



Full wwPDB EM Validation Report ⓘ

Mar 6, 2026 – 08:44 PM UTC

PDB ID : 5WQ7 / pdb_00005wq7
EMDB ID : EMD-6675
Title : CryoEM structure of type II secretion system secretin GspD in E.coli K12
Authors : Yan, Z.; Yin, M.; Li, X.
Deposited on : 2016-11-23
Resolution : 3.04 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:

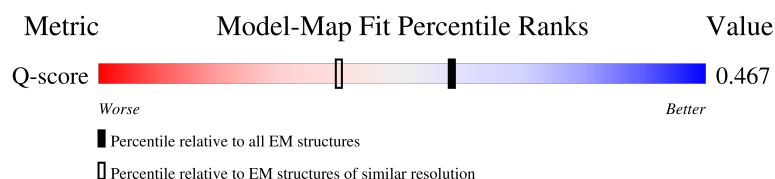
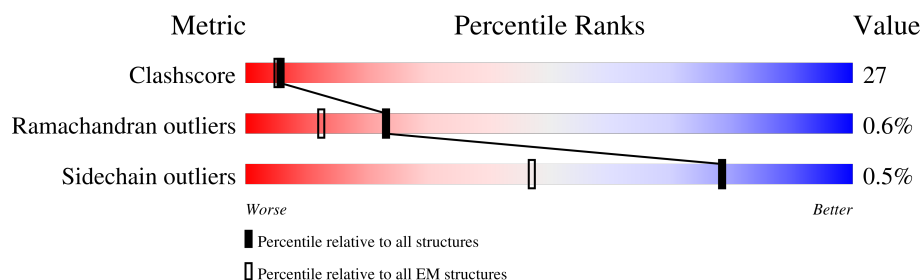
EMDB validation analysis : 0.0.1.dev132
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & 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 3.04 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	13952 (2.54 - 3.54)

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

Mol	Chain	Length	Quality of chain
1	A	627	12% 44% 30% 25%
1	B	627	12% 44% 30% 25%
1	C	627	11% 44% 31% 25%
1	D	627	12% 44% 30% 25%

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Mol	Chain	Length	Quality of chain
1	E	627	
1	F	627	
1	G	627	
1	H	627	
1	I	627	
1	J	627	
1	K	627	
1	L	627	
1	M	627	
1	N	627	
1	O	627	

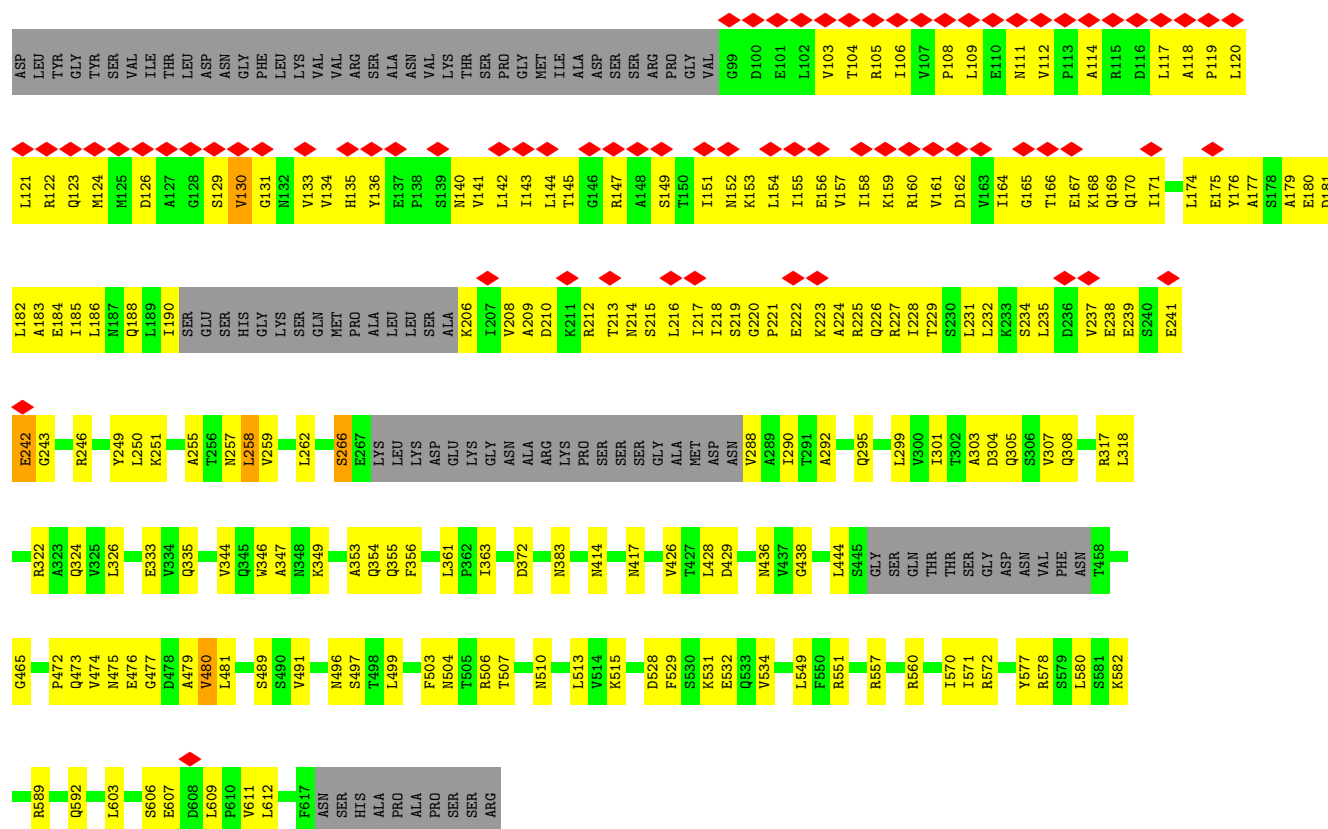
2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 54480 atoms, of which 0 are hydrogens and 0 are deuteriums.

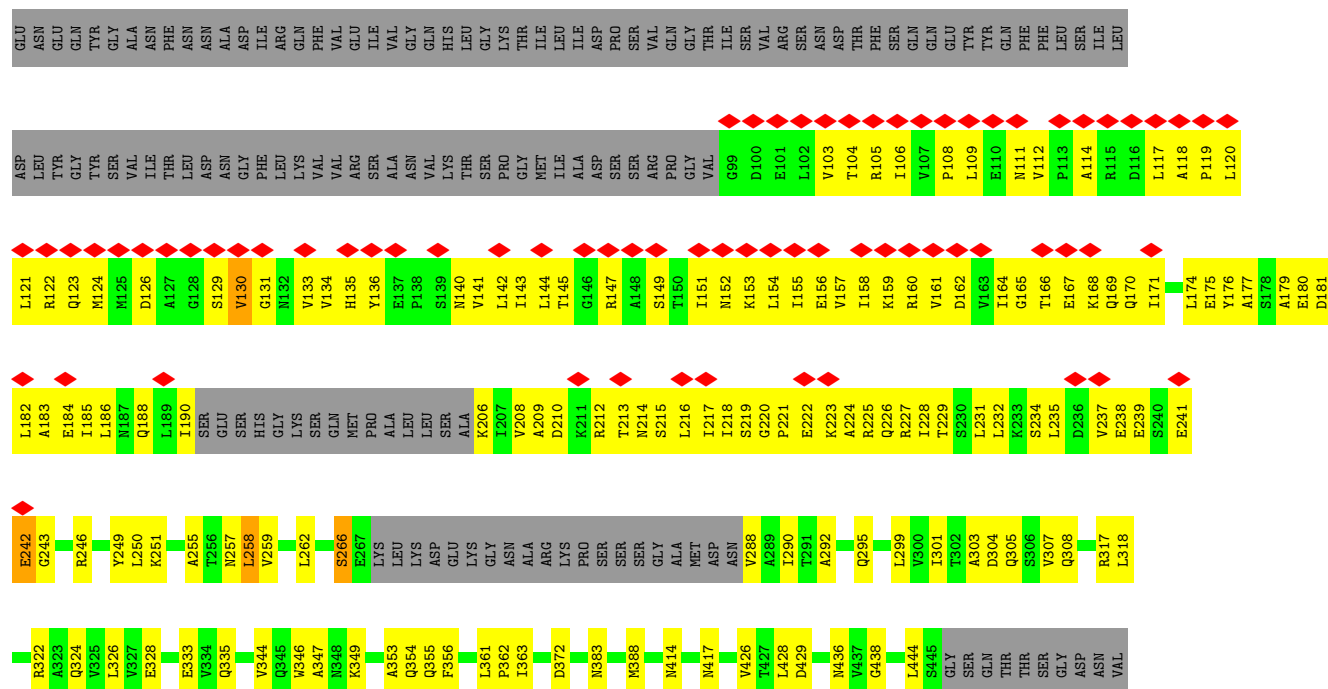
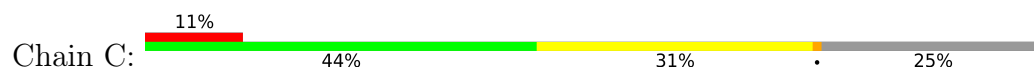
In the tables below, the AltConf column contains the number of residues with at least one atom in alternate conformation and the Trace column contains the number of residues modelled with at most 2 atoms.

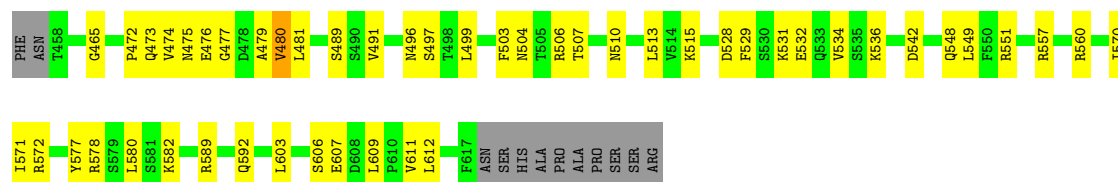
- Molecule 1 is a protein called Putative type II secretion system protein D.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	B	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	C	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	D	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	E	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	F	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	G	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	H	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	I	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	J	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	K	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	L	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	M	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	N	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		
1	O	472	Total	C	N	O	S	0	0
			3632	2279	628	720	5		

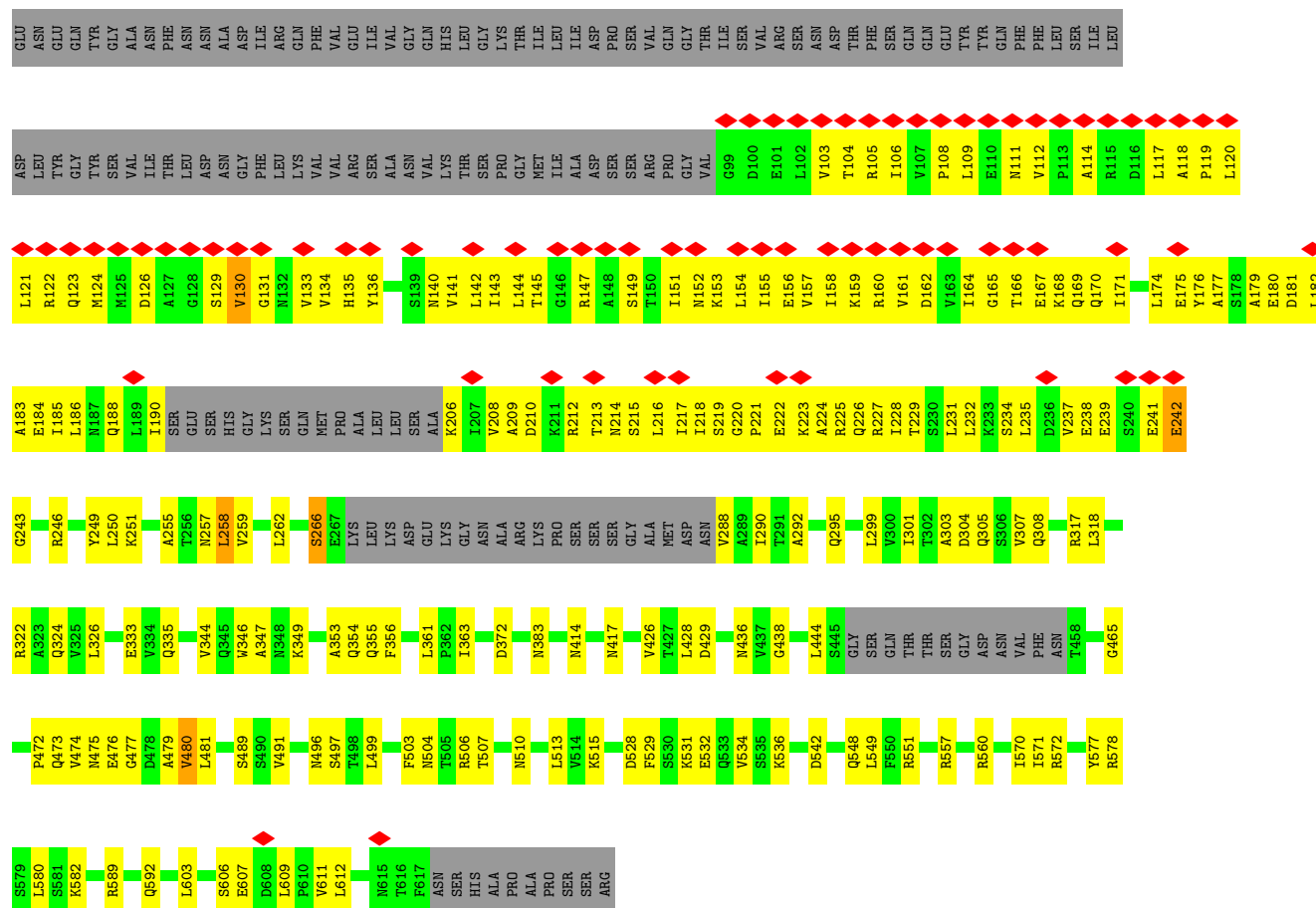
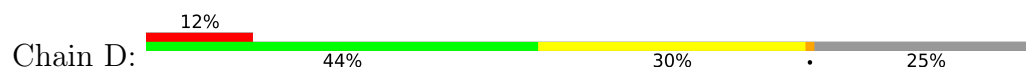


• Molecule 1: Putative type II secretion system protein D

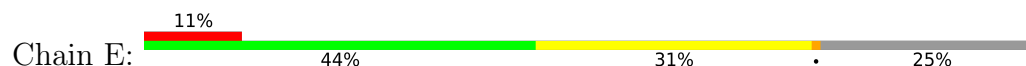


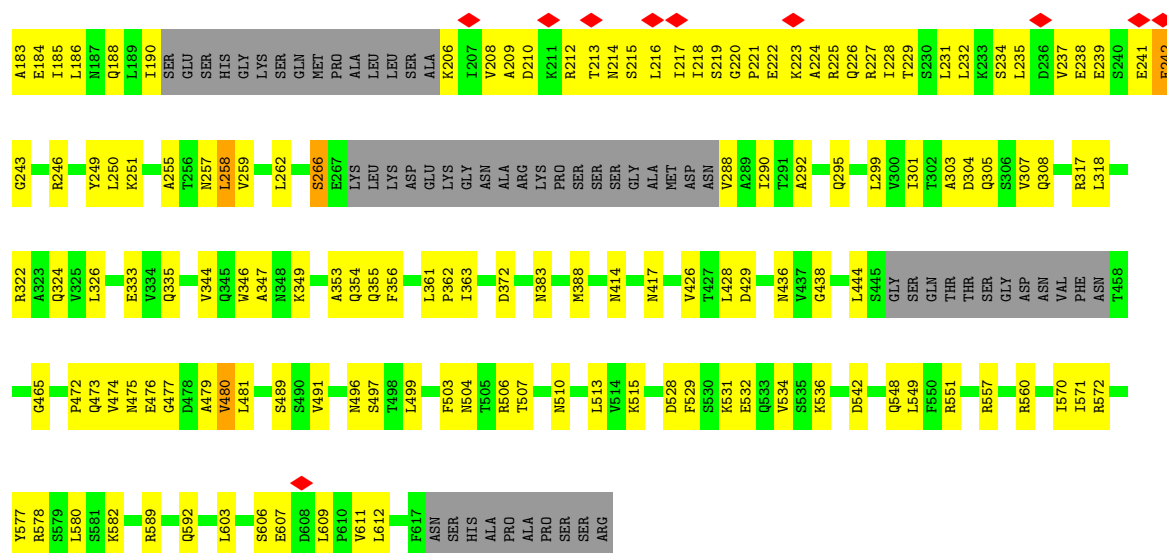


• Molecule 1: Putative type II secretion system protein D

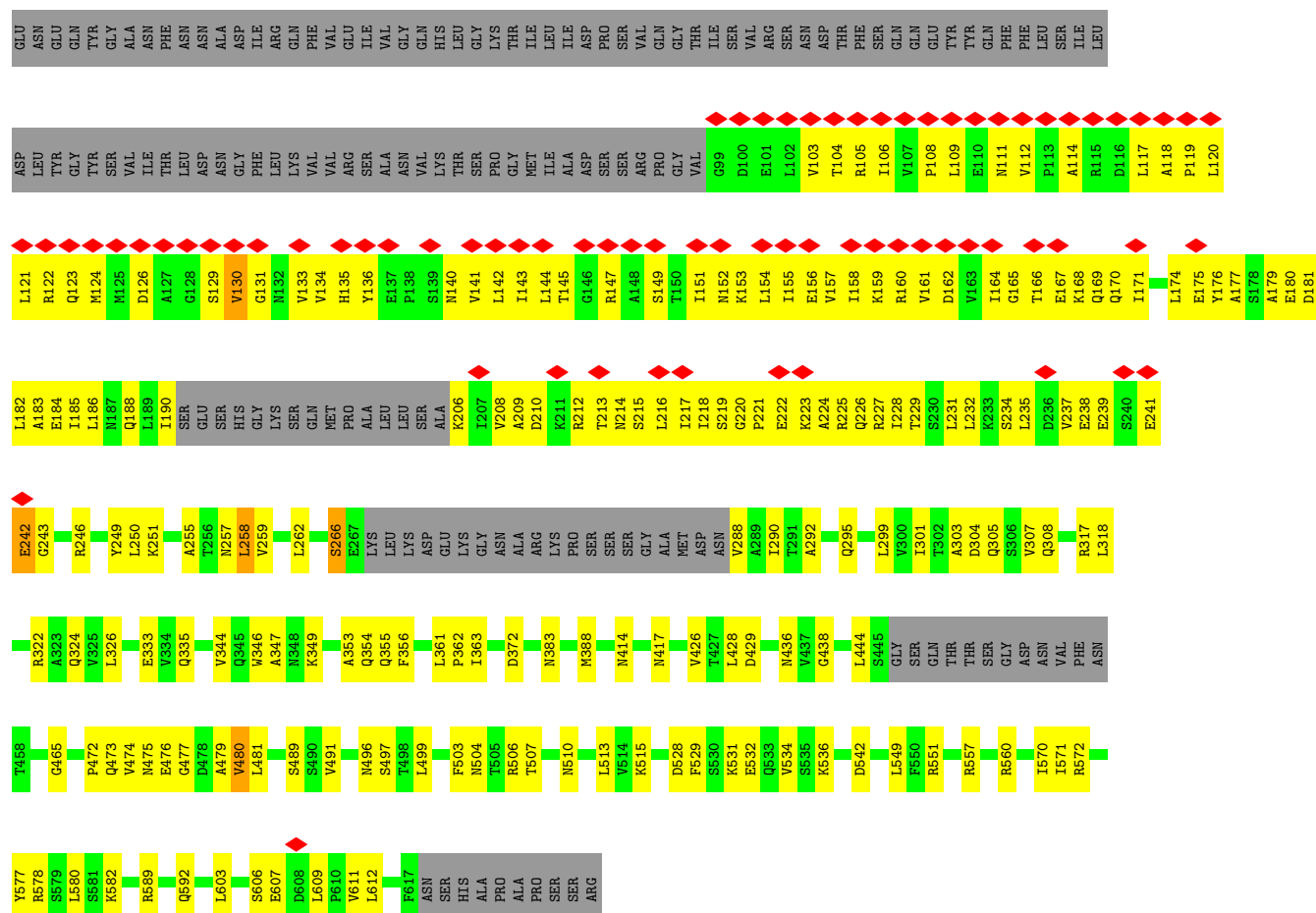


• Molecule 1: Putative type II secretion system protein D

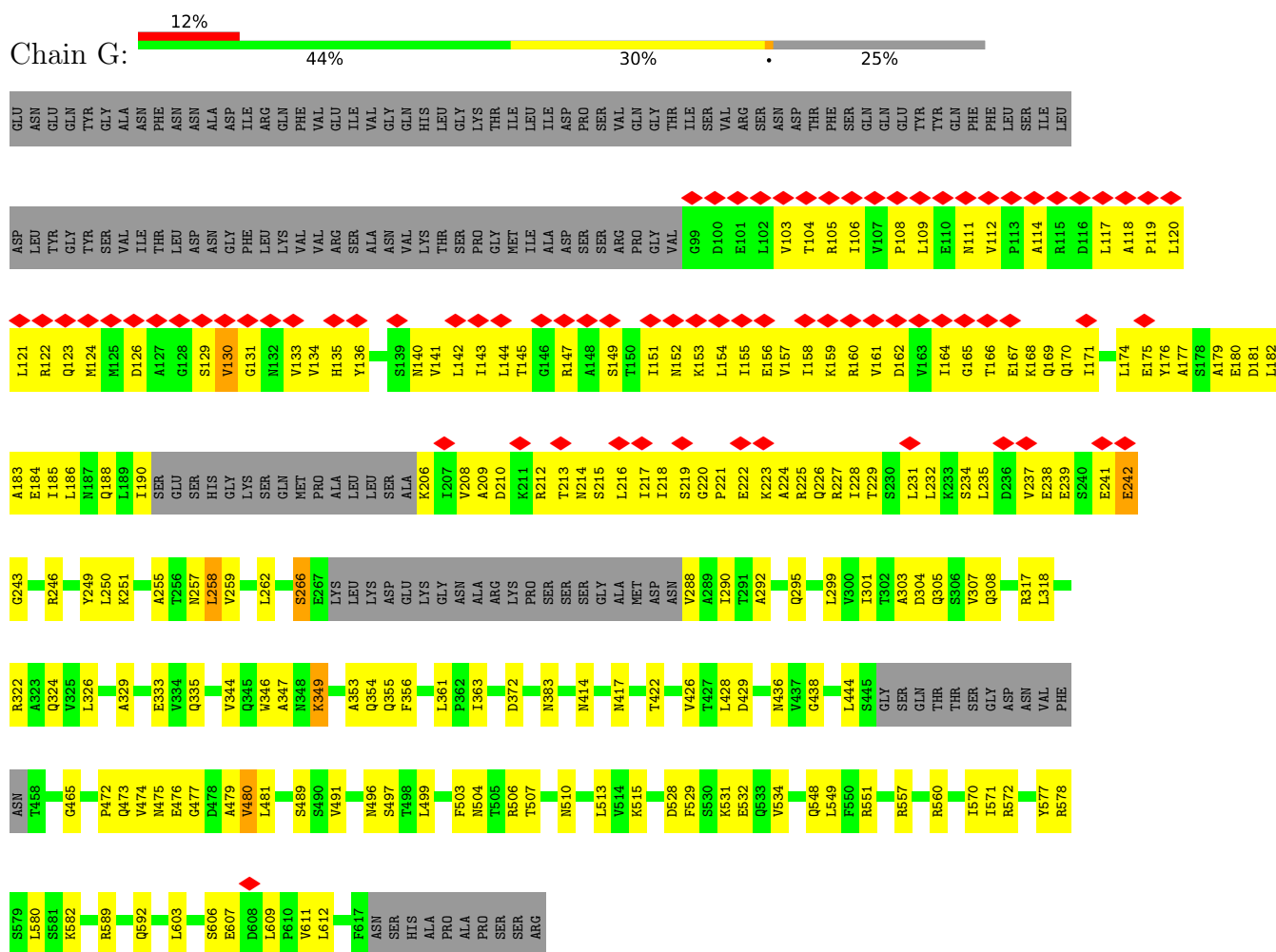




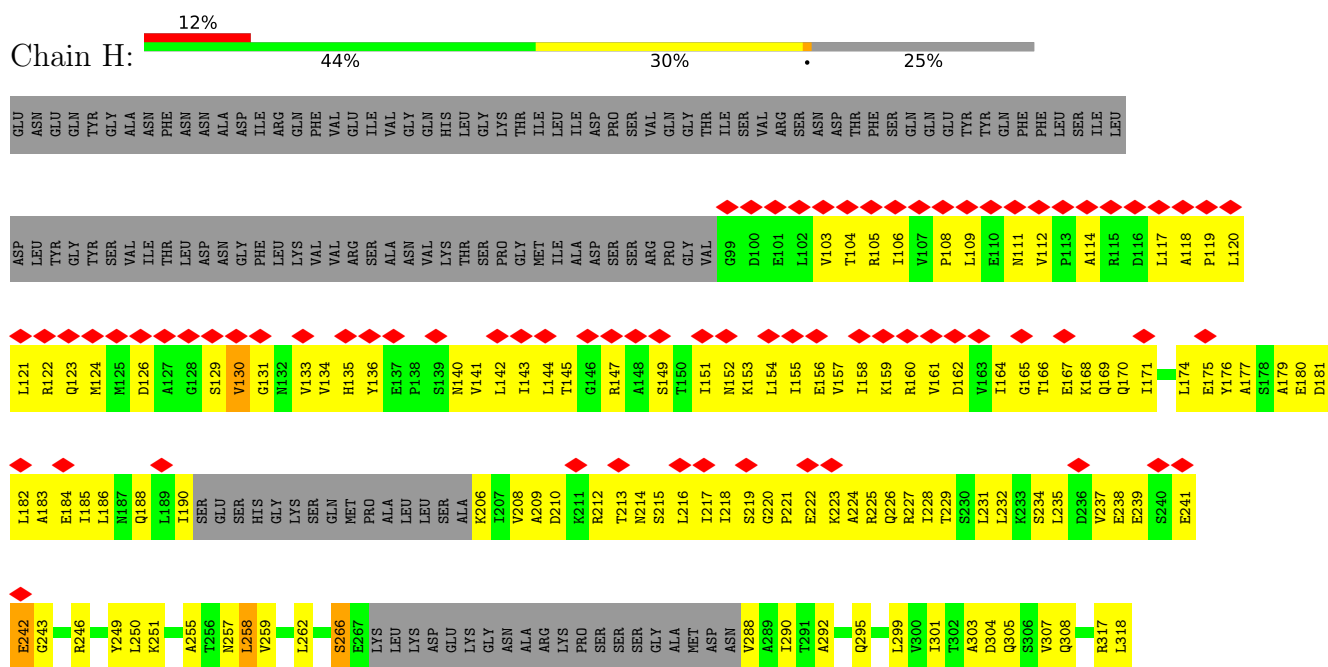
• Molecule 1: Putative type II secretion system protein D



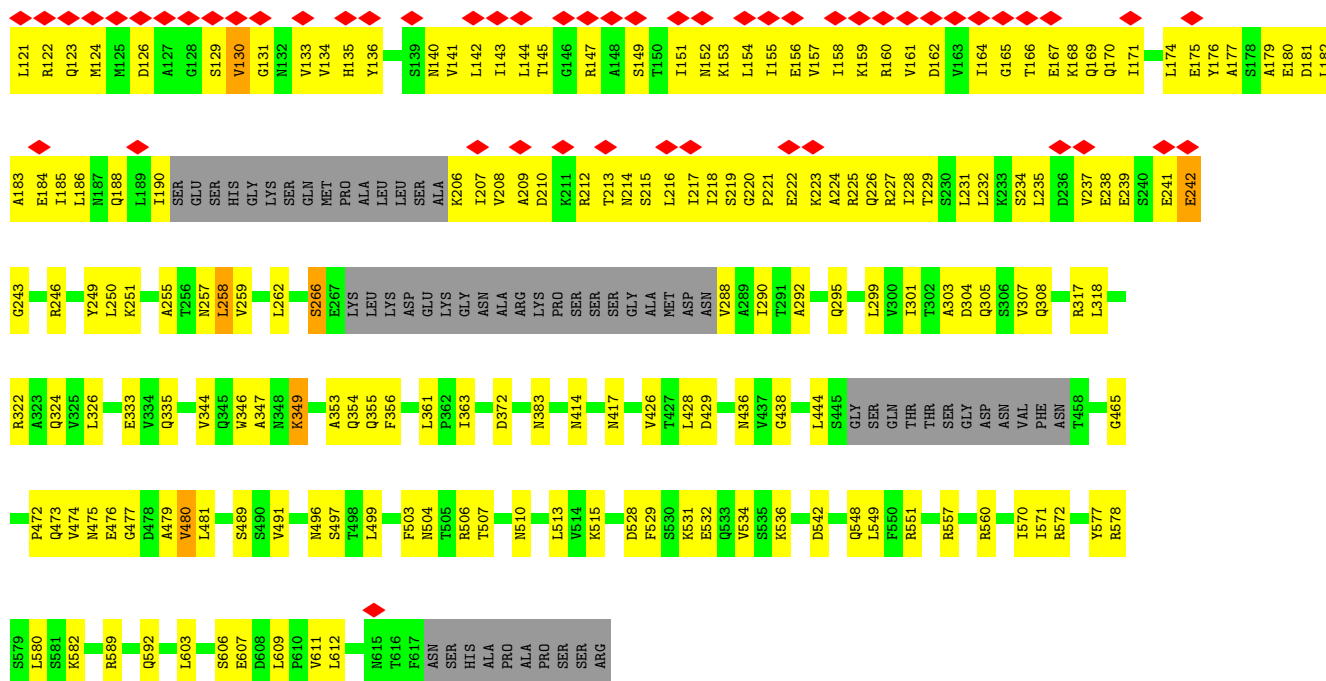
• Molecule 1: Putative type II secretion system protein D



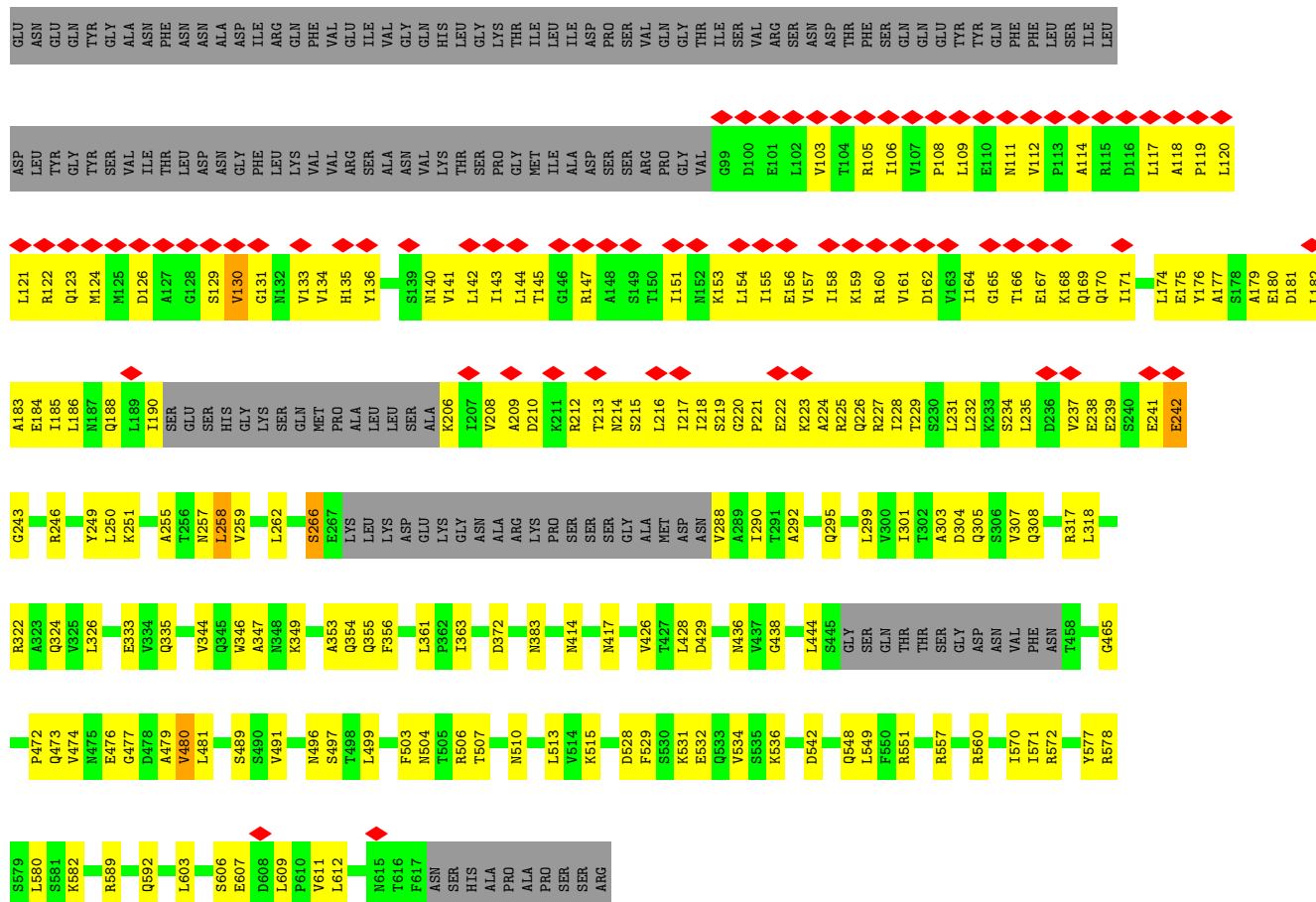
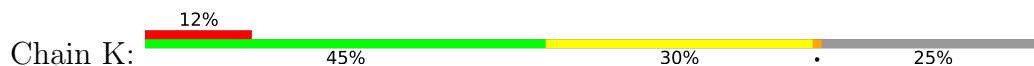
• Molecule 1: Putative type II secretion system protein D



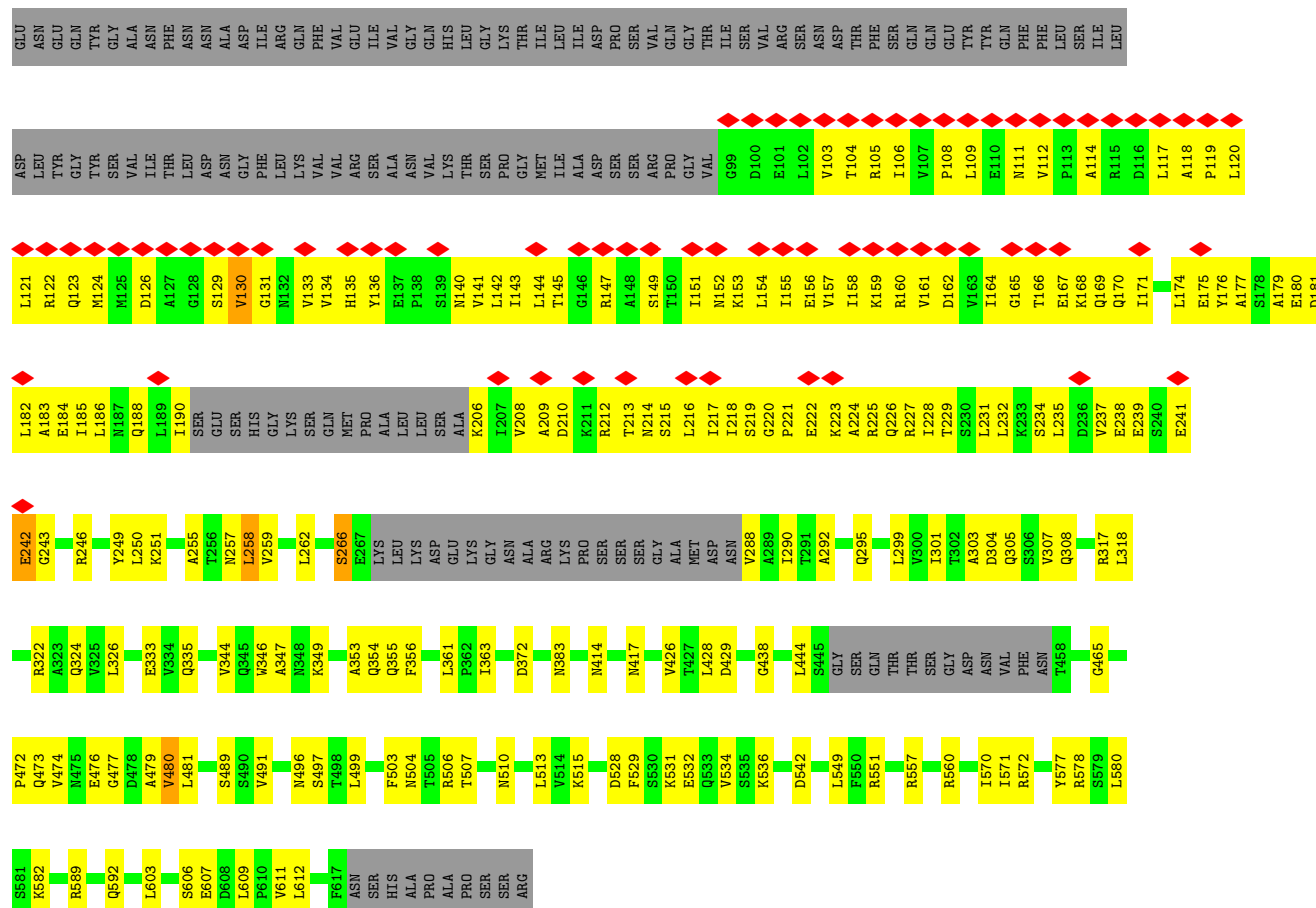
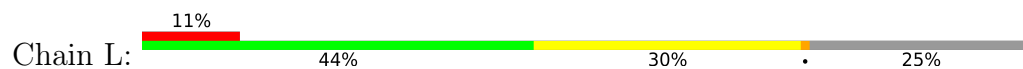




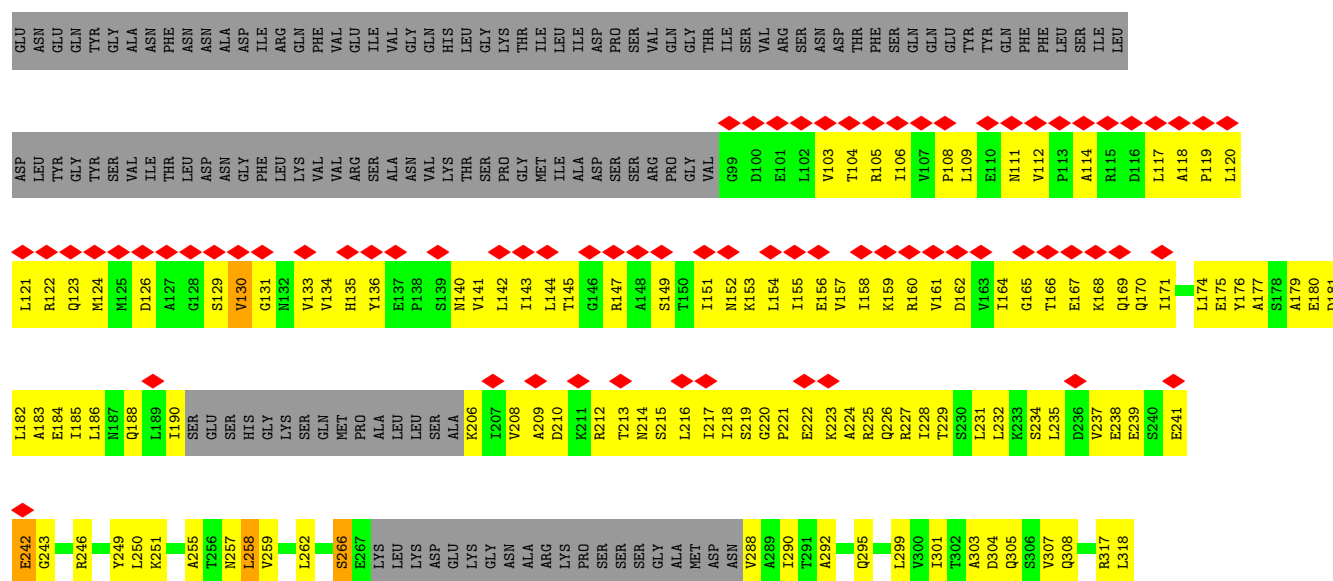
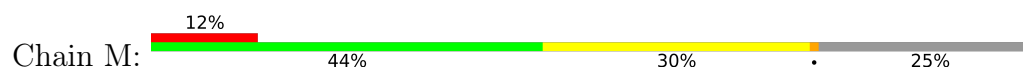
• Molecule 1: Putative type II secretion system protein D

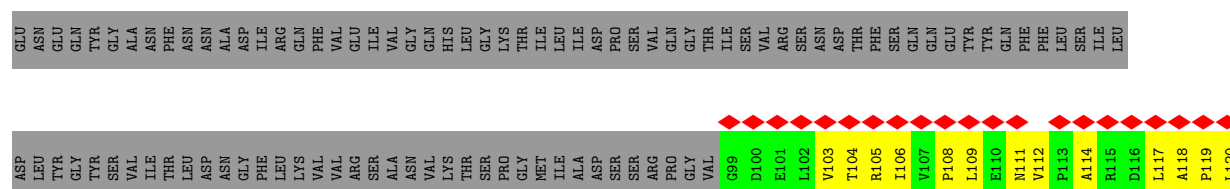


• Molecule 1: Putative type II secretion system protein D



• Molecule 1: Putative type II secretion system protein D





Y577	R578	S579	L580	S581	K582	R589	Q592	K601	A602	L603	S606	E607	D608	L609	P610	V611	L612	F617	ASN	SER	HIS	ALA	ALA	PRO	ALA	PRO	SER	SER	ARG	L121	R122	Q123	M124	D125	D126	A127	G128	S129	V130	G131	N132	V133	V134	H135	Y136	S139	N140	V141	L142	I143	L144	T145	G146	R147	A148	S149	T150	I151	N152	K153	L154	I155	E156	V157	I158	K159	R160	V161	D162	V163	I164	G165	T166	E167	K168	Q169	Q170	I171	L174	E175	Y176	A177	S178	A179	E180	D181	L182																	
G465	P472	V473	V474	N475	E476	G477	D478	A479	V480	L481	S489	S490	V491	M496	S497	T498	L499	F503	N504	T505	T507	N510	L513	V514	K515	D528	F529	S530	K531	E532	Q533	V534	S535	K536	D542	Q548	L549	F550	R551	R557	R560	I570	I571	R572	A183	E184	I185	L186	N187	Q188	L189	I190	SER	GLU	SER	SER	HIS	GLY	LYS	SER	GLN	MET	PRO	ALA	LEU	SER	ALA	K206	I207	V208	A209	D210	K211	R212	T213	N214	S215	L216	I217	I218	S219	G220	P221	E222	K223	A224	R225	Q226	R227	I228	T229	S230	L231	L232	K233	S234	L235	D236	V237	E238	E239	S240	E241	E242
G243	R246	Y249	K251	A255	T256	N257	L258	V259	L262	S266	E267	LYS	LEU	LYS	ASP	GLU	LYS	GLY	ASN	ALA	ARG	LYS	PRO	SER	SER	GLY	ALA	MET	ASP	ASN	V288	A289	I290	T291	A292	Q295	L299	V300	I301	T302	A303	D304	Q305	S306	V307	Q308	R317	L318	G243	R246	Y249	K251	A255	T256	N257	L258	V259	L262	S266	E267	LYS	LEU	LYS	ASP	GLU	LYS	GLY	ASN	ALA	ARG	LYS	PRO	SER	SER	GLY	ALA	MET	ASP	ASN	V288	A289	I290	T291	A292	Q295	L299	V300	I301	T302	A303	D304	Q305	S306	V307	Q308	R317	L318							
R322	A323	Q324	L326	E333	V334	Q335	V344	Q345	R346	A347	N348	K349	A353	Q354	Q355	F356	L361	P362	I363	D372	N383	M388	N414	N417	V426	T427	L428	D429	N436	V437	G438	L444	S445	GLY	SER	GLN	THR	SER	GLY	ASP	ASN	VAL	PHE	ASN	T458	R322	A323	Q324	L326	E333	V334	Q335	V344	Q345	R346	A347	N348	K349	A353	Q354	Q355	F356	L361	P362	I363	D372	N383	M388	N414	N417	V426	T427	L428	D429	N436	V437	G438	L444	S445	GLY	SER	GLN	THR	SER	GLY	ASP	ASN	VAL	PHE	ASN	T458													
G465	P472	V473	V474	N475	E476	G477	D478	A479	V480	L481	S489	S490	V491	M496	S497	T498	L499	F503	N504	T505	T507	N510	L513	V514	K515	D528	F529	S530	K531	E532	Q533	V534	S535	K536	D542	Q548	L549	F550	R551	R557	R560	I570	I571	R572	A183	E184	I185	L186	N187	Q188	L189	I190	SER	GLU	SER	SER	HIS	GLY	LYS	SER	GLN	MET	PRO	ALA	LEU	SER	ALA	K206	I207	V208	A209	D210	K211	R212	T213	N214	S215	L216	I217	I218	S219	G220	P221	E222	K223	A224	R225	Q226	R227	I228	T229	S230	L231	L232	K233	S234	L235	D236	V237	E238	E239	S240	E241	E242
Y577	R578	S579	L580	S581	K582	R589	Q592	K601	A602	L603	S606	E607	D608	L609	P610	V611	L612	F617	ASN	SER	HIS	ALA	ALA	PRO	ALA	PRO	SER	SER	ARG	L121	R122	Q123	M124	D125	D126	A127	G128	S129	V130	G131	N132	V133	V134	H135	Y136	S139	N140	V141	L142	I143	L144	T145	G146	R147	A148	S149	T150	I151	N152	K153	L154	I155	E156	V157	I158	K159	R160	V161	D162	V163	I164	G165	T166	E167	K168	Q169	Q170	I171	L174	E175	Y176	A177	S178	A179	E180	D181	L182																	

4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C15	Depositor
Number of particles used	30659	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	1.6	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor
Maximum map value	0.296	Depositor
Minimum map value	-0.161	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.005	Depositor
Recommended contour level	0.022	Depositor
Map size (Å)	501.6, 501.6, 501.6	wwPDB
Map dimensions	380, 380, 380	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.32, 1.32, 1.32	Depositor

5 Model quality

5.1 Standard geometry

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	$\# Z > 5$	RMSZ	$\# Z > 5$
1	A	0.40	0/3673	0.60	1/4983 (0.0%)
1	B	0.40	0/3673	0.60	1/4983 (0.0%)
1	C	0.40	0/3673	0.60	1/4983 (0.0%)
1	D	0.40	0/3673	0.61	1/4983 (0.0%)
1	E	0.40	0/3673	0.60	1/4983 (0.0%)
1	F	0.40	0/3673	0.60	1/4983 (0.0%)
1	G	0.40	0/3673	0.60	1/4983 (0.0%)
1	H	0.40	0/3673	0.61	1/4983 (0.0%)
1	I	0.40	0/3673	0.61	1/4983 (0.0%)
1	J	0.41	0/3673	0.60	1/4983 (0.0%)
1	K	0.40	0/3673	0.60	1/4983 (0.0%)
1	L	0.41	0/3673	0.61	1/4983 (0.0%)
1	M	0.40	0/3673	0.61	1/4983 (0.0%)
1	N	0.40	0/3673	0.61	1/4983 (0.0%)
1	O	0.40	0/3673	0.61	1/4983 (0.0%)
All	All	0.40	0/55095	0.60	15/74745 (0.0%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	1
1	B	0	1
1	C	0	1
1	D	0	1
1	E	0	1
1	F	0	1
1	G	0	1
1	H	0	1
1	I	0	1
1	J	0	1

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Mol	Chain	#Chirality outliers	#Planarity outliers
1	K	0	1
1	L	0	1
1	M	0	1
1	N	0	1
1	O	0	1
All	All	0	15

There are no bond length outliers.

