



Full wwPDB EM Validation Report ⓘ

Mar 7, 2026 – 04:13 AM UTC

PDB ID : 3H3W / pdb_00003h3w
EMDB ID : EMD-1048
Title : Fitting of the gp6 crystal structure into 3D cryo-EM reconstruction of bacteriophage T4 dome-shaped baseplate
Authors : Aksyuk, A.A.; Leiman, P.G.; Shneider, M.M.; Mesyanzhinov, V.V.; Rossmann, M.G.
Deposited on : 2009-04-17
Resolution : 12.00 Å(reported)
Based on initial model : 3H2T

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

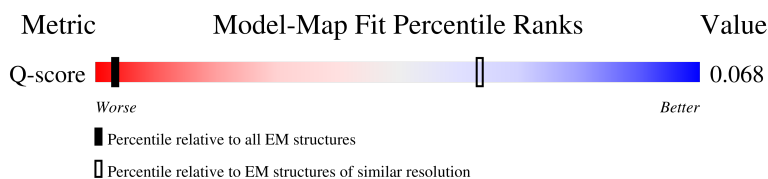
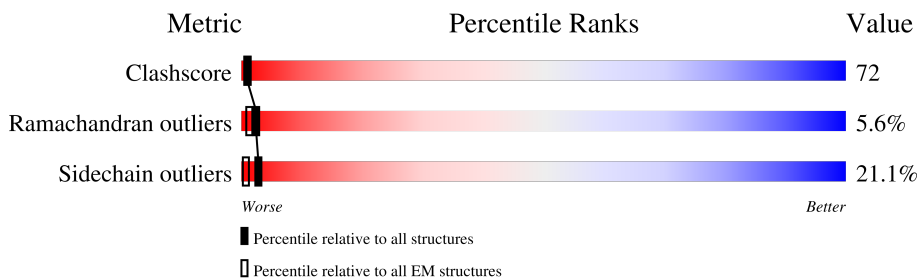
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 12.00 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	93 (11.50 - 12.50)

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	335	32% 32% 47% 16% • •
1	B	335	18% 33% 40% 20% • •
1	C	335	17% 32% 46% 16% • •
1	D	335	31% 33% 39% 20% • •

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
1	E	335	32% 30% 48% 15%
1	F	335	17% 33% 40% 19%
1	G	335	32% 31% 47% 16%
1	H	335	18% 33% 40% 19%
1	I	335	32% 30% 49% 15%
1	J	335	17% 33% 40% 19%
1	K	335	32% 31% 47% 15%
1	L	335	17% 33% 40% 19%

2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 31380 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 Baseplate structural protein Gp6.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	B	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		
1	C	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	D	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		
1	E	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	F	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		
1	G	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	H	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		
1	I	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	J	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		
1	K	326	Total	C	N	O	S	0	0
			2622	1667	424	528	3		
1	L	324	Total	C	N	O	S	0	0
			2608	1658	422	525	3		

There are 96 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	661	LEU	-	expression tag	UNP P19060
A	662	GLU	-	expression tag	UNP P19060
A	663	HIS	-	expression tag	UNP P19060
A	664	HIS	-	expression tag	UNP P19060
A	665	HIS	-	expression tag	UNP P19060
A	666	HIS	-	expression tag	UNP P19060

Continued on next page...

Continued from previous page...

Chain	Residue	Modelled	Actual	Comment	Reference
A	667	HIS	-	expression tag	UNP P19060
A	668	HIS	-	expression tag	UNP P19060
B	661	LEU	-	expression tag	UNP P19060
B	662	GLU	-	expression tag	UNP P19060
B	663	HIS	-	expression tag	UNP P19060
B	664	HIS	-	expression tag	UNP P19060
B	665	HIS	-	expression tag	UNP P19060
B	666	HIS	-	expression tag	UNP P19060
B	667	HIS	-	expression tag	UNP P19060
B	668	HIS	-	expression tag	UNP P19060
C	661	LEU	-	expression tag	UNP P19060
C	662	GLU	-	expression tag	UNP P19060
C	663	HIS	-	expression tag	UNP P19060
C	664	HIS	-	expression tag	UNP P19060
C	665	HIS	-	expression tag	UNP P19060
C	666	HIS	-	expression tag	UNP P19060
C	667	HIS	-	expression tag	UNP P19060
C	668	HIS	-	expression tag	UNP P19060
D	661	LEU	-	expression tag	UNP P19060
D	662	GLU	-	expression tag	UNP P19060
D	663	HIS	-	expression tag	UNP P19060
D	664	HIS	-	expression tag	UNP P19060
D	665	HIS	-	expression tag	UNP P19060
D	666	HIS	-	expression tag	UNP P19060
D	667	HIS	-	expression tag	UNP P19060
D	668	HIS	-	expression tag	UNP P19060
E	661	LEU	-	expression tag	UNP P19060
E	662	GLU	-	expression tag	UNP P19060
E	663	HIS	-	expression tag	UNP P19060
E	664	HIS	-	expression tag	UNP P19060
E	665	HIS	-	expression tag	UNP P19060
E	666	HIS	-	expression tag	UNP P19060
E	667	HIS	-	expression tag	UNP P19060
E	668	HIS	-	expression tag	UNP P19060
F	661	LEU	-	expression tag	UNP P19060
F	662	GLU	-	expression tag	UNP P19060
F	663	HIS	-	expression tag	UNP P19060
F	664	HIS	-	expression tag	UNP P19060
F	665	HIS	-	expression tag	UNP P19060
F	666	HIS	-	expression tag	UNP P19060
F	667	HIS	-	expression tag	UNP P19060
F	668	HIS	-	expression tag	UNP P19060

Continued on next page...

Continued from previous page...

Chain	Residue	Modelled	Actual	Comment	Reference
G	661	LEU	-	expression tag	UNP P19060
G	662	GLU	-	expression tag	UNP P19060
G	663	HIS	-	expression tag	UNP P19060
G	664	HIS	-	expression tag	UNP P19060
G	665	HIS	-	expression tag	UNP P19060
G	666	HIS	-	expression tag	UNP P19060
G	667	HIS	-	expression tag	UNP P19060
G	668	HIS	-	expression tag	UNP P19060
H	661	LEU	-	expression tag	UNP P19060
H	662	GLU	-	expression tag	UNP P19060
H	663	HIS	-	expression tag	UNP P19060
H	664	HIS	-	expression tag	UNP P19060
H	665	HIS	-	expression tag	UNP P19060
H	666	HIS	-	expression tag	UNP P19060
H	667	HIS	-	expression tag	UNP P19060
H	668	HIS	-	expression tag	UNP P19060
I	661	LEU	-	expression tag	UNP P19060
I	662	GLU	-	expression tag	UNP P19060
I	663	HIS	-	expression tag	UNP P19060
I	664	HIS	-	expression tag	UNP P19060
I	665	HIS	-	expression tag	UNP P19060
I	666	HIS	-	expression tag	UNP P19060
I	667	HIS	-	expression tag	UNP P19060
I	668	HIS	-	expression tag	UNP P19060
J	661	LEU	-	expression tag	UNP P19060
J	662	GLU	-	expression tag	UNP P19060
J	663	HIS	-	expression tag	UNP P19060
J	664	HIS	-	expression tag	UNP P19060
J	665	HIS	-	expression tag	UNP P19060
J	666	HIS	-	expression tag	UNP P19060
J	667	HIS	-	expression tag	UNP P19060
J	668	HIS	-	expression tag	UNP P19060
K	661	LEU	-	expression tag	UNP P19060
K	662	GLU	-	expression tag	UNP P19060
K	663	HIS	-	expression tag	UNP P19060
K	664	HIS	-	expression tag	UNP P19060
K	665	HIS	-	expression tag	UNP P19060
K	666	HIS	-	expression tag	UNP P19060
K	667	HIS	-	expression tag	UNP P19060
K	668	HIS	-	expression tag	UNP P19060
L	661	LEU	-	expression tag	UNP P19060
L	662	GLU	-	expression tag	UNP P19060

Continued on next page...

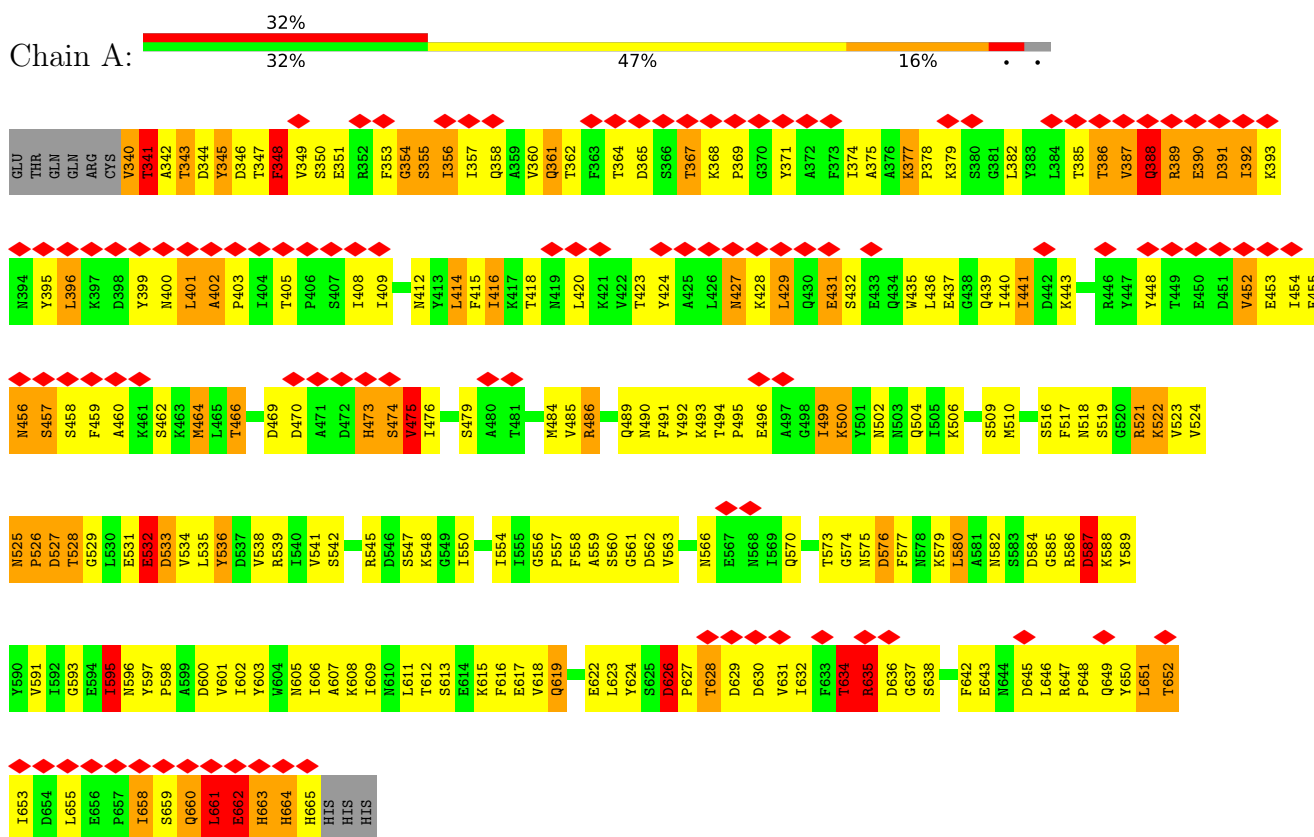
Continued from previous page...

Chain	Residue	Modelled	Actual	Comment	Reference
L	663	HIS	-	expression tag	UNP P19060
L	664	HIS	-	expression tag	UNP P19060
L	665	HIS	-	expression tag	UNP P19060
L	666	HIS	-	expression tag	UNP P19060
L	667	HIS	-	expression tag	UNP P19060
L	668	HIS	-	expression tag	UNP P19060

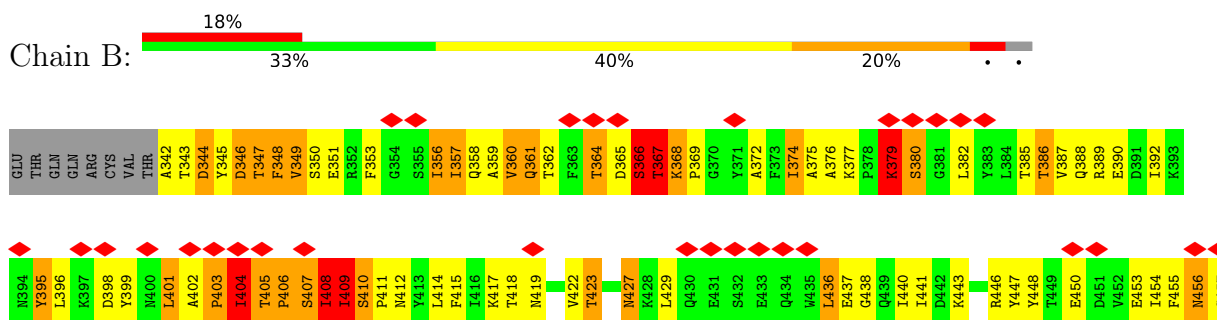
3 Residue-property plots

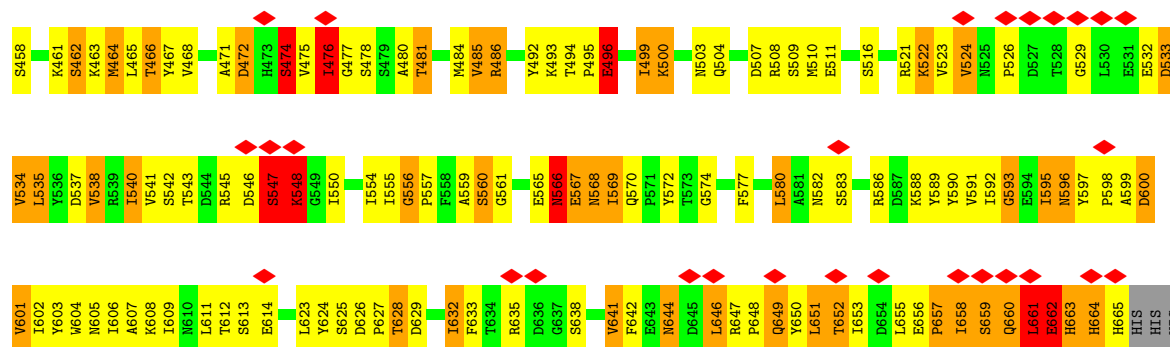
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion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

• Molecule 1: Baseplate structural protein Gp6

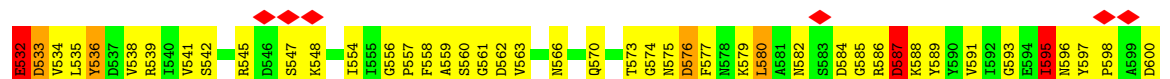
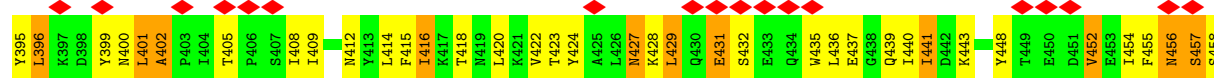
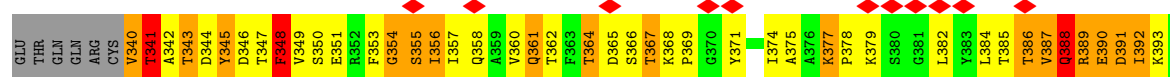


• Molecule 1: Baseplate structural protein Gp6

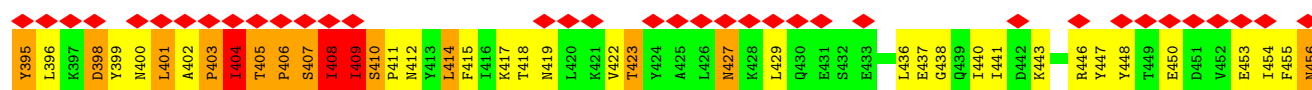


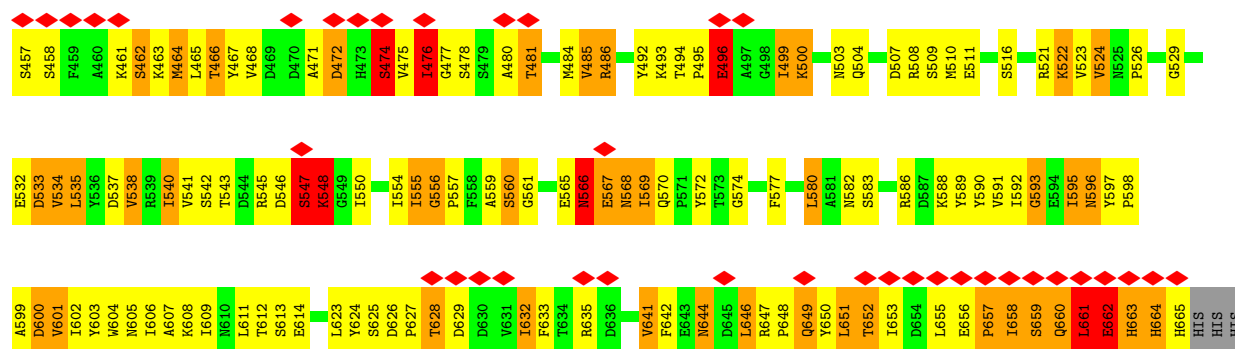


• Molecule 1: Baseplate structural protein Gp6

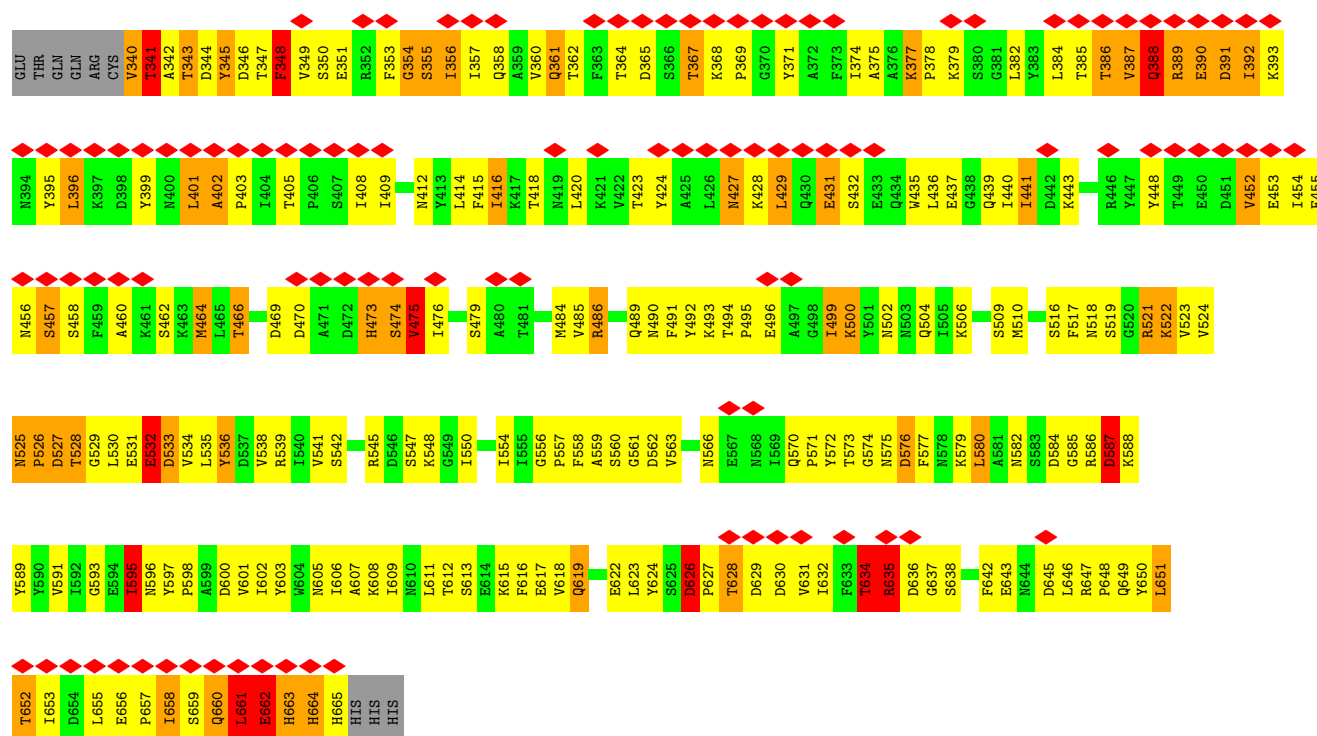


• Molecule 1: Baseplate structural protein Gp6

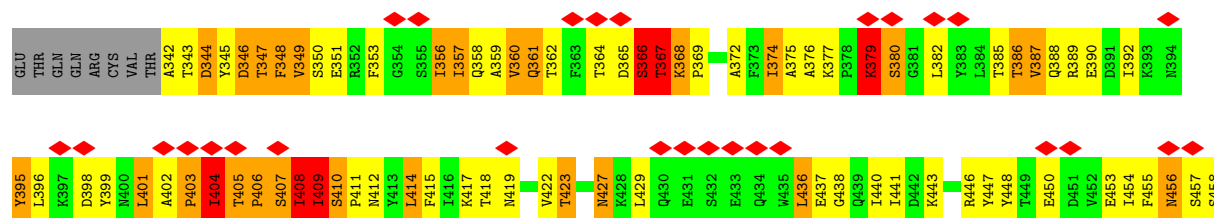


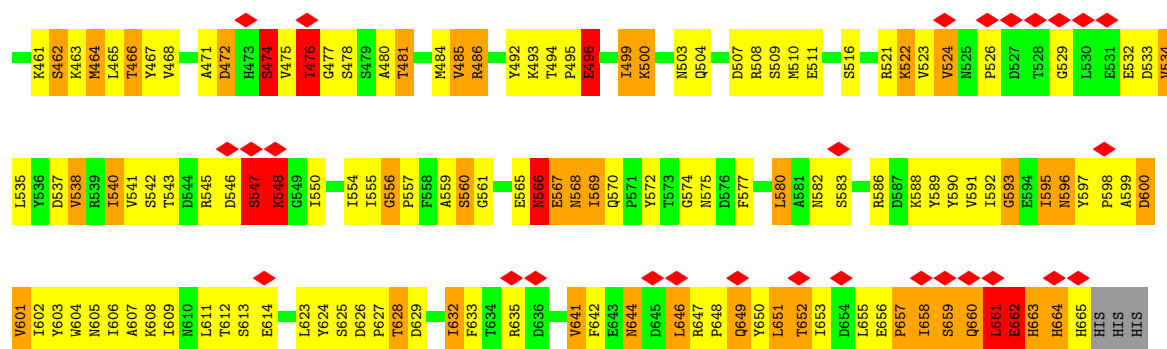


• Molecule 1: Baseplate structural protein Gp6

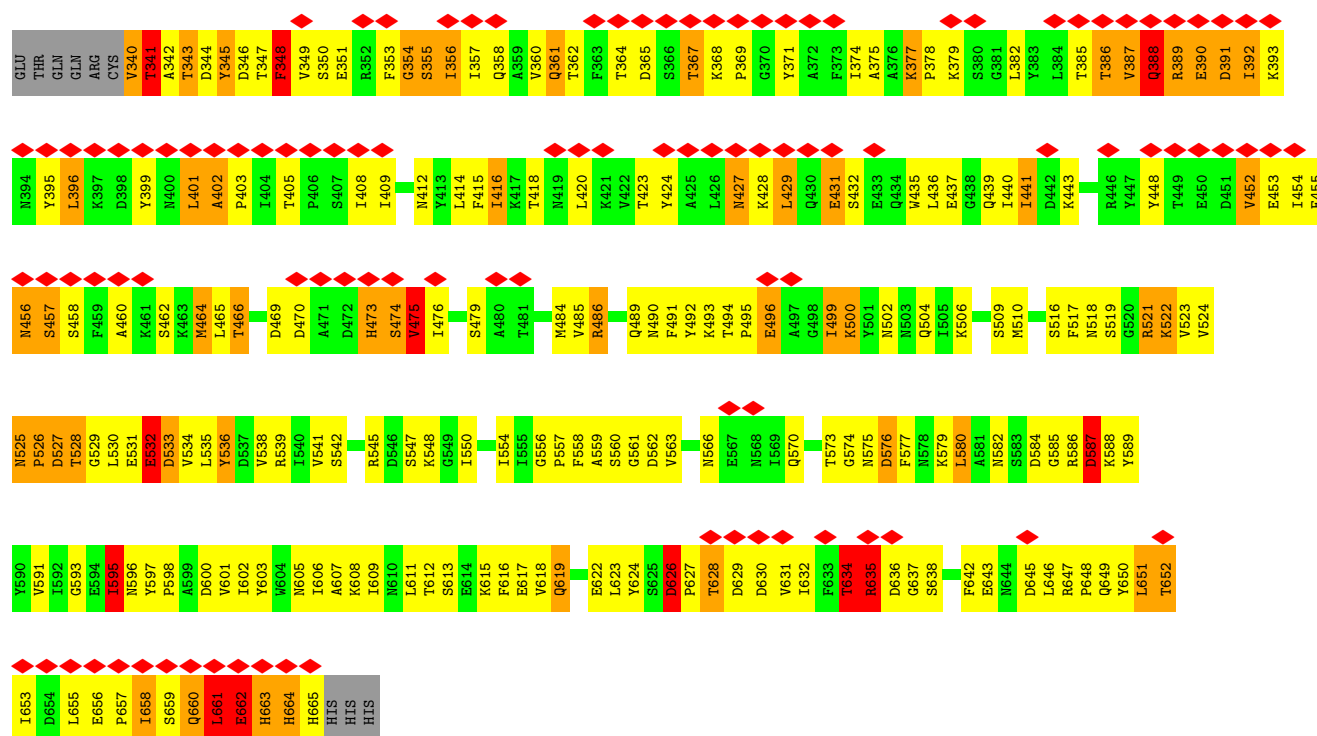


• Molecule 1: Baseplate structural protein Gp6

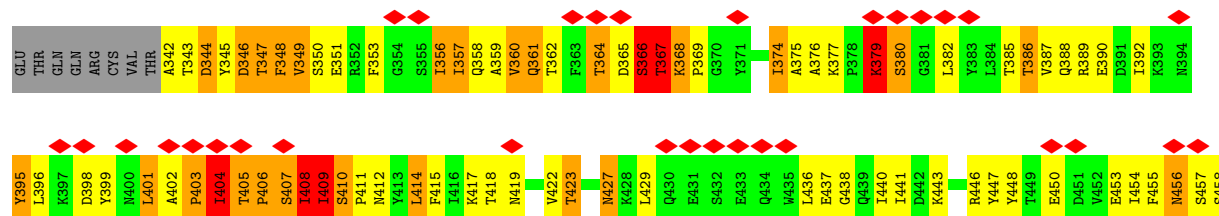


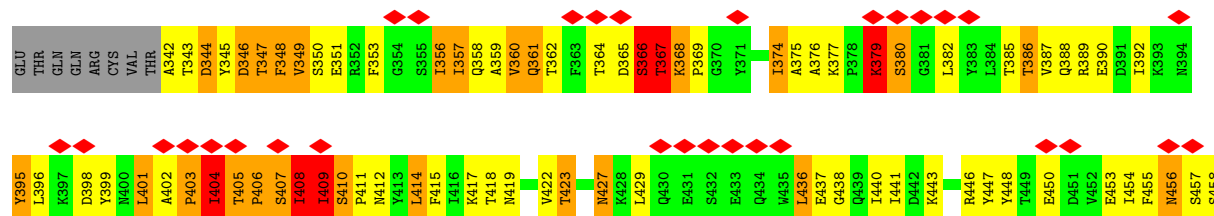


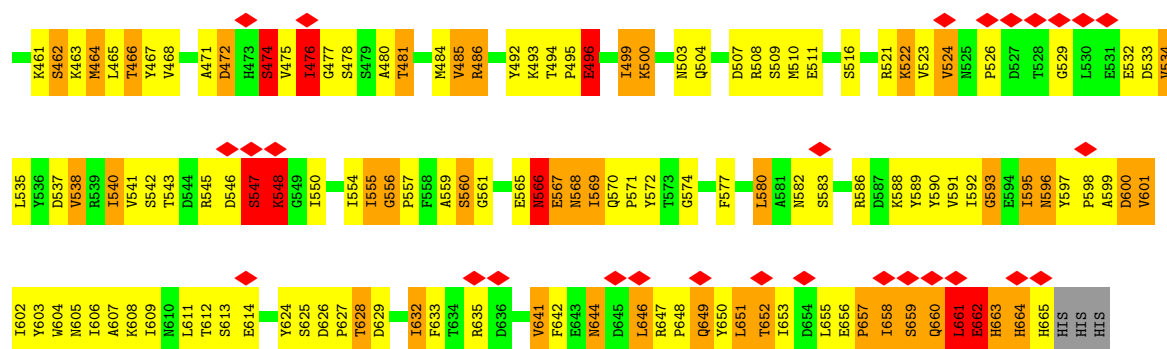
• Molecule 1: Baseplate structural protein Gp6



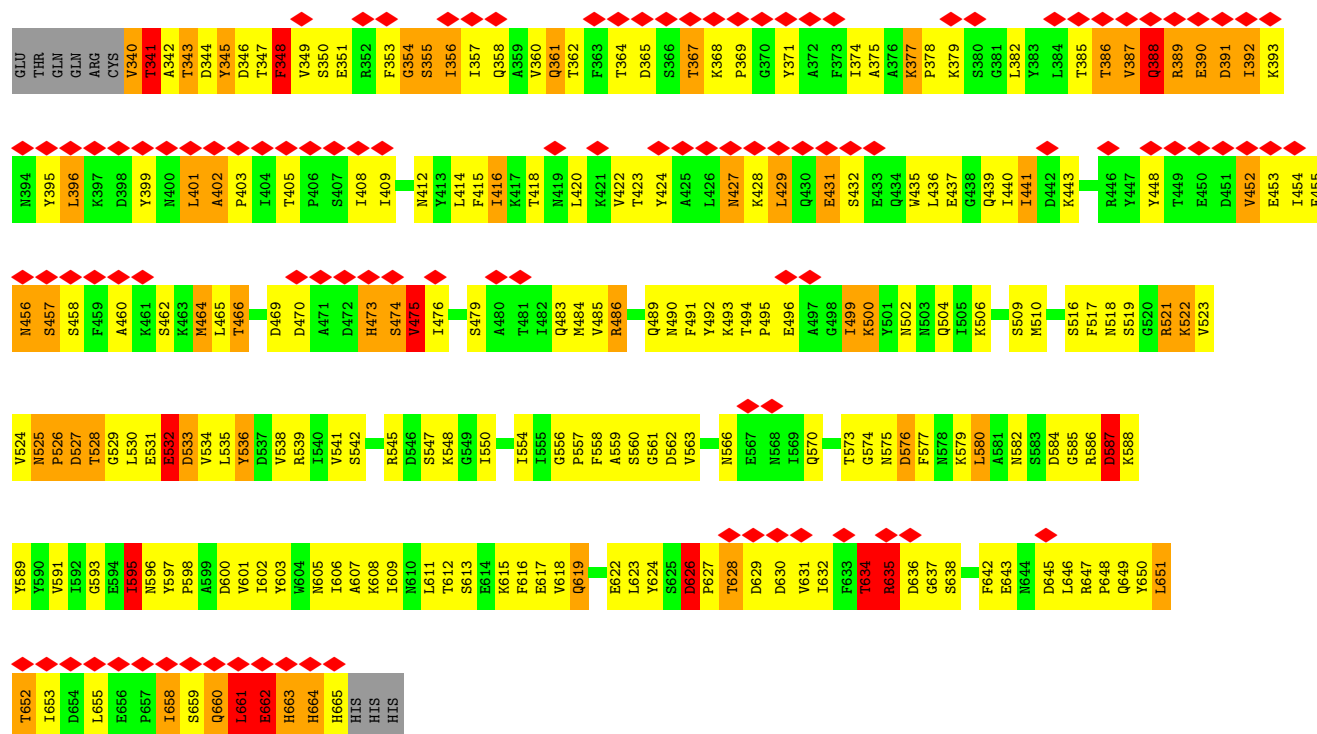
• Molecule 1: Baseplate structural protein Gp6



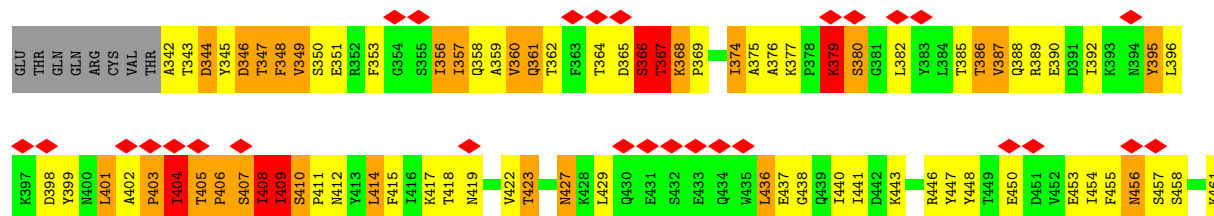


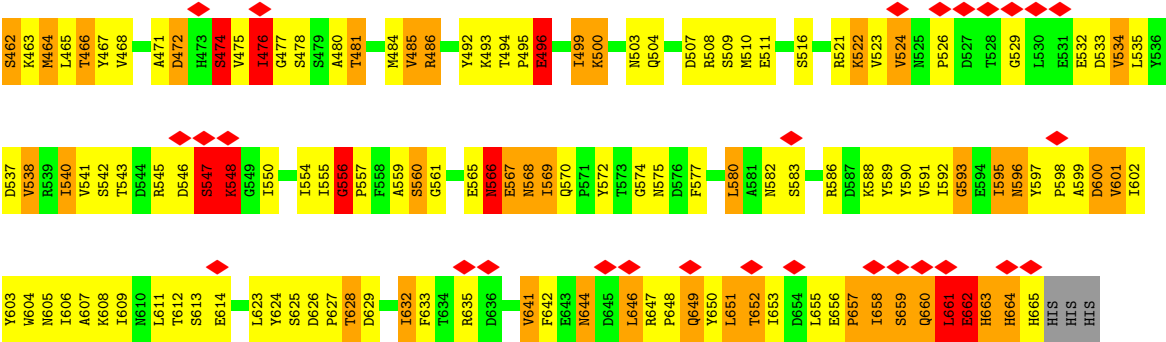


• Molecule 1: Baseplate structural protein Gp6



• Molecule 1: Baseplate structural protein Gp6





4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C6	Depositor
Number of particles used	945	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Not provided	
Microscope	FEI/PHILIPS CM300FEG/T	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	25	Depositor
Minimum defocus (nm)	1200	Depositor
Maximum defocus (nm)	5000	Depositor
Magnification	47000	Depositor
Image detector	KODAK SO-163 FILM	Depositor
Maximum map value	10.345	Depositor
Minimum map value	-7.648	Depositor
Average map value	0.000	Depositor
Map value standard deviation	1.000	Depositor
Recommended contour level	1.45	Depositor
Map size ($\text{\AA}$)	583.8291, 583.8291, 583.8291	wwPDB
Map dimensions	196, 196, 196	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing ($\text{\AA}$)	2.97872, 2.97872, 2.97872	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	0.67	0/2677	1.20	39/3637 (1.1%)
1	B	0.73	3/2663 (0.1%)	1.29	42/3617 (1.2%)
1	C	0.67	0/2677	1.19	39/3637 (1.1%)
1	D	0.73	3/2663 (0.1%)	1.29	43/3617 (1.2%)
1	E	0.67	0/2677	1.20	38/3637 (1.0%)
1	F	0.73	3/2663 (0.1%)	1.29	42/3617 (1.2%)
1	G	0.67	0/2677	1.20	39/3637 (1.1%)
1	H	0.73	3/2663 (0.1%)	1.29	42/3617 (1.2%)
1	I	0.67	0/2677	1.19	38/3637 (1.0%)
1	J	0.73	3/2663 (0.1%)	1.29	43/3617 (1.2%)
1	K	0.67	0/2677	1.20	39/3637 (1.1%)
1	L	0.73	3/2663 (0.1%)	1.29	42/3617 (1.2%)
All	All	0.70	18/32040 (0.1%)	1.25	486/43524 (1.1%)

All (18) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	J	409	ILE	CA-CB	-9.72	1.48	1.55
1	D	409	ILE	CA-CB	-9.70	1.48	1.55
1	L	409	ILE	CA-CB	-9.68	1.48	1.55
1	B	409	ILE	CA-CB	-9.67	1.48	1.55
1	H	409	ILE	CA-CB	-9.64	1.48	1.55
1	F	409	ILE	CA-CB	-9.55	1.48	1.55
1	B	476	ILE	CA-CB	-6.15	1.45	1.54
1	H	476	ILE	CA-CB	-6.14	1.45	1.54
1	J	476	ILE	CA-CB	-6.14	1.45	1.54
1	F	476	ILE	CA-CB	-6.13	1.45	1.54
1	L	476	ILE	CA-CB	-6.13	1.45	1.54
1	D	476	ILE	CA-CB	-6.08	1.45	1.54
1	B	408	ILE	CA-CB	-5.86	1.47	1.54
1	J	408	ILE	CA-CB	-5.84	1.47	1.54
1	H	408	ILE	CA-CB	-5.82	1.47	1.54
1	F	408	ILE	CA-CB	-5.80	1.47	1.54

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	L	408	ILE	CA-CB	-5.79	1.47	1.54
1	D	408	ILE	CA-CB	-5.78	1.47	1.54

