/1
1	HIC	Q	73	1	-	1/5/6/8	0/1/1/1
1	HIC	M	73	1	-	1/5/6/8	0/1/1/1
1	HIC	W	73	1	-	1/5/6/8	0/1/1/1
1	HIC	A	73	1	-	1/5/6/8	0/1/1/1
1	HIC	K	73	1	-	1/5/6/8	0/1/1/1
1	HIC	S	73	1	-	1/5/6/8	0/1/1/1
1	HIC	U	73	1	-	1/5/6/8	0/1/1/1
1	HIC	C	73	1	-	1/5/6/8	0/1/1/1
1	HIC	V	73	1	-	1/5/6/8	0/1/1/1
1	HIC	H	73	1	-	1/5/6/8	0/1/1/1
1	HIC	O	73	1	-	1/5/6/8	0/1/1/1
1	HIC	R	73	1	-	1/5/6/8	0/1/1/1
1	HIC	J	73	1	-	1/5/6/8	0/1/1/1
1	HIC	T	73	1	-	1/5/6/8	0/1/1/1
1	HIC	N	73	1	-	1/5/6/8	0/1/1/1
1	HIC	I	73	1	-	1/5/6/8	0/1/1/1
1	HIC	G	73	1	-	1/5/6/8	0/1/1/1
1	HIC	B	73	1	-	1/5/6/8	0/1/1/1

There are no bond length outliers.

All (92) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	M	73	HIC	NE2-CE1-ND1	-8.78	109.30	112.66
1	C	73	HIC	NE2-CE1-ND1	-8.77	109.31	112.66
1	S	73	HIC	NE2-CE1-ND1	-8.77	109.31	112.66
1	E	73	HIC	NE2-CE1-ND1	-8.75	109.31	112.66
1	W	73	HIC	NE2-CE1-ND1	-8.75	109.31	112.66
1	I	73	HIC	NE2-CE1-ND1	-8.69	109.33	112.66
1	Q	73	HIC	NE2-CE1-ND1	-8.68	109.34	112.66
1	F	73	HIC	NE2-CE1-ND1	-8.68	109.34	112.66
1	D	73	HIC	NE2-CE1-ND1	-8.67	109.34	112.66
1	O	73	HIC	NE2-CE1-ND1	-8.65	109.35	112.66
1	K	73	HIC	NE2-CE1-ND1	-8.64	109.36	112.66
1	T	73	HIC	NE2-CE1-ND1	-8.64	109.36	112.66
1	H	73	HIC	NE2-CE1-ND1	-8.63	109.36	112.66
1	G	73	HIC	NE2-CE1-ND1	-8.63	109.36	112.66
1	B	73	HIC	NE2-CE1-ND1	-8.62	109.36	112.66
1	N	73	HIC	NE2-CE1-ND1	-8.60	109.37	112.66
1	V	73	HIC	NE2-CE1-ND1	-8.60	109.37	112.66

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	R	73	HIC	NE2-CE1-ND1	-8.58	109.38	112.66
1	A	73	HIC	NE2-CE1-ND1	-8.57	109.38	112.66
1	U	73	HIC	NE2-CE1-ND1	-8.57	109.38	112.66
1	L	73	HIC	NE2-CE1-ND1	-8.56	109.38	112.66
1	P	73	HIC	NE2-CE1-ND1	-8.54	109.39	112.66
1	J	73	HIC	NE2-CE1-ND1	-8.54	109.39	112.66
1	A	73	HIC	CD2-NE2-CE1	-2.82	105.60	106.71
1	J	73	HIC	CD2-NE2-CE1	-2.78	105.61	106.71
1	G	73	HIC	CD2-NE2-CE1	-2.75	105.63	106.71
1	L	73	HIC	CD2-NE2-CE1	-2.74	105.63	106.71
1	K	73	HIC	CD2-NE2-CE1	-2.73	105.63	106.71
1	P	73	HIC	CD2-NE2-CE1	-2.71	105.64	106.71
1	B	73	HIC	CD2-NE2-CE1	-2.71	105.64	106.71
1	U	73	HIC	CD2-NE2-CE1	-2.70	105.65	106.71
1	F	73	HIC	CD2-NE2-CE1	-2.69	105.65	106.71
1	I	73	HIC	CD2-NE2-CE1	-2.67	105.66	106.71
1	M	73	HIC	CE1-ND1-CG	2.66	110.07	105.80
1	W	73	HIC	CE1-ND1-CG	2.66	110.07	105.80
1	D	73	HIC	CE1-ND1-CG	2.65	110.05	105.80
1	B	73	HIC	CE1-ND1-CG	2.65	110.04	105.80
1	O	73	HIC	CD2-NE2-CE1	-2.65	105.67	106.71
1	S	73	HIC	CE1-ND1-CG	2.64	110.04	105.80
1	N	73	HIC	CD2-NE2-CE1	-2.64	105.67	106.71
1	C	73	HIC	CE1-ND1-CG	2.64	110.03	105.80
1	D	73	HIC	CD2-NE2-CE1	-2.64	105.67	106.71
1	E	73	HIC	CE1-ND1-CG	2.63	110.02	105.80
1	R	73	HIC	CD2-NE2-CE1	-2.63	105.67	106.71
1	S	73	HIC	CD2-NE2-CE1	-2.63	105.67	106.71
1	T	73	HIC	CE1-ND1-CG	2.63	110.02	105.80
1	G	73	HIC	CE1-ND1-CG	2.63	110.01	105.80
1	V	73	HIC	CD2-NE2-CE1	-2.63	105.67	106.71
1	N	73	HIC	CE1-ND1-CG	2.63	110.01	105.80
1	K	73	HIC	CE1-ND1-CG	2.63	110.01	105.80
1	L	73	HIC	CE1-ND1-CG	2.63	110.01	105.80
1	A	73	HIC	CE1-ND1-CG	2.63	110.01	105.80
1	J	73	HIC	CE1-ND1-CG	2.62	110.01	105.80
1	Q	73	HIC	CE1-ND1-CG	2.62	110.01	105.80
1	T	73	HIC	CD2-NE2-CE1	-2.62	105.67	106.71
1	I	73	HIC	CE1-ND1-CG	2.62	109.99	105.80
1	W	73	HIC	CD2-NE2-CE1	-2.61	105.68	106.71
1	C	73	HIC	CD2-NE2-CE1	-2.61	105.68	106.71
1	F	73	HIC	CE1-ND1-CG	2.61	109.98	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	V	73	HIC	CE1-ND1-CG	2.61	109.98	105.80
1	U	73	HIC	CE1-ND1-CG	2.61	109.98	105.80
1	P	73	HIC	CE1-ND1-CG	2.61	109.98	105.80
1	O	73	HIC	CE1-ND1-CG	2.61	109.98	105.80
1	H	73	HIC	CE1-ND1-CG	2.60	109.97	105.80
1	R	73	HIC	CE1-ND1-CG	2.60	109.97	105.80
1	Q	73	HIC	CD2-NE2-CE1	-2.58	105.69	106.71
1	H	73	HIC	CD2-NE2-CE1	-2.57	105.70	106.71
1	E	73	HIC	CD2-NE2-CE1	-2.56	105.70	106.71
1	M	73	HIC	CD2-NE2-CE1	-2.54	105.71	106.71
1	B	73	HIC	CD2-CG-ND1	-2.17	104.72	109.43
1	D	73	HIC	CD2-CG-ND1	-2.17	104.72	109.43
1	M	73	HIC	CD2-CG-ND1	-2.17	104.72	109.43
1	W	73	HIC	CD2-CG-ND1	-2.16	104.73	109.43
1	N	73	HIC	CD2-CG-ND1	-2.16	104.74	109.43
1	H	73	HIC	CD2-CG-ND1	-2.16	104.75	109.43
1	T	73	HIC	CD2-CG-ND1	-2.16	104.75	109.43
1	Q	73	HIC	CD2-CG-ND1	-2.16	104.75	109.43
1	I	73	HIC	CD2-CG-ND1	-2.15	104.75	109.43
1	K	73	HIC	CD2-CG-ND1	-2.15	104.76	109.43
1	U	73	HIC	CD2-CG-ND1	-2.15	104.76	109.43
1	E	73	HIC	CD2-CG-ND1	-2.15	104.76	109.43
1	V	73	HIC	CD2-CG-ND1	-2.15	104.76	109.43
1	S	73	HIC	CD2-CG-ND1	-2.14	104.77	109.43
1	A	73	HIC	CD2-CG-ND1	-2.14	104.78	109.43
1	J	73	HIC	CD2-CG-ND1	-2.14	104.78	109.43
1	F	73	HIC	CD2-CG-ND1	-2.14	104.79	109.43
1	C	73	HIC	CD2-CG-ND1	-2.14	104.79	109.43
1	G	73	HIC	CD2-CG-ND1	-2.14	104.79	109.43
1	O	73	HIC	CD2-CG-ND1	-2.14	104.79	109.43
1	P	73	HIC	CD2-CG-ND1	-2.13	104.80	109.43
1	L	73	HIC	CD2-CG-ND1	-2.13	104.81	109.43
1	R	73	HIC	CD2-CG-ND1	-2.13	104.81	109.43

There are no chirality outliers.

All (23) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
1	A	73	HIC	O-C-CA-CB
1	B	73	HIC	O-C-CA-CB
1	C	73	HIC	O-C-CA-CB
1	D	73	HIC	O-C-CA-CB

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Mol	Chain	Res	Type	Atoms
1	E	73	HIC	O-C-CA-CB
1	F	73	HIC	O-C-CA-CB
1	G	73	HIC	O-C-CA-CB
1	H	73	HIC	O-C-CA-CB
1	I	73	HIC	O-C-CA-CB
1	J	73	HIC	O-C-CA-CB
1	K	73	HIC	O-C-CA-CB
1	L	73	HIC	O-C-CA-CB
1	M	73	HIC	O-C-CA-CB
1	N	73	HIC	O-C-CA-CB
1	O	73	HIC	O-C-CA-CB
1	P	73	HIC	O-C-CA-CB
1	Q	73	HIC	O-C-CA-CB
1	R	73	HIC	O-C-CA-CB
1	S	73	HIC	O-C-CA-CB
1	T	73	HIC	O-C-CA-CB
1	U	73	HIC	O-C-CA-CB
1	V	73	HIC	O-C-CA-CB
1	W	73	HIC	O-C-CA-CB

There are no ring outliers.

23 monomers are involved in 43 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
1	L	73	HIC	2	0
1	D	73	HIC	2	0
1	F	73	HIC	2	0
1	E	73	HIC	2	0
1	P	73	HIC	1	0
1	Q	73	HIC	2	0
1	M	73	HIC	2	0
1	W	73	HIC	2	0
1	A	73	HIC	2	0
1	K	73	HIC	2	0
1	S	73	HIC	2	0
1	U	73	HIC	2	0
1	C	73	HIC	2	0
1	V	73	HIC	2	0
1	H	73	HIC	2	0
1	O	73	HIC	2	0
1	R	73	HIC	1	0
1	J	73	HIC	2	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
1	T	73	HIC	1	0
1	N	73	HIC	2	0
1	I	73	HIC	2	0
1	G	73	HIC	2	0
1	B	73	HIC	2	0

5.5 Carbohydrates (i)

There are no oligosaccharides in this entry.

5.6 Ligand geometry (i)

23 ligands are modelled in this entry.

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	$\#\|Z\| > 2$
2	ADP	E	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	S	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	Q	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	U	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	W	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	N	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	V	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	G	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	I	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	T	401	-	28,29,29	1.11	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	K	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	L	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	M	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	R	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	C	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.25	6 (13%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
2	ADP	P	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	J	401	-	28,29,29	1.13	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	B	401	-	28,29,29	1.11	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	A	401	-	28,29,29	1.11	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	F	401	-	28,29,29	1.11	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	O	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.25	6 (13%)
2	ADP	H	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)
2	ADP	D	401	-	28,29,29	1.12	2 (7%)	43,45,45	1.24	6 (13%)

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
2	ADP	E	401	-	-	0/16/32/32	0/3/3/3
2	ADP	S	401	-	-	0/16/32/32	0/3/3/3
2	ADP	Q	401	-	-	0/16/32/32	0/3/3/3
2	ADP	U	401	-	-	0/16/32/32	0/3/3/3
2	ADP	W	401	-	-	0/16/32/32	0/3/3/3
2	ADP	N	401	-	-	0/16/32/32	0/3/3/3
2	ADP	V	401	-	-	0/16/32/32	0/3/3/3
2	ADP	G	401	-	-	0/16/32/32	0/3/3/3
2	ADP	I	401	-	-	0/16/32/32	0/3/3/3
2	ADP	T	401	-	-	0/16/32/32	0/3/3/3
2	ADP	K	401	-	-	0/16/32/32	0/3/3/3
2	ADP	L	401	-	-	0/16/32/32	0/3/3/3
2	ADP	M	401	-	-	0/16/32/32	0/3/3/3
2	ADP	R	401	-	-	0/16/32/32	0/3/3/3
2	ADP	C	401	-	-	0/16/32/32	0/3/3/3
2	ADP	P	401	-	-	0/16/32/32	0/3/3/3
2	ADP	J	401	-	-	0/16/32/32	0/3/3/3
2	ADP	B	401	-	-	0/16/32/32	0/3/3/3
2	ADP	A	401	-	-	0/16/32/32	0/3/3/3
2	ADP	F	401	-	-	0/16/32/32	0/3/3/3
2	ADP	O	401	-	-	0/16/32/32	0/3/3/3
2	ADP	H	401	-	-	0/16/32/32	0/3/3/3
2	ADP	D	401	-	-	0/16/32/32	0/3/3/3