All (15) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	H	266	SER	CB-CA-C	-5.24	99.70	110.38
1	L	266	SER	CB-CA-C	-5.24	99.70	110.38
1	B	266	SER	CB-CA-C	-5.22	99.73	110.38
1	D	266	SER	CB-CA-C	-5.22	99.73	110.38
1	O	266	SER	CB-CA-C	-5.22	99.74	110.38
1	A	266	SER	CB-CA-C	-5.21	99.74	110.38
1	K	266	SER	CB-CA-C	-5.21	99.74	110.38
1	N	266	SER	CB-CA-C	-5.21	99.75	110.38
1	E	266	SER	CB-CA-C	-5.21	99.75	110.38
1	I	266	SER	CB-CA-C	-5.21	99.75	110.38
1	M	266	SER	CB-CA-C	-5.21	99.76	110.38
1	F	266	SER	CB-CA-C	-5.20	99.76	110.38
1	G	266	SER	CB-CA-C	-5.20	99.77	110.38
1	C	266	SER	CB-CA-C	-5.20	99.77	110.38
1	J	266	SER	CB-CA-C	-5.19	99.79	110.38

There are no chirality outliers.

All (15) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	166	THR	Peptide
1	B	166	THR	Peptide
1	C	166	THR	Peptide
1	D	166	THR	Peptide
1	E	166	THR	Peptide
1	F	166	THR	Peptide
1	G	166	THR	Peptide
1	H	166	THR	Peptide
1	I	166	THR	Peptide
1	J	166	THR	Peptide

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Mol	Chain	Res	Type	Group
1	K	166	THR	Peptide
1	L	166	THR	Peptide
1	M	166	THR	Peptide
1	N	166	THR	Peptide
1	O	166	THR	Peptide

5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	3632	0	3702	229	0
1	B	3632	0	3702	231	0
1	C	3632	0	3702	231	0
1	D	3632	0	3702	230	0
1	E	3632	0	3702	226	0
1	F	3632	0	3702	224	0
1	G	3632	0	3702	226	0
1	H	3632	0	3702	226	0
1	I	3632	0	3702	231	0
1	J	3632	0	3702	236	0
1	K	3632	0	3702	227	0
1	L	3632	0	3702	225	0
1	M	3632	0	3702	229	0
1	N	3632	0	3702	231	0
1	O	3632	0	3702	232	0
All	All	54480	0	55530	2982	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 27.