All (486) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	548	LYS	N-CA-C	17.52	137.91	113.56
1	D	548	LYS	N-CA-C	17.50	137.88	113.56
1	L	548	LYS	N-CA-C	17.48	137.85	113.56
1	F	548	LYS	N-CA-C	17.46	137.84	113.56
1	H	548	LYS	N-CA-C	17.46	137.83	113.56
1	J	548	LYS	N-CA-C	17.46	137.83	113.56
1	J	405	THR	C-N-CD	-16.64	56.77	125.00
1	H	405	THR	C-N-CD	-16.64	56.77	125.00
1	D	405	THR	C-N-CD	-16.64	56.78	125.00
1	F	405	THR	C-N-CD	-16.64	56.79	125.00
1	L	405	THR	C-N-CD	-16.64	56.80	125.00
1	B	405	THR	C-N-CD	-16.63	56.81	125.00
1	G	646	LEU	N-CA-CB	-14.22	89.45	110.49
1	A	646	LEU	N-CA-CB	-14.21	89.47	110.49
1	E	646	LEU	N-CA-CB	-14.20	89.48	110.49
1	I	646	LEU	N-CA-CB	-14.18	89.50	110.49
1	K	646	LEU	N-CA-CB	-14.18	89.51	110.49
1	C	646	LEU	N-CA-CB	-14.16	89.53	110.49
1	J	534	VAL	N-CA-CB	-12.83	90.06	111.23
1	H	534	VAL	N-CA-CB	-12.81	90.10	111.23
1	D	534	VAL	N-CA-CB	-12.80	90.11	111.23
1	L	534	VAL	N-CA-CB	-12.80	90.11	111.23
1	F	534	VAL	N-CA-CB	-12.79	90.12	111.23
1	B	534	VAL	N-CA-CB	-12.79	90.13	111.23
1	B	556	GLY	CA-C-N	11.74	131.85	119.76
1	B	556	GLY	C-N-CA	11.74	131.85	119.76
1	J	556	GLY	CA-C-N	11.71	131.82	119.76
1	J	556	GLY	C-N-CA	11.71	131.82	119.76
1	D	556	GLY	CA-C-N	11.71	131.82	119.76
1	D	556	GLY	C-N-CA	11.71	131.82	119.76
1	H	556	GLY	CA-C-N	11.71	131.82	119.76
1	H	556	GLY	C-N-CA	11.71	131.82	119.76
1	F	556	GLY	CA-C-N	11.68	131.79	119.76
1	F	556	GLY	C-N-CA	11.68	131.79	119.76
1	L	556	GLY	CA-C-N	11.68	131.79	119.76
1	L	556	GLY	C-N-CA	11.68	131.79	119.76

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	474	SER	N-CA-CB	-10.39	94.03	111.27
1	L	474	SER	N-CA-CB	-10.39	94.03	111.27
1	D	474	SER	N-CA-CB	-10.38	94.04	111.27
1	F	474	SER	N-CA-CB	-10.38	94.05	111.27
1	H	474	SER	N-CA-CB	-10.37	94.06	111.27
1	J	474	SER	N-CA-CB	-10.36	94.07	111.27
1	E	387	VAL	N-CA-C	-10.27	100.14	112.98
1	A	387	VAL	N-CA-C	-10.26	100.16	112.98
1	K	387	VAL	N-CA-C	-10.25	100.17	112.98
1	G	387	VAL	N-CA-C	-10.24	100.18	112.98
1	C	387	VAL	N-CA-C	-10.23	100.20	112.98
1	I	387	VAL	N-CA-C	-10.22	100.20	112.98
1	J	547	SER	N-CA-C	-10.14	94.74	109.96
1	L	547	SER	N-CA-C	-10.12	94.78	109.96
1	D	547	SER	N-CA-C	-10.11	94.79	109.96
1	F	547	SER	N-CA-C	-10.11	94.79	109.96
1	B	547	SER	N-CA-C	-10.11	94.80	109.96
1	H	547	SER	N-CA-C	-10.11	94.80	109.96
1	H	398	ASP	CB-CA-C	9.72	126.38	110.92
1	F	398	ASP	CB-CA-C	9.72	126.37	110.92
1	L	398	ASP	CB-CA-C	9.72	126.37	110.92
1	E	457	SER	N-CA-C	-9.71	94.49	108.96
1	D	398	ASP	CB-CA-C	9.70	126.34	110.92
1	B	398	ASP	CB-CA-C	9.70	126.34	110.92
1	J	398	ASP	CB-CA-C	9.68	126.32	110.92
1	G	457	SER	N-CA-C	-9.68	94.54	108.96
1	C	457	SER	N-CA-C	-9.68	94.54	108.96
1	K	457	SER	N-CA-C	-9.67	94.55	108.96
1	A	457	SER	N-CA-C	-9.66	94.56	108.96
1	I	457	SER	N-CA-C	-9.66	94.57	108.96
1	C	525	ASN	N-CA-C	9.42	116.45	108.07
1	K	525	ASN	N-CA-C	9.41	116.44	108.07
1	E	525	ASN	N-CA-C	9.40	116.44	108.07
1	I	525	ASN	N-CA-C	9.39	116.42	108.07
1	G	525	ASN	N-CA-C	9.38	116.42	108.07
1	A	525	ASN	N-CA-C	9.38	116.42	108.07
1	A	452	VAL	N-CA-C	9.35	119.85	111.81
1	G	452	VAL	N-CA-C	9.32	119.83	111.81
1	E	452	VAL	N-CA-C	9.31	119.82	111.81
1	K	452	VAL	N-CA-C	9.30	119.81	111.81
1	G	645	ASP	CB-CA-C	9.29	126.17	110.74
1	A	645	ASP	CB-CA-C	9.29	126.16	110.74

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	C	452	VAL	N-CA-C	9.27	119.78	111.81
1	E	645	ASP	CB-CA-C	9.27	126.13	110.74
1	I	452	VAL	N-CA-C	9.27	119.78	111.81
1	I	645	ASP	CB-CA-C	9.26	126.12	110.74
1	K	645	ASP	CB-CA-C	9.25	126.10	110.74
1	C	645	ASP	CB-CA-C	9.25	126.09	110.74
1	B	472	ASP	CB-CA-C	8.99	127.34	110.62
1	J	472	ASP	CB-CA-C	8.97	127.31	110.62
1	H	472	ASP	CB-CA-C	8.97	127.30	110.62
1	L	472	ASP	CB-CA-C	8.97	127.30	110.62
1	F	472	ASP	CB-CA-C	8.96	127.29	110.62
1	D	472	ASP	CB-CA-C	8.96	127.29	110.62
1	D	379	LYS	N-CA-C	-8.91	101.49	111.82
1	H	379	LYS	N-CA-C	-8.88	101.52	111.82
1	F	379	LYS	N-CA-C	-8.87	101.53	111.82
1	L	379	LYS	N-CA-C	-8.87	101.53	111.82
1	B	379	LYS	N-CA-C	-8.87	101.53	111.82
1	J	379	LYS	N-CA-C	-8.86	101.54	111.82
1	I	456	ASN	CB-CA-C	-8.78	92.94	110.42
1	E	456	ASN	CB-CA-C	-8.78	92.95	110.42
1	K	456	ASN	CB-CA-C	-8.78	92.95	110.42
1	C	456	ASN	CB-CA-C	-8.78	92.96	110.42
1	A	456	ASN	CB-CA-C	-8.77	92.97	110.42
1	G	456	ASN	CB-CA-C	-8.77	92.97	110.42
1	A	432	SER	N-CA-C	8.72	121.13	110.41
1	K	432	SER	N-CA-C	8.71	121.12	110.41
1	E	432	SER	N-CA-C	8.70	121.11	110.41
1	I	432	SER	N-CA-C	8.70	121.11	110.41
1	C	432	SER	N-CA-C	8.68	121.09	110.41
1	G	432	SER	N-CA-C	8.68	121.08	110.41
1	A	457	SER	N-CA-CB	8.46	124.93	111.56
1	I	457	SER	N-CA-CB	8.46	124.93	111.56
1	E	457	SER	N-CA-CB	8.45	124.90	111.56
1	C	457	SER	N-CA-CB	8.44	124.89	111.56
1	G	457	SER	N-CA-CB	8.43	124.88	111.56
1	K	457	SER	N-CA-CB	8.43	124.88	111.56
1	F	405	THR	CA-C-N	8.34	130.27	119.84
1	F	405	THR	C-N-CA	8.34	130.27	119.84
1	D	405	THR	CA-C-N	8.32	130.24	119.84
1	D	405	THR	C-N-CA	8.32	130.24	119.84
1	J	405	THR	CA-C-N	8.31	130.22	119.84
1	J	405	THR	C-N-CA	8.31	130.22	119.84

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	405	THR	CA-C-N	8.30	130.21	119.84
1	B	405	THR	C-N-CA	8.30	130.21	119.84
1	L	405	THR	CA-C-N	8.29	130.20	119.84
1	L	405	THR	C-N-CA	8.29	130.20	119.84
1	H	405	THR	CA-C-N	8.29	130.20	119.84
1	H	405	THR	C-N-CA	8.29	130.20	119.84
1	L	644	ASN	N-CA-C	8.18	122.53	112.23
1	F	644	ASN	N-CA-C	8.17	122.53	112.23
1	H	644	ASN	N-CA-C	8.17	122.52	112.23
1	B	644	ASN	N-CA-C	8.14	122.49	112.23
1	D	644	ASN	N-CA-C	8.14	122.49	112.23
1	J	644	ASN	N-CA-C	8.14	122.49	112.23
1	D	366	SER	N-CA-C	-7.90	97.77	109.15
1	F	366	SER	N-CA-C	-7.89	97.78	109.15
1	B	366	SER	N-CA-C	-7.89	97.79	109.15
1	L	366	SER	N-CA-C	-7.88	97.80	109.15
1	J	366	SER	N-CA-C	-7.87	97.82	109.15
1	H	366	SER	N-CA-C	-7.87	97.82	109.15
1	C	527	ASP	CA-C-N	-7.85	106.33	122.58
1	C	527	ASP	C-N-CA	-7.85	106.33	122.58
1	I	527	ASP	CA-C-N	-7.85	106.33	122.58
1	I	527	ASP	C-N-CA	-7.85	106.33	122.58
1	G	452	VAL	CB-CA-C	-7.84	102.53	111.55
1	K	527	ASP	CA-C-N	-7.84	106.35	122.58
1	K	527	ASP	C-N-CA	-7.84	106.35	122.58
1	C	452	VAL	CB-CA-C	-7.84	102.53	111.55
1	I	452	VAL	CB-CA-C	-7.84	102.54	111.55
1	A	452	VAL	CB-CA-C	-7.83	102.54	111.55
1	A	527	ASP	CA-C-N	-7.83	106.36	122.58
1	A	527	ASP	C-N-CA	-7.83	106.36	122.58
1	K	452	VAL	CB-CA-C	-7.83	102.55	111.55
1	G	527	ASP	CA-C-N	-7.83	106.38	122.58
1	G	527	ASP	C-N-CA	-7.83	106.38	122.58
1	E	452	VAL	CB-CA-C	-7.82	102.55	111.55
1	E	527	ASP	CA-C-N	-7.81	106.42	122.58
1	E	527	ASP	C-N-CA	-7.81	106.42	122.58
1	D	635	ARG	N-CA-C	-7.71	97.58	108.23
1	F	635	ARG	N-CA-C	-7.71	97.60	108.23
1	B	635	ARG	N-CA-C	-7.70	97.60	108.23
1	J	635	ARG	N-CA-C	-7.70	97.61	108.23
1	L	635	ARG	N-CA-C	-7.69	97.61	108.23
1	H	635	ARG	N-CA-C	-7.69	97.62	108.23

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	E	573	THR	N-CA-C	7.66	119.63	111.28
1	K	573	THR	N-CA-C	7.64	119.61	111.28
1	C	573	THR	N-CA-C	7.61	119.58	111.28
1	I	573	THR	N-CA-C	7.61	119.57	111.28
1	A	573	THR	N-CA-C	7.60	119.56	111.28
1	G	573	THR	N-CA-C	7.58	119.54	111.28
1	A	388	GLN	N-CA-C	7.40	121.28	111.28
1	C	388	GLN	N-CA-C	7.40	121.27	111.28
1	I	388	GLN	N-CA-C	7.39	121.25	111.28
1	K	388	GLN	N-CA-C	7.39	121.25	111.28
1	G	388	GLN	N-CA-C	7.36	121.22	111.28
1	E	388	GLN	N-CA-C	7.34	121.19	111.28
1	L	533	ASP	CB-CA-C	-7.25	99.09	111.41
1	L	566	ASN	N-CA-C	-7.23	98.31	109.52
1	J	533	ASP	CB-CA-C	-7.22	99.13	111.41
1	H	533	ASP	CB-CA-C	-7.22	99.14	111.41
1	F	533	ASP	CB-CA-C	-7.22	99.14	111.41
1	D	566	ASN	N-CA-C	-7.21	98.34	109.52
1	B	533	ASP	CB-CA-C	-7.21	99.15	111.41
1	D	533	ASP	CB-CA-C	-7.21	99.16	111.41
1	H	566	ASN	N-CA-C	-7.20	98.36	109.52
1	J	566	ASN	N-CA-C	-7.20	98.35	109.52
1	B	566	ASN	N-CA-C	-7.20	98.36	109.52
1	F	566	ASN	N-CA-C	-7.18	98.38	109.52
1	L	583	SER	N-CA-C	-7.04	103.95	114.64
1	B	583	SER	N-CA-C	-7.02	103.98	114.64
1	J	583	SER	N-CA-C	-7.01	103.99	114.64
1	H	583	SER	N-CA-C	-7.00	103.99	114.64
1	F	583	SER	N-CA-C	-6.99	104.01	114.64
1	D	583	SER	N-CA-C	-6.98	104.03	114.64
1	I	634	THR	N-CA-C	-6.89	99.18	109.86
1	C	634	THR	N-CA-C	-6.89	99.19	109.86
1	A	634	THR	N-CA-C	-6.88	99.19	109.86
1	K	634	THR	N-CA-C	-6.87	99.21	109.86
1	G	634	THR	N-CA-C	-6.86	99.22	109.86
1	E	634	THR	N-CA-C	-6.86	99.23	109.86
1	H	407	SER	N-CA-C	-6.85	98.08	109.24
1	D	407	SER	N-CA-C	-6.84	98.09	109.24
1	E	574	GLY	N-CA-C	6.83	123.81	111.03
1	F	407	SER	N-CA-C	-6.83	98.10	109.24
1	L	407	SER	N-CA-C	-6.83	98.11	109.24
1	J	407	SER	N-CA-C	-6.83	98.11	109.24

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	407	SER	N-CA-C	-6.82	98.12	109.24
1	A	574	GLY	N-CA-C	6.82	123.78	111.03
1	C	574	GLY	N-CA-C	6.82	123.78	111.03
1	G	574	GLY	N-CA-C	6.82	123.78	111.03
1	K	574	GLY	N-CA-C	6.82	123.78	111.03
1	I	574	GLY	N-CA-C	6.80	123.75	111.03
1	H	410	SER	CA-C-N	6.77	126.75	119.78
1	H	410	SER	C-N-CA	6.77	126.75	119.78
1	L	410	SER	CA-C-N	6.73	126.72	119.78
1	L	410	SER	C-N-CA	6.73	126.72	119.78
1	J	410	SER	CA-C-N	6.73	126.71	119.78
1	J	410	SER	C-N-CA	6.73	126.71	119.78
1	D	410	SER	CA-C-N	6.72	126.70	119.78
1	D	410	SER	C-N-CA	6.72	126.70	119.78
1	B	410	SER	CA-C-N	6.71	126.69	119.78
1	B	410	SER	C-N-CA	6.71	126.69	119.78
1	F	410	SER	CA-C-N	6.71	126.69	119.78
1	F	410	SER	C-N-CA	6.71	126.69	119.78
1	F	367	THR	CB-CA-C	-6.68	97.13	110.42
1	B	367	THR	CB-CA-C	-6.68	97.13	110.42
1	D	367	THR	CB-CA-C	-6.67	97.14	110.42
1	J	367	THR	CB-CA-C	-6.67	97.14	110.42
1	H	367	THR	CB-CA-C	-6.67	97.15	110.42
1	L	367	THR	CB-CA-C	-6.67	97.15	110.42
1	D	404	ILE	N-CA-C	-6.62	100.63	109.30
1	L	404	ILE	N-CA-C	-6.62	100.63	109.30
1	B	404	ILE	N-CA-C	-6.61	100.64	109.30
1	H	404	ILE	N-CA-C	-6.60	100.65	109.30
1	F	404	ILE	N-CA-C	-6.59	100.67	109.30
1	J	404	ILE	N-CA-C	-6.58	100.68	109.30
1	B	657	PRO	CB-CA-C	-6.56	103.44	111.11
1	J	657	PRO	CB-CA-C	-6.54	103.46	111.11
1	F	657	PRO	CB-CA-C	-6.54	103.46	111.11
1	I	626	ASP	CA-C-N	6.54	126.48	120.21
1	I	626	ASP	C-N-CA	6.54	126.48	120.21
1	E	626	ASP	CA-C-N	6.53	126.48	120.21
1	E	626	ASP	C-N-CA	6.53	126.48	120.21
1	K	626	ASP	CA-C-N	6.53	126.48	120.21
1	K	626	ASP	C-N-CA	6.53	126.48	120.21
1	L	657	PRO	CB-CA-C	-6.52	103.48	111.11
1	D	657	PRO	CB-CA-C	-6.52	103.48	111.11
1	H	657	PRO	CB-CA-C	-6.52	103.48	111.11

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	C	626	ASP	CA-C-N	6.50	126.45	120.21
1	C	626	ASP	C-N-CA	6.50	126.45	120.21
1	A	626	ASP	CA-C-N	6.50	126.45	120.21
1	A	626	ASP	C-N-CA	6.50	126.45	120.21
1	G	626	ASP	CA-C-N	6.50	126.45	120.21
1	G	626	ASP	C-N-CA	6.50	126.45	120.21
1	A	368	LYS	CA-C-N	6.39	126.76	119.92
1	A	368	LYS	C-N-CA	6.39	126.76	119.92
1	G	368	LYS	CA-C-N	6.38	126.74	119.92
1	G	368	LYS	C-N-CA	6.38	126.74	119.92
1	C	368	LYS	CA-C-N	6.37	126.74	119.92
1	C	368	LYS	C-N-CA	6.37	126.74	119.92
1	K	368	LYS	CA-C-N	6.37	126.73	119.92
1	K	368	LYS	C-N-CA	6.37	126.73	119.92
1	H	476	ILE	CB-CA-C	-6.35	102.31	112.16
1	D	476	ILE	CB-CA-C	-6.35	102.32	112.16
1	E	368	LYS	CA-C-N	6.34	126.71	119.92
1	E	368	LYS	C-N-CA	6.34	126.71	119.92
1	I	368	LYS	CA-C-N	6.34	126.70	119.92
1	I	368	LYS	C-N-CA	6.34	126.70	119.92
1	B	476	ILE	CB-CA-C	-6.34	102.34	112.16
1	L	476	ILE	CB-CA-C	-6.33	102.34	112.16
1	F	476	ILE	CB-CA-C	-6.33	102.36	112.16
1	J	476	ILE	CB-CA-C	-6.32	102.37	112.16
1	H	641	VAL	N-CA-C	6.22	116.56	108.35
1	F	641	VAL	N-CA-C	6.21	116.54	108.35
1	D	641	VAL	N-CA-C	6.20	116.54	108.35
1	B	641	VAL	N-CA-C	6.19	116.52	108.35
1	J	641	VAL	N-CA-C	6.19	116.52	108.35
1	L	641	VAL	N-CA-C	6.18	116.51	108.35
1	I	661	LEU	CA-C-N	-6.11	109.88	121.54
1	I	661	LEU	C-N-CA	-6.11	109.88	121.54
1	G	661	LEU	CA-C-N	-6.09	109.90	121.54
1	G	661	LEU	C-N-CA	-6.09	109.90	121.54
1	C	661	LEU	CA-C-N	-6.08	109.92	121.54
1	C	661	LEU	C-N-CA	-6.08	109.92	121.54
1	A	661	LEU	CA-C-N	-6.08	109.92	121.54
1	A	661	LEU	C-N-CA	-6.08	109.92	121.54
1	K	661	LEU	CA-C-N	-6.08	109.93	121.54
1	K	661	LEU	C-N-CA	-6.08	109.93	121.54
1	E	588	LYS	N-CA-C	-6.06	97.93	107.93
1	A	588	LYS	N-CA-C	-6.05	97.95	107.93

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	E	661	LEU	CA-C-N	-6.05	109.99	121.54
1	E	661	LEU	C-N-CA	-6.05	109.99	121.54
1	K	588	LYS	N-CA-C	-6.05	97.95	107.93
1	G	588	LYS	N-CA-C	-6.03	97.99	107.93
1	K	630	ASP	N-CA-C	-6.03	105.69	113.16
1	C	588	LYS	N-CA-C	-6.02	98.00	107.93
1	G	630	ASP	N-CA-C	-6.02	105.70	113.16
1	E	630	ASP	N-CA-C	-6.02	105.70	113.16
1	I	588	LYS	N-CA-C	-6.02	98.00	107.93
1	I	630	ASP	N-CA-C	-6.00	105.72	113.16
1	A	630	ASP	N-CA-C	-5.99	105.74	113.16
1	C	630	ASP	N-CA-C	-5.98	105.74	113.16
1	D	644	ASN	CB-CA-C	-5.96	98.23	110.38
1	F	533	ASP	N-CA-C	-5.95	95.39	107.41
1	J	644	ASN	CB-CA-C	-5.95	98.24	110.38
1	F	644	ASN	CB-CA-C	-5.95	98.24	110.38
1	D	533	ASP	N-CA-C	-5.95	95.39	107.41
1	L	644	ASN	CB-CA-C	-5.95	98.24	110.38
1	B	644	ASN	CB-CA-C	-5.95	98.25	110.38
1	H	644	ASN	CB-CA-C	-5.94	98.26	110.38
1	B	533	ASP	N-CA-C	-5.93	95.43	107.41
1	J	533	ASP	N-CA-C	-5.93	95.44	107.41
1	L	533	ASP	N-CA-C	-5.92	95.44	107.41
1	H	533	ASP	N-CA-C	-5.92	95.45	107.41
1	C	532	GLU	CB-CA-C	-5.81	109.89	116.63
1	E	532	GLU	CB-CA-C	-5.80	109.90	116.63
1	I	532	GLU	CB-CA-C	-5.80	109.90	116.63
1	A	532	GLU	CB-CA-C	-5.79	109.91	116.63
1	G	532	GLU	CB-CA-C	-5.79	109.91	116.63
1	K	532	GLU	CB-CA-C	-5.79	109.92	116.63
1	J	635	ARG	CB-CA-C	-5.79	101.94	113.80
1	F	472	ASP	N-CA-C	-5.78	99.94	108.79
1	H	662	GLU	CA-C-N	-5.78	110.50	121.54
1	H	662	GLU	C-N-CA	-5.78	110.50	121.54
1	B	472	ASP	N-CA-C	-5.78	99.94	108.79
1	J	472	ASP	N-CA-C	-5.78	99.95	108.79
1	F	635	ARG	CB-CA-C	-5.78	101.95	113.80
1	L	635	ARG	CB-CA-C	-5.78	101.96	113.80
1	B	635	ARG	CB-CA-C	-5.78	101.96	113.80
1	H	472	ASP	N-CA-C	-5.77	99.96	108.79
1	D	635	ARG	CB-CA-C	-5.77	101.97	113.80
1	H	635	ARG	CB-CA-C	-5.77	101.97	113.80

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	L	662	GLU	CA-C-N	-5.77	110.52	121.54
1	L	662	GLU	C-N-CA	-5.77	110.52	121.54
1	D	472	ASP	N-CA-C	-5.77	99.97	108.79
1	L	472	ASP	N-CA-C	-5.76	99.97	108.79
1	F	662	GLU	CA-C-N	-5.75	110.55	121.54
1	F	662	GLU	C-N-CA	-5.75	110.55	121.54
1	D	662	GLU	CA-C-N	-5.75	110.55	121.54
1	D	662	GLU	C-N-CA	-5.75	110.55	121.54
1	J	662	GLU	CA-C-N	-5.75	110.56	121.54
1	J	662	GLU	C-N-CA	-5.75	110.56	121.54
1	B	662	GLU	CA-C-N	-5.74	110.57	121.54
1	B	662	GLU	C-N-CA	-5.74	110.57	121.54
1	F	401	LEU	CB-CA-C	5.74	119.55	109.80
1	G	476	ILE	CB-CA-C	-5.73	104.53	112.04
1	D	401	LEU	CB-CA-C	5.72	119.53	109.80
1	E	637	GLY	N-CA-C	-5.72	107.77	115.21
1	K	637	GLY	N-CA-C	-5.72	107.77	115.21
1	A	637	GLY	N-CA-C	-5.71	107.78	115.21
1	G	637	GLY	N-CA-C	-5.71	107.79	115.21
1	I	637	GLY	N-CA-C	-5.71	107.79	115.21
1	J	401	LEU	CB-CA-C	5.71	119.50	109.80
1	L	401	LEU	CB-CA-C	5.71	119.50	109.80
1	B	401	LEU	CB-CA-C	5.70	119.49	109.80
1	H	401	LEU	CB-CA-C	5.70	119.49	109.80
1	A	476	ILE	CB-CA-C	-5.70	104.58	112.04
1	I	476	ILE	CB-CA-C	-5.70	104.58	112.04
1	K	476	ILE	CB-CA-C	-5.70	104.58	112.04
1	E	476	ILE	CB-CA-C	-5.69	104.58	112.04
1	C	476	ILE	CB-CA-C	-5.68	104.60	112.04
1	C	637	GLY	N-CA-C	-5.68	107.83	115.21
1	B	405	THR	O-C-N	5.65	127.62	121.23
1	H	405	THR	O-C-N	5.64	127.60	121.23
1	L	405	THR	O-C-N	5.63	127.59	121.23
1	D	405	THR	O-C-N	5.62	127.58	121.23
1	J	405	THR	O-C-N	5.59	127.54	121.23
1	F	405	THR	O-C-N	5.57	127.52	121.23
1	G	539	ARG	N-CA-C	5.53	116.67	108.60
1	C	539	ARG	N-CA-C	5.52	116.66	108.60
1	A	377	LYS	CA-C-N	5.51	125.22	119.82
1	A	377	LYS	C-N-CA	5.51	125.22	119.82
1	K	377	LYS	CA-C-N	5.50	125.21	119.82
1	K	377	LYS	C-N-CA	5.50	125.21	119.82

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	E	539	ARG	N-CA-C	5.50	116.63	108.60
1	A	539	ARG	N-CA-C	5.50	116.63	108.60
1	I	539	ARG	N-CA-C	5.49	116.62	108.60
1	K	539	ARG	N-CA-C	5.49	116.61	108.60
1	G	377	LYS	CA-C-N	5.48	125.19	119.82
1	G	377	LYS	C-N-CA	5.48	125.19	119.82
1	C	377	LYS	CA-C-N	5.48	125.19	119.82
1	C	377	LYS	C-N-CA	5.48	125.19	119.82
1	I	377	LYS	CA-C-N	5.48	125.19	119.82
1	I	377	LYS	C-N-CA	5.48	125.19	119.82
1	E	377	LYS	CA-C-N	5.45	125.16	119.82
1	E	377	LYS	C-N-CA	5.45	125.16	119.82
1	H	347	THR	CB-CA-C	-5.45	102.33	110.88
1	D	347	THR	CB-CA-C	-5.44	102.34	110.88
1	B	347	THR	CB-CA-C	-5.43	102.35	110.88
1	J	347	THR	CB-CA-C	-5.42	102.37	110.88
1	F	347	THR	CB-CA-C	-5.42	102.38	110.88
1	L	347	THR	CB-CA-C	-5.42	102.38	110.88
1	G	595	ILE	CB-CA-C	-5.36	103.55	110.84
1	E	595	ILE	CB-CA-C	-5.36	103.56	110.84
1	K	595	ILE	CB-CA-C	-5.35	103.56	110.84
1	A	595	ILE	CB-CA-C	-5.33	103.59	110.84
1	I	595	ILE	CB-CA-C	-5.33	103.59	110.84
1	C	595	ILE	CB-CA-C	-5.33	103.59	110.84
1	G	628	THR	N-CA-C	-5.32	106.46	113.17
1	I	431	GLU	CB-CA-C	-5.32	101.24	110.45
1	K	431	GLU	CB-CA-C	-5.32	101.25	110.45
1	C	431	GLU	CB-CA-C	-5.32	101.25	110.45
1	A	431	GLU	CB-CA-C	-5.31	101.26	110.45
1	E	431	GLU	CB-CA-C	-5.31	101.26	110.45
1	E	628	THR	N-CA-C	-5.30	106.49	113.17
1	K	628	THR	N-CA-C	-5.30	106.49	113.17
1	G	431	GLU	CB-CA-C	-5.29	101.29	110.45
1	A	628	THR	N-CA-C	-5.29	106.51	113.17
1	I	628	THR	N-CA-C	-5.28	106.52	113.17
1	C	628	THR	N-CA-C	-5.26	106.54	113.17
1	D	395	TYR	N-CA-C	-5.22	106.93	113.72
1	H	450	GLU	N-CA-C	5.21	116.77	111.14
1	H	395	TYR	N-CA-C	-5.21	106.95	113.72
1	F	398	ASP	N-CA-C	-5.20	104.67	111.02
1	F	450	GLU	N-CA-C	5.20	116.76	111.14
1	J	398	ASP	N-CA-C	-5.19	104.69	111.02

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	398	ASP	N-CA-C	-5.19	104.69	111.02
1	H	398	ASP	N-CA-C	-5.19	104.69	111.02
1	B	450	GLU	N-CA-C	5.19	116.74	111.14
1	L	450	GLU	N-CA-C	5.18	116.73	111.14
1	J	395	TYR	N-CA-C	-5.18	106.99	113.72
1	L	395	TYR	N-CA-C	-5.17	106.99	113.72
1	A	386	THR	N-CA-C	-5.17	106.80	113.01
1	F	395	TYR	N-CA-C	-5.17	107.00	113.72
1	G	386	THR	N-CA-C	-5.17	106.80	113.01
1	J	450	GLU	N-CA-C	5.17	116.72	111.14
1	D	450	GLU	N-CA-C	5.17	116.72	111.14
1	L	398	ASP	N-CA-C	-5.17	104.72	111.02
1	B	395	TYR	N-CA-C	-5.16	107.02	113.72
1	D	398	ASP	N-CA-C	-5.16	104.73	111.02
1	I	473	HIS	N-CA-C	-5.15	106.94	113.18
1	L	403	PRO	CB-CA-C	5.15	117.98	111.39
1	E	473	HIS	N-CA-C	-5.14	106.96	113.18
1	I	386	THR	N-CA-C	-5.14	106.84	113.01
1	C	386	THR	N-CA-C	-5.14	106.84	113.01
1	E	386	THR	N-CA-C	-5.14	106.84	113.01
1	K	473	HIS	N-CA-C	-5.14	106.96	113.18
1	A	473	HIS	N-CA-C	-5.13	106.97	113.18
1	B	403	PRO	CB-CA-C	5.13	117.96	111.39
1	G	473	HIS	N-CA-C	-5.13	106.97	113.18
1	K	386	THR	N-CA-C	-5.13	106.86	113.01
1	G	386	THR	CA-C-N	5.12	129.38	121.34
1	G	386	THR	C-N-CA	5.12	129.38	121.34
1	C	473	HIS	N-CA-C	-5.12	106.98	113.18
1	I	386	THR	CA-C-N	5.11	129.36	121.34
1	I	386	THR	C-N-CA	5.11	129.36	121.34
1	H	403	PRO	CB-CA-C	5.11	117.92	111.39
1	D	403	PRO	CB-CA-C	5.10	117.92	111.39
1	I	521	ARG	N-CA-C	5.10	116.69	110.41
1	K	386	THR	CA-C-N	5.09	129.34	121.34
1	K	386	THR	C-N-CA	5.09	129.34	121.34
1	C	386	THR	CA-C-N	5.09	129.34	121.34
1	C	386	THR	C-N-CA	5.09	129.34	121.34
1	E	386	THR	CA-C-N	5.09	129.33	121.34
1	E	386	THR	C-N-CA	5.09	129.33	121.34
1	F	456	ASN	CB-CA-C	-5.09	100.72	110.65
1	B	628	THR	N-CA-C	-5.09	105.43	111.69
1	H	456	ASN	CB-CA-C	-5.09	100.72	110.65

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	L	628	THR	N-CA-C	-5.09	105.43	111.69
1	B	456	ASN	CB-CA-C	-5.09	100.73	110.65
1	F	403	PRO	CB-CA-C	5.09	117.90	111.39
1	J	456	ASN	CB-CA-C	-5.09	100.73	110.65
1	A	386	THR	CA-C-N	5.08	129.31	121.34
1	A	386	THR	C-N-CA	5.08	129.31	121.34
1	J	403	PRO	CB-CA-C	5.08	117.89	111.39
1	D	456	ASN	CB-CA-C	-5.07	100.75	110.65
1	L	456	ASN	CB-CA-C	-5.07	100.76	110.65
1	F	628	THR	N-CA-C	-5.07	105.45	111.69
1	E	521	ARG	N-CA-C	5.07	116.64	110.41
1	F	569	ILE	CB-CA-C	-5.06	102.98	111.29
1	A	521	ARG	N-CA-C	5.06	116.63	110.41
1	D	628	THR	N-CA-C	-5.06	105.47	111.69
1	K	521	ARG	N-CA-C	5.05	116.62	110.41
1	J	628	THR	N-CA-C	-5.05	105.48	111.69
1	H	628	THR	N-CA-C	-5.05	105.48	111.69
1	G	521	ARG	N-CA-C	5.05	116.62	110.41
1	B	569	ILE	CB-CA-C	-5.05	103.01	111.29
1	C	521	ARG	N-CA-C	5.05	116.62	110.41
1	G	663	HIS	CB-CA-C	-5.04	104.10	111.77
1	L	569	ILE	CB-CA-C	-5.04	103.02	111.29
1	D	569	ILE	CB-CA-C	-5.03	103.04	111.29
1	K	663	HIS	CB-CA-C	-5.03	104.12	111.77
1	J	569	ILE	CB-CA-C	-5.03	103.05	111.29
1	C	663	HIS	CB-CA-C	-5.02	104.14	111.77
1	H	569	ILE	CB-CA-C	-5.02	103.05	111.29
1	A	663	HIS	CB-CA-C	-5.02	104.14	111.77
1	D	555	ILE	N-CA-C	5.00	115.12	108.11
1	J	555	ILE	N-CA-C	5.00	115.11	108.11

There are no chirality outliers.

There are no planarity outliers.

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	2622	0	2545	409	0
1	B	2608	0	2530	387	0
1	C	2622	0	2545	411	0
1	D	2608	0	2530	389	0
1	E	2622	0	2543	455	0
1	F	2608	0	2530	348	0
1	G	2622	0	2543	387	0
1	H	2608	0	2530	379	0
1	I	2622	0	2544	392	0
1	J	2608	0	2530	350	0
1	K	2622	0	2543	427	0
1	L	2608	0	2530	350	0
All	All	31380	0	30443	4423	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 72.