All (46) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	J	401	ADP	C5-N7	-3.28	1.33	1.39
2	Q	401	ADP	C5-N7	-3.27	1.33	1.39
2	D	401	ADP	C5-N7	-3.25	1.33	1.39
2	L	401	ADP	C5-N7	-3.25	1.33	1.39
2	I	401	ADP	C5-N7	-3.25	1.33	1.39
2	V	401	ADP	C5-N7	-3.25	1.33	1.39
2	M	401	ADP	C5-N7	-3.24	1.33	1.39
2	A	401	ADP	C5-N7	-3.24	1.33	1.39
2	G	401	ADP	C5-N7	-3.24	1.33	1.39
2	S	401	ADP	C5-N7	-3.23	1.33	1.39
2	K	401	ADP	C5-N7	-3.23	1.33	1.39
2	P	401	ADP	C5-N7	-3.23	1.33	1.39
2	R	401	ADP	C5-N7	-3.23	1.33	1.39
2	C	401	ADP	C5-N7	-3.23	1.33	1.39
2	W	401	ADP	C5-N7	-3.22	1.33	1.39
2	N	401	ADP	C5-N7	-3.22	1.33	1.39
2	O	401	ADP	C5-N7	-3.22	1.33	1.39
2	H	401	ADP	C5-N7	-3.21	1.33	1.39
2	E	401	ADP	C5-N7	-3.21	1.33	1.39
2	T	401	ADP	C5-N7	-3.20	1.33	1.39
2	F	401	ADP	C5-N7	-3.20	1.33	1.39
2	B	401	ADP	C5-N7	-3.20	1.33	1.39
2	U	401	ADP	C5-N7	-3.19	1.33	1.39
2	S	401	ADP	C2-N1	3.16	1.39	1.33
2	E	401	ADP	C2-N1	3.14	1.39	1.33
2	P	401	ADP	C2-N1	3.12	1.39	1.33
2	Q	401	ADP	C2-N1	3.12	1.39	1.33
2	K	401	ADP	C2-N1	3.12	1.39	1.33
2	G	401	ADP	C2-N1	3.12	1.39	1.33
2	J	401	ADP	C2-N1	3.12	1.39	1.33
2	C	401	ADP	C2-N1	3.12	1.39	1.33
2	L	401	ADP	C2-N1	3.12	1.39	1.33
2	O	401	ADP	C2-N1	3.11	1.39	1.33
2	U	401	ADP	C2-N1	3.11	1.39	1.33
2	M	401	ADP	C2-N1	3.11	1.39	1.33
2	W	401	ADP	C2-N1	3.11	1.39	1.33
2	R	401	ADP	C2-N1	3.11	1.39	1.33
2	I	401	ADP	C2-N1	3.10	1.39	1.33
2	T	401	ADP	C2-N1	3.10	1.39	1.33
2	H	401	ADP	C2-N1	3.09	1.39	1.33
2	B	401	ADP	C2-N1	3.08	1.39	1.33
2	N	401	ADP	C2-N1	3.08	1.39	1.33
2	D	401	ADP	C2-N1	3.07	1.39	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	V	401	ADP	C2-N1	3.07	1.39	1.33
2	A	401	ADP	C2-N1	3.06	1.39	1.33
2	F	401	ADP	C2-N1	3.06	1.39	1.33

All (138) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
2	Q	401	ADP	C5-C4-N3	2.57	130.25	126.72
2	O	401	ADP	C4-C5-N7	2.57	113.52	110.58
2	O	401	ADP	C5-C4-N3	2.55	130.23	126.72
2	L	401	ADP	C4-C5-N7	2.55	113.50	110.58
2	Q	401	ADP	C4-C5-N7	2.54	113.49	110.58
2	J	401	ADP	C4-C5-N7	2.54	113.49	110.58
2	D	401	ADP	C5-C4-N3	2.54	130.21	126.72
2	C	401	ADP	C4-C5-N7	2.54	113.49	110.58
2	N	401	ADP	C4-C5-N7	2.54	113.49	110.58
2	G	401	ADP	C5-C4-N3	2.54	130.21	126.72
2	R	401	ADP	C4-C5-N7	2.54	113.48	110.58
2	D	401	ADP	C4-C5-N7	2.54	113.48	110.58
2	G	401	ADP	C4-C5-N7	2.54	113.48	110.58
2	I	401	ADP	C5-C4-N3	2.54	130.21	126.72
2	B	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	J	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	R	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	A	401	ADP	C4-C5-N7	2.53	113.48	110.58
2	T	401	ADP	C4-C5-N7	2.53	113.48	110.58
2	S	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	U	401	ADP	C4-C5-N7	2.53	113.47	110.58
2	L	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	U	401	ADP	C5-C4-N3	2.53	130.20	126.72
2	B	401	ADP	C4-C5-N7	2.52	113.47	110.58
2	C	401	ADP	C5-C4-N3	2.52	130.19	126.72
2	M	401	ADP	C5-C4-N3	2.52	130.18	126.72
2	H	401	ADP	C4-C5-N7	2.52	113.46	110.58
2	T	401	ADP	C5-C4-N3	2.51	130.18	126.72
2	E	401	ADP	C4-C5-N7	2.51	113.45	110.58
2	M	401	ADP	C4-C5-N7	2.51	113.45	110.58
2	V	401	ADP	C4-C5-N7	2.51	113.45	110.58
2	S	401	ADP	C4-C5-N7	2.51	113.45	110.58
2	A	401	ADP	C5-C4-N3	2.51	130.17	126.72
2	N	401	ADP	C5-C4-N3	2.51	130.17	126.72
2	H	401	ADP	C5-C4-N3	2.51	130.17	126.72

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
2	F	401	ADP	C4-C5-N7	2.51	113.45	110.58
2	I	401	ADP	C4-C5-N7	2.50	113.44	110.58
2	V	401	ADP	C5-C4-N3	2.50	130.16	126.72
2	F	401	ADP	C5-C4-N3	2.50	130.16	126.72
2	W	401	ADP	C5-C4-N3	2.50	130.16	126.72
2	K	401	ADP	C4-C5-N7	2.50	113.44	110.58
2	K	401	ADP	C5-C4-N3	2.50	130.15	126.72
2	P	401	ADP	C4-C5-N7	2.49	113.43	110.58
2	E	401	ADP	C5-C4-N3	2.49	130.14	126.72
2	P	401	ADP	C5-C4-N3	2.49	130.14	126.72
2	W	401	ADP	C4-C5-N7	2.48	113.41	110.58
2	R	401	ADP	C4-N9-C8	-2.35	103.28	105.74
2	C	401	ADP	C4-N9-C8	-2.34	103.28	105.74
2	B	401	ADP	C4-N9-C8	-2.33	103.30	105.74
2	E	401	ADP	C4-N9-C8	-2.32	103.30	105.74
2	T	401	ADP	C4-N9-C8	-2.32	103.30	105.74
2	H	401	ADP	C4-N9-C8	-2.32	103.30	105.74
2	J	401	ADP	C4-N9-C8	-2.32	103.30	105.74
2	F	401	ADP	C2-N1-C6	2.32	122.53	118.73
2	W	401	ADP	C4-N9-C8	-2.32	103.31	105.74
2	L	401	ADP	C4-N9-C8	-2.31	103.31	105.74
2	D	401	ADP	C4-N9-C8	-2.31	103.31	105.74
2	G	401	ADP	C2-N1-C6	2.31	122.52	118.73
2	F	401	ADP	C4-N9-C8	-2.31	103.32	105.74
2	A	401	ADP	C4-N9-C8	-2.31	103.32	105.74
2	K	401	ADP	C2-N1-C6	2.31	122.52	118.73
2	N	401	ADP	C4-N9-C8	-2.30	103.32	105.74
2	T	401	ADP	C2-N1-C6	2.30	122.51	118.73
2	Q	401	ADP	C4-N9-C8	-2.30	103.32	105.74
2	P	401	ADP	C4-N9-C8	-2.30	103.32	105.74
2	R	401	ADP	C2-N1-C6	2.30	122.51	118.73
2	S	401	ADP	C4-N9-C8	-2.30	103.33	105.74
2	I	401	ADP	C2-N1-C6	2.30	122.50	118.73
2	N	401	ADP	C2-N1-C6	2.30	122.50	118.73
2	W	401	ADP	C2-N1-C6	2.29	122.50	118.73
2	A	401	ADP	C2-N1-C6	2.29	122.49	118.73
2	O	401	ADP	C2-N1-C6	2.29	122.49	118.73
2	M	401	ADP	C4-N9-C8	-2.29	103.33	105.74
2	D	401	ADP	C2-N1-C6	2.29	122.49	118.73
2	C	401	ADP	C2-N1-C6	2.29	122.48	118.73
2	K	401	ADP	C4-N9-C8	-2.28	103.34	105.74
2	V	401	ADP	C2-N1-C6	2.28	122.48	118.73

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
2	E	401	ADP	C2-N1-C6	2.28	122.48	118.73
2	H	401	ADP	C2-N1-C6	2.28	122.47	118.73
2	V	401	ADP	C4-N9-C8	-2.28	103.34	105.74
2	I	401	ADP	C4-N9-C8	-2.28	103.35	105.74
2	U	401	ADP	C4-N9-C8	-2.28	103.35	105.74
2	J	401	ADP	C2-N1-C6	2.27	122.47	118.73
2	O	401	ADP	C4-N9-C8	-2.27	103.35	105.74
2	B	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	S	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	U	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	M	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	L	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	Q	401	ADP	C2-N1-C6	2.27	122.46	118.73
2	G	401	ADP	C4-N9-C8	-2.26	103.36	105.74
2	P	401	ADP	C2-N1-C6	2.26	122.44	118.73
2	G	401	ADP	N3-C2-N1	-2.23	125.20	128.58
2	K	401	ADP	N3-C2-N1	-2.23	125.21	128.58
2	E	401	ADP	N3-C2-N1	-2.21	125.23	128.58
2	T	401	ADP	N3-C2-N1	-2.21	125.24	128.58
2	C	401	ADP	N3-C2-N1	-2.21	125.24	128.58
2	J	401	ADP	N3-C2-N1	-2.21	125.24	128.58
2	L	401	ADP	N3-C2-N1	-2.21	125.24	128.58
2	W	401	ADP	N3-C2-N1	-2.20	125.25	128.58
2	R	401	ADP	N3-C2-N1	-2.20	125.25	128.58
2	F	401	ADP	N3-C2-N1	-2.20	125.26	128.58
2	N	401	ADP	N3-C2-N1	-2.20	125.26	128.58
2	V	401	ADP	N3-C2-N1	-2.20	125.26	128.58
2	A	401	ADP	N3-C2-N1	-2.19	125.27	128.58
2	I	401	ADP	N3-C2-N1	-2.18	125.28	128.58
2	M	401	ADP	N3-C2-N1	-2.18	125.28	128.58
2	U	401	ADP	N3-C2-N1	-2.18	125.28	128.58
2	O	401	ADP	N3-C2-N1	-2.18	125.28	128.58
2	S	401	ADP	N3-C2-N1	-2.18	125.28	128.58
2	B	401	ADP	N3-C2-N1	-2.18	125.29	128.58
2	H	401	ADP	N3-C2-N1	-2.17	125.29	128.58
2	P	401	ADP	N3-C2-N1	-2.17	125.29	128.58
2	Q	401	ADP	N3-C2-N1	-2.17	125.30	128.58
2	D	401	ADP	N3-C2-N1	-2.17	125.30	128.58
2	M	401	ADP	O3B-PB-O2B	2.10	115.69	107.80
2	L	401	ADP	O3B-PB-O2B	2.10	115.69	107.80
2	T	401	ADP	O3B-PB-O2B	2.10	115.69	107.80
2	W	401	ADP	O3B-PB-O2B	2.10	115.68	107.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
2	G	401	ADP	O3B-PB-O2B	2.10	115.67	107.80
2	B	401	ADP	O3B-PB-O2B	2.10	115.66	107.80
2	R	401	ADP	O3B-PB-O2B	2.09	115.65	107.80
2	P	401	ADP	O3B-PB-O2B	2.09	115.65	107.80
2	E	401	ADP	O3B-PB-O2B	2.09	115.65	107.80
2	J	401	ADP	O3B-PB-O2B	2.09	115.65	107.80
2	K	401	ADP	O3B-PB-O2B	2.09	115.65	107.80
2	H	401	ADP	O3B-PB-O2B	2.09	115.64	107.80
2	C	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	Q	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	V	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	N	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	S	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	I	401	ADP	O3B-PB-O2B	2.09	115.63	107.80
2	D	401	ADP	O3B-PB-O2B	2.09	115.62	107.80
2	O	401	ADP	O3B-PB-O2B	2.08	115.62	107.80
2	U	401	ADP	O3B-PB-O2B	2.08	115.62	107.80
2	A	401	ADP	O3B-PB-O2B	2.08	115.61	107.80
2	F	401	ADP	O3B-PB-O2B	2.08	115.61	107.80

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

23 monomers are involved in 95 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
2	E	401	ADP	4	0
2	S	401	ADP	4	0
2	Q	401	ADP	4	0
2	U	401	ADP	4	0
2	W	401	ADP	4	0
2	N	401	ADP	4	0
2	V	401	ADP	4	0
2	G	401	ADP	4	0
2	I	401	ADP	4	0
2	T	401	ADP	4	0
2	K	401	ADP	4	0
2	L	401	ADP	4	0
2	M	401	ADP	5	0
2	R	401	ADP	4	0
2	C	401	ADP	4	0