All (2982) 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:B:210:ASP:O	1:B:214:ASN:HA	1.50	1.11
1:E:210:ASP:O	1:E:214:ASN:HA	1.50	1.11
1:H:210:ASP:O	1:H:214:ASN:HA	1.50	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:210:ASP:O	1:A:214:ASN:HA	1.50	1.11
1:I:210:ASP:O	1:I:214:ASN:HA	1.50	1.11
1:C:210:ASP:O	1:C:214:ASN:HA	1.50	1.11
1:G:210:ASP:O	1:G:214:ASN:HA	1.50	1.11
1:F:210:ASP:O	1:F:214:ASN:HA	1.50	1.11
1:J:210:ASP:O	1:J:214:ASN:HA	1.50	1.10
1:O:210:ASP:O	1:O:214:ASN:HA	1.51	1.10
1:D:210:ASP:O	1:D:214:ASN:HA	1.50	1.10
1:N:210:ASP:O	1:N:214:ASN:HA	1.50	1.09
1:K:210:ASP:O	1:K:214:ASN:HA	1.50	1.09
1:M:210:ASP:O	1:M:214:ASN:HA	1.50	1.08
1:L:210:ASP:O	1:L:214:ASN:HA	1.50	1.07
1:D:333:GLU:OE1	1:D:506:ARG:NH1	1.90	1.05
1:H:333:GLU:OE1	1:H:506:ARG:NH1	1.90	1.05
1:L:333:GLU:OE1	1:L:506:ARG:NH1	1.90	1.05
1:M:333:GLU:OE1	1:M:506:ARG:NH1	1.90	1.05
1:B:333:GLU:OE1	1:B:506:ARG:NH1	1.90	1.04
1:I:333:GLU:OE1	1:I:506:ARG:NH1	1.90	1.04
1:A:333:GLU:OE1	1:A:506:ARG:NH1	1.90	1.04
1:C:333:GLU:OE1	1:C:506:ARG:NH1	1.90	1.03
1:F:333:GLU:OE1	1:F:506:ARG:NH1	1.90	1.03
1:E:333:GLU:OE1	1:E:506:ARG:NH1	1.90	1.03
1:G:333:GLU:OE1	1:G:506:ARG:NH1	1.90	1.03
1:J:333:GLU:OE1	1:J:506:ARG:NH1	1.90	1.03
1:K:333:GLU:OE1	1:K:506:ARG:NH1	1.90	1.03
1:N:333:GLU:OE1	1:N:506:ARG:NH1	1.90	1.03
1:O:333:GLU:OE1	1:O:506:ARG:NH1	1.90	1.03
1:D:119:PRO:O	1:D:123:GLN:HB2	1.65	0.97
1:E:119:PRO:O	1:E:123:GLN:HB2	1.65	0.97
1:C:119:PRO:O	1:C:123:GLN:HB2	1.65	0.96
1:J:119:PRO:O	1:J:123:GLN:HB2	1.65	0.96
1:I:119:PRO:O	1:I:123:GLN:HB2	1.65	0.96
1:A:119:PRO:O	1:A:123:GLN:HB2	1.65	0.96
1:F:119:PRO:O	1:F:123:GLN:HB2	1.65	0.96
1:K:119:PRO:O	1:K:123:GLN:HB2	1.65	0.95
1:O:119:PRO:O	1:O:123:GLN:HB2	1.65	0.95
1:B:119:PRO:O	1:B:123:GLN:HB2	1.65	0.95
1:H:119:PRO:O	1:H:123:GLN:HB2	1.65	0.95
1:K:212:ARG:HH22	1:L:177:ALA:HB1	1.32	0.95
1:G:119:PRO:O	1:G:123:GLN:HB2	1.65	0.95
1:N:119:PRO:O	1:N:123:GLN:HB2	1.65	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:212:ARG:HH22	1:I:177:ALA:HB1	1.31	0.94
1:N:212:ARG:HH22	1:O:177:ALA:HB1	1.32	0.94
1:L:119:PRO:O	1:L:123:GLN:HB2	1.65	0.94
1:M:119:PRO:O	1:M:123:GLN:HB2	1.65	0.94
1:A:218:ILE:HG12	1:A:228:ILE:HG21	1.49	0.94
1:L:212:ARG:HH22	1:M:177:ALA:HB1	1.32	0.94
1:O:218:ILE:HG12	1:O:228:ILE:HG21	1.49	0.94
1:I:212:ARG:HH22	1:J:177:ALA:HB1	1.31	0.94
1:B:218:ILE:HG12	1:B:228:ILE:HG21	1.49	0.93
1:N:218:ILE:HG12	1:N:228:ILE:HG21	1.49	0.93
1:H:218:ILE:HG12	1:H:228:ILE:HG21	1.49	0.93
1:I:218:ILE:HG12	1:I:228:ILE:HG21	1.49	0.93
1:J:212:ARG:HH22	1:K:177:ALA:HB1	1.32	0.93
1:G:218:ILE:HG12	1:G:228:ILE:HG21	1.49	0.93
1:C:218:ILE:HG12	1:C:228:ILE:HG21	1.49	0.93
1:M:212:ARG:HH22	1:N:177:ALA:HB1	1.31	0.93
1:A:177:ALA:HB1	1:O:212:ARG:HH22	1.32	0.93
1:J:218:ILE:HG12	1:J:228:ILE:HG21	1.49	0.93
1:B:212:ARG:HH22	1:C:177:ALA:HB1	1.32	0.92
1:M:218:ILE:HG12	1:M:228:ILE:HG21	1.49	0.92
1:E:212:ARG:HH22	1:F:177:ALA:HB1	1.32	0.92
1:F:218:ILE:HG12	1:F:228:ILE:HG21	1.49	0.92
1:G:212:ARG:HH22	1:H:177:ALA:HB1	1.32	0.92
1:F:212:ARG:HH22	1:G:177:ALA:HB1	1.32	0.92
1:D:218:ILE:HG12	1:D:228:ILE:HG21	1.49	0.92
1:L:218:ILE:HG12	1:L:228:ILE:HG21	1.49	0.91
1:A:212:ARG:HH22	1:B:177:ALA:HB1	1.31	0.91
1:K:218:ILE:HG12	1:K:228:ILE:HG21	1.49	0.91
1:C:212:ARG:HH22	1:D:177:ALA:HB1	1.32	0.91
1:E:218:ILE:HG12	1:E:228:ILE:HG21	1.49	0.91
1:D:212:ARG:HH22	1:E:177:ALA:HB1	1.31	0.90
1:N:210:ASP:OD2	1:N:213:THR:OG1	1.91	0.89
1:J:210:ASP:OD2	1:J:213:THR:OG1	1.92	0.88
1:C:109:LEU:O	1:C:140:ASN:ND2	2.07	0.88
1:K:210:ASP:OD2	1:K:213:THR:OG1	1.91	0.88
1:I:109:LEU:O	1:I:140:ASN:ND2	2.07	0.88
1:B:109:LEU:O	1:B:140:ASN:ND2	2.07	0.88
1:E:109:LEU:O	1:E:140:ASN:ND2	2.07	0.88
1:G:109:LEU:O	1:G:140:ASN:ND2	2.07	0.88
1:F:109:LEU:O	1:F:140:ASN:ND2	2.07	0.88
1:M:210:ASP:OD2	1:M:213:THR:OG1	1.91	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:210:ASP:OD2	1:O:213:THR:OG1	1.91	0.88
1:A:109:LEU:O	1:A:140:ASN:ND2	2.07	0.88
1:G:210:ASP:OD2	1:G:213:THR:OG1	1.91	0.88
1:J:109:LEU:O	1:J:140:ASN:ND2	2.07	0.88
1:B:210:ASP:OD2	1:B:213:THR:OG1	1.91	0.87
1:H:109:LEU:O	1:H:140:ASN:ND2	2.07	0.87
1:I:210:ASP:OD2	1:I:213:THR:OG1	1.91	0.87
1:C:210:ASP:OD2	1:C:213:THR:OG1	1.92	0.87
1:K:109:LEU:O	1:K:140:ASN:ND2	2.07	0.87
1:M:109:LEU:O	1:M:140:ASN:ND2	2.07	0.87
1:F:210:ASP:OD2	1:F:213:THR:OG1	1.91	0.87
1:D:109:LEU:O	1:D:140:ASN:ND2	2.07	0.87
1:H:210:ASP:OD2	1:H:213:THR:OG1	1.91	0.87
1:N:109:LEU:O	1:N:140:ASN:ND2	2.07	0.87
1:L:109:LEU:O	1:L:140:ASN:ND2	2.07	0.87
1:L:210:ASP:OD2	1:L:213:THR:OG1	1.92	0.87
1:A:210:ASP:OD2	1:A:213:THR:OG1	1.91	0.86
1:O:109:LEU:O	1:O:140:ASN:ND2	2.07	0.86
1:D:210:ASP:OD2	1:D:213:THR:OG1	1.91	0.86
1:O:206:LYS:N	1:O:219:SER:HG	1.74	0.86
1:F:242:GLU:CG	1:F:243:GLY:H	1.89	0.86
1:I:206:LYS:N	1:I:219:SER:HG	1.74	0.86
1:L:206:LYS:N	1:L:219:SER:HG	1.74	0.86
1:J:206:LYS:N	1:J:219:SER:HG	1.74	0.86
1:K:242:GLU:CG	1:K:243:GLY:H	1.89	0.86
1:M:242:GLU:CG	1:M:243:GLY:H	1.89	0.86
1:N:206:LYS:N	1:N:219:SER:HG	1.74	0.85
1:D:242:GLU:CG	1:D:243:GLY:H	1.89	0.85
1:E:210:ASP:OD2	1:E:213:THR:OG1	1.92	0.85
1:H:242:GLU:CG	1:H:243:GLY:H	1.89	0.85
1:A:242:GLU:CG	1:A:243:GLY:H	1.89	0.85
1:K:206:LYS:N	1:K:219:SER:HG	1.75	0.85
1:B:242:GLU:CG	1:B:243:GLY:H	1.89	0.85
1:G:206:LYS:N	1:G:219:SER:HG	1.74	0.85
1:E:206:LYS:N	1:E:219:SER:HG	1.74	0.85
1:I:242:GLU:CG	1:I:243:GLY:H	1.89	0.85
1:J:242:GLU:CG	1:J:243:GLY:H	1.89	0.85
1:C:242:GLU:CG	1:C:243:GLY:H	1.89	0.85
1:N:242:GLU:CG	1:N:243:GLY:H	1.89	0.85
1:G:242:GLU:CG	1:G:243:GLY:H	1.89	0.85
1:O:242:GLU:CG	1:O:243:GLY:H	1.89	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:206:LYS:N	1:M:219:SER:HG	1.74	0.84
1:B:206:LYS:N	1:B:219:SER:HG	1.75	0.84
1:E:242:GLU:CG	1:E:243:GLY:H	1.89	0.84
1:L:242:GLU:CG	1:L:243:GLY:H	1.89	0.84
1:A:206:LYS:N	1:A:219:SER:HG	1.75	0.84
1:D:206:LYS:N	1:D:219:SER:HG	1.75	0.83
1:B:109:LEU:HD23	1:B:114:ALA:HA	1.61	0.83
1:L:109:LEU:HD23	1:L:114:ALA:HA	1.61	0.83
1:C:206:LYS:N	1:C:219:SER:HG	1.75	0.83
1:K:109:LEU:HD23	1:K:114:ALA:HA	1.61	0.83
1:N:109:LEU:HD23	1:N:114:ALA:HA	1.61	0.83
1:C:109:LEU:HD23	1:C:114:ALA:HA	1.61	0.83
1:D:109:LEU:HD23	1:D:114:ALA:HA	1.61	0.83
1:E:109:LEU:HD23	1:E:114:ALA:HA	1.61	0.83
1:I:109:LEU:HD23	1:I:114:ALA:HA	1.61	0.83
1:A:109:LEU:HD23	1:A:114:ALA:HA	1.61	0.82
1:O:109:LEU:HD23	1:O:114:ALA:HA	1.61	0.82
1:H:109:LEU:HD23	1:H:114:ALA:HA	1.61	0.82
1:J:109:LEU:HD23	1:J:114:ALA:HA	1.61	0.82
1:M:109:LEU:HD23	1:M:114:ALA:HA	1.61	0.82
1:F:109:LEU:HD23	1:F:114:ALA:HA	1.61	0.82
1:G:109:LEU:HD23	1:G:114:ALA:HA	1.61	0.82
1:L:117:LEU:HD21	1:L:161:VAL:HG21	1.62	0.82
1:O:117:LEU:HD21	1:O:161:VAL:HG21	1.62	0.81
1:J:117:LEU:HD21	1:J:161:VAL:HG21	1.63	0.81
1:N:117:LEU:HD21	1:N:161:VAL:HG21	1.63	0.81
1:K:258:LEU:HD21	1:K:318:LEU:HD23	1.63	0.81
1:L:258:LEU:HD21	1:L:318:LEU:HD23	1.63	0.81
1:M:117:LEU:HD21	1:M:161:VAL:HG21	1.62	0.81
1:A:117:LEU:HD21	1:A:161:VAL:HG21	1.62	0.81
1:I:258:LEU:HD21	1:I:318:LEU:HD23	1.63	0.81
1:J:258:LEU:HD21	1:J:318:LEU:HD23	1.63	0.81
1:B:117:LEU:HD21	1:B:161:VAL:HG21	1.63	0.81
1:I:117:LEU:HD21	1:I:161:VAL:HG21	1.63	0.81
1:K:117:LEU:HD21	1:K:161:VAL:HG21	1.63	0.81
1:F:206:LYS:N	1:F:219:SER:HG	1.77	0.81
1:G:258:LEU:HD21	1:G:318:LEU:HD23	1.63	0.81
1:H:206:LYS:N	1:H:219:SER:HG	1.78	0.81
1:N:258:LEU:HD21	1:N:318:LEU:HD23	1.63	0.81
1:C:117:LEU:HD21	1:C:161:VAL:HG21	1.63	0.80
1:H:258:LEU:HD21	1:H:318:LEU:HD23	1.63	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:117:LEU:HD21	1:G:161:VAL:HG21	1.62	0.80
1:M:258:LEU:HD21	1:M:318:LEU:HD23	1.63	0.80
1:H:117:LEU:HD21	1:H:161:VAL:HG21	1.62	0.80
1:F:117:LEU:HD21	1:F:161:VAL:HG21	1.63	0.80
1:F:258:LEU:HD21	1:F:318:LEU:HD23	1.63	0.80
1:O:258:LEU:HD21	1:O:318:LEU:HD23	1.63	0.80
1:E:117:LEU:HD21	1:E:161:VAL:HG21	1.63	0.80
1:I:141:VAL:HG21	1:J:164:ILE:HD12	1.64	0.80
1:D:117:LEU:HD21	1:D:161:VAL:HG21	1.63	0.80
1:H:141:VAL:HG21	1:I:164:ILE:HD12	1.64	0.80
1:E:242:GLU:HG2	1:E:243:GLY:N	1.97	0.79
1:O:242:GLU:CG	1:O:243:GLY:N	2.45	0.79
1:J:242:GLU:HG2	1:J:243:GLY:N	1.97	0.79
1:L:242:GLU:HG2	1:L:243:GLY:N	1.97	0.79
1:G:242:GLU:HG2	1:G:243:GLY:N	1.98	0.79
1:J:141:VAL:HG21	1:K:164:ILE:HD12	1.64	0.79
1:M:242:GLU:HG2	1:M:243:GLY:N	1.97	0.79
1:D:258:LEU:HD21	1:D:318:LEU:HD23	1.63	0.79
1:H:242:GLU:HG2	1:H:243:GLY:N	1.97	0.79
1:A:258:LEU:HD21	1:A:318:LEU:HD23	1.63	0.79
1:B:258:LEU:HD21	1:B:318:LEU:HD23	1.63	0.79
1:E:258:LEU:HD21	1:E:318:LEU:HD23	1.63	0.79
1:F:242:GLU:HG2	1:F:243:GLY:N	1.97	0.79
1:G:141:VAL:HG21	1:H:164:ILE:HD12	1.64	0.79
1:I:242:GLU:CG	1:I:243:GLY:N	2.45	0.79
1:O:242:GLU:HG2	1:O:243:GLY:N	1.98	0.79
1:C:242:GLU:CG	1:C:243:GLY:N	2.45	0.79
1:C:258:LEU:HD21	1:C:318:LEU:HD23	1.63	0.79
1:G:212:ARG:NE	1:H:181:ASP:OD2	2.16	0.79
1:I:242:GLU:HG2	1:I:243:GLY:N	1.97	0.79
1:A:181:ASP:OD2	1:O:212:ARG:NE	2.16	0.79
1:C:242:GLU:HG2	1:C:243:GLY:N	1.97	0.79
1:F:212:ARG:NE	1:G:181:ASP:OD2	2.16	0.79
1:N:212:ARG:NE	1:O:181:ASP:OD2	2.16	0.79
1:H:212:ARG:NE	1:I:181:ASP:OD2	2.16	0.79
1:L:242:GLU:CG	1:L:243:GLY:N	2.45	0.78
1:M:212:ARG:NE	1:N:181:ASP:OD2	2.16	0.78
1:M:242:GLU:CG	1:M:243:GLY:N	2.45	0.78
1:A:212:ARG:NE	1:B:181:ASP:OD2	2.16	0.78
1:D:242:GLU:CG	1:D:243:GLY:N	2.45	0.78
1:F:242:GLU:CG	1:F:243:GLY:N	2.45	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:242:GLU:HG2	1:B:243:GLY:N	1.97	0.78
1:N:242:GLU:HG2	1:N:243:GLY:N	1.97	0.78
1:K:141:VAL:HG21	1:L:164:ILE:HD12	1.64	0.78
1:N:141:VAL:HG21	1:O:164:ILE:HD12	1.64	0.78
1:M:141:VAL:HG21	1:N:164:ILE:HD12	1.64	0.78
1:A:164:ILE:HD12	1:O:141:VAL:HG21	1.64	0.78
1:C:212:ARG:NE	1:D:181:ASP:OD2	2.16	0.78
1:E:212:ARG:NE	1:F:181:ASP:OD2	2.16	0.78
1:F:141:VAL:HG21	1:G:164:ILE:HD12	1.64	0.78
1:K:242:GLU:HG2	1:K:243:GLY:N	1.97	0.78
1:L:212:ARG:NE	1:M:181:ASP:OD2	2.16	0.78
1:N:242:GLU:CG	1:N:243:GLY:N	2.45	0.78
1:A:141:VAL:HG21	1:B:164:ILE:HD12	1.64	0.78
1:A:242:GLU:CG	1:A:243:GLY:N	2.45	0.78
1:B:212:ARG:NE	1:C:181:ASP:OD2	2.16	0.78
1:I:212:ARG:NE	1:J:181:ASP:OD2	2.16	0.78
1:D:212:ARG:NE	1:E:181:ASP:OD2	2.16	0.77
1:B:141:VAL:HG21	1:C:164:ILE:HD12	1.64	0.77
1:C:141:VAL:HG21	1:D:164:ILE:HD12	1.64	0.77
1:D:141:VAL:HG21	1:E:164:ILE:HD12	1.64	0.77
1:J:242:GLU:CG	1:J:243:GLY:N	2.45	0.77
1:L:141:VAL:HG21	1:M:164:ILE:HD12	1.64	0.77
1:A:242:GLU:HG2	1:A:243:GLY:N	1.97	0.77
1:D:242:GLU:HG2	1:D:243:GLY:N	1.98	0.77
1:K:212:ARG:NE	1:L:181:ASP:OD2	2.16	0.77
1:E:141:VAL:HG21	1:F:164:ILE:HD12	1.64	0.77
1:E:295:GLN:HG2	1:F:257:ASN:HD21	1.50	0.77
1:G:242:GLU:CG	1:G:243:GLY:N	2.45	0.77
1:G:295:GLN:HG2	1:H:257:ASN:HD21	1.50	0.77
1:J:295:GLN:HG2	1:K:257:ASN:HD21	1.50	0.76
1:B:242:GLU:CG	1:B:243:GLY:N	2.45	0.76
1:J:212:ARG:NE	1:K:181:ASP:OD2	2.16	0.76
1:C:295:GLN:HG2	1:D:257:ASN:HD21	1.50	0.76
1:L:295:GLN:HG2	1:M:257:ASN:HD21	1.50	0.76
1:H:295:GLN:HG2	1:I:257:ASN:HD21	1.50	0.76
1:I:295:GLN:HG2	1:J:257:ASN:HD21	1.50	0.76
1:B:295:GLN:HG2	1:C:257:ASN:HD21	1.50	0.76
1:K:609:LEU:HG	1:M:528:ASP:OD2	1.86	0.76
1:B:609:LEU:HG	1:D:528:ASP:OD2	1.86	0.76
1:K:242:GLU:CG	1:K:243:GLY:N	2.45	0.76
1:A:257:ASN:HD21	1:O:295:GLN:HG2	1.50	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:609:LEU:HG	1:C:528:ASP:OD2	1.86	0.75
1:B:528:ASP:OD2	1:O:609:LEU:HG	1.86	0.75
1:C:609:LEU:HG	1:E:528:ASP:OD2	1.86	0.75
1:F:609:LEU:HG	1:H:528:ASP:OD2	1.86	0.75
1:I:609:LEU:HG	1:K:528:ASP:OD2	1.86	0.75
1:J:609:LEU:HG	1:L:528:ASP:OD2	1.86	0.75
1:L:609:LEU:HG	1:N:528:ASP:OD2	1.86	0.75
1:D:609:LEU:HG	1:F:528:ASP:OD2	1.86	0.75
1:H:609:LEU:HG	1:J:528:ASP:OD2	1.86	0.75
1:A:108:PRO:HB3	1:A:141:VAL:HG12	1.69	0.75
1:B:108:PRO:HB3	1:B:141:VAL:HG12	1.69	0.75
1:C:108:PRO:HB3	1:C:141:VAL:HG12	1.69	0.75
1:E:609:LEU:HG	1:G:528:ASP:OD2	1.86	0.75
1:O:108:PRO:HB3	1:O:141:VAL:HG12	1.69	0.75
1:A:528:ASP:OD2	1:N:609:LEU:HG	1.86	0.75
1:D:295:GLN:HG2	1:E:257:ASN:HD21	1.50	0.75
1:D:108:PRO:HB3	1:D:141:VAL:HG12	1.69	0.75
1:G:609:LEU:HG	1:I:528:ASP:OD2	1.86	0.75
1:N:295:GLN:HG2	1:O:257:ASN:HD21	1.50	0.75
1:F:295:GLN:HG2	1:G:257:ASN:HD21	1.50	0.75
1:G:295:GLN:NE2	1:H:322:ARG:HH12	1.85	0.75
1:N:295:GLN:NE2	1:O:322:ARG:HH12	1.85	0.75
1:C:114:ALA:HB1	1:C:142:LEU:HD12	1.69	0.75
1:M:295:GLN:HG2	1:N:257:ASN:HD21	1.50	0.75
1:N:108:PRO:HB3	1:N:141:VAL:HG12	1.69	0.75
1:D:295:GLN:NE2	1:E:322:ARG:HH12	1.85	0.74
1:I:114:ALA:HB1	1:I:142:LEU:HD12	1.69	0.74
1:L:108:PRO:HB3	1:L:141:VAL:HG12	1.69	0.74
1:M:108:PRO:HB3	1:M:141:VAL:HG12	1.69	0.74
1:A:114:ALA:HB1	1:A:142:LEU:HD12	1.69	0.74
1:A:295:GLN:NE2	1:B:322:ARG:HH12	1.85	0.74
1:E:114:ALA:HB1	1:E:142:LEU:HD12	1.69	0.74
1:G:114:ALA:HB1	1:G:142:LEU:HD12	1.69	0.74
1:E:108:PRO:HB3	1:E:141:VAL:HG12	1.69	0.74
1:E:295:GLN:NE2	1:F:322:ARG:HH12	1.85	0.74
1:H:295:GLN:NE2	1:I:322:ARG:HH12	1.85	0.74
1:K:295:GLN:HG2	1:L:257:ASN:HD21	1.50	0.74
1:M:609:LEU:HG	1:O:528:ASP:OD2	1.86	0.74
1:B:326:LEU:HD22	1:B:580:LEU:HD21	1.69	0.74
1:C:326:LEU:HD22	1:C:580:LEU:HD21	1.69	0.74
1:J:295:GLN:NE2	1:K:322:ARG:HH12	1.85	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:114:ALA:HB1	1:K:142:LEU:HD12	1.69	0.74
1:D:326:LEU:HD22	1:D:580:LEU:HD21	1.69	0.74
1:K:108:PRO:HB3	1:K:141:VAL:HG12	1.69	0.74
1:A:295:GLN:HG2	1:B:257:ASN:HD21	1.50	0.74
1:H:242:GLU:CG	1:H:243:GLY:N	2.45	0.74
1:M:114:ALA:HB1	1:M:142:LEU:HD12	1.69	0.74
1:L:295:GLN:NE2	1:M:322:ARG:HH12	1.85	0.74
1:B:295:GLN:NE2	1:C:322:ARG:HH12	1.85	0.74
1:H:111:ASN:ND2	1:H:162:ASP:O	2.21	0.74
1:H:114:ALA:HB1	1:H:142:LEU:HD12	1.69	0.74
1:A:326:LEU:HD22	1:A:580:LEU:HD21	1.69	0.74
1:G:111:ASN:ND2	1:G:162:ASP:O	2.21	0.74
1:K:295:GLN:NE2	1:L:322:ARG:HH12	1.85	0.74
1:F:108:PRO:HB3	1:F:141:VAL:HG12	1.69	0.73
1:F:114:ALA:HB1	1:F:142:LEU:HD12	1.69	0.73
1:J:108:PRO:HB3	1:J:141:VAL:HG12	1.69	0.73
1:O:114:ALA:HB1	1:O:142:LEU:HD12	1.69	0.73
1:B:114:ALA:HB1	1:B:142:LEU:HD12	1.69	0.73
1:E:112:VAL:O	1:E:140:ASN:ND2	2.21	0.73
1:N:114:ALA:HB1	1:N:142:LEU:HD12	1.69	0.73
1:A:111:ASN:ND2	1:A:162:ASP:O	2.21	0.73
1:D:114:ALA:HB1	1:D:142:LEU:HD12	1.69	0.73
1:I:295:GLN:NE2	1:J:322:ARG:HH12	1.85	0.73
1:L:326:LEU:HD22	1:L:580:LEU:HD21	1.69	0.73
1:M:295:GLN:NE2	1:N:322:ARG:HH12	1.85	0.73
1:M:324:GLN:NE2	1:N:513:LEU:H	1.87	0.73
1:A:324:GLN:NE2	1:B:513:LEU:H	1.87	0.73
1:D:324:GLN:NE2	1:E:513:LEU:H	1.87	0.73
1:E:326:LEU:HD22	1:E:580:LEU:HD21	1.69	0.73
1:F:295:GLN:NE2	1:G:322:ARG:HH12	1.85	0.73
1:K:326:LEU:HD22	1:K:580:LEU:HD21	1.69	0.73
1:A:513:LEU:H	1:O:324:GLN:NE2	1.87	0.73
1:B:111:ASN:ND2	1:B:162:ASP:O	2.21	0.73
1:B:112:VAL:O	1:B:140:ASN:ND2	2.21	0.73
1:G:108:PRO:HB3	1:G:141:VAL:HG12	1.69	0.73
1:I:108:PRO:HB3	1:I:141:VAL:HG12	1.69	0.73
1:B:103:VAL:HG21	1:B:105:ARG:HH21	1.54	0.73
1:F:111:ASN:ND2	1:F:162:ASP:O	2.21	0.73
1:G:324:GLN:NE2	1:H:513:LEU:H	1.87	0.73
1:H:108:PRO:HB3	1:H:141:VAL:HG12	1.69	0.73
1:I:208:VAL:HB	1:I:217:ILE:HB	1.71	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:114:ALA:HB1	1:J:142:LEU:HD12	1.69	0.73
1:L:112:VAL:O	1:L:140:ASN:ND2	2.21	0.73
1:B:208:VAL:HB	1:B:217:ILE:HB	1.71	0.73
1:J:324:GLN:NE2	1:K:513:LEU:H	1.87	0.73
1:L:103:VAL:HG21	1:L:105:ARG:HH21	1.54	0.73
1:M:326:LEU:HD22	1:M:580:LEU:HD21	1.69	0.73
1:A:208:VAL:HB	1:A:217:ILE:HB	1.71	0.73
1:C:103:VAL:HG21	1:C:105:ARG:HH21	1.54	0.73
1:H:208:VAL:HB	1:H:217:ILE:HB	1.71	0.73
1:J:208:VAL:HB	1:J:217:ILE:HB	1.71	0.73
1:J:326:LEU:HD22	1:J:580:LEU:HD21	1.69	0.73
1:A:181:ASP:O	1:A:184:GLU:N	2.22	0.73
1:B:324:GLN:NE2	1:C:513:LEU:H	1.87	0.73
1:C:208:VAL:HB	1:C:217:ILE:HB	1.71	0.73
1:C:295:GLN:NE2	1:D:322:ARG:HH12	1.85	0.73
1:C:324:GLN:NE2	1:D:513:LEU:H	1.87	0.73
1:O:181:ASP:O	1:O:184:GLU:N	2.22	0.73
1:G:326:LEU:HD22	1:G:580:LEU:HD21	1.69	0.73
1:H:103:VAL:HG21	1:H:105:ARG:HH21	1.54	0.73
1:H:326:LEU:HD22	1:H:580:LEU:HD21	1.69	0.73
1:J:181:ASP:O	1:J:184:GLU:N	2.22	0.73
1:K:103:VAL:HG21	1:K:105:ARG:HH21	1.54	0.73
1:M:103:VAL:HG21	1:M:105:ARG:HH21	1.54	0.73
1:O:326:LEU:HD22	1:O:580:LEU:HD21	1.69	0.73
1:C:111:ASN:ND2	1:C:162:ASP:O	2.21	0.72
1:F:324:GLN:NE2	1:G:513:LEU:H	1.87	0.72
1:I:326:LEU:HD22	1:I:580:LEU:HD21	1.69	0.72
1:A:103:VAL:HG21	1:A:105:ARG:HH21	1.54	0.72
1:B:181:ASP:O	1:B:184:GLU:N	2.22	0.72
1:D:103:VAL:HG21	1:D:105:ARG:HH21	1.54	0.72
1:D:208:VAL:HB	1:D:217:ILE:HB	1.71	0.72
1:D:262:LEU:HD12	1:D:299:LEU:HD21	1.72	0.72
1:E:181:ASP:O	1:E:184:GLU:N	2.22	0.72
1:F:181:ASP:O	1:F:184:GLU:N	2.22	0.72
1:G:181:ASP:O	1:G:184:GLU:N	2.22	0.72
1:G:262:LEU:HD12	1:G:299:LEU:HD21	1.71	0.72
1:H:112:VAL:O	1:H:140:ASN:ND2	2.21	0.72
1:H:262:LEU:HD12	1:H:299:LEU:HD21	1.71	0.72
1:K:208:VAL:HB	1:K:217:ILE:HB	1.71	0.72
1:N:181:ASP:O	1:N:184:GLU:N	2.22	0.72
1:N:324:GLN:NE2	1:O:513:LEU:H	1.87	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:208:VAL:HB	1:O:217:ILE:HB	1.71	0.72
1:A:112:VAL:O	1:A:140:ASN:ND2	2.21	0.72
1:A:262:LEU:HD12	1:A:299:LEU:HD21	1.72	0.72
1:C:262:LEU:HD12	1:C:299:LEU:HD21	1.72	0.72
1:D:181:ASP:O	1:D:184:GLU:N	2.22	0.72
1:E:262:LEU:HD12	1:E:299:LEU:HD21	1.71	0.72
1:F:262:LEU:HD12	1:F:299:LEU:HD21	1.71	0.72
1:G:208:VAL:HB	1:G:217:ILE:HB	1.71	0.72
1:H:181:ASP:O	1:H:184:GLU:N	2.22	0.72
1:I:181:ASP:O	1:I:184:GLU:N	2.22	0.72
1:L:324:GLN:NE2	1:M:513:LEU:H	1.87	0.72
1:C:181:ASP:O	1:C:184:GLU:N	2.22	0.72
1:E:208:VAL:HB	1:E:217:ILE:HB	1.71	0.72
1:E:324:GLN:NE2	1:F:513:LEU:H	1.87	0.72
1:I:324:GLN:NE2	1:J:513:LEU:H	1.87	0.72
1:K:181:ASP:O	1:K:184:GLU:N	2.22	0.72
1:K:206:LYS:N	1:K:219:SER:OG	2.23	0.72
1:K:324:GLN:NE2	1:L:513:LEU:H	1.87	0.72
1:L:114:ALA:HB1	1:L:142:LEU:HD12	1.69	0.72
1:N:326:LEU:HD22	1:N:580:LEU:HD21	1.69	0.72
1:O:262:LEU:HD12	1:O:299:LEU:HD21	1.72	0.72
1:F:103:VAL:HG21	1:F:105:ARG:HH21	1.54	0.72
1:J:206:LYS:N	1:J:219:SER:OG	2.23	0.72
1:K:210:ASP:O	1:K:214:ASN:CA	2.36	0.72
1:L:208:VAL:HB	1:L:217:ILE:HB	1.71	0.72
1:F:208:VAL:HB	1:F:217:ILE:HB	1.71	0.72
1:F:326:LEU:HD22	1:F:580:LEU:HD21	1.69	0.72
1:J:103:VAL:HG21	1:J:105:ARG:HH21	1.54	0.72
1:N:208:VAL:HB	1:N:217:ILE:HB	1.71	0.72
1:A:322:ARG:HH12	1:O:295:GLN:NE2	1.85	0.72
1:B:262:LEU:HD12	1:B:299:LEU:HD21	1.71	0.72
1:E:288:VAL:HG21	1:E:307:VAL:HG11	1.71	0.72
1:F:288:VAL:HG21	1:F:307:VAL:HG11	1.71	0.72
1:H:288:VAL:HG21	1:H:307:VAL:HG11	1.71	0.72
1:I:288:VAL:HG21	1:I:307:VAL:HG11	1.71	0.72
1:M:208:VAL:HB	1:M:217:ILE:HB	1.71	0.72
1:H:324:GLN:NE2	1:I:513:LEU:H	1.87	0.72
1:J:262:LEU:HD12	1:J:299:LEU:HD21	1.71	0.72
1:M:111:ASN:ND2	1:M:162:ASP:O	2.21	0.72
1:M:181:ASP:O	1:M:184:GLU:N	2.22	0.72
1:B:206:LYS:N	1:B:219:SER:OG	2.23	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:112:VAL:O	1:D:140:ASN:ND2	2.21	0.72
1:E:111:ASN:ND2	1:E:162:ASP:O	2.21	0.72
1:G:288:VAL:HG21	1:G:307:VAL:HG11	1.71	0.72
1:I:262:LEU:HD12	1:I:299:LEU:HD21	1.71	0.72
1:D:206:LYS:N	1:D:219:SER:OG	2.23	0.72
1:M:262:LEU:HD12	1:M:299:LEU:HD21	1.71	0.72
1:N:103:VAL:HG21	1:N:105:ARG:HH21	1.54	0.72
1:D:288:VAL:HG21	1:D:307:VAL:HG11	1.72	0.71
1:E:206:LYS:N	1:E:219:SER:OG	2.23	0.71
1:L:111:ASN:ND2	1:L:162:ASP:O	2.21	0.71
1:L:206:LYS:N	1:L:219:SER:OG	2.23	0.71
1:D:111:ASN:ND2	1:D:162:ASP:O	2.21	0.71
1:E:242:GLU:CG	1:E:243:GLY:N	2.45	0.71
1:I:103:VAL:HG21	1:I:105:ARG:HH21	1.54	0.71
1:J:288:VAL:HG21	1:J:307:VAL:HG11	1.72	0.71
1:K:262:LEU:HD12	1:K:299:LEU:HD21	1.71	0.71
1:L:262:LEU:HD12	1:L:299:LEU:HD21	1.71	0.71
1:C:206:LYS:N	1:C:219:SER:OG	2.23	0.71
1:E:103:VAL:HG21	1:E:105:ARG:HH21	1.54	0.71
1:L:181:ASP:O	1:L:184:GLU:N	2.22	0.71
1:N:206:LYS:N	1:N:219:SER:OG	2.23	0.71
1:D:210:ASP:O	1:D:214:ASN:CA	2.36	0.71
1:K:112:VAL:O	1:K:140:ASN:ND2	2.21	0.71
1:M:589:ARG:HB2	1:M:612:LEU:HD12	1.73	0.71
1:N:262:LEU:HD12	1:N:299:LEU:HD21	1.72	0.71
1:O:103:VAL:HG21	1:O:105:ARG:HH21	1.54	0.71
1:O:206:LYS:N	1:O:219:SER:OG	2.23	0.71
1:O:210:ASP:O	1:O:214:ASN:CA	2.36	0.71
1:F:206:LYS:N	1:F:219:SER:OG	2.23	0.71
1:L:589:ARG:HB2	1:L:612:LEU:HD12	1.73	0.71
1:C:288:VAL:HG21	1:C:307:VAL:HG11	1.71	0.71
1:C:589:ARG:HB2	1:C:612:LEU:HD12	1.73	0.71
1:D:589:ARG:HB2	1:D:612:LEU:HD12	1.73	0.71
1:I:206:LYS:N	1:I:219:SER:OG	2.23	0.71
1:K:111:ASN:ND2	1:K:162:ASP:O	2.21	0.71
1:K:589:ARG:HB2	1:K:612:LEU:HD12	1.73	0.71
1:N:255:ALA:O	1:N:259:VAL:HG23	1.91	0.71
1:N:589:ARG:HB2	1:N:612:LEU:HD12	1.73	0.71
1:B:589:ARG:HB2	1:B:612:LEU:HD12	1.73	0.71
1:G:206:LYS:N	1:G:219:SER:OG	2.23	0.71
1:K:288:VAL:HG21	1:K:307:VAL:HG11	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:112:VAL:O	1:O:140:ASN:ND2	2.21	0.71
1:O:255:ALA:O	1:O:259:VAL:HG23	1.91	0.71
1:D:255:ALA:O	1:D:259:VAL:HG23	1.91	0.71
1:E:255:ALA:O	1:E:259:VAL:HG23	1.91	0.71
1:O:589:ARG:HB2	1:O:612:LEU:HD12	1.73	0.71
1:A:589:ARG:HB2	1:A:612:LEU:HD12	1.73	0.71
1:E:589:ARG:HB2	1:E:612:LEU:HD12	1.73	0.71
1:J:111:ASN:ND2	1:J:162:ASP:O	2.21	0.71
1:M:288:VAL:HG21	1:M:307:VAL:HG11	1.71	0.71
1:N:288:VAL:HG21	1:N:307:VAL:HG11	1.71	0.71
1:A:288:VAL:HG21	1:A:307:VAL:HG11	1.71	0.71
1:F:255:ALA:O	1:F:259:VAL:HG23	1.91	0.71
1:J:589:ARG:HB2	1:J:612:LEU:HD12	1.73	0.71
1:M:255:ALA:O	1:M:259:VAL:HG23	1.91	0.71
1:O:288:VAL:HG21	1:O:307:VAL:HG11	1.71	0.71
1:M:206:LYS:N	1:M:219:SER:OG	2.23	0.70
1:C:255:ALA:O	1:C:259:VAL:HG23	1.91	0.70
1:G:103:VAL:HG21	1:G:105:ARG:HH21	1.54	0.70
1:G:112:VAL:O	1:G:140:ASN:ND2	2.21	0.70
1:G:255:ALA:O	1:G:259:VAL:HG23	1.91	0.70
1:H:206:LYS:N	1:H:219:SER:OG	2.23	0.70
1:F:589:ARG:HB2	1:F:612:LEU:HD12	1.73	0.70
1:J:210:ASP:O	1:J:214:ASN:CA	2.36	0.70
1:L:288:VAL:HG21	1:L:307:VAL:HG11	1.71	0.70
1:B:288:VAL:HG21	1:B:307:VAL:HG11	1.71	0.70
1:E:210:ASP:O	1:E:214:ASN:CA	2.36	0.70
1:H:210:ASP:O	1:H:214:ASN:CA	2.36	0.70
1:H:255:ALA:O	1:H:259:VAL:HG23	1.91	0.70
1:I:589:ARG:HB2	1:I:612:LEU:HD12	1.73	0.70
1:A:255:ALA:O	1:A:259:VAL:HG23	1.91	0.70
1:B:255:ALA:O	1:B:259:VAL:HG23	1.91	0.70
1:G:589:ARG:HB2	1:G:612:LEU:HD12	1.73	0.70
1:I:255:ALA:O	1:I:259:VAL:HG23	1.91	0.70
1:A:206:LYS:N	1:A:219:SER:OG	2.23	0.70
1:G:210:ASP:O	1:G:214:ASN:CA	2.36	0.70
1:H:170:GLN:HB3	1:H:225:ARG:NH1	2.07	0.70
1:J:174:LEU:O	1:J:214:ASN:ND2	2.25	0.70
1:J:255:ALA:O	1:J:259:VAL:HG23	1.91	0.70
1:L:255:ALA:O	1:L:259:VAL:HG23	1.91	0.70
1:N:174:LEU:O	1:N:214:ASN:ND2	2.25	0.70
1:C:210:ASP:O	1:C:214:ASN:CA	2.36	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:170:GLN:HB3	1:E:225:ARG:NH1	2.07	0.70
1:F:170:GLN:HB3	1:F:225:ARG:NH1	2.07	0.70
1:H:589:ARG:HB2	1:H:612:LEU:HD12	1.73	0.70
1:K:174:LEU:O	1:K:214:ASN:ND2	2.25	0.70
1:O:174:LEU:O	1:O:214:ASN:ND2	2.25	0.70
1:D:170:GLN:HB3	1:D:225:ARG:NH1	2.07	0.70
1:H:242:GLU:HG3	1:H:243:GLY:H	1.57	0.70
1:N:111:ASN:ND2	1:N:162:ASP:O	2.21	0.70
1:G:170:GLN:HB3	1:G:225:ARG:NH1	2.07	0.69
1:O:170:GLN:HB3	1:O:225:ARG:NH1	2.07	0.69
1:G:242:GLU:HG3	1:G:243:GLY:H	1.57	0.69
1:I:170:GLN:HB3	1:I:225:ARG:NH1	2.07	0.69
1:J:242:GLU:HG3	1:J:243:GLY:H	1.57	0.69
1:L:174:LEU:O	1:L:214:ASN:ND2	2.25	0.69
1:A:170:GLN:HB3	1:A:225:ARG:NH1	2.07	0.69
1:E:242:GLU:HG3	1:E:243:GLY:H	1.57	0.69
1:K:255:ALA:O	1:K:259:VAL:HG23	1.91	0.69
1:M:174:LEU:O	1:M:214:ASN:ND2	2.25	0.69
1:N:170:GLN:HB3	1:N:225:ARG:NH1	2.07	0.69
1:B:222:GLU:HG2	1:B:225:ARG:HD2	1.75	0.69
1:C:112:VAL:O	1:C:140:ASN:ND2	2.21	0.69
1:D:174:LEU:O	1:D:214:ASN:ND2	2.25	0.69
1:J:222:GLU:HG2	1:J:225:ARG:HD2	1.75	0.69
1:K:222:GLU:HG2	1:K:225:ARG:HD2	1.75	0.69
1:M:170:GLN:HB3	1:M:225:ARG:NH1	2.07	0.69
1:A:222:GLU:HG2	1:A:225:ARG:HD2	1.75	0.69
1:E:174:LEU:O	1:E:214:ASN:ND2	2.25	0.69
1:F:174:LEU:O	1:F:214:ASN:ND2	2.25	0.69
1:I:174:LEU:O	1:I:214:ASN:ND2	2.25	0.69
1:N:210:ASP:O	1:N:214:ASN:CA	2.36	0.69
1:A:174:LEU:O	1:A:214:ASN:ND2	2.25	0.69
1:C:174:LEU:O	1:C:214:ASN:ND2	2.25	0.69
1:G:174:LEU:O	1:G:214:ASN:ND2	2.25	0.69
1:G:186:LEU:HD22	1:G:231:LEU:HD23	1.75	0.69
1:L:170:GLN:HB3	1:L:225:ARG:NH1	2.07	0.69
1:B:174:LEU:O	1:B:214:ASN:ND2	2.25	0.69
1:C:295:GLN:HG2	1:D:257:ASN:ND2	2.08	0.69
1:D:222:GLU:HG2	1:D:225:ARG:HD2	1.75	0.69
1:E:222:GLU:HG2	1:E:225:ARG:HD2	1.75	0.69
1:G:222:GLU:HG2	1:G:225:ARG:HD2	1.75	0.69
1:I:186:LEU:HD22	1:I:231:LEU:HD23	1.75	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:222:GLU:HG2	1:I:225:ARG:HD2	1.75	0.69
1:L:222:GLU:HG2	1:L:225:ARG:HD2	1.75	0.69
1:N:112:VAL:O	1:N:140:ASN:ND2	2.21	0.69
1:C:222:GLU:HG2	1:C:225:ARG:HD2	1.75	0.69
1:D:295:GLN:HG2	1:E:257:ASN:ND2	2.08	0.69
1:E:186:LEU:HD22	1:E:231:LEU:HD23	1.75	0.69
1:H:174:LEU:O	1:H:214:ASN:ND2	2.25	0.69
1:H:222:GLU:HG2	1:H:225:ARG:HD2	1.75	0.69
1:J:295:GLN:HG2	1:K:257:ASN:ND2	2.08	0.69
1:K:295:GLN:HG2	1:L:257:ASN:ND2	2.08	0.69
1:O:111:ASN:ND2	1:O:162:ASP:O	2.21	0.69
1:O:222:GLU:HG2	1:O:225:ARG:HD2	1.75	0.69
1:B:170:GLN:HB3	1:B:225:ARG:NH1	2.07	0.69
1:B:295:GLN:HG2	1:C:257:ASN:ND2	2.08	0.69
1:C:170:GLN:HB3	1:C:225:ARG:NH1	2.07	0.69
1:F:210:ASP:O	1:F:214:ASN:CA	2.36	0.69
1:J:112:VAL:O	1:J:140:ASN:ND2	2.21	0.69
1:L:295:GLN:HG2	1:M:257:ASN:ND2	2.08	0.69
1:N:222:GLU:HG2	1:N:225:ARG:HD2	1.75	0.69
1:A:242:GLU:HG3	1:A:243:GLY:H	1.57	0.69
1:E:295:GLN:HG2	1:F:257:ASN:ND2	2.08	0.69
1:F:222:GLU:HG2	1:F:225:ARG:HD2	1.75	0.69
1:K:170:GLN:HB3	1:K:225:ARG:NH1	2.07	0.69
1:M:222:GLU:HG2	1:M:225:ARG:HD2	1.75	0.68
1:C:186:LEU:HD22	1:C:231:LEU:HD23	1.75	0.68
1:F:186:LEU:HD22	1:F:231:LEU:HD23	1.75	0.68
1:I:111:ASN:ND2	1:I:162:ASP:O	2.21	0.68
1:J:170:GLN:HB3	1:J:225:ARG:NH1	2.07	0.68
1:O:242:GLU:HG3	1:O:243:GLY:H	1.57	0.68
1:A:295:GLN:HG2	1:B:257:ASN:ND2	2.08	0.68
1:D:186:LEU:HD22	1:D:231:LEU:HD23	1.75	0.68
1:F:242:GLU:HG3	1:F:243:GLY:H	1.57	0.68
1:H:186:LEU:HD22	1:H:231:LEU:HD23	1.75	0.68
1:I:210:ASP:O	1:I:214:ASN:CA	2.36	0.68
1:I:295:GLN:HG2	1:J:257:ASN:ND2	2.08	0.68
1:K:186:LEU:HD22	1:K:231:LEU:HD23	1.75	0.68
1:M:242:GLU:HG3	1:M:243:GLY:H	1.57	0.68
1:M:295:GLN:HG2	1:N:257:ASN:ND2	2.08	0.68
1:C:242:GLU:HG3	1:C:243:GLY:H	1.57	0.68
1:F:295:GLN:HG2	1:G:257:ASN:ND2	2.08	0.68
1:K:242:GLU:HG3	1:K:243:GLY:H	1.57	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:186:LEU:HD22	1:A:231:LEU:HD23	1.75	0.68
1:J:186:LEU:HD22	1:J:231:LEU:HD23	1.75	0.68
1:L:242:GLU:HG3	1:L:243:GLY:H	1.57	0.68
1:A:257:ASN:ND2	1:O:295:GLN:HG2	2.08	0.68
1:L:186:LEU:HD22	1:L:231:LEU:HD23	1.75	0.68
1:B:242:GLU:HG3	1:B:243:GLY:H	1.57	0.67
1:C:119:PRO:O	1:C:123:GLN:CB	2.42	0.67
1:F:112:VAL:O	1:F:140:ASN:ND2	2.21	0.67
1:H:119:PRO:O	1:H:123:GLN:CB	2.42	0.67
1:H:295:GLN:HG2	1:I:257:ASN:ND2	2.08	0.67
1:N:295:GLN:HG2	1:O:257:ASN:ND2	2.08	0.67
1:B:210:ASP:O	1:B:214:ASN:CA	2.36	0.67
1:F:135:HIS:CD2	1:F:136:TYR:H	2.13	0.67
1:I:112:VAL:O	1:I:140:ASN:ND2	2.21	0.67
1:G:135:HIS:CD2	1:G:136:TYR:H	2.13	0.67
1:N:186:LEU:HD22	1:N:231:LEU:HD23	1.75	0.67
1:E:119:PRO:O	1:E:123:GLN:CB	2.42	0.67
1:E:135:HIS:CD2	1:E:136:TYR:H	2.13	0.67
1:H:135:HIS:CD2	1:H:136:TYR:H	2.13	0.67
1:I:135:HIS:CD2	1:I:136:TYR:H	2.13	0.67
1:L:210:ASP:O	1:L:214:ASN:CA	2.36	0.67
1:N:242:GLU:HG3	1:N:243:GLY:H	1.57	0.67
1:A:241:GLU:O	1:A:242:GLU:HB3	1.95	0.67
1:B:186:LEU:HD22	1:B:231:LEU:HD23	1.75	0.67
1:B:324:GLN:HE21	1:C:513:LEU:H	1.43	0.67
1:D:135:HIS:CD2	1:D:136:TYR:H	2.13	0.67
1:M:606:SER:HB3	1:O:532:GLU:HB2	1.77	0.67
1:D:242:GLU:HG3	1:D:243:GLY:H	1.57	0.67
1:G:295:GLN:HG2	1:H:257:ASN:ND2	2.08	0.67
1:A:532:GLU:HB2	1:N:606:SER:HB3	1.77	0.67
1:F:119:PRO:O	1:F:123:GLN:CB	2.42	0.67
1:I:242:GLU:HG3	1:I:243:GLY:H	1.57	0.67
1:J:135:HIS:CD2	1:J:136:TYR:H	2.13	0.67
1:L:606:SER:HB3	1:N:532:GLU:HB2	1.77	0.67
1:N:135:HIS:CD2	1:N:136:TYR:H	2.13	0.67
1:O:135:HIS:CD2	1:O:136:TYR:H	2.13	0.67
1:A:513:LEU:H	1:O:324:GLN:HE21	1.43	0.67
1:B:241:GLU:O	1:B:242:GLU:HB3	1.95	0.67
1:M:186:LEU:HD22	1:M:231:LEU:HD23	1.75	0.67
1:M:210:ASP:O	1:M:214:ASN:CA	2.36	0.67
1:O:241:GLU:O	1:O:242:GLU:HB3	1.95	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:186:LEU:HD22	1:O:231:LEU:HD23	1.75	0.67
1:K:241:GLU:O	1:K:242:GLU:HB3	1.95	0.66
1:K:606:SER:HB3	1:M:532:GLU:HB2	1.77	0.66
1:M:135:HIS:CD2	1:M:136:TYR:H	2.13	0.66
1:A:135:HIS:CD2	1:A:136:TYR:H	2.13	0.66
1:C:135:HIS:CD2	1:C:136:TYR:H	2.13	0.66
1:D:324:GLN:HE21	1:E:513:LEU:H	1.43	0.66
1:B:532:GLU:HB2	1:O:606:SER:HB3	1.77	0.66
1:J:606:SER:HB3	1:L:532:GLU:HB2	1.77	0.66
1:K:119:PRO:O	1:K:123:GLN:CB	2.42	0.66
1:K:135:HIS:CD2	1:K:136:TYR:H	2.13	0.66
1:I:119:PRO:O	1:I:123:GLN:CB	2.42	0.66
1:L:241:GLU:O	1:L:242:GLU:HB3	1.95	0.66
1:D:119:PRO:O	1:D:123:GLN:CB	2.42	0.66
1:D:606:SER:HB3	1:F:532:GLU:HB2	1.77	0.66
1:I:606:SER:HB3	1:K:532:GLU:HB2	1.77	0.66
1:J:241:GLU:O	1:J:242:GLU:HB3	1.95	0.66
1:N:241:GLU:O	1:N:242:GLU:HB3	1.95	0.66
1:L:135:HIS:CD2	1:L:136:TYR:H	2.13	0.66
1:M:112:VAL:O	1:M:140:ASN:ND2	2.21	0.66
1:E:324:GLN:HE21	1:F:513:LEU:H	1.43	0.66
1:C:606:SER:HB3	1:E:532:GLU:HB2	1.77	0.66
1:A:606:SER:HB3	1:C:532:GLU:HB2	1.77	0.66
1:B:135:HIS:CD2	1:B:136:TYR:H	2.13	0.66
1:E:606:SER:HB3	1:G:532:GLU:HB2	1.77	0.66
1:H:606:SER:HB3	1:J:532:GLU:HB2	1.77	0.66
1:B:119:PRO:O	1:B:123:GLN:CB	2.42	0.66
1:C:241:GLU:O	1:C:242:GLU:HB3	1.95	0.66
1:F:241:GLU:O	1:F:242:GLU:HB3	1.95	0.66
1:G:241:GLU:O	1:G:242:GLU:HB3	1.95	0.66
1:H:241:GLU:O	1:H:242:GLU:HB3	1.95	0.65
1:L:114:ALA:N	1:L:140:ASN:OD1	2.28	0.65
1:N:324:GLN:HE21	1:O:513:LEU:H	1.43	0.65
1:O:119:PRO:O	1:O:123:GLN:CB	2.42	0.65
1:D:212:ARG:NH2	1:E:177:ALA:HB1	2.10	0.65
1:E:242:GLU:HG2	1:E:243:GLY:H	1.60	0.65
1:C:324:GLN:HE21	1:D:513:LEU:H	1.43	0.65
1:E:241:GLU:O	1:E:242:GLU:HB3	1.95	0.65
1:L:324:GLN:HE21	1:M:513:LEU:H	1.43	0.65
1:M:241:GLU:O	1:M:242:GLU:HB3	1.95	0.65
1:B:606:SER:HB3	1:D:532:GLU:HB2	1.77	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:212:ARG:NH2	1:D:177:ALA:HB1	2.10	0.65
1:E:212:ARG:NH2	1:F:177:ALA:HB1	2.10	0.65
1:G:324:GLN:HE21	1:H:513:LEU:H	1.43	0.65
1:C:242:GLU:HG2	1:C:243:GLY:H	1.60	0.65
1:G:606:SER:HB3	1:I:532:GLU:HB2	1.77	0.65
1:M:324:GLN:HE21	1:N:513:LEU:H	1.43	0.65
1:O:179:ALA:HB1	1:O:209:ALA:HB1	1.79	0.65
1:F:606:SER:HB3	1:H:532:GLU:HB2	1.77	0.65
1:I:324:GLN:HE21	1:J:513:LEU:H	1.43	0.65
1:K:114:ALA:N	1:K:140:ASN:OD1	2.28	0.65
1:L:179:ALA:HB1	1:L:209:ALA:HB1	1.79	0.65
1:M:179:ALA:HB1	1:M:209:ALA:HB1	1.79	0.65
1:G:119:PRO:O	1:G:123:GLN:CB	2.42	0.65
1:I:241:GLU:O	1:I:242:GLU:HB3	1.95	0.65
1:N:179:ALA:HB1	1:N:209:ALA:HB1	1.79	0.65
1:A:324:GLN:HE21	1:B:513:LEU:H	1.43	0.65
1:J:114:ALA:N	1:J:140:ASN:OD1	2.28	0.65
1:J:179:ALA:HB1	1:J:209:ALA:HB1	1.79	0.65
1:B:212:ARG:NH2	1:C:177:ALA:HB1	2.10	0.64
1:D:241:GLU:O	1:D:242:GLU:HB3	1.95	0.64
1:E:181:ASP:O	1:E:184:GLU:HG2	1.98	0.64
1:F:181:ASP:O	1:F:184:GLU:HG2	1.98	0.64
1:F:212:ARG:NH2	1:G:177:ALA:HB1	2.10	0.64
1:F:324:GLN:HE21	1:G:513:LEU:H	1.43	0.64
1:K:179:ALA:HB1	1:K:209:ALA:HB1	1.79	0.64
1:K:181:ASP:O	1:K:184:GLU:HG2	1.98	0.64
1:A:179:ALA:HB1	1:A:209:ALA:HB1	1.79	0.64
1:C:114:ALA:N	1:C:140:ASN:OD1	2.28	0.64
1:E:114:ALA:N	1:E:140:ASN:OD1	2.28	0.64
1:F:114:ALA:N	1:F:140:ASN:OD1	2.28	0.64
1:I:114:ALA:N	1:I:140:ASN:OD1	2.28	0.64
1:B:179:ALA:HB1	1:B:209:ALA:HB1	1.79	0.64
1:D:114:ALA:N	1:D:140:ASN:OD1	2.28	0.64
1:G:114:ALA:N	1:G:140:ASN:OD1	2.28	0.64
1:G:181:ASP:O	1:G:184:GLU:HG2	1.98	0.64
1:H:114:ALA:N	1:H:140:ASN:OD1	2.28	0.64
1:J:181:ASP:O	1:J:184:GLU:HG2	1.98	0.64
1:K:324:GLN:HE21	1:L:513:LEU:H	1.43	0.64
1:B:114:ALA:N	1:B:140:ASN:OD1	2.28	0.64
1:D:181:ASP:O	1:D:184:GLU:HG2	1.98	0.64
1:M:181:ASP:O	1:M:184:GLU:HG2	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:181:ASP:O	1:N:184:GLU:HG2	1.97	0.64
1:A:210:ASP:O	1:A:214:ASN:CA	2.36	0.64
1:C:179:ALA:HB1	1:C:209:ALA:HB1	1.79	0.64
1:K:213:THR:HG22	1:L:239:GLU:HB3	1.79	0.64
1:K:259:VAL:CG2	1:K:292:ALA:HB2	2.28	0.64
1:L:181:ASP:O	1:L:184:GLU:HG2	1.98	0.64
1:A:114:ALA:N	1:A:140:ASN:OD1	2.28	0.64
1:A:181:ASP:O	1:A:184:GLU:HG2	1.98	0.64
1:J:213:THR:HG22	1:K:239:GLU:HB3	1.79	0.64
1:J:324:GLN:HE21	1:K:513:LEU:H	1.43	0.64
1:N:259:VAL:CG2	1:N:292:ALA:HB2	2.28	0.64
1:B:213:THR:HG22	1:C:239:GLU:HB3	1.79	0.64
1:E:213:THR:HG22	1:F:239:GLU:HB3	1.79	0.64
1:I:179:ALA:HB1	1:I:209:ALA:HB1	1.79	0.64
1:M:242:GLU:HG2	1:M:243:GLY:H	1.60	0.64
1:A:206:LYS:HZ1	1:B:190:ILE:HB	1.62	0.64
1:C:170:GLN:HB3	1:C:225:ARG:HH12	1.63	0.64
1:C:181:ASP:O	1:C:184:GLU:HG2	1.98	0.64
1:D:170:GLN:HB3	1:D:225:ARG:HH12	1.63	0.64
1:I:181:ASP:O	1:I:184:GLU:HG2	1.98	0.64
1:J:259:VAL:CG2	1:J:292:ALA:HB2	2.28	0.64
1:L:213:THR:HG22	1:M:239:GLU:HB3	1.79	0.64
1:O:181:ASP:O	1:O:184:GLU:HG2	1.98	0.64
1:A:213:THR:HG22	1:B:239:GLU:HB3	1.79	0.64
1:G:479:ALA:HA	1:G:515:LYS:HA	1.80	0.64
1:H:181:ASP:O	1:H:184:GLU:HG2	1.98	0.64
1:H:259:VAL:CG2	1:H:292:ALA:HB2	2.28	0.64
1:I:213:THR:HG22	1:J:239:GLU:HB3	1.79	0.64
1:I:259:VAL:CG2	1:I:292:ALA:HB2	2.28	0.64
1:J:479:ALA:HA	1:J:515:LYS:HA	1.80	0.64
1:K:479:ALA:HA	1:K:515:LYS:HA	1.80	0.64
1:L:259:VAL:CG2	1:L:292:ALA:HB2	2.28	0.64
1:O:114:ALA:N	1:O:140:ASN:OD1	2.28	0.64
1:D:213:THR:HG22	1:E:239:GLU:HB3	1.79	0.64
1:F:479:ALA:HA	1:F:515:LYS:HA	1.80	0.64
1:K:426:VAL:O	1:L:510:ASN:ND2	2.27	0.64
1:A:190:ILE:HB	1:O:206:LYS:HZ1	1.63	0.63
1:A:212:ARG:NH2	1:B:177:ALA:HB1	2.10	0.63
1:G:259:VAL:CG2	1:G:292:ALA:HB2	2.28	0.63
1:M:479:ALA:HA	1:M:515:LYS:HA	1.80	0.63
1:N:119:PRO:O	1:N:123:GLN:CB	2.42	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:181:ASP:O	1:B:184:GLU:HG2	1.98	0.63
1:G:212:ARG:NH2	1:H:177:ALA:HB1	2.10	0.63
1:H:479:ALA:HA	1:H:515:LYS:HA	1.80	0.63
1:I:479:ALA:HA	1:I:515:LYS:HA	1.80	0.63
1:M:259:VAL:CG2	1:M:292:ALA:HB2	2.28	0.63
1:B:259:VAL:CG2	1:B:292:ALA:HB2	2.28	0.63
1:H:206:LYS:HZ1	1:I:190:ILE:HB	1.62	0.63
1:L:354:GLN:HE21	1:L:356:PHE:HE2	1.47	0.63
1:M:213:THR:HG22	1:N:239:GLU:HB3	1.79	0.63
1:B:170:GLN:HB3	1:B:225:ARG:HH12	1.63	0.63
1:C:206:LYS:HZ1	1:D:190:ILE:HB	1.62	0.63