All (4423) 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:366:SER:CB	1:K:403:PRO:HB3	1.21	1.65
1:A:664:HIS:NE2	1:K:348:PHE:CZ	1.70	1.60
1:I:400:ASN:ND2	1:K:664:HIS:CE1	1.69	1.58
1:B:364:THR:CG2	1:K:401:LEU:CD1	1.79	1.58
1:E:348:PHE:CZ	1:G:664:HIS:NE2	1.68	1.57
1:E:401:LEU:CD1	1:H:364:THR:CG2	1.78	1.56
1:E:403:PRO:HB3	1:H:366:SER:CB	1.18	1.54
1:E:348:PHE:HZ	1:G:664:HIS:CD2	1.24	1.53
1:E:401:LEU:CD1	1:H:364:THR:HG21	1.34	1.53
1:A:664:HIS:CD2	1:K:348:PHE:HZ	1.28	1.51
1:B:364:THR:HG21	1:K:401:LEU:CD1	1.34	1.51
1:D:398:ASP:OD2	1:E:663:HIS:CD2	1.65	1.49
1:E:403:PRO:CB	1:H:366:SER:HB3	1.42	1.49
1:B:366:SER:HB3	1:K:403:PRO:CB	1.45	1.47
1:E:403:PRO:CA	1:H:366:SER:HB3	1.46	1.44
1:D:400:ASN:ND2	1:E:662:GLU:HG2	1.32	1.44
1:A:403:PRO:CG	1:C:366:SER:HB2	1.45	1.42
1:B:366:SER:HB3	1:K:403:PRO:CA	1.48	1.42
1:E:348:PHE:HZ	1:G:664:HIS:NE2	1.02	1.41
1:D:400:ASN:HD22	1:E:662:GLU:CG	1.33	1.39
1:E:403:PRO:HB3	1:H:366:SER:CA	1.50	1.39
1:I:400:ASN:HD21	1:K:664:HIS:CD2	1.40	1.39

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:400:ASN:ND2	1:K:664:HIS:ND1	1.62	1.39
1:A:661:LEU:HD13	1:A:662:GLU:N	1.06	1.38
1:B:366:SER:CB	1:K:403:PRO:CB	1.99	1.38
1:D:661:LEU:C	1:D:661:LEU:HD13	1.48	1.38
1:K:661:LEU:HD13	1:K:662:GLU:N	1.06	1.38
1:B:366:SER:CA	1:K:403:PRO:HB3	1.52	1.38
1:C:661:LEU:HD13	1:C:662:GLU:N	1.06	1.38
1:A:559:ALA:C	1:A:561:GLY:HA3	1.49	1.37
1:F:661:LEU:HD13	1:F:661:LEU:C	1.48	1.37
1:E:403:PRO:CB	1:H:366:SER:CB	1.97	1.36
1:I:400:ASN:ND2	1:K:664:HIS:CG	1.92	1.36
1:C:559:ALA:C	1:C:561:GLY:HA3	1.49	1.35
1:H:661:LEU:HD13	1:H:661:LEU:C	1.48	1.35
1:I:661:LEU:HD13	1:I:662:GLU:N	1.06	1.35
1:G:559:ALA:C	1:G:561:GLY:HA3	1.49	1.35
1:E:559:ALA:C	1:E:561:GLY:HA3	1.49	1.34
1:G:661:LEU:HD13	1:G:662:GLU:N	1.06	1.34
1:I:559:ALA:C	1:I:561:GLY:HA3	1.49	1.34
1:K:559:ALA:C	1:K:561:GLY:HA3	1.49	1.34
1:E:661:LEU:HD13	1:E:662:GLU:N	1.06	1.33
1:A:664:HIS:NE2	1:K:348:PHE:HZ	1.02	1.32
1:D:398:ASP:OD2	1:E:663:HIS:CG	1.82	1.32
1:F:408:ILE:C	1:F:408:ILE:HD12	1.56	1.30
1:D:408:ILE:C	1:D:408:ILE:HD12	1.56	1.30
1:I:525:ASN:CB	1:I:526:PRO:HD2	1.59	1.30
1:K:401:LEU:C	1:K:401:LEU:HD23	1.55	1.29
1:E:401:LEU:HD23	1:E:401:LEU:C	1.55	1.29
1:L:661:LEU:C	1:L:661:LEU:HD13	1.48	1.29
1:G:401:LEU:HD23	1:G:401:LEU:C	1.55	1.29
1:G:525:ASN:CB	1:G:526:PRO:HD2	1.59	1.29
1:J:408:ILE:C	1:J:408:ILE:HD12	1.56	1.28
1:E:385:THR:OG1	1:E:387:VAL:HG23	1.33	1.28
1:A:401:LEU:HD23	1:A:401:LEU:C	1.55	1.28
1:B:661:LEU:C	1:B:661:LEU:HD13	1.48	1.28
1:H:408:ILE:C	1:H:408:ILE:HD12	1.56	1.28
1:L:408:ILE:C	1:L:408:ILE:HD12	1.56	1.27
1:L:409:ILE:C	1:L:409:ILE:HD12	1.59	1.27
1:A:525:ASN:CB	1:A:526:PRO:HD2	1.58	1.26
1:A:401:LEU:CD1	1:C:364:THR:HG23	1.64	1.26
1:I:401:LEU:HD23	1:I:401:LEU:C	1.55	1.26
1:A:661:LEU:HD13	1:A:661:LEU:C	1.58	1.26

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:409:ILE:HD12	1:H:409:ILE:C	1.59	1.26
1:D:646:LEU:HD23	1:D:646:LEU:O	1.36	1.25
1:B:408:ILE:C	1:B:408:ILE:HD12	1.56	1.25
1:J:661:LEU:C	1:J:661:LEU:HD13	1.48	1.25
1:C:385:THR:OG1	1:C:387:VAL:HG23	1.33	1.25
1:D:400:ASN:ND2	1:E:662:GLU:CG	1.91	1.24
1:E:525:ASN:CB	1:E:526:PRO:HD2	1.59	1.24
1:J:409:ILE:C	1:J:409:ILE:HD12	1.59	1.24
1:G:385:THR:OG1	1:G:387:VAL:HG23	1.33	1.24
1:D:409:ILE:C	1:D:409:ILE:HD12	1.59	1.23
1:G:559:ALA:O	1:G:561:GLY:HA3	1.40	1.23
1:J:646:LEU:O	1:J:646:LEU:HD23	1.36	1.23
1:A:385:THR:OG1	1:A:387:VAL:HG23	1.33	1.22
1:C:401:LEU:C	1:C:401:LEU:HD23	1.55	1.22
1:E:559:ALA:O	1:E:561:GLY:HA3	1.40	1.22
1:F:646:LEU:HD23	1:F:646:LEU:O	1.36	1.22
1:I:559:ALA:O	1:I:561:GLY:HA3	1.39	1.22
1:L:661:LEU:O	1:L:663:HIS:N	1.72	1.22
1:B:646:LEU:O	1:B:646:LEU:HD23	1.36	1.22
1:G:401:LEU:HD23	1:G:402:ALA:N	1.54	1.22
1:C:401:LEU:HD23	1:C:402:ALA:N	1.54	1.22
1:K:629:ASP:HA	1:L:664:HIS:CE1	1.75	1.22
1:A:401:LEU:HD23	1:A:402:ALA:N	1.54	1.22
1:I:629:ASP:HA	1:J:664:HIS:CE1	1.75	1.22
1:C:629:ASP:HA	1:D:664:HIS:CE1	1.75	1.21
1:K:525:ASN:CB	1:K:526:PRO:HD2	1.59	1.21
1:A:629:ASP:HA	1:B:664:HIS:CE1	1.75	1.21
1:J:661:LEU:O	1:J:663:HIS:N	1.72	1.21
1:K:385:THR:OG1	1:K:387:VAL:HG23	1.33	1.21
1:B:661:LEU:O	1:B:663:HIS:N	1.72	1.21
1:C:661:LEU:HD13	1:C:661:LEU:C	1.58	1.21
1:D:661:LEU:O	1:D:663:HIS:N	1.72	1.21
1:E:348:PHE:CZ	1:G:664:HIS:CD2	2.16	1.21
1:E:401:LEU:HD23	1:E:402:ALA:N	1.54	1.21
1:H:646:LEU:O	1:H:646:LEU:HD23	1.36	1.21
1:I:385:THR:OG1	1:I:387:VAL:HG23	1.33	1.21
1:B:409:ILE:C	1:B:409:ILE:HD12	1.59	1.20
1:I:401:LEU:HD23	1:I:402:ALA:N	1.54	1.20
1:E:629:ASP:HA	1:F:664:HIS:CE1	1.75	1.20
1:L:646:LEU:HD23	1:L:646:LEU:O	1.36	1.20
1:A:664:HIS:CD2	1:K:348:PHE:CZ	2.19	1.20

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:340:VAL:C	1:C:345:TYR:HE1	1.49	1.20
1:E:661:LEU:HD13	1:E:661:LEU:C	1.58	1.20
1:E:661:LEU:CD1	1:E:662:GLU:N	2.03	1.20
1:F:661:LEU:O	1:F:663:HIS:N	1.72	1.20
1:J:367:THR:O	1:J:369:PRO:HD3	1.42	1.20
1:B:367:THR:O	1:B:369:PRO:HD3	1.42	1.20
1:G:661:LEU:CD1	1:G:662:GLU:N	2.03	1.20
1:A:661:LEU:CD1	1:A:662:GLU:N	2.03	1.20
1:C:661:LEU:CD1	1:C:662:GLU:N	2.03	1.20
1:D:367:THR:O	1:D:369:PRO:HD3	1.42	1.20
1:G:340:VAL:C	1:G:345:TYR:HE1	1.49	1.20
1:A:340:VAL:HA	1:A:345:TYR:OH	1.43	1.19
1:C:502:ASN:HD22	1:C:635:ARG:HB3	1.08	1.19
1:C:525:ASN:CB	1:C:526:PRO:HD2	1.59	1.19
1:K:401:LEU:HD23	1:K:402:ALA:N	1.54	1.19
1:E:340:VAL:C	1:E:345:TYR:HE1	1.49	1.19
1:G:629:ASP:HA	1:H:664:HIS:CE1	1.75	1.19
1:G:525:ASN:O	1:G:528:THR:HG23	1.43	1.19
1:I:661:LEU:CD1	1:I:662:GLU:N	2.03	1.19
1:K:661:LEU:HD13	1:K:661:LEU:C	1.58	1.19
1:A:340:VAL:C	1:A:345:TYR:HE1	1.49	1.19
1:A:403:PRO:HG3	1:C:366:SER:HB2	1.24	1.19
1:H:661:LEU:O	1:H:663:HIS:N	1.72	1.19
1:K:661:LEU:CD1	1:K:662:GLU:N	2.03	1.19
1:I:340:VAL:C	1:I:345:TYR:HE1	1.49	1.19
1:K:340:VAL:C	1:K:345:TYR:HE1	1.49	1.19
1:K:559:ALA:O	1:K:561:GLY:HA3	1.39	1.19
1:F:409:ILE:C	1:F:409:ILE:HD12	1.59	1.18
1:J:366:SER:C	1:J:367:THR:HG22	1.68	1.18
1:K:340:VAL:HA	1:K:345:TYR:OH	1.43	1.18
1:L:367:THR:O	1:L:369:PRO:HD3	1.42	1.18
1:H:367:THR:O	1:H:369:PRO:HD3	1.42	1.18
1:C:525:ASN:O	1:C:528:THR:HG23	1.43	1.18
1:D:409:ILE:HD12	1:D:409:ILE:O	1.44	1.18
1:L:409:ILE:HD12	1:L:409:ILE:O	1.44	1.18
1:C:559:ALA:O	1:C:561:GLY:HA3	1.39	1.17
1:G:661:LEU:HD13	1:G:661:LEU:C	1.58	1.17
1:E:340:VAL:HA	1:E:345:TYR:OH	1.43	1.17
1:E:525:ASN:O	1:E:528:THR:HG23	1.43	1.17
1:B:409:ILE:HD12	1:B:409:ILE:O	1.44	1.17
1:I:340:VAL:HA	1:I:345:TYR:OH	1.43	1.17

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:409:ILE:HD12	1:J:409:ILE:O	1.44	1.17
1:K:635:ARG:HG2	1:K:636:ASP:H	1.09	1.17
1:A:525:ASN:O	1:A:528:THR:HG23	1.43	1.16
1:G:340:VAL:HA	1:G:345:TYR:OH	1.43	1.16
1:K:525:ASN:O	1:K:528:THR:HG23	1.43	1.16
1:A:403:PRO:CB	1:C:366:SER:HB2	1.76	1.16
1:A:635:ARG:HG2	1:A:636:ASP:H	1.09	1.16
1:E:401:LEU:HD11	1:H:364:THR:CG2	1.55	1.16
1:D:647:ARG:NH1	1:D:649:GLN:OE1	1.78	1.16
1:F:647:ARG:NH1	1:F:649:GLN:OE1	1.78	1.16
1:I:525:ASN:O	1:I:528:THR:HG23	1.43	1.16
1:I:661:LEU:HD13	1:I:661:LEU:C	1.58	1.16
1:A:403:PRO:CD	1:C:366:SER:HB3	1.76	1.16
1:A:559:ALA:O	1:A:561:GLY:HA3	1.40	1.16
1:E:525:ASN:HB3	1:E:526:PRO:HD2	1.25	1.16
1:A:403:PRO:HD3	1:C:366:SER:CB	1.74	1.15
1:A:502:ASN:HD22	1:A:635:ARG:HB3	1.08	1.15
1:G:525:ASN:HB3	1:G:526:PRO:HD2	1.25	1.15
1:A:403:PRO:CD	1:C:366:SER:CB	2.23	1.15
1:F:367:THR:O	1:F:369:PRO:HD3	1.42	1.15
1:A:353:PHE:O	1:A:355:SER:N	1.80	1.15
1:E:353:PHE:O	1:E:355:SER:N	1.80	1.15
1:H:409:ILE:HD12	1:H:409:ILE:O	1.44	1.15
1:I:353:PHE:O	1:I:355:SER:N	1.80	1.15
1:J:647:ARG:NH1	1:J:649:GLN:OE1	1.78	1.15
1:I:502:ASN:HD22	1:I:635:ARG:HB3	1.08	1.15
1:C:340:VAL:HA	1:C:345:TYR:OH	1.43	1.14
1:A:525:ASN:HB3	1:A:526:PRO:HD2	1.25	1.14
1:E:385:THR:HG21	1:E:389:ARG:CA	1.78	1.14
1:L:647:ARG:NH1	1:L:649:GLN:OE1	1.78	1.14
1:C:385:THR:HG21	1:C:389:ARG:CA	1.78	1.14
1:I:385:THR:HG21	1:I:389:ARG:CA	1.78	1.14
1:B:366:SER:C	1:B:367:THR:HG22	1.68	1.14
1:F:409:ILE:HD12	1:F:409:ILE:O	1.44	1.14
1:G:502:ASN:HD22	1:G:635:ARG:HB3	1.08	1.14
1:K:353:PHE:O	1:K:355:SER:N	1.80	1.14
1:K:385:THR:HG21	1:K:389:ARG:CA	1.78	1.14
1:L:366:SER:C	1:L:367:THR:HG22	1.68	1.14
1:B:364:THR:CG2	1:K:401:LEU:HD11	1.58	1.13
1:I:635:ARG:HG2	1:I:636:ASP:H	1.09	1.13
1:C:353:PHE:O	1:C:355:SER:N	1.80	1.13

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:398:ASP:OD2	1:E:663:HIS:CB	1.97	1.13
1:G:385:THR:HG21	1:G:389:ARG:CA	1.78	1.13
1:B:647:ARG:NH1	1:B:649:GLN:OE1	1.78	1.13
1:G:353:PHE:O	1:G:355:SER:N	1.80	1.12
1:H:647:ARG:NH1	1:H:649:GLN:OE1	1.78	1.12
1:J:353:PHE:HZ	1:J:395:TYR:CE2	1.67	1.12
1:E:401:LEU:HD12	1:H:364:THR:CG2	1.63	1.12
1:I:631:VAL:HG21	1:J:476:ILE:CG2	1.80	1.12
1:G:525:ASN:CG	1:G:526:PRO:HD2	1.74	1.12
1:G:631:VAL:HG21	1:H:476:ILE:CG2	1.80	1.12
1:H:366:SER:C	1:H:367:THR:HG22	1.68	1.12
1:J:367:THR:HG23	1:J:368:LYS:H	1.00	1.12
1:C:525:ASN:CG	1:C:526:PRO:HD2	1.74	1.11
1:C:631:VAL:HG21	1:D:476:ILE:CG2	1.80	1.11
1:G:502:ASN:ND2	1:G:635:ARG:HB3	1.66	1.11
1:H:353:PHE:HZ	1:H:395:TYR:CE2	1.67	1.11
1:A:385:THR:HG21	1:A:389:ARG:CA	1.78	1.11
1:B:353:PHE:HZ	1:B:395:TYR:CE2	1.67	1.11
1:D:353:PHE:HZ	1:D:395:TYR:CE2	1.67	1.11
1:G:385:THR:CG2	1:G:389:ARG:H	1.63	1.11
1:K:631:VAL:HG21	1:L:476:ILE:CG2	1.80	1.11
1:D:398:ASP:CG	1:E:663:HIS:CD2	2.29	1.11
1:D:565:GLU:HB3	1:D:570:GLN:HE21	0.95	1.11
1:E:401:LEU:CG	1:H:364:THR:CG2	2.29	1.11
1:E:502:ASN:ND2	1:E:635:ARG:HB3	1.66	1.11
1:C:385:THR:CG2	1:C:389:ARG:H	1.63	1.11
1:C:525:ASN:HB3	1:C:526:PRO:HD2	1.25	1.11
1:E:385:THR:CG2	1:E:389:ARG:H	1.63	1.11
1:E:631:VAL:HG21	1:F:476:ILE:CG2	1.80	1.11
1:E:635:ARG:HG2	1:E:636:ASP:H	1.09	1.11
1:F:353:PHE:HZ	1:F:395:TYR:CE2	1.67	1.11
1:I:502:ASN:ND2	1:I:635:ARG:HB3	1.66	1.11
1:A:525:ASN:CG	1:A:526:PRO:HD2	1.74	1.10
1:A:631:VAL:HG21	1:B:476:ILE:CG2	1.80	1.10
1:E:502:ASN:HD22	1:E:635:ARG:HB3	1.08	1.10
1:G:635:ARG:HG2	1:G:636:ASP:H	1.09	1.10
1:K:525:ASN:HB3	1:K:526:PRO:HD2	1.25	1.10
1:I:340:VAL:C	1:I:345:TYR:CE1	2.30	1.10
1:I:385:THR:CG2	1:I:389:ARG:H	1.63	1.10
1:L:353:PHE:HZ	1:L:395:TYR:CE2	1.67	1.10
1:E:525:ASN:CG	1:E:526:PRO:HD2	1.74	1.10

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:400:ASN:ND2	1:K:664:HIS:NE2	1.98	1.10
1:K:340:VAL:C	1:K:345:TYR:CE1	2.30	1.10
1:K:525:ASN:CG	1:K:526:PRO:HD2	1.74	1.10
1:D:408:ILE:HD12	1:D:408:ILE:O	1.52	1.10
1:G:340:VAL:C	1:G:345:TYR:CE1	2.30	1.10
1:K:502:ASN:HD22	1:K:635:ARG:HB3	1.08	1.10
1:A:340:VAL:C	1:A:345:TYR:CE1	2.30	1.09
1:B:364:THR:HG21	1:K:401:LEU:CG	1.82	1.09
1:I:525:ASN:CG	1:I:526:PRO:HD2	1.74	1.09
1:K:502:ASN:ND2	1:K:635:ARG:HB3	1.66	1.09
1:A:385:THR:CG2	1:A:389:ARG:H	1.63	1.09
1:K:385:THR:CG2	1:K:389:ARG:H	1.63	1.09
1:A:502:ASN:ND2	1:A:635:ARG:HB3	1.65	1.09
1:B:367:THR:HG23	1:B:368:LYS:H	1.00	1.09
1:B:364:THR:CG2	1:K:401:LEU:CG	2.30	1.09
1:D:366:SER:C	1:D:367:THR:HG22	1.68	1.09
1:E:340:VAL:C	1:E:345:TYR:CE1	2.30	1.09
1:F:366:SER:C	1:F:367:THR:HG22	1.68	1.09
1:F:565:GLU:HB3	1:F:570:GLN:HE21	0.95	1.08
1:C:502:ASN:ND2	1:C:635:ARG:HB3	1.66	1.08
1:J:565:GLU:HB3	1:J:570:GLN:HE21	0.95	1.08
1:C:340:VAL:C	1:C:345:TYR:CE1	2.30	1.08
1:E:401:LEU:CG	1:H:364:THR:HG21	1.82	1.08
1:D:399:TYR:HA	1:E:664:HIS:HB2	1.36	1.07
1:A:403:PRO:HD3	1:C:366:SER:HB3	1.08	1.07
1:L:408:ILE:HD12	1:L:408:ILE:O	1.52	1.07
1:C:635:ARG:HG2	1:C:636:ASP:H	1.09	1.07
1:D:367:THR:HG23	1:D:368:LYS:H	1.00	1.07
1:F:408:ILE:HD12	1:F:408:ILE:O	1.52	1.07
1:J:408:ILE:HD12	1:J:408:ILE:O	1.52	1.07
1:C:385:THR:OG1	1:C:387:VAL:CG2	2.03	1.07
1:F:367:THR:HG23	1:F:368:LYS:H	1.00	1.07
1:H:408:ILE:HD12	1:H:408:ILE:O	1.52	1.07
1:A:385:THR:OG1	1:A:387:VAL:CG2	2.03	1.07
1:B:565:GLU:HB3	1:B:570:GLN:HE21	0.94	1.07
1:E:348:PHE:CE2	1:G:664:HIS:NE2	2.23	1.07
1:B:408:ILE:HD12	1:B:408:ILE:O	1.52	1.06
1:I:525:ASN:HB3	1:I:526:PRO:HD2	1.25	1.06
1:K:385:THR:OG1	1:K:387:VAL:CG2	2.03	1.06
1:A:401:LEU:HD12	1:C:364:THR:HG23	1.32	1.06
1:F:569:ILE:O	1:F:569:ILE:HG22	1.55	1.06

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:403:PRO:CG	1:C:366:SER:CB	2.32	1.06
1:D:398:ASP:HA	1:E:663:HIS:HB3	1.32	1.06
1:H:367:THR:HG23	1:H:368:LYS:H	1.00	1.06
1:E:385:THR:OG1	1:E:387:VAL:CG2	2.03	1.06
1:I:385:THR:OG1	1:I:387:VAL:CG2	2.03	1.06
1:B:364:THR:CG2	1:K:401:LEU:HD12	1.63	1.06
1:E:661:LEU:O	1:E:662:GLU:O	1.74	1.06
1:G:385:THR:OG1	1:G:387:VAL:CG2	2.03	1.06
1:A:401:LEU:HD11	1:C:364:THR:HG23	1.35	1.05
1:C:661:LEU:O	1:C:662:GLU:O	1.74	1.05
1:L:367:THR:HG23	1:L:368:LYS:H	1.00	1.05
1:A:559:ALA:C	1:A:561:GLY:CA	2.30	1.05
1:H:569:ILE:O	1:H:569:ILE:HG22	1.55	1.05
1:E:559:ALA:C	1:E:561:GLY:CA	2.30	1.05
1:H:565:GLU:HB3	1:H:570:GLN:HE21	0.95	1.05
1:L:565:GLU:HB3	1:L:570:GLN:HE21	0.95	1.05
1:G:661:LEU:O	1:G:662:GLU:O	1.74	1.05
1:E:403:PRO:HA	1:H:366:SER:HB3	1.39	1.04
1:G:525:ASN:CG	1:G:526:PRO:CD	2.30	1.04
1:C:525:ASN:CG	1:C:526:PRO:CD	2.30	1.04
1:A:525:ASN:CG	1:A:526:PRO:CD	2.30	1.04
1:C:559:ALA:C	1:C:561:GLY:CA	2.30	1.04
1:K:525:ASN:CG	1:K:526:PRO:CD	2.30	1.04
1:A:661:LEU:O	1:A:662:GLU:O	1.74	1.04
1:D:399:TYR:HA	1:E:664:HIS:CB	1.88	1.04
1:G:353:PHE:CD1	1:G:395:TYR:CE2	2.46	1.04
1:I:559:ALA:C	1:I:561:GLY:CA	2.30	1.04
1:K:559:ALA:C	1:K:561:GLY:CA	2.30	1.04
1:D:569:ILE:HG22	1:D:569:ILE:O	1.55	1.04
1:E:353:PHE:CD1	1:E:395:TYR:CE2	2.46	1.04
1:E:525:ASN:CG	1:E:526:PRO:CD	2.30	1.04
1:I:353:PHE:CD1	1:I:395:TYR:CE2	2.46	1.04
1:G:559:ALA:C	1:G:561:GLY:CA	2.30	1.03
1:H:409:ILE:C	1:H:409:ILE:CD1	2.29	1.03
1:C:353:PHE:CD1	1:C:395:TYR:CE2	2.46	1.03
1:F:660:GLN:HA	1:F:661:LEU:HB2	1.41	1.03
1:I:525:ASN:CG	1:I:526:PRO:CD	2.30	1.03
1:B:364:THR:HG22	1:K:401:LEU:CD1	1.87	1.03
1:H:661:LEU:C	1:H:661:LEU:CD1	2.30	1.03
1:I:661:LEU:O	1:I:662:GLU:O	1.74	1.03
1:K:661:LEU:O	1:K:662:GLU:O	1.74	1.03

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:353:PHE:CD1	1:A:395:TYR:CE2	2.46	1.03
1:E:635:ARG:CG	1:E:636:ASP:H	1.70	1.03
1:J:569:ILE:HG22	1:J:569:ILE:O	1.55	1.03
1:A:401:LEU:C	1:A:401:LEU:CD2	2.30	1.02
1:A:635:ARG:CG	1:A:636:ASP:H	1.70	1.02
1:A:664:HIS:NE2	1:K:348:PHE:CE2	2.27	1.02
1:B:364:THR:HG22	1:K:401:LEU:HD12	1.38	1.02
1:E:401:LEU:HD12	1:H:364:THR:HG22	1.38	1.02
1:K:353:PHE:CD1	1:K:395:TYR:CE2	2.46	1.02
1:F:565:GLU:CB	1:F:570:GLN:HE21	1.73	1.02
1:G:661:LEU:CD1	1:G:661:LEU:C	2.29	1.02
1:J:409:ILE:C	1:J:409:ILE:CD1	2.29	1.02
1:L:660:GLN:HA	1:L:661:LEU:HB2	1.41	1.02
1:B:565:GLU:CB	1:B:570:GLN:HE21	1.73	1.02
1:A:401:LEU:CD1	1:C:364:THR:CG2	2.37	1.02
1:E:661:LEU:CD1	1:E:662:GLU:HB2	1.89	1.02
1:F:661:LEU:C	1:F:661:LEU:CD1	2.30	1.02
1:G:661:LEU:CD1	1:G:662:GLU:HB2	1.89	1.02
1:H:565:GLU:CB	1:H:570:GLN:HE21	1.73	1.02
1:I:661:LEU:CD1	1:I:662:GLU:HB2	1.89	1.02
1:L:409:ILE:C	1:L:409:ILE:CD1	2.29	1.02
1:H:660:GLN:HA	1:H:661:LEU:HB2	1.41	1.02
1:I:661:LEU:CD1	1:I:661:LEU:C	2.29	1.02
1:J:565:GLU:CB	1:J:570:GLN:HE21	1.73	1.02
1:E:401:LEU:C	1:E:401:LEU:CD2	2.30	1.01
1:J:367:THR:HG23	1:J:368:LYS:N	1.70	1.01
1:K:401:LEU:C	1:K:401:LEU:CD2	2.30	1.01
1:K:661:LEU:CD1	1:K:662:GLU:HB2	1.89	1.01
1:J:661:LEU:C	1:J:661:LEU:CD1	2.30	1.01
1:J:661:LEU:O	1:J:661:LEU:HD22	1.60	1.01
1:B:409:ILE:C	1:B:409:ILE:CD1	2.29	1.01
1:C:661:LEU:CD1	1:C:662:GLU:HB2	1.89	1.01
1:F:661:LEU:HD13	1:F:661:LEU:O	1.60	1.01
1:H:661:LEU:HD13	1:H:661:LEU:O	1.60	1.01
1:H:661:LEU:O	1:H:661:LEU:HD22	1.60	1.01
1:L:565:GLU:CB	1:L:570:GLN:HE21	1.73	1.01
1:B:569:ILE:HG22	1:B:569:ILE:O	1.55	1.01
1:F:403:PRO:O	1:F:405:THR:CG2	2.09	1.01
1:F:661:LEU:O	1:F:661:LEU:HD22	1.60	1.01
1:H:403:PRO:O	1:H:405:THR:CG2	2.09	1.01
1:L:569:ILE:HG22	1:L:569:ILE:O	1.55	1.01

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:403:PRO:O	1:D:405:THR:CG2	2.09	1.01
1:D:661:LEU:O	1:D:661:LEU:HD22	1.60	1.01
1:D:662:GLU:O	1:D:663:HIS:HB2	1.60	1.01
1:I:400:ASN:ND2	1:K:664:HIS:CD2	2.11	1.01
1:J:403:PRO:O	1:J:405:THR:CG2	2.09	1.01
1:K:635:ARG:CG	1:K:636:ASP:H	1.70	1.01
1:L:661:LEU:O	1:L:661:LEU:HD22	1.60	1.01
1:D:565:GLU:CB	1:D:570:GLN:HE21	1.73	1.00
1:H:662:GLU:O	1:H:663:HIS:HB2	1.60	1.00
1:L:367:THR:HG23	1:L:368:LYS:N	1.70	1.00
1:A:661:LEU:CD1	1:A:662:GLU:HB2	1.89	1.00
1:F:408:ILE:C	1:F:408:ILE:CD1	2.30	1.00
1:F:662:GLU:O	1:F:663:HIS:HB2	1.60	1.00
1:I:401:LEU:C	1:I:401:LEU:CD2	2.30	1.00
1:E:401:LEU:CD1	1:H:364:THR:HG22	1.85	1.00
1:G:401:LEU:C	1:G:401:LEU:CD2	2.30	1.00
1:L:403:PRO:O	1:L:405:THR:CG2	2.09	1.00
1:A:403:PRO:CD	1:C:366:SER:HB2	1.90	1.00
1:B:366:SER:HB3	1:K:403:PRO:HA	1.39	1.00
1:J:661:LEU:HD13	1:J:661:LEU:O	1.60	1.00
1:D:409:ILE:C	1:D:409:ILE:CD1	2.29	1.00
1:G:665:HIS:CE1	1:H:342:ALA:HB3	1.97	0.99
1:B:661:LEU:C	1:B:661:LEU:CD1	2.30	0.99
1:D:399:TYR:HA	1:E:664:HIS:CG	1.88	0.99
1:D:660:GLN:HA	1:D:661:LEU:HB2	1.41	0.99
1:H:367:THR:HG23	1:H:368:LYS:N	1.70	0.99
1:J:660:GLN:HA	1:J:661:LEU:HB2	1.41	0.99
1:B:660:GLN:HA	1:B:661:LEU:HB2	1.41	0.99
1:C:635:ARG:CG	1:C:636:ASP:H	1.70	0.99
1:C:661:LEU:C	1:C:661:LEU:CD1	2.29	0.99
1:E:665:HIS:CE1	1:F:342:ALA:HB3	1.98	0.99
1:G:385:THR:HG23	1:G:389:ARG:H	1.27	0.99
1:J:408:ILE:C	1:J:408:ILE:CD1	2.30	0.99
1:B:403:PRO:O	1:B:405:THR:CG2	2.09	0.99
1:B:662:GLU:O	1:B:663:HIS:HB2	1.60	0.99
1:A:665:HIS:CE1	1:B:342:ALA:HB3	1.97	0.99
1:K:385:THR:HG23	1:K:389:ARG:H	1.27	0.99
1:L:408:ILE:C	1:L:408:ILE:CD1	2.30	0.99
1:B:661:LEU:HD13	1:B:661:LEU:O	1.60	0.99
1:B:661:LEU:O	1:B:661:LEU:HD22	1.60	0.99
1:K:665:HIS:CE1	1:L:342:ALA:HB3	1.98	0.99

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:661:LEU:HD13	1:D:661:LEU:O	1.60	0.99
1:I:385:THR:HG23	1:I:389:ARG:H	1.27	0.99
1:D:367:THR:HG23	1:D:368:LYS:N	1.70	0.98
1:D:398:ASP:HA	1:E:663:HIS:CB	1.92	0.98
1:L:353:PHE:CZ	1:L:395:TYR:CE2	2.51	0.98
1:F:353:PHE:CZ	1:F:395:TYR:CE2	2.51	0.98
1:G:635:ARG:CG	1:G:636:ASP:H	1.70	0.98
1:L:662:GLU:O	1:L:663:HIS:HB2	1.60	0.98
1:D:398:ASP:CG	1:E:663:HIS:HD2	1.64	0.98
1:E:340:VAL:C	1:E:341:THR:HG23	1.89	0.98
1:I:635:ARG:CG	1:I:636:ASP:H	1.70	0.98
1:I:665:HIS:CE1	1:J:342:ALA:HB3	1.98	0.98
1:L:661:LEU:C	1:L:661:LEU:CD1	2.30	0.98
1:L:661:LEU:HD13	1:L:661:LEU:O	1.60	0.98
1:B:353:PHE:CZ	1:B:395:TYR:CE2	2.51	0.98
1:C:665:HIS:CE1	1:D:342:ALA:HB3	1.97	0.98
1:B:403:PRO:O	1:B:405:THR:HG23	1.64	0.98
1:B:408:ILE:C	1:B:408:ILE:CD1	2.30	0.97
1:C:385:THR:HG21	1:C:389:ARG:C	1.89	0.97
1:E:385:THR:HG21	1:E:389:ARG:C	1.89	0.97
1:G:385:THR:HG21	1:G:389:ARG:C	1.89	0.97
1:L:565:GLU:HB3	1:L:570:GLN:NE2	1.79	0.97
1:C:389:ARG:HA	1:C:392:ILE:HG12	1.46	0.97
1:C:401:LEU:C	1:C:401:LEU:CD2	2.30	0.97
1:F:367:THR:HG23	1:F:368:LYS:N	1.70	0.97
1:J:353:PHE:CZ	1:J:395:TYR:CE2	2.51	0.97
1:D:353:PHE:CZ	1:D:395:TYR:CE2	2.51	0.97
1:G:340:VAL:C	1:G:341:THR:HG23	1.89	0.97
1:J:565:GLU:HB3	1:J:570:GLN:NE2	1.79	0.97
1:D:403:PRO:O	1:D:405:THR:HG23	1.64	0.97
1:F:403:PRO:O	1:F:405:THR:HG23	1.64	0.97
1:A:385:THR:HG23	1:A:389:ARG:H	1.27	0.97
1:A:661:LEU:C	1:A:661:LEU:CD1	2.29	0.97
1:B:565:GLU:HB3	1:B:570:GLN:NE2	1.79	0.97
1:A:389:ARG:HA	1:A:392:ILE:HG12	1.46	0.97
1:I:385:THR:HG21	1:I:389:ARG:C	1.89	0.97
1:E:403:PRO:HB3	1:H:366:SER:HB2	1.43	0.97
1:H:408:ILE:C	1:H:408:ILE:CD1	2.30	0.97
1:H:353:PHE:CZ	1:H:395:TYR:CE2	2.51	0.97
1:D:398:ASP:CA	1:E:663:HIS:HB3	1.94	0.97
1:D:660:GLN:HA	1:D:661:LEU:CB	1.95	0.97

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:403:PRO:O	1:H:405:THR:HG23	1.64	0.97
1:F:409:ILE:C	1:F:409:ILE:CD1	2.29	0.96
1:J:662:GLU:O	1:J:663:HIS:HB2	1.60	0.96
1:I:340:VAL:C	1:I:341:THR:HG23	1.89	0.96
1:I:389:ARG:HA	1:I:392:ILE:HG12	1.46	0.96
1:L:660:GLN:HA	1:L:661:LEU:CB	1.95	0.96
1:B:660:GLN:HA	1:B:661:LEU:CB	1.95	0.96
1:D:343:THR:O	1:D:347:THR:HG23	1.65	0.96
1:E:385:THR:HG23	1:E:389:ARG:H	1.27	0.96
1:H:660:GLN:HA	1:H:661:LEU:CB	1.95	0.96
1:L:403:PRO:O	1:L:405:THR:HG23	1.64	0.96
1:A:340:VAL:C	1:A:341:THR:HG23	1.89	0.96
1:B:367:THR:HG23	1:B:368:LYS:N	1.70	0.96
1:J:660:GLN:HA	1:J:661:LEU:CB	1.95	0.96
1:L:343:THR:O	1:L:347:THR:HG23	1.65	0.96
1:C:340:VAL:HA	1:C:345:TYR:CZ	2.01	0.96
1:D:398:ASP:OD2	1:E:663:HIS:HD2	1.32	0.96
1:I:353:PHE:CD1	1:I:395:TYR:HE2	1.84	0.96
1:H:565:GLU:HB3	1:H:570:GLN:NE2	1.79	0.96
1:J:403:PRO:O	1:J:405:THR:HG23	1.64	0.96
1:K:340:VAL:C	1:K:341:THR:HG23	1.89	0.96
1:C:661:LEU:HD13	1:C:662:GLU:H	1.26	0.96
1:D:565:GLU:HB3	1:D:570:GLN:NE2	1.79	0.96
1:F:503:ASN:HD21	1:F:633:PHE:H	0.96	0.96
1:F:565:GLU:HB3	1:F:570:GLN:NE2	1.79	0.96
1:I:401:LEU:HD23	1:I:402:ALA:CB	1.96	0.96
1:K:525:ASN:CB	1:K:526:PRO:CD	2.44	0.96
1:E:661:LEU:HD13	1:E:662:GLU:H	1.26	0.95
1:F:660:GLN:HA	1:F:661:LEU:CB	1.95	0.95
1:A:385:THR:HG21	1:A:389:ARG:C	1.89	0.95
1:B:503:ASN:HD21	1:B:633:PHE:H	0.96	0.95
1:C:401:LEU:CD2	1:C:402:ALA:N	2.29	0.95
1:K:385:THR:HG21	1:K:389:ARG:C	1.89	0.95
1:F:343:THR:O	1:F:347:THR:HG23	1.65	0.95
1:I:525:ASN:CB	1:I:526:PRO:CD	2.44	0.95
1:A:525:ASN:CB	1:A:526:PRO:CD	2.44	0.95
1:B:343:THR:O	1:B:347:THR:HG23	1.65	0.95
1:G:389:ARG:HA	1:G:392:ILE:HG12	1.46	0.95
1:H:343:THR:O	1:H:347:THR:HG23	1.65	0.95
1:K:401:LEU:HD23	1:K:402:ALA:CB	1.96	0.95
1:C:385:THR:HG23	1:C:389:ARG:H	1.27	0.95