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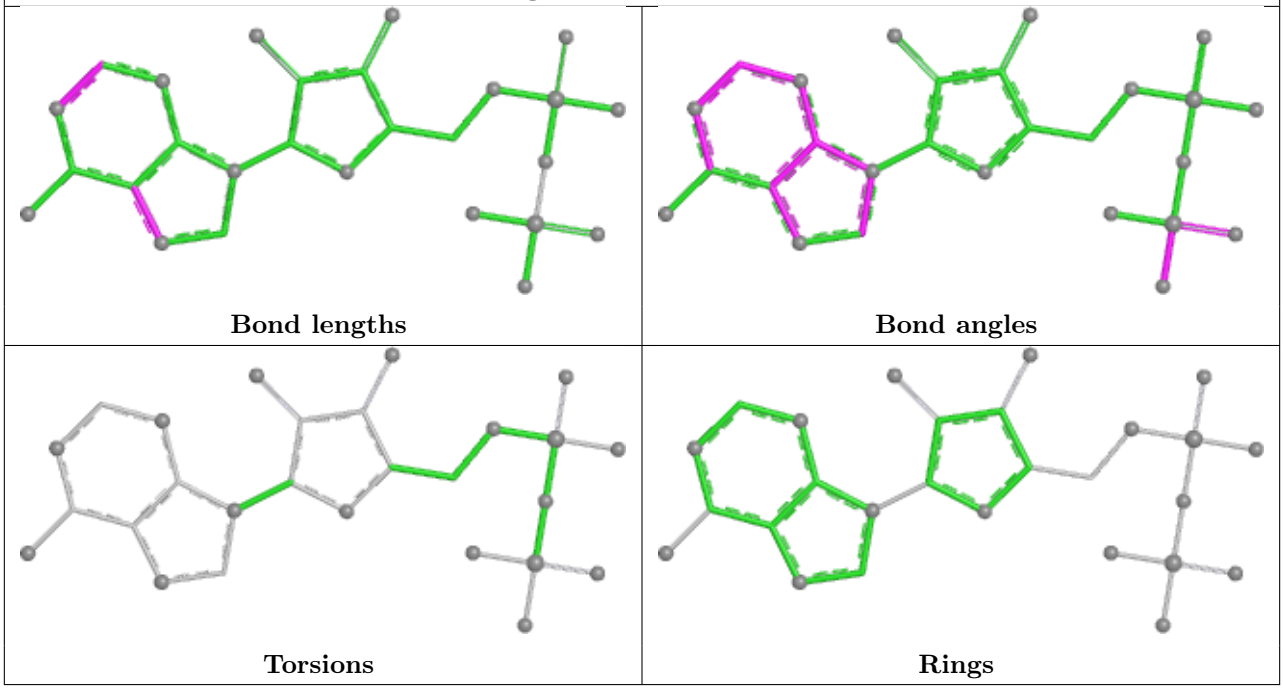
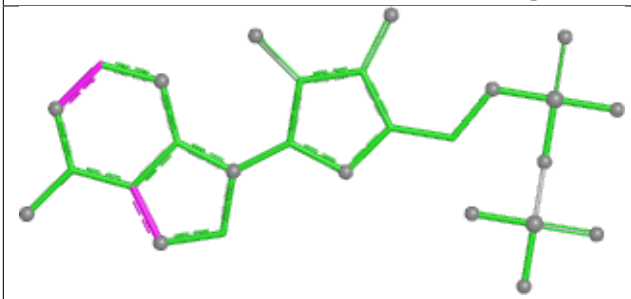
Mol	Chain	Res	Type	Clashes	Symm-Clashes
2	P	401	ADP	4	0
2	J	401	ADP	4	0
2	B	401	ADP	4	0
2	A	401	ADP	4	0
2	F	401	ADP	5	0
2	O	401	ADP	5	0
2	H	401	ADP	4	0
2	D	401	ADP	4	0

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 then 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.

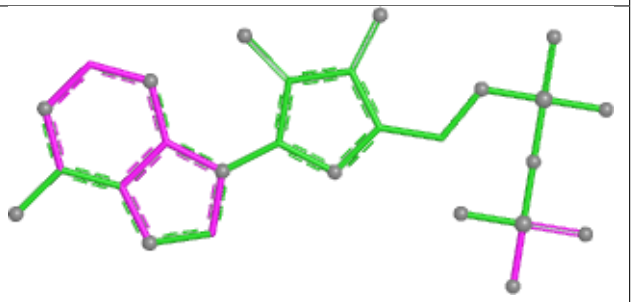
Ligand ADP E 401



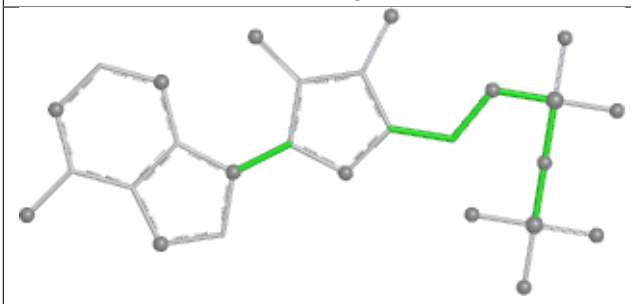
Ligand ADP S 401



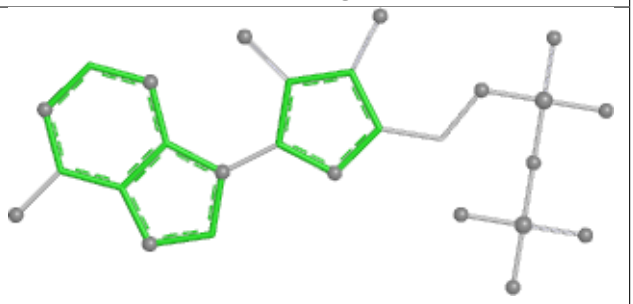
Bond lengths



Bond angles

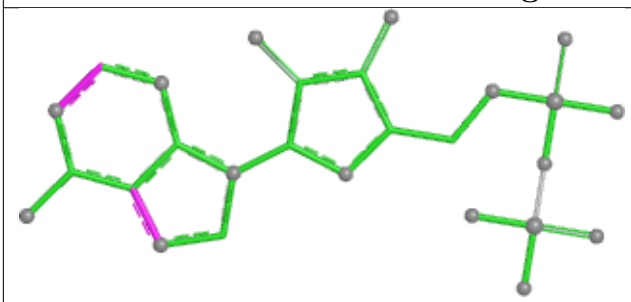


Torsions

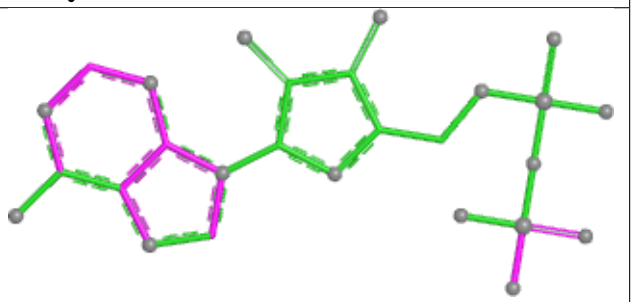


Rings

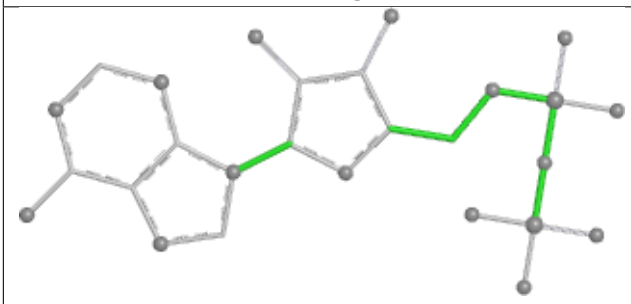
Ligand ADP Q 401



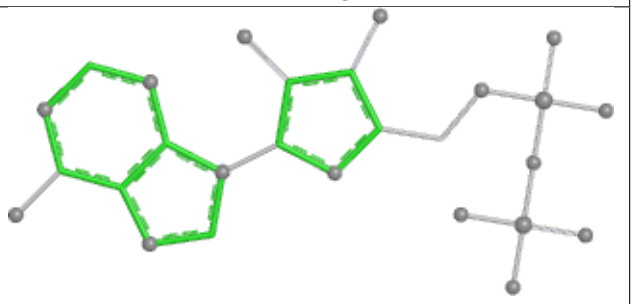
Bond lengths



Bond angles

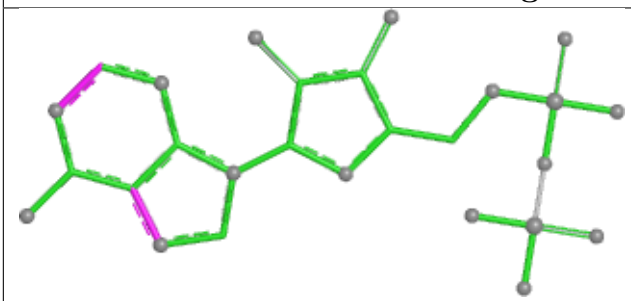


Torsions

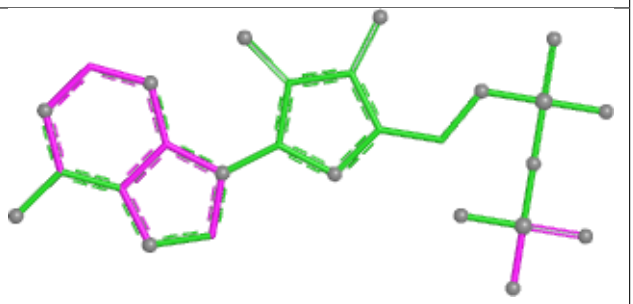


Rings

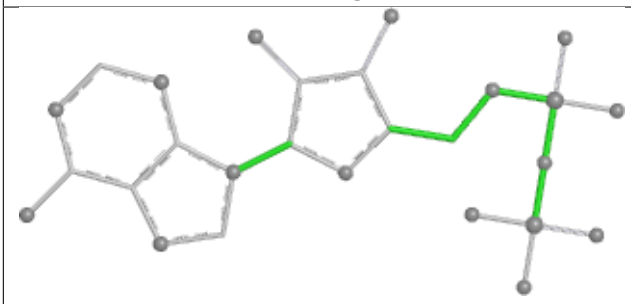
Ligand ADP U 401



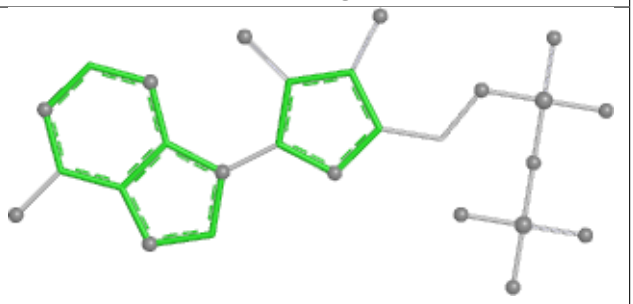
Bond lengths



Bond angles

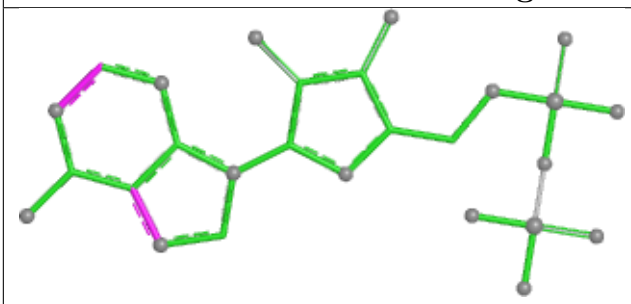


Torsions

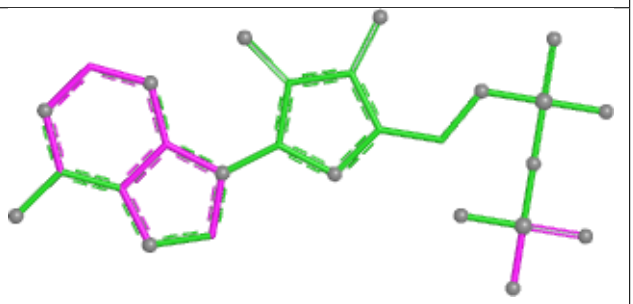


Rings

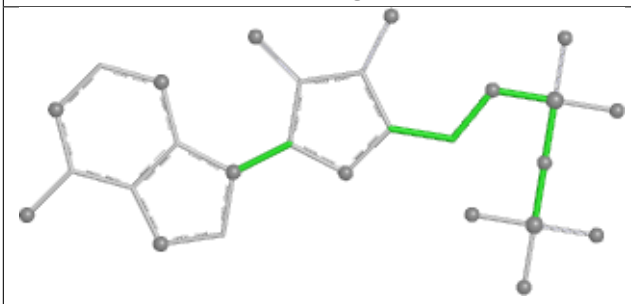
Ligand ADP W 401



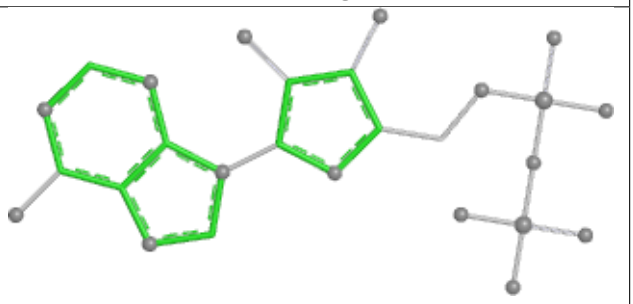
Bond lengths



Bond angles



Torsions



Rings

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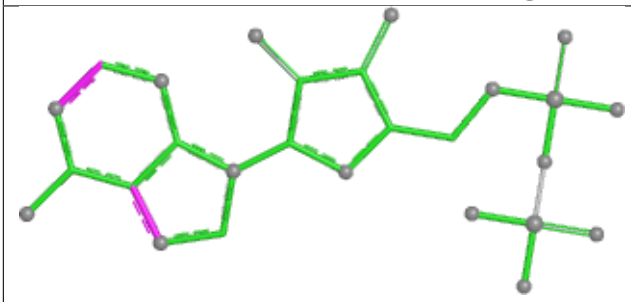
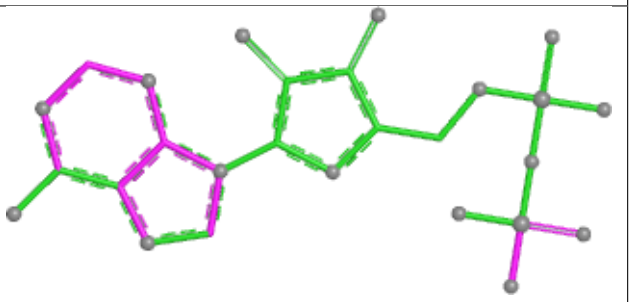
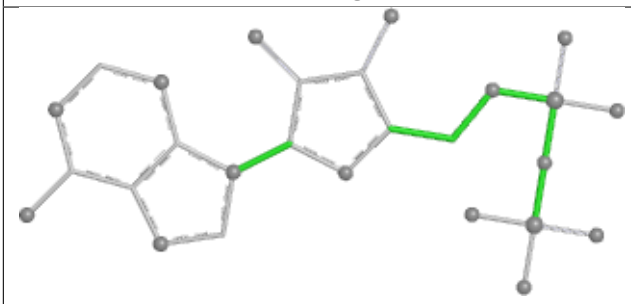
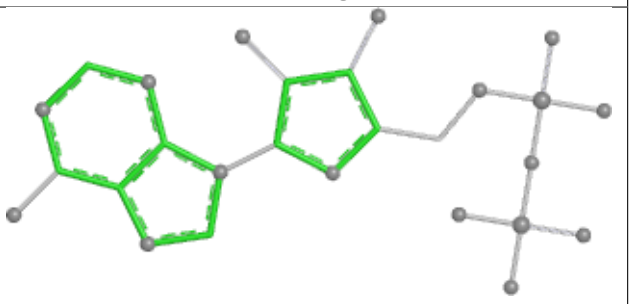
 Bond lengths	 Bond angles
 Torsions	 Rings