1:F:206:LYS:HZ1	1:G:190:ILE:HB	1.62	0.63
1:G:179:ALA:HB1	1:G:209:ALA:HB1	1.79	0.63
1:H:179:ALA:HB1	1:H:209:ALA:HB1	1.79	0.63
1:L:426:VAL:O	1:M:510:ASN:ND2	2.27	0.63
1:N:170:GLN:HB3	1:N:225:ARG:HH12	1.63	0.63
1:N:479:ALA:HA	1:N:515:LYS:HA	1.80	0.63
1:O:259:VAL:CG2	1:O:292:ALA:HB2	2.28	0.63
1:B:206:LYS:HZ1	1:C:190:ILE:HB	1.63	0.63
1:C:213:THR:HG22	1:D:239:GLU:HB3	1.79	0.63
1:D:179:ALA:HB1	1:D:209:ALA:HB1	1.79	0.63
1:D:479:ALA:HA	1:D:515:LYS:HA	1.80	0.63
1:E:259:VAL:CG2	1:E:292:ALA:HB2	2.28	0.63
1:E:479:ALA:HA	1:E:515:LYS:HA	1.81	0.63
1:F:213:THR:HG22	1:G:239:GLU:HB3	1.79	0.63
1:G:206:LYS:HZ1	1:H:190:ILE:HB	1.63	0.63
1:H:213:THR:HG22	1:I:239:GLU:HB3	1.79	0.63
1:J:242:GLU:HG2	1:J:243:GLY:H	1.60	0.63
1:L:206:LYS:HZ1	1:M:190:ILE:HB	1.62	0.63
1:N:354:GLN:HE21	1:N:356:PHE:HE2	1.47	0.63
1:D:354:GLN:HE21	1:D:356:PHE:HE2	1.47	0.63
1:I:354:GLN:HE21	1:I:356:PHE:HE2	1.47	0.63
1:K:206:LYS:HZ1	1:L:190:ILE:HB	1.63	0.63
1:L:119:PRO:O	1:L:123:GLN:CB	2.42	0.63
1:L:479:ALA:HA	1:L:515:LYS:HA	1.80	0.63
1:N:114:ALA:N	1:N:140:ASN:OD1	2.28	0.63
1:O:170:GLN:HB3	1:O:225:ARG:HH12	1.63	0.63
1:A:177:ALA:HB1	1:O:212:ARG:NH2	2.10	0.63
1:A:259:VAL:CG2	1:A:292:ALA:HB2	2.28	0.63
1:A:354:GLN:HE21	1:A:356:PHE:HE2	1.47	0.63
1:C:479:ALA:HA	1:C:515:LYS:HA	1.80	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:179:ALA:HB1	1:E:209:ALA:HB1	1.79	0.63
1:F:179:ALA:HB1	1:F:209:ALA:HB1	1.79	0.63
1:F:354:GLN:HE21	1:F:356:PHE:HE2	1.47	0.63
1:M:170:GLN:HB3	1:M:225:ARG:HH12	1.63	0.63
1:N:206:LYS:HZ1	1:O:190:ILE:HB	1.63	0.63
1:A:479:ALA:HA	1:A:515:LYS:HA	1.80	0.63
1:G:117:LEU:HD22	1:G:120:LEU:HD13	1.81	0.63
1:I:117:LEU:HD22	1:I:120:LEU:HD13	1.81	0.63
1:M:119:PRO:O	1:M:123:GLN:CB	2.42	0.63
1:C:354:GLN:HE21	1:C:356:PHE:HE2	1.47	0.63
1:L:170:GLN:HB3	1:L:225:ARG:HH12	1.63	0.63
1:M:426:VAL:O	1:N:510:ASN:ND2	2.27	0.63
1:N:212:ARG:NH2	1:O:177:ALA:HB1	2.10	0.63
1:N:213:THR:HG22	1:O:239:GLU:HB3	1.79	0.63
1:D:259:VAL:CG2	1:D:292:ALA:HB2	2.28	0.62
1:E:206:LYS:HZ1	1:F:190:ILE:HB	1.63	0.62
1:H:324:GLN:HE21	1:I:513:LEU:H	1.43	0.62
1:I:170:GLN:HB3	1:I:225:ARG:HH12	1.63	0.62
1:D:206:LYS:HZ1	1:E:190:ILE:HB	1.62	0.62
1:E:117:LEU:HD22	1:E:120:LEU:HD13	1.81	0.62
1:G:213:THR:HG22	1:H:239:GLU:HB3	1.79	0.62
1:H:170:GLN:HB3	1:H:225:ARG:HH12	1.63	0.62
1:O:479:ALA:HA	1:O:515:LYS:HA	1.80	0.62
1:A:170:GLN:HB3	1:A:225:ARG:HH12	1.63	0.62
1:A:239:GLU:HB3	1:O:213:THR:HG22	1.79	0.62
1:B:479:ALA:HA	1:B:515:LYS:HA	1.80	0.62
1:H:242:GLU:HG2	1:H:243:GLY:H	1.60	0.62
1:J:119:PRO:O	1:J:123:GLN:CB	2.42	0.62
1:N:426:VAL:O	1:O:510:ASN:ND2	2.27	0.62
1:G:354:GLN:HE21	1:G:356:PHE:HE2	1.47	0.62
1:J:354:GLN:HE21	1:J:356:PHE:HE2	1.47	0.62
1:K:170:GLN:HB3	1:K:225:ARG:HH12	1.63	0.62
1:M:212:ARG:NH2	1:N:177:ALA:HB1	2.10	0.62
1:C:259:VAL:CG2	1:C:292:ALA:HB2	2.28	0.62
1:K:117:LEU:HD22	1:K:120:LEU:HD13	1.81	0.62
1:K:168:LYS:N	1:K:220:GLY:O	2.30	0.62
1:L:153:LYS:O	1:L:157:VAL:HG23	2.00	0.62
1:M:153:LYS:O	1:M:157:VAL:HG23	2.00	0.62
1:B:354:GLN:HE21	1:B:356:PHE:HE2	1.47	0.62
1:G:266:SER:HB3	1:G:290:ILE:CD1	2.30	0.62
1:K:153:LYS:O	1:K:157:VAL:HG23	2.00	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:121:LEU:HD22	1:L:154:LEU:HD23	1.82	0.62
1:H:212:ARG:NH2	1:I:177:ALA:HB1	2.10	0.62
1:J:170:GLN:HB3	1:J:225:ARG:HH12	1.63	0.62
1:K:121:LEU:HD22	1:K:154:LEU:HD23	1.82	0.62
1:L:266:SER:HB3	1:L:290:ILE:CD1	2.30	0.62
1:M:114:ALA:N	1:M:140:ASN:OD1	2.28	0.62
1:M:121:LEU:HD22	1:M:154:LEU:HD23	1.82	0.62
1:J:121:LEU:HD22	1:J:154:LEU:HD23	1.82	0.62
1:J:266:SER:HB3	1:J:290:ILE:CD1	2.30	0.62
1:A:317:ARG:O	1:O:249:TYR:OH	2.17	0.62
1:F:259:VAL:CG2	1:F:292:ALA:HB2	2.28	0.62
1:F:266:SER:HB3	1:F:290:ILE:CD1	2.30	0.62
1:H:266:SER:HB3	1:H:290:ILE:CD1	2.30	0.62
1:J:153:LYS:O	1:J:157:VAL:HG23	2.00	0.62
1:K:354:GLN:HE21	1:K:356:PHE:HE2	1.47	0.62
1:L:212:ARG:NH2	1:M:177:ALA:HB1	2.10	0.62
1:M:266:SER:HB3	1:M:290:ILE:CD1	2.30	0.62
1:N:121:LEU:HD22	1:N:154:LEU:HD23	1.82	0.62
1:N:249:TYR:OH	1:O:317:ARG:O	2.17	0.62
1:O:354:GLN:HE21	1:O:356:PHE:HE2	1.47	0.62
1:C:117:LEU:HD22	1:C:120:LEU:HD13	1.81	0.62
1:D:153:LYS:O	1:D:157:VAL:HG23	2.00	0.62
1:E:153:LYS:O	1:E:157:VAL:HG23	2.00	0.62
1:F:153:LYS:O	1:F:157:VAL:HG23	2.00	0.62
1:G:170:GLN:HB3	1:G:225:ARG:HH12	1.63	0.62
1:H:121:LEU:HD22	1:H:154:LEU:HD23	1.82	0.62
1:N:153:LYS:O	1:N:157:VAL:HG23	2.00	0.62
1:A:119:PRO:O	1:A:123:GLN:CB	2.42	0.61
1:B:153:LYS:O	1:B:157:VAL:HG23	2.00	0.61
1:C:153:LYS:O	1:C:157:VAL:HG23	2.00	0.61
1:I:121:LEU:HD22	1:I:154:LEU:HD23	1.82	0.61
1:K:266:SER:HB3	1:K:290:ILE:CD1	2.30	0.61
1:N:266:SER:HB3	1:N:290:ILE:CD1	2.30	0.61
1:B:176:TYR:CD1	1:B:238:GLU:OE1	2.54	0.61
1:F:170:GLN:HB3	1:F:225:ARG:HH12	1.63	0.61
1:G:121:LEU:HD22	1:G:154:LEU:HD23	1.82	0.61
1:H:153:LYS:O	1:H:157:VAL:HG23	2.00	0.61
1:J:206:LYS:HZ1	1:K:190:ILE:HB	1.63	0.61
1:K:212:ARG:NH2	1:L:177:ALA:HB1	2.10	0.61
1:M:176:TYR:CD1	1:M:238:GLU:OE1	2.53	0.61
1:F:246:ARG:NH1	1:F:308:GLN:HG3	2.16	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:153:LYS:O	1:G:157:VAL:HG23	2.00	0.61
1:H:246:ARG:NH1	1:H:308:GLN:HG3	2.16	0.61
1:M:206:LYS:HZ1	1:N:190:ILE:HB	1.64	0.61
1:A:153:LYS:O	1:A:157:VAL:HG23	2.00	0.61
1:A:176:TYR:CD1	1:A:238:GLU:OE1	2.54	0.61
1:D:246:ARG:NH1	1:D:308:GLN:HG3	2.16	0.61
1:E:266:SER:HB3	1:E:290:ILE:CD1	2.30	0.61
1:G:246:ARG:NH1	1:G:308:GLN:HG3	2.16	0.61
1:I:212:ARG:NH2	1:J:177:ALA:HB1	2.10	0.61
1:I:249:TYR:OH	1:J:317:ARG:O	2.17	0.61
1:I:266:SER:HB3	1:I:290:ILE:CD1	2.30	0.61
1:J:212:ARG:NH2	1:K:177:ALA:HB1	2.10	0.61
1:M:354:GLN:HE21	1:M:356:PHE:HE2	1.47	0.61
1:E:168:LYS:N	1:E:220:GLY:O	2.31	0.61
1:E:246:ARG:NH1	1:E:308:GLN:HG3	2.16	0.61
1:I:153:LYS:O	1:I:157:VAL:HG23	2.00	0.61
1:K:242:GLU:HG2	1:K:243:GLY:H	1.60	0.61
1:D:168:LYS:N	1:D:220:GLY:O	2.31	0.61
1:F:117:LEU:HD22	1:F:120:LEU:HD13	1.81	0.61
1:F:121:LEU:HD22	1:F:154:LEU:HD23	1.82	0.61
1:F:168:LYS:N	1:F:220:GLY:O	2.30	0.61
1:H:176:TYR:CD1	1:H:238:GLU:OE1	2.54	0.61
1:J:168:LYS:N	1:J:220:GLY:O	2.31	0.61
1:M:117:LEU:HD22	1:M:120:LEU:HD13	1.81	0.61
1:O:117:LEU:HD22	1:O:120:LEU:HD13	1.81	0.61
1:O:121:LEU:HD22	1:O:154:LEU:HD23	1.82	0.61
1:D:266:SER:HB3	1:D:290:ILE:CD1	2.30	0.61
1:D:426:VAL:O	1:E:510:ASN:ND2	2.27	0.61
1:E:170:GLN:HB3	1:E:225:ARG:HH12	1.63	0.61
1:F:479:ALA:O	1:F:480:VAL:HG13	2.01	0.61
1:G:176:TYR:CD1	1:G:238:GLU:OE1	2.54	0.61
1:I:246:ARG:NH1	1:I:308:GLN:HG3	2.16	0.61
1:J:246:ARG:NH1	1:J:308:GLN:HG3	2.16	0.61
1:L:479:ALA:O	1:L:480:VAL:HG13	2.01	0.61
1:M:249:TYR:OH	1:N:317:ARG:O	2.17	0.61
1:A:121:LEU:HD22	1:A:154:LEU:HD23	1.82	0.61
1:C:176:TYR:CD1	1:C:238:GLU:OE1	2.54	0.61
1:H:117:LEU:HD22	1:H:120:LEU:HD13	1.81	0.61
1:L:176:TYR:CD1	1:L:238:GLU:OE1	2.54	0.61
1:N:117:LEU:HD22	1:N:120:LEU:HD13	1.81	0.61
1:N:176:TYR:CD1	1:N:238:GLU:OE1	2.54	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:117:LEU:HD22	1:A:120:LEU:HD13	1.81	0.61
1:A:266:SER:HB3	1:A:290:ILE:CD1	2.30	0.61
1:B:266:SER:HB3	1:B:290:ILE:CD1	2.30	0.61
1:D:120:LEU:HD22	1:D:157:VAL:HG11	1.83	0.61
1:D:479:ALA:O	1:D:480:VAL:HG13	2.01	0.61
1:E:354:GLN:HE21	1:E:356:PHE:HE2	1.47	0.61
1:H:354:GLN:HE21	1:H:356:PHE:HE2	1.47	0.61
1:H:479:ALA:O	1:H:480:VAL:HG13	2.01	0.61
1:J:479:ALA:O	1:J:480:VAL:HG13	2.01	0.61
1:N:479:ALA:O	1:N:480:VAL:HG13	2.01	0.61
1:B:246:ARG:NH1	1:B:308:GLN:HG3	2.16	0.61
1:B:249:TYR:OH	1:C:317:ARG:O	2.17	0.61
1:C:168:LYS:N	1:C:220:GLY:O	2.31	0.61
1:C:266:SER:HB3	1:C:290:ILE:CD1	2.30	0.61
1:G:168:LYS:N	1:G:220:GLY:O	2.31	0.61
1:O:266:SER:HB3	1:O:290:ILE:CD1	2.30	0.61
1:C:246:ARG:NH1	1:C:308:GLN:HG3	2.16	0.60
1:D:117:LEU:HD22	1:D:120:LEU:HD13	1.81	0.60
1:E:176:TYR:CD1	1:E:238:GLU:OE1	2.54	0.60
1:G:120:LEU:HD22	1:G:157:VAL:HG11	1.83	0.60
1:I:206:LYS:HZ1	1:J:190:ILE:HB	1.65	0.60
1:K:176:TYR:CD1	1:K:238:GLU:OE1	2.54	0.60
1:O:120:LEU:HD22	1:O:157:VAL:HG11	1.83	0.60
1:O:153:LYS:O	1:O:157:VAL:HG23	2.00	0.60
1:O:176:TYR:CD1	1:O:238:GLU:OE1	2.54	0.60
1:B:121:LEU:HD22	1:B:154:LEU:HD23	1.82	0.60
1:B:479:ALA:O	1:B:480:VAL:HG13	2.01	0.60
1:C:120:LEU:HD22	1:C:157:VAL:HG11	1.83	0.60
1:C:479:ALA:O	1:C:480:VAL:HG13	2.01	0.60
1:E:121:LEU:HD22	1:E:154:LEU:HD23	1.82	0.60
1:H:120:LEU:HD22	1:H:157:VAL:HG11	1.83	0.60
1:K:246:ARG:NH1	1:K:308:GLN:HG3	2.16	0.60
1:A:426:VAL:O	1:B:510:ASN:ND2	2.27	0.60
1:D:121:LEU:HD22	1:D:154:LEU:HD23	1.82	0.60
1:E:249:TYR:OH	1:F:317:ARG:O	2.17	0.60
1:E:295:GLN:CG	1:F:257:ASN:ND2	2.65	0.60
1:I:295:GLN:CG	1:J:257:ASN:ND2	2.65	0.60
1:J:117:LEU:HD22	1:J:120:LEU:HD13	1.81	0.60
1:J:223:LYS:HA	1:J:226:GLN:HG2	1.83	0.60
1:A:120:LEU:HD22	1:A:157:VAL:HG11	1.83	0.60
1:A:295:GLN:CG	1:B:257:ASN:ND2	2.65	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:479:ALA:O	1:A:480:VAL:HG13	2.01	0.60
1:D:295:GLN:CG	1:E:257:ASN:ND2	2.65	0.60
1:E:167:GLU:HA	1:E:221:PRO:HA	1.84	0.60
1:E:426:VAL:O	1:F:510:ASN:ND2	2.27	0.60
1:F:176:TYR:CD1	1:F:238:GLU:OE1	2.54	0.60
1:F:223:LYS:HA	1:F:226:GLN:HG2	1.83	0.60
1:I:176:TYR:CD1	1:I:238:GLU:OE1	2.54	0.60
1:I:223:LYS:HA	1:I:226:GLN:HG2	1.83	0.60
1:J:176:TYR:CD1	1:J:238:GLU:OE1	2.54	0.60
1:J:295:GLN:CG	1:K:257:ASN:ND2	2.65	0.60
1:L:117:LEU:HD22	1:L:120:LEU:HD13	1.81	0.60
1:D:176:TYR:CD1	1:D:238:GLU:OE1	2.54	0.60
1:E:223:LYS:HA	1:E:226:GLN:HG2	1.83	0.60
1:F:167:GLU:HA	1:F:221:PRO:HA	1.84	0.60
1:F:295:GLN:CG	1:G:257:ASN:ND2	2.65	0.60
1:K:223:LYS:HA	1:K:226:GLN:HG2	1.83	0.60
1:M:295:GLN:CG	1:N:257:ASN:ND2	2.65	0.60
1:A:510:ASN:ND2	1:O:426:VAL:O	2.27	0.60
1:D:167:GLU:HA	1:D:221:PRO:HA	1.84	0.60
1:E:355:GLN:HB2	1:E:363:ILE:HG22	1.84	0.60
1:G:167:GLU:HA	1:G:221:PRO:HA	1.84	0.60
1:G:223:LYS:HA	1:G:226:GLN:HG2	1.83	0.60
1:L:246:ARG:NH1	1:L:308:GLN:HG3	2.16	0.60
1:L:295:GLN:CG	1:M:257:ASN:ND2	2.65	0.60
1:N:120:LEU:HD22	1:N:157:VAL:HG11	1.83	0.60
1:B:117:LEU:HD22	1:B:120:LEU:HD13	1.81	0.60
1:B:295:GLN:CG	1:C:257:ASN:ND2	2.65	0.60
1:E:120:LEU:HD22	1:E:157:VAL:HG11	1.83	0.60
1:H:223:LYS:HA	1:H:226:GLN:HG2	1.84	0.60
1:H:295:GLN:CG	1:I:257:ASN:ND2	2.65	0.60
1:I:167:GLU:HA	1:I:221:PRO:HA	1.84	0.60
1:J:167:GLU:HA	1:J:221:PRO:HA	1.84	0.60
1:A:168:LYS:N	1:A:220:GLY:O	2.30	0.60
1:B:426:VAL:O	1:C:510:ASN:ND2	2.27	0.60
1:C:121:LEU:HD22	1:C:154:LEU:HD23	1.82	0.60
1:C:167:GLU:HA	1:C:221:PRO:HA	1.84	0.60
1:I:120:LEU:HD22	1:I:157:VAL:HG11	1.83	0.60
1:K:295:GLN:CG	1:L:257:ASN:ND2	2.65	0.60
1:A:246:ARG:NH1	1:A:308:GLN:HG3	2.16	0.60
1:B:168:LYS:N	1:B:220:GLY:O	2.31	0.60
1:D:223:LYS:HA	1:D:226:GLN:HG2	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:167:GLU:HA	1:H:221:PRO:HA	1.84	0.60
1:I:479:ALA:O	1:I:480:VAL:HG13	2.01	0.60
1:K:167:GLU:HA	1:K:221:PRO:HA	1.84	0.60
1:N:246:ARG:NH1	1:N:308:GLN:HG3	2.16	0.60
1:O:479:ALA:O	1:O:480:VAL:HG13	2.01	0.60
1:D:355:GLN:HB2	1:D:363:ILE:HG22	1.84	0.60
1:F:120:LEU:HD22	1:F:157:VAL:HG11	1.83	0.60
1:F:355:GLN:HB2	1:F:363:ILE:HG22	1.84	0.60
1:L:223:LYS:HA	1:L:226:GLN:HG2	1.84	0.60
1:M:246:ARG:NH1	1:M:308:GLN:HG3	2.16	0.60
1:A:257:ASN:ND2	1:O:295:GLN:CG	2.65	0.59
1:I:168:LYS:N	1:I:220:GLY:O	2.31	0.59
1:K:479:ALA:O	1:K:480:VAL:HG13	2.01	0.59
1:M:479:ALA:O	1:M:480:VAL:HG13	2.01	0.59
1:B:355:GLN:HB2	1:B:363:ILE:HG22	1.84	0.59
1:C:295:GLN:CG	1:D:257:ASN:ND2	2.65	0.59
1:C:355:GLN:HB2	1:C:363:ILE:HG22	1.84	0.59
1:H:355:GLN:HB2	1:H:363:ILE:HG22	1.84	0.59
1:L:167:GLU:HA	1:L:221:PRO:HA	1.84	0.59
1:M:120:LEU:HD22	1:M:157:VAL:HG11	1.83	0.59
1:N:295:GLN:CG	1:O:257:ASN:ND2	2.65	0.59
1:O:246:ARG:NH1	1:O:308:GLN:HG3	2.16	0.59
1:E:479:ALA:O	1:E:480:VAL:HG13	2.01	0.59
1:F:426:VAL:O	1:G:510:ASN:ND2	2.27	0.59
1:I:355:GLN:HB2	1:I:363:ILE:HG22	1.84	0.59
1:B:167:GLU:HA	1:B:221:PRO:HA	1.84	0.59
1:C:157:VAL:HG22	1:C:160:ARG:NH2	2.18	0.59
1:C:168:LYS:HD2	1:C:169:GLN:H	1.68	0.59
1:G:355:GLN:HB2	1:G:363:ILE:HG22	1.84	0.59
1:G:479:ALA:O	1:G:480:VAL:HG13	2.01	0.59
1:N:210:ASP:HB3	1:N:215:SER:H	1.68	0.59
1:A:210:ASP:HB3	1:A:215:SER:H	1.68	0.59
1:A:355:GLN:HB2	1:A:363:ILE:HG22	1.84	0.59
1:K:157:VAL:HG22	1:K:160:ARG:NH2	2.18	0.59
1:L:249:TYR:OH	1:M:317:ARG:O	2.17	0.59
1:M:167:GLU:HA	1:M:221:PRO:HA	1.84	0.59
1:O:157:VAL:HG22	1:O:160:ARG:NH2	2.18	0.59
1:B:120:LEU:HD22	1:B:157:VAL:HG11	1.83	0.59
1:B:168:LYS:HD2	1:B:169:GLN:H	1.68	0.59
1:D:168:LYS:HD2	1:D:169:GLN:H	1.68	0.59
1:H:157:VAL:HG22	1:H:160:ARG:NH2	2.18	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:120:LEU:HD22	1:J:157:VAL:HG11	1.83	0.59
1:L:168:LYS:HD2	1:L:169:GLN:H	1.68	0.59
1:M:168:LYS:HD2	1:M:169:GLN:H	1.68	0.59
1:M:210:ASP:HB3	1:M:215:SER:H	1.68	0.59
1:O:210:ASP:HB3	1:O:215:SER:H	1.68	0.59
1:C:223:LYS:HA	1:C:226:GLN:HG2	1.83	0.59
1:D:372:ASP:OD2	1:D:383:ASN:ND2	2.36	0.59
1:E:372:ASP:OD2	1:E:383:ASN:ND2	2.36	0.59
1:F:157:VAL:HG22	1:F:160:ARG:NH2	2.18	0.59
1:G:295:GLN:CG	1:H:257:ASN:ND2	2.65	0.59
1:I:157:VAL:HG22	1:I:160:ARG:NH2	2.18	0.59
1:M:223:LYS:HA	1:M:226:GLN:HG2	1.83	0.59
1:N:157:VAL:HG22	1:N:160:ARG:NH2	2.18	0.59
1:B:223:LYS:HA	1:B:226:GLN:HG2	1.83	0.59
1:K:168:LYS:HD2	1:K:169:GLN:H	1.68	0.59
1:L:120:LEU:HD22	1:L:157:VAL:HG11	1.83	0.59
1:N:167:GLU:HA	1:N:221:PRO:HA	1.84	0.59
1:N:223:LYS:HA	1:N:226:GLN:HG2	1.83	0.59
1:A:167:GLU:HA	1:A:221:PRO:HA	1.84	0.59
1:C:426:VAL:O	1:D:510:ASN:ND2	2.27	0.59
1:F:372:ASP:OD2	1:F:383:ASN:ND2	2.36	0.59
1:G:157:VAL:HG22	1:G:160:ARG:NH2	2.18	0.59
1:H:249:TYR:OH	1:I:317:ARG:O	2.17	0.59
1:L:157:VAL:HG22	1:L:160:ARG:NH2	2.18	0.59
1:L:210:ASP:HB3	1:L:215:SER:H	1.68	0.59
1:D:157:VAL:HG22	1:D:160:ARG:NH2	2.18	0.59
1:I:168:LYS:HD2	1:I:169:GLN:H	1.68	0.59
1:N:168:LYS:HD2	1:N:169:GLN:H	1.68	0.59
1:O:167:GLU:HA	1:O:221:PRO:HA	1.84	0.59
1:A:249:TYR:OH	1:B:317:ARG:O	2.17	0.58
1:B:157:VAL:HG22	1:B:160:ARG:NH2	2.18	0.58
1:B:210:ASP:HB3	1:B:215:SER:H	1.68	0.58
1:C:372:ASP:OD2	1:C:383:ASN:ND2	2.36	0.58
1:O:168:LYS:N	1:O:220:GLY:O	2.31	0.58
1:A:157:VAL:HG22	1:A:160:ARG:NH2	2.18	0.58
1:G:372:ASP:OD2	1:G:383:ASN:ND2	2.36	0.58
1:J:168:LYS:HD2	1:J:169:GLN:H	1.68	0.58
1:J:355:GLN:HB2	1:J:363:ILE:HG22	1.84	0.58
1:K:372:ASP:OD2	1:K:383:ASN:ND2	2.36	0.58
1:N:372:ASP:OD2	1:N:383:ASN:ND2	2.36	0.58
1:O:355:GLN:HB2	1:O:363:ILE:HG22	1.84	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:372:ASP:OD2	1:O:383:ASN:ND2	2.36	0.58
1:A:223:LYS:HA	1:A:226:GLN:HG2	1.83	0.58
1:E:168:LYS:HD2	1:E:169:GLN:H	1.68	0.58
1:G:168:LYS:HD2	1:G:169:GLN:H	1.68	0.58
1:J:157:VAL:HG22	1:J:160:ARG:NH2	2.18	0.58
1:J:372:ASP:OD2	1:J:383:ASN:ND2	2.36	0.58
1:K:355:GLN:HB2	1:K:363:ILE:HG22	1.84	0.58
1:O:223:LYS:HA	1:O:226:GLN:HG2	1.83	0.58
1:D:210:ASP:HB3	1:D:215:SER:H	1.68	0.58
1:K:120:LEU:HD22	1:K:157:VAL:HG11	1.83	0.58
1:N:355:GLN:HB2	1:N:363:ILE:HG22	1.84	0.58
1:A:372:ASP:OD2	1:A:383:ASN:ND2	2.36	0.58
1:A:479:ALA:O	1:A:570:ILE:HD11	2.04	0.58
1:K:210:ASP:HB3	1:K:215:SER:H	1.68	0.58
1:M:372:ASP:OD2	1:M:383:ASN:ND2	2.36	0.58
1:O:479:ALA:O	1:O:570:ILE:HD11	2.04	0.58
1:A:168:LYS:HD2	1:A:169:GLN:H	1.68	0.58
1:B:479:ALA:O	1:B:570:ILE:HD11	2.04	0.58
1:G:426:VAL:O	1:H:510:ASN:ND2	2.27	0.58
1:H:479:ALA:O	1:H:570:ILE:HD11	2.04	0.58
1:I:578:ARG:O	1:I:582:LYS:N	2.32	0.58
1:L:355:GLN:HB2	1:L:363:ILE:HG22	1.84	0.58
1:E:157:VAL:HG22	1:E:160:ARG:NH2	2.18	0.58
1:F:104:THR:OG1	1:F:144:LEU:O	2.21	0.58
1:J:210:ASP:HB3	1:J:215:SER:H	1.68	0.58
1:M:355:GLN:HB2	1:M:363:ILE:HG22	1.84	0.58
1:N:479:ALA:O	1:N:570:ILE:HD11	2.04	0.58
1:B:354:GLN:NE2	1:B:356:PHE:HE2	2.02	0.58
1:C:210:ASP:HB3	1:C:215:SER:H	1.68	0.58
1:D:479:ALA:O	1:D:570:ILE:HD11	2.04	0.58
1:G:578:ARG:O	1:G:582:LYS:N	2.32	0.58
1:H:372:ASP:OD2	1:H:383:ASN:ND2	2.36	0.58
1:I:372:ASP:OD2	1:I:383:ASN:ND2	2.36	0.58
1:C:354:GLN:NE2	1:C:356:PHE:HE2	2.02	0.58
1:C:479:ALA:O	1:C:570:ILE:HD11	2.04	0.58
1:E:479:ALA:O	1:E:570:ILE:HD11	2.04	0.58
1:H:168:LYS:HD2	1:H:169:GLN:H	1.68	0.58
1:H:168:LYS:N	1:H:220:GLY:O	2.31	0.58
1:M:157:VAL:HG22	1:M:160:ARG:NH2	2.18	0.58
1:M:479:ALA:O	1:M:570:ILE:HD11	2.04	0.58
1:A:578:ARG:O	1:A:582:LYS:N	2.32	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:354:GLN:NE2	1:D:356:PHE:HE2	2.02	0.58
1:I:479:ALA:O	1:I:570:ILE:HD11	2.04	0.58
1:L:372:ASP:OD2	1:L:383:ASN:ND2	2.36	0.58
1:O:168:LYS:HD2	1:O:169:GLN:H	1.68	0.58
1:B:372:ASP:OD2	1:B:383:ASN:ND2	2.36	0.57
1:E:210:ASP:HB3	1:E:215:SER:H	1.68	0.57
1:L:479:ALA:O	1:L:570:ILE:HD11	2.04	0.57
1:D:249:TYR:OH	1:E:317:ARG:O	2.17	0.57
1:G:210:ASP:HB3	1:G:215:SER:H	1.68	0.57
1:G:479:ALA:O	1:G:570:ILE:HD11	2.04	0.57
1:K:479:ALA:O	1:K:570:ILE:HD11	2.04	0.57
1:M:354:GLN:NE2	1:M:356:PHE:HE2	2.02	0.57
1:F:168:LYS:HD2	1:F:169:GLN:H	1.68	0.57
1:J:479:ALA:O	1:J:570:ILE:HD11	2.04	0.57
1:N:354:GLN:NE2	1:N:356:PHE:HE2	2.02	0.57
1:C:231:LEU:O	1:C:234:SER:HB2	2.05	0.57
1:E:231:LEU:O	1:E:234:SER:HB2	2.05	0.57
1:E:354:GLN:NE2	1:E:356:PHE:HE2	2.02	0.57
1:N:168:LYS:N	1:N:220:GLY:O	2.31	0.57
1:N:185:ILE:HG13	1:N:186:LEU:N	2.20	0.57
1:O:185:ILE:HG13	1:O:186:LEU:N	2.20	0.57
1:A:185:ILE:HG13	1:A:186:LEU:N	2.20	0.57
1:A:231:LEU:O	1:A:234:SER:HB2	2.05	0.57
1:B:231:LEU:O	1:B:234:SER:HB2	2.05	0.57
1:D:231:LEU:O	1:D:234:SER:HB2	2.05	0.57
1:F:231:LEU:O	1:F:234:SER:HB2	2.05	0.57
1:H:210:ASP:HB3	1:H:215:SER:H	1.68	0.57
1:M:231:LEU:O	1:M:234:SER:HB2	2.05	0.57
1:L:185:ILE:HG13	1:L:186:LEU:N	2.20	0.57
1:O:231:LEU:O	1:O:234:SER:HB2	2.05	0.57
1:A:354:GLN:NE2	1:A:356:PHE:HE2	2.02	0.57
1:F:479:ALA:O	1:F:570:ILE:HD11	2.04	0.57
1:I:104:THR:OG1	1:I:144:LEU:O	2.21	0.57
1:L:354:GLN:NE2	1:L:356:PHE:HE2	2.02	0.57
1:I:231:LEU:O	1:I:234:SER:HB2	2.05	0.57
1:J:185:ILE:HG13	1:J:186:LEU:N	2.20	0.57
1:J:231:LEU:O	1:J:234:SER:HB2	2.05	0.57
1:N:231:LEU:O	1:N:234:SER:HB2	2.05	0.57
1:O:354:GLN:NE2	1:O:356:PHE:HE2	2.02	0.57
1:G:176:TYR:CE1	1:G:238:GLU:OE1	2.58	0.57
1:J:176:TYR:CE1	1:J:238:GLU:OE1	2.58	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:176:TYR:CE1	1:L:238:GLU:OE1	2.58	0.57
1:C:185:ILE:HG13	1:C:186:LEU:N	2.20	0.56
1:F:176:TYR:CE1	1:F:238:GLU:OE1	2.58	0.56
1:I:210:ASP:HB3	1:I:215:SER:H	1.68	0.56
1:M:185:ILE:HG13	1:M:186:LEU:N	2.20	0.56
1:H:354:GLN:NE2	1:H:356:PHE:HE2	2.02	0.56
1:H:426:VAL:O	1:I:510:ASN:ND2	2.27	0.56
1:I:176:TYR:CE1	1:I:238:GLU:OE1	2.58	0.56
1:K:185:ILE:HG13	1:K:186:LEU:N	2.20	0.56
1:K:249:TYR:OH	1:L:317:ARG:O	2.17	0.56
1:A:344:VAL:HB	1:A:549:LEU:O	2.06	0.56
1:G:231:LEU:O	1:G:234:SER:HB2	2.05	0.56
1:H:176:TYR:CE1	1:H:238:GLU:OE1	2.58	0.56
1:H:185:ILE:HG13	1:H:186:LEU:N	2.20	0.56
1:I:354:GLN:NE2	1:I:356:PHE:HE2	2.02	0.56
1:K:176:TYR:CE1	1:K:238:GLU:OE1	2.58	0.56
1:L:231:LEU:O	1:L:234:SER:HB2	2.05	0.56
1:M:176:TYR:CE1	1:M:238:GLU:OE1	2.58	0.56
1:B:185:ILE:HG13	1:B:186:LEU:N	2.20	0.56
1:D:104:THR:OG1	1:D:144:LEU:O	2.21	0.56
1:F:210:ASP:HB3	1:F:215:SER:H	1.68	0.56
1:J:344:VAL:HB	1:J:549:LEU:O	2.06	0.56
1:L:578:ARG:O	1:L:582:LYS:N	2.32	0.56
1:N:344:VAL:HB	1:N:549:LEU:O	2.06	0.56
1:O:578:ARG:O	1:O:582:LYS:N	2.32	0.56
1:D:185:ILE:HG13	1:D:186:LEU:N	2.20	0.56
1:E:344:VAL:HB	1:E:549:LEU:O	2.06	0.56
1:F:185:ILE:HG13	1:F:186:LEU:N	2.20	0.56
1:F:354:GLN:NE2	1:F:356:PHE:HE2	2.02	0.56
1:H:344:VAL:HB	1:H:549:LEU:O	2.06	0.56
1:I:344:VAL:HB	1:I:549:LEU:O	2.06	0.56
1:K:344:VAL:HB	1:K:549:LEU:O	2.06	0.56
1:L:344:VAL:HB	1:L:549:LEU:O	2.06	0.56
1:M:344:VAL:HB	1:M:549:LEU:O	2.06	0.56
1:N:176:TYR:CE1	1:N:238:GLU:OE1	2.58	0.56
1:O:344:VAL:HB	1:O:549:LEU:O	2.06	0.56
1:C:258:LEU:CD2	1:C:318:LEU:HD23	2.36	0.56
1:H:231:LEU:O	1:H:234:SER:HB2	2.05	0.56
1:E:176:TYR:CE1	1:E:238:GLU:OE1	2.58	0.56
1:G:124:MET:HG2	1:G:154:LEU:HG	1.88	0.56
1:J:124:MET:HG2	1:J:154:LEU:HG	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:354:GLN:NE2	1:J:356:PHE:HE2	2.02	0.56
1:D:344:VAL:HB	1:D:549:LEU:O	2.06	0.56
1:E:161:VAL:HA	1:E:164:ILE:HG12	1.88	0.56
1:G:182:LEU:HD21	1:G:216:LEU:HD22	1.88	0.56
1:G:185:ILE:HG13	1:G:186:LEU:N	2.20	0.56
1:G:354:GLN:NE2	1:G:356:PHE:HE2	2.02	0.56
1:I:124:MET:HG2	1:I:154:LEU:HG	1.88	0.56
1:I:161:VAL:HA	1:I:164:ILE:HG12	1.88	0.56
1:I:182:LEU:HD21	1:I:216:LEU:HD22	1.88	0.56
1:J:182:LEU:HD21	1:J:216:LEU:HD22	1.88	0.56
1:K:231:LEU:O	1:K:234:SER:HB2	2.05	0.56
1:K:354:GLN:NE2	1:K:356:PHE:HE2	2.02	0.56
1:M:182:LEU:HD21	1:M:216:LEU:HD22	1.88	0.56
1:F:161:VAL:HA	1:F:164:ILE:HG12	1.88	0.56
1:F:182:LEU:HD21	1:F:216:LEU:HD22	1.88	0.56
1:G:344:VAL:HB	1:G:549:LEU:O	2.06	0.56
1:H:124:MET:HG2	1:H:154:LEU:HG	1.88	0.56
1:H:161:VAL:HA	1:H:164:ILE:HG12	1.88	0.56
1:I:185:ILE:HG13	1:I:186:LEU:N	2.20	0.56
1:J:249:TYR:OH	1:K:317:ARG:O	2.17	0.56
1:L:124:MET:HG2	1:L:154:LEU:HG	1.88	0.56
1:L:182:LEU:HD21	1:L:216:LEU:HD22	1.88	0.56
1:N:182:LEU:HD21	1:N:216:LEU:HD22	1.88	0.56
1:B:176:TYR:CE1	1:B:238:GLU:OE1	2.58	0.56
1:C:176:TYR:CE1	1:C:238:GLU:OE1	2.58	0.56
1:D:258:LEU:CD2	1:D:318:LEU:HD23	2.36	0.56
1:E:185:ILE:HG13	1:E:186:LEU:N	2.20	0.56
1:F:249:TYR:OH	1:G:317:ARG:O	2.17	0.56
1:G:249:TYR:OH	1:H:317:ARG:O	2.17	0.56
1:A:182:LEU:HD21	1:A:216:LEU:HD22	1.88	0.55
1:B:344:VAL:HB	1:B:549:LEU:O	2.06	0.55
1:E:603:LEU:HD13	1:G:551:ARG:NH1	2.21	0.55
1:F:344:VAL:HB	1:F:549:LEU:O	2.06	0.55
1:H:182:LEU:HD21	1:H:216:LEU:HD22	1.88	0.55
1:K:124:MET:HG2	1:K:154:LEU:HG	1.88	0.55
1:M:124:MET:HG2	1:M:154:LEU:HG	1.88	0.55
1:A:176:TYR:CE1	1:A:238:GLU:OE1	2.58	0.55
1:D:161:VAL:HA	1:D:164:ILE:HG12	1.88	0.55
1:D:182:LEU:HD21	1:D:216:LEU:HD22	1.88	0.55
1:J:161:VAL:HA	1:J:164:ILE:HG12	1.88	0.55
1:K:182:LEU:HD21	1:K:216:LEU:HD22	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:168:LYS:N	1:M:220:GLY:O	2.31	0.55
1:E:124:MET:HG2	1:E:154:LEU:HG	1.88	0.55
1:G:161:VAL:HA	1:G:164:ILE:HG12	1.88	0.55
1:O:182:LEU:HD21	1:O:216:LEU:HD22	1.88	0.55
1:C:161:VAL:HA	1:C:164:ILE:HG12	1.88	0.55
1:C:182:LEU:HD21	1:C:216:LEU:HD22	1.88	0.55
1:C:603:LEU:HD13	1:E:551:ARG:NH1	2.21	0.55
1:D:603:LEU:HD13	1:F:551:ARG:NH1	2.21	0.55
1:F:266:SER:HB3	1:F:290:ILE:HD11	1.89	0.55
1:G:603:LEU:HD13	1:I:551:ARG:NH1	2.21	0.55
1:B:161:VAL:HA	1:B:164:ILE:HG12	1.88	0.55
1:B:182:LEU:HD21	1:B:216:LEU:HD22	1.88	0.55
1:D:176:TYR:CE1	1:D:238:GLU:OE1	2.58	0.55
1:E:182:LEU:HD21	1:E:216:LEU:HD22	1.88	0.55
1:E:266:SER:HB3	1:E:290:ILE:HD11	1.89	0.55
1:J:105:ARG:NE	1:J:151:ILE:HD12	2.22	0.55
1:K:161:VAL:HA	1:K:164:ILE:HG12	1.88	0.55
1:L:161:VAL:HA	1:L:164:ILE:HG12	1.88	0.55
1:O:176:TYR:CE1	1:O:238:GLU:OE1	2.58	0.55
1:G:266:SER:HB3	1:G:290:ILE:HD11	1.89	0.55
1:M:603:LEU:HD13	1:O:551:ARG:NH1	2.21	0.55
1:D:124:MET:HG2	1:D:154:LEU:HG	1.88	0.55
1:F:124:MET:HG2	1:F:154:LEU:HG	1.88	0.55
1:J:151:ILE:O	1:J:155:ILE:HG12	2.07	0.55
1:K:153:LYS:HA	1:K:156:GLU:HB2	1.89	0.55
1:K:496:ASN:OD1	1:K:497:SER:N	2.40	0.55
1:K:603:LEU:HD13	1:M:551:ARG:NH1	2.21	0.55
1:A:551:ARG:NH1	1:N:603:LEU:HD13	2.21	0.55
1:B:151:ILE:O	1:B:155:ILE:HG12	2.07	0.55
1:B:603:LEU:HD13	1:D:551:ARG:NH1	2.21	0.55
1:C:344:VAL:HB	1:C:549:LEU:O	2.06	0.55
1:F:151:ILE:O	1:F:155:ILE:HG12	2.07	0.55
1:F:603:LEU:HD13	1:H:551:ARG:NH1	2.21	0.55
1:K:258:LEU:CD2	1:K:318:LEU:HD23	2.36	0.55
1:M:258:LEU:CD2	1:M:318:LEU:HD23	2.36	0.55
1:M:496:ASN:OD1	1:M:497:SER:N	2.40	0.55
1:O:266:SER:HB3	1:O:290:ILE:HD11	1.89	0.55
1:B:551:ARG:NH1	1:O:603:LEU:HD13	2.21	0.55
1:D:266:SER:HB3	1:D:290:ILE:HD11	1.89	0.55
1:I:426:VAL:O	1:J:510:ASN:ND2	2.27	0.55
1:J:258:LEU:CD2	1:J:318:LEU:HD23	2.36	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:496:ASN:OD1	1:J:497:SER:N	2.40	0.55
1:L:258:LEU:CD2	1:L:318:LEU:HD23	2.36	0.55
1:N:124:MET:HG2	1:N:154:LEU:HG	1.88	0.55
1:O:124:MET:HG2	1:O:154:LEU:HG	1.88	0.55
1:B:496:ASN:OD1	1:B:497:SER:N	2.40	0.55
1:F:496:ASN:OD1	1:F:497:SER:N	2.40	0.55
1:H:258:LEU:CD2	1:H:318:LEU:HD23	2.36	0.55
1:J:153:LYS:HA	1:J:156:GLU:HB2	1.89	0.55
1:J:426:VAL:O	1:K:510:ASN:ND2	2.27	0.55
1:M:414:ASN:HB3	1:N:534:VAL:HG22	1.90	0.55
1:N:266:SER:HB3	1:N:290:ILE:HD11	1.89	0.55
1:A:124:MET:HG2	1:A:154:LEU:HG	1.88	0.54
1:A:151:ILE:O	1:A:155:ILE:HG12	2.07	0.54
1:A:266:SER:HB3	1:A:290:ILE:HD11	1.89	0.54
1:C:151:ILE:O	1:C:155:ILE:HG12	2.07	0.54
1:C:496:ASN:OD1	1:C:497:SER:N	2.40	0.54
1:E:151:ILE:O	1:E:155:ILE:HG12	2.07	0.54
1:E:258:LEU:CD2	1:E:318:LEU:HD23	2.36	0.54
1:E:496:ASN:OD1	1:E:497:SER:N	2.40	0.54
1:G:105:ARG:NE	1:G:151:ILE:HD12	2.22	0.54
1:G:288:VAL:HG22	1:G:303:ALA:CB	2.38	0.54
1:I:105:ARG:NE	1:I:151:ILE:HD12	2.22	0.54
1:I:151:ILE:O	1:I:155:ILE:HG12	2.07	0.54
1:I:496:ASN:OD1	1:I:497:SER:N	2.40	0.54
1:K:414:ASN:HB3	1:L:534:VAL:HG22	1.90	0.54
1:L:603:LEU:HD13	1:N:551:ARG:NH1	2.21	0.54
1:N:161:VAL:HA	1:N:164:ILE:HG12	1.88	0.54
1:O:161:VAL:HA	1:O:164:ILE:HG12	1.88	0.54
1:A:161:VAL:HA	1:A:164:ILE:HG12	1.88	0.54
1:E:105:ARG:NE	1:E:151:ILE:HD12	2.22	0.54
1:E:118:ALA:HA	1:E:136:TYR:OH	2.07	0.54
1:F:105:ARG:NE	1:F:151:ILE:HD12	2.22	0.54
1:H:414:ASN:HB3	1:I:534:VAL:HG22	1.90	0.54
1:I:603:LEU:HD13	1:K:551:ARG:NH1	2.21	0.54
1:K:105:ARG:NE	1:K:151:ILE:HD12	2.22	0.54
1:K:151:ILE:O	1:K:155:ILE:HG12	2.07	0.54
1:M:118:ALA:HA	1:M:136:TYR:OH	2.07	0.54
1:M:161:VAL:HA	1:M:164:ILE:HG12	1.88	0.54
1:N:258:LEU:CD2	1:N:318:LEU:HD23	2.36	0.54
1:N:496:ASN:OD1	1:N:497:SER:N	2.40	0.54
1:A:603:LEU:HD13	1:C:551:ARG:NH1	2.21	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:288:VAL:HG22	1:C:303:ALA:CB	2.38	0.54
1:D:105:ARG:NE	1:D:151:ILE:HD12	2.22	0.54
1:D:118:ALA:HA	1:D:136:TYR:OH	2.07	0.54
1:J:414:ASN:HB3	1:K:534:VAL:HG22	1.90	0.54
1:N:153:LYS:HA	1:N:156:GLU:HB2	1.89	0.54
1:N:414:ASN:HB3	1:O:534:VAL:HG22	1.90	0.54
1:A:288:VAL:HG22	1:A:303:ALA:CB	2.38	0.54
1:B:124:MET:HG2	1:B:154:LEU:HG	1.88	0.54
1:B:288:VAL:HG22	1:B:303:ALA:CB	2.38	0.54
1:C:105:ARG:NE	1:C:151:ILE:HD12	2.22	0.54
1:F:130:VAL:HG22	1:F:131:GLY:H	1.73	0.54
1:G:151:ILE:O	1:G:155:ILE:HG12	2.07	0.54
1:H:578:ARG:O	1:H:582:LYS:N	2.32	0.54
1:I:221:PRO:HG2	1:I:224:ALA:H	1.73	0.54
1:I:414:ASN:HB3	1:J:534:VAL:HG22	1.90	0.54
1:L:118:ALA:HA	1:L:136:TYR:OH	2.07	0.54
1:C:124:MET:HG2	1:C:154:LEU:HG	1.88	0.54
1:D:288:VAL:HG22	1:D:303:ALA:CB	2.38	0.54
1:G:130:VAL:HG22	1:G:131:GLY:H	1.73	0.54
1:G:221:PRO:HG2	1:G:224:ALA:H	1.73	0.54
1:H:221:PRO:HG2	1:H:224:ALA:H	1.73	0.54
1:H:266:SER:HB3	1:H:290:ILE:HD11	1.89	0.54
1:L:414:ASN:HB3	1:M:534:VAL:HG22	1.90	0.54
1:M:105:ARG:NE	1:M:151:ILE:HD12	2.22	0.54
1:M:266:SER:HB3	1:M:290:ILE:HD11	1.89	0.54
1:N:118:ALA:HA	1:N:136:TYR:OH	2.07	0.54
1:O:288:VAL:HG22	1:O:303:ALA:CB	2.38	0.54
1:O:473:GLN:HB3	1:O:481:LEU:HB3	1.90	0.54
1:A:414:ASN:HB3	1:B:534:VAL:HG22	1.90	0.54
1:B:105:ARG:NE	1:B:151:ILE:HD12	2.22	0.54
1:B:130:VAL:HG22	1:B:131:GLY:H	1.73	0.54
1:B:212:ARG:HH12	1:C:177:ALA:CB	2.21	0.54
1:B:266:SER:HB3	1:B:290:ILE:HD11	1.89	0.54
1:C:212:ARG:HH12	1:D:177:ALA:CB	2.21	0.54
1:D:212:ARG:HH12	1:E:177:ALA:CB	2.21	0.54
1:E:212:ARG:HH12	1:F:177:ALA:CB	2.21	0.54
1:E:288:VAL:HG22	1:E:303:ALA:CB	2.38	0.54
1:H:288:VAL:HG22	1:H:303:ALA:CB	2.38	0.54
1:M:153:LYS:HA	1:M:156:GLU:HB2	1.89	0.54
1:N:105:ARG:NE	1:N:151:ILE:HD12	2.22	0.54
1:O:130:VAL:HG22	1:O:131:GLY:H	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:473:GLN:HB3	1:A:481:LEU:HB3	1.90	0.54
1:A:534:VAL:HG22	1:O:414:ASN:HB3	1.90	0.54
1:C:118:ALA:HA	1:C:136:TYR:OH	2.07	0.54
1:D:224:ALA:HA	1:D:227:ARG:HG2	1.90	0.54
1:E:130:VAL:HG22	1:E:131:GLY:H	1.73	0.54
1:E:153:LYS:HA	1:E:156:GLU:HB2	1.89	0.54
1:F:288:VAL:HG22	1:F:303:ALA:CB	2.38	0.54
1:F:473:GLN:HB3	1:F:481:LEU:HB3	1.90	0.54
1:H:153:LYS:HA	1:H:156:GLU:HB2	1.89	0.54
1:K:118:ALA:HA	1:K:136:TYR:OH	2.07	0.54
1:L:153:LYS:HA	1:L:156:GLU:HB2	1.89	0.54
1:L:266:SER:HB3	1:L:290:ILE:HD11	1.89	0.54
1:L:496:ASN:OD1	1:L:497:SER:N	2.40	0.54
1:N:130:VAL:HG22	1:N:131:GLY:H	1.73	0.54
1:N:473:GLN:HB3	1:N:481:LEU:HB3	1.90	0.54
1:A:224:ALA:HA	1:A:227:ARG:HG2	1.90	0.54
1:B:224:ALA:HA	1:B:227:ARG:HG2	1.90	0.54
1:C:153:LYS:HA	1:C:156:GLU:HB2	1.89	0.54
1:C:249:TYR:OH	1:D:317:ARG:O	2.17	0.54
1:D:130:VAL:HG22	1:D:131:GLY:H	1.73	0.54
1:F:118:ALA:HA	1:F:136:TYR:OH	2.07	0.54
1:F:221:PRO:HG2	1:F:224:ALA:H	1.73	0.54
1:G:118:ALA:HA	1:G:136:TYR:OH	2.07	0.54
1:G:153:LYS:HA	1:G:156:GLU:HB2	1.89	0.54
1:G:414:ASN:HB3	1:H:534:VAL:HG22	1.90	0.54
1:H:118:ALA:HA	1:H:136:TYR:OH	2.07	0.54
1:H:603:LEU:HD13	1:J:551:ARG:NH1	2.21	0.54
1:I:288:VAL:HG22	1:I:303:ALA:CB	2.38	0.54
1:L:224:ALA:HA	1:L:227:ARG:HG2	1.90	0.54
1:N:224:ALA:HA	1:N:227:ARG:HG2	1.90	0.54
1:O:118:ALA:HA	1:O:136:TYR:OH	2.07	0.54
1:O:151:ILE:O	1:O:155:ILE:HG12	2.07	0.54
1:O:153:LYS:HA	1:O:156:GLU:HB2	1.89	0.54
1:O:496:ASN:OD1	1:O:497:SER:N	2.40	0.54
1:A:130:VAL:HG22	1:A:131:GLY:H	1.73	0.54
1:A:496:ASN:OD1	1:A:497:SER:N	2.40	0.54
1:C:224:ALA:HA	1:C:227:ARG:HG2	1.90	0.54
1:C:266:SER:HB3	1:C:290:ILE:HD11	1.89	0.54
1:E:216:LEU:HD23	1:E:232:LEU:HD11	1.90	0.54
1:E:224:ALA:HA	1:E:227:ARG:HG2	1.90	0.54
1:F:134:VAL:HG22	1:F:144:LEU:HD22	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:414:ASN:HB3	1:G:534:VAL:HG22	1.90	0.54
1:G:216:LEU:HD23	1:G:232:LEU:HD11	1.90	0.54
1:G:473:GLN:HB3	1:G:481:LEU:HB3	1.90	0.54
1:G:496:ASN:OD1	1:G:497:SER:N	2.40	0.54
1:H:134:VAL:HG22	1:H:144:LEU:HD22	1.90	0.54
1:H:151:ILE:O	1:H:155:ILE:HG12	2.07	0.54
1:I:118:ALA:HA	1:I:136:TYR:OH	2.07	0.54
1:J:221:PRO:HG2	1:J:224:ALA:H	1.73	0.54
1:J:288:VAL:HG22	1:J:303:ALA:CB	2.38	0.54
1:K:224:ALA:HA	1:K:227:ARG:HG2	1.90	0.54
1:K:288:VAL:HG22	1:K:303:ALA:CB	2.38	0.54
1:L:151:ILE:O	1:L:155:ILE:HG12	2.07	0.54
1:A:105:ARG:NE	1:A:151:ILE:HD12	2.22	0.54
1:A:118:ALA:HA	1:A:136:TYR:OH	2.07	0.54
1:A:134:VAL:HG22	1:A:144:LEU:HD22	1.90	0.54
1:B:414:ASN:HB3	1:C:534:VAL:HG22	1.90	0.54
1:D:216:LEU:HD23	1:D:232:LEU:HD11	1.90	0.54
1:D:578:ARG:O	1:D:582:LYS:N	2.32	0.54
1:E:221:PRO:HG2	1:E:224:ALA:H	1.73	0.54
1:E:414:ASN:HB3	1:F:534:VAL:HG22	1.90	0.54
1:E:473:GLN:HB3	1:E:481:LEU:HB3	1.90	0.54
1:F:212:ARG:HH12	1:G:177:ALA:CB	2.21	0.54
1:F:216:LEU:HD23	1:F:232:LEU:HD11	1.91	0.54
1:H:105:ARG:NE	1:H:151:ILE:HD12	2.22	0.54
1:H:130:VAL:HG22	1:H:131:GLY:H	1.73	0.54
1:K:266:SER:HB3	1:K:290:ILE:HD11	1.89	0.54
1:L:105:ARG:NE	1:L:151:ILE:HD12	2.22	0.54
1:L:212:ARG:HH12	1:M:177:ALA:CB	2.21	0.54
1:M:224:ALA:HA	1:M:227:ARG:HG2	1.90	0.54
1:N:288:VAL:HG22	1:N:303:ALA:CB	2.38	0.54
1:O:224:ALA:HA	1:O:227:ARG:HG2	1.90	0.54
1:O:258:LEU:CD2	1:O:318:LEU:HD23	2.36	0.54
1:A:212:ARG:HH12	1:B:177:ALA:CB	2.21	0.53
1:C:216:LEU:HD23	1:C:232:LEU:HD11	1.90	0.53
1:D:134:VAL:HG22	1:D:144:LEU:HD22	1.90	0.53
1:D:496:ASN:OD1	1:D:497:SER:N	2.40	0.53
1:E:304:ASP:OD1	1:E:305:GLN:N	2.42	0.53
1:G:104:THR:OG1	1:G:144:LEU:O	2.21	0.53
1:K:212:ARG:HH12	1:L:177:ALA:CB	2.21	0.53
1:K:304:ASP:OD1	1:K:305:GLN:N	2.42	0.53
1:M:212:ARG:HH12	1:N:177:ALA:CB	2.21	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:473:GLN:HB3	1:M:481:LEU:HB3	1.90	0.53
1:N:134:VAL:HG22	1:N:144:LEU:HD22	1.90	0.53
1:N:578:ARG:O	1:N:582:LYS:N	2.32	0.53
1:O:105:ARG:NE	1:O:151:ILE:HD12	2.22	0.53
1:B:153:LYS:HA	1:B:156:GLU:HB2	1.89	0.53
1:D:304:ASP:OD1	1:D:305:GLN:N	2.42	0.53
1:F:258:LEU:CD2	1:F:318:LEU:HD23	2.36	0.53
1:F:304:ASP:OD1	1:F:305:GLN:N	2.42	0.53
1:H:216:LEU:HD23	1:H:232:LEU:HD11	1.90	0.53
1:K:578:ARG:O	1:K:582:LYS:N	2.32	0.53
1:M:130:VAL:HG22	1:M:131:GLY:H	1.73	0.53
1:O:221:PRO:HG2	1:O:224:ALA:H	1.73	0.53
1:A:177:ALA:CB	1:O:212:ARG:HH12	2.21	0.53
1:B:473:GLN:HB3	1:B:481:LEU:HB3	1.90	0.53
1:C:578:ARG:O	1:C:582:LYS:N	2.32	0.53
1:D:151:ILE:O	1:D:155:ILE:HG12	2.07	0.53
1:D:414:ASN:HB3	1:E:534:VAL:HG22	1.90	0.53
1:D:473:GLN:HB3	1:D:481:LEU:HB3	1.90	0.53
1:E:578:ARG:O	1:E:582:LYS:N	2.32	0.53
1:H:473:GLN:HB3	1:H:481:LEU:HB3	1.90	0.53
1:J:212:ARG:HH12	1:K:177:ALA:CB	2.21	0.53
1:J:603:LEU:HD13	1:L:551:ARG:NH1	2.21	0.53
1:L:168:LYS:N	1:L:220:GLY:O	2.30	0.53
1:M:288:VAL:HG22	1:M:303:ALA:CB	2.38	0.53
1:N:221:PRO:HG2	1:N:224:ALA:H	1.73	0.53
1:C:304:ASP:OD1	1:C:305:GLN:N	2.42	0.53
1:D:153:LYS:HA	1:D:156:GLU:HB2	1.89	0.53
1:D:221:PRO:HG2	1:D:224:ALA:H	1.73	0.53
1:F:224:ALA:HA	1:F:227:ARG:HG2	1.90	0.53
1:F:578:ARG:O	1:F:582:LYS:N	2.32	0.53
1:H:224:ALA:HA	1:H:227:ARG:HG2	1.90	0.53
1:I:153:LYS:HA	1:I:156:GLU:HB2	1.89	0.53
1:I:212:ARG:HH12	1:J:177:ALA:CB	2.21	0.53
1:J:118:ALA:HA	1:J:136:TYR:OH	2.07	0.53
1:J:134:VAL:HG22	1:J:144:LEU:HD22	1.90	0.53
1:J:266:SER:HB3	1:J:290:ILE:HD11	1.89	0.53
1:L:304:ASP:OD1	1:L:305:GLN:N	2.42	0.53
1:N:212:ARG:HH12	1:O:177:ALA:CB	2.21	0.53
1:O:304:ASP:OD1	1:O:305:GLN:N	2.42	0.53
1:B:216:LEU:HD23	1:B:232:LEU:HD11	1.90	0.53
1:C:134:VAL:HG22	1:C:144:LEU:HD22	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:153:LYS:HA	1:F:156:GLU:HB2	1.89	0.53
1:G:157:VAL:HG22	1:G:160:ARG:HH21	1.74	0.53
1:G:224:ALA:HA	1:G:227:ARG:HG2	1.90	0.53
1:G:304:ASP:OD1	1:G:305:GLN:N	2.42	0.53
1:I:216:LEU:HD23	1:I:232:LEU:HD11	1.90	0.53
1:I:224:ALA:HA	1:I:227:ARG:HG2	1.90	0.53
1:I:266:SER:HB3	1:I:290:ILE:HD11	1.89	0.53
1:I:589:ARG:HG3	1:I:611:VAL:HG22	1.91	0.53
1:J:224:ALA:HA	1:J:227:ARG:HG2	1.90	0.53
1:J:304:ASP:OD1	1:J:305:GLN:N	2.42	0.53
1:J:589:ARG:HG3	1:J:611:VAL:HG22	1.91	0.53
1:L:134:VAL:HG22	1:L:144:LEU:HD22	1.90	0.53
1:L:288:VAL:HG22	1:L:303:ALA:CB	2.38	0.53
1:N:259:VAL:HG21	1:N:292:ALA:HB2	1.90	0.53
1:A:174:LEU:HG	1:A:216:LEU:N	2.24	0.53
1:B:134:VAL:HG22	1:B:144:LEU:HD22	1.90	0.53
1:H:212:ARG:HH12	1:I:177:ALA:CB	2.21	0.53
1:I:258:LEU:CD2	1:I:318:LEU:HD23	2.36	0.53
1:K:259:VAL:HG21	1:K:292:ALA:HB2	1.90	0.53
1:M:151:ILE:O	1:M:155:ILE:HG12	2.07	0.53
1:N:151:ILE:O	1:N:155:ILE:HG12	2.07	0.53
1:N:174:LEU:HG	1:N:216:LEU:N	2.24	0.53
1:A:221:PRO:HG2	1:A:224:ALA:H	1.73	0.53
1:B:118:ALA:HA	1:B:136:TYR:OH	2.07	0.53
1:C:414:ASN:HB3	1:D:534:VAL:HG22	1.90	0.53