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:401:LEU:CD2	1:G:402:ALA:N	2.30	0.95
1:I:401:LEU:CD2	1:I:402:ALA:N	2.30	0.95
1:L:503:ASN:HD21	1:L:633:PHE:H	0.96	0.95
1:C:353:PHE:CD1	1:C:395:TYR:HE2	1.83	0.95
1:D:400:ASN:HD22	1:E:662:GLU:HG3	1.28	0.95
1:J:343:THR:O	1:J:347:THR:HG23	1.65	0.95
1:K:661:LEU:C	1:K:661:LEU:CD1	2.29	0.95
1:A:340:VAL:HA	1:A:345:TYR:CZ	2.01	0.94
1:E:340:VAL:HA	1:E:345:TYR:CZ	2.01	0.94
1:E:401:LEU:CD2	1:E:402:ALA:N	2.30	0.94
1:G:401:LEU:HD23	1:G:402:ALA:CB	1.96	0.94
1:K:340:VAL:HA	1:K:345:TYR:CZ	2.01	0.94
1:C:340:VAL:C	1:C:341:THR:HG23	1.89	0.94
1:E:401:LEU:HD23	1:E:402:ALA:CB	1.96	0.94
1:J:503:ASN:HD21	1:J:633:PHE:H	0.96	0.94
1:E:401:LEU:HD23	1:E:402:ALA:CA	1.97	0.94
1:G:525:ASN:CB	1:G:526:PRO:CD	2.44	0.94
1:K:389:ARG:HA	1:K:392:ILE:HG12	1.46	0.94
1:C:525:ASN:CB	1:C:526:PRO:CD	2.44	0.94
1:I:340:VAL:HA	1:I:345:TYR:CZ	2.01	0.94
1:C:401:LEU:HD23	1:C:402:ALA:CB	1.96	0.94
1:D:661:LEU:C	1:D:661:LEU:CD1	2.30	0.94
1:I:385:THR:HG1	1:I:387:VAL:HG23	1.32	0.94
1:A:401:LEU:CD2	1:A:402:ALA:N	2.30	0.94
1:D:503:ASN:HD21	1:D:633:PHE:H	0.96	0.94
1:E:525:ASN:CB	1:E:526:PRO:CD	2.45	0.94
1:A:385:THR:CG2	1:A:389:ARG:N	2.31	0.94
1:A:401:LEU:HD23	1:A:402:ALA:CB	1.96	0.94
1:B:366:SER:HB2	1:K:403:PRO:HB3	1.45	0.94
1:G:340:VAL:HA	1:G:345:TYR:CZ	2.01	0.94
1:G:353:PHE:CD1	1:G:395:TYR:HE2	1.84	0.94
1:H:503:ASN:HD21	1:H:633:PHE:H	0.96	0.94
1:I:401:LEU:HD23	1:I:402:ALA:CA	1.97	0.94
1:K:401:LEU:HD23	1:K:402:ALA:CA	1.97	0.94
1:K:661:LEU:O	1:K:662:GLU:C	2.08	0.94
1:D:408:ILE:C	1:D:408:ILE:CD1	2.30	0.94
1:K:353:PHE:CD1	1:K:395:TYR:HE2	1.84	0.94
1:C:401:LEU:HD23	1:C:402:ALA:CA	1.97	0.93
1:E:389:ARG:HA	1:E:392:ILE:HG12	1.46	0.93
1:E:385:THR:CG2	1:E:389:ARG:N	2.31	0.93
1:B:364:THR:HG23	1:K:401:LEU:HG	1.50	0.93

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:566:ASN:N	1:H:570:GLN:HE22	1.66	0.93
1:G:401:LEU:HD23	1:G:402:ALA:CA	1.97	0.93
1:K:385:THR:CG2	1:K:389:ARG:N	2.31	0.93
1:E:401:LEU:HG	1:H:364:THR:HG23	1.49	0.93
1:K:401:LEU:CD2	1:K:402:ALA:N	2.30	0.93
1:A:401:LEU:HD23	1:A:402:ALA:CA	1.97	0.93
1:F:557:PRO:HG2	1:F:580:LEU:HD11	1.51	0.93
1:B:557:PRO:HG2	1:B:580:LEU:HD11	1.51	0.93
1:G:661:LEU:O	1:G:662:GLU:C	2.08	0.93
1:J:566:ASN:N	1:J:570:GLN:HE22	1.66	0.93
1:E:401:LEU:CD1	1:H:364:THR:HG23	1.98	0.92
1:G:661:LEU:HD13	1:G:662:GLU:H	1.26	0.92
1:C:661:LEU:O	1:C:662:GLU:C	2.08	0.92
1:L:566:ASN:N	1:L:570:GLN:HE22	1.66	0.92
1:A:387:VAL:HG21	1:A:390:GLU:OE1	1.70	0.92
1:A:661:LEU:O	1:A:662:GLU:C	2.08	0.92
1:G:385:THR:CG2	1:G:389:ARG:N	2.31	0.92
1:I:661:LEU:O	1:I:662:GLU:C	2.08	0.92
1:A:353:PHE:CD1	1:A:395:TYR:HE2	1.84	0.92
1:C:387:VAL:HG21	1:C:390:GLU:OE1	1.69	0.92
1:K:387:VAL:HG21	1:K:390:GLU:OE1	1.70	0.92
1:C:385:THR:CG2	1:C:389:ARG:N	2.31	0.92
1:L:647:ARG:CG	1:L:650:TYR:CD1	2.53	0.92
1:B:647:ARG:CG	1:B:650:TYR:CD1	2.53	0.92
1:A:403:PRO:CB	1:C:366:SER:CB	2.48	0.92
1:A:661:LEU:HD13	1:A:662:GLU:H	1.26	0.92
1:E:403:PRO:HB3	1:H:366:SER:HA	1.52	0.92
1:F:647:ARG:CG	1:F:650:TYR:CD1	2.53	0.92
1:H:647:ARG:CG	1:H:650:TYR:CD1	2.53	0.92
1:I:385:THR:CG2	1:I:389:ARG:N	2.31	0.92
1:K:401:LEU:CD2	1:K:402:ALA:HB2	2.00	0.92
1:A:401:LEU:CD2	1:A:402:ALA:HB2	2.00	0.91
1:B:566:ASN:N	1:B:570:GLN:HE22	1.66	0.91
1:D:647:ARG:CG	1:D:650:TYR:CD1	2.53	0.91
1:H:557:PRO:HG2	1:H:580:LEU:HD11	1.51	0.91
1:L:557:PRO:HG2	1:L:580:LEU:HD11	1.51	0.91
1:A:403:PRO:HB3	1:C:366:SER:CB	2.00	0.91
1:C:436:LEU:HD22	1:C:655:LEU:HD21	1.53	0.91
1:I:401:LEU:CD2	1:I:402:ALA:HB2	2.00	0.91
1:A:403:PRO:HB3	1:C:366:SER:HB2	1.52	0.91
1:C:401:LEU:CD2	1:C:402:ALA:HB2	2.00	0.91

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:557:PRO:HG2	1:J:580:LEU:HD11	1.52	0.91
1:E:519:SER:HB2	1:E:562:ASP:OD1	1.71	0.91
1:C:519:SER:HB2	1:C:562:ASP:OD1	1.71	0.91
1:K:436:LEU:HD22	1:K:655:LEU:HD21	1.52	0.91
1:E:661:LEU:C	1:E:661:LEU:CD1	2.29	0.91
1:I:387:VAL:HG21	1:I:390:GLU:OE1	1.70	0.91
1:A:635:ARG:CG	1:A:636:ASP:N	2.30	0.91
1:B:364:THR:HG23	1:K:401:LEU:CD1	2.01	0.91
1:G:387:VAL:HG21	1:G:390:GLU:OE1	1.70	0.91
1:F:566:ASN:N	1:F:570:GLN:HE22	1.66	0.91
1:G:401:LEU:CD2	1:G:402:ALA:HB2	2.00	0.91
1:G:519:SER:HB2	1:G:562:ASP:OD1	1.71	0.91
1:K:635:ARG:CG	1:K:636:ASP:N	2.30	0.91
1:C:517:PHE:HD2	1:C:536:TYR:HE2	1.18	0.91
1:E:401:LEU:CD2	1:E:402:ALA:HB2	2.00	0.91
1:E:661:LEU:O	1:E:662:GLU:C	2.08	0.91
1:G:517:PHE:HD2	1:G:536:TYR:HE2	1.18	0.91
1:D:566:ASN:N	1:D:570:GLN:HE22	1.66	0.91
1:J:647:ARG:CG	1:J:650:TYR:CD1	2.53	0.91
1:I:436:LEU:HD22	1:I:655:LEU:HD21	1.53	0.90
1:E:387:VAL:HG21	1:E:390:GLU:OE1	1.70	0.90
1:I:519:SER:HB2	1:I:562:ASP:OD1	1.71	0.90
1:B:364:THR:CG2	1:K:401:LEU:HG	1.97	0.90
1:D:557:PRO:HG2	1:D:580:LEU:HD11	1.51	0.90
1:I:635:ARG:HH22	1:J:661:LEU:HD12	1.36	0.90
1:K:661:LEU:HD13	1:K:662:GLU:H	1.26	0.90
1:B:409:ILE:HD13	1:B:410:SER:O	1.72	0.90
1:D:409:ILE:HD13	1:D:410:SER:O	1.72	0.90
1:D:400:ASN:CG	1:E:662:GLU:HG2	1.97	0.90
1:E:353:PHE:CD1	1:E:395:TYR:HE2	1.84	0.90
1:C:635:ARG:HH22	1:D:661:LEU:HD12	1.36	0.90
1:E:517:PHE:HD2	1:E:536:TYR:HE2	1.18	0.90
1:E:635:ARG:HH22	1:F:661:LEU:HD12	1.36	0.90
1:J:409:ILE:HD13	1:J:410:SER:O	1.72	0.90
1:K:385:THR:HG1	1:K:387:VAL:HG23	1.37	0.90
1:L:409:ILE:HD13	1:L:410:SER:O	1.72	0.90
1:A:436:LEU:HD22	1:A:655:LEU:HD21	1.53	0.90
1:C:629:ASP:HA	1:D:664:HIS:NE2	1.87	0.90
1:E:631:VAL:HG21	1:F:476:ILE:HG21	1.53	0.90
1:E:664:HIS:O	1:E:665:HIS:CG	2.25	0.90
1:G:385:THR:HG1	1:G:387:VAL:HG23	1.34	0.90

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:635:ARG:CG	1:I:636:ASP:N	2.30	0.90
1:I:664:HIS:O	1:I:665:HIS:CG	2.25	0.90
1:B:364:THR:HG23	1:K:401:LEU:CG	2.02	0.90
1:G:664:HIS:O	1:G:665:HIS:CG	2.25	0.90
1:H:409:ILE:HD13	1:H:410:SER:O	1.72	0.90
1:I:517:PHE:HD2	1:I:536:TYR:HE2	1.18	0.90
1:I:629:ASP:HA	1:J:664:HIS:NE2	1.87	0.90
1:I:661:LEU:HD13	1:I:662:GLU:H	1.26	0.90
1:B:366:SER:CA	1:K:403:PRO:CB	2.41	0.90
1:C:353:PHE:CZ	1:C:395:TYR:CD2	2.60	0.90
1:K:629:ASP:HA	1:L:664:HIS:NE2	1.87	0.90
1:K:631:VAL:HG21	1:L:476:ILE:HG21	1.53	0.90
1:E:401:LEU:HG	1:H:364:THR:CG2	1.97	0.89
1:F:409:ILE:HD13	1:F:410:SER:O	1.72	0.89
1:K:664:HIS:O	1:K:665:HIS:CG	2.25	0.89
1:A:385:THR:HG1	1:A:387:VAL:HG23	1.36	0.89
1:B:366:SER:HA	1:K:403:PRO:HB3	1.52	0.89
1:G:353:PHE:CZ	1:G:395:TYR:CD2	2.61	0.89
1:I:353:PHE:CZ	1:I:395:TYR:CD2	2.61	0.89
1:B:647:ARG:HG2	1:B:650:TYR:CD1	2.08	0.89
1:C:631:VAL:HG21	1:D:476:ILE:HG21	1.53	0.89
1:A:519:SER:HB2	1:A:562:ASP:OD1	1.71	0.89
1:D:496:GLU:H	1:D:496:GLU:CD	1.80	0.89
1:J:366:SER:O	1:J:367:THR:HG22	1.73	0.89
1:B:496:GLU:CD	1:B:496:GLU:H	1.80	0.89
1:L:496:GLU:H	1:L:496:GLU:CD	1.80	0.89
1:F:366:SER:O	1:F:367:THR:HG22	1.73	0.89
1:G:436:LEU:HD22	1:G:655:LEU:HD21	1.52	0.89
1:K:517:PHE:HD2	1:K:536:TYR:HE2	1.18	0.89
1:E:353:PHE:CZ	1:E:395:TYR:CD2	2.61	0.89
1:G:635:ARG:CG	1:G:636:ASP:N	2.30	0.89
1:H:366:SER:O	1:H:367:THR:HG22	1.73	0.89
1:E:401:LEU:CG	1:H:364:THR:HG23	2.00	0.89
1:E:661:LEU:HG	1:F:456:ASN:HB3	1.55	0.89
1:A:664:HIS:O	1:A:665:HIS:CG	2.25	0.89
1:D:366:SER:O	1:D:367:THR:HG22	1.73	0.89
1:H:662:GLU:O	1:H:663:HIS:CB	2.20	0.89
1:K:353:PHE:CZ	1:K:395:TYR:CD2	2.60	0.89
1:K:519:SER:HB2	1:K:562:ASP:OD1	1.71	0.89
1:L:366:SER:O	1:L:367:THR:HG22	1.73	0.89
1:A:353:PHE:CZ	1:A:395:TYR:CD2	2.61	0.88

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:517:PHE:HD2	1:A:536:TYR:HE2	1.18	0.88
1:C:635:ARG:CG	1:C:636:ASP:N	2.30	0.88
1:C:664:HIS:O	1:C:665:HIS:CG	2.25	0.88
1:G:635:ARG:HH22	1:H:661:LEU:HD12	1.36	0.88
1:K:340:VAL:O	1:K:345:TYR:HE1	1.55	0.88
1:E:340:VAL:O	1:E:345:TYR:HE1	1.55	0.88
1:G:661:LEU:HG	1:H:456:ASN:HB3	1.55	0.88
1:G:340:VAL:O	1:G:345:TYR:HE1	1.55	0.88
1:L:647:ARG:HG2	1:L:650:TYR:CD1	2.08	0.88
1:A:629:ASP:HA	1:B:664:HIS:NE2	1.87	0.88
1:A:635:ARG:HH22	1:B:661:LEU:HD12	1.36	0.88
1:C:340:VAL:O	1:C:345:TYR:HE1	1.55	0.88
1:E:436:LEU:HD22	1:E:655:LEU:HD21	1.53	0.88
1:F:366:SER:C	1:F:367:THR:CG2	2.42	0.88
1:H:657:PRO:C	1:H:658:ILE:HD13	1.99	0.88
1:J:496:GLU:CD	1:J:496:GLU:H	1.80	0.88
1:D:657:PRO:C	1:D:658:ILE:HD13	1.99	0.88
1:E:401:LEU:HD23	1:E:402:ALA:HB2	1.55	0.88
1:H:366:SER:C	1:H:367:THR:CG2	2.42	0.88
1:L:657:PRO:C	1:L:658:ILE:HD13	1.99	0.88
1:I:401:LEU:HD23	1:I:402:ALA:HB2	1.55	0.88
1:I:661:LEU:HG	1:J:456:ASN:HB3	1.55	0.88
1:C:401:LEU:HD23	1:C:402:ALA:HB2	1.56	0.88
1:F:657:PRO:C	1:F:658:ILE:HD13	1.99	0.88
1:G:631:VAL:HG21	1:H:476:ILE:HG21	1.53	0.88
1:A:340:VAL:O	1:A:345:TYR:HE1	1.55	0.87
1:G:401:LEU:HD23	1:G:402:ALA:HB2	1.56	0.87
1:J:657:PRO:C	1:J:658:ILE:HD13	1.99	0.87
1:E:629:ASP:HA	1:F:664:HIS:NE2	1.87	0.87
1:I:631:VAL:HG21	1:J:476:ILE:HG21	1.53	0.87
1:C:631:VAL:HG21	1:D:476:ILE:HG22	1.57	0.87
1:D:647:ARG:HG2	1:D:650:TYR:CD1	2.08	0.87
1:H:496:GLU:H	1:H:496:GLU:CD	1.80	0.87
1:E:635:ARG:CG	1:E:636:ASP:N	2.30	0.87
1:G:629:ASP:HA	1:H:664:HIS:NE2	1.87	0.87
1:I:340:VAL:CA	1:I:345:TYR:CE1	2.58	0.87
1:I:347:THR:OG1	1:I:348:PHE:N	2.05	0.87
1:A:519:SER:HB2	1:A:562:ASP:CG	2.00	0.87
1:A:631:VAL:HG21	1:B:476:ILE:HG21	1.53	0.87
1:A:635:ARG:HG2	1:A:636:ASP:N	1.90	0.87
1:G:660:GLN:HG3	1:G:661:LEU:N	1.90	0.87

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:647:ARG:HG2	1:J:650:TYR:CD1	2.08	0.87
1:K:340:VAL:CA	1:K:345:TYR:CE1	2.58	0.87
1:B:657:PRO:C	1:B:658:ILE:HD13	1.99	0.87
1:C:519:SER:HB2	1:C:562:ASP:CG	2.00	0.87
1:G:519:SER:HB2	1:G:562:ASP:CG	2.00	0.87
1:I:340:VAL:O	1:I:345:TYR:HE1	1.55	0.87
1:A:340:VAL:CA	1:A:345:TYR:CE1	2.58	0.87
1:D:367:THR:CG2	1:D:368:LYS:H	1.87	0.87
1:G:341:THR:HG1	1:G:344:ASP:HB2	1.39	0.87
1:I:519:SER:HB2	1:I:562:ASP:CG	2.00	0.87
1:C:340:VAL:CA	1:C:345:TYR:CE1	2.58	0.87
1:F:496:GLU:H	1:F:496:GLU:CD	1.80	0.87
1:B:366:SER:O	1:B:367:THR:HG22	1.73	0.86
1:F:647:ARG:HG2	1:F:650:TYR:CD1	2.08	0.86
1:C:660:GLN:HG3	1:C:661:LEU:N	1.90	0.86
1:H:647:ARG:HG2	1:H:650:TYR:CD1	2.08	0.86
1:G:340:VAL:CA	1:G:345:TYR:CE1	2.58	0.86
1:K:519:SER:HB2	1:K:562:ASP:CG	2.00	0.86
1:A:661:LEU:HG	1:B:456:ASN:HB3	1.55	0.86
1:I:631:VAL:HG21	1:J:476:ILE:HG22	1.57	0.86
1:K:661:LEU:HG	1:L:456:ASN:HB3	1.55	0.86
1:E:519:SER:HB2	1:E:562:ASP:CG	2.00	0.86
1:B:364:THR:HG21	1:K:401:LEU:HD11	0.87	0.86
1:K:635:ARG:HH22	1:L:661:LEU:HD12	1.37	0.86
1:A:560:SER:N	1:A:561:GLY:CA	2.39	0.86
1:K:347:THR:OG1	1:K:348:PHE:N	2.05	0.86
1:A:660:GLN:HG3	1:A:661:LEU:N	1.90	0.86
1:K:341:THR:HG1	1:K:344:ASP:HB2	1.41	0.86
1:K:631:VAL:HG21	1:L:476:ILE:HG22	1.57	0.86
1:L:664:HIS:CG	1:L:664:HIS:O	2.29	0.86
1:A:401:LEU:HD12	1:C:364:THR:CG2	2.05	0.85
1:I:353:PHE:CE1	1:I:395:TYR:CE2	2.65	0.85
1:C:661:LEU:HG	1:D:456:ASN:HB3	1.55	0.85
1:E:340:VAL:CA	1:E:345:TYR:CE1	2.58	0.85
1:K:401:LEU:HD23	1:K:402:ALA:HB2	1.55	0.85
1:L:353:PHE:HZ	1:L:395:TYR:HE2	1.21	0.85
1:E:560:SER:N	1:E:561:GLY:CA	2.39	0.85
1:E:401:LEU:HD11	1:H:364:THR:HG21	0.86	0.85
1:A:341:THR:O	1:A:345:TYR:CD1	2.30	0.85
1:A:347:THR:OG1	1:A:348:PHE:N	2.05	0.85
1:J:662:GLU:O	1:J:663:HIS:CB	2.20	0.85

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:353:PHE:HZ	1:B:395:TYR:HE2	1.21	0.85
1:C:341:THR:O	1:C:345:TYR:CD1	2.30	0.85
1:B:664:HIS:CG	1:B:664:HIS:O	2.29	0.85
1:E:353:PHE:CE1	1:E:395:TYR:CE2	2.65	0.85
1:E:660:GLN:HG3	1:E:661:LEU:N	1.90	0.85
1:J:345:TYR:O	1:J:345:TYR:CD1	2.30	0.85
1:L:345:TYR:O	1:L:345:TYR:CD1	2.30	0.85
1:B:662:GLU:O	1:B:663:HIS:CB	2.20	0.85
1:D:400:ASN:ND2	1:E:662:GLU:CD	2.34	0.85
1:G:385:THR:HA	1:G:387:VAL:HG22	1.59	0.85
1:I:660:GLN:HG3	1:I:661:LEU:N	1.90	0.85
1:J:353:PHE:HZ	1:J:395:TYR:HE2	1.21	0.85
1:D:345:TYR:O	1:D:345:TYR:CD1	2.30	0.85
1:H:345:TYR:CD1	1:H:345:TYR:O	2.30	0.85
1:I:635:ARG:HG2	1:I:636:ASP:N	1.90	0.85
1:D:662:GLU:O	1:D:663:HIS:CB	2.20	0.84
1:G:631:VAL:HG21	1:H:476:ILE:HG22	1.57	0.84
1:E:341:THR:O	1:E:345:TYR:CD1	2.30	0.84
1:E:385:THR:HA	1:E:387:VAL:HG22	1.59	0.84
1:F:345:TYR:CD1	1:F:345:TYR:O	2.30	0.84
1:I:341:THR:O	1:I:345:TYR:CD1	2.30	0.84
1:I:385:THR:HA	1:I:387:VAL:HG22	1.59	0.84
1:C:353:PHE:CE1	1:C:395:TYR:CE2	2.65	0.84
1:C:560:SER:N	1:C:561:GLY:CA	2.39	0.84
1:G:353:PHE:CE1	1:G:395:TYR:CE2	2.65	0.84
1:K:341:THR:O	1:K:345:TYR:CD1	2.30	0.84
1:L:367:THR:CG2	1:L:368:LYS:H	1.87	0.84
1:B:345:TYR:O	1:B:345:TYR:CD1	2.30	0.84
1:F:664:HIS:O	1:F:664:HIS:CG	2.29	0.84
1:E:385:THR:HG1	1:E:387:VAL:HG23	1.37	0.84
1:E:403:PRO:CB	1:H:366:SER:CA	2.39	0.84
1:H:664:HIS:CG	1:H:664:HIS:O	2.29	0.84
1:K:660:GLN:HG3	1:K:661:LEU:N	1.90	0.84
1:B:567:GLU:HA	1:B:570:GLN:OE1	1.78	0.84
1:C:385:THR:HG1	1:C:387:VAL:HG23	1.39	0.84
1:D:664:HIS:CG	1:D:664:HIS:O	2.29	0.84
1:I:340:VAL:CA	1:I:345:TYR:HE1	1.90	0.84
1:C:347:THR:OG1	1:C:348:PHE:N	2.05	0.84
1:C:635:ARG:HG2	1:C:636:ASP:N	1.90	0.84
1:A:631:VAL:HG21	1:B:476:ILE:HG22	1.57	0.84
1:F:567:GLU:HA	1:F:570:GLN:OE1	1.78	0.84

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:662:GLU:O	1:F:663:HIS:CB	2.20	0.84
1:J:366:SER:C	1:J:367:THR:CG2	2.42	0.84
1:K:353:PHE:CE1	1:K:395:TYR:CE2	2.65	0.84
1:A:353:PHE:CE1	1:A:395:TYR:CE2	2.65	0.84
1:A:401:LEU:HD11	1:C:364:THR:CG2	2.01	0.84
1:B:543:THR:HG22	1:B:577:PHE:HB3	1.60	0.84
1:H:353:PHE:HZ	1:H:395:TYR:HE2	1.21	0.84
1:H:543:THR:HG22	1:H:577:PHE:HB3	1.60	0.84
1:D:353:PHE:HZ	1:D:395:TYR:HE2	1.21	0.83
1:I:557:PRO:HB3	1:I:587:ASP:HA	1.60	0.83
1:K:340:VAL:CA	1:K:345:TYR:HE1	1.90	0.83
1:G:341:THR:O	1:G:345:TYR:CD1	2.30	0.83
1:L:662:GLU:O	1:L:663:HIS:CB	2.20	0.83
1:D:366:SER:C	1:D:367:THR:CG2	2.42	0.83
1:F:582:ASN:HD21	1:F:586:ARG:HB2	1.44	0.83
1:G:347:THR:OG1	1:G:348:PHE:N	2.05	0.83
1:H:567:GLU:HA	1:H:570:GLN:OE1	1.78	0.83
1:I:532:GLU:N	1:I:532:GLU:CD	2.36	0.83
1:B:582:ASN:HD21	1:B:586:ARG:HB2	1.43	0.83
1:C:340:VAL:CA	1:C:345:TYR:HE1	1.90	0.83
1:D:582:ASN:HD21	1:D:586:ARG:HB2	1.44	0.83
1:E:557:PRO:HB3	1:E:587:ASP:HA	1.60	0.83
1:J:664:HIS:CG	1:J:664:HIS:O	2.29	0.83
1:F:353:PHE:HZ	1:F:395:TYR:HE2	1.21	0.83
1:F:543:THR:HG22	1:F:577:PHE:HB3	1.60	0.83
1:A:532:GLU:N	1:A:532:GLU:CD	2.37	0.83
1:E:345:TYR:N	1:E:345:TYR:HD1	1.77	0.83
1:G:532:GLU:CD	1:G:532:GLU:N	2.36	0.83
1:K:635:ARG:HG2	1:K:636:ASP:N	1.90	0.83
1:L:543:THR:HG22	1:L:577:PHE:HB3	1.60	0.83
1:E:340:VAL:CA	1:E:345:TYR:HE1	1.90	0.83
1:E:631:VAL:HG21	1:F:476:ILE:HG22	1.57	0.83
1:G:345:TYR:N	1:G:345:TYR:HD1	1.77	0.83
1:E:341:THR:HG1	1:E:344:ASP:HB2	1.41	0.83
1:L:567:GLU:HA	1:L:570:GLN:OE1	1.78	0.83
1:A:341:THR:HG1	1:A:344:ASP:HB2	1.42	0.83
1:G:560:SER:N	1:G:561:GLY:CA	2.39	0.83
1:H:367:THR:CG2	1:H:368:LYS:H	1.87	0.83
1:J:345:TYR:CE1	1:J:349:VAL:HG23	2.14	0.83
1:K:532:GLU:CD	1:K:532:GLU:N	2.36	0.83
1:C:532:GLU:CD	1:C:532:GLU:N	2.37	0.82

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:567:GLU:HA	1:J:570:GLN:OE1	1.78	0.82
1:A:345:TYR:N	1:A:345:TYR:HD1	1.77	0.82
1:B:379:LYS:HG3	1:B:380:SER:N	1.94	0.82
1:D:345:TYR:CE1	1:D:349:VAL:HG23	2.14	0.82
1:E:347:THR:OG1	1:E:348:PHE:N	2.05	0.82
1:G:340:VAL:CA	1:G:345:TYR:HE1	1.90	0.82
1:G:557:PRO:HB3	1:G:587:ASP:HA	1.60	0.82
1:H:503:ASN:ND2	1:H:633:PHE:H	1.77	0.82
1:K:385:THR:HA	1:K:387:VAL:HG22	1.59	0.82
1:D:379:LYS:HG3	1:D:380:SER:N	1.94	0.82
1:H:582:ASN:HD21	1:H:586:ARG:HB2	1.44	0.82
1:J:582:ASN:HD21	1:J:586:ARG:HB2	1.43	0.82
1:J:659:SER:O	1:J:660:GLN:HG3	1.80	0.82
1:K:661:LEU:HD13	1:K:662:GLU:CA	2.10	0.82
1:B:345:TYR:CE1	1:B:349:VAL:HG23	2.14	0.82
1:L:582:ASN:HD21	1:L:586:ARG:HB2	1.43	0.82
1:I:661:LEU:HD13	1:I:662:GLU:CA	2.10	0.82
1:A:385:THR:HA	1:A:387:VAL:HG22	1.59	0.82
1:B:366:SER:C	1:B:367:THR:CG2	2.42	0.82
1:D:646:LEU:HD23	1:D:646:LEU:C	2.05	0.82
1:F:659:SER:O	1:F:660:GLN:HG3	1.80	0.82
1:A:340:VAL:CA	1:A:345:TYR:HE1	1.90	0.82
1:A:401:LEU:HD23	1:A:402:ALA:HB2	1.55	0.82
1:D:567:GLU:HA	1:D:570:GLN:OE1	1.78	0.82
1:F:345:TYR:CE1	1:F:349:VAL:HG23	2.14	0.82
1:I:396:LEU:HD22	1:I:408:ILE:HD13	1.62	0.82
1:F:646:LEU:HD23	1:F:646:LEU:C	2.05	0.82
1:C:557:PRO:HB3	1:C:587:ASP:HA	1.60	0.82
1:D:543:THR:HG22	1:D:577:PHE:HB3	1.60	0.82
1:G:396:LEU:HD22	1:G:408:ILE:HD13	1.62	0.82
1:L:345:TYR:CE1	1:L:349:VAL:HG23	2.14	0.82
1:A:661:LEU:HD13	1:A:662:GLU:CA	2.10	0.81
1:J:503:ASN:ND2	1:J:633:PHE:H	1.77	0.81
1:C:635:ARG:NH2	1:D:661:LEU:HD12	1.95	0.81
1:D:503:ASN:ND2	1:D:633:PHE:H	1.77	0.81
1:F:379:LYS:HG3	1:F:380:SER:N	1.94	0.81
1:G:661:LEU:HD13	1:G:662:GLU:CA	2.10	0.81
1:H:646:LEU:HD23	1:H:646:LEU:C	2.05	0.81
1:I:341:THR:HG1	1:I:344:ASP:HB2	1.43	0.81
1:J:543:THR:HG22	1:J:577:PHE:HB3	1.60	0.81
1:C:340:VAL:HA	1:C:345:TYR:CE1	2.15	0.81

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:385:THR:HA	1:C:387:VAL:HG22	1.59	0.81
1:D:399:TYR:CA	1:E:664:HIS:CG	2.61	0.81
1:D:659:SER:O	1:D:660:GLN:HG3	1.80	0.81
1:G:340:VAL:HA	1:G:345:TYR:CE1	2.15	0.81
1:H:659:SER:O	1:H:660:GLN:HG3	1.80	0.81
1:L:595:ILE:H	1:L:595:ILE:HD12	1.45	0.81
1:H:345:TYR:CE1	1:H:349:VAL:HG23	2.14	0.81
1:E:532:GLU:CD	1:E:532:GLU:N	2.37	0.81
1:I:340:VAL:HA	1:I:345:TYR:CE1	2.15	0.81
1:A:635:ARG:NH2	1:B:661:LEU:HD12	1.95	0.81
1:E:403:PRO:CA	1:H:366:SER:CB	2.42	0.81
1:I:560:SER:N	1:I:561:GLY:CA	2.39	0.81
1:A:557:PRO:HB3	1:A:587:ASP:HA	1.60	0.81
1:C:345:TYR:N	1:C:345:TYR:HD1	1.77	0.81
1:J:646:LEU:HD23	1:J:646:LEU:C	2.05	0.81
1:K:396:LEU:HD22	1:K:408:ILE:HD13	1.62	0.81
1:L:646:LEU:HD23	1:L:646:LEU:C	2.05	0.81
1:I:345:TYR:N	1:I:345:TYR:HD1	1.77	0.81
1:L:379:LYS:HG3	1:L:380:SER:N	1.94	0.81
1:B:646:LEU:HD23	1:B:646:LEU:C	2.05	0.81
1:E:396:LEU:HD22	1:E:408:ILE:HD13	1.62	0.81
1:F:595:ILE:HD12	1:F:595:ILE:H	1.45	0.81
1:K:557:PRO:HB3	1:K:587:ASP:HA	1.60	0.81
1:L:659:SER:O	1:L:660:GLN:CG	2.29	0.81
1:A:340:VAL:HA	1:A:345:TYR:CE1	2.15	0.80
1:F:659:SER:O	1:F:660:GLN:CG	2.29	0.80
1:B:366:SER:O	1:B:367:THR:CG2	2.29	0.80
1:B:595:ILE:H	1:B:595:ILE:HD12	1.45	0.80
1:B:659:SER:O	1:B:660:GLN:HG3	1.80	0.80
1:C:661:LEU:HD13	1:C:662:GLU:CA	2.10	0.80
1:E:403:PRO:CB	1:H:366:SER:HA	2.10	0.80
1:E:635:ARG:NH2	1:F:661:LEU:HD12	1.95	0.80
1:K:560:SER:N	1:K:561:GLY:CA	2.39	0.80
1:D:659:SER:O	1:D:660:GLN:CG	2.29	0.80
1:G:388:GLN:O	1:G:389:ARG:CG	2.30	0.80
1:J:595:ILE:H	1:J:595:ILE:HD12	1.45	0.80
1:A:525:ASN:OD1	1:A:526:PRO:CD	2.30	0.80
1:E:661:LEU:HD13	1:E:662:GLU:CA	2.10	0.80
1:K:525:ASN:OD1	1:K:526:PRO:CD	2.30	0.80
1:J:659:SER:O	1:J:660:GLN:CG	2.29	0.80
1:L:366:SER:O	1:L:367:THR:CG2	2.29	0.80

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:659:SER:O	1:H:660:GLN:CG	2.29	0.80
1:J:567:GLU:O	1:J:568:ASN:CB	2.30	0.80
1:J:661:LEU:O	1:J:661:LEU:CD2	2.29	0.80
1:K:340:VAL:HA	1:K:345:TYR:CE1	2.15	0.80
1:K:388:GLN:O	1:K:389:ARG:CG	2.30	0.80
1:L:659:SER:O	1:L:660:GLN:HG3	1.80	0.80
1:B:366:SER:O	1:B:367:THR:CB	2.30	0.80
1:C:385:THR:CG2	1:C:389:ARG:CA	2.60	0.80
1:E:353:PHE:CE1	1:E:395:TYR:CD2	2.70	0.80
1:I:525:ASN:OD1	1:I:526:PRO:CD	2.30	0.80
1:L:661:LEU:O	1:L:661:LEU:CD2	2.29	0.80
1:A:629:ASP:HA	1:B:664:HIS:HE1	1.47	0.80
1:B:367:THR:CG2	1:B:368:LYS:H	1.87	0.80
1:D:366:SER:O	1:D:367:THR:CB	2.30	0.80
1:D:567:GLU:O	1:D:568:ASN:CB	2.30	0.80
1:E:385:THR:CG2	1:E:389:ARG:CA	2.60	0.80
1:G:635:ARG:NH2	1:H:661:LEU:HD12	1.95	0.80
1:H:379:LYS:HG3	1:H:380:SER:N	1.94	0.80
1:I:353:PHE:CE1	1:I:395:TYR:CD2	2.70	0.80
1:I:635:ARG:NH2	1:J:661:LEU:HD12	1.95	0.80
1:J:379:LYS:HG3	1:J:380:SER:N	1.94	0.80
1:K:353:PHE:CE1	1:K:395:TYR:CD2	2.70	0.80
1:L:503:ASN:HD21	1:L:633:PHE:N	1.79	0.80
1:A:385:THR:CG2	1:A:389:ARG:CA	2.60	0.80
1:A:396:LEU:HD22	1:A:408:ILE:HD13	1.62	0.80
1:B:659:SER:O	1:B:660:GLN:CG	2.29	0.80
1:D:595:ILE:H	1:D:595:ILE:HD12	1.45	0.80
1:K:635:ARG:NH2	1:L:661:LEU:HD12	1.96	0.80
1:C:396:LEU:HD22	1:C:408:ILE:HD13	1.62	0.80
1:G:353:PHE:CE1	1:G:395:TYR:CD2	2.70	0.80
1:G:525:ASN:OD1	1:G:526:PRO:CD	2.30	0.80
1:H:661:LEU:O	1:H:661:LEU:CD2	2.29	0.80
1:E:340:VAL:HA	1:E:345:TYR:CE1	2.15	0.79
1:F:567:GLU:O	1:F:568:ASN:CB	2.30	0.79
1:E:388:GLN:O	1:E:389:ARG:CB	2.30	0.79
1:E:388:GLN:O	1:E:389:ARG:CG	2.30	0.79
1:H:595:ILE:H	1:H:595:ILE:HD12	1.46	0.79
1:L:503:ASN:ND2	1:L:633:PHE:H	1.77	0.79
1:F:661:LEU:O	1:F:661:LEU:CD1	2.29	0.79
1:I:440:ILE:HD11	1:I:653:ILE:HD13	1.64	0.79
1:A:353:PHE:CE1	1:A:395:TYR:CD2	2.70	0.79