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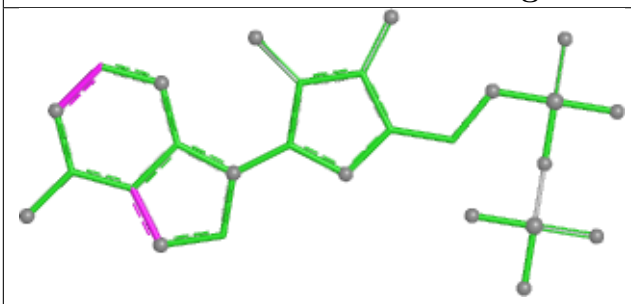
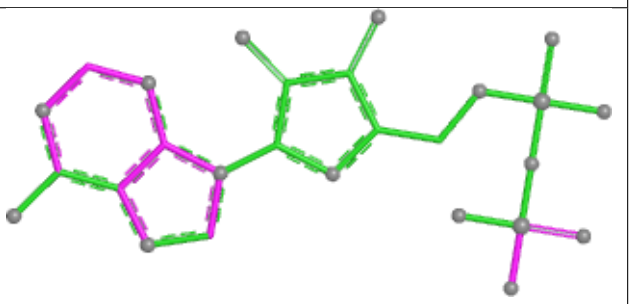
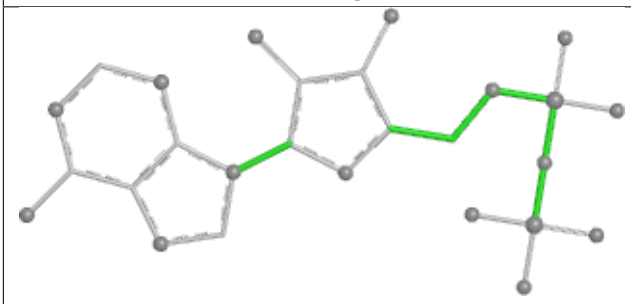
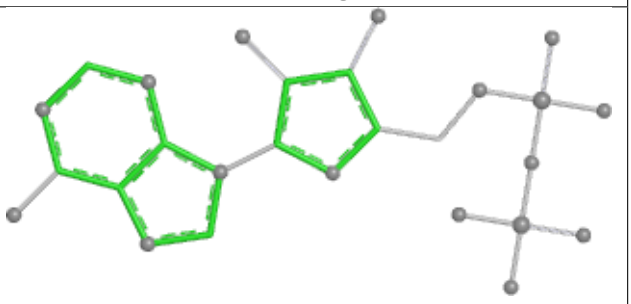
 Bond lengths	 Bond angles
 Torsions	 Rings

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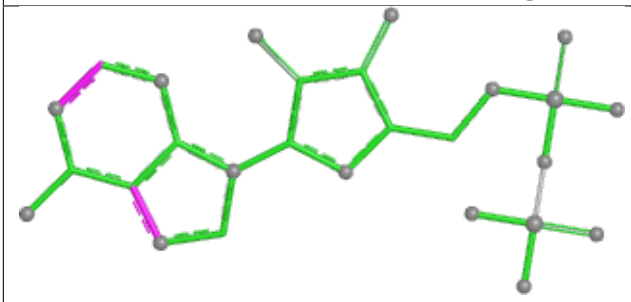
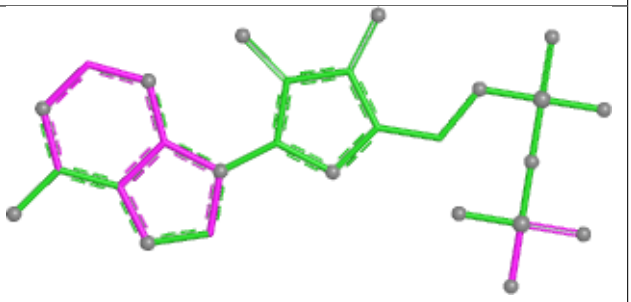
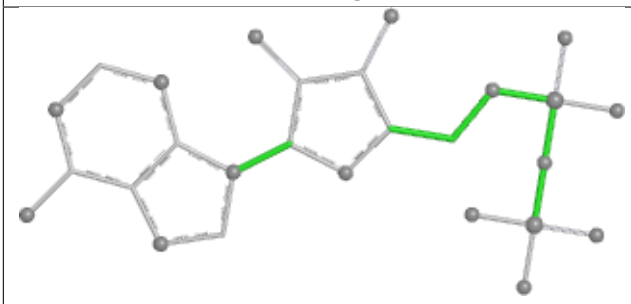
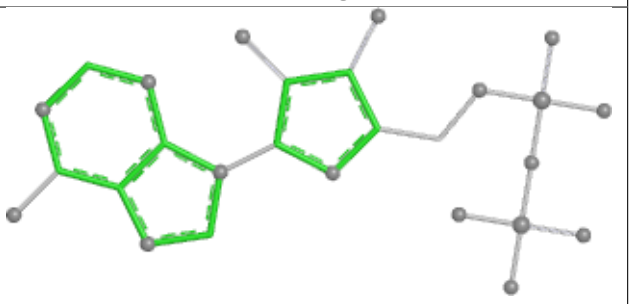
 Bond lengths	 Bond angles
 Torsions	 Rings

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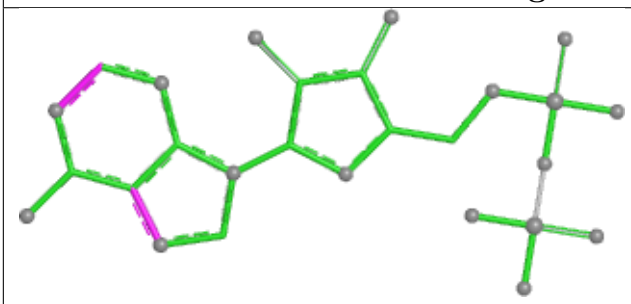
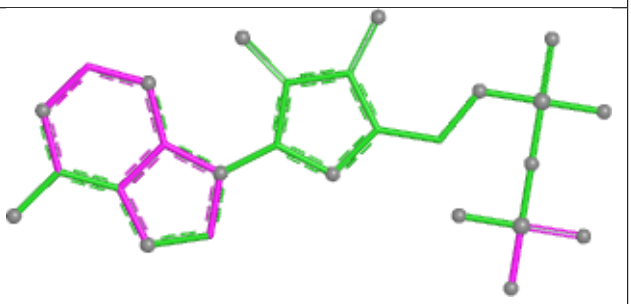
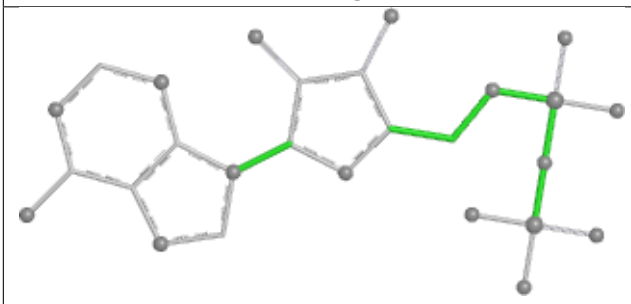
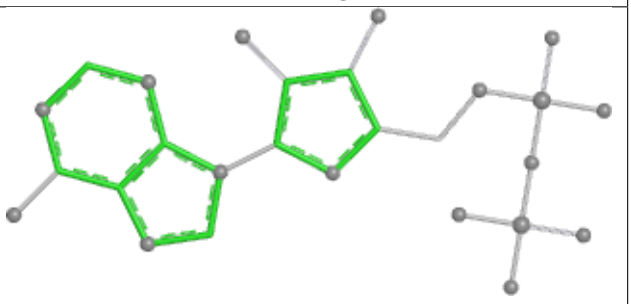
 Bond lengths	 Bond angles
 Torsions	 Rings

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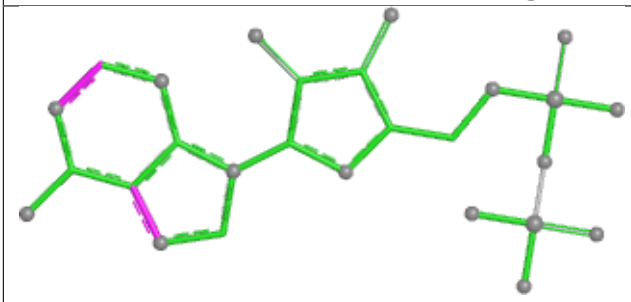
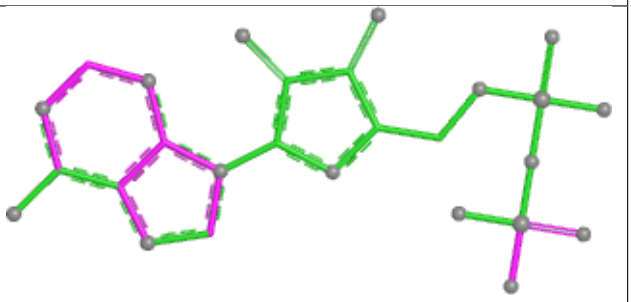
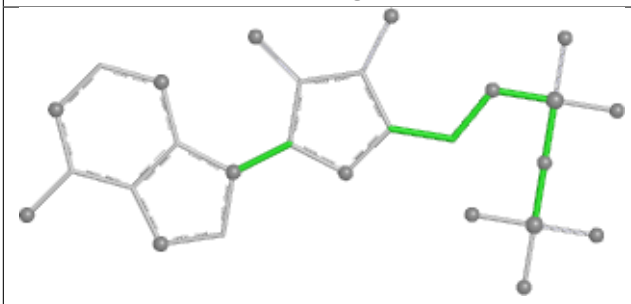
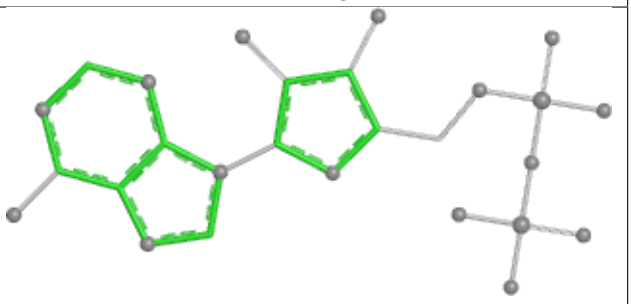
 Bond lengths	 Bond angles
 Torsions	 Rings

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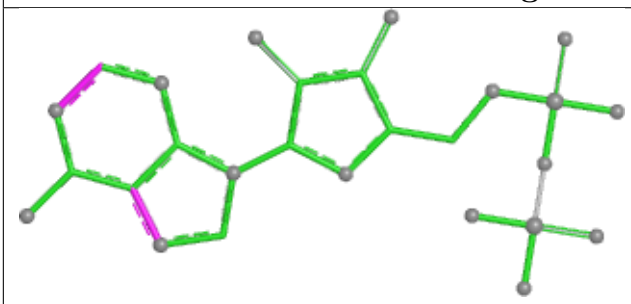
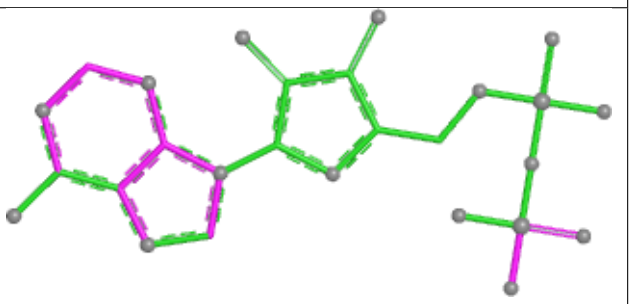
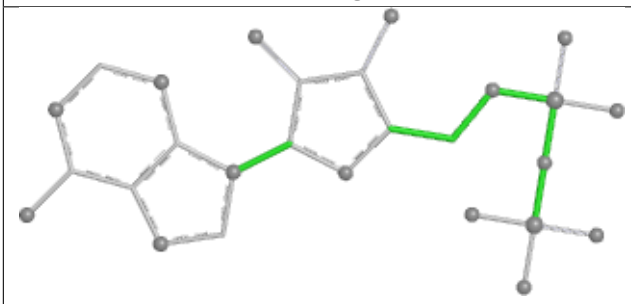
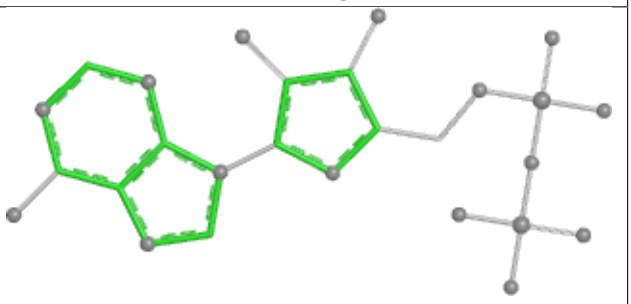
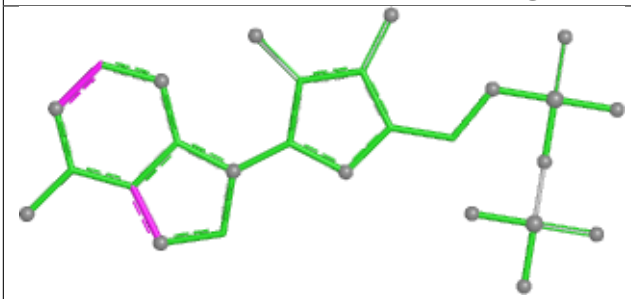
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand ADP L 401