1:D:259:VAL:HG21	1:D:292:ALA:HB2	1.90	0.53
1:G:589:ARG:HG3	1:G:611:VAL:HG22	1.91	0.53
1:H:496:ASN:OD1	1:H:497:SER:N	2.40	0.53
1:J:216:LEU:HD23	1:J:232:LEU:HD11	1.90	0.53
1:M:259:VAL:HG21	1:M:292:ALA:HB2	1.90	0.53
1:N:304:ASP:OD1	1:N:305:GLN:N	2.42	0.53
1:O:157:VAL:HG22	1:O:160:ARG:HH21	1.74	0.53
1:A:153:LYS:HA	1:A:156:GLU:HB2	1.89	0.53
1:A:216:LEU:HD23	1:A:232:LEU:HD11	1.90	0.53
1:A:304:ASP:OD1	1:A:305:GLN:N	2.42	0.53
1:B:242:GLU:HG2	1:B:243:GLY:H	1.60	0.53
1:C:174:LEU:HG	1:C:216:LEU:N	2.24	0.53
1:C:221:PRO:HG2	1:C:224:ALA:H	1.73	0.53
1:C:473:GLN:HB3	1:C:481:LEU:HB3	1.90	0.53
1:D:288:VAL:HG13	1:D:303:ALA:HB2	1.91	0.53
1:G:212:ARG:HH12	1:H:177:ALA:CB	2.21	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:174:LEU:HG	1:H:216:LEU:N	2.24	0.53
1:I:130:VAL:HG22	1:I:131:GLY:H	1.72	0.53
1:I:174:LEU:HG	1:I:216:LEU:N	2.24	0.53
1:J:578:ARG:O	1:J:582:LYS:N	2.32	0.53
1:K:130:VAL:HG22	1:K:131:GLY:H	1.73	0.53
1:K:221:PRO:HG2	1:K:224:ALA:H	1.73	0.53
1:L:589:ARG:HG3	1:L:611:VAL:HG22	1.91	0.53
1:O:109:LEU:HG	1:O:112:VAL:O	2.09	0.53
1:A:288:VAL:HG13	1:A:303:ALA:HB2	1.91	0.53
1:C:130:VAL:HG22	1:C:131:GLY:H	1.73	0.53
1:C:259:VAL:HG21	1:C:292:ALA:HB2	1.90	0.53
1:E:259:VAL:HG21	1:E:292:ALA:HB2	1.90	0.53
1:F:288:VAL:HG13	1:F:303:ALA:HB2	1.91	0.53
1:G:174:LEU:HG	1:G:216:LEU:N	2.24	0.53
1:G:288:VAL:HG13	1:G:303:ALA:HB2	1.91	0.53
1:H:304:ASP:OD1	1:H:305:GLN:N	2.42	0.53
1:H:589:ARG:HG3	1:H:611:VAL:HG22	1.91	0.53
1:I:134:VAL:HG22	1:I:144:LEU:HD22	1.90	0.53
1:I:288:VAL:HG13	1:I:303:ALA:HB2	1.91	0.53
1:K:216:LEU:HD23	1:K:232:LEU:HD11	1.90	0.53
1:L:157:VAL:HG22	1:L:160:ARG:HH21	1.74	0.53
1:L:216:LEU:HD23	1:L:232:LEU:HD11	1.90	0.53
1:L:221:PRO:HG2	1:L:224:ALA:H	1.73	0.53
1:L:473:GLN:HB3	1:L:481:LEU:HB3	1.90	0.53
1:M:221:PRO:HG2	1:M:224:ALA:H	1.73	0.53
1:O:216:LEU:HD23	1:O:232:LEU:HD11	1.90	0.53
1:B:157:VAL:HG22	1:B:160:ARG:HH21	1.74	0.53
1:B:288:VAL:HG13	1:B:303:ALA:HB2	1.91	0.53
1:F:589:ARG:HG3	1:F:611:VAL:HG22	1.91	0.53
1:J:160:ARG:O	1:J:164:ILE:HG12	2.09	0.53
1:J:174:LEU:HG	1:J:216:LEU:N	2.24	0.53
1:K:589:ARG:HG3	1:K:611:VAL:HG22	1.91	0.53
1:L:109:LEU:HG	1:L:112:VAL:O	2.09	0.53
1:L:130:VAL:HG22	1:L:131:GLY:H	1.73	0.53
1:M:301:ILE:HG22	1:M:308:GLN:HE22	1.74	0.53
1:N:216:LEU:HD23	1:N:232:LEU:HD11	1.90	0.53
1:A:160:ARG:HG2	1:A:164:ILE:HD13	1.91	0.52
1:B:304:ASP:OD1	1:B:305:GLN:N	2.42	0.52
1:C:109:LEU:HG	1:C:112:VAL:O	2.09	0.52
1:C:160:ARG:HG2	1:C:164:ILE:HD13	1.91	0.52
1:C:288:VAL:HG13	1:C:303:ALA:HB2	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:134:VAL:HG22	1:G:144:LEU:HD22	1.90	0.52
1:H:301:ILE:HG22	1:H:308:GLN:HE22	1.74	0.52
1:I:157:VAL:HG22	1:I:160:ARG:HH21	1.74	0.52
1:I:473:GLN:HB3	1:I:481:LEU:HB3	1.90	0.52
1:J:109:LEU:HG	1:J:112:VAL:O	2.09	0.52
1:J:288:VAL:HG13	1:J:303:ALA:HB2	1.91	0.52
1:K:160:ARG:O	1:K:164:ILE:HG12	2.09	0.52
1:L:174:LEU:HG	1:L:216:LEU:N	2.24	0.52
1:L:242:GLU:HG2	1:L:243:GLY:H	1.60	0.52
1:M:216:LEU:HD23	1:M:232:LEU:HD11	1.90	0.52
1:O:301:ILE:HG22	1:O:308:GLN:HE22	1.74	0.52
1:B:174:LEU:HG	1:B:216:LEU:N	2.24	0.52
1:D:157:VAL:HG22	1:D:160:ARG:HH21	1.74	0.52
1:E:288:VAL:HG13	1:E:303:ALA:HB2	1.91	0.52
1:F:174:LEU:HG	1:F:216:LEU:N	2.24	0.52
1:H:288:VAL:HG13	1:H:303:ALA:HB2	1.91	0.52
1:I:109:LEU:HG	1:I:112:VAL:O	2.09	0.52
1:I:160:ARG:O	1:I:164:ILE:HG12	2.09	0.52
1:L:160:ARG:O	1:L:164:ILE:HG12	2.09	0.52
1:L:259:VAL:HG21	1:L:292:ALA:HB2	1.90	0.52
1:N:160:ARG:HG2	1:N:164:ILE:HD13	1.91	0.52
1:O:134:VAL:HG22	1:O:144:LEU:HD22	1.90	0.52
1:O:288:VAL:HG13	1:O:303:ALA:HB2	1.91	0.52
1:A:258:LEU:CD2	1:A:318:LEU:HD23	2.36	0.52
1:B:221:PRO:HG2	1:B:224:ALA:H	1.73	0.52
1:E:134:VAL:HG22	1:E:144:LEU:HD22	1.90	0.52
1:E:174:LEU:HG	1:E:216:LEU:N	2.24	0.52
1:F:109:LEU:HG	1:F:112:VAL:O	2.09	0.52
1:K:174:LEU:HG	1:K:216:LEU:N	2.24	0.52
1:M:160:ARG:O	1:M:164:ILE:HG12	2.09	0.52
1:A:259:VAL:HG21	1:A:292:ALA:HB2	1.90	0.52
1:B:109:LEU:HG	1:B:112:VAL:O	2.09	0.52
1:B:301:ILE:HG22	1:B:308:GLN:HE22	1.74	0.52
1:D:160:ARG:HG2	1:D:164:ILE:HD13	1.91	0.52
1:E:160:ARG:HG2	1:E:164:ILE:HD13	1.92	0.52
1:F:301:ILE:HG22	1:F:308:GLN:HE22	1.74	0.52
1:G:109:LEU:HG	1:G:112:VAL:O	2.09	0.52
1:G:160:ARG:HG2	1:G:164:ILE:HD13	1.91	0.52
1:G:258:LEU:CD2	1:G:318:LEU:HD23	2.36	0.52
1:I:304:ASP:OD1	1:I:305:GLN:N	2.42	0.52
1:K:134:VAL:HG22	1:K:144:LEU:HD22	1.90	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:288:VAL:HG13	1:K:303:ALA:HB2	1.91	0.52
1:M:109:LEU:HG	1:M:112:VAL:O	2.09	0.52
1:M:589:ARG:HG3	1:M:611:VAL:HG22	1.91	0.52
1:O:259:VAL:HG21	1:O:292:ALA:HB2	1.90	0.52
1:B:578:ARG:O	1:B:582:LYS:N	2.32	0.52
1:D:301:ILE:HG22	1:D:308:GLN:HE22	1.74	0.52
1:H:160:ARG:O	1:H:164:ILE:HG12	2.09	0.52
1:J:259:VAL:HG21	1:J:292:ALA:HB2	1.90	0.52
1:K:301:ILE:HG22	1:K:308:GLN:HE22	1.74	0.52
1:M:134:VAL:HG22	1:M:144:LEU:HD22	1.90	0.52
1:N:288:VAL:HG13	1:N:303:ALA:HB2	1.91	0.52
1:A:157:VAL:HG22	1:A:160:ARG:HH21	1.74	0.52
1:E:301:ILE:HG22	1:E:308:GLN:HE22	1.74	0.52
1:F:160:ARG:HG2	1:F:164:ILE:HD13	1.91	0.52
1:G:477:GLY:O	1:G:572:ARG:NH2	2.43	0.52
1:H:259:VAL:HG21	1:H:292:ALA:HB2	1.90	0.52
1:L:477:GLY:O	1:L:572:ARG:NH2	2.43	0.52
1:M:304:ASP:OD1	1:M:305:GLN:N	2.42	0.52
1:O:174:LEU:HG	1:O:216:LEU:N	2.24	0.52
1:B:160:ARG:HG2	1:B:164:ILE:HD13	1.91	0.52
1:B:259:VAL:HG21	1:B:292:ALA:HB2	1.90	0.52
1:C:157:VAL:HG22	1:C:160:ARG:HH21	1.74	0.52
1:J:130:VAL:HG22	1:J:131:GLY:H	1.72	0.52
1:J:301:ILE:HG22	1:J:308:GLN:HE22	1.75	0.52
1:J:473:GLN:HB3	1:J:481:LEU:HB3	1.90	0.52
1:L:288:VAL:HG13	1:L:303:ALA:HB2	1.91	0.52
1:N:109:LEU:HG	1:N:112:VAL:O	2.09	0.52
1:A:109:LEU:HG	1:A:112:VAL:O	2.09	0.52
1:A:301:ILE:HG22	1:A:308:GLN:HE22	1.74	0.52
1:B:589:ARG:HG3	1:B:611:VAL:HG22	1.91	0.52
1:C:104:THR:OG1	1:C:144:LEU:O	2.21	0.52
1:E:109:LEU:HG	1:E:112:VAL:O	2.09	0.52
1:F:259:VAL:HG21	1:F:292:ALA:HB2	1.90	0.52
1:I:160:ARG:HG2	1:I:164:ILE:HD13	1.91	0.52
1:I:259:VAL:HG21	1:I:292:ALA:HB2	1.90	0.52
1:K:473:GLN:HB3	1:K:481:LEU:HB3	1.90	0.52
1:M:288:VAL:HG13	1:M:303:ALA:HB2	1.91	0.52
1:N:160:ARG:O	1:N:164:ILE:HG12	2.09	0.52
1:O:160:ARG:HG2	1:O:164:ILE:HD13	1.91	0.52
1:C:589:ARG:HG3	1:C:611:VAL:HG22	1.91	0.52
1:F:157:VAL:HG22	1:F:160:ARG:HH21	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:259:VAL:HG21	1:G:292:ALA:HB2	1.90	0.52
1:J:104:THR:OG1	1:J:144:LEU:O	2.21	0.52
1:M:160:ARG:HG2	1:M:164:ILE:HD13	1.91	0.52
1:M:174:LEU:HG	1:M:216:LEU:N	2.24	0.52
1:O:589:ARG:HG3	1:O:611:VAL:HG22	1.91	0.52
1:A:242:GLU:HG2	1:A:243:GLY:H	1.60	0.52
1:E:589:ARG:HG3	1:E:611:VAL:HG22	1.91	0.52
1:I:301:ILE:HG22	1:I:308:GLN:HE22	1.74	0.52
1:K:135:HIS:CE1	1:L:120:LEU:HD12	2.45	0.52
1:K:160:ARG:HG2	1:K:164:ILE:HD13	1.91	0.52
1:K:477:GLY:O	1:K:572:ARG:NH2	2.43	0.52
1:L:135:HIS:CE1	1:M:120:LEU:HD12	2.45	0.52
1:A:104:THR:OG1	1:A:144:LEU:O	2.21	0.51
1:C:135:HIS:CE1	1:D:120:LEU:HD12	2.45	0.51
1:C:301:ILE:HG22	1:C:308:GLN:HE22	1.74	0.51
1:D:135:HIS:CE1	1:E:120:LEU:HD12	2.45	0.51
1:D:174:LEU:HG	1:D:216:LEU:N	2.24	0.51
1:F:477:GLY:O	1:F:572:ARG:NH2	2.43	0.51
1:G:160:ARG:O	1:G:164:ILE:HG12	2.10	0.51
1:H:109:LEU:HG	1:H:112:VAL:O	2.09	0.51
1:I:135:HIS:CE1	1:J:120:LEU:HD12	2.45	0.51
1:J:135:HIS:CE1	1:K:120:LEU:HD12	2.45	0.51
1:J:157:VAL:HG22	1:J:160:ARG:HH21	1.74	0.51
1:K:157:VAL:HG22	1:K:160:ARG:HH21	1.74	0.51
1:L:301:ILE:HG22	1:L:308:GLN:HE22	1.74	0.51
1:M:135:HIS:CE1	1:N:120:LEU:HD12	2.45	0.51
1:N:157:VAL:HG22	1:N:160:ARG:HH21	1.74	0.51
1:O:242:GLU:HG2	1:O:243:GLY:H	1.60	0.51
1:D:109:LEU:HG	1:D:112:VAL:O	2.09	0.51
1:D:589:ARG:HG3	1:D:611:VAL:HG22	1.91	0.51
1:N:589:ARG:HG3	1:N:611:VAL:HG22	1.91	0.51
1:B:160:ARG:O	1:B:164:ILE:HG12	2.09	0.51
1:F:160:ARG:O	1:F:164:ILE:HG12	2.09	0.51
1:G:301:ILE:HG22	1:G:308:GLN:HE22	1.74	0.51
1:H:135:HIS:CE1	1:I:120:LEU:HD12	2.45	0.51
1:I:206:LYS:HZ3	1:J:190:ILE:C	2.19	0.51
1:J:262:LEU:CD1	1:J:299:LEU:HD21	2.40	0.51
1:K:109:LEU:HG	1:K:112:VAL:O	2.09	0.51
1:L:160:ARG:HG2	1:L:164:ILE:HD13	1.91	0.51
1:O:104:THR:OG1	1:O:144:LEU:O	2.22	0.51
1:A:180:GLU:O	1:A:183:ALA:HB3	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:135:HIS:CE1	1:C:120:LEU:HD12	2.45	0.51
1:E:135:HIS:CE1	1:F:120:LEU:HD12	2.45	0.51
1:F:180:GLU:O	1:F:183:ALA:HB3	2.11	0.51
1:G:180:GLU:O	1:G:183:ALA:HB3	2.11	0.51
1:A:589:ARG:HG3	1:A:611:VAL:HG22	1.91	0.51
1:L:117:LEU:HB3	1:L:121:LEU:HG	1.92	0.51
1:N:104:THR:OG1	1:N:144:LEU:O	2.21	0.51
1:N:135:HIS:CE1	1:O:120:LEU:HD12	2.45	0.51
1:A:135:HIS:CE1	1:B:120:LEU:HD12	2.45	0.51
1:B:180:GLU:O	1:B:183:ALA:HB3	2.11	0.51
1:D:171:ILE:HG22	1:D:217:ILE:HG23	1.93	0.51
1:D:242:GLU:HG2	1:D:243:GLY:H	1.60	0.51
1:E:157:VAL:HG22	1:E:160:ARG:HH21	1.74	0.51
1:H:180:GLU:O	1:H:183:ALA:HB3	2.11	0.51
1:K:117:LEU:HB3	1:K:121:LEU:HG	1.92	0.51
1:M:117:LEU:HB3	1:M:121:LEU:HG	1.92	0.51
1:N:301:ILE:HG22	1:N:308:GLN:HE22	1.74	0.51
1:C:117:LEU:HB3	1:C:121:LEU:HG	1.92	0.51
1:C:160:ARG:O	1:C:164:ILE:HG12	2.10	0.51
1:C:171:ILE:HG22	1:C:217:ILE:HG23	1.93	0.51
1:H:262:LEU:CD1	1:H:299:LEU:HD21	2.40	0.51
1:L:335:GLN:HE21	1:L:560:ARG:HE	1.59	0.51
1:O:160:ARG:O	1:O:164:ILE:HG12	2.10	0.51
1:O:180:GLU:O	1:O:183:ALA:HB3	2.11	0.51
1:B:171:ILE:HG22	1:B:217:ILE:HG23	1.93	0.51
1:B:262:LEU:CD1	1:B:299:LEU:HD21	2.40	0.51
1:D:117:LEU:HB3	1:D:121:LEU:HG	1.92	0.51
1:E:171:ILE:HG22	1:E:217:ILE:HG23	1.93	0.51
1:E:180:GLU:O	1:E:183:ALA:HB3	2.11	0.51
1:F:135:HIS:CE1	1:G:120:LEU:HD12	2.45	0.51
1:H:160:ARG:HG2	1:H:164:ILE:HD13	1.91	0.51
1:J:117:LEU:HB3	1:J:121:LEU:HG	1.92	0.51
1:J:335:GLN:HE21	1:J:560:ARG:HE	1.59	0.51
1:K:335:GLN:HE21	1:K:560:ARG:HE	1.59	0.51
1:N:262:LEU:CD1	1:N:299:LEU:HD21	2.40	0.51
1:A:477:GLY:O	1:A:572:ARG:NH2	2.43	0.51
1:B:117:LEU:HB3	1:B:121:LEU:HG	1.92	0.51
1:B:258:LEU:CD2	1:B:318:LEU:HD23	2.36	0.51
1:G:135:HIS:CE1	1:H:120:LEU:HD12	2.45	0.51
1:I:180:GLU:O	1:I:183:ALA:HB3	2.11	0.51
1:M:335:GLN:HE21	1:M:560:ARG:HE	1.59	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:117:LEU:HB3	1:N:121:LEU:HG	1.92	0.51
1:D:180:GLU:O	1:D:183:ALA:HB3	2.11	0.51
1:E:160:ARG:O	1:E:164:ILE:HG12	2.09	0.51
1:H:157:VAL:HG22	1:H:160:ARG:HH21	1.74	0.51
1:J:180:GLU:O	1:J:183:ALA:HB3	2.11	0.51
1:J:477:GLY:O	1:J:572:ARG:NH2	2.43	0.51
1:N:335:GLN:HE21	1:N:560:ARG:HE	1.59	0.50
1:A:120:LEU:HD12	1:O:135:HIS:CE1	2.45	0.50
1:B:335:GLN:HE21	1:B:560:ARG:HE	1.59	0.50
1:E:262:LEU:CD1	1:E:299:LEU:HD21	2.40	0.50
1:E:477:GLY:O	1:E:572:ARG:NH2	2.43	0.50
1:I:335:GLN:HE21	1:I:560:ARG:HE	1.59	0.50
1:K:180:GLU:O	1:K:183:ALA:HB3	2.11	0.50
1:A:335:GLN:HE21	1:A:560:ARG:HE	1.59	0.50
1:M:578:ARG:O	1:M:582:LYS:N	2.32	0.50
1:N:180:GLU:O	1:N:183:ALA:HB3	2.11	0.50
1:O:335:GLN:HE21	1:O:560:ARG:HE	1.59	0.50
1:A:160:ARG:O	1:A:164:ILE:HG12	2.09	0.50
1:A:171:ILE:HG22	1:A:217:ILE:HG23	1.93	0.50
1:C:180:GLU:O	1:C:183:ALA:HB3	2.11	0.50
1:F:171:ILE:HG22	1:F:217:ILE:HG23	1.93	0.50
1:F:242:GLU:HG2	1:F:243:GLY:H	1.60	0.50
1:G:335:GLN:HE21	1:G:560:ARG:HE	1.59	0.50
1:H:335:GLN:HE21	1:H:560:ARG:HE	1.59	0.50
1:I:117:LEU:HB3	1:I:121:LEU:HG	1.92	0.50
1:O:117:LEU:HB3	1:O:121:LEU:HG	1.92	0.50
1:B:322:ARG:HG3	1:B:428:LEU:HD22	1.94	0.50
1:C:335:GLN:HE21	1:C:560:ARG:HE	1.59	0.50
1:E:322:ARG:HG3	1:E:428:LEU:HD22	1.94	0.50
1:G:288:VAL:HG22	1:G:303:ALA:HB1	1.94	0.50
1:J:160:ARG:HG2	1:J:164:ILE:HD13	1.91	0.50
1:L:180:GLU:O	1:L:183:ALA:HB3	2.11	0.50
1:E:117:LEU:HB3	1:E:121:LEU:HG	1.92	0.50
1:F:262:LEU:CD1	1:F:299:LEU:HD21	2.40	0.50
1:F:288:VAL:HG22	1:F:303:ALA:HB1	1.94	0.50
1:A:262:LEU:CD1	1:A:299:LEU:HD21	2.40	0.50
1:D:160:ARG:O	1:D:164:ILE:HG12	2.10	0.50
1:E:288:VAL:HG22	1:E:303:ALA:HB1	1.94	0.50
1:F:335:GLN:HE21	1:F:560:ARG:HE	1.59	0.50
1:H:322:ARG:HG3	1:H:428:LEU:HD22	1.94	0.50
1:K:262:LEU:CD1	1:K:299:LEU:HD21	2.40	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:322:ARG:HG3	1:N:428:LEU:HD22	1.94	0.50
1:A:117:LEU:HB3	1:A:121:LEU:HG	1.92	0.50
1:A:218:ILE:HD13	1:A:228:ILE:HD13	1.94	0.50
1:B:218:ILE:HD13	1:B:228:ILE:HD13	1.94	0.50
1:G:117:LEU:HB3	1:G:121:LEU:HG	1.92	0.50
1:H:117:LEU:HB3	1:H:121:LEU:HG	1.92	0.50
1:H:288:VAL:HG22	1:H:303:ALA:HB1	1.94	0.50
1:N:218:ILE:HD13	1:N:228:ILE:HD13	1.94	0.50
1:O:477:GLY:O	1:O:572:ARG:NH2	2.43	0.50
1:E:335:GLN:HE21	1:E:560:ARG:HE	1.59	0.50
1:M:104:THR:OG1	1:M:144:LEU:O	2.21	0.50
1:M:180:GLU:O	1:M:183:ALA:HB3	2.11	0.50
1:M:262:LEU:CD1	1:M:299:LEU:HD21	2.40	0.50
1:O:218:ILE:HD13	1:O:228:ILE:HD13	1.94	0.50
1:C:218:ILE:HD13	1:C:228:ILE:HD13	1.94	0.49
1:I:477:GLY:O	1:I:572:ARG:NH2	2.43	0.49
1:M:157:VAL:HG22	1:M:160:ARG:HH21	1.74	0.49
1:D:218:ILE:HD13	1:D:228:ILE:HD13	1.94	0.49
1:D:288:VAL:HG22	1:D:303:ALA:HB1	1.94	0.49
1:F:322:ARG:HG3	1:F:428:LEU:HD22	1.94	0.49
1:I:262:LEU:CD1	1:I:299:LEU:HD21	2.40	0.49
1:M:218:ILE:HD13	1:M:228:ILE:HD13	1.94	0.49
1:O:322:ARG:HG3	1:O:428:LEU:HD22	1.94	0.49
1:A:322:ARG:HG3	1:A:428:LEU:HD22	1.94	0.49
1:D:477:GLY:O	1:D:572:ARG:NH2	2.43	0.49
1:G:250:LEU:HD13	1:G:258:LEU:HD12	1.95	0.49
1:G:322:ARG:HG3	1:G:428:LEU:HD22	1.94	0.49
1:L:444:LEU:HB2	1:M:499:LEU:HD21	1.94	0.49
1:M:444:LEU:HB2	1:N:499:LEU:HD21	1.94	0.49
1:N:242:GLU:HG2	1:N:243:GLY:H	1.60	0.49
1:C:322:ARG:HG3	1:C:428:LEU:HD22	1.94	0.49
1:G:171:ILE:HG22	1:G:217:ILE:HG23	1.93	0.49
1:J:322:ARG:HG3	1:J:428:LEU:HD22	1.94	0.49
1:K:171:ILE:HG22	1:K:217:ILE:HG23	1.93	0.49
1:K:218:ILE:HD13	1:K:228:ILE:HD13	1.94	0.49
1:L:171:ILE:HG22	1:L:217:ILE:HG23	1.93	0.49
1:L:322:ARG:HG3	1:L:428:LEU:HD22	1.94	0.49
1:N:250:LEU:HD13	1:N:258:LEU:HD12	1.95	0.49
1:O:171:ILE:HG22	1:O:217:ILE:HG23	1.93	0.49
1:C:262:LEU:CD1	1:C:299:LEU:HD21	2.41	0.49
1:D:322:ARG:HG3	1:D:428:LEU:HD22	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:335:GLN:HE21	1:D:560:ARG:HE	1.59	0.49
1:F:117:LEU:HB3	1:F:121:LEU:HG	1.92	0.49
1:L:218:ILE:HD13	1:L:228:ILE:HD13	1.94	0.49
1:O:250:LEU:HD13	1:O:258:LEU:HD12	1.95	0.49
1:A:499:LEU:HD21	1:O:444:LEU:HB2	1.94	0.49
1:C:477:GLY:O	1:C:572:ARG:NH2	2.43	0.49
1:F:250:LEU:HD13	1:F:258:LEU:HD12	1.95	0.49
1:H:250:LEU:HD13	1:H:258:LEU:HD12	1.95	0.49
1:I:171:ILE:HG22	1:I:217:ILE:HG23	1.93	0.49
1:J:171:ILE:HG22	1:J:217:ILE:HG23	1.93	0.49
1:M:171:ILE:HG22	1:M:217:ILE:HG23	1.93	0.49
1:O:262:LEU:CD1	1:O:299:LEU:HD21	2.40	0.49
1:E:444:LEU:HB2	1:F:499:LEU:HD21	1.94	0.49
1:G:262:LEU:CD1	1:G:299:LEU:HD21	2.40	0.49
1:I:288:VAL:HG22	1:I:303:ALA:HB1	1.94	0.49
1:J:218:ILE:HD13	1:J:228:ILE:HD13	1.94	0.49
1:J:444:LEU:HB2	1:K:499:LEU:HD21	1.94	0.49
1:N:444:LEU:HB2	1:O:499:LEU:HD21	1.94	0.49
1:E:218:ILE:HD13	1:E:228:ILE:HD13	1.94	0.49
1:M:250:LEU:HD13	1:M:258:LEU:HD12	1.95	0.49
1:A:250:LEU:HD13	1:A:258:LEU:HD12	1.95	0.49
1:F:218:ILE:HD13	1:F:228:ILE:HD13	1.94	0.49
1:F:444:LEU:HB2	1:G:499:LEU:HD21	1.94	0.49
1:H:171:ILE:HG22	1:H:217:ILE:HG23	1.93	0.49
1:I:218:ILE:HD13	1:I:228:ILE:HD13	1.94	0.49
1:I:250:LEU:HD13	1:I:258:LEU:HD12	1.95	0.49
1:K:444:LEU:HB2	1:L:499:LEU:HD21	1.94	0.49
1:N:171:ILE:HG22	1:N:217:ILE:HG23	1.93	0.49
1:O:429:ASP:HA	1:O:472:PRO:HG2	1.95	0.49
1:B:491:VAL:HG22	1:B:503:PHE:HE1	1.78	0.49
1:C:288:VAL:HG22	1:C:303:ALA:HB1	1.94	0.49
1:D:444:LEU:HB2	1:E:499:LEU:HD21	1.94	0.49
1:D:491:VAL:HG22	1:D:503:PHE:HE1	1.78	0.49
1:G:218:ILE:HD13	1:G:228:ILE:HD13	1.94	0.49
1:G:607:GLU:OE2	1:I:557:ARG:HB2	2.13	0.49
1:H:218:ILE:HD13	1:H:228:ILE:HD13	1.94	0.49
1:K:491:VAL:HG22	1:K:503:PHE:HE1	1.78	0.49
1:L:607:GLU:OE2	1:N:557:ARG:HB2	2.13	0.49
1:A:288:VAL:HG22	1:A:303:ALA:HB1	1.94	0.48
1:A:429:ASP:HA	1:A:472:PRO:HG2	1.95	0.48
1:B:557:ARG:HB2	1:O:607:GLU:OE2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:429:ASP:HA	1:G:472:PRO:HG2	1.95	0.48
1:H:607:GLU:OE2	1:J:557:ARG:HB2	2.13	0.48
1:I:444:LEU:HB2	1:J:499:LEU:HD21	1.94	0.48
1:I:491:VAL:HG22	1:I:503:PHE:HE1	1.78	0.48
1:I:607:GLU:OE2	1:K:557:ARG:HB2	2.13	0.48
1:K:429:ASP:HA	1:K:472:PRO:HG2	1.95	0.48
1:L:288:VAL:HG22	1:L:303:ALA:HB1	1.94	0.48
1:M:288:VAL:HG22	1:M:303:ALA:HB1	1.94	0.48
1:N:477:GLY:O	1:N:572:ARG:NH2	2.43	0.48
1:O:288:VAL:HG22	1:O:303:ALA:HB1	1.94	0.48
1:C:607:GLU:OE2	1:E:557:ARG:HB2	2.13	0.48
1:F:607:GLU:OE2	1:H:557:ARG:HB2	2.13	0.48
1:H:429:ASP:HA	1:H:472:PRO:HG2	1.95	0.48
1:J:607:GLU:OE2	1:L:557:ARG:HB2	2.13	0.48
1:L:429:ASP:HA	1:L:472:PRO:HG2	1.95	0.48
1:M:491:VAL:HG22	1:M:503:PHE:HE1	1.78	0.48
1:B:477:GLY:O	1:B:572:ARG:NH2	2.43	0.48
1:E:250:LEU:HD13	1:E:258:LEU:HD12	1.95	0.48
1:E:607:GLU:OE2	1:G:557:ARG:HB2	2.13	0.48
1:H:246:ARG:HH22	1:H:305:GLN:HE22	1.61	0.48
1:I:322:ARG:HG3	1:I:428:LEU:HD22	1.94	0.48
1:L:250:LEU:HD13	1:L:258:LEU:HD12	1.95	0.48
1:M:322:ARG:HG3	1:M:428:LEU:HD22	1.94	0.48
1:N:288:VAL:HG22	1:N:303:ALA:HB1	1.94	0.48
1:O:491:VAL:HG22	1:O:503:PHE:HE1	1.78	0.48
1:A:444:LEU:HB2	1:B:499:LEU:HD21	1.94	0.48
1:D:607:GLU:OE2	1:F:557:ARG:HB2	2.13	0.48
1:F:429:ASP:HA	1:F:472:PRO:HG2	1.95	0.48
1:K:288:VAL:HG22	1:K:303:ALA:HB1	1.94	0.48
1:K:607:GLU:OE2	1:M:557:ARG:HB2	2.13	0.48
1:A:607:GLU:OE2	1:C:557:ARG:HB2	2.13	0.48
1:E:136:TYR:CE1	1:E:142:LEU:HG	2.49	0.48
1:F:246:ARG:HH22	1:F:305:GLN:HE22	1.61	0.48
1:J:250:LEU:HD13	1:J:258:LEU:HD12	1.95	0.48
1:B:250:LEU:HD13	1:B:258:LEU:HD12	1.95	0.48
1:B:288:VAL:HG22	1:B:303:ALA:HB1	1.94	0.48
1:E:592:GLN:OE1	1:E:611:VAL:HG23	2.14	0.48
1:F:491:VAL:HG22	1:F:503:PHE:HE1	1.78	0.48
1:G:444:LEU:HB2	1:H:499:LEU:HD21	1.94	0.48
1:H:477:GLY:O	1:H:572:ARG:NH2	2.43	0.48
1:H:592:GLN:OE1	1:H:611:VAL:HG23	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:592:GLN:OE1	1:I:611:VAL:HG23	2.14	0.48
1:K:322:ARG:HG3	1:K:428:LEU:HD22	1.94	0.48
1:L:212:ARG:HH21	1:M:181:ASP:HB2	1.79	0.48
1:N:429:ASP:HA	1:N:472:PRO:HG2	1.95	0.48
1:B:136:TYR:CE1	1:B:142:LEU:HG	2.49	0.48
1:C:136:TYR:CE1	1:C:142:LEU:HG	2.49	0.48
1:D:592:GLN:OE1	1:D:611:VAL:HG23	2.14	0.48
1:E:429:ASP:HA	1:E:472:PRO:HG2	1.95	0.48
1:F:212:ARG:HH21	1:G:181:ASP:HB2	1.79	0.48
1:F:491:VAL:HG22	1:F:503:PHE:CE1	2.49	0.48
1:G:592:GLN:OE1	1:G:611:VAL:HG23	2.14	0.48
1:H:491:VAL:HG22	1:H:503:PHE:CE1	2.49	0.48
1:J:171:ILE:HB	1:J:217:ILE:HG13	1.96	0.48
1:K:171:ILE:HB	1:K:217:ILE:HG13	1.96	0.48
1:N:212:ARG:HH21	1:O:181:ASP:HB2	1.79	0.48
1:B:429:ASP:HA	1:B:472:PRO:HG2	1.95	0.48
1:B:444:LEU:HB2	1:C:499:LEU:HD21	1.94	0.48
1:C:491:VAL:HG22	1:C:503:PHE:CE1	2.49	0.48
1:F:592:GLN:OE1	1:F:611:VAL:HG23	2.14	0.48
1:G:491:VAL:HG22	1:G:503:PHE:HE1	1.78	0.48
1:I:108:PRO:HA	1:I:141:VAL:HA	1.96	0.48
1:J:108:PRO:HA	1:J:141:VAL:HA	1.96	0.48
1:J:592:GLN:OE1	1:J:611:VAL:HG23	2.14	0.48
1:K:491:VAL:HG22	1:K:503:PHE:CE1	2.49	0.48
1:L:491:VAL:HG22	1:L:503:PHE:CE1	2.49	0.48
1:M:491:VAL:HG22	1:M:503:PHE:CE1	2.49	0.48
1:A:557:ARG:HB2	1:N:607:GLU:OE2	2.13	0.48
1:D:109:LEU:HD13	1:D:162:ASP:HA	1.96	0.48
1:H:109:LEU:HD13	1:H:162:ASP:HA	1.96	0.48
1:H:136:TYR:CE1	1:H:142:LEU:HG	2.49	0.48
1:I:120:LEU:O	1:I:123:GLN:HB3	2.14	0.48
1:I:429:ASP:HA	1:I:472:PRO:HG2	1.95	0.48
1:J:288:VAL:HG22	1:J:303:ALA:HB1	1.94	0.48
1:J:429:ASP:HA	1:J:472:PRO:HG2	1.95	0.48
1:K:136:TYR:CE1	1:K:142:LEU:HG	2.49	0.48
1:L:491:VAL:HG22	1:L:503:PHE:HE1	1.78	0.48
1:M:171:ILE:HB	1:M:217:ILE:HG13	1.96	0.48
1:M:607:GLU:OE2	1:O:557:ARG:HB2	2.13	0.48
1:N:120:LEU:O	1:N:123:GLN:HB3	2.14	0.48
1:N:171:ILE:HB	1:N:217:ILE:HG13	1.96	0.48
1:N:491:VAL:HG22	1:N:503:PHE:CE1	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:109:LEU:HD13	1:A:162:ASP:HA	1.96	0.48
1:A:212:ARG:HH21	1:B:181:ASP:HB2	1.79	0.48
1:A:491:VAL:HG22	1:A:503:PHE:CE1	2.49	0.48
1:C:109:LEU:HD13	1:C:162:ASP:HA	1.96	0.48
1:F:136:TYR:CE1	1:F:142:LEU:HG	2.49	0.48
1:G:120:LEU:O	1:G:123:GLN:HB3	2.14	0.48
1:H:108:PRO:HA	1:H:141:VAL:HA	1.96	0.48
1:I:171:ILE:HB	1:I:217:ILE:HG13	1.96	0.48
1:J:212:ARG:HH21	1:K:181:ASP:HB2	1.79	0.48
1:J:246:ARG:HH22	1:J:305:GLN:HE22	1.61	0.48
1:K:250:LEU:HD13	1:K:258:LEU:HD12	1.95	0.48
1:L:171:ILE:HB	1:L:217:ILE:HG13	1.96	0.48
1:M:206:LYS:HZ3	1:N:190:ILE:C	2.22	0.48
1:O:491:VAL:HG22	1:O:503:PHE:CE1	2.49	0.48
1:B:109:LEU:HD13	1:B:162:ASP:HA	1.96	0.47
1:C:212:ARG:HH21	1:D:181:ASP:HB2	1.79	0.47
1:C:444:LEU:HB2	1:D:499:LEU:HD21	1.94	0.47
1:D:212:ARG:HH21	1:E:181:ASP:HB2	1.79	0.47
1:D:250:LEU:HD13	1:D:258:LEU:HD12	1.95	0.47
1:E:491:VAL:HG22	1:E:503:PHE:CE1	2.49	0.47
1:F:108:PRO:HA	1:F:141:VAL:HA	1.96	0.47
1:G:109:LEU:HD13	1:G:162:ASP:HA	1.96	0.47
1:G:171:ILE:HB	1:G:217:ILE:HG13	1.96	0.47
1:H:171:ILE:HB	1:H:217:ILE:HG13	1.96	0.47
1:H:444:LEU:HB2	1:I:499:LEU:HD21	1.94	0.47
1:J:491:VAL:HG22	1:J:503:PHE:CE1	2.49	0.47
1:K:108:PRO:HA	1:K:141:VAL:HA	1.96	0.47
1:B:607:GLU:OE2	1:D:557:ARG:HB2	2.13	0.47
1:C:106:ILE:HG22	1:C:143:ILE:HA	1.96	0.47
1:D:429:ASP:HA	1:D:472:PRO:HG2	1.95	0.47
1:D:491:VAL:HG22	1:D:503:PHE:CE1	2.49	0.47
1:E:109:LEU:HD13	1:E:162:ASP:HA	1.96	0.47
1:G:108:PRO:HA	1:G:141:VAL:HA	1.96	0.47
1:G:136:TYR:CE1	1:G:142:LEU:HG	2.49	0.47
1:I:212:ARG:HH21	1:J:181:ASP:HB2	1.79	0.47
1:J:109:LEU:HD13	1:J:162:ASP:HA	1.96	0.47
1:K:109:LEU:HD13	1:K:162:ASP:HA	1.96	0.47
1:K:592:GLN:OE1	1:K:611:VAL:HG23	2.14	0.47
1:L:120:LEU:O	1:L:123:GLN:HB3	2.14	0.47
1:N:491:VAL:HG22	1:N:503:PHE:HE1	1.78	0.47
1:A:120:LEU:O	1:A:123:GLN:HB3	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:592:GLN:OE1	1:A:611:VAL:HG23	2.14	0.47
1:B:106:ILE:HG22	1:B:143:ILE:HA	1.96	0.47
1:C:246:ARG:HH22	1:C:305:GLN:HE22	1.61	0.47
1:D:108:PRO:HA	1:D:141:VAL:HA	1.96	0.47
1:D:120:LEU:O	1:D:123:GLN:HB3	2.14	0.47
1:E:108:PRO:HA	1:E:141:VAL:HA	1.96	0.47
1:F:120:LEU:O	1:F:123:GLN:HB3	2.14	0.47
1:H:491:VAL:HG22	1:H:503:PHE:HE1	1.78	0.47
1:I:109:LEU:HD13	1:I:162:ASP:HA	1.96	0.47
1:K:120:LEU:O	1:K:123:GLN:HB3	2.14	0.47
1:L:104:THR:OG1	1:L:144:LEU:O	2.21	0.47
1:L:136:TYR:CE1	1:L:142:LEU:HG	2.49	0.47
1:L:262:LEU:CD1	1:L:299:LEU:HD21	2.40	0.47
1:M:429:ASP:HA	1:M:472:PRO:HG2	1.95	0.47
1:O:136:TYR:CE1	1:O:142:LEU:HG	2.49	0.47
1:O:171:ILE:HB	1:O:217:ILE:HG13	1.96	0.47
1:B:491:VAL:HG22	1:B:503:PHE:CE1	2.49	0.47
1:C:108:PRO:HA	1:C:141:VAL:HA	1.96	0.47
1:C:120:LEU:O	1:C:123:GLN:HB3	2.14	0.47
1:C:250:LEU:HD13	1:C:258:LEU:HD12	1.95	0.47
1:D:246:ARG:HH22	1:D:305:GLN:HE22	1.61	0.47
1:E:246:ARG:HH22	1:E:305:GLN:HE22	1.61	0.47
1:G:212:ARG:HH21	1:H:181:ASP:HB2	1.79	0.47
1:M:477:GLY:O	1:M:572:ARG:NH2	2.43	0.47
1:O:109:LEU:HD13	1:O:162:ASP:HA	1.96	0.47
1:A:106:ILE:HG22	1:A:143:ILE:HA	1.96	0.47
1:B:108:PRO:HA	1:B:141:VAL:HA	1.96	0.47
1:C:592:GLN:OE1	1:C:611:VAL:HG23	2.14	0.47
1:D:106:ILE:HG22	1:D:143:ILE:HA	1.97	0.47
1:D:136:TYR:CE1	1:D:142:LEU:HG	2.49	0.47
1:F:109:LEU:HD13	1:F:162:ASP:HA	1.96	0.47
1:F:171:ILE:HB	1:F:217:ILE:HG13	1.96	0.47
1:L:108:PRO:HA	1:L:141:VAL:HA	1.96	0.47
1:L:592:GLN:OE1	1:L:611:VAL:HG23	2.14	0.47
1:M:109:LEU:HD13	1:M:162:ASP:HA	1.96	0.47
1:N:109:LEU:HD13	1:N:162:ASP:HA	1.96	0.47
1:A:108:PRO:HA	1:A:141:VAL:HA	1.96	0.47
1:E:120:LEU:O	1:E:123:GLN:HB3	2.14	0.47
1:E:491:VAL:HG22	1:E:503:PHE:HE1	1.78	0.47
1:M:136:TYR:CE1	1:M:142:LEU:HG	2.49	0.47
1:O:108:PRO:HA	1:O:141:VAL:HA	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:171:ILE:HB	1:A:217:ILE:HG13	1.96	0.47
1:A:491:VAL:HG22	1:A:503:PHE:HE1	1.78	0.47
1:B:171:ILE:HB	1:B:217:ILE:HG13	1.96	0.47
1:B:592:GLN:OE1	1:B:611:VAL:HG23	2.14	0.47
1:C:182:LEU:HA	1:C:185:ILE:HG12	1.97	0.47
1:C:429:ASP:HA	1:C:472:PRO:HG2	1.95	0.47
1:D:182:LEU:HA	1:D:185:ILE:HG12	1.97	0.47
1:D:235:LEU:HD23	1:D:237:VAL:HG12	1.97	0.47
1:G:242:GLU:HG2	1:G:243:GLY:H	1.60	0.47
1:G:246:ARG:HH22	1:G:305:GLN:HE22	1.61	0.47
1:G:491:VAL:HG22	1:G:503:PHE:CE1	2.49	0.47
1:H:212:ARG:HH21	1:I:181:ASP:HB2	1.79	0.47
1:I:129:SER:HB3	1:I:147:ARG:HH22	1.80	0.47
1:I:491:VAL:HG22	1:I:503:PHE:CE1	2.49	0.47
1:J:136:TYR:CE1	1:J:142:LEU:HG	2.49	0.47
1:M:108:PRO:HA	1:M:141:VAL:HA	1.96	0.47
1:M:592:GLN:OE1	1:M:611:VAL:HG23	2.14	0.47
1:N:108:PRO:HA	1:N:141:VAL:HA	1.96	0.47
1:N:136:TYR:CE1	1:N:142:LEU:HG	2.49	0.47
1:O:106:ILE:HG22	1:O:143:ILE:HA	1.96	0.47
1:O:592:GLN:OE1	1:O:611:VAL:HG23	2.14	0.47
1:A:136:TYR:CE1	1:A:142:LEU:HG	2.49	0.47
1:C:491:VAL:HG22	1:C:503:PHE:HE1	1.78	0.47
1:D:171:ILE:HB	1:D:217:ILE:HG13	1.96	0.47
1:E:106:ILE:HG22	1:E:143:ILE:HA	1.97	0.47
1:E:171:ILE:HB	1:E:217:ILE:HG13	1.96	0.47
1:E:206:LYS:HZ3	1:F:190:ILE:C	2.23	0.47
1:J:118:ALA:HB1	1:J:122:ARG:NH2	2.30	0.47
1:J:491:VAL:HG22	1:J:503:PHE:HE1	1.78	0.47
1:L:109:LEU:HD13	1:L:162:ASP:HA	1.96	0.47
1:M:246:ARG:HH22	1:M:305:GLN:HE22	1.61	0.47
1:N:118:ALA:HB1	1:N:122:ARG:NH2	2.30	0.47
1:N:246:ARG:HH22	1:N:305:GLN:HE22	1.61	0.47
1:A:182:LEU:HA	1:A:185:ILE:HG12	1.97	0.47
1:B:120:LEU:O	1:B:123:GLN:HB3	2.14	0.47
1:B:182:LEU:HA	1:B:185:ILE:HG12	1.97	0.47
1:C:235:LEU:HD23	1:C:237:VAL:HG12	1.97	0.47
1:E:212:ARG:HH21	1:F:181:ASP:HB2	1.79	0.47
1:E:235:LEU:HD23	1:E:237:VAL:HG12	1.97	0.47
1:H:235:LEU:HD23	1:H:237:VAL:HG12	1.97	0.47
1:I:136:TYR:CE1	1:I:142:LEU:HG	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:120:LEU:O	1:J:123:GLN:HB3	2.14	0.47
1:K:212:ARG:HH21	1:L:181:ASP:HB2	1.79	0.47
1:L:129:SER:HB3	1:L:147:ARG:HH22	1.80	0.47
1:M:118:ALA:HB1	1:M:122:ARG:NH2	2.30	0.47
1:M:120:LEU:O	1:M:123:GLN:HB3	2.14	0.47
1:M:129:SER:HB3	1:M:147:ARG:HH22	1.80	0.47
1:C:118:ALA:HB1	1:C:122:ARG:NH2	2.30	0.47
1:D:118:ALA:HB1	1:D:122:ARG:NH2	2.30	0.47
1:F:235:LEU:HD23	1:F:237:VAL:HG12	1.97	0.47
1:G:235:LEU:HD23	1:G:237:VAL:HG12	1.97	0.47
1:H:120:LEU:O	1:H:123:GLN:HB3	2.14	0.47
1:J:129:SER:HB3	1:J:147:ARG:HH22	1.80	0.47
1:K:118:ALA:HB1	1:K:122:ARG:NH2	2.30	0.47
1:N:592:GLN:OE1	1:N:611:VAL:HG23	2.14	0.47
1:O:182:LEU:HA	1:O:185:ILE:HG12	1.97	0.47
1:A:220:GLY:HA3	1:A:224:ALA:HB3	1.98	0.46
1:A:246:ARG:HH22	1:A:305:GLN:HE22	1.61	0.46
1:E:182:LEU:HA	1:E:185:ILE:HG12	1.97	0.46
1:F:106:ILE:HG22	1:F:143:ILE:HA	1.96	0.46
1:G:182:LEU:HA	1:G:185:ILE:HG12	1.97	0.46
1:I:224:ALA:HA	1:I:227:ARG:NE	2.30	0.46
1:J:106:ILE:HG22	1:J:143:ILE:HA	1.96	0.46
1:L:111:ASN:HB3	1:L:165:GLY:O	2.16	0.46
1:M:111:ASN:HB3	1:M:165:GLY:O	2.16	0.46
1:A:181:ASP:HB2	1:O:212:ARG:HH21	1.79	0.46
1:B:220:GLY:HA3	1:B:224:ALA:HB3	1.98	0.46
1:D:129:SER:HB3	1:D:147:ARG:HH22	1.80	0.46
1:E:111:ASN:HB3	1:E:165:GLY:O	2.16	0.46
1:E:118:ALA:HB1	1:E:122:ARG:NH2	2.30	0.46
1:F:182:LEU:HA	1:F:185:ILE:HG12	1.97	0.46
1:G:106:ILE:HG22	1:G:143:ILE:HA	1.96	0.46
1:G:224:ALA:HA	1:G:227:ARG:NE	2.31	0.46
1:I:242:GLU:HG2	1:I:243:GLY:H	1.60	0.46
1:K:106:ILE:HG22	1:K:143:ILE:HA	1.96	0.46
1:K:206:LYS:HZ3	1:L:190:ILE:C	2.23	0.46
1:L:118:ALA:HB1	1:L:122:ARG:NH2	2.30	0.46
1:M:212:ARG:HH21	1:N:181:ASP:HB2	1.79	0.46
1:M:220:GLY:HA3	1:M:224:ALA:HB3	1.98	0.46
1:N:220:GLY:HA3	1:N:224:ALA:HB3	1.98	0.46
1:O:118:ALA:HB1	1:O:122:ARG:NH2	2.30	0.46
1:A:111:ASN:HB3	1:A:165:GLY:O	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:235:LEU:HD23	1:A:237:VAL:HG12	1.97	0.46
1:B:111:ASN:HB3	1:B:165:GLY:O	2.16	0.46
1:B:118:ALA:HB1	1:B:122:ARG:NH2	2.30	0.46
1:C:171:ILE:HB	1:C:217:ILE:HG13	1.96	0.46
1:D:133:VAL:HB	1:D:145:THR:HB	1.98	0.46
1:D:262:LEU:CD1	1:D:299:LEU:HD21	2.40	0.46
1:E:133:VAL:HB	1:E:145:THR:HB	1.98	0.46
1:F:129:SER:HB3	1:F:147:ARG:HH22	1.80	0.46
1:K:111:ASN:HB3	1:K:165:GLY:O	2.16	0.46
1:K:224:ALA:HA	1:K:227:ARG:NE	2.30	0.46
1:K:251:LYS:HB3	1:L:476:GLU:HB2	1.98	0.46
1:M:251:LYS:HB3	1:N:476:GLU:HB2	1.98	0.46
1:N:111:ASN:HB3	1:N:165:GLY:O	2.16	0.46
1:N:224:ALA:HA	1:N:227:ARG:NE	2.30	0.46
1:O:220:GLY:HA3	1:O:224:ALA:HB3	1.98	0.46
1:O:224:ALA:HA	1:O:227:ARG:NE	2.30	0.46
1:A:476:GLU:HB2	1:O:251:LYS:HB3	1.98	0.46
1:B:129:SER:HB3	1:B:147:ARG:HH22	1.80	0.46
1:B:235:LEU:HD23	1:B:237:VAL:HG12	1.97	0.46
1:B:246:ARG:HH22	1:B:305:GLN:HE22	1.61	0.46
1:C:220:GLY:HA3	1:C:224:ALA:HB3	1.98	0.46
1:I:118:ALA:HB1	1:I:122:ARG:NH2	2.30	0.46
1:I:235:LEU:HD23	1:I:237:VAL:HG12	1.97	0.46
1:I:251:LYS:HB3	1:J:476:GLU:HB2	1.98	0.46
1:J:111:ASN:HB3	1:J:165:GLY:O	2.16	0.46
1:K:246:ARG:HH22	1:K:305:GLN:HE22	1.61	0.46
1:N:106:ILE:HG22	1:N:143:ILE:HA	1.97	0.46
1:N:182:LEU:HA	1:N:185:ILE:HG12	1.97	0.46
1:A:224:ALA:HA	1:A:227:ARG:NE	2.30	0.46
1:C:133:VAL:HB	1:C:145:THR:HB	1.98	0.46
1:E:259:VAL:HG22	1:E:299:LEU:CD1	2.46	0.46
1:F:111:ASN:HB3	1:F:165:GLY:O	2.16	0.46
1:F:118:ALA:HB1	1:F:122:ARG:NH2	2.30	0.46
1:F:259:VAL:HG22	1:F:299:LEU:CD1	2.46	0.46
1:G:489:SER:HA	1:G:504:ASN:O	2.16	0.46
1:L:220:GLY:HA3	1:L:224:ALA:HB3	1.98	0.46
1:L:259:VAL:HG22	1:L:299:LEU:CD1	2.46	0.46
1:M:224:ALA:HA	1:M:227:ARG:NE	2.31	0.46
1:O:111:ASN:HB3	1:O:165:GLY:O	2.16	0.46
1:O:259:VAL:HG22	1:O:299:LEU:CD1	2.46	0.46
1:B:259:VAL:HG22	1:B:299:LEU:CD1	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:220:GLY:HA3	1:D:224:ALA:HB3	1.97	0.46
1:E:224:ALA:HA	1:E:227:ARG:NE	2.30	0.46
1:F:133:VAL:HB	1:F:145:THR:HB	1.98	0.46
1:F:489:SER:HA	1:F:504:ASN:O	2.16	0.46
1:H:106:ILE:HG22	1:H:143:ILE:HA	1.96	0.46
1:I:106:ILE:HG22	1:I:143:ILE:HA	1.97	0.46
1:I:111:ASN:HB3	1:I:165:GLY:O	2.16	0.46
1:I:259:VAL:HG22	1:I:299:LEU:CD1	2.46	0.46
1:L:106:ILE:HG22	1:L:143:ILE:HA	1.96	0.46
1:L:224:ALA:HA	1:L:227:ARG:NE	2.30	0.46
1:O:120:LEU:O	1:O:123:GLN:HB3	2.14	0.46
1:A:489:SER:HA	1:A:504:ASN:O	2.16	0.46
1:B:212:ARG:HH21	1:C:181:ASP:HB2	1.79	0.46
1:B:224:ALA:HA	1:B:227:ARG:NE	2.31	0.46
1:D:259:VAL:HG22	1:D:299:LEU:CD1	2.46	0.46
1:E:220:GLY:HA3	1:E:224:ALA:HB3	1.98	0.46
1:G:111:ASN:HB3	1:G:165:GLY:O	2.16	0.46
1:G:129:SER:HB3	1:G:147:ARG:HH22	1.80	0.46
1:H:111:ASN:HB3	1:H:165:GLY:O	2.16	0.46
1:I:220:GLY:HA3	1:I:224:ALA:HB3	1.98	0.46
1:K:220:GLY:HA3	1:K:224:ALA:HB3	1.98	0.46
1:O:235:LEU:HD23	1:O:237:VAL:HG12	1.97	0.46
1:A:129:SER:HB3	1:A:147:ARG:HH22	1.80	0.46
1:B:170:GLN:HG3	1:B:218:ILE:HG23	1.98	0.46
1:C:170:GLN:HG3	1:C:218:ILE:HG23	1.98	0.46
1:C:224:ALA:HA	1:C:227:ARG:NE	2.30	0.46
1:C:259:VAL:HG22	1:C:299:LEU:CD1	2.46	0.46
1:F:224:ALA:HA	1:F:227:ARG:NE	2.30	0.46
1:G:118:ALA:HB1	1:G:122:ARG:NH2	2.30	0.46
1:G:220:GLY:HA3	1:G:224:ALA:HB3	1.98	0.46
1:G:259:VAL:HG22	1:G:299:LEU:CD1	2.46	0.46
1:H:220:GLY:HA3	1:H:224:ALA:HB3	1.98	0.46
1:J:220:GLY:HA3	1:J:224:ALA:HB3	1.98	0.46
1:J:235:LEU:HD23	1:J:237:VAL:HG12	1.97	0.46
1:J:438:GLY:HA3	1:J:465:GLY:HA3	1.98	0.46
1:K:259:VAL:HG22	1:K:299:LEU:CD1	2.46	0.46
1:N:147:ARG:O	1:N:151:ILE:HG12	2.16	0.46
1:N:259:VAL:HG22	1:N:299:LEU:CD1	2.46	0.46
1:O:129:SER:HB3	1:O:147:ARG:HH22	1.80	0.46
1:O:489:SER:HA	1:O:504:ASN:O	2.16	0.46
1:D:170:GLN:HG3	1:D:218:ILE:HG23	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:220:GLY:HA3	1:F:224:ALA:HB3	1.98	0.46
1:G:133:VAL:HB	1:G:145:THR:HB	1.98	0.46
1:G:474:VAL:HG22	1:G:480:VAL:HG12	1.98	0.46
1:H:118:ALA:HB1	1:H:122:ARG:NH2	2.30	0.46
1:H:129:SER:HB3	1:H:147:ARG:HH22	1.80	0.46
1:H:182:LEU:HA	1:H:185:ILE:HG12	1.97	0.46
1:H:224:ALA:HA	1:H:227:ARG:NE	2.30	0.46
1:I:246:ARG:HH22	1:I:305:GLN:HE22	1.61	0.46
1:I:438:GLY:HA3	1:I:465:GLY:HA3	1.98	0.46
1:J:224:ALA:HA	1:J:227:ARG:NE	2.30	0.46
1:K:129:SER:HB3	1:K:147:ARG:HH22	1.80	0.46
1:K:235:LEU:HD23	1:K:237:VAL:HG12	1.97	0.46
1:L:147:ARG:O	1:L:151:ILE:HG12	2.16	0.46
1:L:246:ARG:HH22	1:L:305:GLN:HE22	1.61	0.46
1:M:182:LEU:HA	1:M:185:ILE:HG12	1.97	0.46
1:A:118:ALA:HB1	1:A:122:ARG:NH2	2.30	0.46
1:B:133:VAL:HB	1:B:145:THR:HB	1.98	0.46
1:B:474:VAL:HG22	1:B:480:VAL:HG12	1.98	0.46
1:B:489:SER:HA	1:B:504:ASN:O	2.16	0.46
1:D:111:ASN:HB3	1:D:165:GLY:O	2.16	0.46
1:D:224:ALA:HA	1:D:227:ARG:NE	2.31	0.46
1:H:489:SER:HA	1:H:504:ASN:O	2.16	0.46
1:K:438:GLY:HA3	1:K:465:GLY:HA3	1.98	0.46
1:M:106:ILE:HG22	1:M:143:ILE:HA	1.96	0.46
1:N:474:VAL:HG22	1:N:480:VAL:HG12	1.98	0.46
1:O:474:VAL:HG22	1:O:480:VAL:HG12	1.98	0.46
1:A:170:GLN:HG3	1:A:218:ILE:HG23	1.98	0.45
1:B:251:LYS:HB3	1:C:476:GLU:HB2	1.98	0.45
1:C:111:ASN:HB3	1:C:165:GLY:O	2.16	0.45
1:D:474:VAL:HG22	1:D:480:VAL:HG12	1.98	0.45
1:E:170:GLN:HG3	1:E:218:ILE:HG23	1.98	0.45
1:E:489:SER:HA	1:E:504:ASN:O	2.16	0.45
1:G:251:LYS:HB3	1:H:476:GLU:HB2	1.98	0.45
1:H:259:VAL:HG22	1:H:299:LEU:CD1	2.46	0.45
1:H:474:VAL:HG22	1:H:480:VAL:HG12	1.98	0.45
1:I:182:LEU:HA	1:I:185:ILE:HG12	1.97	0.45
1:K:489:SER:HA	1:K:504:ASN:O	2.16	0.45
1:L:182:LEU:HA	1:L:185:ILE:HG12	1.97	0.45
1:N:129:SER:HB3	1:N:147:ARG:HH22	1.80	0.45
1:N:206:LYS:HZ3	1:O:190:ILE:C	2.24	0.45
1:O:246:ARG:HH22	1:O:305:GLN:HE22	1.61	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:147:ARG:O	1:A:151:ILE:HG12	2.16	0.45
1:A:474:VAL:HG22	1:A:480:VAL:HG12	1.98	0.45
1:C:129:SER:HB3	1:C:147:ARG:HH22	1.80	0.45
1:C:438:GLY:HA3	1:C:465:GLY:HA3	1.98	0.45
1:D:438:GLY:HA3	1:D:465:GLY:HA3	1.98	0.45
1:E:474:VAL:HG22	1:E:480:VAL:HG12	1.98	0.45
1:F:474:VAL:HG22	1:F:480:VAL:HG12	1.98	0.45
1:G:222:GLU:O	1:G:225:ARG:HB2	2.17	0.45
1:H:133:VAL:HB	1:H:145:THR:HB	1.98	0.45
1:I:118:ALA:HB3	1:I:119:PRO:HD3	1.98	0.45
1:I:475:ASN:HA	1:I:476:GLU:HA	1.78	0.45
1:J:118:ALA:HB3	1:J:119:PRO:HD3	1.98	0.45
1:J:182:LEU:HA	1:J:185:ILE:HG12	1.97	0.45
1:J:474:VAL:HG22	1:J:480:VAL:HG12	1.98	0.45
1:K:222:GLU:O	1:K:225:ARG:HB2	2.17	0.45
1:K:474:VAL:HG22	1:K:480:VAL:HG12	1.98	0.45
1:L:347:ALA:HA	1:L:353:ALA:HA	1.99	0.45
1:L:489:SER:HA	1:L:504:ASN:O	2.16	0.45
1:M:235:LEU:HD23	1:M:237:VAL:HG12	1.97	0.45
1:M:347:ALA:HA	1:M:353:ALA:HA	1.99	0.45
1:N:235:LEU:HD23	1:N:237:VAL:HG12	1.97	0.45
1:N:489:SER:HA	1:N:504:ASN:O	2.16	0.45
1:C:222:GLU:O	1:C:225:ARG:HB2	2.17	0.45
1:D:222:GLU:O	1:D:225:ARG:HB2	2.17	0.45
1:I:133:VAL:HB	1:I:145:THR:HB	1.98	0.45
1:J:206:LYS:HZ3	1:K:190:ILE:C	2.24	0.45
1:J:347:ALA:HA	1:J:353:ALA:HA	1.99	0.45
1:K:182:LEU:HA	1:K:185:ILE:HG12	1.97	0.45
1:A:133:VAL:HB	1:A:145:THR:HB	1.98	0.45
1:A:190:ILE:C	1:O:206:LYS:HZ3	2.24	0.45
1:B:161:VAL:HA	1:B:164:ILE:CG1	2.47	0.45
1:F:170:GLN:HG3	1:F:218:ILE:HG23	1.98	0.45