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:661:LEU:O	1:B:661:LEU:CD2	2.29	0.79
1:I:388:GLN:O	1:I:389:ARG:CG	2.30	0.79
1:A:388:GLN:O	1:A:389:ARG:CG	2.30	0.79
1:D:661:LEU:O	1:D:661:LEU:CD2	2.29	0.79
1:F:366:SER:O	1:F:367:THR:CG2	2.29	0.79
1:F:503:ASN:HD21	1:F:633:PHE:N	1.79	0.79
1:L:567:GLU:O	1:L:568:ASN:CB	2.30	0.79
1:C:353:PHE:CE1	1:C:395:TYR:CD2	2.70	0.79
1:D:366:SER:O	1:D:367:THR:CG2	2.29	0.79
1:D:503:ASN:HD21	1:D:633:PHE:N	1.79	0.79
1:J:366:SER:O	1:J:367:THR:CG2	2.29	0.79
1:C:385:THR:HG21	1:C:389:ARG:HA	1.64	0.79
1:C:388:GLN:O	1:C:389:ARG:CG	2.30	0.79
1:F:661:LEU:O	1:F:661:LEU:CD2	2.30	0.79
1:H:496:GLU:CD	1:H:496:GLU:N	2.41	0.79
1:K:345:TYR:N	1:K:345:TYR:HD1	1.77	0.79
1:K:388:GLN:O	1:K:389:ARG:CB	2.30	0.79
1:E:440:ILE:HD11	1:E:653:ILE:HD13	1.64	0.79
1:G:361:GLN:HG2	1:G:455:PHE:CG	2.18	0.79
1:G:385:THR:HG21	1:G:389:ARG:HA	1.64	0.79
1:H:366:SER:O	1:H:367:THR:CB	2.30	0.79
1:I:388:GLN:O	1:I:389:ARG:CB	2.30	0.79
1:J:496:GLU:CD	1:J:496:GLU:N	2.41	0.79
1:L:408:ILE:O	1:L:408:ILE:CD1	2.30	0.79
1:A:385:THR:CA	1:A:387:VAL:HG22	2.13	0.79
1:B:567:GLU:O	1:B:568:ASN:CB	2.30	0.79
1:B:647:ARG:HD2	1:B:650:TYR:CE1	2.19	0.79
1:K:385:THR:CA	1:K:387:VAL:HG22	2.13	0.79
1:L:366:SER:C	1:L:367:THR:CG2	2.42	0.79
1:D:374:ILE:HG12	1:D:408:ILE:HA	1.65	0.78
1:E:403:PRO:CG	1:H:366:SER:HA	2.13	0.78
1:I:385:THR:CA	1:I:387:VAL:HG22	2.13	0.78
1:J:503:ASN:HD21	1:J:633:PHE:N	1.79	0.78
1:L:496:GLU:CD	1:L:496:GLU:N	2.41	0.78
1:A:385:THR:HG21	1:A:389:ARG:HA	1.64	0.78
1:C:388:GLN:O	1:C:389:ARG:CB	2.30	0.78
1:C:525:ASN:OD1	1:C:526:PRO:CD	2.30	0.78
1:G:388:GLN:O	1:G:389:ARG:CB	2.30	0.78
1:A:457:SER:H	1:A:634:THR:CG2	1.97	0.78
1:G:440:ILE:HD11	1:G:653:ILE:HD13	1.64	0.78
1:H:503:ASN:HD21	1:H:633:PHE:N	1.79	0.78

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:661:LEU:O	1:L:661:LEU:CD1	2.29	0.78
1:B:374:ILE:HG12	1:B:408:ILE:HA	1.65	0.78
1:C:361:GLN:HG2	1:C:455:PHE:CG	2.19	0.78
1:C:457:SER:H	1:C:634:THR:CG2	1.97	0.78
1:I:361:GLN:HG2	1:I:455:PHE:CG	2.19	0.78
1:J:647:ARG:HD2	1:J:650:TYR:CE1	2.19	0.78
1:B:503:ASN:ND2	1:B:633:PHE:H	1.77	0.78
1:E:361:GLN:HG2	1:E:455:PHE:CG	2.19	0.78
1:F:503:ASN:ND2	1:F:633:PHE:H	1.77	0.78
1:H:374:ILE:HG12	1:H:408:ILE:HA	1.65	0.78
1:I:385:THR:HG21	1:I:389:ARG:HA	1.64	0.78
1:D:408:ILE:O	1:D:408:ILE:CD1	2.30	0.78
1:E:525:ASN:OD1	1:E:526:PRO:CD	2.30	0.78
1:E:629:ASP:HA	1:F:664:HIS:HE1	1.47	0.78
1:J:569:ILE:O	1:J:569:ILE:CG2	2.30	0.78
1:B:496:GLU:CD	1:B:496:GLU:N	2.41	0.78
1:A:388:GLN:O	1:A:389:ARG:CB	2.30	0.78
1:F:374:ILE:HG12	1:F:408:ILE:HA	1.65	0.78
1:G:524:VAL:HG22	1:G:529:GLY:O	1.84	0.78
1:H:567:GLU:O	1:H:568:ASN:CB	2.30	0.78
1:I:353:PHE:CE2	1:I:395:TYR:HD2	2.02	0.78
1:K:361:GLN:HG2	1:K:455:PHE:CG	2.18	0.78
1:C:440:ILE:HD11	1:C:653:ILE:HD13	1.64	0.78
1:H:647:ARG:HD2	1:H:650:TYR:CE1	2.19	0.78
1:J:374:ILE:HG12	1:J:408:ILE:HA	1.65	0.78
1:K:353:PHE:CE2	1:K:395:TYR:HD2	2.02	0.78
1:C:385:THR:CA	1:C:387:VAL:HG22	2.14	0.78
1:E:345:TYR:CD1	1:E:345:TYR:N	2.51	0.78
1:J:367:THR:CG2	1:J:368:LYS:H	1.88	0.78
1:L:374:ILE:HG12	1:L:408:ILE:HA	1.65	0.78
1:D:647:ARG:HD2	1:D:650:TYR:CE1	2.19	0.77
1:L:647:ARG:HD2	1:L:650:TYR:CE1	2.19	0.77
1:G:385:THR:CA	1:G:387:VAL:HG22	2.13	0.77
1:J:366:SER:O	1:J:367:THR:CB	2.30	0.77
1:K:440:ILE:HD11	1:K:653:ILE:HD13	1.64	0.77
1:D:496:GLU:CD	1:D:496:GLU:N	2.41	0.77
1:E:385:THR:HG21	1:E:389:ARG:HA	1.64	0.77
1:H:366:SER:O	1:H:367:THR:CG2	2.29	0.77
1:A:361:GLN:HG2	1:A:455:PHE:CG	2.19	0.77
1:A:403:PRO:HB3	1:C:366:SER:OG	1.85	0.77
1:A:440:ILE:HD11	1:A:653:ILE:HD13	1.64	0.77

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:385:THR:CA	1:E:387:VAL:HG22	2.14	0.77
1:E:353:PHE:CE2	1:E:395:TYR:HD2	2.02	0.77
1:F:647:ARG:HD2	1:F:650:TYR:CE1	2.18	0.77
1:H:403:PRO:O	1:H:405:THR:HG22	1.85	0.77
1:I:524:VAL:HG22	1:I:529:GLY:O	1.84	0.77
1:A:340:VAL:CA	1:A:345:TYR:OH	2.30	0.77
1:B:661:LEU:O	1:B:661:LEU:CD1	2.29	0.77
1:A:353:PHE:CE2	1:A:395:TYR:HD2	2.02	0.77
1:A:524:VAL:HG22	1:A:529:GLY:O	1.84	0.77
1:C:355:SER:OG	1:C:356:ILE:HG23	1.85	0.77
1:D:367:THR:O	1:D:369:PRO:CD	2.30	0.77
1:H:661:LEU:O	1:H:661:LEU:CD1	2.29	0.77
1:L:569:ILE:O	1:L:569:ILE:CG2	2.30	0.77
1:E:355:SER:OG	1:E:356:ILE:HG23	1.85	0.77
1:E:524:VAL:HG22	1:E:529:GLY:O	1.84	0.77
1:K:524:VAL:HG22	1:K:529:GLY:O	1.84	0.77
1:G:353:PHE:CE2	1:G:395:TYR:HD2	2.02	0.77
1:L:403:PRO:O	1:L:405:THR:HG22	1.85	0.77
1:A:345:TYR:CD1	1:A:345:TYR:N	2.51	0.76
1:C:353:PHE:CE2	1:C:395:TYR:HD2	2.02	0.76
1:F:367:THR:CG2	1:F:368:LYS:H	1.88	0.76
1:G:355:SER:OG	1:G:356:ILE:HG23	1.85	0.76
1:K:457:SER:H	1:K:634:THR:CG2	1.97	0.76
1:B:366:SER:CB	1:K:403:PRO:CA	2.44	0.76
1:B:367:THR:O	1:B:369:PRO:CD	2.30	0.76
1:D:408:ILE:O	1:D:409:ILE:CG2	2.34	0.76
1:J:403:PRO:O	1:J:405:THR:HG22	1.85	0.76
1:B:366:SER:HA	1:K:403:PRO:CG	2.15	0.76
1:C:629:ASP:HA	1:D:664:HIS:HE1	1.47	0.76
1:J:367:THR:O	1:J:369:PRO:CD	2.30	0.76
1:L:408:ILE:O	1:L:409:ILE:CG2	2.34	0.76
1:B:408:ILE:O	1:B:409:ILE:CG2	2.34	0.76
1:C:524:VAL:HG22	1:C:529:GLY:O	1.84	0.76
1:D:408:ILE:HD12	1:D:409:ILE:N	2.01	0.76
1:D:661:LEU:HD13	1:D:662:GLU:N	2.01	0.76
1:E:457:SER:H	1:E:634:THR:CG2	1.97	0.76
1:F:661:LEU:HD13	1:F:662:GLU:N	2.01	0.76
1:L:345:TYR:HE1	1:L:349:VAL:CG2	1.99	0.76
1:L:366:SER:O	1:L:367:THR:CB	2.30	0.76
1:A:355:SER:OG	1:A:356:ILE:HG23	1.85	0.76
1:B:408:ILE:HD12	1:B:409:ILE:N	2.01	0.76

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:408:ILE:O	1:F:409:ILE:CG2	2.34	0.76
1:F:496:GLU:CD	1:F:496:GLU:N	2.41	0.76
1:G:345:TYR:CD1	1:G:345:TYR:N	2.51	0.76
1:G:401:LEU:O	1:G:402:ALA:HB3	1.84	0.76
1:A:401:LEU:O	1:A:402:ALA:HB3	1.84	0.76
1:B:345:TYR:HE1	1:B:349:VAL:CG2	1.99	0.76
1:B:427:ASN:ND2	1:B:427:ASN:H	1.84	0.76
1:C:416:ILE:HG13	1:C:448:TYR:OH	1.86	0.76
1:F:403:PRO:O	1:F:405:THR:HG22	1.85	0.76
1:H:342:ALA:C	1:H:346:ASP:OD2	2.29	0.76
1:A:560:SER:N	1:A:561:GLY:HA3	2.01	0.76
1:B:342:ALA:O	1:B:346:ASP:CG	2.29	0.76
1:C:340:VAL:CA	1:C:345:TYR:OH	2.30	0.76
1:F:342:ALA:C	1:F:346:ASP:OD2	2.29	0.76
1:J:408:ILE:O	1:J:408:ILE:CD1	2.30	0.76
1:K:385:THR:HG21	1:K:389:ARG:HA	1.64	0.76
1:L:342:ALA:C	1:L:346:ASP:OD2	2.29	0.76
1:D:568:ASN:CG	1:D:569:ILE:H	1.94	0.75
1:D:661:LEU:O	1:D:661:LEU:CD1	2.29	0.75
1:J:408:ILE:HD12	1:J:409:ILE:N	2.01	0.75
1:K:401:LEU:O	1:K:402:ALA:HB3	1.84	0.75
1:A:587:ASP:OD2	1:A:587:ASP:C	2.30	0.75
1:B:366:SER:HA	1:K:403:PRO:CB	2.11	0.75
1:F:344:ASP:C	1:F:344:ASP:OD1	2.29	0.75
1:H:408:ILE:O	1:H:409:ILE:CG2	2.34	0.75
1:K:416:ILE:HG13	1:K:448:TYR:OH	1.86	0.75
1:A:416:ILE:HG13	1:A:448:TYR:OH	1.86	0.75
1:B:403:PRO:O	1:B:405:THR:HG22	1.85	0.75
1:B:644:ASN:O	1:B:644:ASN:ND2	2.20	0.75
1:F:342:ALA:O	1:F:346:ASP:CG	2.29	0.75
1:F:366:SER:O	1:F:367:THR:CB	2.30	0.75
1:H:342:ALA:O	1:H:346:ASP:CG	2.30	0.75
1:I:355:SER:OG	1:I:356:ILE:HG23	1.85	0.75
1:J:408:ILE:O	1:J:409:ILE:CG2	2.34	0.75
1:K:355:SER:OG	1:K:356:ILE:HG23	1.85	0.75
1:L:344:ASP:C	1:L:344:ASP:OD1	2.29	0.75
1:B:344:ASP:C	1:B:344:ASP:OD1	2.29	0.75
1:D:398:ASP:OD2	1:E:663:HIS:HB3	1.86	0.75
1:E:340:VAL:CA	1:E:345:TYR:OH	2.30	0.75
1:K:587:ASP:C	1:K:587:ASP:OD2	2.30	0.75
1:A:587:ASP:CG	1:A:587:ASP:O	2.30	0.75

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:400:ASN:HB3	1:E:662:GLU:HA	1.66	0.75
1:H:344:ASP:C	1:H:344:ASP:OD1	2.29	0.75
1:H:345:TYR:HE1	1:H:349:VAL:CG2	1.99	0.75
1:I:526:PRO:O	1:I:528:THR:HG22	1.87	0.75
1:K:340:VAL:CA	1:K:345:TYR:OH	2.30	0.75
1:L:367:THR:O	1:L:369:PRO:CD	2.30	0.75
1:C:401:LEU:O	1:C:402:ALA:HB3	1.85	0.75
1:C:526:PRO:O	1:C:528:THR:HG22	1.87	0.75
1:D:427:ASN:ND2	1:D:427:ASN:H	1.84	0.75
1:E:416:ILE:HG13	1:E:448:TYR:OH	1.86	0.75
1:G:416:ILE:HG13	1:G:448:TYR:OH	1.86	0.75
1:H:367:THR:O	1:H:369:PRO:CD	2.30	0.75
1:H:661:LEU:HD13	1:H:662:GLU:N	2.01	0.75
1:J:342:ALA:C	1:J:346:ASP:OD2	2.29	0.75
1:J:345:TYR:HE1	1:J:349:VAL:CG2	1.99	0.75
1:J:659:SER:O	1:J:660:GLN:CD	2.30	0.75
1:K:340:VAL:O	1:K:345:TYR:CE1	2.37	0.75
1:K:526:PRO:O	1:K:528:THR:HG22	1.87	0.75
1:L:342:ALA:O	1:L:346:ASP:CG	2.29	0.75
1:L:568:ASN:CG	1:L:569:ILE:H	1.94	0.75
1:L:661:LEU:HD13	1:L:662:GLU:N	2.01	0.75
1:D:345:TYR:HE1	1:D:349:VAL:CG2	1.99	0.75
1:F:427:ASN:ND2	1:F:427:ASN:H	1.84	0.75
1:F:644:ASN:ND2	1:F:644:ASN:O	2.20	0.75
1:G:526:PRO:O	1:G:528:THR:HG22	1.87	0.75
1:I:401:LEU:O	1:I:402:ALA:HB3	1.84	0.75
1:J:342:ALA:O	1:J:346:ASP:CG	2.29	0.75
1:K:345:TYR:CD1	1:K:345:TYR:N	2.52	0.75
1:B:568:ASN:CG	1:B:569:ILE:H	1.94	0.75
1:C:341:THR:HG1	1:C:344:ASP:HB2	1.48	0.75
1:E:347:THR:O	1:E:349:VAL:N	2.20	0.75
1:E:401:LEU:O	1:E:402:ALA:HB3	1.85	0.75
1:H:408:ILE:HD12	1:H:409:ILE:N	2.01	0.75
1:L:408:ILE:HD12	1:L:409:ILE:N	2.01	0.75
1:D:344:ASP:OD1	1:D:344:ASP:C	2.29	0.75
1:D:403:PRO:O	1:D:405:THR:HG22	1.85	0.75
1:F:345:TYR:HE1	1:F:349:VAL:CG2	1.99	0.75
1:J:568:ASN:CG	1:J:569:ILE:H	1.94	0.75
1:L:659:SER:O	1:L:660:GLN:CD	2.30	0.75
1:B:342:ALA:C	1:B:346:ASP:OD2	2.29	0.74
1:B:569:ILE:O	1:B:569:ILE:CG2	2.30	0.74

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:342:ALA:C	1:D:346:ASP:OD2	2.29	0.74
1:E:587:ASP:C	1:E:587:ASP:OD2	2.30	0.74
1:E:587:ASP:CG	1:E:587:ASP:O	2.30	0.74
1:I:340:VAL:O	1:I:345:TYR:CE1	2.37	0.74
1:J:344:ASP:C	1:J:344:ASP:OD1	2.29	0.74
1:L:427:ASN:ND2	1:L:427:ASN:H	1.84	0.74
1:L:644:ASN:O	1:L:644:ASN:ND2	2.20	0.74
1:A:526:PRO:O	1:A:528:THR:HG22	1.87	0.74
1:G:347:THR:O	1:G:349:VAL:N	2.20	0.74
1:H:659:SER:O	1:H:660:GLN:CD	2.30	0.74
1:E:526:PRO:O	1:E:528:THR:HG22	1.87	0.74
1:H:567:GLU:O	1:H:568:ASN:CG	2.30	0.74
1:L:567:GLU:O	1:L:568:ASN:CG	2.30	0.74
1:C:587:ASP:C	1:C:587:ASP:OD2	2.30	0.74
1:F:408:ILE:HD12	1:F:409:ILE:N	2.01	0.74
1:G:385:THR:HG21	1:G:389:ARG:N	2.01	0.74
1:J:661:LEU:O	1:J:661:LEU:CD1	2.29	0.74
1:B:567:GLU:O	1:B:568:ASN:CG	2.30	0.74
1:D:342:ALA:O	1:D:346:ASP:CG	2.29	0.74
1:F:567:GLU:O	1:F:568:ASN:CG	2.30	0.74
1:G:457:SER:H	1:G:634:THR:CG2	1.97	0.74
1:G:629:ASP:HA	1:H:664:HIS:HE1	1.47	0.74
1:B:661:LEU:HD13	1:B:662:GLU:N	2.01	0.74
1:D:644:ASN:O	1:D:644:ASN:ND2	2.20	0.74
1:I:416:ILE:HG13	1:I:448:TYR:OH	1.86	0.74
1:J:345:TYR:CD1	1:J:345:TYR:C	2.65	0.74
1:A:347:THR:O	1:A:349:VAL:N	2.20	0.74
1:B:345:TYR:CD1	1:B:345:TYR:C	2.65	0.74
1:C:347:THR:O	1:C:349:VAL:N	2.20	0.74
1:D:398:ASP:C	1:E:663:HIS:HB3	2.13	0.74
1:D:437:GLU:O	1:D:441:ILE:HG22	1.88	0.74
1:H:345:TYR:CD1	1:H:345:TYR:C	2.65	0.74
1:I:457:SER:H	1:I:634:THR:CG2	1.97	0.74
1:D:659:SER:O	1:D:660:GLN:CD	2.30	0.74
1:K:587:ASP:O	1:K:587:ASP:CG	2.30	0.74
1:B:659:SER:O	1:B:660:GLN:CD	2.30	0.74
1:D:345:TYR:CD1	1:D:345:TYR:C	2.65	0.74
1:F:437:GLU:O	1:F:441:ILE:HG22	1.88	0.74
1:F:568:ASN:CG	1:F:569:ILE:H	1.94	0.74
1:I:345:TYR:CD1	1:I:345:TYR:N	2.51	0.74
1:J:567:GLU:O	1:J:568:ASN:CG	2.30	0.74

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:503:ASN:HD21	1:B:633:PHE:N	1.79	0.73
1:F:345:TYR:CD1	1:F:345:TYR:C	2.65	0.73
1:H:437:GLU:O	1:H:441:ILE:HG22	1.88	0.73
1:I:587:ASP:CG	1:I:587:ASP:O	2.30	0.73
1:J:661:LEU:HD13	1:J:662:GLU:N	2.01	0.73
1:L:437:GLU:O	1:L:441:ILE:HG22	1.88	0.73
1:H:568:ASN:CG	1:H:569:ILE:H	1.94	0.73
1:I:347:THR:O	1:I:349:VAL:N	2.20	0.73
1:K:347:THR:O	1:K:349:VAL:N	2.20	0.73
1:F:659:SER:O	1:F:660:GLN:CD	2.30	0.73
1:J:437:GLU:O	1:J:441:ILE:HG22	1.88	0.73
1:E:635:ARG:HG2	1:E:636:ASP:N	1.90	0.73
1:G:587:ASP:OD2	1:G:587:ASP:C	2.30	0.73
1:L:345:TYR:CD1	1:L:345:TYR:C	2.65	0.73
1:D:567:GLU:O	1:D:568:ASN:CG	2.30	0.73
1:G:665:HIS:CE1	1:H:342:ALA:CB	2.71	0.73
1:I:665:HIS:CE1	1:J:342:ALA:CB	2.71	0.73
1:K:629:ASP:HB3	1:K:631:VAL:O	1.89	0.73
1:C:587:ASP:O	1:C:587:ASP:CG	2.30	0.73
1:E:665:HIS:CE1	1:F:342:ALA:CB	2.71	0.73
1:G:587:ASP:CG	1:G:587:ASP:O	2.30	0.73
1:H:408:ILE:O	1:H:408:ILE:CD1	2.30	0.73
1:H:644:ASN:O	1:H:644:ASN:ND2	2.20	0.73
1:I:340:VAL:CA	1:I:345:TYR:OH	2.30	0.73
1:J:644:ASN:O	1:J:644:ASN:ND2	2.20	0.73
1:A:518:ASN:OD1	1:A:533:ASP:HB2	1.88	0.73
1:B:408:ILE:O	1:B:409:ILE:HG22	1.89	0.73
1:C:665:HIS:CE1	1:D:342:ALA:CB	2.71	0.73
1:G:340:VAL:CA	1:G:345:TYR:OH	2.30	0.73
1:I:385:THR:HG21	1:I:389:ARG:N	2.01	0.73
1:I:629:ASP:HB3	1:I:631:VAL:O	1.89	0.73
1:A:629:ASP:HB3	1:A:631:VAL:O	1.89	0.73
1:G:340:VAL:O	1:G:345:TYR:CE1	2.37	0.73
1:E:629:ASP:HB3	1:E:631:VAL:O	1.88	0.73
1:F:408:ILE:O	1:F:409:ILE:HG22	1.89	0.73
1:H:427:ASN:H	1:H:427:ASN:ND2	1.84	0.73
1:L:408:ILE:O	1:L:409:ILE:HG22	1.89	0.73
1:A:665:HIS:CE1	1:B:342:ALA:CB	2.71	0.73
1:C:518:ASN:OD1	1:C:533:ASP:HB2	1.89	0.73
1:G:635:ARG:HG2	1:G:636:ASP:N	1.90	0.73
1:I:587:ASP:OD2	1:I:587:ASP:C	2.30	0.73

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:340:VAL:O	1:A:345:TYR:CE1	2.37	0.72
1:B:437:GLU:O	1:B:441:ILE:HG22	1.88	0.72
1:D:398:ASP:OD1	1:E:663:HIS:HD2	1.71	0.72
1:K:665:HIS:CE1	1:L:342:ALA:CB	2.71	0.72
1:E:340:VAL:O	1:E:345:TYR:CE1	2.37	0.72
1:C:345:TYR:HD1	1:C:345:TYR:H	1.36	0.72
1:F:367:THR:O	1:F:369:PRO:CD	2.30	0.72
1:F:408:ILE:O	1:F:408:ILE:CD1	2.30	0.72
1:H:408:ILE:O	1:H:409:ILE:HG22	1.89	0.72
1:J:427:ASN:ND2	1:J:427:ASN:H	1.84	0.72
1:G:629:ASP:HB3	1:G:631:VAL:O	1.89	0.72
1:I:345:TYR:HD1	1:I:345:TYR:H	1.36	0.72
1:K:518:ASN:OD1	1:K:533:ASP:HB2	1.88	0.72
1:E:518:ASN:OD1	1:E:533:ASP:HB2	1.89	0.72
1:E:664:HIS:O	1:E:665:HIS:ND1	2.23	0.72
1:A:340:VAL:C	1:A:341:THR:CG2	2.62	0.72
1:A:385:THR:HG21	1:A:389:ARG:N	2.01	0.72
1:C:340:VAL:O	1:C:345:TYR:CE1	2.37	0.72
1:C:385:THR:HG21	1:C:389:ARG:N	2.01	0.72
1:E:385:THR:HG23	1:E:389:ARG:N	2.02	0.72
1:G:518:ASN:OD1	1:G:533:ASP:HB2	1.89	0.72
1:I:664:HIS:O	1:I:665:HIS:ND1	2.23	0.72
1:J:408:ILE:O	1:J:409:ILE:HG22	1.89	0.72
1:L:408:ILE:C	1:L:409:ILE:HG23	2.15	0.72
1:C:664:HIS:O	1:C:665:HIS:ND1	2.23	0.72
1:A:517:PHE:CD2	1:A:536:TYR:HE2	2.07	0.72
1:I:518:ASN:OD1	1:I:533:ASP:HB2	1.89	0.72
1:K:385:THR:HG23	1:K:389:ARG:N	2.02	0.72
1:I:385:THR:HG23	1:I:389:ARG:N	2.02	0.71
1:J:408:ILE:C	1:J:409:ILE:HG23	2.15	0.71
1:A:664:HIS:O	1:A:665:HIS:ND1	2.23	0.71
1:C:345:TYR:CD1	1:C:345:TYR:N	2.51	0.71
1:C:629:ASP:HB3	1:C:631:VAL:O	1.89	0.71
1:K:664:HIS:O	1:K:665:HIS:ND1	2.23	0.71
1:B:408:ILE:O	1:B:408:ILE:CD1	2.30	0.71
1:A:340:VAL:O	1:A:344:ASP:HB2	1.91	0.71
1:E:340:VAL:O	1:E:344:ASP:HB2	1.91	0.71
1:K:345:TYR:HD1	1:K:345:TYR:H	1.36	0.71
1:D:398:ASP:OD1	1:E:663:HIS:CD2	2.43	0.71
1:G:345:TYR:HD1	1:G:345:TYR:H	1.36	0.71
1:D:408:ILE:O	1:D:409:ILE:HG22	1.89	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:664:HIS:O	1:G:665:HIS:ND1	2.23	0.71
1:K:353:PHE:CZ	1:K:395:TYR:HD2	2.09	0.71
1:C:340:VAL:O	1:C:344:ASP:HB2	1.91	0.71
1:D:398:ASP:O	1:E:663:HIS:N	2.20	0.71
1:E:358:GLN:HB2	1:E:379:LYS:HA	1.73	0.71
1:B:408:ILE:C	1:B:409:ILE:HG23	2.15	0.71
1:I:340:VAL:O	1:I:344:ASP:HB2	1.91	0.71
1:K:340:VAL:O	1:K:344:ASP:HB2	1.91	0.71
1:G:661:LEU:CD1	1:G:662:GLU:CB	2.68	0.71
1:K:353:PHE:CG	1:K:395:TYR:HE2	2.09	0.71
1:E:517:PHE:CD2	1:E:536:TYR:HE2	2.07	0.71
1:F:385:THR:HG23	1:F:388:GLN:H	1.56	0.71
1:I:629:ASP:HA	1:J:664:HIS:HE1	1.47	0.71
1:B:385:THR:HG23	1:B:388:GLN:H	1.56	0.70
1:C:661:LEU:HD12	1:C:662:GLU:HB2	1.73	0.70
1:I:353:PHE:CG	1:I:395:TYR:HE2	2.09	0.70
1:A:353:PHE:CG	1:A:395:TYR:HE2	2.09	0.70
1:G:340:VAL:O	1:G:344:ASP:HB2	1.91	0.70
1:K:385:THR:HG21	1:K:389:ARG:N	2.01	0.70
1:L:385:THR:HG23	1:L:388:GLN:H	1.56	0.70
1:A:358:GLN:HB2	1:A:379:LYS:HA	1.73	0.70
1:E:522:LYS:N	1:E:522:LYS:HD3	2.07	0.70
1:F:408:ILE:C	1:F:409:ILE:HG23	2.15	0.70
1:I:358:GLN:HB2	1:I:379:LYS:HA	1.73	0.70
1:A:522:LYS:HD3	1:A:522:LYS:N	2.07	0.70
1:G:353:PHE:CG	1:G:395:TYR:HE2	2.09	0.70
1:G:401:LEU:O	1:G:402:ALA:CB	2.40	0.70
1:H:385:THR:HG23	1:H:388:GLN:H	1.56	0.70
1:J:385:THR:HG23	1:J:388:GLN:H	1.56	0.70
1:B:345:TYR:CE1	1:B:349:VAL:CG2	2.74	0.70
1:C:340:VAL:C	1:C:341:THR:CG2	2.62	0.70
1:D:408:ILE:C	1:D:409:ILE:HG23	2.15	0.70
1:E:401:LEU:O	1:E:402:ALA:CB	2.40	0.70
1:I:388:GLN:O	1:I:389:ARG:HB2	1.91	0.70
1:K:522:LYS:HD3	1:K:522:LYS:N	2.07	0.70
1:A:387:VAL:HG21	1:A:390:GLU:CD	2.17	0.70
1:A:661:LEU:HD12	1:A:662:GLU:HB2	1.73	0.70
1:E:353:PHE:CG	1:E:395:TYR:HE2	2.09	0.70
1:H:408:ILE:C	1:H:409:ILE:HG23	2.15	0.70
1:K:387:VAL:HG21	1:K:390:GLU:CD	2.17	0.70
1:K:661:LEU:CD1	1:K:662:GLU:CB	2.68	0.70