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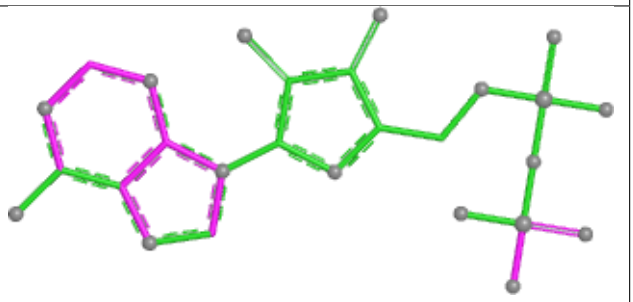

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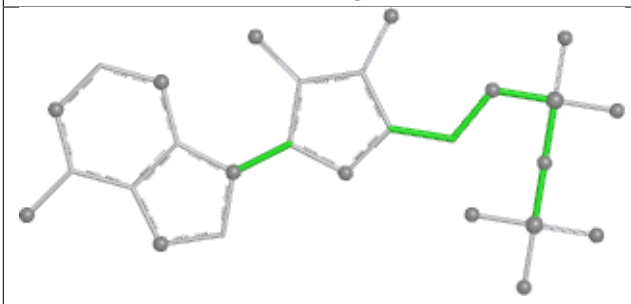

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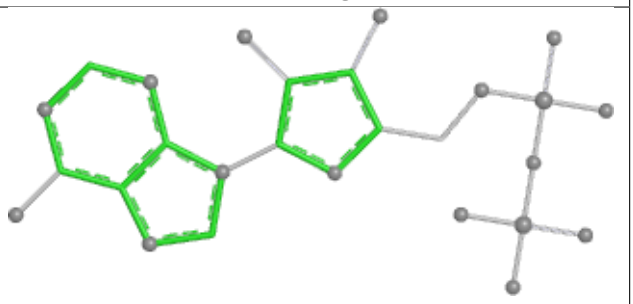
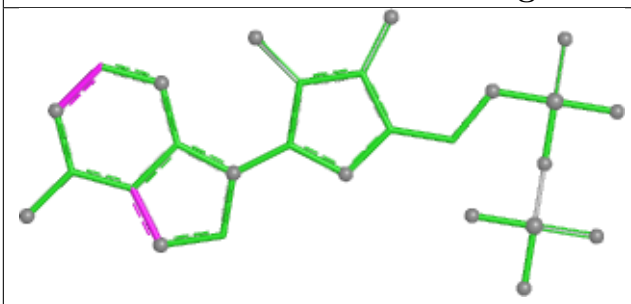


Rings

Ligand ADP M 401



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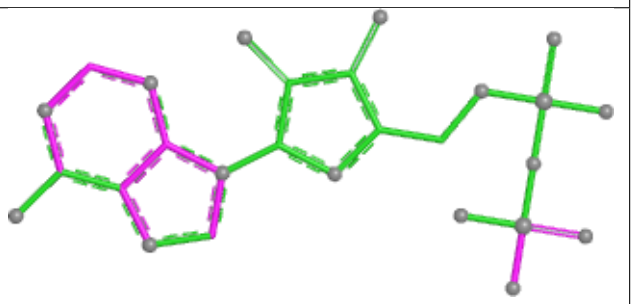
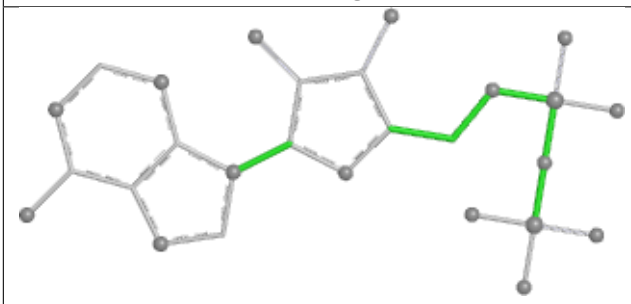


Bond angles



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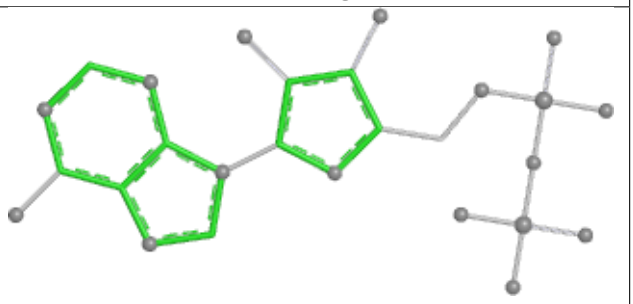

Rings

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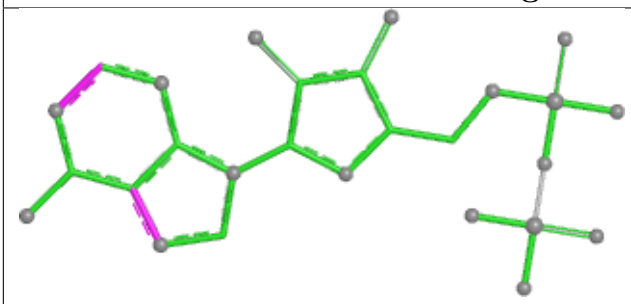

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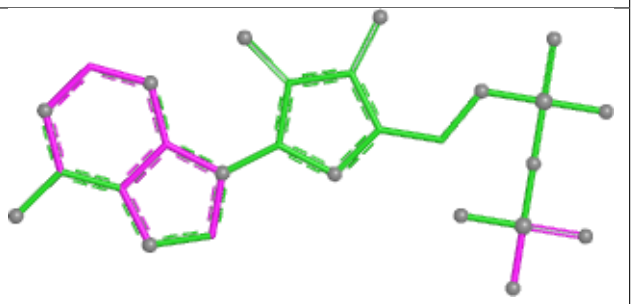

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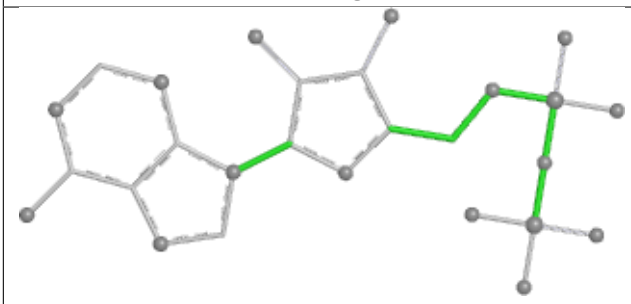

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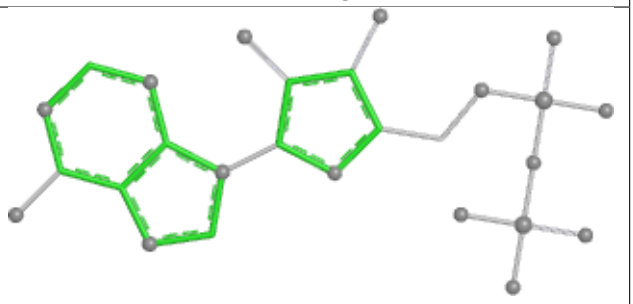

Rings

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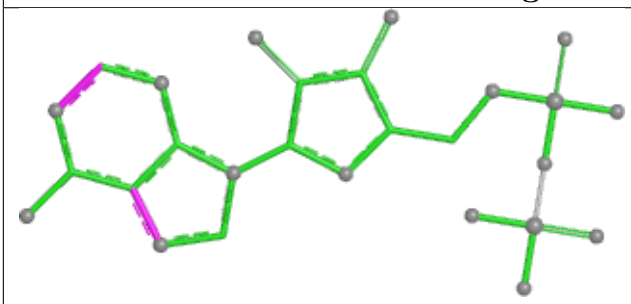

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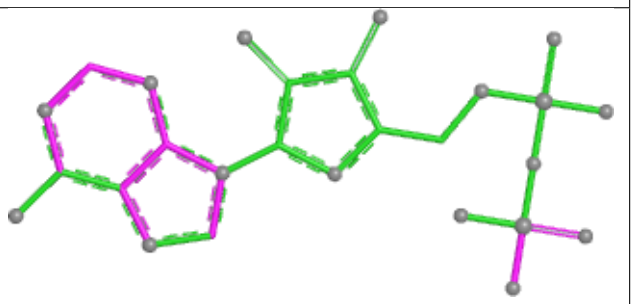

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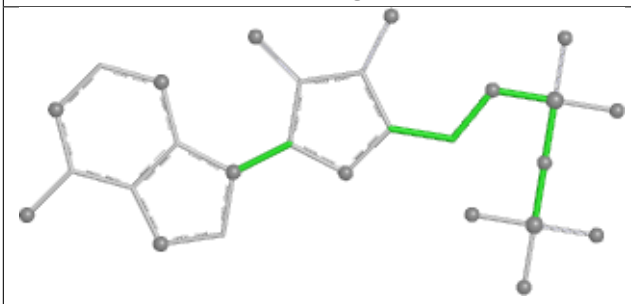

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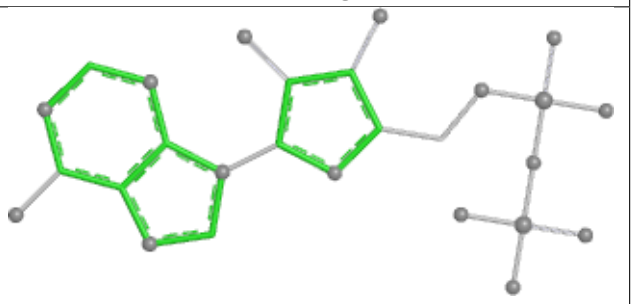

Rings

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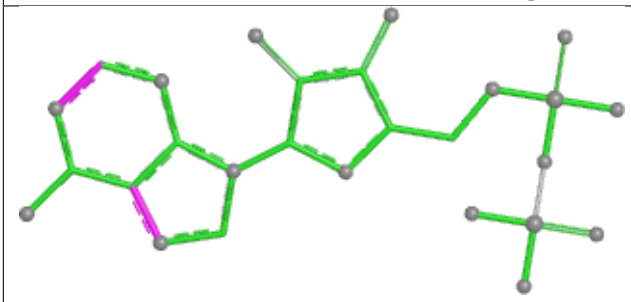
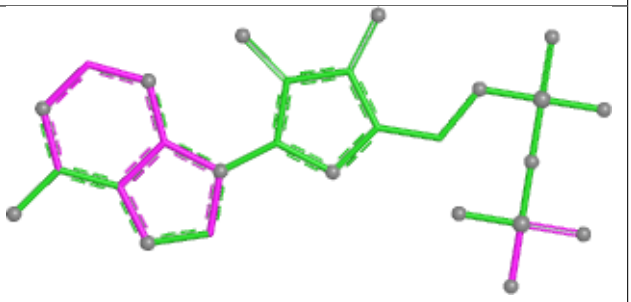
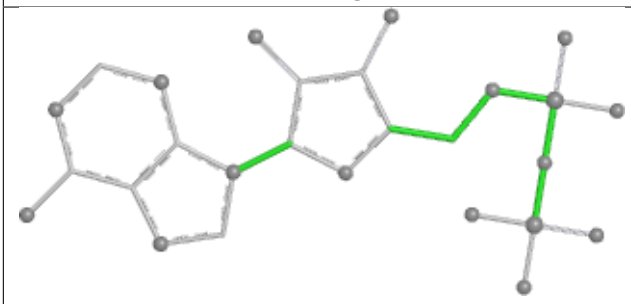
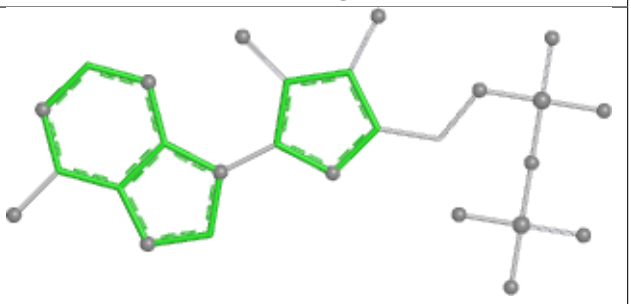
 Bond lengths	 Bond angles
 Torsions	 Rings