1:H:118:ALA:HB3	1:H:119:PRO:HD3	1.98	0.45
1:H:222:GLU:O	1:H:225:ARG:HB2	2.17	0.45
1:H:438:GLY:HA3	1:H:465:GLY:HA3	1.98	0.45
1:I:474:VAL:HG22	1:I:480:VAL:HG12	1.98	0.45
1:J:147:ARG:O	1:J:151:ILE:HG12	2.16	0.45
1:J:259:VAL:HG22	1:J:299:LEU:CD1	2.46	0.45
1:K:109:LEU:HD21	1:K:112:VAL:HB	1.99	0.45
1:M:474:VAL:HG22	1:M:480:VAL:HG12	1.98	0.45
1:B:206:LYS:HZ3	1:C:190:ILE:C	2.24	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:222:GLU:O	1:B:225:ARG:HB2	2.17	0.45
1:C:474:VAL:HG22	1:C:480:VAL:HG12	1.98	0.45
1:D:489:SER:HA	1:D:504:ASN:O	2.16	0.45
1:D:529:PHE:HZ	1:D:531:LYS:HE3	1.82	0.45
1:E:222:GLU:O	1:E:225:ARG:HB2	2.17	0.45
1:E:438:GLY:HA3	1:E:465:GLY:HA3	1.98	0.45
1:H:104:THR:OG1	1:H:144:LEU:O	2.21	0.45
1:I:489:SER:HA	1:I:504:ASN:O	2.16	0.45
1:K:347:ALA:HA	1:K:353:ALA:HA	1.99	0.45
1:L:133:VAL:HB	1:L:145:THR:HB	1.98	0.45
1:L:438:GLY:HA3	1:L:465:GLY:HA3	1.98	0.45
1:L:474:VAL:HG22	1:L:480:VAL:HG12	1.98	0.45
1:M:133:VAL:HB	1:M:145:THR:HB	1.98	0.45
1:N:161:VAL:HA	1:N:164:ILE:CG1	2.46	0.45
1:O:347:ALA:HA	1:O:353:ALA:HA	1.99	0.45
1:A:161:VAL:HA	1:A:164:ILE:CG1	2.47	0.45
1:A:222:GLU:O	1:A:225:ARG:HB2	2.17	0.45
1:A:529:PHE:HZ	1:A:531:LYS:HE3	1.82	0.45
1:B:438:GLY:HA3	1:B:465:GLY:HA3	1.98	0.45
1:G:170:GLN:HG3	1:G:218:ILE:HG23	1.98	0.45
1:I:147:ARG:O	1:I:151:ILE:HG12	2.16	0.45
1:J:222:GLU:O	1:J:225:ARG:HB2	2.17	0.45
1:K:133:VAL:HB	1:K:145:THR:HB	1.98	0.45
1:L:222:GLU:O	1:L:225:ARG:HB2	2.17	0.45
1:L:235:LEU:HD23	1:L:237:VAL:HG12	1.97	0.45
1:M:220:GLY:HA3	1:M:224:ALA:CB	2.47	0.45
1:M:529:PHE:HZ	1:M:531:LYS:HE3	1.82	0.45
1:N:251:LYS:HB3	1:O:476:GLU:HB2	1.98	0.45
1:N:347:ALA:HA	1:N:353:ALA:HA	1.99	0.45
1:O:161:VAL:HA	1:O:164:ILE:CG1	2.47	0.45
1:O:170:GLN:HG3	1:O:218:ILE:HG23	1.98	0.45
1:B:529:PHE:HZ	1:B:531:LYS:HE3	1.82	0.45
1:D:147:ARG:O	1:D:151:ILE:HG12	2.16	0.45
1:D:220:GLY:HA3	1:D:224:ALA:CB	2.47	0.45
1:G:109:LEU:HD21	1:G:112:VAL:HB	1.99	0.45
1:H:109:LEU:HD21	1:H:112:VAL:HB	1.99	0.45
1:H:161:VAL:HA	1:H:164:ILE:CG1	2.46	0.45
1:H:220:GLY:HA3	1:H:224:ALA:CB	2.47	0.45
1:I:161:VAL:HA	1:I:164:ILE:CG1	2.46	0.45
1:J:133:VAL:HB	1:J:145:THR:HB	1.98	0.45
1:K:170:GLN:HG3	1:K:218:ILE:HG23	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:220:GLY:HA3	1:K:224:ALA:CB	2.47	0.45
1:K:529:PHE:HZ	1:K:531:LYS:HE3	1.82	0.45
1:L:220:GLY:HA3	1:L:224:ALA:CB	2.47	0.45
1:L:251:LYS:HB3	1:M:476:GLU:HB2	1.98	0.45
1:M:161:VAL:HA	1:M:164:ILE:CG1	2.47	0.45
1:N:109:LEU:HD11	1:N:112:VAL:HB	1.99	0.45
1:N:529:PHE:HZ	1:N:531:LYS:HE3	1.82	0.45
1:O:109:LEU:HD21	1:O:112:VAL:HB	1.99	0.45
1:A:438:GLY:HA3	1:A:465:GLY:HA3	1.98	0.45
1:B:346:TRP:O	1:B:346:TRP:CE3	2.70	0.45
1:C:489:SER:HA	1:C:504:ASN:O	2.16	0.45
1:D:251:LYS:HB3	1:E:476:GLU:HB2	1.98	0.45
1:F:222:GLU:O	1:F:225:ARG:HB2	2.17	0.45
1:G:118:ALA:HB3	1:G:119:PRO:HD3	1.98	0.45
1:G:147:ARG:O	1:G:151:ILE:HG12	2.16	0.45
1:G:220:GLY:HA3	1:G:224:ALA:CB	2.47	0.45
1:I:220:GLY:HA3	1:I:224:ALA:CB	2.47	0.45
1:I:347:ALA:HA	1:I:353:ALA:HA	1.99	0.45
1:J:109:LEU:HD21	1:J:112:VAL:HB	1.99	0.45
1:J:170:GLN:HG3	1:J:218:ILE:HG23	1.98	0.45
1:J:220:GLY:HA3	1:J:224:ALA:CB	2.47	0.45
1:J:251:LYS:HB3	1:K:476:GLU:HB2	1.98	0.45
1:K:118:ALA:HB3	1:K:119:PRO:HD3	1.98	0.45
1:K:147:ARG:O	1:K:151:ILE:HG12	2.16	0.45
1:M:109:LEU:HD11	1:M:112:VAL:HB	1.99	0.45
1:M:259:VAL:HG22	1:M:299:LEU:CD1	2.46	0.45
1:N:109:LEU:HD21	1:N:112:VAL:HB	1.99	0.45
1:O:109:LEU:HD11	1:O:112:VAL:HB	1.99	0.45
1:A:259:VAL:HG22	1:A:299:LEU:CD1	2.46	0.45
1:A:347:ALA:HA	1:A:353:ALA:HA	1.99	0.45
1:B:104:THR:OG1	1:B:144:LEU:O	2.21	0.45
1:E:147:ARG:O	1:E:151:ILE:HG12	2.16	0.45
1:F:438:GLY:HA3	1:F:465:GLY:HA3	1.98	0.45
1:F:529:PHE:HZ	1:F:531:LYS:HE3	1.82	0.45
1:H:346:TRP:O	1:H:346:TRP:CE3	2.70	0.45
1:I:346:TRP:CE3	1:I:346:TRP:O	2.70	0.45
1:J:489:SER:HA	1:J:504:ASN:O	2.16	0.45
1:J:529:PHE:HZ	1:J:531:LYS:HE3	1.82	0.45
1:L:109:LEU:HD11	1:L:112:VAL:HB	1.99	0.45
1:O:222:GLU:O	1:O:225:ARG:HB2	2.17	0.45
1:A:118:ALA:HB3	1:A:119:PRO:HD3	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:346:TRP:O	1:C:346:TRP:CE3	2.70	0.45
1:E:129:SER:HB3	1:E:147:ARG:HH22	1.80	0.45
1:E:251:LYS:HB3	1:F:476:GLU:HB2	1.98	0.45
1:H:170:GLN:HG3	1:H:218:ILE:HG23	1.98	0.45
1:I:109:LEU:HD11	1:I:112:VAL:HB	1.99	0.45
1:L:109:LEU:HD21	1:L:112:VAL:HB	1.99	0.45
1:L:170:GLN:HG3	1:L:218:ILE:HG23	1.98	0.45
1:N:133:VAL:HB	1:N:145:THR:HB	1.98	0.45
1:N:220:GLY:HA3	1:N:224:ALA:CB	2.47	0.45
1:N:414:ASN:HB3	1:O:534:VAL:CG2	2.47	0.45
1:O:438:GLY:HA3	1:O:465:GLY:HA3	1.98	0.45
1:A:109:LEU:HD11	1:A:112:VAL:HB	1.99	0.44
1:B:147:ARG:O	1:B:151:ILE:HG12	2.16	0.44
1:B:151:ILE:HA	1:B:154:LEU:HB2	2.00	0.44
1:C:414:ASN:HB3	1:D:534:VAL:CG2	2.47	0.44
1:E:346:TRP:O	1:E:346:TRP:CE3	2.70	0.44
1:E:414:ASN:HB3	1:F:534:VAL:CG2	2.47	0.44
1:F:346:TRP:CE3	1:F:346:TRP:O	2.70	0.44
1:H:347:ALA:HA	1:H:353:ALA:HA	1.99	0.44
1:H:529:PHE:HZ	1:H:531:LYS:HE3	1.82	0.44
1:I:170:GLN:HG3	1:I:218:ILE:HG23	1.98	0.44
1:J:109:LEU:HD11	1:J:112:VAL:HB	1.99	0.44
1:J:414:ASN:HB3	1:K:534:VAL:CG2	2.47	0.44
1:K:109:LEU:HD11	1:K:112:VAL:HB	1.99	0.44
1:M:147:ARG:O	1:M:151:ILE:HG12	2.16	0.44
1:M:346:TRP:O	1:M:346:TRP:CE3	2.70	0.44
1:M:489:SER:HA	1:M:504:ASN:O	2.16	0.44
1:O:118:ALA:HB3	1:O:119:PRO:HD3	1.98	0.44
1:O:133:VAL:HB	1:O:145:THR:HB	1.98	0.44
1:A:109:LEU:HD21	1:A:112:VAL:HB	1.99	0.44
1:A:534:VAL:CG2	1:O:414:ASN:HB3	2.47	0.44
1:E:151:ILE:HA	1:E:154:LEU:HB2	2.00	0.44
1:E:220:GLY:HA3	1:E:224:ALA:CB	2.47	0.44
1:F:109:LEU:HD21	1:F:112:VAL:HB	1.99	0.44
1:F:251:LYS:HB3	1:G:476:GLU:HB2	1.98	0.44
1:H:109:LEU:HD11	1:H:112:VAL:HB	1.99	0.44
1:H:147:ARG:O	1:H:151:ILE:HG12	2.16	0.44
1:H:251:LYS:HB3	1:I:476:GLU:HB2	1.98	0.44
1:I:225:ARG:O	1:I:229:THR:HG23	2.18	0.44
1:J:161:VAL:HA	1:J:164:ILE:CG1	2.47	0.44
1:J:346:TRP:O	1:J:346:TRP:CE3	2.70	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:475:ASN:HA	1:J:476:GLU:HA	1.78	0.44
1:M:414:ASN:HB3	1:N:534:VAL:CG2	2.47	0.44
1:M:438:GLY:HA3	1:M:465:GLY:HA3	1.98	0.44
1:N:170:GLN:HG3	1:N:218:ILE:HG23	1.98	0.44
1:B:118:ALA:HB3	1:B:119:PRO:HD3	1.98	0.44
1:B:347:ALA:HA	1:B:353:ALA:HA	1.99	0.44
1:C:118:ALA:HB3	1:C:119:PRO:HD3	1.98	0.44
1:C:220:GLY:HA3	1:C:224:ALA:CB	2.47	0.44
1:D:151:ILE:HA	1:D:154:LEU:HB2	2.00	0.44
1:E:161:VAL:HA	1:E:164:ILE:CG1	2.46	0.44
1:F:220:GLY:HA3	1:F:224:ALA:CB	2.47	0.44
1:G:161:VAL:HA	1:G:164:ILE:CG1	2.46	0.44
1:G:346:TRP:O	1:G:346:TRP:CE3	2.70	0.44
1:G:347:ALA:HA	1:G:353:ALA:HA	1.99	0.44
1:J:225:ARG:O	1:J:229:THR:HG23	2.18	0.44
1:K:414:ASN:HB3	1:L:534:VAL:CG2	2.47	0.44
1:L:414:ASN:HB3	1:M:534:VAL:CG2	2.47	0.44
1:N:222:GLU:O	1:N:225:ARG:HB2	2.17	0.44
1:N:346:TRP:CE3	1:N:346:TRP:O	2.70	0.44
1:O:220:GLY:HA3	1:O:224:ALA:CB	2.47	0.44
1:A:220:GLY:HA3	1:A:224:ALA:CB	2.47	0.44
1:A:251:LYS:HB3	1:B:476:GLU:HB2	1.98	0.44
1:C:151:ILE:HA	1:C:154:LEU:HB2	2.00	0.44
1:D:118:ALA:HB3	1:D:119:PRO:HD3	1.98	0.44
1:D:161:VAL:HA	1:D:164:ILE:CG1	2.47	0.44
1:G:151:ILE:HA	1:G:154:LEU:HB2	2.00	0.44
1:G:324:GLN:HB2	1:G:571:ILE:HB	1.99	0.44
1:G:438:GLY:HA3	1:G:465:GLY:HA3	1.98	0.44
1:I:529:PHE:HZ	1:I:531:LYS:HE3	1.82	0.44
1:K:161:VAL:HA	1:K:164:ILE:CG1	2.47	0.44
1:N:118:ALA:HB3	1:N:119:PRO:HD3	1.98	0.44
1:O:151:ILE:HA	1:O:154:LEU:HB2	2.00	0.44
1:O:346:TRP:CE3	1:O:346:TRP:O	2.70	0.44
1:B:220:GLY:HA3	1:B:224:ALA:CB	2.47	0.44
1:C:347:ALA:HA	1:C:353:ALA:HA	1.99	0.44
1:E:225:ARG:O	1:E:229:THR:HG23	2.18	0.44
1:E:529:PHE:HZ	1:E:531:LYS:HE3	1.82	0.44
1:G:109:LEU:HD11	1:G:112:VAL:HB	1.99	0.44
1:H:324:GLN:HB2	1:H:571:ILE:HB	1.99	0.44
1:K:225:ARG:O	1:K:229:THR:HG23	2.18	0.44
1:L:118:ALA:HB3	1:L:119:PRO:HD3	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:346:TRP:CE3	1:L:346:TRP:O	2.70	0.44
1:N:324:GLN:HB2	1:N:571:ILE:HB	1.99	0.44
1:N:438:GLY:HA3	1:N:465:GLY:HA3	1.98	0.44
1:A:151:ILE:HA	1:A:154:LEU:HB2	2.00	0.44
1:B:109:LEU:HD11	1:B:112:VAL:HB	1.99	0.44
1:C:147:ARG:O	1:C:151:ILE:HG12	2.16	0.44
1:C:251:LYS:HB3	1:D:476:GLU:HB2	1.98	0.44
1:F:147:ARG:O	1:F:151:ILE:HG12	2.16	0.44
1:F:225:ARG:O	1:F:229:THR:HG23	2.18	0.44
1:G:529:PHE:HZ	1:G:531:LYS:HE3	1.82	0.44
1:I:109:LEU:HD21	1:I:112:VAL:HB	1.99	0.44
1:K:335:GLN:HB2	1:K:417:ASN:HB3	2.00	0.44
1:M:170:GLN:HG3	1:M:218:ILE:HG23	1.98	0.44
1:M:324:GLN:HB2	1:M:571:ILE:HB	1.99	0.44
1:N:151:ILE:HA	1:N:154:LEU:HB2	2.00	0.44
1:O:147:ARG:O	1:O:151:ILE:HG12	2.16	0.44
1:A:346:TRP:CE3	1:A:346:TRP:O	2.70	0.44
1:A:414:ASN:HB3	1:B:534:VAL:CG2	2.47	0.44
1:D:221:PRO:O	1:D:225:ARG:HG3	2.18	0.44
1:D:225:ARG:O	1:D:229:THR:HG23	2.18	0.44
1:F:118:ALA:HB3	1:F:119:PRO:HD3	1.98	0.44
1:F:151:ILE:HA	1:F:154:LEU:HB2	2.00	0.44
1:F:324:GLN:HB2	1:F:571:ILE:HB	1.99	0.44
1:F:414:ASN:HB3	1:G:534:VAL:CG2	2.47	0.44
1:H:151:ILE:HA	1:H:154:LEU:HB2	2.00	0.44
1:H:225:ARG:O	1:H:229:THR:HG23	2.18	0.44
1:I:222:GLU:O	1:I:225:ARG:HB2	2.17	0.44
1:L:221:PRO:O	1:L:225:ARG:HG3	2.18	0.44
1:M:109:LEU:HD21	1:M:112:VAL:HB	1.99	0.44
1:M:222:GLU:O	1:M:225:ARG:HB2	2.17	0.44
1:N:259:VAL:HG22	1:N:299:LEU:HD11	2.00	0.44
1:O:225:ARG:O	1:O:229:THR:HG23	2.18	0.44
1:B:259:VAL:HG22	1:B:299:LEU:HD11	2.00	0.44
1:B:414:ASN:HB3	1:C:534:VAL:CG2	2.47	0.44
1:F:109:LEU:HD11	1:F:112:VAL:HB	1.99	0.44
1:F:347:ALA:HA	1:F:353:ALA:HA	1.99	0.44
1:H:210:ASP:OD1	1:H:212:ARG:N	2.51	0.44
1:I:324:GLN:HB2	1:I:571:ILE:HB	1.99	0.44
1:J:221:PRO:O	1:J:225:ARG:HG3	2.18	0.44
1:M:259:VAL:HG22	1:M:299:LEU:HD11	2.00	0.44
1:M:335:GLN:HB2	1:M:417:ASN:HB3	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:324:GLN:HB2	1:O:571:ILE:HB	1.99	0.44
1:A:225:ARG:O	1:A:229:THR:HG23	2.18	0.44
1:A:361:LEU:HD12	1:A:361:LEU:O	2.18	0.44
1:C:259:VAL:HG22	1:C:299:LEU:HD11	2.00	0.44
1:C:529:PHE:HZ	1:C:531:LYS:HE3	1.82	0.44
1:D:347:ALA:HA	1:D:353:ALA:HA	1.99	0.44
1:E:109:LEU:HD21	1:E:112:VAL:HB	1.99	0.44
1:E:118:ALA:HB3	1:E:119:PRO:HD3	1.98	0.44
1:E:206:LYS:O	1:E:218:ILE:HD12	2.18	0.44
1:G:414:ASN:HB3	1:H:534:VAL:CG2	2.47	0.44
1:I:361:LEU:HD12	1:I:361:LEU:O	2.18	0.44
1:K:210:ASP:OD1	1:K:212:ARG:N	2.51	0.44
1:K:259:VAL:HG22	1:K:299:LEU:HD11	2.00	0.44
1:K:346:TRP:O	1:K:346:TRP:CE3	2.70	0.44
1:L:225:ARG:O	1:L:229:THR:HG23	2.18	0.44
1:L:324:GLN:HB2	1:L:571:ILE:HB	1.99	0.44
1:L:335:GLN:HB2	1:L:417:ASN:HB3	2.00	0.44
1:M:118:ALA:HB3	1:M:119:PRO:HD3	1.98	0.44
1:N:221:PRO:O	1:N:225:ARG:HG3	2.18	0.44
1:O:259:VAL:HG22	1:O:299:LEU:HD11	2.00	0.44
1:A:210:ASP:OD1	1:A:212:ARG:N	2.51	0.43
1:B:109:LEU:HD21	1:B:112:VAL:HB	1.99	0.43
1:C:210:ASP:OD1	1:C:212:ARG:N	2.51	0.43
1:E:324:GLN:HB2	1:E:571:ILE:HB	1.99	0.43
1:E:347:ALA:HA	1:E:353:ALA:HA	1.99	0.43
1:F:221:PRO:O	1:F:225:ARG:HG3	2.18	0.43
1:H:414:ASN:HB3	1:I:534:VAL:CG2	2.47	0.43
1:H:475:ASN:HA	1:H:476:GLU:HA	1.78	0.43
1:I:335:GLN:HB2	1:I:417:ASN:HB3	2.00	0.43
1:J:259:VAL:HG22	1:J:299:LEU:HD11	2.00	0.43
1:J:335:GLN:HB2	1:J:417:ASN:HB3	2.00	0.43
1:K:361:LEU:HD12	1:K:361:LEU:O	2.18	0.43
1:L:161:VAL:HA	1:L:164:ILE:CG1	2.46	0.43
1:L:259:VAL:HG22	1:L:299:LEU:HD11	2.00	0.43
1:N:335:GLN:HB2	1:N:417:ASN:HB3	2.00	0.43
1:O:221:PRO:HD2	1:O:224:ALA:HB2	2.00	0.43
1:A:221:PRO:HD2	1:A:224:ALA:HB2	2.00	0.43
1:B:221:PRO:HD2	1:B:224:ALA:HB2	2.00	0.43
1:B:221:PRO:O	1:B:225:ARG:HG3	2.18	0.43
1:C:109:LEU:HD11	1:C:112:VAL:HB	1.99	0.43
1:C:225:ARG:O	1:C:229:THR:HG23	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:361:LEU:HD12	1:C:361:LEU:O	2.18	0.43
1:D:206:LYS:O	1:D:218:ILE:HD12	2.18	0.43
1:D:346:TRP:CE3	1:D:346:TRP:O	2.70	0.43
1:G:225:ARG:O	1:G:229:THR:HG23	2.18	0.43
1:I:221:PRO:O	1:I:225:ARG:HG3	2.18	0.43
1:J:324:GLN:HB2	1:J:571:ILE:HB	1.99	0.43
1:L:529:PHE:HZ	1:L:531:LYS:HE3	1.82	0.43
1:N:225:ARG:O	1:N:229:THR:HG23	2.18	0.43
1:O:221:PRO:O	1:O:225:ARG:HG3	2.18	0.43
1:A:259:VAL:HG22	1:A:299:LEU:HD11	2.00	0.43
1:C:221:PRO:HD2	1:C:224:ALA:HB2	2.00	0.43
1:D:106:ILE:HG13	1:D:106:ILE:O	2.19	0.43
1:G:361:LEU:HD12	1:G:361:LEU:O	2.18	0.43
1:H:335:GLN:HB2	1:H:417:ASN:HB3	2.00	0.43
1:I:206:LYS:O	1:I:218:ILE:HD12	2.18	0.43
1:I:210:ASP:OD1	1:I:212:ARG:N	2.51	0.43
1:L:151:ILE:HA	1:L:154:LEU:HB2	2.00	0.43
1:L:210:ASP:OD1	1:L:212:ARG:N	2.51	0.43
1:M:151:ILE:HA	1:M:154:LEU:HB2	2.00	0.43
1:N:106:ILE:O	1:N:106:ILE:HG13	2.19	0.43
1:N:361:LEU:HD12	1:N:361:LEU:O	2.18	0.43
1:O:106:ILE:HG13	1:O:106:ILE:O	2.19	0.43
1:O:529:PHE:HZ	1:O:531:LYS:HE3	1.82	0.43
1:A:106:ILE:O	1:A:106:ILE:HG13	2.19	0.43
1:D:109:LEU:HD11	1:D:112:VAL:HB	1.99	0.43
1:D:414:ASN:HB3	1:E:534:VAL:CG2	2.47	0.43
1:E:109:LEU:HD11	1:E:112:VAL:HB	1.99	0.43
1:F:161:VAL:HA	1:F:164:ILE:CG1	2.47	0.43
1:F:206:LYS:O	1:F:218:ILE:HD12	2.18	0.43
1:G:221:PRO:O	1:G:225:ARG:HG3	2.18	0.43
1:I:414:ASN:HB3	1:J:534:VAL:CG2	2.47	0.43
1:J:151:ILE:HA	1:J:154:LEU:HB2	2.00	0.43
1:K:151:ILE:HA	1:K:154:LEU:HB2	2.00	0.43
1:M:106:ILE:HG13	1:M:106:ILE:O	2.19	0.43
1:N:221:PRO:HD2	1:N:224:ALA:HB2	2.00	0.43
1:B:106:ILE:HG13	1:B:106:ILE:O	2.19	0.43
1:C:161:VAL:HA	1:C:164:ILE:CG1	2.47	0.43
1:C:324:GLN:HB2	1:C:571:ILE:HB	1.99	0.43
1:D:335:GLN:HB2	1:D:417:ASN:HB3	2.00	0.43
1:D:361:LEU:HD12	1:D:361:LEU:O	2.18	0.43
1:E:335:GLN:HB2	1:E:417:ASN:HB3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:361:LEU:HD12	1:E:361:LEU:O	2.18	0.43
1:G:210:ASP:OD1	1:G:212:ARG:N	2.51	0.43
1:H:206:LYS:O	1:H:218:ILE:HD12	2.18	0.43
1:H:221:PRO:O	1:H:225:ARG:HG3	2.18	0.43
1:H:259:VAL:HG22	1:H:299:LEU:HD11	2.00	0.43
1:K:324:GLN:HB2	1:K:571:ILE:HB	1.99	0.43
1:M:221:PRO:HD2	1:M:224:ALA:HB2	2.00	0.43
1:A:221:PRO:O	1:A:225:ARG:HG3	2.18	0.43
1:A:324:GLN:HB2	1:A:571:ILE:HB	1.99	0.43
1:C:109:LEU:HD21	1:C:112:VAL:HB	1.99	0.43
1:E:259:VAL:HG22	1:E:299:LEU:HD11	2.00	0.43
1:G:221:PRO:HD2	1:G:224:ALA:HB2	2.00	0.43
1:I:151:ILE:HA	1:I:154:LEU:HB2	2.00	0.43
1:I:259:VAL:HG22	1:I:299:LEU:HD11	2.00	0.43
1:L:106:ILE:HG13	1:L:106:ILE:O	2.19	0.43
1:N:262:LEU:HD23	1:N:262:LEU:HA	1.83	0.43
1:O:124:MET:HB3	1:O:154:LEU:HD21	2.01	0.43
1:B:124:MET:HB3	1:B:154:LEU:HD21	2.01	0.43
1:C:106:ILE:O	1:C:106:ILE:HG13	2.19	0.43
1:C:124:MET:HB3	1:C:154:LEU:HD21	2.01	0.43
1:D:109:LEU:HD21	1:D:112:VAL:HB	1.99	0.43
1:D:124:MET:HB3	1:D:154:LEU:HD21	2.01	0.43
1:D:259:VAL:HG22	1:D:299:LEU:HD11	2.00	0.43
1:D:324:GLN:HB2	1:D:571:ILE:HB	1.99	0.43
1:F:361:LEU:HD12	1:F:361:LEU:O	2.18	0.43
1:H:117:LEU:HD13	1:H:157:VAL:HG12	2.01	0.43
1:H:221:PRO:HD2	1:H:224:ALA:HB2	2.00	0.43
1:I:117:LEU:HD13	1:I:157:VAL:HG12	2.01	0.43
1:I:232:LEU:HD23	1:I:232:LEU:HA	1.90	0.43
1:J:206:LYS:O	1:J:218:ILE:HD12	2.18	0.43
1:J:210:ASP:OD1	1:J:212:ARG:N	2.51	0.43
1:J:221:PRO:HD2	1:J:224:ALA:HB2	2.00	0.43
1:K:106:ILE:O	1:K:106:ILE:HG13	2.19	0.43
1:M:221:PRO:O	1:M:225:ARG:HG3	2.18	0.43
1:M:347:ALA:HB2	1:M:363:ILE:HG12	2.01	0.43
1:M:361:LEU:HD12	1:M:361:LEU:O	2.18	0.43
1:O:335:GLN:HB2	1:O:417:ASN:HB3	2.00	0.43
1:C:206:LYS:HZ3	1:D:190:ILE:C	2.27	0.43
1:C:221:PRO:O	1:C:225:ARG:HG3	2.18	0.43
1:E:210:ASP:OD1	1:E:212:ARG:N	2.51	0.43
1:F:259:VAL:HG22	1:F:299:LEU:HD11	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:117:LEU:HD13	1:G:157:VAL:HG12	2.01	0.43
1:I:221:PRO:HD2	1:I:224:ALA:HB2	2.00	0.43
1:J:117:LEU:HD13	1:J:157:VAL:HG12	2.01	0.43
1:L:206:LYS:HZ3	1:M:190:ILE:C	2.26	0.43
1:N:124:MET:HB3	1:N:154:LEU:HD21	2.01	0.43
1:N:210:ASP:OD1	1:N:212:ARG:N	2.51	0.43
1:O:206:LYS:O	1:O:218:ILE:HD12	2.18	0.43
1:A:124:MET:HB3	1:A:154:LEU:HD21	2.01	0.43
1:A:206:LYS:O	1:A:218:ILE:HD12	2.18	0.43
1:A:335:GLN:HB2	1:A:417:ASN:HB3	2.00	0.43
1:A:475:ASN:HA	1:A:476:GLU:HA	1.78	0.43
1:B:225:ARG:O	1:B:229:THR:HG23	2.18	0.43
1:B:324:GLN:HB2	1:B:571:ILE:HB	1.99	0.43
1:C:206:LYS:O	1:C:218:ILE:HD12	2.18	0.43
1:E:221:PRO:O	1:E:225:ARG:HG3	2.18	0.43
1:G:259:VAL:HG22	1:G:299:LEU:HD11	2.00	0.43
1:I:171:ILE:HD11	1:J:234:SER:HB3	2.01	0.43
1:K:221:PRO:O	1:K:225:ARG:HG3	2.18	0.43
1:L:221:PRO:HD2	1:L:224:ALA:HB2	2.00	0.43
1:L:347:ALA:HB2	1:L:363:ILE:HG12	2.01	0.43
1:L:361:LEU:HD12	1:L:361:LEU:O	2.18	0.43
1:M:225:ARG:O	1:M:229:THR:HG23	2.18	0.43
1:A:171:ILE:HD11	1:B:234:SER:HB3	2.01	0.43
1:B:361:LEU:HD12	1:B:361:LEU:O	2.18	0.43
1:B:475:ASN:HA	1:B:476:GLU:HA	1.78	0.43
1:C:335:GLN:HB2	1:C:417:ASN:HB3	2.00	0.43
1:D:221:PRO:HD2	1:D:224:ALA:HB2	2.00	0.43
1:F:335:GLN:HB2	1:F:417:ASN:HB3	2.00	0.43
1:J:171:ILE:HD11	1:K:234:SER:HB3	2.01	0.43
1:N:206:LYS:O	1:N:218:ILE:HD12	2.18	0.43
1:A:234:SER:HB3	1:O:171:ILE:HD11	2.01	0.42
1:E:221:PRO:HD2	1:E:224:ALA:HB2	2.00	0.42
1:N:347:ALA:HB2	1:N:363:ILE:HG12	2.01	0.42
1:O:109:LEU:CD1	1:O:162:ASP:HA	2.50	0.42
1:B:171:ILE:HD11	1:C:234:SER:HB3	2.01	0.42
1:D:123:GLN:HA	1:D:126:ASP:HB3	2.02	0.42
1:E:124:MET:HB3	1:E:154:LEU:HD21	2.01	0.42
1:F:124:MET:HB3	1:F:154:LEU:HD21	2.01	0.42
1:G:335:GLN:HB2	1:G:417:ASN:HB3	2.00	0.42
1:H:171:ILE:HD11	1:I:234:SER:HB3	2.01	0.42
1:L:109:LEU:CD1	1:L:162:ASP:HA	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:124:MET:HB3	1:L:154:LEU:HD21	2.01	0.42
1:O:475:ASN:HA	1:O:476:GLU:HA	1.78	0.42
1:B:117:LEU:HD13	1:B:157:VAL:HG12	2.01	0.42
1:B:123:GLN:HA	1:B:126:ASP:HB3	2.02	0.42
1:B:206:LYS:O	1:B:218:ILE:HD12	2.18	0.42
1:D:206:LYS:HZ3	1:E:190:ILE:C	2.27	0.42
1:D:344:VAL:O	1:D:354:GLN:NE2	2.53	0.42
1:E:106:ILE:HG13	1:E:106:ILE:O	2.19	0.42
1:F:109:LEU:CD1	1:F:162:ASP:HA	2.50	0.42
1:F:117:LEU:HD13	1:F:157:VAL:HG12	2.01	0.42
1:F:210:ASP:OD1	1:F:212:ARG:N	2.51	0.42
1:G:106:ILE:HG13	1:G:106:ILE:O	2.19	0.42
1:K:347:ALA:HB2	1:K:363:ILE:HG12	2.01	0.42
1:M:124:MET:HB3	1:M:154:LEU:HD21	2.01	0.42
1:M:206:LYS:O	1:M:218:ILE:HD12	2.18	0.42
1:O:361:LEU:HD12	1:O:361:LEU:O	2.18	0.42
1:A:344:VAL:O	1:A:354:GLN:NE2	2.53	0.42
1:B:344:VAL:O	1:B:354:GLN:NE2	2.53	0.42
1:D:210:ASP:OD1	1:D:212:ARG:N	2.51	0.42
1:F:344:VAL:O	1:F:354:GLN:NE2	2.53	0.42
1:H:106:ILE:O	1:H:106:ILE:HG13	2.19	0.42
1:H:344:VAL:O	1:H:354:GLN:NE2	2.53	0.42
1:J:361:LEU:HD12	1:J:361:LEU:O	2.18	0.42
1:K:117:LEU:HD13	1:K:157:VAL:HG12	2.01	0.42
1:K:171:ILE:HD11	1:L:234:SER:HB3	2.01	0.42
1:M:210:ASP:OD1	1:M:212:ARG:N	2.51	0.42
1:O:123:GLN:HA	1:O:126:ASP:HB3	2.02	0.42
1:O:210:ASP:OD1	1:O:212:ARG:N	2.51	0.42
1:A:117:LEU:HD13	1:A:157:VAL:HG12	2.01	0.42
1:B:335:GLN:HB2	1:B:417:ASN:HB3	2.00	0.42
1:C:117:LEU:HD13	1:C:157:VAL:HG12	2.01	0.42
1:C:475:ASN:HA	1:C:476:GLU:HA	1.78	0.42
1:E:109:LEU:CD1	1:E:162:ASP:HA	2.50	0.42
1:F:123:GLN:HA	1:F:126:ASP:HB3	2.02	0.42
1:F:262:LEU:HD23	1:F:262:LEU:HA	1.83	0.42
1:G:109:LEU:CD1	1:G:162:ASP:HA	2.50	0.42
1:J:106:ILE:HG13	1:J:106:ILE:O	2.19	0.42
1:B:210:ASP:OD1	1:B:212:ARG:N	2.51	0.42
1:D:109:LEU:CD1	1:D:162:ASP:HA	2.50	0.42
1:D:135:HIS:HE1	1:E:120:LEU:HB2	1.85	0.42
1:E:344:VAL:O	1:E:354:GLN:NE2	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:347:ALA:HB2	1:F:363:ILE:HG12	2.01	0.42
1:G:206:LYS:O	1:G:218:ILE:HD12	2.18	0.42
1:G:344:VAL:O	1:G:354:GLN:NE2	2.53	0.42
1:H:361:LEU:O	1:H:361:LEU:HD12	2.18	0.42
1:O:344:VAL:O	1:O:354:GLN:NE2	2.53	0.42
1:B:135:HIS:HE1	1:C:120:LEU:HB2	1.85	0.42
1:C:171:ILE:HD11	1:D:234:SER:HB3	2.02	0.42
1:D:347:ALA:HB2	1:D:363:ILE:HG12	2.01	0.42
1:G:124:MET:HB3	1:G:154:LEU:HD21	2.01	0.42
1:G:475:ASN:HA	1:G:476:GLU:HA	1.78	0.42
1:H:144:LEU:HD11	1:H:158:ILE:CD1	2.50	0.42
1:J:144:LEU:HD11	1:J:158:ILE:CD1	2.50	0.42
1:J:344:VAL:O	1:J:354:GLN:NE2	2.53	0.42
1:K:124:MET:HB3	1:K:154:LEU:HD21	2.01	0.42
1:K:221:PRO:HD2	1:K:224:ALA:HB2	2.00	0.42
1:N:171:ILE:HD11	1:O:234:SER:HB3	2.01	0.42
1:O:117:LEU:HD13	1:O:157:VAL:HG12	2.01	0.42
1:O:262:LEU:HD23	1:O:262:LEU:HA	1.83	0.42
1:O:571:ILE:CG2	1:O:577:TYR:HB2	2.50	0.42
1:A:109:LEU:CD1	1:A:162:ASP:HA	2.50	0.42
1:A:571:ILE:CG2	1:A:577:TYR:HB2	2.50	0.42
1:C:135:HIS:HE1	1:D:120:LEU:HB2	1.85	0.42
1:E:135:HIS:HE1	1:F:120:LEU:HB2	1.85	0.42
1:E:347:ALA:HB2	1:E:363:ILE:HG12	2.01	0.42
1:G:171:ILE:HD11	1:H:234:SER:HB3	2.01	0.42
1:H:109:LEU:CD1	1:H:162:ASP:HA	2.50	0.42
1:I:144:LEU:HD11	1:I:158:ILE:CD1	2.50	0.42
1:I:344:VAL:O	1:I:354:GLN:NE2	2.53	0.42
1:K:109:LEU:CD1	1:K:162:ASP:HA	2.50	0.42
1:K:206:LYS:O	1:K:218:ILE:HD12	2.18	0.42
1:L:144:LEU:HD11	1:L:158:ILE:CD1	2.50	0.42
1:A:120:LEU:HB2	1:O:135:HIS:HE1	1.85	0.42
1:C:109:LEU:CD1	1:C:162:ASP:HA	2.50	0.42
1:C:344:VAL:O	1:C:354:GLN:NE2	2.53	0.42
1:D:117:LEU:HD13	1:D:157:VAL:HG12	2.01	0.42
1:D:184:GLU:O	1:D:188:GLN:HG2	2.20	0.42
1:E:117:LEU:HD13	1:E:157:VAL:HG12	2.01	0.42
1:E:184:GLU:O	1:E:188:GLN:HG2	2.20	0.42
1:F:221:PRO:HD2	1:F:224:ALA:HB2	2.00	0.42
1:F:571:ILE:CG2	1:F:577:TYR:HB2	2.50	0.42
1:I:109:LEU:CD1	1:I:162:ASP:HA	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:184:GLU:O	1:J:188:GLN:HG2	2.20	0.42
1:L:135:HIS:HE1	1:M:120:LEU:HB2	1.85	0.42
1:L:344:VAL:O	1:L:354:GLN:NE2	2.53	0.42
1:M:109:LEU:CD1	1:M:162:ASP:HA	2.50	0.42
1:M:184:GLU:O	1:M:188:GLN:HG2	2.20	0.42
1:N:109:LEU:CD1	1:N:162:ASP:HA	2.50	0.42
1:N:184:GLU:O	1:N:188:GLN:HG2	2.20	0.42
1:N:571:ILE:CG2	1:N:577:TYR:HB2	2.50	0.42
1:O:347:ALA:HB2	1:O:363:ILE:HG12	2.01	0.42
1:A:436:ASN:HB2	1:B:507:THR:HB	2.02	0.42
1:C:184:GLU:O	1:C:188:GLN:HG2	2.20	0.42
1:C:347:ALA:HB2	1:C:363:ILE:HG12	2.01	0.42
1:C:571:ILE:CG2	1:C:577:TYR:HB2	2.50	0.42
1:D:171:ILE:HD11	1:E:234:SER:HB3	2.01	0.42
1:H:124:MET:HB3	1:H:154:LEU:HD21	2.01	0.42
1:I:106:ILE:O	1:I:106:ILE:HG13	2.19	0.42
1:I:124:MET:HB3	1:I:154:LEU:HD21	2.01	0.42
1:K:144:LEU:HD11	1:K:158:ILE:CD1	2.50	0.42
1:K:344:VAL:O	1:K:354:GLN:NE2	2.53	0.42
1:L:206:LYS:O	1:L:218:ILE:HD12	2.18	0.42
1:M:135:HIS:HE1	1:N:120:LEU:HB2	1.85	0.42
1:N:475:ASN:HA	1:N:476:GLU:HA	1.78	0.42
1:B:118:ALA:HB1	1:B:122:ARG:HH21	1.85	0.41
1:B:184:GLU:O	1:B:188:GLN:HG2	2.20	0.41
1:D:475:ASN:HA	1:D:476:GLU:HA	1.78	0.41
1:D:571:ILE:CG2	1:D:577:TYR:HB2	2.50	0.41
1:F:106:ILE:O	1:F:106:ILE:HG13	2.19	0.41
1:F:118:ALA:HB1	1:F:122:ARG:HH21	1.85	0.41
1:F:184:GLU:O	1:F:188:GLN:HG2	2.20	0.41
1:G:123:GLN:HA	1:G:126:ASP:HB3	2.02	0.41
1:G:144:LEU:HD11	1:G:158:ILE:CD1	2.50	0.41
1:H:232:LEU:HD23	1:H:232:LEU:HA	1.90	0.41
1:J:159:LYS:O	1:J:162:ASP:HB3	2.20	0.41
1:K:118:ALA:HB1	1:K:122:ARG:HH21	1.85	0.41
1:K:135:HIS:HE1	1:L:120:LEU:HB2	1.85	0.41
1:M:344:VAL:O	1:M:354:GLN:NE2	2.53	0.41
1:N:144:LEU:HD11	1:N:158:ILE:CD1	2.50	0.41
1:N:344:VAL:O	1:N:354:GLN:NE2	2.53	0.41
1:A:507:THR:HB	1:O:436:ASN:HB2	2.03	0.41
1:B:175:GLU:HG3	1:B:176:TYR:CD2	2.56	0.41
1:B:436:ASN:HB2	1:C:507:THR:HB	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:159:LYS:O	1:D:162:ASP:HB3	2.20	0.41
1:D:175:GLU:HG3	1:D:176:TYR:CD2	2.55	0.41
1:F:144:LEU:HD11	1:F:158:ILE:CD1	2.50	0.41
1:F:175:GLU:HG3	1:F:176:TYR:CD2	2.55	0.41
1:G:175:GLU:HG3	1:G:176:TYR:CD2	2.56	0.41
1:G:347:ALA:HB2	1:G:363:ILE:HG12	2.01	0.41
1:H:571:ILE:CG2	1:H:577:TYR:HB2	2.50	0.41
1:I:123:GLN:HA	1:I:126:ASP:HB3	2.02	0.41
1:I:184:GLU:O	1:I:188:GLN:HG2	2.20	0.41
1:J:118:ALA:HB1	1:J:122:ARG:HH21	1.85	0.41
1:K:571:ILE:CG2	1:K:577:TYR:HB2	2.50	0.41
1:L:118:ALA:HB1	1:L:122:ARG:HH21	1.85	0.41
1:L:123:GLN:HA	1:L:126:ASP:HB3	2.02	0.41
1:L:171:ILE:HD11	1:M:234:SER:HB3	2.01	0.41
1:L:571:ILE:CG2	1:L:577:TYR:HB2	2.50	0.41
1:M:144:LEU:HD11	1:M:158:ILE:CD1	2.50	0.41
1:M:175:GLU:HG3	1:M:176:TYR:CD2	2.55	0.41
1:M:571:ILE:CG2	1:M:577:TYR:HB2	2.50	0.41
1:N:117:LEU:HD13	1:N:157:VAL:HG12	2.01	0.41
1:N:175:GLU:HG3	1:N:176:TYR:CD2	2.56	0.41
1:A:118:ALA:HB1	1:A:122:ARG:HH21	1.85	0.41
1:A:184:GLU:O	1:A:188:GLN:HG2	2.20	0.41
1:B:571:ILE:CG2	1:B:577:TYR:HB2	2.50	0.41
1:E:118:ALA:HB1	1:E:122:ARG:HH21	1.85	0.41
1:F:135:HIS:HE1	1:G:120:LEU:HB2	1.85	0.41
1:F:475:ASN:HA	1:F:476:GLU:HA	1.78	0.41
1:I:159:LYS:O	1:I:162:ASP:HB3	2.20	0.41
1:J:109:LEU:CD1	1:J:162:ASP:HA	2.50	0.41
1:J:124:MET:HB3	1:J:154:LEU:HD21	2.01	0.41
1:J:212:ARG:CZ	1:K:181:ASP:OD2	2.69	0.41
1:J:347:ALA:HB2	1:J:363:ILE:HG12	2.01	0.41
1:L:117:LEU:HD13	1:L:157:VAL:HG12	2.01	0.41
1:N:123:GLN:HA	1:N:126:ASP:HB3	2.02	0.41
1:O:175:GLU:HG3	1:O:176:TYR:CD2	2.56	0.41
1:B:109:LEU:CD1	1:B:162:ASP:HA	2.50	0.41
1:E:159:LYS:O	1:E:162:ASP:HB3	2.20	0.41
1:E:571:ILE:CG2	1:E:577:TYR:HB2	2.50	0.41
1:G:184:GLU:O	1:G:188:GLN:HG2	2.20	0.41
1:H:347:ALA:HB2	1:H:363:ILE:HG12	2.01	0.41
1:H:436:ASN:HB2	1:I:507:THR:HB	2.02	0.41
1:N:135:HIS:HE1	1:O:120:LEU:HB2	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:212:ARG:CZ	1:O:181:ASP:OD2	2.69	0.41
1:O:159:LYS:O	1:O:162:ASP:HB3	2.20	0.41
1:B:159:LYS:O	1:B:162:ASP:HB3	2.20	0.41
1:C:436:ASN:HB2	1:D:507:THR:HB	2.02	0.41
1:E:123:GLN:HA	1:E:126:ASP:HB3	2.02	0.41
1:E:475:ASN:HA	1:E:476:GLU:HA	1.78	0.41
1:F:171:ILE:HD11	1:G:234:SER:HB3	2.01	0.41
1:G:571:ILE:CG2	1:G:577:TYR:HB2	2.50	0.41
1:I:347:ALA:HB2	1:I:363:ILE:HG12	2.01	0.41
1:O:184:GLU:O	1:O:188:GLN:HG2	2.20	0.41
1:A:123:GLN:HA	1:A:126:ASP:HB3	2.02	0.41
1:B:212:ARG:CZ	1:C:181:ASP:OD2	2.69	0.41
1:B:347:ALA:HB2	1:B:363:ILE:HG12	2.01	0.41
1:C:159:LYS:O	1:C:162:ASP:HB3	2.20	0.41
1:E:171:ILE:HD11	1:F:234:SER:HB3	2.01	0.41
1:G:149:SER:O	1:G:152:ASN:HB2	2.21	0.41
1:I:548:GLN:HA	1:I:551:ARG:HG2	2.03	0.41
1:J:571:ILE:CG2	1:J:577:TYR:HB2	2.50	0.41
1:K:159:LYS:O	1:K:162:ASP:HB3	2.20	0.41
1:L:149:SER:O	1:L:152:ASN:HB2	2.21	0.41
1:L:159:LYS:O	1:L:162:ASP:HB3	2.20	0.41
1:M:171:ILE:HD11	1:N:234:SER:HB3	2.01	0.41
1:N:159:LYS:O	1:N:162:ASP:HB3	2.20	0.41
1:N:436:ASN:HB2	1:O:507:THR:HB	2.02	0.41
1:C:149:SER:O	1:C:152:ASN:HB2	2.21	0.41
1:C:167:GLU:HA	1:C:221:PRO:CA	2.50	0.41
1:C:175:GLU:HG3	1:C:176:TYR:CD2	2.56	0.41
1:D:436:ASN:HB2	1:E:507:THR:HB	2.02	0.41
1:H:159:LYS:O	1:H:162:ASP:HB3	2.20	0.41
1:H:175:GLU:HG3	1:H:176:TYR:CD2	2.56	0.41
1:I:121:LEU:HD23	1:I:121:LEU:HA	1.86	0.41
1:I:436:ASN:HB2	1:J:507:THR:HB	2.03	0.41
1:J:135:HIS:HE1	1:K:120:LEU:HB2	1.85	0.41
1:K:184:GLU:O	1:K:188:GLN:HG2	2.20	0.41
1:M:118:ALA:HB1	1:M:122:ARG:HH21	1.85	0.41
1:M:212:ARG:CZ	1:N:181:ASP:OD2	2.69	0.41
1:A:108:PRO:HB3	1:A:141:VAL:HA	2.03	0.41
1:A:144:LEU:HD11	1:A:158:ILE:CD1	2.50	0.41
1:A:149:SER:O	1:A:152:ASN:HB2	2.21	0.41
1:A:183:ALA:O	1:A:184:GLU:C	2.64	0.41
1:B:183:ALA:O	1:B:184:GLU:C	2.64	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:262:LEU:HD23	1:B:262:LEU:HA	1.83	0.41
1:C:123:GLN:HA	1:C:126:ASP:HB3	2.02	0.41
1:D:167:GLU:HA	1:D:221:PRO:CA	2.51	0.41
1:G:436:ASN:HB2	1:H:507:THR:HB	2.02	0.41
1:H:548:GLN:HA	1:H:551:ARG:HG2	2.03	0.41
1:I:118:ALA:HB1	1:I:122:ARG:HH21	1.85	0.41
1:J:149:SER:O	1:J:152:ASN:HB2	2.21	0.41
1:J:548:GLN:HA	1:J:551:ARG:HG2	2.03	0.41
1:K:108:PRO:HB3	1:K:141:VAL:HA	2.03	0.41
1:L:184:GLU:O	1:L:188:GLN:HG2	2.20	0.41
1:M:117:LEU:HD13	1:M:157:VAL:HG12	2.01	0.41
1:N:149:SER:O	1:N:152:ASN:HB2	2.21	0.41
1:O:144:LEU:HD11	1:O:158:ILE:CD1	2.50	0.41
1:A:135:HIS:HE1	1:B:120:LEU:HB2	1.85	0.41
1:B:144:LEU:HD11	1:B:158:ILE:CD1	2.50	0.41
1:B:167:GLU:HA	1:B:221:PRO:CA	2.50	0.41
1:B:206:LYS:NZ	1:C:190:ILE:C	2.79	0.41
1:C:144:LEU:HD11	1:C:158:ILE:CD1	2.50	0.41
1:D:149:SER:O	1:D:152:ASN:HB2	2.21	0.41
1:D:258:LEU:HD21	1:D:318:LEU:CD2	2.43	0.41
1:D:536:LYS:HE3	1:D:542:ASP:OD1	2.21	0.41
1:E:167:GLU:HA	1:E:221:PRO:CA	2.51	0.41
1:E:175:GLU:HG3	1:E:176:TYR:CD2	2.55	0.41
1:E:548:GLN:HA	1:E:551:ARG:HG2	2.03	0.41
1:F:149:SER:O	1:F:152:ASN:HB2	2.21	0.41
1:F:159:LYS:O	1:F:162:ASP:HB3	2.20	0.41
1:I:212:ARG:CZ	1:J:181:ASP:OD2	2.69	0.41
1:J:108:PRO:HB3	1:J:141:VAL:HA	2.03	0.41
1:J:436:ASN:HB2	1:K:507:THR:HB	2.02	0.41
1:K:167:GLU:HA	1:K:221:PRO:CA	2.51	0.41
1:L:175:GLU:HG3	1:L:176:TYR:CD2	2.56	0.41
1:M:108:PRO:HB3	1:M:141:VAL:HA	2.03	0.41
1:M:288:VAL:HG22	1:M:303:ALA:HB2	2.03	0.41
1:N:121:LEU:HD21	1:N:157:VAL:HB	2.03	0.41
1:O:108:PRO:HB3	1:O:141:VAL:HA	2.03	0.41
1:O:183:ALA:O	1:O:184:GLU:C	2.64	0.41
1:A:175:GLU:HG3	1:A:176:TYR:CD2	2.56	0.41
1:A:206:LYS:NZ	1:B:190:ILE:C	2.79	0.41
1:A:347:ALA:HB2	1:A:363:ILE:HG12	2.01	0.41
1:C:258:LEU:HD21	1:C:318:LEU:CD2	2.43	0.41
1:D:144:LEU:HD11	1:D:158:ILE:CD1	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:548:GLN:HA	1:D:551:ARG:HG2	2.03	0.41
1:H:149:SER:O	1:H:152:ASN:HB2	2.21	0.41
1:I:571:ILE:CG2	1:I:577:TYR:HB2	2.50	0.41
1:J:206:LYS:NZ	1:K:190:ILE:C	2.79	0.41
1:K:121:LEU:HD21	1:K:157:VAL:HB	2.03	0.41
1:K:206:LYS:NZ	1:L:190:ILE:C	2.79	0.41
1:L:108:PRO:HB3	1:L:141:VAL:HA	2.03	0.41
1:L:121:LEU:HD21	1:L:157:VAL:HB	2.03	0.41
1:M:123:GLN:HA	1:M:126:ASP:HB3	2.02	0.41
1:M:536:LYS:HE3	1:M:542:ASP:OD1	2.21	0.41
1:O:232:LEU:HD23	1:O:232:LEU:HA	1.90	0.41
1:A:190:ILE:C	1:O:206:LYS:NZ	2.79	0.40
1:A:212:ARG:CZ	1:B:181:ASP:OD2	2.69	0.40
1:B:108:PRO:HB3	1:B:141:VAL:HA	2.03	0.40
1:C:206:LYS:NZ	1:D:190:ILE:C	2.79	0.40
1:E:536:LYS:HE3	1:E:542:ASP:OD1	2.21	0.40
1:F:436:ASN:HB2	1:G:507:THR:HB	2.02	0.40
1:H:123:GLN:HA	1:H:126:ASP:HB3	2.02	0.40
1:J:288:VAL:HG22	1:J:303:ALA:HB2	2.03	0.40
1:M:121:LEU:HD21	1:M:157:VAL:HB	2.03	0.40
1:M:232:LEU:HA	1:M:232:LEU:HD23	1.90	0.40
1:N:108:PRO:HB3	1:N:141:VAL:HA	2.03	0.40
1:N:536:LYS:HE3	1:N:542:ASP:OD1	2.21	0.40
1:O:118:ALA:HB1	1:O:122:ARG:HH21	1.85	0.40
1:O:536:LYS:HE3	1:O:542:ASP:OD1	2.21	0.40
1:B:149:SER:O	1:B:152:ASN:HB2	2.21	0.40
1:C:226:GLN:HG3	1:C:227:ARG:N	2.37	0.40
1:E:362:PRO:HD2	1:E:388:MET:SD	2.62	0.40
1:E:436:ASN:HB2	1:F:507:THR:HB	2.02	0.40
1:F:108:PRO:HB3	1:F:141:VAL:HA	2.03	0.40
1:G:108:PRO:HB3	1:G:141:VAL:HA	2.03	0.40
1:G:135:HIS:HE1	1:H:120:LEU:HB2	1.85	0.40
1:H:226:GLN:HG3	1:H:227:ARG:N	2.37	0.40
1:I:108:PRO:HB3	1:I:141:VAL:HA	2.03	0.40
1:I:175:GLU:HG3	1:I:176:TYR:CD2	2.56	0.40
1:J:123:GLN:HA	1:J:126:ASP:HB3	2.02	0.40
1:J:226:GLN:HG3	1:J:227:ARG:N	2.37	0.40
1:K:175:GLU:HG3	1:K:176:TYR:CD2	2.56	0.40
1:K:436:ASN:HB2	1:L:507:THR:HB	2.02	0.40
1:L:536:LYS:HE3	1:L:542:ASP:OD1	2.21	0.40
1:M:159:LYS:O	1:M:162:ASP:HB3	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:206:LYS:NZ	1:N:190:ILE:C	2.79	0.40
1:M:436:ASN:HB2	1:N:507:THR:HB	2.02	0.40
1:A:121:LEU:HD21	1:A:157:VAL:HB	2.03	0.40
1:A:167:GLU:HA	1:A:221:PRO:CA	2.51	0.40
1:A:226:GLN:HG3	1:A:227:ARG:N	2.37	0.40
1:B:226:GLN:HG3	1:B:227:ARG:N	2.37	0.40
1:C:362:PRO:HD2	1:C:388:MET:SD	2.62	0.40
1:C:536:LYS:HE3	1:C:542:ASP:OD1	2.21	0.40
1:C:548:GLN:HA	1:C:551:ARG:HG2	2.03	0.40
1:D:118:ALA:HB1	1:D:122:ARG:HH21	1.85	0.40
1:D:121:LEU:HD21	1:D:157:VAL:HB	2.03	0.40
1:E:108:PRO:HB3	1:E:141:VAL:HA	2.03	0.40
1:E:144:LEU:HD11	1:E:158:ILE:CD1	2.50	0.40
1:F:536:LYS:HE3	1:F:542:ASP:OD1	2.21	0.40
1:G:159:LYS:O	1:G:162:ASP:HB3	2.20	0.40
1:G:428:LEU:HD11	1:H:481:LEU:HD11	2.04	0.40
1:H:108:PRO:HB3	1:H:141:VAL:HA	2.03	0.40
1:H:135:HIS:HE1	1:I:120:LEU:HB2	1.85	0.40
1:H:184:GLU:O	1:H:188:GLN:HG2	2.20	0.40
1:H:206:LYS:NZ	1:I:190:ILE:C	2.79	0.40
1:H:428:LEU:HD11	1:I:481:LEU:HD11	2.04	0.40
1:I:428:LEU:HD11	1:J:481:LEU:HD11	2.04	0.40
1:J:175:GLU:HG3	1:J:176:TYR:CD2	2.55	0.40
1:J:183:ALA:O	1:J:184:GLU:C	2.64	0.40
1:J:262:LEU:HD23	1:J:262:LEU:HA	1.83	0.40
1:J:349:LYS:HE3	1:J:349:LYS:HB2	1.90	0.40
1:J:536:LYS:HE3	1:J:542:ASP:OD1	2.21	0.40
1:K:123:GLN:HA	1:K:126:ASP:HB3	2.02	0.40
1:K:548:GLN:HA	1:K:551:ARG:HG2	2.03	0.40
1:N:183:ALA:O	1:N:184:GLU:C	2.64	0.40
1:O:121:LEU:HD21	1:O:157:VAL:HB	2.03	0.40
1:O:149:SER:O	1:O:152:ASN:HB2	2.21	0.40
1:O:362:PRO:HD2	1:O:388:MET:SD	2.62	0.40
1:O:548:GLN:HA	1:O:551:ARG:HG2	2.03	0.40
1:B:258:LEU:HD21	1:B:318:LEU:CD2	2.43	0.40
1:D:226:GLN:HG3	1:D:227:ARG:N	2.37	0.40
1:G:226:GLN:HG3	1:G:227:ARG:N	2.37	0.40
1:G:232:LEU:HD23	1:G:232:LEU:HA	1.90	0.40
1:I:135:HIS:HE1	1:J:120:LEU:HB2	1.85	0.40
1:I:183:ALA:O	1:I:184:GLU:C	2.64	0.40
1:I:226:GLN:HG3	1:I:227:ARG:N	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:536:LYS:HE3	1:I:542:ASP:OD1	2.21	0.40
1:J:167:GLU:HA	1:J:221:PRO:CA	2.51	0.40
1:K:536:LYS:HE3	1:K:542:ASP:OD1	2.21	0.40
1:L:226:GLN:HG3	1:L:227:ARG:N	2.37	0.40
1:M:149:SER:O	1:M:152:ASN:HB2	2.21	0.40
1:M:329:ALA:O	1:M:422:THR:HA	2.22	0.40
1:M:475:ASN:HA	1:M:476:GLU:HA	1.78	0.40
1:A:362:PRO:HD2	1:A:388:MET:SD	2.62	0.40
1:A:428:LEU:HD11	1:B:481:LEU:HD11	2.04	0.40
1:C:108:PRO:HB3	1:C:141:VAL:HA	2.03	0.40
1:C:326:LEU:HD21	1:C:328:GLU:OE2	2.22	0.40
1:D:108:PRO:HB3	1:D:141:VAL:HA	2.03	0.40
1:D:206:LYS:NZ	1:E:190:ILE:C	2.79	0.40
1:E:149:SER:O	1:E:152:ASN:HB2	2.21	0.40
1:F:362:PRO:HD2	1:F:388:MET:SD	2.62	0.40
1:G:118:ALA:HB1	1:G:122:ARG:HH21	1.85	0.40
1:G:329:ALA:O	1:G:422:THR:HA	2.22	0.40
1:G:349:LYS:HE3	1:G:349:LYS:HB2	1.90	0.40
1:G:548:GLN:HA	1:G:551:ARG:HG2	2.03	0.40
1:H:167:GLU:HA	1:H:221:PRO:CA	2.50	0.40
1:I:207:ILE:HG23	1:I:216:LEU:HD11	2.04	0.40
1:J:207:ILE:HG23	1:J:216:LEU:HD11	2.04	0.40
1:J:428:LEU:HD11	1:K:481:LEU:HD11	2.04	0.40
1:L:212:ARG:CZ	1:M:181:ASP:OD2	2.69	0.40
1:N:136:TYR:HE1	1:N:142:LEU:HG	1.87	0.40
1:N:206:LYS:NZ	1:O:190:ILE:C	2.79	0.40
1:N:288:VAL:HG22	1:N:303:ALA:HB2	2.03	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	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	B	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	C	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	D	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	E	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	F	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	G	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	H	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	I	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	J	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	K	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	L	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	M	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	N	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
1	O	464/627 (74%)	442 (95%)	19 (4%)	3 (1%)	21	53
All	All	6960/9405 (74%)	6630 (95%)	285 (4%)	45 (1%)	23	53