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:385:THR:HG23	1:A:389:ARG:N	2.02	0.70
1:B:364:THR:HG23	1:K:401:LEU:HD12	1.68	0.70
1:E:388:GLN:O	1:E:389:ARG:HG3	1.92	0.70
1:I:401:LEU:O	1:I:402:ALA:CB	2.40	0.70
1:I:440:ILE:HD11	1:I:653:ILE:CD1	2.21	0.70
1:K:385:THR:CG2	1:K:389:ARG:CA	2.60	0.70
1:A:506:LYS:O	1:A:509:SER:HB3	1.92	0.70
1:C:353:PHE:CZ	1:C:395:TYR:HD2	2.09	0.70
1:C:353:PHE:CG	1:C:395:TYR:HE2	2.09	0.70
1:C:388:GLN:O	1:C:389:ARG:HG3	1.92	0.70
1:D:345:TYR:CE1	1:D:349:VAL:CG2	2.74	0.70
1:D:567:GLU:O	1:D:568:ASN:HB3	1.92	0.70
1:E:440:ILE:HD11	1:E:653:ILE:CD1	2.21	0.70
1:G:388:GLN:O	1:G:389:ARG:HB2	1.91	0.70
1:G:440:ILE:HD11	1:G:653:ILE:CD1	2.21	0.70
1:G:516:SER:HB2	1:G:617:GLU:CD	2.17	0.70
1:G:517:PHE:CD2	1:G:536:TYR:HE2	2.07	0.70
1:K:358:GLN:HB2	1:K:379:LYS:HA	1.73	0.70
1:B:567:GLU:O	1:B:568:ASN:HB3	1.92	0.70
1:D:385:THR:HG23	1:D:388:GLN:H	1.56	0.70
1:G:401:LEU:HD23	1:G:401:LEU:O	1.92	0.70
1:G:522:LYS:HD3	1:G:522:LYS:N	2.07	0.70
1:K:440:ILE:HD11	1:K:653:ILE:CD1	2.21	0.70
1:K:516:SER:HB2	1:K:617:GLU:CD	2.17	0.70
1:C:388:GLN:O	1:C:389:ARG:HB2	1.91	0.70
1:E:342:ALA:HB1	1:E:362:THR:HB	1.74	0.70
1:I:506:LYS:O	1:I:509:SER:HB3	1.92	0.70
1:K:457:SER:H	1:K:634:THR:HG23	1.56	0.70
1:L:365:ASP:C	1:L:367:THR:HG22	2.17	0.70
1:C:358:GLN:HB2	1:C:379:LYS:HA	1.73	0.69
1:E:516:SER:HB2	1:E:617:GLU:CD	2.17	0.69
1:H:345:TYR:CE1	1:H:349:VAL:CG2	2.74	0.69
1:H:647:ARG:HH11	1:H:649:GLN:CD	2.00	0.69
1:I:516:SER:HB2	1:I:617:GLU:CD	2.17	0.69
1:K:388:GLN:O	1:K:389:ARG:HB2	1.91	0.69
1:K:506:LYS:O	1:K:509:SER:HB3	1.92	0.69
1:G:358:GLN:HB2	1:G:379:LYS:HA	1.73	0.69
1:I:661:LEU:CD1	1:I:662:GLU:CB	2.68	0.69
1:A:661:LEU:CD1	1:A:662:GLU:CB	2.68	0.69
1:C:342:ALA:HB1	1:C:362:THR:HB	1.74	0.69
1:G:353:PHE:CZ	1:G:395:TYR:HD2	2.09	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:345:TYR:CE1	1:J:349:VAL:CG2	2.74	0.69
1:K:340:VAL:C	1:K:341:THR:CG2	2.62	0.69
1:L:345:TYR:CE1	1:L:349:VAL:CG2	2.74	0.69
1:C:440:ILE:HD11	1:C:653:ILE:CD1	2.21	0.69
1:D:365:ASP:C	1:D:367:THR:HG22	2.17	0.69
1:E:403:PRO:CB	1:H:366:SER:HB2	2.09	0.69
1:G:385:THR:HG23	1:G:389:ARG:N	2.02	0.69
1:G:661:LEU:HD12	1:G:662:GLU:HB2	1.73	0.69
1:I:522:LYS:HD3	1:I:522:LYS:N	2.07	0.69
1:J:401:LEU:HG	1:J:403:PRO:HD2	1.75	0.69
1:A:345:TYR:HD1	1:A:345:TYR:H	1.36	0.69
1:A:516:SER:HB2	1:A:617:GLU:CD	2.17	0.69
1:E:661:LEU:HD12	1:E:662:GLU:HB2	1.73	0.69
1:G:342:ALA:HB1	1:G:362:THR:HB	1.74	0.69
1:G:347:THR:O	1:G:350:SER:N	2.26	0.69
1:I:347:THR:O	1:I:350:SER:N	2.26	0.69
1:A:440:ILE:HD11	1:A:653:ILE:CD1	2.21	0.69
1:C:506:LYS:O	1:C:509:SER:HB3	1.92	0.69
1:C:516:SER:HB2	1:C:617:GLU:CD	2.17	0.69
1:D:566:ASN:N	1:D:570:GLN:NE2	2.41	0.69
1:F:345:TYR:CE1	1:F:349:VAL:CG2	2.74	0.69
1:F:647:ARG:HH11	1:F:649:GLN:CD	1.99	0.69
1:G:457:SER:H	1:G:634:THR:HG23	1.56	0.69
1:I:342:ALA:HB1	1:I:362:THR:HB	1.74	0.69
1:A:347:THR:O	1:A:350:SER:N	2.26	0.69
1:A:401:LEU:HD23	1:A:401:LEU:O	1.92	0.69
1:B:365:ASP:C	1:B:367:THR:HG22	2.17	0.69
1:B:523:VAL:HG12	1:B:532:GLU:H	1.58	0.69
1:C:347:THR:O	1:C:350:SER:N	2.26	0.69
1:C:387:VAL:HG21	1:C:390:GLU:CD	2.16	0.69
1:C:661:LEU:CD1	1:C:662:GLU:CB	2.68	0.69
1:E:388:GLN:O	1:E:389:ARG:HB2	1.91	0.69
1:E:506:LYS:O	1:E:509:SER:HB3	1.92	0.69
1:H:365:ASP:C	1:H:367:THR:HG22	2.17	0.69
1:I:457:SER:H	1:I:634:THR:HG23	1.56	0.69
1:K:401:LEU:O	1:K:402:ALA:CB	2.40	0.69
1:L:401:LEU:HG	1:L:403:PRO:HD2	1.75	0.69
1:L:567:GLU:O	1:L:568:ASN:HB3	1.92	0.69
1:A:388:GLN:O	1:A:389:ARG:HB2	1.91	0.69
1:C:401:LEU:O	1:C:402:ALA:CB	2.40	0.69
1:E:340:VAL:C	1:E:341:THR:CG2	2.62	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:345:TYR:HD1	1:E:345:TYR:H	1.36	0.69
1:I:517:PHE:CD2	1:I:536:TYR:HE2	2.07	0.69
1:K:347:THR:O	1:K:350:SER:N	2.26	0.69
1:C:457:SER:H	1:C:634:THR:HG23	1.56	0.69
1:C:522:LYS:HD3	1:C:522:LYS:N	2.07	0.69
1:F:660:GLN:CA	1:F:661:LEU:CB	2.71	0.69
1:G:387:VAL:HG21	1:G:390:GLU:CD	2.17	0.69
1:H:401:LEU:HG	1:H:403:PRO:HD2	1.75	0.69
1:I:387:VAL:HG21	1:I:390:GLU:CD	2.17	0.69
1:J:567:GLU:O	1:J:568:ASN:HB3	1.92	0.69
1:G:506:LYS:O	1:G:509:SER:HB3	1.92	0.68
1:I:563:VAL:HG22	1:I:611:LEU:HD21	1.75	0.68
1:K:629:ASP:HA	1:L:664:HIS:HE1	1.47	0.68
1:E:563:VAL:HG22	1:E:611:LEU:HD21	1.75	0.68
1:I:388:GLN:O	1:I:389:ARG:HG3	1.92	0.68
1:I:661:LEU:HD12	1:I:662:GLU:HB2	1.73	0.68
1:D:523:VAL:HG12	1:D:532:GLU:H	1.58	0.68
1:E:347:THR:O	1:E:350:SER:N	2.26	0.68
1:E:531:GLU:C	1:E:532:GLU:CD	2.61	0.68
1:G:388:GLN:O	1:G:389:ARG:HG3	1.92	0.68
1:H:523:VAL:HG12	1:H:532:GLU:H	1.58	0.68
1:K:563:VAL:HG22	1:K:611:LEU:HD21	1.75	0.68
1:A:401:LEU:O	1:A:402:ALA:CB	2.40	0.68
1:C:401:LEU:HD23	1:C:401:LEU:O	1.92	0.68
1:E:387:VAL:HG21	1:E:390:GLU:CD	2.17	0.68
1:A:531:GLU:C	1:A:532:GLU:CD	2.61	0.68
1:A:563:VAL:HG22	1:A:611:LEU:HD21	1.75	0.68
1:C:563:VAL:HG22	1:C:611:LEU:HD21	1.75	0.68
1:E:385:THR:HG21	1:E:389:ARG:N	2.01	0.68
1:F:521:ARG:HH22	1:F:559:ALA:HB3	1.59	0.68
1:G:563:VAL:HG22	1:G:611:LEU:HD21	1.75	0.68
1:K:385:THR:HG22	1:K:392:ILE:HG23	1.76	0.68
1:K:388:GLN:O	1:K:389:ARG:HG3	1.92	0.68
1:B:657:PRO:C	1:B:658:ILE:CD1	2.67	0.68
1:D:521:ARG:HH22	1:D:559:ALA:HB3	1.59	0.68
1:D:657:PRO:C	1:D:658:ILE:CD1	2.67	0.68
1:F:365:ASP:C	1:F:367:THR:HG22	2.17	0.68
1:F:401:LEU:HG	1:F:403:PRO:HD2	1.75	0.68
1:I:385:THR:HG22	1:I:392:ILE:HG23	1.76	0.68
1:J:365:ASP:C	1:J:367:THR:HG22	2.17	0.68
1:K:342:ALA:HB1	1:K:362:THR:HB	1.74	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:395:TYR:HB2	1:K:408:ILE:HG12	1.76	0.68
1:A:388:GLN:O	1:A:389:ARG:HG3	1.92	0.68
1:B:521:ARG:HH22	1:B:559:ALA:HB3	1.59	0.68
1:J:647:ARG:HG3	1:J:650:TYR:CD1	2.29	0.68
1:K:531:GLU:C	1:K:532:GLU:CD	2.61	0.68
1:K:661:LEU:HD12	1:K:662:GLU:HB2	1.73	0.68
1:E:436:LEU:HD23	1:E:436:LEU:C	2.19	0.68
1:G:531:GLU:C	1:G:532:GLU:CD	2.61	0.68
1:I:395:TYR:HB2	1:I:408:ILE:HG12	1.76	0.68
1:I:401:LEU:HD23	1:I:401:LEU:O	1.92	0.68
1:I:401:LEU:CG	1:I:402:ALA:N	2.57	0.68
1:J:523:VAL:HG12	1:J:532:GLU:H	1.58	0.68
1:A:342:ALA:HB1	1:A:362:THR:HB	1.74	0.68
1:A:385:THR:HG22	1:A:392:ILE:HG23	1.76	0.68
1:C:531:GLU:C	1:C:532:GLU:CD	2.61	0.68
1:D:647:ARG:HG3	1:D:650:TYR:CD1	2.29	0.68
1:E:395:TYR:HB2	1:E:408:ILE:HG12	1.76	0.68
1:H:521:ARG:HH22	1:H:559:ALA:HB3	1.59	0.68
1:H:657:PRO:C	1:H:658:ILE:CD1	2.67	0.68
1:K:401:LEU:HD23	1:K:401:LEU:O	1.92	0.68
1:F:523:VAL:HG12	1:F:532:GLU:H	1.58	0.67
1:I:531:GLU:C	1:I:532:GLU:CD	2.61	0.67
1:K:401:LEU:CG	1:K:402:ALA:N	2.57	0.67
1:B:401:LEU:HG	1:B:403:PRO:HD2	1.75	0.67
1:D:401:LEU:HG	1:D:403:PRO:HD2	1.75	0.67
1:E:629:ASP:CA	1:F:664:HIS:NE2	2.57	0.67
1:G:440:ILE:HG13	1:G:441:ILE:N	2.09	0.67
1:I:436:LEU:HD23	1:I:436:LEU:C	2.19	0.67
1:I:440:ILE:HG13	1:I:441:ILE:N	2.09	0.67
1:D:647:ARG:HH11	1:D:649:GLN:CD	2.00	0.67
1:F:567:GLU:O	1:F:568:ASN:HB3	1.92	0.67
1:L:566:ASN:N	1:L:570:GLN:NE2	2.41	0.67
1:C:395:TYR:HB2	1:C:408:ILE:HG12	1.76	0.67
1:G:401:LEU:CG	1:G:402:ALA:N	2.57	0.67
1:H:647:ARG:HG3	1:H:650:TYR:CD1	2.29	0.67
1:I:353:PHE:CZ	1:I:395:TYR:HD2	2.09	0.67
1:L:521:ARG:HH22	1:L:559:ALA:HB3	1.59	0.67
1:A:340:VAL:O	1:A:341:THR:HG23	1.95	0.67
1:C:440:ILE:HG13	1:C:441:ILE:N	2.09	0.67
1:G:385:THR:CG2	1:G:389:ARG:CA	2.60	0.67
1:H:567:GLU:O	1:H:568:ASN:HB3	1.92	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:521:ARG:HH22	1:J:559:ALA:HB3	1.59	0.67
1:L:647:ARG:HG3	1:L:650:TYR:CD1	2.29	0.67
1:F:647:ARG:HG3	1:F:650:TYR:CD1	2.29	0.67
1:G:340:VAL:O	1:G:341:THR:HG23	1.95	0.67
1:K:340:VAL:O	1:K:341:THR:HG23	1.95	0.67
1:K:517:PHE:CD2	1:K:536:TYR:HE2	2.07	0.67
1:A:395:TYR:HB2	1:A:408:ILE:HG12	1.76	0.67
1:C:401:LEU:CG	1:C:402:ALA:N	2.57	0.67
1:D:540:ILE:HG22	1:D:555:ILE:HG13	1.77	0.67
1:D:660:GLN:CA	1:D:661:LEU:CB	2.71	0.67
1:E:661:LEU:CD1	1:E:662:GLU:CB	2.68	0.67
1:G:385:THR:HG22	1:G:392:ILE:HG23	1.76	0.67
1:G:436:LEU:HD23	1:G:436:LEU:C	2.19	0.67
1:J:647:ARG:HH11	1:J:649:GLN:CD	1.99	0.67
1:J:657:PRO:C	1:J:658:ILE:CD1	2.67	0.67
1:K:436:LEU:C	1:K:436:LEU:HD23	2.19	0.67
1:L:523:VAL:HG12	1:L:532:GLU:H	1.58	0.67
1:A:401:LEU:CG	1:A:402:ALA:N	2.57	0.67
1:K:440:ILE:HG13	1:K:441:ILE:N	2.09	0.67
1:L:657:PRO:C	1:L:658:ILE:CD1	2.67	0.67
1:A:556:GLY:HA3	1:A:589:TYR:CD1	2.30	0.67
1:B:566:ASN:N	1:B:570:GLN:NE2	2.41	0.67
1:C:424:TYR:CE2	1:C:429:LEU:HD12	2.30	0.67
1:E:440:ILE:HG13	1:E:441:ILE:N	2.09	0.67
1:F:566:ASN:C	1:F:570:GLN:HE22	2.03	0.67
1:F:657:PRO:C	1:F:658:ILE:CD1	2.67	0.67
1:I:424:TYR:CE2	1:I:429:LEU:HD12	2.30	0.67
1:L:647:ARG:HD2	1:L:650:TYR:CZ	2.30	0.67
1:C:556:GLY:HA3	1:C:589:TYR:CD1	2.30	0.67
1:D:396:LEU:CD2	1:D:406:PRO:HG2	2.25	0.67
1:E:340:VAL:O	1:E:341:THR:HG23	1.95	0.67
1:E:385:THR:HG22	1:E:392:ILE:HG23	1.76	0.67
1:E:556:GLY:HA3	1:E:589:TYR:CD1	2.30	0.67
1:F:540:ILE:HG22	1:F:555:ILE:HG13	1.77	0.67
1:J:647:ARG:HD2	1:J:650:TYR:CZ	2.30	0.67
1:B:647:ARG:HH11	1:B:649:GLN:CD	1.99	0.66
1:D:408:ILE:C	1:D:409:ILE:CG2	2.68	0.66
1:D:647:ARG:HD2	1:D:650:TYR:CZ	2.30	0.66
1:E:424:TYR:CE2	1:E:429:LEU:HD12	2.30	0.66
1:G:629:ASP:CA	1:H:664:HIS:NE2	2.57	0.66
1:H:540:ILE:HG22	1:H:555:ILE:HG13	1.77	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:540:ILE:HG22	1:J:555:ILE:HG13	1.77	0.66
1:K:424:TYR:CE2	1:K:429:LEU:HD12	2.30	0.66
1:K:556:GLY:HA3	1:K:589:TYR:CD1	2.30	0.66
1:L:647:ARG:HH11	1:L:649:GLN:CD	2.00	0.66
1:C:525:ASN:CG	1:C:526:PRO:HD3	2.20	0.66
1:E:401:LEU:CG	1:E:402:ALA:N	2.57	0.66
1:F:566:ASN:N	1:F:570:GLN:NE2	2.41	0.66
1:G:340:VAL:C	1:G:341:THR:CG2	2.62	0.66
1:H:647:ARG:HD2	1:H:650:TYR:CZ	2.30	0.66
1:L:510:MET:HE2	1:L:542:SER:HB3	1.78	0.66
1:A:424:TYR:CE2	1:A:429:LEU:HD12	2.30	0.66
1:C:436:LEU:C	1:C:436:LEU:HD23	2.19	0.66
1:D:510:MET:HE2	1:D:542:SER:HB3	1.78	0.66
1:F:365:ASP:OD1	1:F:367:THR:HG21	1.95	0.66
1:G:424:TYR:CE2	1:G:429:LEU:HD12	2.30	0.66
1:G:454:ILE:CG2	1:G:455:PHE:N	2.58	0.66
1:J:566:ASN:C	1:J:570:GLN:HE22	2.03	0.66
1:L:408:ILE:C	1:L:409:ILE:CG2	2.68	0.66
1:L:566:ASN:C	1:L:570:GLN:HE22	2.03	0.66
1:A:436:LEU:HD23	1:A:436:LEU:C	2.19	0.66
1:B:408:ILE:C	1:B:409:ILE:CG2	2.68	0.66
1:B:510:MET:HE2	1:B:542:SER:HB3	1.78	0.66
1:F:408:ILE:C	1:F:409:ILE:CG2	2.68	0.66
1:I:340:VAL:O	1:I:341:THR:HG23	1.95	0.66
1:A:661:LEU:HD11	1:A:662:GLU:HB2	1.78	0.66
1:B:396:LEU:CD2	1:B:406:PRO:HG2	2.25	0.66
1:C:385:THR:HG22	1:C:392:ILE:HG23	1.76	0.66
1:C:629:ASP:CA	1:D:664:HIS:NE2	2.57	0.66
1:D:365:ASP:OD1	1:D:367:THR:HG21	1.95	0.66
1:G:395:TYR:HB2	1:G:408:ILE:HG12	1.76	0.66
1:H:408:ILE:C	1:H:409:ILE:CG2	2.68	0.66
1:B:647:ARG:HD2	1:B:650:TYR:CZ	2.30	0.66
1:C:342:ALA:CB	1:C:362:THR:HG22	2.26	0.66
1:F:396:LEU:CD2	1:F:406:PRO:HG2	2.25	0.66
1:J:510:MET:HE2	1:J:542:SER:HB3	1.78	0.66
1:K:661:LEU:HD11	1:K:662:GLU:HB2	1.78	0.66
1:A:342:ALA:CB	1:A:362:THR:HG22	2.26	0.66
1:B:365:ASP:OD1	1:B:367:THR:HG21	1.96	0.66
1:B:366:SER:O	1:B:367:THR:HB	1.96	0.66
1:C:454:ILE:CG2	1:C:455:PHE:N	2.58	0.66
1:E:454:ILE:CG2	1:E:455:PHE:N	2.59	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:510:MET:HE2	1:F:542:SER:HB3	1.78	0.66
1:G:525:ASN:CG	1:G:526:PRO:HD3	2.20	0.66
1:I:595:ILE:HD13	1:I:597:TYR:CE2	2.31	0.66
1:I:661:LEU:HD11	1:I:662:GLU:HB2	1.78	0.66
1:K:595:ILE:HD13	1:K:597:TYR:CE2	2.31	0.66
1:B:540:ILE:HG22	1:B:555:ILE:HG13	1.77	0.66
1:J:348:PHE:O	1:J:349:VAL:C	2.39	0.66
1:C:661:LEU:HD11	1:C:662:GLU:HB2	1.78	0.66
1:E:353:PHE:CZ	1:E:395:TYR:HD2	2.09	0.66
1:F:366:SER:O	1:F:367:THR:HB	1.96	0.66
1:G:401:LEU:CD2	1:G:402:ALA:CA	2.74	0.66
1:H:348:PHE:O	1:H:351:GLU:N	2.21	0.66
1:I:454:ILE:CG2	1:I:455:PHE:N	2.58	0.66
1:I:629:ASP:CA	1:J:664:HIS:NE2	2.57	0.66
1:L:540:ILE:HG22	1:L:555:ILE:HG13	1.77	0.66
1:B:566:ASN:C	1:B:570:GLN:HE22	2.03	0.66
1:B:647:ARG:HG3	1:B:650:TYR:CD1	2.29	0.66
1:C:595:ILE:HD13	1:C:597:TYR:CE2	2.31	0.66
1:D:566:ASN:C	1:D:570:GLN:HE22	2.03	0.66
1:E:341:THR:OG1	1:E:344:ASP:HB2	1.96	0.66
1:G:556:GLY:HA3	1:G:589:TYR:CD1	2.30	0.66
1:H:569:ILE:O	1:H:569:ILE:CG2	2.30	0.66
1:I:342:ALA:CB	1:I:362:THR:HG22	2.26	0.66
1:J:566:ASN:N	1:J:570:GLN:NE2	2.41	0.66
1:A:629:ASP:CA	1:B:664:HIS:NE2	2.57	0.65
1:E:457:SER:H	1:E:634:THR:HG23	1.56	0.65
1:H:557:PRO:HG2	1:H:580:LEU:CD1	2.26	0.65
1:I:556:GLY:HA3	1:I:589:TYR:CD1	2.30	0.65
1:A:403:PRO:HG3	1:C:366:SER:CB	2.13	0.65
1:F:647:ARG:HD2	1:F:650:TYR:CZ	2.30	0.65
1:H:566:ASN:C	1:H:570:GLN:HE22	2.03	0.65
1:A:525:ASN:CG	1:A:526:PRO:HD3	2.20	0.65
1:C:392:ILE:O	1:C:396:LEU:HD23	1.97	0.65
1:E:548:LYS:O	1:E:548:LYS:HG2	1.96	0.65
1:G:595:ILE:HD13	1:G:597:TYR:CE2	2.31	0.65
1:J:557:PRO:HG2	1:J:580:LEU:CD1	2.26	0.65
1:K:454:ILE:CG2	1:K:455:PHE:N	2.58	0.65
1:L:365:ASP:OD1	1:L:367:THR:HG21	1.96	0.65
1:L:396:LEU:CD2	1:L:406:PRO:HG2	2.25	0.65
1:A:440:ILE:HG13	1:A:441:ILE:N	2.09	0.65
1:A:595:ILE:HD13	1:A:597:TYR:CE2	2.31	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:340:VAL:O	1:C:341:THR:HG23	1.95	0.65
1:H:348:PHE:O	1:H:349:VAL:C	2.39	0.65
1:H:510:MET:HE2	1:H:542:SER:HB3	1.78	0.65
1:K:342:ALA:CB	1:K:362:THR:HG22	2.26	0.65
1:E:392:ILE:O	1:E:396:LEU:HD23	1.97	0.65
1:E:634:THR:OG1	1:E:635:ARG:N	2.30	0.65
1:H:365:ASP:OD1	1:H:367:THR:HG21	1.96	0.65
1:H:366:SER:O	1:H:367:THR:HB	1.96	0.65
1:J:348:PHE:O	1:J:351:GLU:N	2.21	0.65
1:J:396:LEU:CD2	1:J:406:PRO:HG2	2.25	0.65
1:J:408:ILE:C	1:J:409:ILE:CG2	2.68	0.65
1:K:629:ASP:CA	1:L:664:HIS:NE2	2.57	0.65
1:C:341:THR:OG1	1:C:344:ASP:HB2	1.96	0.65
1:E:595:ILE:HD13	1:E:597:TYR:CE2	2.31	0.65
1:F:348:PHE:O	1:F:351:GLU:N	2.21	0.65
1:H:343:THR:O	1:H:347:THR:CG2	2.44	0.65
1:H:396:LEU:CD2	1:H:406:PRO:HG2	2.25	0.65
1:B:348:PHE:O	1:B:349:VAL:C	2.39	0.65
1:F:557:PRO:HG2	1:F:580:LEU:CD1	2.26	0.65
1:G:341:THR:OG1	1:G:344:ASP:HB2	1.96	0.65
1:G:519:SER:CB	1:G:562:ASP:OD1	2.44	0.65
1:I:343:THR:CG2	1:I:344:ASP:N	2.60	0.65
1:I:525:ASN:CG	1:I:526:PRO:HD3	2.20	0.65
1:J:646:LEU:O	1:J:646:LEU:CD2	2.30	0.65
1:L:659:SER:O	1:L:660:GLN:NE2	2.30	0.65
1:A:341:THR:OG1	1:A:344:ASP:HB2	1.96	0.65
1:D:398:ASP:O	1:E:663:HIS:CB	2.41	0.65
1:G:392:ILE:O	1:G:396:LEU:HD23	1.97	0.65
1:I:341:THR:OG1	1:I:344:ASP:HB2	1.96	0.65
1:J:440:ILE:HG22	1:J:471:ALA:HB3	1.79	0.65
1:L:366:SER:O	1:L:367:THR:HB	1.96	0.65
1:B:659:SER:O	1:B:660:GLN:NE2	2.30	0.65
1:C:385:THR:HG23	1:C:389:ARG:N	2.02	0.65
1:D:366:SER:O	1:D:367:THR:HB	1.96	0.65
1:E:401:LEU:HD23	1:E:401:LEU:O	1.92	0.65
1:H:566:ASN:O	1:H:570:GLN:NE2	2.30	0.65
1:J:366:SER:O	1:J:367:THR:HB	1.96	0.65
1:K:548:LYS:HG2	1:K:548:LYS:O	1.96	0.65
1:L:440:ILE:HG22	1:L:471:ALA:HB3	1.79	0.65
1:B:440:ILE:HG22	1:B:471:ALA:HB3	1.79	0.65
1:C:634:THR:OG1	1:C:635:ARG:N	2.30	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:398:ASP:HA	1:E:663:HIS:HB2	1.79	0.65
1:D:659:SER:O	1:D:660:GLN:NE2	2.30	0.65
1:F:566:ASN:O	1:F:570:GLN:NE2	2.30	0.65
1:J:659:SER:O	1:J:660:GLN:NE2	2.30	0.65
1:C:548:LYS:HG2	1:C:548:LYS:O	1.96	0.64
1:E:545:ARG:HB3	1:E:597:TYR:CD2	2.32	0.64
1:J:365:ASP:OD1	1:J:367:THR:HG21	1.96	0.64
1:K:519:SER:CB	1:K:562:ASP:OD1	2.44	0.64
1:L:557:PRO:HG2	1:L:580:LEU:CD1	2.26	0.64
1:D:398:ASP:CG	1:E:663:HIS:HB3	2.22	0.64
1:D:566:ASN:O	1:D:570:GLN:NE2	2.30	0.64
1:E:342:ALA:CB	1:E:362:THR:HG22	2.26	0.64
1:G:545:ARG:HB3	1:G:597:TYR:CD2	2.32	0.64
1:I:548:LYS:HG2	1:I:548:LYS:O	1.96	0.64
1:L:348:PHE:O	1:L:349:VAL:C	2.39	0.64
1:A:392:ILE:O	1:A:396:LEU:HD23	1.97	0.64
1:G:342:ALA:CB	1:G:362:THR:HG22	2.26	0.64
1:H:422:VAL:CG1	1:H:475:VAL:HG13	2.27	0.64
1:J:566:ASN:O	1:J:570:GLN:NE2	2.30	0.64
1:J:660:GLN:CA	1:J:661:LEU:CB	2.71	0.64
1:K:341:THR:OG1	1:K:344:ASP:HB2	1.96	0.64
1:A:519:SER:CB	1:A:562:ASP:OD1	2.44	0.64
1:C:343:THR:CG2	1:C:344:ASP:N	2.60	0.64
1:E:525:ASN:CG	1:E:526:PRO:HD3	2.20	0.64
1:F:659:SER:O	1:F:660:GLN:NE2	2.30	0.64
1:G:343:THR:CG2	1:G:344:ASP:N	2.60	0.64
1:A:454:ILE:CG2	1:A:455:PHE:N	2.58	0.64
1:C:517:PHE:CD2	1:C:536:TYR:HE2	2.07	0.64
1:F:422:VAL:CG1	1:F:475:VAL:HG13	2.27	0.64
1:H:440:ILE:HG22	1:H:471:ALA:HB3	1.79	0.64
1:H:566:ASN:N	1:H:570:GLN:NE2	2.41	0.64
1:A:597:TYR:HB2	1:A:598:PRO:HD3	1.80	0.64
1:F:440:ILE:HG22	1:F:471:ALA:HB3	1.79	0.64
1:I:392:ILE:O	1:I:396:LEU:HD23	1.97	0.64
1:I:545:ARG:HB3	1:I:597:TYR:CD2	2.32	0.64
1:L:345:TYR:HE1	1:L:349:VAL:HG21	1.62	0.64
1:C:518:ASN:HA	1:C:535:LEU:HD22	1.80	0.64
1:D:453:GLU:CD	1:D:641:VAL:HG23	2.23	0.64
1:D:557:PRO:HG2	1:D:580:LEU:CD1	2.26	0.64
1:G:597:TYR:HB2	1:G:598:PRO:HD3	1.80	0.64
1:H:344:ASP:OD1	1:H:345:TYR:N	2.31	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:634:THR:OG1	1:K:635:ARG:N	2.30	0.64
1:L:422:VAL:CG1	1:L:475:VAL:HG13	2.27	0.64
1:L:566:ASN:O	1:L:570:GLN:NE2	2.30	0.64
1:B:345:TYR:HE1	1:B:349:VAL:HG21	1.62	0.64
1:B:557:PRO:HG2	1:B:580:LEU:CD1	2.26	0.64
1:E:518:ASN:HA	1:E:535:LEU:HD22	1.80	0.64
1:G:661:LEU:HD11	1:G:662:GLU:HB2	1.78	0.64
1:H:659:SER:O	1:H:660:GLN:NE2	2.30	0.64
1:J:422:VAL:CG1	1:J:475:VAL:HG13	2.27	0.64
1:K:392:ILE:O	1:K:396:LEU:HD23	1.97	0.64
1:B:422:VAL:CG1	1:B:475:VAL:HG13	2.27	0.64
1:D:343:THR:O	1:D:347:THR:CG2	2.44	0.64
1:D:348:PHE:O	1:D:349:VAL:C	2.39	0.64
1:G:401:LEU:CD2	1:G:402:ALA:CB	2.69	0.64
1:L:646:LEU:O	1:L:646:LEU:CD2	2.30	0.64
1:A:343:THR:HG22	1:A:344:ASP:N	2.13	0.64
1:A:548:LYS:HG2	1:A:548:LYS:O	1.96	0.64
1:B:566:ASN:O	1:B:570:GLN:NE2	2.30	0.64
1:C:401:LEU:CD2	1:C:402:ALA:CA	2.74	0.64
1:G:343:THR:HG22	1:G:344:ASP:N	2.13	0.64
1:G:385:THR:HG21	1:G:392:ILE:H	1.63	0.64
1:J:404:ILE:HD13	1:J:404:ILE:H	1.63	0.64
1:L:343:THR:O	1:L:347:THR:CG2	2.44	0.64
1:D:465:LEU:HD21	1:D:480:ALA:HB2	1.81	0.63
1:I:385:THR:HG21	1:I:392:ILE:H	1.63	0.63
1:I:519:SER:CB	1:I:562:ASP:OD1	2.44	0.63
1:I:634:THR:OG1	1:I:635:ARG:N	2.30	0.63
1:K:545:ARG:HB3	1:K:597:TYR:CD2	2.32	0.63
1:L:465:LEU:HD21	1:L:480:ALA:HB2	1.81	0.63
1:A:545:ARG:HB3	1:A:597:TYR:CD2	2.32	0.63
1:C:385:THR:HG21	1:C:392:ILE:H	1.63	0.63
1:D:440:ILE:HG22	1:D:471:ALA:HB3	1.79	0.63
1:G:518:ASN:HA	1:G:535:LEU:HD22	1.80	0.63
1:J:344:ASP:OD1	1:J:345:TYR:N	2.31	0.63
1:J:345:TYR:HE1	1:J:349:VAL:HG21	1.62	0.63
1:J:647:ARG:CD	1:J:650:TYR:CE1	2.81	0.63
1:K:597:TYR:HB2	1:K:598:PRO:HD3	1.80	0.63
1:L:344:ASP:OD1	1:L:345:TYR:N	2.31	0.63
1:C:545:ARG:HB3	1:C:597:TYR:CD2	2.32	0.63
1:D:404:ILE:HD13	1:D:404:ILE:H	1.63	0.63
1:D:422:VAL:CG1	1:D:475:VAL:HG13	2.27	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:403:PRO:HA	1:H:366:SER:CB	2.22	0.63
1:F:344:ASP:OD1	1:F:345:TYR:N	2.31	0.63
1:F:453:GLU:CD	1:F:641:VAL:HG23	2.23	0.63
1:H:499:ILE:HD12	1:H:500:LYS:N	2.14	0.63
1:B:366:SER:HB2	1:K:403:PRO:CB	2.12	0.63
1:B:396:LEU:O	1:B:399:TYR:O	2.17	0.63
1:B:465:LEU:HD21	1:B:480:ALA:HB2	1.81	0.63
1:D:344:ASP:OD1	1:D:345:TYR:N	2.31	0.63
1:E:385:THR:HG21	1:E:392:ILE:H	1.63	0.63
1:H:647:ARG:CD	1:H:650:TYR:CE1	2.81	0.63
1:I:343:THR:HG22	1:I:344:ASP:N	2.13	0.63
1:I:385:THR:CG2	1:I:389:ARG:CA	2.60	0.63
1:A:385:THR:CG2	1:A:392:ILE:H	2.12	0.63
1:C:517:PHE:HD2	1:C:536:TYR:CE2	2.09	0.63
1:C:597:TYR:HB2	1:C:598:PRO:HD3	1.80	0.63
1:E:340:VAL:HG12	1:E:345:TYR:OH	1.99	0.63
1:H:453:GLU:CD	1:H:641:VAL:HG23	2.23	0.63
1:J:499:ILE:HD12	1:J:500:LYS:N	2.14	0.63
1:L:453:GLU:CD	1:L:641:VAL:HG23	2.23	0.63
1:A:524:VAL:HG13	1:A:529:GLY:O	1.99	0.63
1:C:519:SER:CB	1:C:562:ASP:OD1	2.44	0.63
1:E:597:TYR:HB2	1:E:598:PRO:HD3	1.80	0.63
1:G:340:VAL:HG12	1:G:345:TYR:OH	1.99	0.63
1:I:385:THR:CG2	1:I:392:ILE:H	2.12	0.63
1:E:385:THR:CG2	1:E:392:ILE:H	2.12	0.63
1:G:548:LYS:O	1:G:548:LYS:HG2	1.96	0.63
1:I:378:PRO:O	1:I:379:LYS:HB3	1.99	0.63
1:I:597:TYR:HB2	1:I:598:PRO:HD3	1.80	0.63
1:J:453:GLU:CD	1:J:641:VAL:HG23	2.23	0.63
1:K:386:THR:C	1:K:388:GLN:H	2.07	0.63
1:L:396:LEU:O	1:L:399:TYR:O	2.17	0.63
1:L:404:ILE:HD13	1:L:404:ILE:H	1.63	0.63
1:B:646:LEU:C	1:B:646:LEU:CD2	2.72	0.63
1:D:345:TYR:HE1	1:D:349:VAL:HG21	1.62	0.63
1:E:343:THR:CG2	1:E:344:ASP:N	2.60	0.63
1:E:519:SER:CB	1:E:562:ASP:OD1	2.44	0.63
1:G:378:PRO:O	1:G:379:LYS:HB3	1.99	0.63
1:I:525:ASN:OD1	1:I:526:PRO:HD3	1.99	0.63
1:J:396:LEU:O	1:J:399:TYR:O	2.17	0.63
1:L:499:ILE:HD12	1:L:500:LYS:N	2.14	0.63
1:E:661:LEU:HD11	1:E:662:GLU:HB2	1.78	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:465:LEU:HD21	1:F:480:ALA:HB2	1.81	0.63
1:F:647:ARG:CD	1:F:650:TYR:CE1	2.81	0.63
1:L:647:ARG:CD	1:L:650:TYR:CE1	2.81	0.63
1:A:340:VAL:HG12	1:A:345:TYR:OH	1.99	0.62
1:B:344:ASP:OD1	1:B:345:TYR:N	2.31	0.62
1:B:453:GLU:CD	1:B:641:VAL:HG23	2.23	0.62
1:C:340:VAL:HG12	1:C:345:TYR:OH	1.99	0.62
1:C:378:PRO:O	1:C:379:LYS:HB3	1.99	0.62
1:C:524:VAL:HG13	1:C:529:GLY:O	1.99	0.62
1:B:499:ILE:HD12	1:B:500:LYS:N	2.14	0.62
1:C:385:THR:CG2	1:C:392:ILE:H	2.12	0.62
1:F:348:PHE:O	1:F:349:VAL:C	2.39	0.62
1:H:404:ILE:H	1:H:404:ILE:HD13	1.63	0.62
1:H:465:LEU:HD21	1:H:480:ALA:HB2	1.81	0.62
1:I:533:ASP:OD2	1:I:533:ASP:N	2.32	0.62
1:K:340:VAL:HG12	1:K:345:TYR:OH	1.99	0.62
1:K:385:THR:HG21	1:K:392:ILE:H	1.63	0.62
1:A:634:THR:OG1	1:A:635:ARG:N	2.30	0.62
1:D:499:ILE:HD12	1:D:500:LYS:N	2.14	0.62
1:D:646:LEU:C	1:D:646:LEU:CD2	2.72	0.62
1:E:349:VAL:C	1:E:351:GLU:H	2.08	0.62
1:E:408:ILE:H	1:E:408:ILE:HD12	1.64	0.62
1:F:499:ILE:HD12	1:F:500:LYS:N	2.14	0.62
1:G:385:THR:CG2	1:G:392:ILE:H	2.12	0.62
1:H:660:GLN:CA	1:H:661:LEU:CB	2.71	0.62
1:K:517:PHE:HD2	1:K:536:TYR:CE2	2.09	0.62
1:K:525:ASN:OD1	1:K:526:PRO:HD3	1.98	0.62
1:A:517:PHE:HD2	1:A:536:TYR:CE2	2.09	0.62
1:A:518:ASN:HA	1:A:535:LEU:HD22	1.80	0.62
1:A:533:ASP:OD2	1:A:533:ASP:N	2.32	0.62
1:C:343:THR:HG22	1:C:344:ASP:N	2.13	0.62
1:C:415:PHE:HB2	1:C:485:VAL:HB	1.81	0.62