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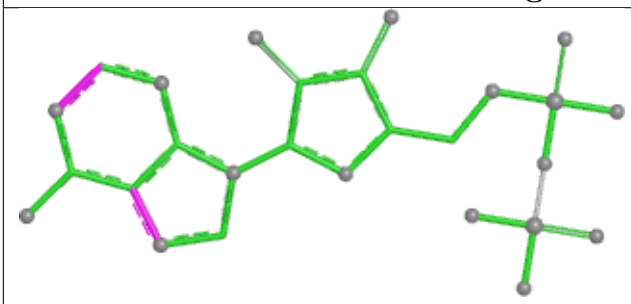
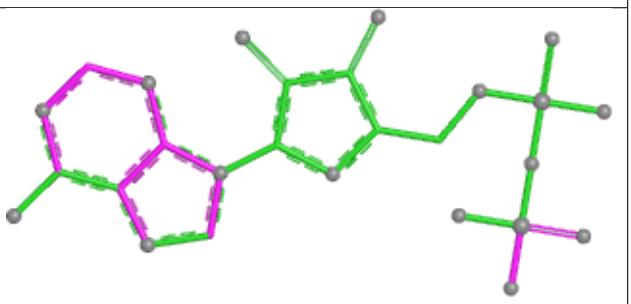
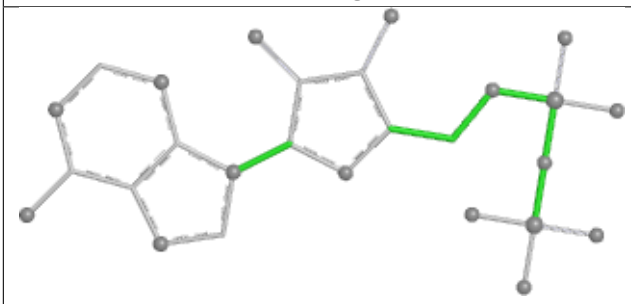
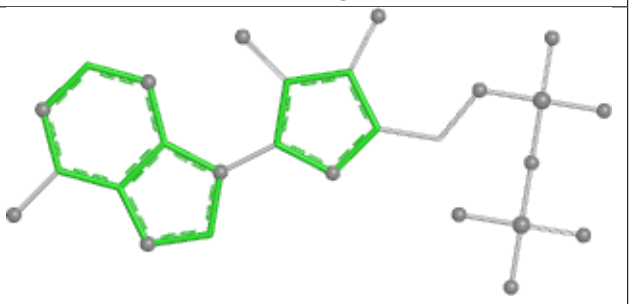
 Bond lengths	 Bond angles
 Torsions	 Rings

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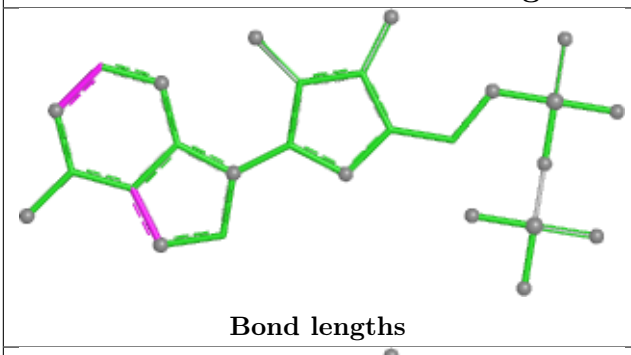
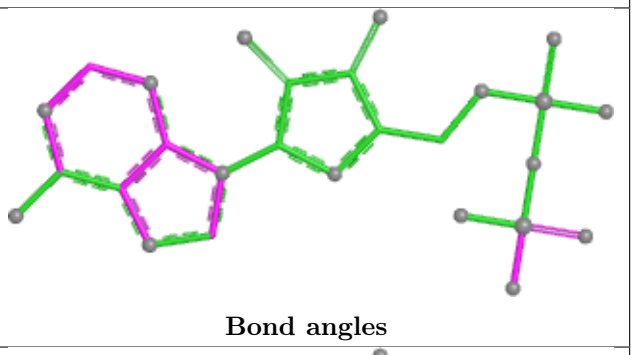
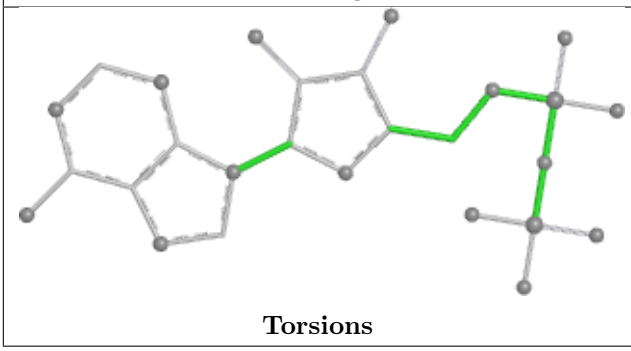