All (45) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	480	VAL
1	B	480	VAL
1	C	480	VAL
1	D	480	VAL
1	E	480	VAL
1	F	480	VAL
1	G	480	VAL
1	H	480	VAL
1	I	480	VAL
1	J	480	VAL
1	K	480	VAL
1	L	480	VAL
1	M	480	VAL
1	N	480	VAL
1	O	480	VAL
1	A	242	GLU
1	B	242	GLU
1	C	242	GLU

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Mol	Chain	Res	Type
1	D	242	GLU
1	E	242	GLU
1	F	242	GLU
1	G	242	GLU
1	H	242	GLU
1	I	242	GLU
1	J	242	GLU
1	K	242	GLU
1	L	242	GLU
1	M	242	GLU
1	N	242	GLU
1	O	242	GLU
1	A	130	VAL
1	B	130	VAL
1	C	130	VAL
1	D	130	VAL
1	E	130	VAL
1	F	130	VAL
1	G	130	VAL
1	H	130	VAL
1	I	130	VAL
1	J	130	VAL
1	K	130	VAL
1	L	130	VAL
1	M	130	VAL
1	N	130	VAL
1	O	130	VAL

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	B	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	C	405/537 (75%)	403 (100%)	2 (0%)	81	86

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	D	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	E	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	F	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	G	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	H	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	I	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	J	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	K	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	L	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	M	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	N	405/537 (75%)	403 (100%)	2 (0%)	81	86
1	O	405/537 (75%)	403 (100%)	2 (0%)	81	86
All	All	6075/8055 (75%)	6045 (100%)	30 (0%)	78	86