1:D:376:ALA:O	1:D:411:PRO:HD3	2.00	0.62
1:E:358:GLN:HB2	1:E:379:LYS:CA	2.30	0.62
1:F:646:LEU:C	1:F:646:LEU:CD2	2.72	0.62
1:G:356:ILE:O	1:G:356:ILE:HG12	1.98	0.62
1:G:408:ILE:HD12	1:G:408:ILE:H	1.64	0.62
1:G:533:ASP:OD2	1:G:533:ASP:N	2.32	0.62
1:J:465:LEU:HD21	1:J:480:ALA:HB2	1.81	0.62
1:A:378:PRO:O	1:A:379:LYS:HB3	1.99	0.62
1:A:387:VAL:C	1:A:388:GLN:CD	2.68	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:376:ALA:O	1:B:411:PRO:HD3	1.99	0.62
1:B:565:GLU:C	1:B:570:GLN:NE2	2.58	0.62
1:C:387:VAL:C	1:C:388:GLN:CD	2.68	0.62
1:D:347:THR:OG1	1:D:348:PHE:N	2.30	0.62
1:D:647:ARG:CD	1:D:650:TYR:CE1	2.81	0.62
1:E:343:THR:HG22	1:E:344:ASP:N	2.13	0.62
1:E:389:ARG:O	1:E:392:ILE:N	2.31	0.62
1:I:340:VAL:CG1	1:I:345:TYR:OH	2.48	0.62
1:K:387:VAL:C	1:K:388:GLN:CD	2.68	0.62
1:L:565:GLU:C	1:L:570:GLN:NE2	2.58	0.62
1:A:385:THR:HG21	1:A:392:ILE:H	1.63	0.62
1:A:408:ILE:H	1:A:408:ILE:HD12	1.64	0.62
1:C:340:VAL:CG1	1:C:345:TYR:OH	2.48	0.62
1:E:533:ASP:OD2	1:E:533:ASP:N	2.32	0.62
1:F:404:ILE:HD13	1:F:404:ILE:H	1.63	0.62
1:G:415:PHE:HB2	1:G:485:VAL:HB	1.81	0.62
1:H:660:GLN:CA	1:H:661:LEU:HB2	2.26	0.62
1:I:340:VAL:HG12	1:I:345:TYR:OH	1.99	0.62
1:I:518:ASN:HA	1:I:535:LEU:HD22	1.80	0.62
1:K:525:ASN:CG	1:K:526:PRO:HD3	2.20	0.62
1:D:415:PHE:HB2	1:D:485:VAL:HB	1.82	0.62
1:G:341:THR:O	1:G:345:TYR:CE1	2.53	0.62
1:G:389:ARG:O	1:G:392:ILE:N	2.31	0.62
1:H:565:GLU:C	1:H:570:GLN:NE2	2.58	0.62
1:H:647:ARG:HG2	1:H:650:TYR:CE1	2.35	0.62
1:I:386:THR:C	1:I:388:GLN:H	2.07	0.62
1:J:347:THR:OG1	1:J:348:PHE:N	2.30	0.62
1:J:660:GLN:CA	1:J:661:LEU:HB2	2.26	0.62
1:K:340:VAL:CG1	1:K:345:TYR:OH	2.48	0.62
1:K:353:PHE:CG	1:K:395:TYR:CE2	2.87	0.62
1:K:524:VAL:HG13	1:K:529:GLY:O	1.99	0.62
1:L:646:LEU:C	1:L:646:LEU:CD2	2.72	0.62
1:B:647:ARG:CD	1:B:650:TYR:CE1	2.81	0.62
1:D:396:LEU:O	1:D:399:TYR:O	2.17	0.62
1:G:525:ASN:OD1	1:G:526:PRO:HD3	1.98	0.62
1:L:647:ARG:HG2	1:L:650:TYR:CE1	2.35	0.62
1:E:401:LEU:CD2	1:E:402:ALA:CA	2.74	0.62
1:E:525:ASN:OD1	1:E:526:PRO:HD3	1.99	0.62
1:G:349:VAL:C	1:G:351:GLU:H	2.08	0.62
1:I:408:ILE:H	1:I:408:ILE:HD12	1.64	0.62
1:J:565:GLU:C	1:J:570:GLN:NE2	2.58	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:647:ARG:HG2	1:J:650:TYR:CE1	2.35	0.62
1:K:356:ILE:O	1:K:356:ILE:HG12	1.98	0.62
1:K:387:VAL:C	1:K:388:GLN:CG	2.73	0.62
1:A:340:VAL:CG1	1:A:345:TYR:OH	2.48	0.62
1:B:404:ILE:HD13	1:B:404:ILE:H	1.63	0.62
1:B:647:ARG:HG2	1:B:650:TYR:CE1	2.35	0.62
1:B:660:GLN:CA	1:B:661:LEU:CB	2.71	0.62
1:C:349:VAL:C	1:C:351:GLU:H	2.08	0.62
1:C:408:ILE:H	1:C:408:ILE:HD12	1.64	0.62
1:D:565:GLU:C	1:D:570:GLN:NE2	2.58	0.62
1:F:646:LEU:O	1:F:646:LEU:CD2	2.30	0.62
1:F:647:ARG:HG2	1:F:650:TYR:CE1	2.35	0.62
1:G:358:GLN:HB2	1:G:379:LYS:CA	2.30	0.62
1:G:524:VAL:HG13	1:G:529:GLY:O	1.99	0.62
1:H:376:ALA:O	1:H:411:PRO:HD3	2.00	0.62
1:H:396:LEU:O	1:H:399:TYR:O	2.17	0.62
1:I:356:ILE:O	1:I:356:ILE:HG12	1.99	0.62
1:J:376:ALA:O	1:J:411:PRO:HD3	2.00	0.62
1:K:343:THR:HG22	1:K:344:ASP:N	2.13	0.62
1:K:378:PRO:O	1:K:379:LYS:HB3	1.99	0.62
1:K:385:THR:CG2	1:K:392:ILE:H	2.12	0.62
1:A:341:THR:O	1:A:345:TYR:CE1	2.53	0.61
1:C:341:THR:O	1:C:345:TYR:CE1	2.53	0.61
1:C:353:PHE:CG	1:C:395:TYR:CE2	2.87	0.61
1:C:525:ASN:OD1	1:C:526:PRO:HD3	1.98	0.61
1:E:415:PHE:HB2	1:E:485:VAL:HB	1.81	0.61
1:E:427:ASN:ND2	1:E:660:GLN:HB2	2.15	0.61
1:E:521:ARG:O	1:E:533:ASP:HA	2.00	0.61
1:G:386:THR:C	1:G:388:GLN:H	2.07	0.61
1:G:521:ARG:O	1:G:533:ASP:HA	2.00	0.61
1:L:660:GLN:CA	1:L:661:LEU:HB2	2.25	0.61
1:A:415:PHE:HB2	1:A:485:VAL:HB	1.81	0.61
1:A:427:ASN:ND2	1:A:660:GLN:HB2	2.15	0.61
1:A:521:ARG:O	1:A:533:ASP:HA	2.00	0.61
1:C:401:LEU:CD2	1:C:402:ALA:CB	2.69	0.61
1:C:427:ASN:ND2	1:C:660:GLN:HB2	2.15	0.61
1:E:340:VAL:CG1	1:E:345:TYR:OH	2.48	0.61
1:F:345:TYR:HE1	1:F:349:VAL:HG21	1.62	0.61
1:F:396:LEU:O	1:F:399:TYR:O	2.17	0.61
1:F:537:ASP:O	1:F:538:VAL:HG23	2.00	0.61
1:I:341:THR:O	1:I:345:TYR:CE1	2.53	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:387:VAL:C	1:I:388:GLN:CD	2.68	0.61
1:L:347:THR:OG1	1:L:348:PHE:N	2.30	0.61
1:B:646:LEU:O	1:B:646:LEU:CD2	2.30	0.61
1:C:389:ARG:CA	1:C:392:ILE:HG12	2.28	0.61
1:E:341:THR:O	1:E:345:TYR:CE1	2.53	0.61
1:E:524:VAL:HG13	1:E:529:GLY:O	1.99	0.61
1:F:343:THR:O	1:F:347:THR:CG2	2.44	0.61
1:F:565:GLU:C	1:F:570:GLN:NE2	2.58	0.61
1:G:427:ASN:ND2	1:G:660:GLN:HB2	2.15	0.61
1:H:345:TYR:HE1	1:H:349:VAL:HG21	1.62	0.61
1:H:347:THR:OG1	1:H:348:PHE:N	2.30	0.61
1:I:385:THR:HG1	1:I:387:VAL:CG2	1.98	0.61
1:I:500:LYS:HB3	1:I:600:ASP:O	2.00	0.61
1:K:521:ARG:O	1:K:533:ASP:HA	2.00	0.61
1:A:343:THR:CG2	1:A:344:ASP:N	2.60	0.61
1:C:533:ASP:OD2	1:C:533:ASP:N	2.32	0.61
1:D:647:ARG:HG2	1:D:650:TYR:CE1	2.35	0.61
1:F:568:ASN:OD1	1:F:569:ILE:N	2.30	0.61
1:I:342:ALA:HB1	1:I:362:THR:CB	2.30	0.61
1:I:349:VAL:C	1:I:351:GLU:H	2.08	0.61
1:I:387:VAL:C	1:I:388:GLN:CG	2.73	0.61
1:I:387:VAL:O	1:I:388:GLN:CD	2.44	0.61
1:I:517:PHE:HD2	1:I:536:TYR:CE2	2.09	0.61
1:K:408:ILE:H	1:K:408:ILE:HD12	1.64	0.61
1:K:415:PHE:HB2	1:K:485:VAL:HB	1.81	0.61
1:K:500:LYS:HB3	1:K:600:ASP:O	2.00	0.61
1:K:518:ASN:HA	1:K:535:LEU:HD22	1.80	0.61
1:L:415:PHE:HB2	1:L:485:VAL:HB	1.82	0.61
1:A:342:ALA:HB1	1:A:362:THR:CB	2.30	0.61
1:A:387:VAL:O	1:A:388:GLN:CD	2.44	0.61
1:B:347:THR:OG1	1:B:348:PHE:N	2.30	0.61
1:E:356:ILE:HG12	1:E:356:ILE:O	1.98	0.61
1:E:378:PRO:O	1:E:379:LYS:HB3	1.99	0.61
1:E:387:VAL:C	1:E:388:GLN:CD	2.68	0.61
1:G:634:THR:OG1	1:G:635:ARG:N	2.30	0.61
1:I:415:PHE:HB2	1:I:485:VAL:HB	1.81	0.61
1:I:436:LEU:HD22	1:I:655:LEU:CD2	2.29	0.61
1:I:524:VAL:HG13	1:I:529:GLY:O	1.99	0.61
1:K:358:GLN:HB2	1:K:379:LYS:CA	2.30	0.61
1:K:533:ASP:OD2	1:K:533:ASP:N	2.32	0.61
1:A:353:PHE:CG	1:A:395:TYR:CE2	2.87	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:387:VAL:C	1:A:388:GLN:CG	2.73	0.61
1:B:348:PHE:O	1:B:351:GLU:N	2.21	0.61
1:G:340:VAL:CG1	1:G:345:TYR:OH	2.48	0.61
1:G:387:VAL:C	1:G:388:GLN:CD	2.68	0.61
1:G:387:VAL:O	1:G:388:GLN:CD	2.44	0.61
1:J:537:ASP:O	1:J:538:VAL:HG23	2.00	0.61
1:K:342:ALA:HB1	1:K:362:THR:CB	2.30	0.61
1:K:387:VAL:O	1:K:388:GLN:CD	2.44	0.61
1:A:349:VAL:C	1:A:351:GLU:H	2.08	0.61
1:A:401:LEU:CD2	1:A:402:ALA:CA	2.74	0.61
1:C:521:ARG:O	1:C:533:ASP:HA	2.00	0.61
1:D:348:PHE:O	1:D:351:GLU:N	2.21	0.61
1:E:500:LYS:HB3	1:E:600:ASP:O	2.00	0.61
1:G:342:ALA:HB1	1:G:362:THR:CB	2.30	0.61
1:G:517:PHE:HD2	1:G:536:TYR:CE2	2.09	0.61
1:K:401:LEU:CD2	1:K:402:ALA:CA	2.74	0.61
1:K:427:ASN:ND2	1:K:660:GLN:HB2	2.15	0.61
1:L:376:ALA:O	1:L:411:PRO:HD3	2.00	0.61
1:A:647:ARG:HG3	1:A:650:TYR:CD1	2.36	0.61
1:C:358:GLN:HB2	1:C:379:LYS:CA	2.30	0.61
1:E:386:THR:C	1:E:388:GLN:H	2.07	0.61
1:G:436:LEU:HD22	1:G:655:LEU:CD2	2.29	0.61
1:G:500:LYS:HB3	1:G:600:ASP:O	2.00	0.61
1:A:389:ARG:CA	1:A:392:ILE:HG12	2.28	0.61
1:A:457:SER:H	1:A:634:THR:HG23	1.56	0.61
1:A:500:LYS:HB3	1:A:600:ASP:O	2.00	0.61
1:B:415:PHE:HB2	1:B:485:VAL:HB	1.82	0.61
1:E:517:PHE:HD2	1:E:536:TYR:CE2	2.09	0.61
1:G:504:GLN:HE22	1:G:598:PRO:HA	1.66	0.61
1:I:353:PHE:CG	1:I:395:TYR:CE2	2.87	0.61
1:I:504:GLN:HE22	1:I:598:PRO:HA	1.66	0.61
1:I:521:ARG:O	1:I:533:ASP:HA	2.00	0.61
1:K:349:VAL:C	1:K:351:GLU:H	2.08	0.61
1:L:537:ASP:O	1:L:538:VAL:HG23	2.00	0.61
1:A:525:ASN:OD1	1:A:526:PRO:HD3	1.98	0.61
1:B:644:ASN:ND2	1:B:644:ASN:C	2.59	0.61
1:C:386:THR:C	1:C:388:GLN:H	2.07	0.61
1:C:387:VAL:O	1:C:388:GLN:CD	2.44	0.61
1:D:568:ASN:CG	1:D:569:ILE:N	2.59	0.61
1:E:387:VAL:C	1:E:388:GLN:CG	2.73	0.61
1:I:427:ASN:ND2	1:I:660:GLN:HB2	2.15	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:415:PHE:HB2	1:J:485:VAL:HB	1.81	0.61
1:K:343:THR:CG2	1:K:344:ASP:N	2.60	0.61
1:K:389:ARG:CA	1:K:392:ILE:HG12	2.28	0.61
1:K:647:ARG:HG3	1:K:650:TYR:CD1	2.36	0.61
1:A:358:GLN:HB2	1:A:379:LYS:CA	2.30	0.60
1:C:443:LYS:HE3	1:C:470:ASP:O	2.01	0.60
1:C:647:ARG:HG3	1:C:650:TYR:CD1	2.36	0.60
1:F:376:ALA:O	1:F:411:PRO:HD3	2.00	0.60
1:E:387:VAL:O	1:E:388:GLN:CD	2.44	0.60
1:E:443:LYS:HE3	1:E:470:ASP:O	2.01	0.60
1:F:644:ASN:ND2	1:F:644:ASN:C	2.59	0.60
1:K:347:THR:O	1:K:348:PHE:C	2.44	0.60
1:K:443:LYS:HE3	1:K:470:ASP:O	2.01	0.60
1:A:353:PHE:CZ	1:A:395:TYR:HD2	2.09	0.60
1:A:386:THR:C	1:A:388:GLN:H	2.07	0.60
1:C:342:ALA:HB1	1:C:362:THR:CB	2.30	0.60
1:E:429:LEU:HD22	1:E:431:GLU:O	2.01	0.60
1:E:457:SER:H	1:E:634:THR:HG21	1.66	0.60
1:H:415:PHE:HB2	1:H:485:VAL:HB	1.81	0.60
1:H:568:ASN:CG	1:H:569:ILE:N	2.59	0.60
1:I:358:GLN:HB2	1:I:379:LYS:CA	2.30	0.60
1:K:341:THR:O	1:K:345:TYR:CE1	2.53	0.60
1:K:429:LEU:HD22	1:K:431:GLU:O	2.01	0.60
1:A:504:GLN:HE22	1:A:598:PRO:HA	1.66	0.60
1:E:342:ALA:HB1	1:E:362:THR:CB	2.30	0.60
1:E:504:GLN:HE22	1:E:598:PRO:HA	1.66	0.60
1:H:537:ASP:O	1:H:538:VAL:HG23	2.01	0.60
1:I:647:ARG:HG3	1:I:650:TYR:CD1	2.36	0.60
1:J:593:GLY:HA3	1:J:603:TYR:O	2.02	0.60
1:C:387:VAL:C	1:C:388:GLN:CG	2.73	0.60
1:D:537:ASP:O	1:D:538:VAL:HG23	2.00	0.60
1:D:644:ASN:ND2	1:D:644:ASN:C	2.59	0.60
1:F:347:THR:OG1	1:F:348:PHE:N	2.30	0.60
1:I:443:LYS:HE3	1:I:470:ASP:O	2.01	0.60
1:B:418:THR:HG21	1:B:464:MET:HE1	1.84	0.60
1:E:647:ARG:HG3	1:E:650:TYR:CD1	2.36	0.60
1:G:647:ARG:HG3	1:G:650:TYR:CD1	2.36	0.60
1:J:646:LEU:C	1:J:646:LEU:CD2	2.72	0.60
1:J:664:HIS:O	1:J:664:HIS:CD2	2.54	0.60
1:L:656:GLU:OE1	1:L:664:HIS:O	2.20	0.60
1:B:664:HIS:O	1:B:664:HIS:CD2	2.54	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:385:THR:HB	1:C:391:ASP:CG	2.27	0.60
1:C:500:LYS:HB3	1:C:600:ASP:O	2.00	0.60
1:C:504:GLN:HE22	1:C:598:PRO:HA	1.66	0.60
1:G:387:VAL:C	1:G:388:GLN:CG	2.73	0.60
1:G:429:LEU:HD22	1:G:431:GLU:O	2.02	0.60
1:L:644:ASN:ND2	1:L:644:ASN:C	2.59	0.60
1:A:429:LEU:HD22	1:A:431:GLU:O	2.02	0.60
1:A:443:LYS:HE3	1:A:470:ASP:O	2.02	0.60
1:C:342:ALA:HB1	1:C:362:THR:O	2.02	0.60
1:D:398:ASP:CG	1:E:663:HIS:CG	2.71	0.60
1:D:400:ASN:HD21	1:E:662:GLU:CD	2.10	0.60
1:I:347:THR:O	1:I:348:PHE:C	2.44	0.60
1:J:343:THR:O	1:J:347:THR:CG2	2.44	0.60
1:B:656:GLU:OE1	1:B:664:HIS:O	2.20	0.60
1:F:593:GLY:HA3	1:F:603:TYR:O	2.02	0.60
1:F:647:ARG:CG	1:F:650:TYR:CE1	2.85	0.60
1:G:443:LYS:HE3	1:G:470:ASP:O	2.01	0.60
1:G:457:SER:H	1:G:634:THR:HG21	1.66	0.60
1:K:342:ALA:HB1	1:K:362:THR:O	2.02	0.60
1:C:566:ASN:O	1:C:570:GLN:NE2	2.35	0.60
1:D:350:SER:HA	1:D:357:ILE:HD11	1.84	0.60
1:D:409:ILE:HD12	1:D:410:SER:N	2.17	0.60
1:D:418:THR:HG21	1:D:464:MET:HE1	1.83	0.60
1:E:342:ALA:HB1	1:E:362:THR:O	2.02	0.60
1:F:415:PHE:HB2	1:F:485:VAL:HB	1.81	0.60
1:H:644:ASN:ND2	1:H:644:ASN:C	2.59	0.60
1:J:409:ILE:HD12	1:J:410:SER:N	2.16	0.60
1:J:464:MET:HG3	1:J:465:LEU:N	2.17	0.60
1:K:436:LEU:HD22	1:K:655:LEU:CD2	2.29	0.60
1:L:418:THR:HG21	1:L:464:MET:HE1	1.84	0.60
1:L:593:GLY:HA3	1:L:603:TYR:O	2.02	0.60
1:L:664:HIS:O	1:L:664:HIS:CD2	2.55	0.60
1:B:343:THR:O	1:B:347:THR:CG2	2.44	0.59
1:B:359:ALA:HB1	1:B:455:PHE:CE2	2.37	0.59
1:B:568:ASN:CG	1:B:569:ILE:N	2.59	0.59
1:C:356:ILE:O	1:C:356:ILE:HG12	1.98	0.59
1:D:647:ARG:CG	1:D:650:TYR:CE1	2.85	0.59
1:D:656:GLU:OE1	1:D:664:HIS:O	2.20	0.59
1:G:385:THR:HG1	1:G:387:VAL:CG2	2.01	0.59
1:K:385:THR:HB	1:K:391:ASP:CG	2.27	0.59
1:L:359:ALA:HB1	1:L:455:PHE:CE2	2.37	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:464:MET:HG3	1:L:465:LEU:N	2.17	0.59
1:D:359:ALA:HB1	1:D:455:PHE:CE2	2.37	0.59
1:H:359:ALA:HB1	1:H:455:PHE:CE2	2.37	0.59
1:H:409:ILE:HD12	1:H:410:SER:N	2.17	0.59
1:H:568:ASN:OD1	1:H:569:ILE:N	2.30	0.59
1:J:550:ILE:HD13	1:J:596:ASN:HA	1.84	0.59
1:B:537:ASP:O	1:B:538:VAL:HG23	2.00	0.59
1:B:593:GLY:HA3	1:B:603:TYR:O	2.02	0.59
1:B:660:GLN:CA	1:B:661:LEU:HB2	2.26	0.59
1:C:429:LEU:HD22	1:C:431:GLU:O	2.02	0.59
1:D:664:HIS:O	1:D:664:HIS:CD2	2.55	0.59
1:E:385:THR:O	1:E:388:GLN:N	2.35	0.59
1:F:359:ALA:HB1	1:F:455:PHE:CE2	2.37	0.59
1:G:509:SER:OG	1:G:626:ASP:N	2.34	0.59
1:I:385:THR:O	1:I:388:GLN:N	2.35	0.59
1:A:342:ALA:HB1	1:A:362:THR:O	2.02	0.59
1:A:356:ILE:O	1:A:356:ILE:HG12	1.98	0.59
1:B:347:THR:O	1:B:348:PHE:C	2.45	0.59
1:B:557:PRO:HG3	1:B:589:TYR:CZ	2.38	0.59
1:G:347:THR:O	1:G:348:PHE:C	2.44	0.59
1:G:385:THR:O	1:G:388:GLN:N	2.35	0.59
1:H:464:MET:HG3	1:H:465:LEU:N	2.17	0.59
1:H:593:GLY:HA3	1:H:603:TYR:O	2.02	0.59
1:A:457:SER:H	1:A:634:THR:HG21	1.66	0.59
1:B:440:ILE:HG22	1:B:471:ALA:CB	2.33	0.59
1:E:436:LEU:HD22	1:E:655:LEU:CD2	2.29	0.59
1:F:464:MET:HG3	1:F:465:LEU:N	2.17	0.59
1:F:557:PRO:HG3	1:F:589:TYR:CZ	2.38	0.59
1:F:664:HIS:O	1:F:664:HIS:CD2	2.54	0.59
1:H:418:THR:HG21	1:H:464:MET:HE1	1.84	0.59
1:H:647:ARG:CG	1:H:650:TYR:CE1	2.85	0.59
1:I:385:THR:HB	1:I:391:ASP:CG	2.27	0.59
1:I:389:ARG:O	1:I:392:ILE:N	2.31	0.59
1:J:644:ASN:ND2	1:J:644:ASN:C	2.59	0.59
1:L:647:ARG:CG	1:L:650:TYR:CE1	2.85	0.59
1:F:656:GLU:OE1	1:F:664:HIS:O	2.20	0.59
1:H:646:LEU:O	1:H:646:LEU:CD2	2.30	0.59
1:I:342:ALA:HB1	1:I:362:THR:O	2.02	0.59
1:I:429:LEU:HD22	1:I:431:GLU:O	2.01	0.59
1:J:656:GLU:OE1	1:J:664:HIS:O	2.20	0.59
1:L:550:ILE:HD13	1:L:596:ASN:HA	1.84	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:557:PRO:HG3	1:L:589:TYR:CZ	2.38	0.59
1:A:385:THR:HB	1:A:391:ASP:CG	2.27	0.59
1:B:568:ASN:OD1	1:B:569:ILE:N	2.30	0.59
1:D:402:ALA:N	1:D:403:PRO:CD	2.66	0.59
1:D:440:ILE:HG22	1:D:471:ALA:CB	2.33	0.59
1:F:350:SER:HA	1:F:357:ILE:HD11	1.84	0.59
1:F:440:ILE:HG22	1:F:471:ALA:CB	2.33	0.59
1:F:509:SER:OG	1:F:626:ASP:N	2.32	0.59
1:G:353:PHE:CG	1:G:395:TYR:CE2	2.87	0.59
1:G:385:THR:HB	1:G:391:ASP:CG	2.27	0.59
1:G:566:ASN:O	1:G:570:GLN:NE2	2.35	0.59
1:L:440:ILE:HG22	1:L:471:ALA:CB	2.33	0.59
1:C:509:SER:OG	1:C:626:ASP:N	2.34	0.59
1:F:568:ASN:CG	1:F:569:ILE:N	2.59	0.59
1:H:664:HIS:O	1:H:664:HIS:CD2	2.54	0.59
1:J:418:THR:HG21	1:J:464:MET:HE1	1.84	0.59
1:K:504:GLN:HE22	1:K:598:PRO:HA	1.66	0.59
1:L:350:SER:HA	1:L:357:ILE:HD11	1.84	0.59
1:B:350:SER:HA	1:B:357:ILE:HD11	1.84	0.59
1:B:367:THR:CG2	1:B:368:LYS:N	2.44	0.59
1:D:593:GLY:HA3	1:D:603:TYR:O	2.02	0.59
1:E:385:THR:HB	1:E:391:ASP:CG	2.27	0.59
1:H:557:PRO:HG3	1:H:589:TYR:CZ	2.38	0.59
1:J:359:ALA:HB1	1:J:455:PHE:CE2	2.37	0.59
1:L:346:ASP:OD2	1:L:361:GLN:HA	2.03	0.59
1:A:385:THR:CG2	1:A:389:ARG:C	2.73	0.59
1:B:550:ILE:HD13	1:B:596:ASN:HA	1.84	0.59
1:E:347:THR:O	1:E:348:PHE:C	2.44	0.59
1:C:347:THR:O	1:C:348:PHE:C	2.44	0.58
1:G:342:ALA:HB1	1:G:362:THR:O	2.02	0.58
1:J:647:ARG:CG	1:J:650:TYR:CE1	2.85	0.58
1:A:385:THR:O	1:A:388:GLN:N	2.35	0.58
1:C:385:THR:O	1:C:388:GLN:N	2.35	0.58
1:C:389:ARG:O	1:C:392:ILE:N	2.31	0.58
1:C:617:GLU:HG3	1:C:618:VAL:HG13	1.85	0.58
1:D:464:MET:HG3	1:D:465:LEU:N	2.17	0.58
1:F:346:ASP:OD2	1:F:361:GLN:HA	2.03	0.58
1:F:550:ILE:HD13	1:F:596:ASN:HA	1.84	0.58
1:H:347:THR:O	1:H:348:PHE:C	2.46	0.58
1:H:402:ALA:N	1:H:403:PRO:CD	2.66	0.58
1:H:440:ILE:HG22	1:H:471:ALA:CB	2.33	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:550:ILE:HD13	1:H:596:ASN:HA	1.84	0.58
1:L:409:ILE:HD12	1:L:410:SER:N	2.17	0.58
1:A:385:THR:OG1	1:A:389:ARG:N	2.37	0.58
1:A:436:LEU:HD22	1:A:655:LEU:CD2	2.29	0.58
1:D:396:LEU:HD21	1:D:406:PRO:HG2	1.85	0.58
1:H:509:SER:OG	1:H:626:ASP:N	2.32	0.58
1:I:524:VAL:HG13	1:I:529:GLY:H	1.69	0.58
1:J:346:ASP:OD2	1:J:361:GLN:HA	2.03	0.58
1:J:557:PRO:HG3	1:J:589:TYR:CZ	2.38	0.58
1:K:385:THR:O	1:K:388:GLN:N	2.35	0.58
1:A:524:VAL:HG13	1:A:529:GLY:H	1.69	0.58
1:B:647:ARG:CG	1:B:650:TYR:CE1	2.85	0.58
1:C:664:HIS:O	1:C:665:HIS:CB	2.51	0.58
1:D:346:ASP:OD2	1:D:361:GLN:HA	2.03	0.58
1:E:524:VAL:HG13	1:E:529:GLY:H	1.68	0.58
1:G:385:THR:OG1	1:G:389:ARG:N	2.37	0.58
1:G:524:VAL:HG13	1:G:529:GLY:H	1.69	0.58
1:H:646:LEU:C	1:H:646:LEU:CD2	2.72	0.58
1:J:440:ILE:HG22	1:J:471:ALA:CB	2.33	0.58
1:K:524:VAL:HG13	1:K:529:GLY:H	1.69	0.58
1:K:664:HIS:O	1:K:665:HIS:CB	2.51	0.58
1:L:402:ALA:N	1:L:403:PRO:CD	2.66	0.58
1:A:385:THR:HG1	1:A:387:VAL:CG2	2.05	0.58
1:A:664:HIS:O	1:A:665:HIS:CB	2.51	0.58
1:C:385:THR:OG1	1:C:389:ARG:N	2.37	0.58
1:D:557:PRO:HG3	1:D:589:TYR:CZ	2.38	0.58
1:F:418:THR:HG21	1:F:464:MET:HE1	1.84	0.58
1:I:457:SER:H	1:I:634:THR:HG21	1.66	0.58
1:K:389:ARG:O	1:K:392:ILE:N	2.31	0.58
1:K:499:ILE:HD11	1:K:623:LEU:HD23	1.86	0.58
1:L:504:GLN:NE2	1:L:597:TYR:O	2.37	0.58
1:A:408:ILE:HD12	1:A:408:ILE:N	2.19	0.58
1:B:346:ASP:OD2	1:B:361:GLN:HA	2.03	0.58
1:B:402:ALA:N	1:B:403:PRO:CD	2.66	0.58
1:E:389:ARG:CA	1:E:392:ILE:HG12	2.28	0.58
1:I:385:THR:OG1	1:I:389:ARG:N	2.37	0.58
1:I:499:ILE:HD11	1:I:623:LEU:HD23	1.86	0.58
1:K:408:ILE:HD12	1:K:408:ILE:N	2.19	0.58
1:C:385:THR:CB	1:C:389:ARG:H	2.17	0.58
1:E:353:PHE:CG	1:E:395:TYR:CE2	2.87	0.58
1:G:486:ARG:NH1	1:G:500:LYS:O	2.36	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:656:GLU:OE1	1:H:664:HIS:O	2.20	0.58
1:I:401:LEU:CD2	1:I:402:ALA:CA	2.74	0.58
1:K:617:GLU:HG3	1:K:618:VAL:HG13	1.85	0.58
1:D:398:ASP:CG	1:E:663:HIS:CB	2.77	0.58
1:D:504:GLN:NE2	1:D:597:TYR:O	2.37	0.58
1:F:389:ARG:NH2	1:F:410:SER:OG	2.37	0.58
1:G:408:ILE:HD12	1:G:408:ILE:N	2.19	0.58
1:G:499:ILE:HD11	1:G:623:LEU:HD23	1.86	0.58
1:J:350:SER:HA	1:J:357:ILE:HD11	1.84	0.58
1:J:396:LEU:HD21	1:J:406:PRO:HG2	1.85	0.58
1:K:635:ARG:HG3	1:K:636:ASP:N	2.19	0.58
1:L:409:ILE:CD1	1:L:410:SER:O	2.50	0.58
1:L:593:GLY:HA3	1:L:604:TRP:HA	1.86	0.58
1:A:347:THR:O	1:A:348:PHE:C	2.44	0.58
1:A:566:ASN:O	1:A:570:GLN:NE2	2.35	0.58
1:A:635:ARG:HG3	1:A:636:ASP:N	2.19	0.58
1:C:424:TYR:CB	1:C:475:VAL:H	2.17	0.58
1:D:389:ARG:NH2	1:D:410:SER:OG	2.37	0.58
1:D:593:GLY:HA3	1:D:604:TRP:HA	1.86	0.58
1:G:525:ASN:HB3	1:G:526:PRO:CD	2.17	0.58
1:H:346:ASP:OD2	1:H:361:GLN:HA	2.03	0.58
1:C:408:ILE:HD12	1:C:408:ILE:N	2.19	0.58
1:C:524:VAL:HG13	1:C:529:GLY:H	1.69	0.58
1:E:566:ASN:O	1:E:570:GLN:NE2	2.35	0.58
1:H:350:SER:HA	1:H:357:ILE:HD11	1.84	0.58
1:K:385:THR:OG1	1:K:389:ARG:N	2.37	0.58
1:L:565:GLU:C	1:L:570:GLN:HE22	2.12	0.58
1:A:499:ILE:HD11	1:A:623:LEU:HD23	1.86	0.57
1:A:536:TYR:CD1	1:A:586:ARG:HD3	2.39	0.57
1:B:565:GLU:C	1:B:570:GLN:HE22	2.12	0.57
1:C:536:TYR:CD1	1:C:586:ARG:HD3	2.39	0.57
1:E:493:LYS:HE2	1:E:494:THR:O	2.04	0.57
1:F:402:ALA:N	1:F:403:PRO:CD	2.66	0.57
1:F:504:GLN:NE2	1:F:597:TYR:O	2.37	0.57
1:L:389:ARG:NH2	1:L:410:SER:OG	2.37	0.57
1:L:396:LEU:HD13	1:L:406:PRO:HD2	1.86	0.57
1:A:389:ARG:O	1:A:392:ILE:N	2.31	0.57
1:C:486:ARG:NH1	1:C:500:LYS:O	2.36	0.57
1:E:385:THR:OG1	1:E:389:ARG:N	2.37	0.57
1:E:408:ILE:HD12	1:E:408:ILE:N	2.19	0.57
1:F:396:LEU:HD21	1:F:406:PRO:HG2	1.85	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:593:GLY:HA3	1:F:604:TRP:HA	1.86	0.57
1:G:424:TYR:CB	1:G:475:VAL:H	2.17	0.57
1:H:389:ARG:NH2	1:H:410:SER:OG	2.37	0.57
1:H:656:GLU:OE1	1:H:664:HIS:HB2	2.04	0.57
1:I:617:GLU:HG3	1:I:618:VAL:HG13	1.86	0.57
1:J:565:GLU:C	1:J:570:GLN:HE22	2.12	0.57
1:A:424:TYR:CB	1:A:475:VAL:H	2.17	0.57
1:B:396:LEU:HD21	1:B:406:PRO:HG2	1.85	0.57
1:C:499:ILE:HD11	1:C:623:LEU:HD23	1.86	0.57
1:D:347:THR:O	1:D:348:PHE:C	2.45	0.57
1:D:396:LEU:HD13	1:D:406:PRO:HD2	1.87	0.57
1:E:385:THR:CG2	1:E:392:ILE:HG23	2.35	0.57
1:F:657:PRO:O	1:F:658:ILE:CD1	2.53	0.57
1:H:504:GLN:NE2	1:H:597:TYR:O	2.37	0.57
1:I:424:TYR:CB	1:I:475:VAL:H	2.17	0.57
1:J:389:ARG:NH2	1:J:410:SER:OG	2.37	0.57
1:J:593:GLY:HA3	1:J:604:TRP:HA	1.86	0.57
1:K:457:SER:H	1:K:634:THR:HG21	1.66	0.57
1:K:509:SER:OG	1:K:626:ASP:N	2.34	0.57
1:K:536:TYR:CD1	1:K:586:ARG:HD3	2.39	0.57
1:L:656:GLU:OE1	1:L:664:HIS:HB2	2.04	0.57
1:B:389:ARG:NH2	1:B:410:SER:OG	2.37	0.57
1:B:464:MET:HG3	1:B:465:LEU:N	2.17	0.57
1:B:657:PRO:O	1:B:658:ILE:CD1	2.53	0.57
1:E:424:TYR:CB	1:E:475:VAL:H	2.17	0.57
1:F:409:ILE:CD1	1:F:410:SER:N	2.67	0.57
1:F:565:GLU:C	1:F:570:GLN:HE22	2.12	0.57
1:F:656:GLU:OE1	1:F:664:HIS:HB2	2.04	0.57
1:J:656:GLU:OE1	1:J:664:HIS:HB2	2.04	0.57
1:K:385:THR:CG2	1:K:389:ARG:C	2.73	0.57
1:K:424:TYR:CB	1:K:475:VAL:H	2.17	0.57
1:A:509:SER:OG	1:A:626:ASP:N	2.34	0.57
1:A:617:GLU:HG3	1:A:618:VAL:HG13	1.86	0.57
1:B:409:ILE:HD12	1:B:410:SER:N	2.16	0.57
1:C:436:LEU:HD22	1:C:655:LEU:CD2	2.29	0.57
1:D:656:GLU:OE1	1:D:664:HIS:HB2	2.04	0.57
1:H:565:GLU:C	1:H:570:GLN:HE22	2.12	0.57
1:I:385:THR:HG21	1:I:389:ARG:O	2.04	0.57
1:I:664:HIS:O	1:I:665:HIS:CB	2.51	0.57
1:K:385:THR:HG21	1:K:389:ARG:O	2.04	0.57
1:K:566:ASN:O	1:K:570:GLN:NE2	2.35	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:493:LYS:HE2	1:C:494:THR:O	2.04	0.57
1:D:550:ILE:HD13	1:D:596:ASN:HA	1.84	0.57
1:D:565:GLU:C	1:D:570:GLN:HE22	2.12	0.57
1:D:657:PRO:O	1:D:658:ILE:CD1	2.53	0.57
1:E:499:ILE:HD11	1:E:623:LEU:HD23	1.86	0.57
1:F:656:GLU:OE1	1:F:664:HIS:CB	2.53	0.57
1:G:493:LYS:HE2	1:G:494:THR:O	2.04	0.57
1:H:657:PRO:O	1:H:658:ILE:CD1	2.53	0.57
1:I:566:ASN:O	1:I:570:GLN:NE2	2.35	0.57
1:L:656:GLU:OE1	1:L:664:HIS:CB	2.53	0.57
1:B:504:GLN:NE2	1:B:597:TYR:O	2.37	0.57
1:B:656:GLU:OE1	1:B:664:HIS:HB2	2.04	0.57
1:E:486:ARG:NH1	1:E:500:LYS:O	2.36	0.57
1:F:346:ASP:OD1	1:F:362:THR:OG1	2.23	0.57
1:G:385:THR:HG21	1:G:389:ARG:O	2.04	0.57
1:I:408:ILE:HD12	1:I:408:ILE:N	2.19	0.57
1:I:493:LYS:HE2	1:I:494:THR:O	2.04	0.57
1:J:347:THR:O	1:J:348:PHE:C	2.46	0.57
1:J:656:GLU:OE1	1:J:664:HIS:CB	2.53	0.57
1:K:493:LYS:HE2	1:K:494:THR:O	2.04	0.57
1:L:409:ILE:CD1	1:L:410:SER:N	2.67	0.57
1:C:635:ARG:HH22	1:D:661:LEU:CD1	2.14	0.57
1:G:379:LYS:O	1:G:379:LYS:HG2	2.05	0.57
1:I:536:TYR:CD1	1:I:586:ARG:HD3	2.39	0.57
1:B:509:SER:OG	1:B:626:ASP:N	2.32	0.57
1:C:388:GLN:C	1:C:389:ARG:HG3	2.30	0.57
1:D:656:GLU:OE1	1:D:664:HIS:CB	2.53	0.57
1:E:536:TYR:CD1	1:E:586:ARG:HD3	2.39	0.57
1:G:536:TYR:CD1	1:G:586:ARG:HD3	2.39	0.57
1:I:385:THR:CG2	1:I:392:ILE:HG23	2.35	0.57
1:J:657:PRO:O	1:J:658:ILE:CD1	2.53	0.57