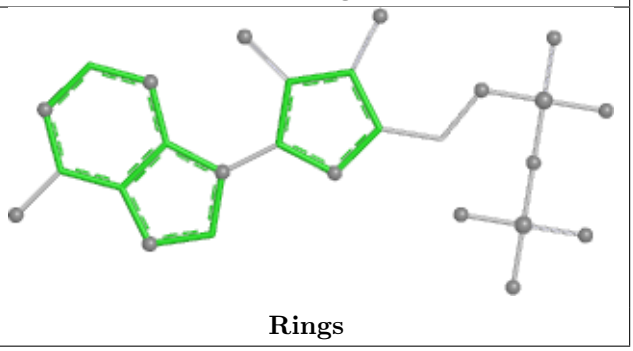
Bond lengths



Bond angles

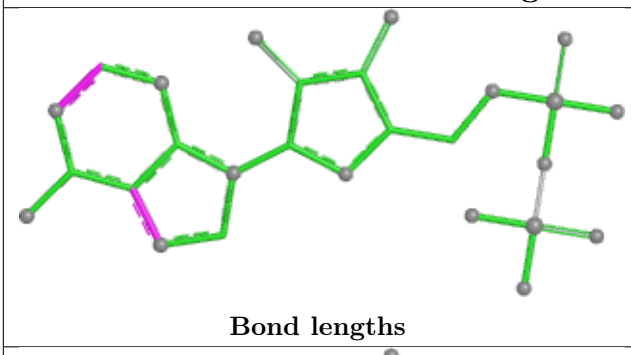


Torsions

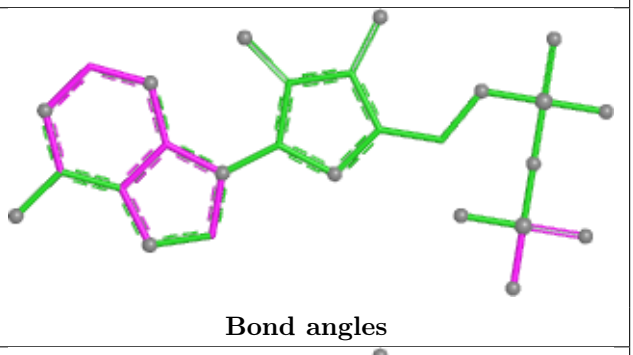


Rings

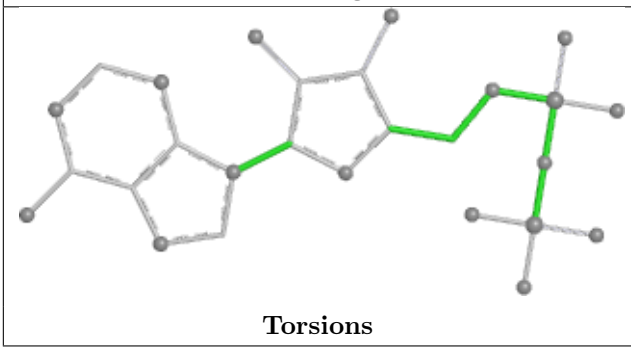
Ligand ADP A 401



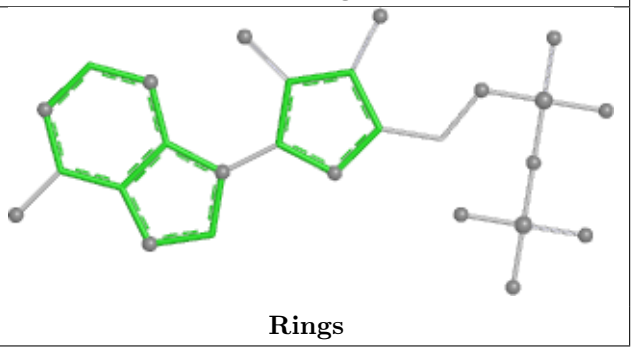
Bond lengths



Bond angles

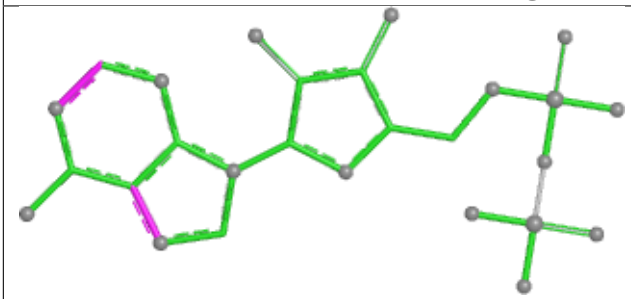


Torsions

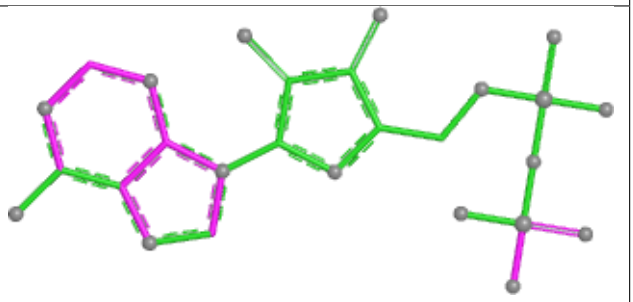


Rings

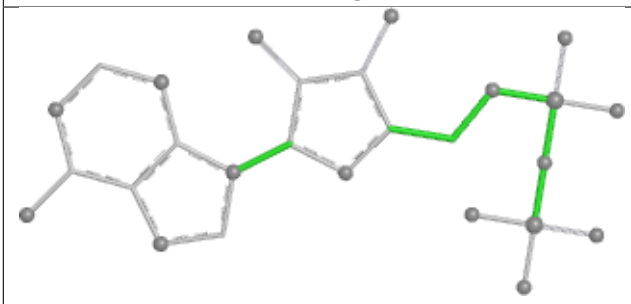
Ligand ADP F 401



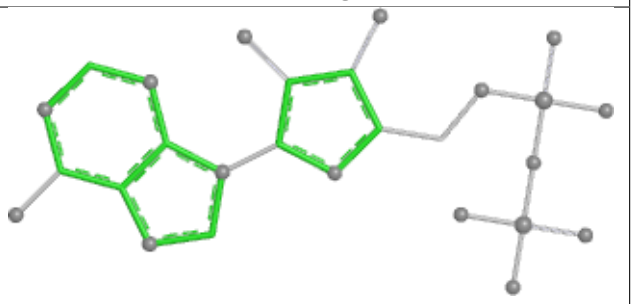
Bond lengths



Bond angles

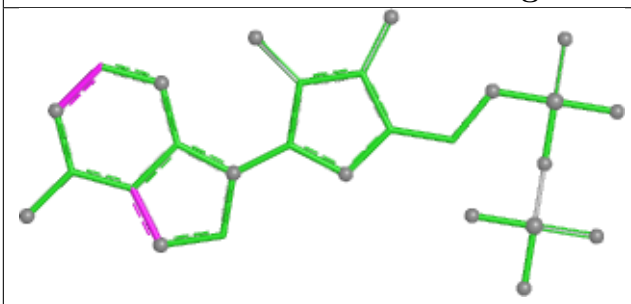


Torsions

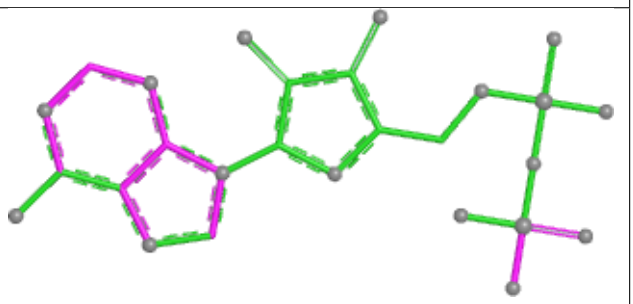


Rings

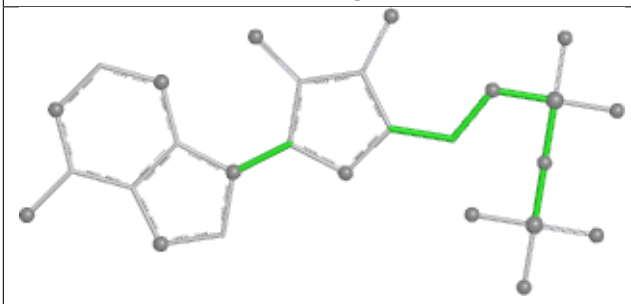
Ligand ADP O 401



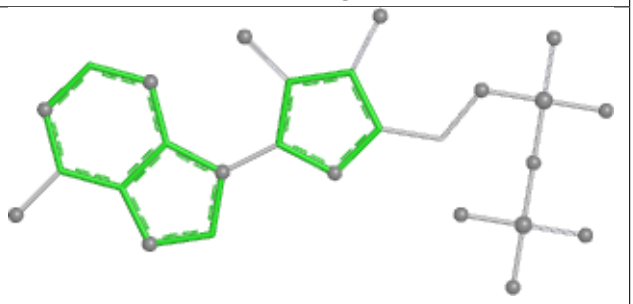
Bond lengths



Bond angles

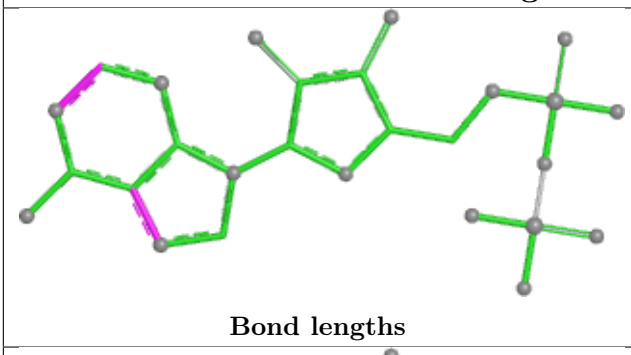


Torsions

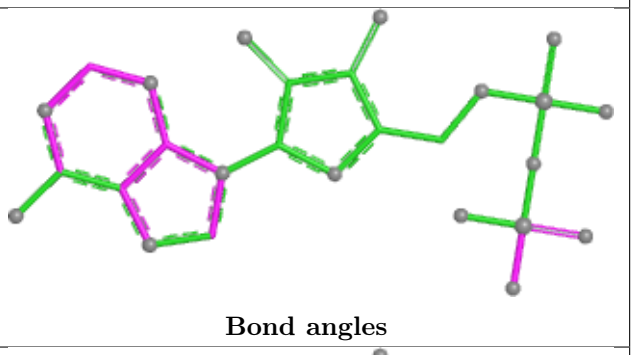


Rings

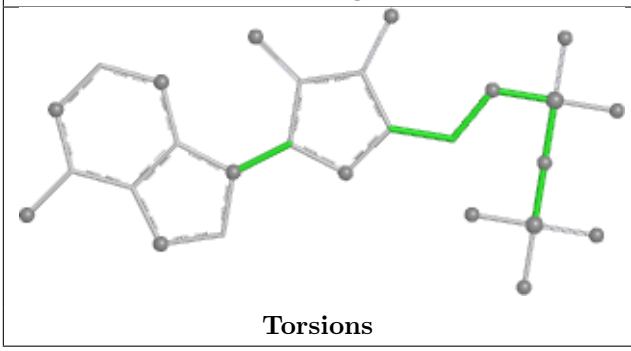
Ligand ADP H 401



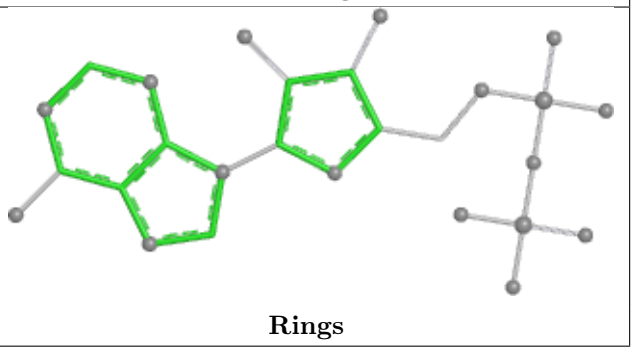
Bond lengths



Bond angles

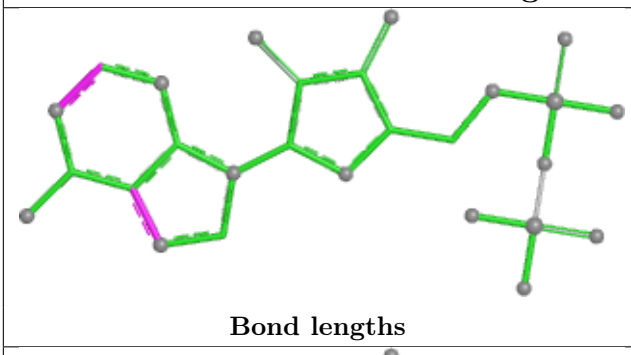


Torsions

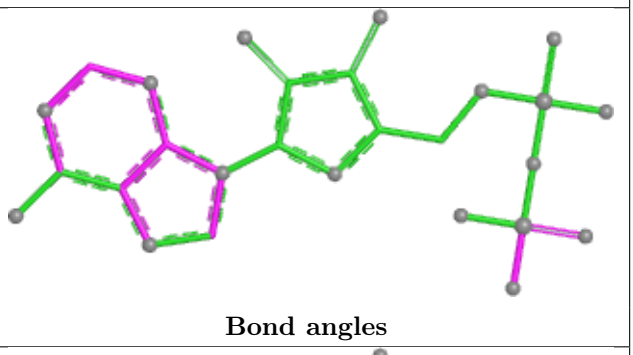


Rings

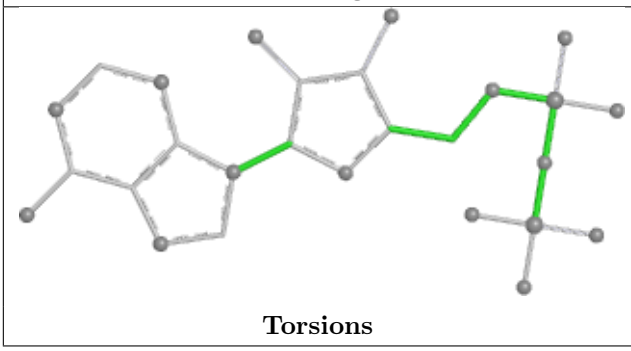
Ligand ADP D 401



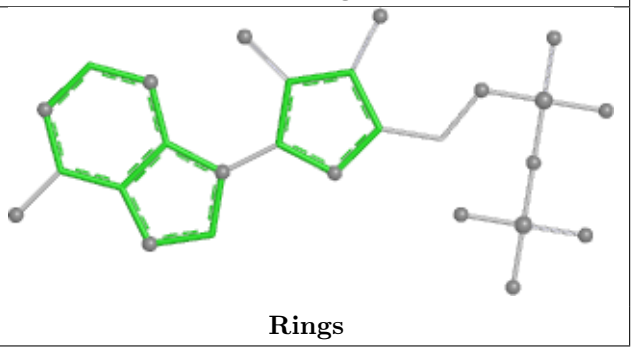
Bond lengths



Bond angles



Torsions



Rings

5.7 Other polymers

There are no such residues in this entry.

5.8 Polymer linkage issues (i)

There are no chain breaks in this entry.

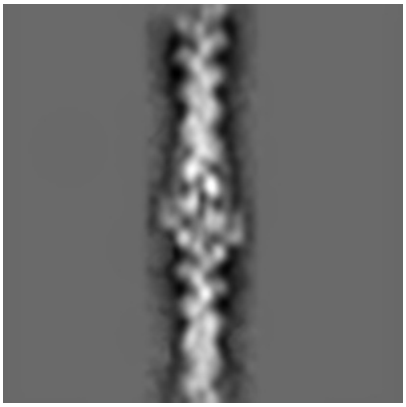
6 Map visualisation

This section contains visualisations of the EMDB entry EMD-3578. These allow visual inspection of the internal detail of the map and identification of artifacts.

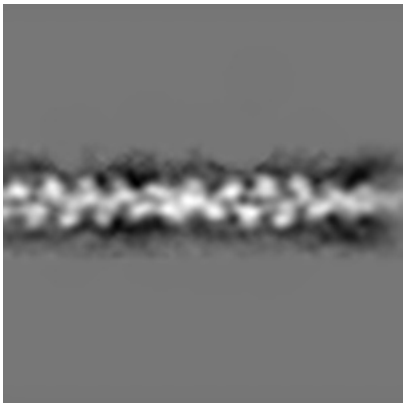
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections

6.1.1 Primary map



X



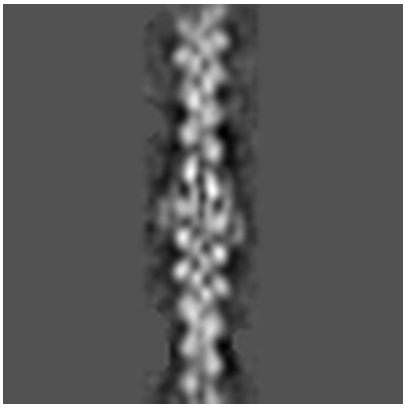
Y

Z

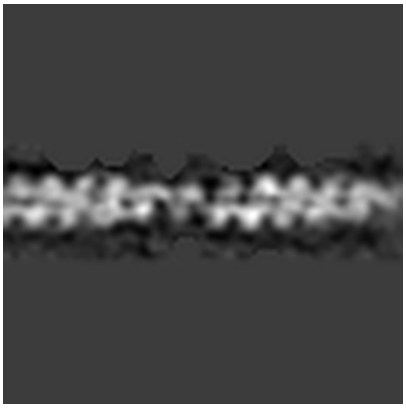
The images above show the map projected in three orthogonal directions.

6.2 Central slices

6.2.1 Primary map



X Index: 48



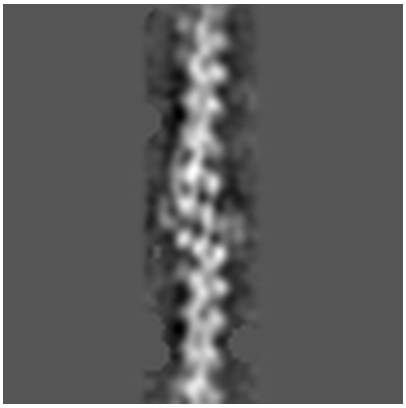
Y Index: 48

Z Index: 48

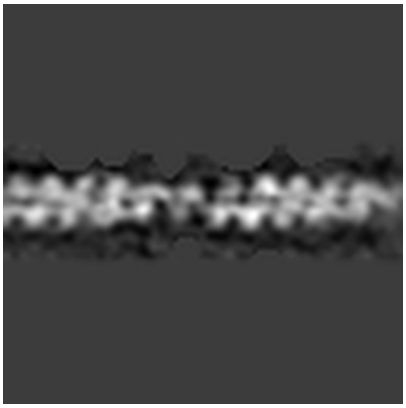
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices (i)

6.3.1 Primary map



X Index: 50



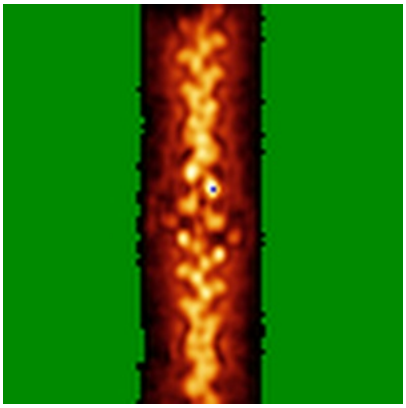
Y Index: 48

Z Index: 27

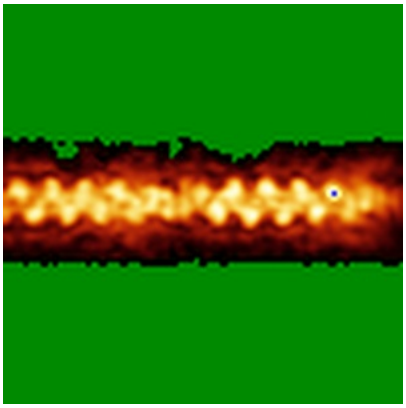
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) (i)

6.4.1 Primary map



X



Y

Z

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

6.5 Orthogonal surface views (i)

This section was not generated.

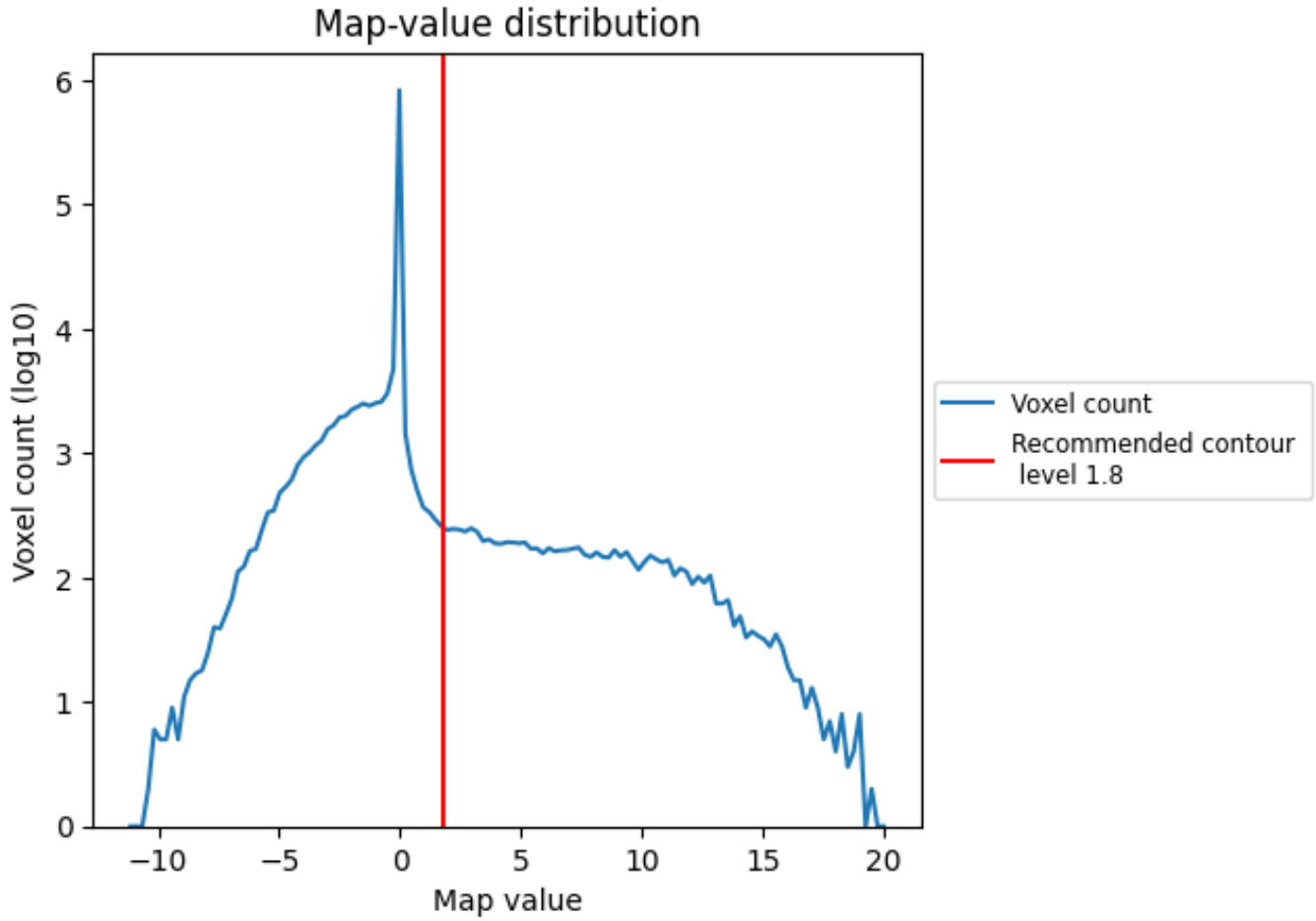
6.6 Mask visualisation (i)