All (30) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	258	LEU
1	A	349	LYS
1	B	258	LEU
1	B	349	LYS
1	C	258	LEU
1	C	349	LYS
1	D	258	LEU
1	D	349	LYS
1	E	258	LEU
1	E	349	LYS
1	F	258	LEU
1	F	349	LYS
1	G	258	LEU
1	G	349	LYS
1	H	258	LEU
1	H	349	LYS
1	I	258	LEU
1	I	349	LYS
1	J	258	LEU
1	J	349	LYS
1	K	258	LEU

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Mol	Chain	Res	Type
1	K	349	LYS
1	L	258	LEU
1	L	349	LYS
1	M	258	LEU
1	M	349	LYS
1	N	258	LEU
1	N	349	LYS
1	O	258	LEU
1	O	349	LYS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (184) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	135	HIS
1	A	187	ASN
1	A	214	ASN
1	A	257	ASN
1	A	295	GLN
1	A	305	GLN
1	A	308	GLN
1	A	324	GLN
1	A	354	GLN
1	A	368	GLN
1	A	473	GLN
1	A	475	ASN
1	B	135	HIS
1	B	187	ASN
1	B	214	ASN
1	B	257	ASN
1	B	295	GLN
1	B	297	ASN
1	B	305	GLN
1	B	308	GLN
1	B	324	GLN
1	B	358	ASN
1	B	368	GLN
1	B	473	GLN
1	B	475	ASN
1	C	135	HIS
1	C	187	ASN
1	C	214	ASN
1	C	257	ASN

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Mol	Chain	Res	Type
1	C	295	GLN
1	C	305	GLN
1	C	308	GLN
1	C	324	GLN
1	C	358	ASN
1	C	368	GLN
1	C	475	ASN
1	D	135	HIS
1	D	187	ASN
1	D	214	ASN
1	D	257	ASN
1	D	295	GLN
1	D	305	GLN
1	D	308	GLN
1	D	324	GLN
1	D	354	GLN
1	D	358	ASN
1	D	368	GLN
1	D	473	GLN
1	D	475	ASN
1	E	135	HIS
1	E	187	ASN
1	E	214	ASN
1	E	257	ASN
1	E	295	GLN
1	E	297	ASN
1	E	305	GLN
1	E	308	GLN
1	E	324	GLN
1	E	354	GLN
1	E	358	ASN
1	E	368	GLN
1	E	473	GLN
1	E	475	ASN
1	F	135	HIS
1	F	187	ASN
1	F	214	ASN
1	F	257	ASN
1	F	295	GLN
1	F	297	ASN
1	F	305	GLN
1	F	308	GLN

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Mol	Chain	Res	Type
1	F	324	GLN
1	F	354	GLN
1	F	358	ASN
1	F	368	GLN
1	F	475	ASN
1	G	135	HIS
1	G	187	ASN
1	G	214	ASN
1	G	257	ASN
1	G	295	GLN
1	G	297	ASN
1	G	305	GLN
1	G	308	GLN
1	G	324	GLN
1	G	358	ASN
1	G	368	GLN
1	G	475	ASN
1	H	135	HIS
1	H	187	ASN
1	H	214	ASN
1	H	257	ASN
1	H	295	GLN
1	H	305	GLN
1	H	308	GLN
1	H	324	GLN
1	H	358	ASN
1	H	368	GLN
1	H	475	ASN
1	I	135	HIS
1	I	187	ASN
1	I	214	ASN
1	I	257	ASN
1	I	295	GLN
1	I	305	GLN
1	I	308	GLN
1	I	324	GLN
1	I	354	GLN
1	I	358	ASN
1	I	368	GLN
1	I	475	ASN
1	J	135	HIS
1	J	187	ASN

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Mol	Chain	Res	Type
1	J	214	ASN
1	J	257	ASN
1	J	295	GLN
1	J	305	GLN
1	J	308	GLN
1	J	324	GLN
1	J	368	GLN
1	J	475	ASN
1	K	135	HIS
1	K	187	ASN
1	K	214	ASN
1	K	257	ASN
1	K	295	GLN
1	K	297	ASN
1	K	305	GLN
1	K	308	GLN
1	K	324	GLN
1	K	354	GLN
1	K	358	ASN
1	K	368	GLN
1	K	473	GLN
1	K	475	ASN
1	L	135	HIS
1	L	187	ASN
1	L	214	ASN
1	L	257	ASN
1	L	295	GLN
1	L	305	GLN
1	L	308	GLN
1	L	324	GLN
1	L	354	GLN
1	L	358	ASN
1	L	368	GLN
1	L	475	ASN
1	M	135	HIS
1	M	187	ASN
1	M	214	ASN
1	M	257	ASN
1	M	295	GLN
1	M	297	ASN
1	M	305	GLN
1	M	308	GLN

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Mol	Chain	Res	Type
1	M	324	GLN
1	M	354	GLN
1	M	358	ASN
1	M	368	GLN
1	M	473	GLN
1	N	135	HIS
1	N	187	ASN
1	N	214	ASN
1	N	257	ASN
1	N	295	GLN
1	N	297	ASN
1	N	305	GLN
1	N	308	GLN
1	N	324	GLN
1	N	358	ASN
1	N	368	GLN
1	N	475	ASN
1	O	135	HIS
1	O	187	ASN
1	O	214	ASN
1	O	257	ASN
1	O	295	GLN
1	O	305	GLN
1	O	308	GLN
1	O	324	GLN
1	O	354	GLN
1	O	358	ASN
1	O	368	GLN
1	O	475	ASN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates ⓘ

There are no oligosaccharides in this entry.

5.6 Ligand geometry

There are no ligands in this entry.

5.7 Other polymers

There are no such residues in this entry.

5.8 Polymer linkage issues

There are no chain breaks in this entry.

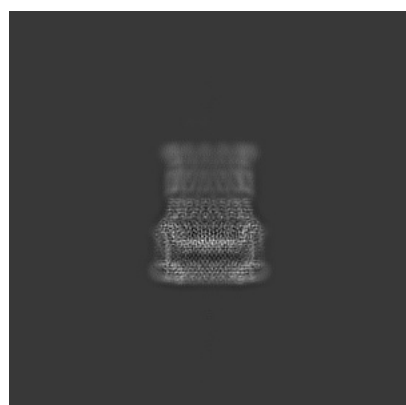
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-6675. These allow visual inspection of the internal detail of the map and identification of artifacts.

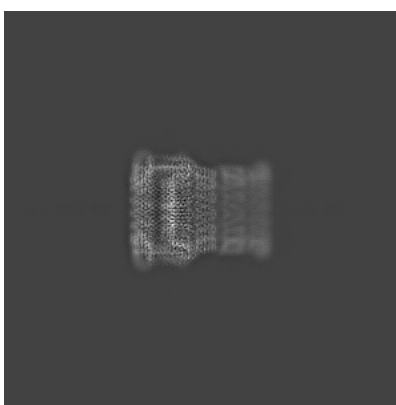
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

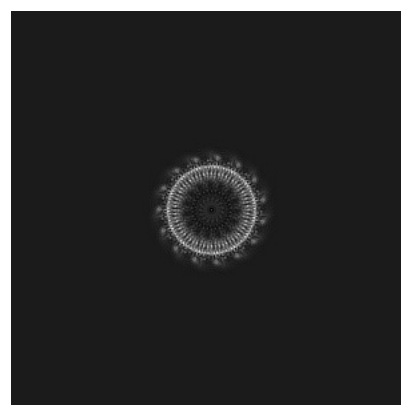
6.1.1 Primary map



X



Y



Z

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

6.2 Central slices [i](#)

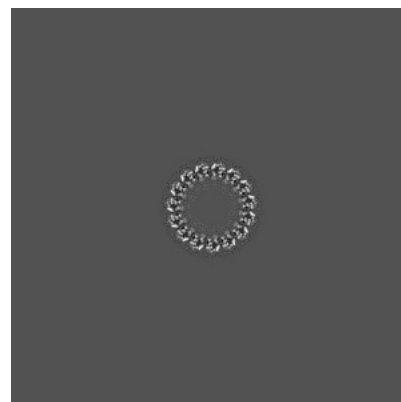
6.2.1 Primary map



X Index: 190



Y Index: 190

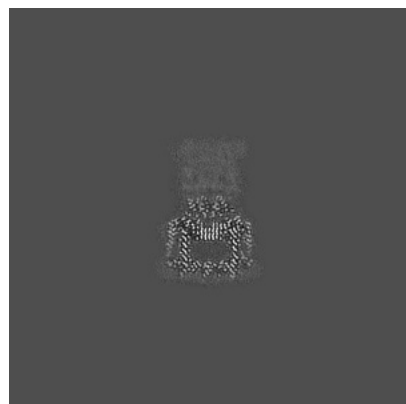


Z Index: 190

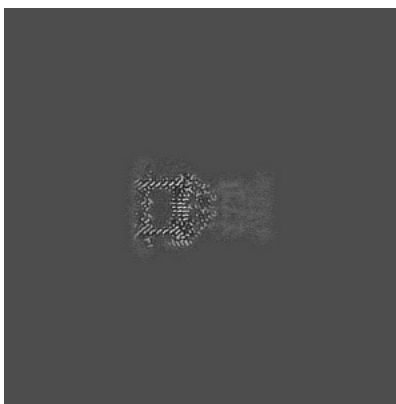
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

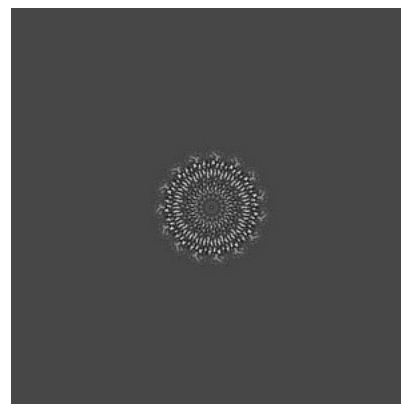
6.3.1 Primary map



X Index: 156



Y Index: 155

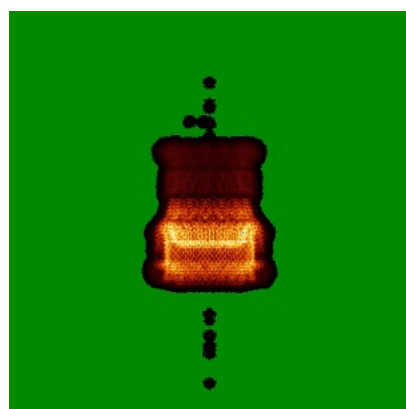


Z Index: 160

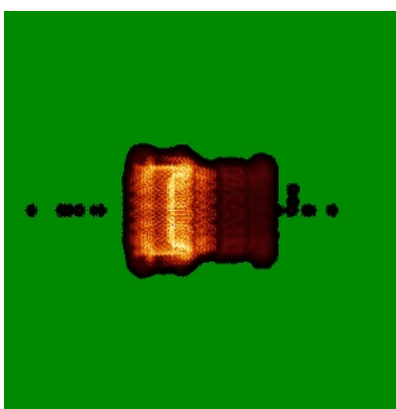
The images above show the largest variance slices of the map in three orthogonal directions.

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

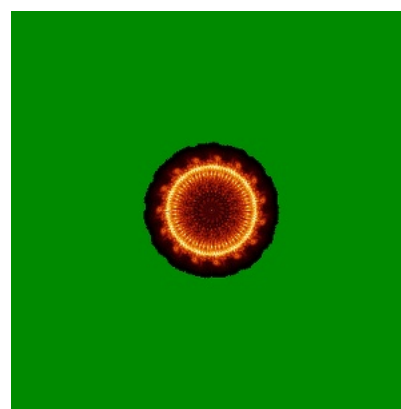
6.4.1 Primary map



X



Y

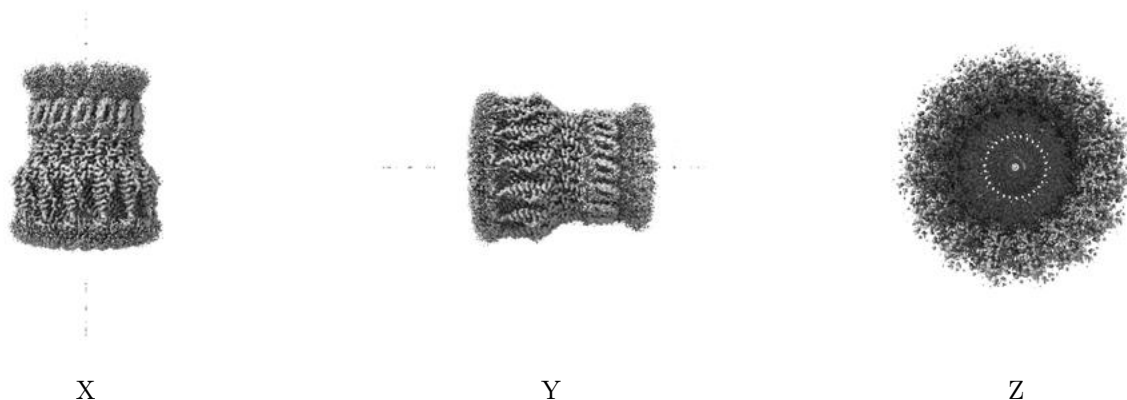


Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.022. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

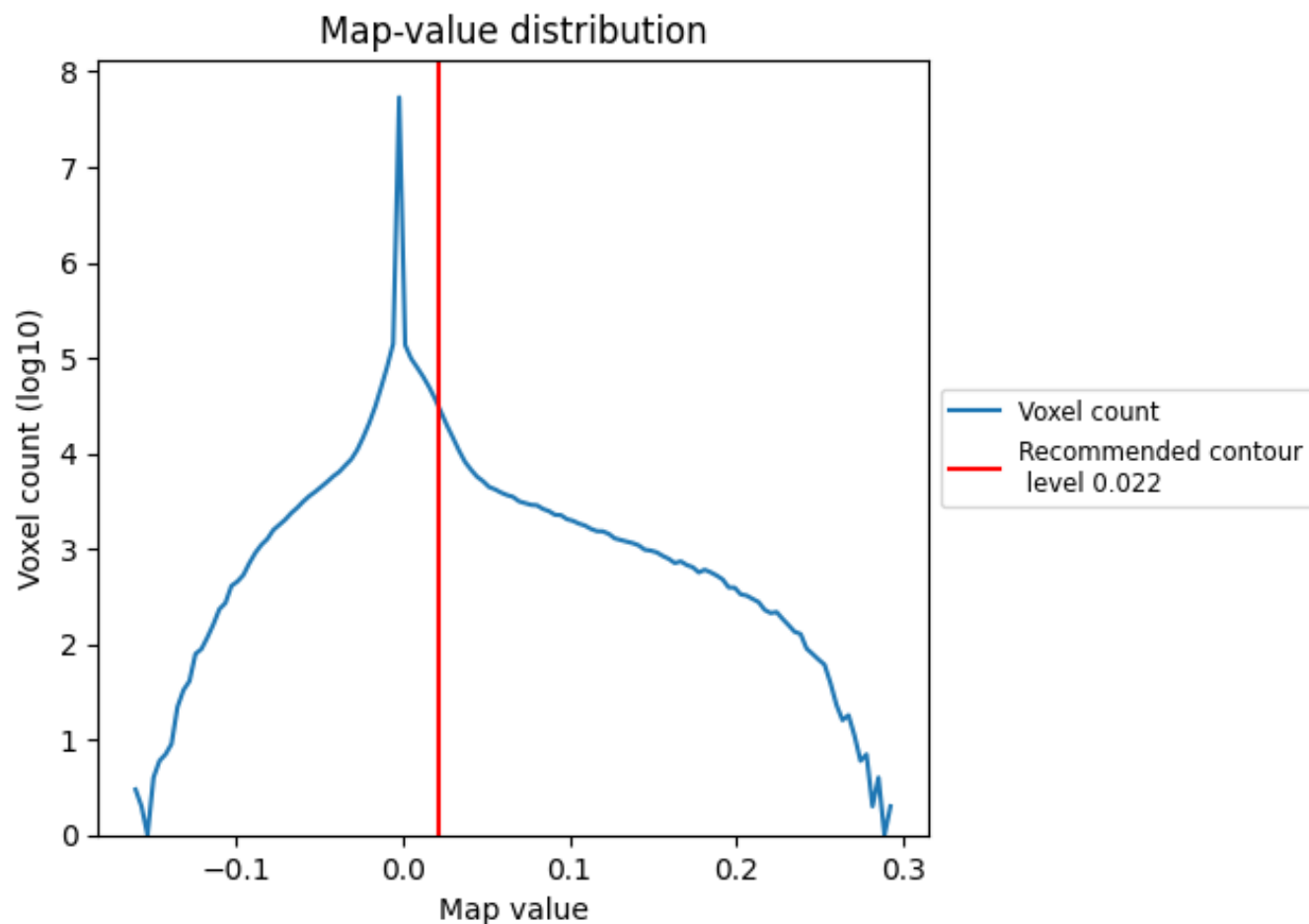
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

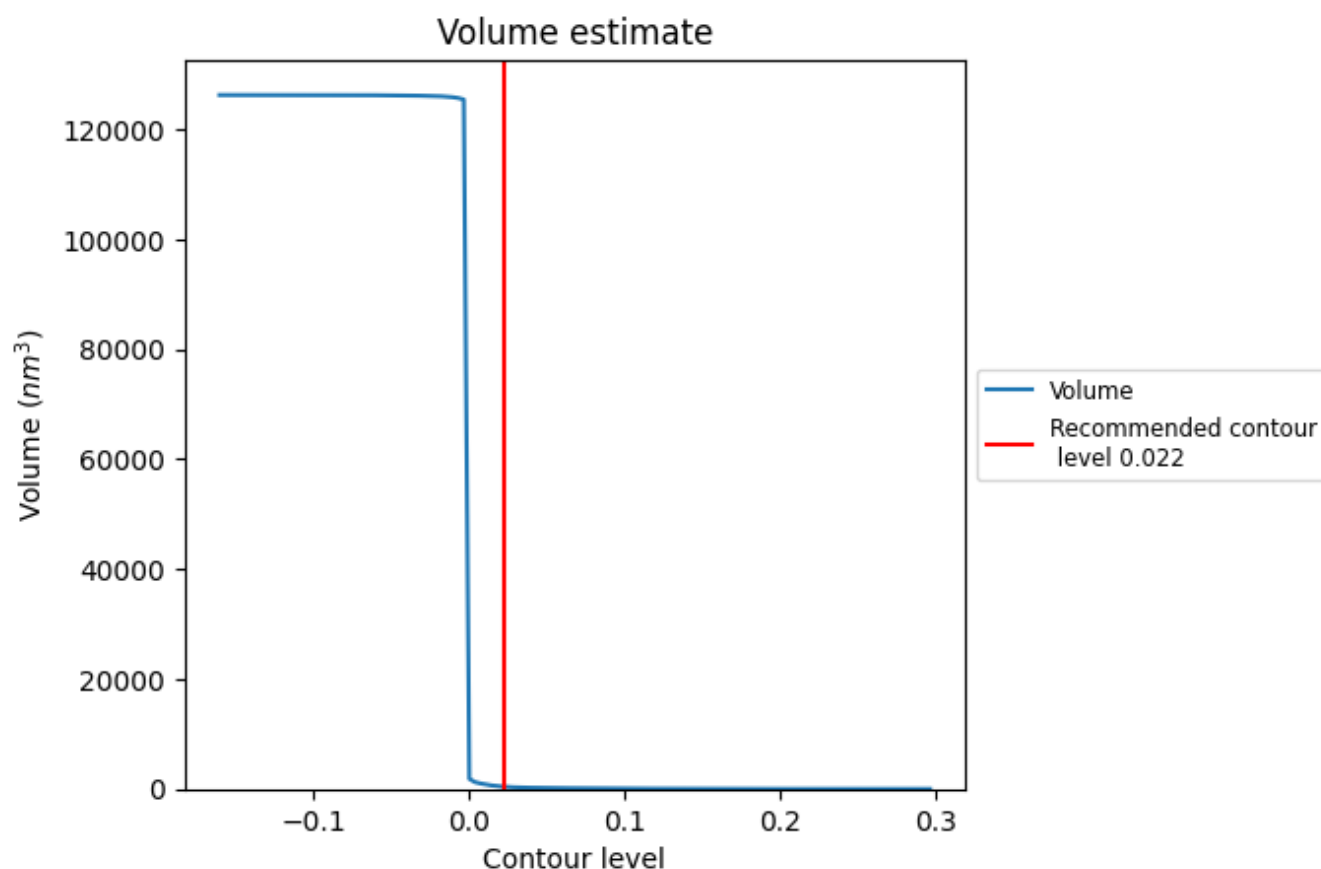
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

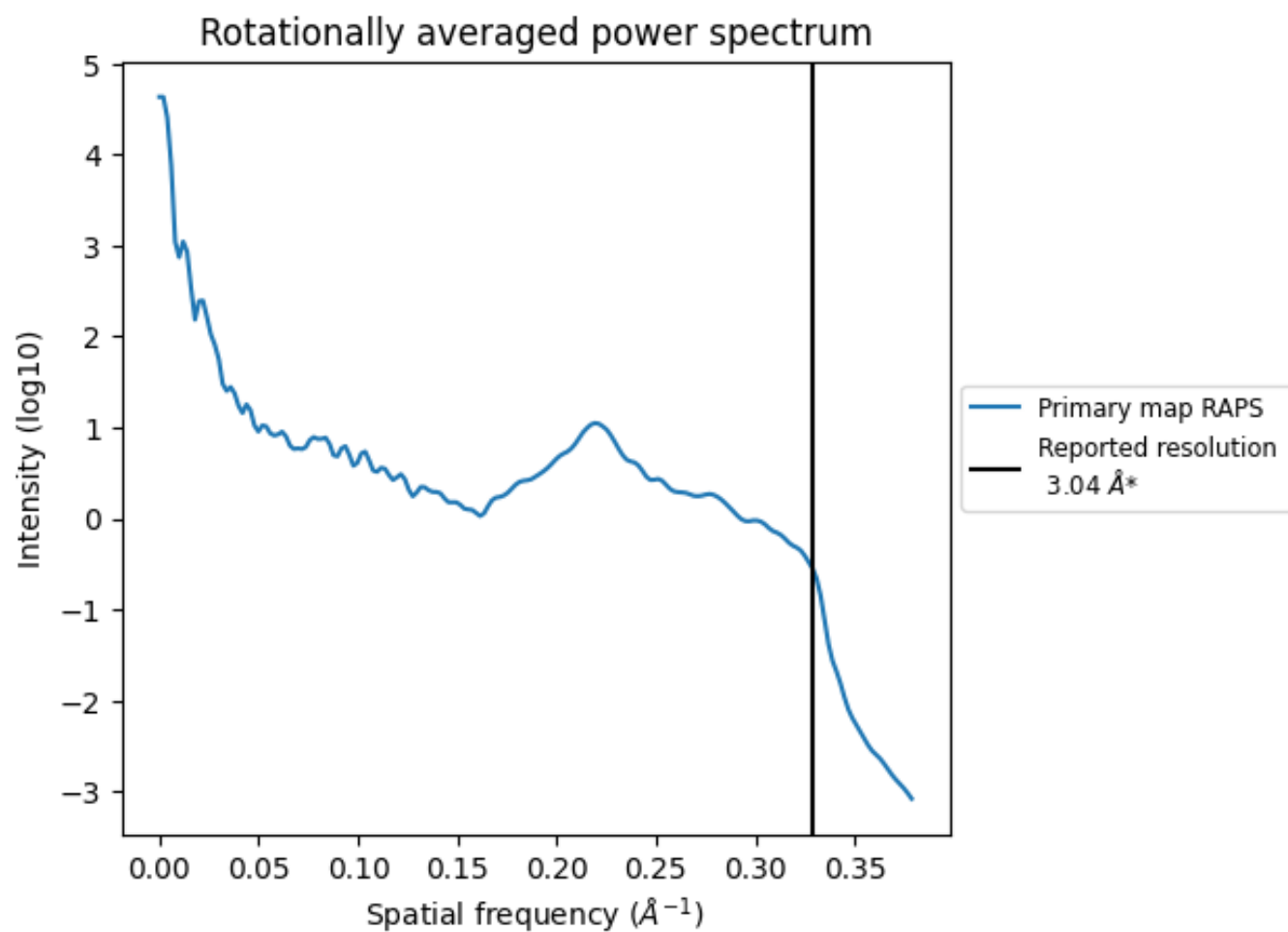
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 416 nm³; this corresponds to an approximate mass of 376 kDa.

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

7.3 Rotationally averaged power spectrum ⓘ



*Reported resolution corresponds to spatial frequency of 0.329 Å⁻¹

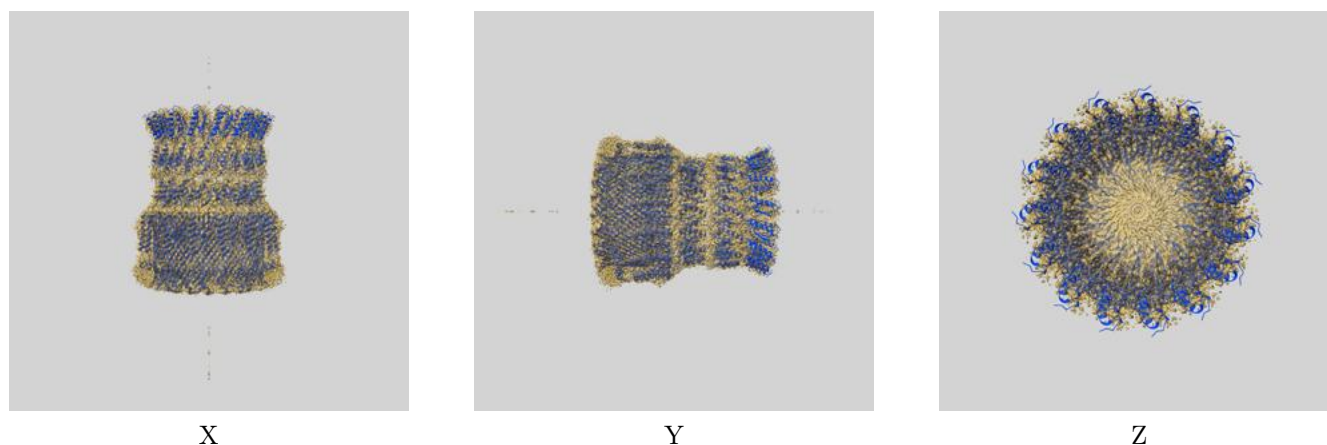
8 Fourier-Shell correlation

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

9 Map-model fit [i](#)

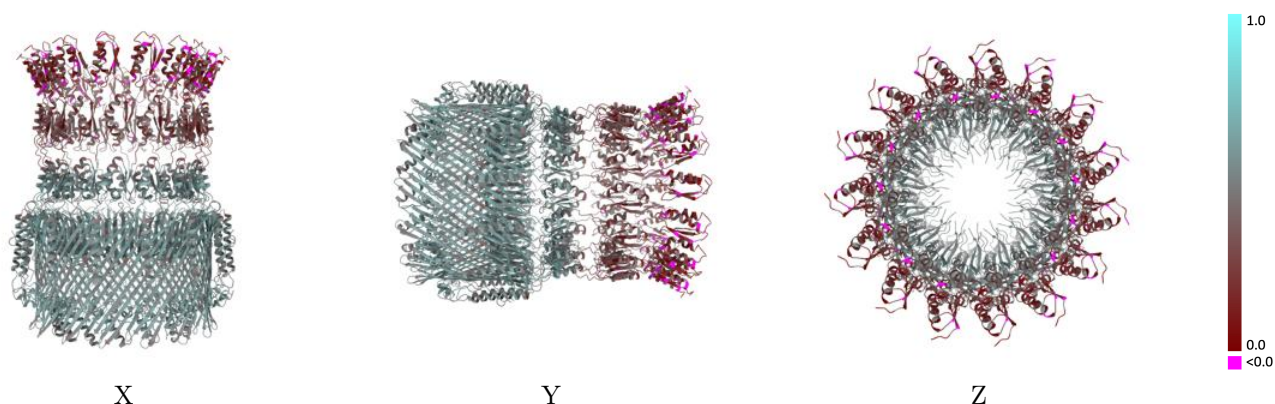
This section contains information regarding the fit between EMDB map EMD-6675 and PDB model 5WQ7. Per-residue inclusion information can be found in [section 3](#) on [page 5](#).

9.1 Map-model overlay [i](#)



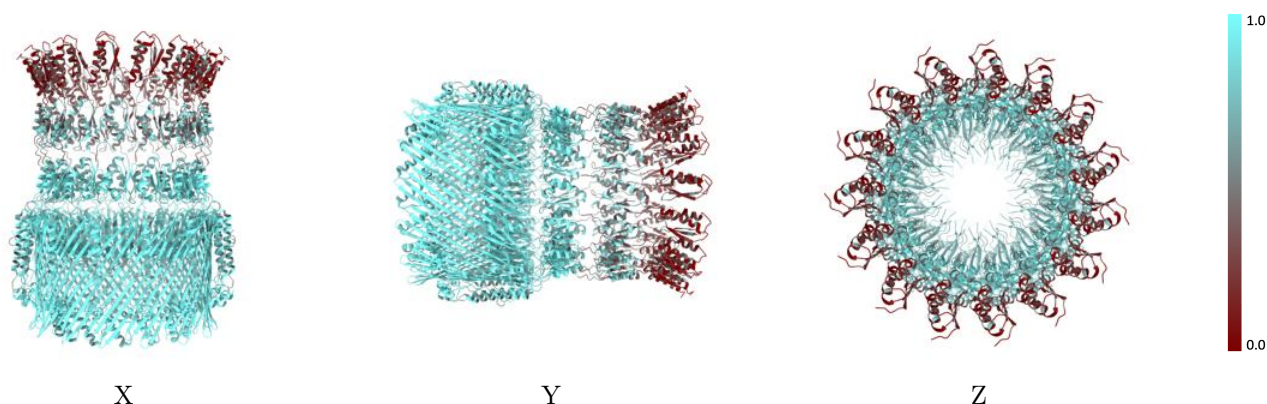
The images above show the 3D surface view of the map at the recommended contour level 0.022 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



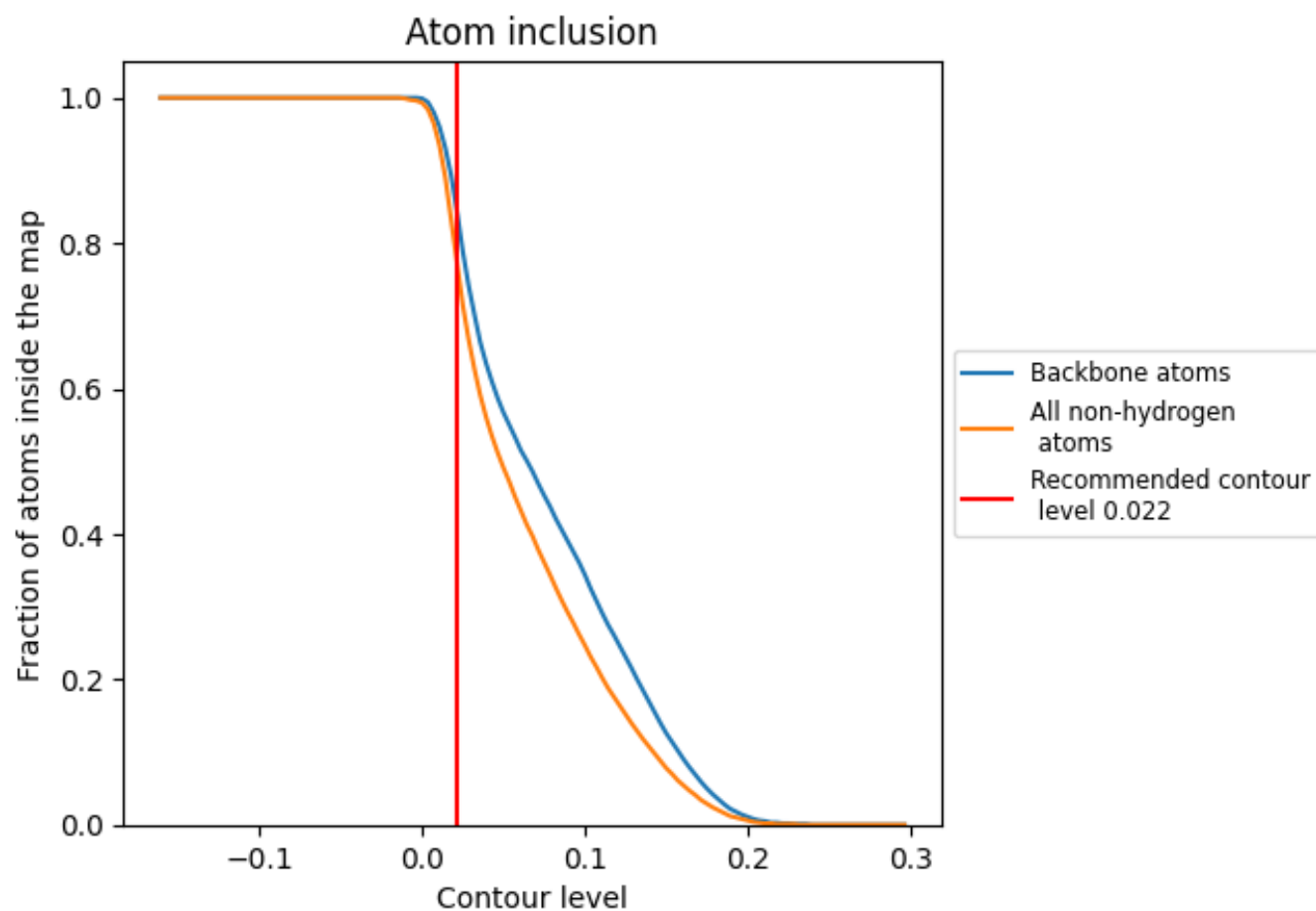
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions at which atoms can be resolved. Low Q-score values may therefore be expected for many entries.

9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.022).

9.4 Atom inclusion [i](#)



At the recommended contour level, 84% of all backbone atoms, 76% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.022) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.7640	0.4670
A	0.7640	0.4680
B	0.7630	0.4670
C	0.7650	0.4660
D	0.7650	0.4680
E	0.7620	0.4660
F	0.7650	0.4650
G	0.7640	0.4670
H	0.7650	0.4680
I	0.7650	0.4660
J	0.7620	0.4650
K	0.7660	0.4660
L	0.7640	0.4680
M	0.7630	0.4670
N	0.7590	0.4690
O	0.7630	0.4680