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

7 Map analysis

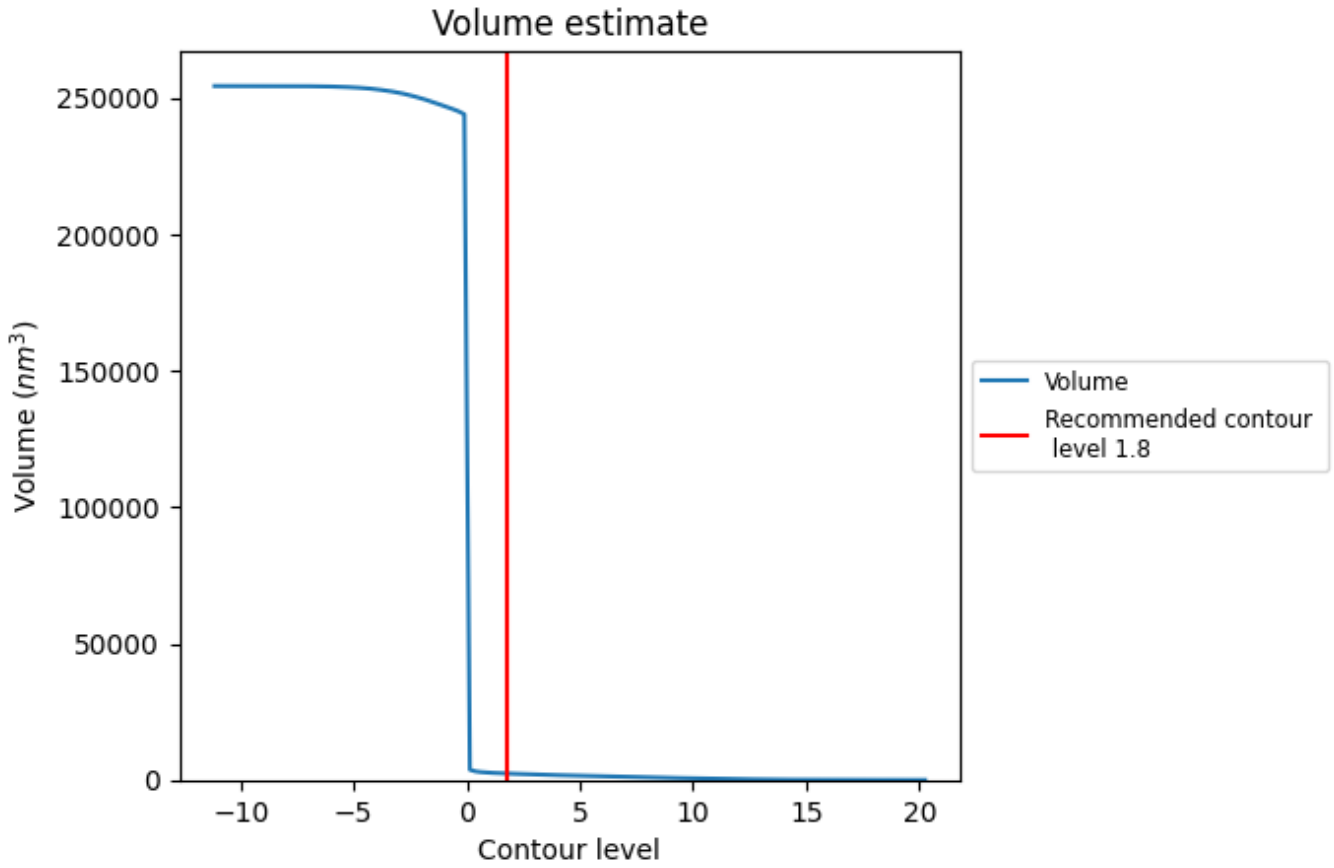
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution



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.

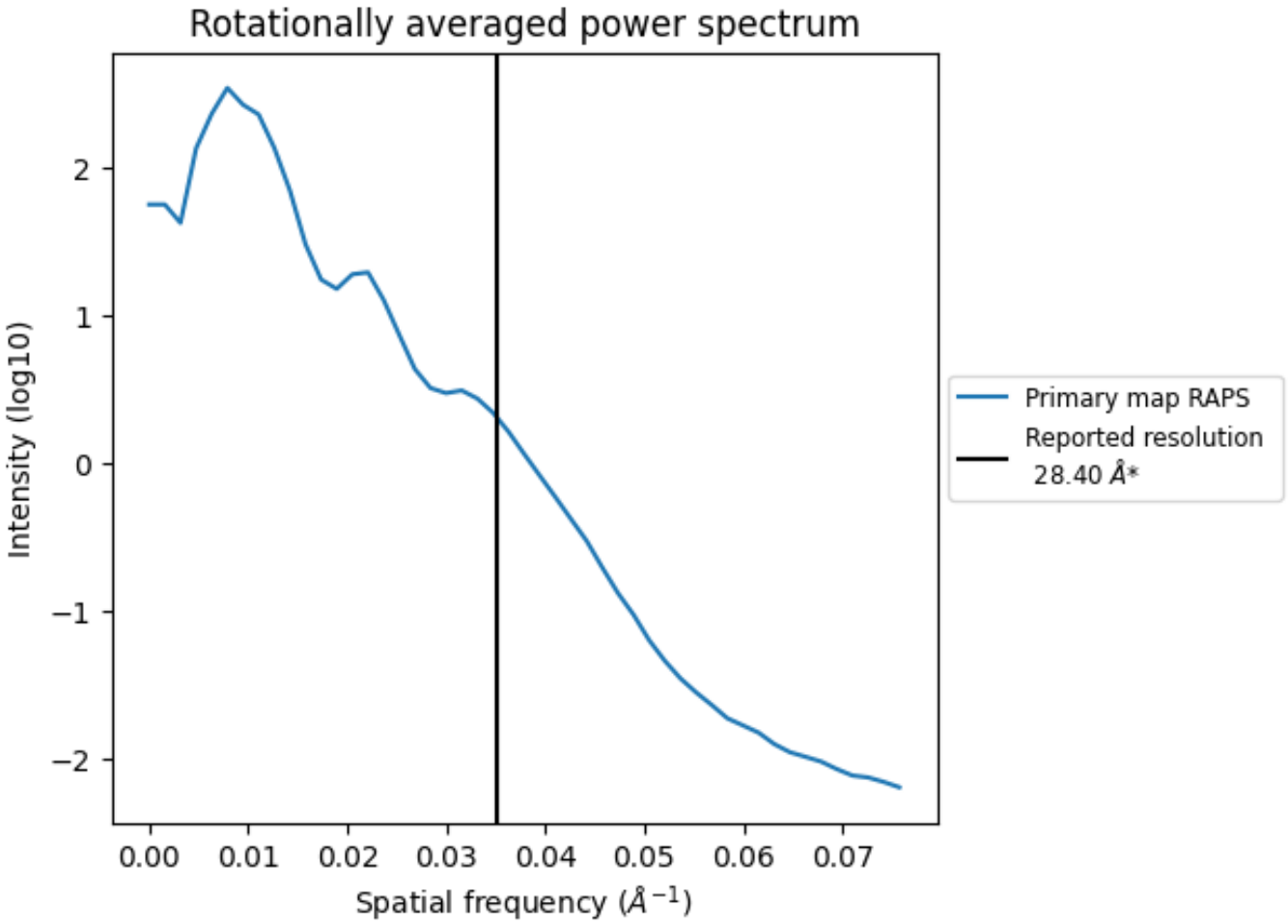
7.2 Volume estimate



The volume at the recommended contour level is 2359 nm^3; this corresponds to an approximate mass of 2131 kDa.

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

7.3 Rotationally averaged power spectrum (i)



*Reported resolution corresponds to spatial frequency of 0.035 Å^{-1}

8 Fourier-Shell correlation

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

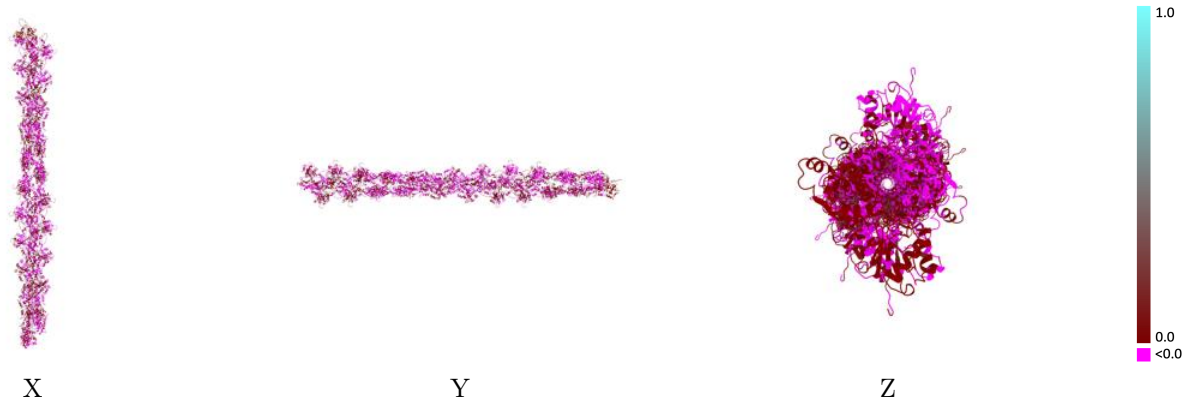
9 Map-model fit

This section contains information regarding the fit between EMDB map EMD-3578 and PDB model 5MVY. Per-residue inclusion information can be found in section 3 on page 7.

9.1 Map-model overlay

This section was not generated.

9.2 Q-score mapped to coordinate model

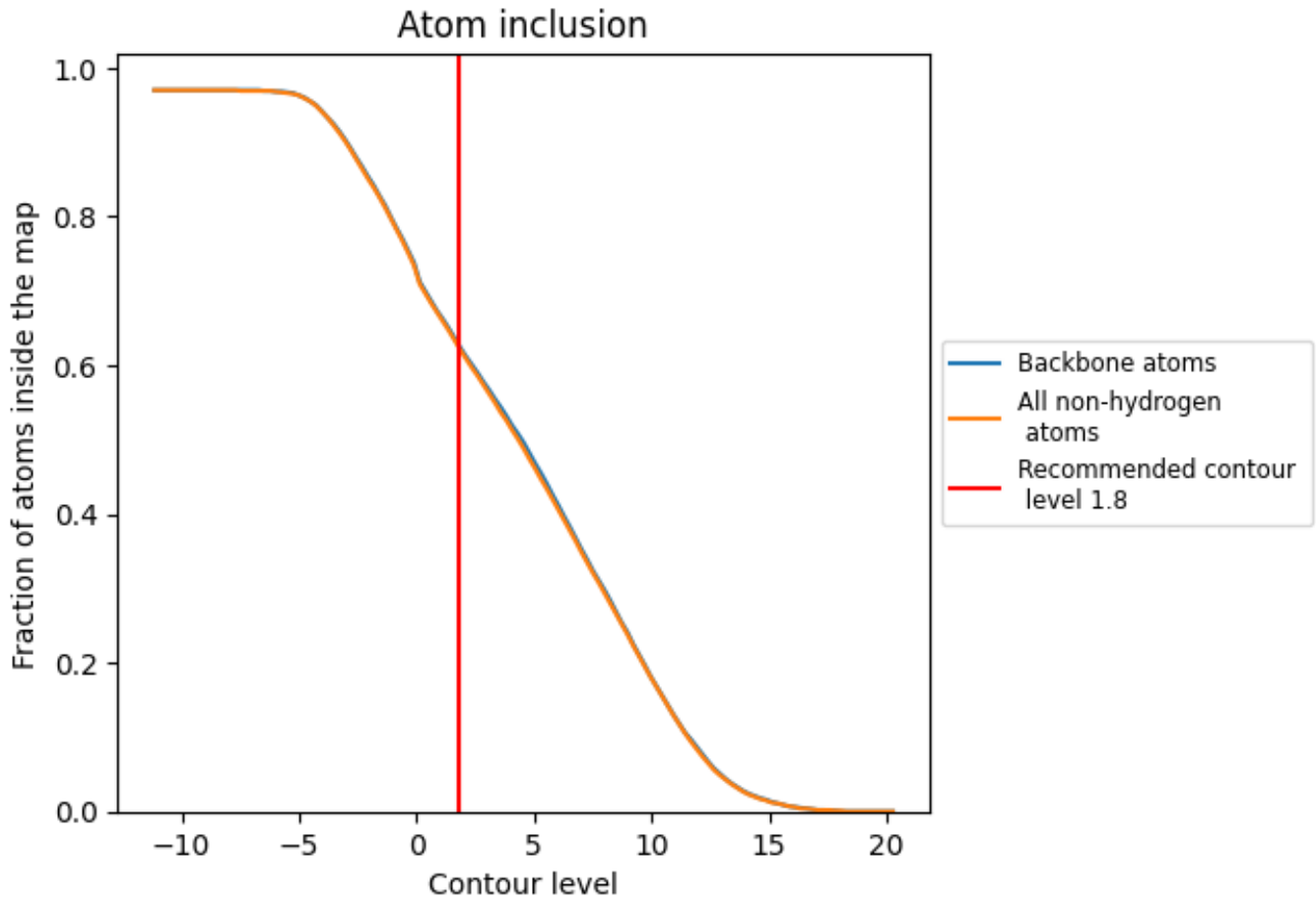


The images above show the model with each residue coloured according its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

This section was not generated.

9.4 Atom inclusion (i)



At the recommended contour level, 63% of all backbone atoms, 62% 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 (1.8) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.6240	0.0030
A	0.2000	-0.0060
B	0.2890	-0.0190
C	0.8970	0.0240
D	0.4420	-0.0100
E	0.9360	0.0230
F	0.5210	-0.0000
G	0.7920	0.0050
H	0.6140	0.0040
I	0.6360	0.0150
J	0.6210	-0.0000
K	0.4950	-0.0010
L	0.6010	-0.0240
M	0.5040	-0.0030
N	0.6580	-0.0170
O	0.4470	0.0040
P	0.8110	-0.0100
Q	0.5490	-0.0000
R	0.9060	0.0090
S	0.5970	0.0070
T	0.8150	0.0150
U	0.6530	0.0220
V	0.6700	0.0120
W	0.6930	0.0230