1:A:385:THR:CB	1:A:389:ARG:H	2.17	0.57
1:C:385:THR:CG2	1:C:392:ILE:HG23	2.35	0.57
1:E:382:LEU:HD21	1:E:643:GLU:HG3	1.87	0.57
1:F:405:THR:O	1:F:406:PRO:O	2.23	0.57
1:G:385:THR:CG2	1:G:392:ILE:HG23	2.35	0.57
1:H:656:GLU:OE1	1:H:664:HIS:CB	2.53	0.57
1:J:504:GLN:NE2	1:J:597:TYR:O	2.37	0.57
1:K:385:THR:CG2	1:K:392:ILE:HG23	2.35	0.57
1:L:396:LEU:HD21	1:L:406:PRO:HG2	1.85	0.57
1:A:493:LYS:HE2	1:A:494:THR:O	2.04	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:346:ASP:OD1	1:B:362:THR:OG1	2.23	0.56
1:E:388:GLN:C	1:E:389:ARG:HG3	2.30	0.56
1:G:617:GLU:HG3	1:G:618:VAL:HG13	1.85	0.56
1:H:396:LEU:HD21	1:H:406:PRO:HG2	1.85	0.56
1:K:379:LYS:HG2	1:K:379:LYS:O	2.05	0.56
1:K:454:ILE:HG22	1:K:455:PHE:N	2.20	0.56
1:B:405:THR:O	1:B:406:PRO:O	2.23	0.56
1:D:568:ASN:OD1	1:D:569:ILE:N	2.30	0.56
1:G:388:GLN:O	1:G:389:ARG:NE	2.39	0.56
1:G:389:ARG:CA	1:G:392:ILE:HG12	2.28	0.56
1:G:664:HIS:O	1:G:665:HIS:CB	2.51	0.56
1:H:409:ILE:CD1	1:H:410:SER:N	2.67	0.56
1:H:593:GLY:HA3	1:H:604:TRP:HA	1.86	0.56
1:J:402:ALA:N	1:J:403:PRO:CD	2.66	0.56
1:L:568:ASN:OD1	1:L:569:ILE:N	2.30	0.56
1:A:385:THR:CG2	1:A:392:ILE:HG23	2.35	0.56
1:A:388:GLN:C	1:A:389:ARG:HG3	2.30	0.56
1:A:525:ASN:HB3	1:A:526:PRO:CD	2.17	0.56
1:B:499:ILE:HG23	1:B:602:ILE:HB	1.87	0.56
1:C:427:ASN:ND2	1:C:427:ASN:H	2.03	0.56
1:D:409:ILE:CD1	1:D:410:SER:O	2.50	0.56
1:D:661:LEU:C	1:D:661:LEU:HD22	2.30	0.56
1:E:353:PHE:O	1:E:354:GLY:C	2.48	0.56
1:E:635:ARG:HH22	1:F:661:LEU:CD1	2.14	0.56
1:F:661:LEU:C	1:F:661:LEU:HD22	2.30	0.56
1:H:405:THR:O	1:H:406:PRO:O	2.23	0.56
1:K:388:GLN:O	1:K:389:ARG:NE	2.39	0.56
1:L:568:ASN:CG	1:L:569:ILE:N	2.59	0.56
1:B:656:GLU:OE1	1:B:664:HIS:CB	2.53	0.56
1:C:385:THR:HG21	1:C:389:ARG:O	2.04	0.56
1:C:457:SER:H	1:C:634:THR:HG21	1.66	0.56
1:D:346:ASP:OD1	1:D:362:THR:OG1	2.23	0.56
1:E:385:THR:HG21	1:E:389:ARG:O	2.04	0.56
1:G:403:PRO:CD	1:J:366:SER:HA	2.24	0.56
1:H:596:ASN:HD22	1:H:599:ALA:H	1.54	0.56
1:J:396:LEU:HD13	1:J:406:PRO:HD2	1.86	0.56
1:J:509:SER:OG	1:J:626:ASP:N	2.32	0.56
1:K:382:LEU:HD21	1:K:643:GLU:HG3	1.87	0.56
1:K:387:VAL:HG21	1:K:390:GLU:HB2	1.88	0.56
1:K:486:ARG:NH1	1:K:500:LYS:O	2.36	0.56
1:A:486:ARG:NH1	1:A:500:LYS:O	2.36	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:409:ILE:HD13	1:B:410:SER:C	2.31	0.56
1:D:499:ILE:HG23	1:D:602:ILE:HB	1.87	0.56
1:I:385:THR:CB	1:I:389:ARG:H	2.17	0.56
1:K:385:THR:CB	1:K:389:ARG:H	2.17	0.56
1:K:525:ASN:HB3	1:K:526:PRO:CD	2.17	0.56
1:L:657:PRO:O	1:L:658:ILE:CD1	2.53	0.56
1:A:342:ALA:CB	1:A:362:THR:CG2	2.84	0.56
1:A:388:GLN:O	1:A:389:ARG:NE	2.38	0.56
1:B:409:ILE:CD1	1:B:410:SER:N	2.67	0.56
1:B:409:ILE:CD1	1:B:410:SER:O	2.50	0.56
1:B:593:GLY:HA3	1:B:604:TRP:HA	1.86	0.56
1:D:646:LEU:O	1:D:646:LEU:CD2	2.30	0.56
1:E:389:ARG:HA	1:E:392:ILE:CG1	2.29	0.56
1:F:396:LEU:HD13	1:F:406:PRO:HD2	1.86	0.56
1:G:388:GLN:C	1:G:389:ARG:HG3	2.30	0.56
1:H:346:ASP:OD1	1:H:362:THR:OG1	2.23	0.56
1:H:409:ILE:CD1	1:H:410:SER:C	2.79	0.56
1:I:387:VAL:HG21	1:I:390:GLU:HB2	1.88	0.56
1:I:509:SER:OG	1:I:626:ASP:N	2.34	0.56
1:L:346:ASP:OD1	1:L:362:THR:OG1	2.23	0.56
1:A:387:VAL:HG21	1:A:390:GLU:HB2	1.88	0.56
1:B:396:LEU:HD13	1:B:406:PRO:HD2	1.86	0.56
1:C:387:VAL:CG2	1:C:390:GLU:CD	2.79	0.56
1:D:409:ILE:CD1	1:D:410:SER:N	2.67	0.56
1:E:664:HIS:O	1:E:665:HIS:CB	2.51	0.56
1:F:365:ASP:CG	1:F:367:THR:HG21	2.31	0.56
1:G:560:SER:N	1:G:561:GLY:HA2	2.20	0.56
1:K:387:VAL:CG2	1:K:390:GLU:CD	2.79	0.56
1:K:388:GLN:C	1:K:389:ARG:HG3	2.30	0.56
1:L:405:THR:O	1:L:406:PRO:O	2.23	0.56
1:E:379:LYS:HG2	1:E:379:LYS:O	2.05	0.56
1:E:509:SER:OG	1:E:626:ASP:N	2.34	0.56
1:F:596:ASN:HD22	1:F:599:ALA:H	1.54	0.56
1:G:387:VAL:HG21	1:G:390:GLU:HB2	1.88	0.56
1:I:379:LYS:O	1:I:379:LYS:HG2	2.05	0.56
1:I:389:ARG:HA	1:I:392:ILE:CG1	2.29	0.56
1:J:405:THR:O	1:J:406:PRO:O	2.23	0.56
1:A:379:LYS:O	1:A:379:LYS:HG2	2.05	0.56
1:E:387:VAL:HG21	1:E:390:GLU:HB2	1.88	0.56
1:E:617:GLU:HG3	1:E:618:VAL:HG13	1.85	0.56
1:H:409:ILE:HD13	1:H:410:SER:C	2.31	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:448:TYR:CD1	1:K:642:PHE:HB2	2.41	0.56
1:L:596:ASN:HD22	1:L:599:ALA:H	1.54	0.56
1:A:427:ASN:ND2	1:A:427:ASN:H	2.03	0.56
1:B:365:ASP:CG	1:B:367:THR:HG21	2.31	0.56
1:B:366:SER:CB	1:K:403:PRO:HA	2.22	0.56
1:B:658:ILE:HD13	1:B:658:ILE:N	2.21	0.56
1:C:388:GLN:O	1:C:389:ARG:NE	2.39	0.56
1:D:359:ALA:O	1:D:376:ALA:HA	2.06	0.56
1:D:647:ARG:HD3	1:D:649:GLN:HG3	1.88	0.56
1:E:401:LEU:HD12	1:H:364:THR:HG23	1.67	0.56
1:E:448:TYR:CD1	1:E:642:PHE:HB2	2.41	0.56
1:H:396:LEU:HD13	1:H:406:PRO:HD2	1.87	0.56
1:I:342:ALA:CB	1:I:362:THR:CG2	2.84	0.56
1:I:388:GLN:C	1:I:389:ARG:HG3	2.30	0.56
1:I:427:ASN:ND2	1:I:427:ASN:H	2.03	0.56
1:J:409:ILE:CD1	1:J:410:SER:C	2.79	0.56
1:A:454:ILE:HG22	1:A:455:PHE:N	2.20	0.55
1:B:596:ASN:HD22	1:B:599:ALA:H	1.54	0.55
1:C:353:PHE:O	1:C:354:GLY:C	2.48	0.55
1:E:385:THR:HG1	1:E:387:VAL:CG2	2.05	0.55
1:E:427:ASN:ND2	1:E:427:ASN:H	2.03	0.55
1:F:409:ILE:CD1	1:F:410:SER:C	2.79	0.55
1:I:388:GLN:O	1:I:389:ARG:NE	2.39	0.55
1:J:596:ASN:HD22	1:J:599:ALA:H	1.54	0.55
1:K:385:THR:HG1	1:K:387:VAL:CG2	2.05	0.55
1:L:359:ALA:O	1:L:376:ALA:HA	2.06	0.55
1:A:382:LEU:HD21	1:A:643:GLU:HG3	1.87	0.55
1:B:359:ALA:O	1:B:376:ALA:HA	2.06	0.55
1:D:405:THR:O	1:D:406:PRO:O	2.23	0.55
1:G:342:ALA:CB	1:G:362:THR:CG2	2.84	0.55
1:H:365:ASP:CG	1:H:367:THR:HG21	2.31	0.55
1:I:342:ALA:HB1	1:I:362:THR:CG2	2.37	0.55
1:I:358:GLN:OE1	1:I:377:LYS:HE3	2.07	0.55
1:I:387:VAL:CG2	1:I:390:GLU:CD	2.79	0.55
1:I:448:TYR:CD1	1:I:642:PHE:HB2	2.41	0.55
1:J:346:ASP:OD1	1:J:362:THR:OG1	2.23	0.55
1:J:409:ILE:HD13	1:J:410:SER:C	2.31	0.55
1:K:358:GLN:OE1	1:K:377:LYS:HE3	2.07	0.55
1:L:409:ILE:CD1	1:L:410:SER:C	2.79	0.55
1:A:560:SER:N	1:A:561:GLY:HA2	2.20	0.55
1:B:409:ILE:CD1	1:B:410:SER:C	2.79	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:382:LEU:HD21	1:C:643:GLU:HG3	1.87	0.55
1:D:356:ILE:O	1:D:379:LYS:HB3	2.06	0.55
1:D:365:ASP:CG	1:D:367:THR:HG21	2.31	0.55
1:D:509:SER:OG	1:D:626:ASP:N	2.32	0.55
1:D:658:ILE:HD13	1:D:658:ILE:N	2.21	0.55
1:G:382:LEU:HD21	1:G:643:GLU:HG3	1.87	0.55
1:G:454:ILE:HG22	1:G:455:PHE:N	2.20	0.55
1:J:409:ILE:CD1	1:J:410:SER:N	2.67	0.55
1:J:568:ASN:CG	1:J:569:ILE:N	2.59	0.55
1:K:342:ALA:CB	1:K:362:THR:CG2	2.84	0.55
1:A:385:THR:HG21	1:A:389:ARG:O	2.05	0.55
1:A:448:TYR:CD1	1:A:642:PHE:HB2	2.41	0.55
1:D:398:ASP:OD2	1:E:663:HIS:HB2	2.00	0.55
1:D:409:ILE:CD1	1:D:410:SER:C	2.79	0.55
1:D:409:ILE:HD13	1:D:410:SER:C	2.31	0.55
1:E:387:VAL:CG2	1:E:390:GLU:CD	2.79	0.55
1:E:388:GLN:O	1:E:389:ARG:NE	2.39	0.55
1:E:526:PRO:O	1:E:528:THR:CG2	2.55	0.55
1:I:560:SER:N	1:I:561:GLY:HA2	2.20	0.55
1:J:356:ILE:O	1:J:379:LYS:HB3	2.06	0.55
1:A:358:GLN:OE1	1:A:377:LYS:HE3	2.07	0.55
1:B:661:LEU:C	1:B:663:HIS:N	2.61	0.55
1:C:342:ALA:CB	1:C:362:THR:CG2	2.84	0.55
1:C:448:TYR:CD1	1:C:642:PHE:HB2	2.41	0.55
1:C:454:ILE:HG22	1:C:455:PHE:N	2.20	0.55
1:D:398:ASP:O	1:E:663:HIS:HB3	2.02	0.55
1:G:427:ASN:ND2	1:G:427:ASN:H	2.03	0.55
1:H:499:ILE:HG23	1:H:602:ILE:HB	1.87	0.55
1:I:382:LEU:HD21	1:I:643:GLU:HG3	1.87	0.55
1:I:486:ARG:NH1	1:I:500:LYS:O	2.36	0.55
1:I:532:GLU:N	1:I:532:GLU:OE2	2.39	0.55
1:K:427:ASN:ND2	1:K:427:ASN:H	2.03	0.55
1:E:342:ALA:CB	1:E:362:THR:CG2	2.84	0.55
1:E:403:PRO:HG3	1:H:366:SER:HA	1.87	0.55
1:F:409:ILE:HD13	1:F:410:SER:C	2.31	0.55
1:K:389:ARG:HA	1:K:392:ILE:CG1	2.29	0.55
1:C:379:LYS:HG2	1:C:379:LYS:O	2.05	0.55
1:C:387:VAL:HG21	1:C:390:GLU:HB2	1.88	0.55
1:E:348:PHE:CZ	1:G:664:HIS:HD2	2.13	0.55
1:F:499:ILE:HG23	1:F:602:ILE:HB	1.87	0.55
1:F:647:ARG:HD3	1:F:649:GLN:HG3	1.88	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:532:GLU:N	1:K:532:GLU:OE2	2.39	0.55
1:K:560:SER:N	1:K:561:GLY:HA2	2.20	0.55
1:L:347:THR:O	1:L:348:PHE:C	2.46	0.55
1:A:387:VAL:CG2	1:A:390:GLU:CD	2.79	0.55
1:C:342:ALA:HB1	1:C:362:THR:CG2	2.37	0.55
1:E:525:ASN:HB3	1:E:526:PRO:CD	2.17	0.55
1:G:342:ALA:HB1	1:G:362:THR:CG2	2.37	0.55
1:G:385:THR:CB	1:G:389:ARG:H	2.17	0.55
1:G:448:TYR:CD1	1:G:642:PHE:HB2	2.41	0.55
1:G:526:PRO:O	1:G:528:THR:CG2	2.55	0.55
1:I:437:GLU:C	1:I:439:GLN:H	2.15	0.55
1:J:499:ILE:HG23	1:J:602:ILE:HB	1.87	0.55
1:J:647:ARG:HD3	1:J:649:GLN:HG3	1.88	0.55
1:A:437:GLU:C	1:A:439:GLN:H	2.15	0.55
1:B:365:ASP:O	1:B:367:THR:N	2.40	0.55
1:D:648:PRO:HD2	1:D:649:GLN:HG2	1.89	0.55
1:F:534:VAL:O	1:F:534:VAL:HG12	2.07	0.55
1:G:358:GLN:OE1	1:G:377:LYS:HE3	2.07	0.55
1:I:454:ILE:HG22	1:I:455:PHE:N	2.20	0.55
1:K:631:VAL:CG2	1:L:476:ILE:CG2	2.72	0.55
1:L:648:PRO:HD2	1:L:649:GLN:HG2	1.89	0.55
1:B:648:PRO:HD2	1:B:649:GLN:HG2	1.89	0.55
1:G:437:GLU:C	1:G:439:GLN:H	2.15	0.55
1:G:532:GLU:N	1:G:532:GLU:OE2	2.39	0.55
1:H:365:ASP:OD1	1:H:367:THR:CG2	2.55	0.55
1:L:658:ILE:HD13	1:L:658:ILE:N	2.21	0.55
1:A:403:PRO:N	1:C:366:SER:HB3	2.21	0.54
1:B:647:ARG:HD3	1:B:649:GLN:HG3	1.88	0.54
1:C:532:GLU:N	1:C:532:GLU:OE2	2.40	0.54
1:C:560:SER:N	1:C:561:GLY:HA2	2.20	0.54
1:D:596:ASN:HD22	1:D:599:ALA:H	1.54	0.54
1:G:631:VAL:CG2	1:H:476:ILE:HG22	2.34	0.54
1:H:356:ILE:O	1:H:379:LYS:HB3	2.07	0.54
1:I:385:THR:CG2	1:I:389:ARG:C	2.73	0.54
1:K:342:ALA:HB1	1:K:362:THR:CG2	2.37	0.54
1:L:365:ASP:OD1	1:L:367:THR:CG2	2.55	0.54
1:L:365:ASP:CG	1:L:367:THR:HG21	2.31	0.54
1:L:365:ASP:O	1:L:367:THR:N	2.40	0.54
1:L:409:ILE:HD13	1:L:410:SER:C	2.31	0.54
1:L:499:ILE:HG23	1:L:602:ILE:HB	1.88	0.54
1:A:532:GLU:N	1:A:532:GLU:OE2	2.39	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:605:ASN:HB3	1:A:608:LYS:HG3	1.89	0.54
1:C:526:PRO:O	1:C:528:THR:CG2	2.55	0.54
1:G:389:ARG:HA	1:G:392:ILE:CG1	2.29	0.54
1:J:359:ALA:O	1:J:376:ALA:HA	2.06	0.54
1:K:437:GLU:C	1:K:439:GLN:H	2.15	0.54
1:L:509:SER:OG	1:L:626:ASP:N	2.32	0.54
1:A:342:ALA:HB1	1:A:362:THR:CG2	2.37	0.54
1:C:605:ASN:HB3	1:C:608:LYS:HG3	1.89	0.54
1:E:342:ALA:HB1	1:E:362:THR:CG2	2.37	0.54
1:E:385:THR:CG2	1:E:389:ARG:C	2.73	0.54
1:E:532:GLU:N	1:E:532:GLU:OE2	2.39	0.54
1:F:347:THR:O	1:F:348:PHE:C	2.46	0.54
1:F:658:ILE:HD13	1:F:658:ILE:N	2.21	0.54
1:H:658:ILE:HD13	1:H:658:ILE:N	2.21	0.54
1:L:356:ILE:O	1:L:379:LYS:HB3	2.06	0.54
1:A:389:ARG:HA	1:A:392:ILE:CG1	2.29	0.54
1:C:385:THR:CG2	1:C:389:ARG:C	2.73	0.54
1:D:567:GLU:C	1:D:568:ASN:OD1	2.51	0.54
1:D:657:PRO:O	1:D:658:ILE:HD12	2.08	0.54
1:F:359:ALA:O	1:F:376:ALA:HA	2.06	0.54
1:H:365:ASP:O	1:H:367:THR:N	2.40	0.54
1:J:534:VAL:O	1:J:534:VAL:HG12	2.07	0.54
1:K:631:VAL:CG2	1:L:476:ILE:HG22	2.34	0.54
1:L:534:VAL:O	1:L:534:VAL:HG12	2.07	0.54
1:B:427:ASN:ND2	1:B:427:ASN:N	2.54	0.54
1:B:567:GLU:C	1:B:568:ASN:OD1	2.51	0.54
1:E:358:GLN:OE1	1:E:377:LYS:HE3	2.07	0.54
1:E:385:THR:CB	1:E:389:ARG:H	2.17	0.54
1:F:567:GLU:C	1:F:568:ASN:OD1	2.51	0.54
1:G:635:ARG:HH22	1:H:661:LEU:CD1	2.15	0.54
1:I:631:VAL:CG2	1:J:476:ILE:HG22	2.34	0.54
1:J:365:ASP:CG	1:J:367:THR:HG21	2.31	0.54
1:J:657:PRO:O	1:J:658:ILE:HD12	2.08	0.54
1:B:356:ILE:O	1:B:379:LYS:HB3	2.07	0.54
1:C:437:GLU:C	1:C:439:GLN:H	2.15	0.54
1:D:365:ASP:O	1:D:367:THR:N	2.40	0.54
1:E:560:SER:N	1:E:561:GLY:HA2	2.20	0.54
1:H:647:ARG:HD3	1:H:649:GLN:HG3	1.88	0.54
1:I:353:PHE:CE2	1:I:395:TYR:CD2	2.86	0.54
1:K:526:PRO:O	1:K:528:THR:CG2	2.55	0.54
1:K:605:ASN:HB3	1:K:608:LYS:HG3	1.89	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:358:GLN:OE1	1:C:377:LYS:HE3	2.07	0.54
1:F:365:ASP:O	1:F:367:THR:N	2.40	0.54
1:H:359:ALA:O	1:H:376:ALA:HA	2.07	0.54
1:I:526:PRO:O	1:I:528:THR:CG2	2.55	0.54
1:L:647:ARG:HD3	1:L:649:GLN:HG3	1.88	0.54
1:B:534:VAL:O	1:B:534:VAL:HG12	2.07	0.54
1:E:437:GLU:C	1:E:439:GLN:H	2.15	0.54
1:H:657:PRO:O	1:H:658:ILE:HD12	2.08	0.54
1:J:648:PRO:HD2	1:J:649:GLN:HG2	1.89	0.54
1:L:657:PRO:O	1:L:658:ILE:HD12	2.08	0.54
1:A:536:TYR:HB2	1:A:586:ARG:NH1	2.23	0.54
1:B:365:ASP:OD1	1:B:367:THR:CG2	2.55	0.54
1:D:595:ILE:HD12	1:D:595:ILE:N	2.21	0.54
1:I:389:ARG:CA	1:I:392:ILE:HG12	2.28	0.54
1:B:359:ALA:HB1	1:B:455:PHE:HE2	1.73	0.54
1:E:635:ARG:HG3	1:E:636:ASP:N	2.19	0.54
1:F:356:ILE:O	1:F:379:LYS:HB3	2.06	0.54
1:F:409:ILE:CD1	1:F:410:SER:O	2.50	0.54
1:G:387:VAL:CG2	1:G:390:GLU:CD	2.79	0.54
1:G:605:ASN:HB3	1:G:608:LYS:HG3	1.89	0.54
1:G:635:ARG:HG3	1:G:636:ASP:N	2.19	0.54
1:H:596:ASN:ND2	1:H:599:ALA:H	2.06	0.54
1:L:567:GLU:C	1:L:568:ASN:OD1	2.51	0.54
1:A:526:PRO:O	1:A:528:THR:CG2	2.55	0.53
1:C:389:ARG:HA	1:C:392:ILE:CG1	2.29	0.53
1:D:365:ASP:OD1	1:D:367:THR:CG2	2.55	0.53
1:D:396:LEU:HD22	1:D:406:PRO:HG2	1.90	0.53
1:D:534:VAL:O	1:D:534:VAL:HG12	2.07	0.53
1:I:631:VAL:CG2	1:J:476:ILE:CG2	2.72	0.53
1:J:365:ASP:OD1	1:J:367:THR:CG2	2.55	0.53
1:J:658:ILE:HD13	1:J:658:ILE:N	2.21	0.53
1:K:412:ASN:O	1:K:638:SER:HA	2.08	0.53
1:B:526:PRO:HA	1:B:529:GLY:O	2.09	0.53
1:B:657:PRO:O	1:B:658:ILE:HD12	2.08	0.53
1:D:359:ALA:HB1	1:D:455:PHE:HE2	1.73	0.53
1:D:526:PRO:HA	1:D:529:GLY:O	2.09	0.53
1:D:596:ASN:HB3	1:D:601:VAL:HG23	1.90	0.53
1:F:359:ALA:HB1	1:F:455:PHE:HE2	1.73	0.53
1:F:526:PRO:HA	1:F:529:GLY:O	2.08	0.53
1:H:567:GLU:C	1:H:568:ASN:OD1	2.51	0.53
1:J:365:ASP:O	1:J:367:THR:N	2.40	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:531:GLU:C	1:A:532:GLU:OE1	2.52	0.53
1:F:559:ALA:O	1:F:561:GLY:N	2.42	0.53
1:F:661:LEU:C	1:F:663:HIS:N	2.61	0.53
1:H:559:ALA:O	1:H:561:GLY:N	2.41	0.53
1:I:353:PHE:O	1:I:354:GLY:C	2.48	0.53
1:I:605:ASN:HB3	1:I:608:LYS:HG3	1.89	0.53
1:I:635:ARG:HH22	1:J:661:LEU:CD1	2.14	0.53
1:K:382:LEU:HD12	1:K:647:ARG:HH21	1.74	0.53
1:K:536:TYR:HB2	1:K:586:ARG:NH1	2.23	0.53
1:L:526:PRO:HA	1:L:529:GLY:O	2.08	0.53
1:L:559:ALA:O	1:L:561:GLY:N	2.42	0.53
1:A:631:VAL:CG2	1:B:476:ILE:CG2	2.72	0.53
1:B:542:SER:HB2	1:B:595:ILE:HG12	1.91	0.53
1:E:412:ASN:O	1:E:638:SER:HA	2.09	0.53
1:E:536:TYR:HB2	1:E:586:ARG:NH1	2.23	0.53
1:F:365:ASP:OD1	1:F:367:THR:CG2	2.55	0.53
1:H:526:PRO:HA	1:H:529:GLY:O	2.09	0.53
1:H:534:VAL:HG12	1:H:534:VAL:O	2.07	0.53
1:H:596:ASN:HB3	1:H:601:VAL:HG23	1.90	0.53
1:I:412:ASN:O	1:I:638:SER:HA	2.08	0.53
1:K:531:GLU:C	1:K:532:GLU:OE1	2.52	0.53
1:K:635:ARG:HH22	1:L:661:LEU:CD1	2.15	0.53
1:L:542:SER:HB2	1:L:595:ILE:HG12	1.91	0.53
1:A:382:LEU:HD12	1:A:647:ARG:HH21	1.74	0.53
1:B:476:ILE:HG22	1:B:477:GLY:N	2.18	0.53
1:E:605:ASN:HB3	1:E:608:LYS:HG3	1.89	0.53
1:G:428:LYS:HD2	1:G:473:HIS:HD2	1.74	0.53
1:H:648:PRO:HD2	1:H:649:GLN:HG2	1.89	0.53
1:I:385:THR:C	1:I:387:VAL:N	2.66	0.53
1:J:542:SER:HB2	1:J:595:ILE:HG12	1.91	0.53
1:J:596:ASN:ND2	1:J:599:ALA:H	2.06	0.53
1:C:382:LEU:HD12	1:C:647:ARG:HH21	1.74	0.53
1:D:400:ASN:H	1:E:664:HIS:HB2	1.74	0.53
1:F:657:PRO:O	1:F:658:ILE:HD12	2.08	0.53
1:G:353:PHE:O	1:G:354:GLY:C	2.48	0.53
1:G:412:ASN:O	1:G:638:SER:HA	2.08	0.53
1:J:567:GLU:C	1:J:568:ASN:OD1	2.51	0.53
1:A:385:THR:C	1:A:387:VAL:N	2.66	0.53
1:A:428:LYS:HD2	1:A:473:HIS:HD2	1.74	0.53
1:A:619:GLN:HA	1:A:619:GLN:HE21	1.74	0.53
1:B:661:LEU:C	1:B:661:LEU:HD22	2.30	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:619:GLN:HA	1:C:619:GLN:HE21	1.74	0.53
1:D:559:ALA:O	1:D:561:GLY:N	2.41	0.53
1:F:453:GLU:O	1:F:454:ILE:HG13	2.09	0.53
1:G:343:THR:O	1:G:347:THR:HG23	2.09	0.53
1:G:536:TYR:HB2	1:G:586:ARG:NH1	2.23	0.53
1:H:409:ILE:CD1	1:H:410:SER:O	2.50	0.53
1:I:635:ARG:HG3	1:I:636:ASP:N	2.19	0.53
1:L:348:PHE:O	1:L:351:GLU:N	2.21	0.53
1:A:631:VAL:CG2	1:B:476:ILE:HG22	2.34	0.53
1:B:596:ASN:ND2	1:B:599:ALA:H	2.06	0.53
1:B:596:ASN:HB3	1:B:601:VAL:HG23	1.90	0.53
1:C:536:TYR:HB2	1:C:586:ARG:NH1	2.23	0.53
1:D:343:THR:HB	1:D:347:THR:CG2	2.39	0.53
1:D:542:SER:HB2	1:D:595:ILE:HG12	1.91	0.53
1:D:596:ASN:ND2	1:D:599:ALA:H	2.06	0.53
1:F:648:PRO:HD2	1:F:649:GLN:HG2	1.89	0.53
1:G:385:THR:CG2	1:G:389:ARG:C	2.73	0.53
1:I:343:THR:O	1:I:347:THR:HG23	2.09	0.53
1:J:559:ALA:O	1:J:561:GLY:N	2.41	0.53
1:K:428:LYS:HD2	1:K:473:HIS:HD2	1.74	0.53
1:C:412:ASN:O	1:C:638:SER:HA	2.08	0.53
1:E:454:ILE:HG22	1:E:455:PHE:N	2.20	0.53
1:G:531:GLU:C	1:G:532:GLU:OE1	2.52	0.53
1:H:453:GLU:O	1:H:454:ILE:HG13	2.09	0.53
1:I:382:LEU:HD12	1:I:647:ARG:HH21	1.74	0.53
1:I:536:TYR:HB2	1:I:586:ARG:NH1	2.23	0.53
1:B:396:LEU:HD22	1:B:406:PRO:HG2	1.90	0.53
1:D:547:SER:O	1:D:548:LYS:HG2	2.09	0.53
1:E:651:LEU:CD2	1:E:653:ILE:HG13	2.39	0.53
1:F:596:ASN:HB3	1:F:601:VAL:HG23	1.90	0.53
1:H:343:THR:HB	1:H:347:THR:CG2	2.39	0.53
1:J:343:THR:HB	1:J:347:THR:CG2	2.39	0.53
1:A:343:THR:O	1:A:347:THR:HG23	2.09	0.52
1:B:440:ILE:HD11	1:B:653:ILE:HD13	1.91	0.52
1:B:547:SER:O	1:B:548:LYS:HG2	2.09	0.52
1:B:559:ALA:O	1:B:561:GLY:N	2.41	0.52
1:E:631:VAL:CG2	1:F:476:ILE:HG22	2.34	0.52
1:F:343:THR:HB	1:F:347:THR:CG2	2.39	0.52
1:F:409:ILE:HD12	1:F:410:SER:N	2.17	0.52
1:G:382:LEU:HD12	1:G:647:ARG:HH21	1.74	0.52
1:G:385:THR:CB	1:G:387:VAL:CG2	2.88	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:619:GLN:HE21	1:G:619:GLN:HA	1.74	0.52
1:I:385:THR:CB	1:I:387:VAL:CG2	2.88	0.52
1:I:525:ASN:HB3	1:I:526:PRO:CD	2.17	0.52
1:L:343:THR:HB	1:L:347:THR:CG2	2.39	0.52
1:L:408:ILE:HD12	1:L:409:ILE:CA	2.40	0.52
1:C:452:VAL:HG22	1:C:458:SER:O	2.10	0.52
1:E:556:GLY:HA3	1:E:589:TYR:HD1	1.74	0.52
1:F:345:TYR:C	1:F:345:TYR:HD1	2.18	0.52
1:G:353:PHE:CE2	1:G:395:TYR:CD2	2.86	0.52
1:H:542:SER:HB2	1:H:595:ILE:HG12	1.91	0.52
1:I:491:PHE:CZ	1:I:616:PHE:HB2	2.45	0.52
1:K:343:THR:O	1:K:347:THR:HG23	2.09	0.52
1:L:547:SER:O	1:L:548:LYS:HG2	2.09	0.52
1:C:631:VAL:CG2	1:D:476:ILE:HG22	2.34	0.52
1:D:400:ASN:CB	1:E:662:GLU:HG2	2.39	0.52
1:D:453:GLU:O	1:D:454:ILE:HG13	2.09	0.52
1:E:403:PRO:HB3	1:H:366:SER:C	2.29	0.52
1:H:565:GLU:CB	1:H:570:GLN:NE2	2.55	0.52
1:I:619:GLN:HA	1:I:619:GLN:HE21	1.74	0.52
1:I:651:LEU:CD2	1:I:653:ILE:HG13	2.39	0.52
1:J:453:GLU:O	1:J:454:ILE:HG13	2.09	0.52
1:J:568:ASN:OD1	1:J:569:ILE:N	2.30	0.52
1:K:556:GLY:HA3	1:K:589:TYR:HD1	1.74	0.52
1:K:619:GLN:HA	1:K:619:GLN:HE21	1.74	0.52
1:D:440:ILE:HD11	1:D:653:ILE:HD13	1.91	0.52
1:E:382:LEU:HD12	1:E:647:ARG:HH21	1.74	0.52
1:E:531:GLU:C	1:E:532:GLU:OE1	2.52	0.52
1:F:408:ILE:HD12	1:F:409:ILE:CA	2.40	0.52
1:F:476:ILE:HG22	1:F:477:GLY:N	2.18	0.52
1:F:547:SER:O	1:F:548:LYS:HG2	2.09	0.52
1:I:428:LYS:HD2	1:I:473:HIS:HD2	1.74	0.52
1:J:359:ALA:HB1	1:J:455:PHE:HE2	1.73	0.52
1:L:440:ILE:HD11	1:L:653:ILE:HD13	1.91	0.52
1:L:596:ASN:ND2	1:L:599:ALA:H	2.07	0.52
1:A:635:ARG:HH22	1:B:661:LEU:CD1	2.14	0.52
1:C:531:GLU:C	1:C:532:GLU:OE1	2.52	0.52
1:C:651:LEU:CD2	1:C:653:ILE:HG13	2.39	0.52
1:E:343:THR:O	1:E:347:THR:HG23	2.09	0.52
1:E:619:GLN:HA	1:E:619:GLN:HE21	1.74	0.52
1:F:472:ASP:OD2	1:F:474:SER:HB3	2.10	0.52
1:F:542:SER:HB2	1:F:595:ILE:HG12	1.91	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:452:VAL:HG22	1:G:458:SER:O	2.10	0.52
1:G:651:LEU:CD2	1:G:653:ILE:HG13	2.39	0.52
1:I:400:ASN:CG	1:K:664:HIS:CE1	2.75	0.52
1:J:472:ASP:OD2	1:J:474:SER:HB3	2.10	0.52
1:J:596:ASN:HB3	1:J:601:VAL:HG23	1.90	0.52
1:L:359:ALA:HB1	1:L:455:PHE:HE2	1.73	0.52
1:L:453:GLU:O	1:L:454:ILE:HG13	2.09	0.52
1:L:486:ARG:HG2	1:L:499:ILE:CD1	2.40	0.52
1:A:385:THR:CB	1:A:387:VAL:CG2	2.88	0.52
1:A:412:ASN:O	1:A:638:SER:HA	2.08	0.52
1:C:343:THR:O	1:C:347:THR:HG23	2.09	0.52
1:D:660:GLN:CA	1:D:661:LEU:HB2	2.26	0.52
1:E:353:PHE:C	1:E:355:SER:N	2.65	0.52
1:F:486:ARG:HG2	1:F:499:ILE:CD1	2.40	0.52
1:H:396:LEU:HD22	1:H:406:PRO:HG2	1.90	0.52
1:K:385:THR:CB	1:K:387:VAL:CG2	2.88	0.52
1:L:661:LEU:C	1:L:663:HIS:N	2.61	0.52
1:A:491:PHE:CZ	1:A:616:PHE:HB2	2.45	0.52
1:C:385:THR:C	1:C:387:VAL:H	2.16	0.52
1:D:486:ARG:HG2	1:D:499:ILE:CD1	2.40	0.52
1:F:596:ASN:ND2	1:F:599:ALA:H	2.06	0.52
1:L:595:ILE:HD12	1:L:595:ILE:N	2.21	0.52
1:A:385:THR:C	1:A:387:VAL:H	2.16	0.52
1:B:408:ILE:HD12	1:B:409:ILE:CA	2.40	0.52
1:C:533:ASP:O	1:C:535:LEU:HD23	2.10	0.52
1:C:556:GLY:HA3	1:C:589:TYR:HD1	1.74	0.52
1:E:342:ALA:CB	1:E:362:THR:O	2.58	0.52
1:F:348:PHE:O	1:F:350:SER:N	2.43	0.52
1:F:382:LEU:HD12	1:F:646:LEU:HD13	1.92	0.52
1:F:440:ILE:HD11	1:F:653:ILE:HD13	1.91	0.52
1:F:660:GLN:CA	1:F:661:LEU:HB2	2.26	0.52
1:G:448:TYR:CE1	1:G:642:PHE:HB2	2.45	0.52
1:G:491:PHE:CZ	1:G:616:PHE:HB2	2.45	0.52
1:J:526:PRO:HA	1:J:529:GLY:O	2.09	0.52
1:L:596:ASN:HB3	1:L:601:VAL:HG23	1.90	0.52
1:A:651:LEU:CD2	1:A:653:ILE:HG13	2.39	0.52
1:B:343:THR:HB	1:B:347:THR:CG2	2.39	0.52
1:B:453:GLU:O	1:B:454:ILE:HG13	2.09	0.52
1:E:386:THR:C	1:E:388:GLN:N	2.67	0.52
1:I:531:GLU:C	1:I:532:GLU:OE1	2.52	0.52
1:J:348:PHE:O	1:J:350:SER:N	2.43	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:486:ARG:HG2	1:J:499:ILE:CD1	2.40	0.52
1:K:554:ILE:HG22	1:K:591:VAL:HA	1.92	0.52
1:A:342:ALA:CB	1:A:362:THR:O	2.58	0.52
1:B:472:ASP:OD2	1:B:474:SER:HB3	2.10	0.52
1:B:628:THR:HG22	1:B:629:ASP:OD1	2.10	0.52
1:C:385:THR:C	1:C:387:VAL:N	2.66	0.52
1:E:428:LYS:HD2	1:E:473:HIS:HD2	1.74	0.52
1:E:448:TYR:CE1	1:E:642:PHE:HB2	2.45	0.52
1:F:661:LEU:O	1:F:661:LEU:CG	2.58	0.52
1:H:359:ALA:HB1	1:H:455:PHE:HE2	1.73	0.52
1:J:396:LEU:HD22	1:J:406:PRO:HG2	1.90	0.52
1:J:409:ILE:CD1	1:J:410:SER:O	2.50	0.52
1:A:342:ALA:HA	1:A:362:THR:HG21	1.93	0.51
1:B:348:PHE:O	1:B:350:SER:N	2.43	0.51
1:B:523:VAL:HG11	1:B:532:GLU:HB2	1.92	0.51
1:C:342:ALA:CB	1:C:362:THR:O	2.58	0.51
1:C:448:TYR:CE1	1:C:642:PHE:HB2	2.45	0.51
1:C:491:PHE:CZ	1:C:616:PHE:HB2	2.45	0.51
1:D:427:ASN:ND2	1:D:427:ASN:N	2.54	0.51
1:D:523:VAL:HG11	1:D:532:GLU:HB2	1.92	0.51
1:E:491:PHE:CZ	1:E:616:PHE:HB2	2.45	0.51
1:G:385:THR:C	1:G:387:VAL:N	2.66	0.51
1:G:556:GLY:HA3	1:G:589:TYR:HD1	1.74	0.51
1:H:382:LEU:HD12	1:H:646:LEU:HD13	1.92	0.51
1:H:486:ARG:HG2	1:H:499:ILE:CD1	2.40	0.51
1:I:448:TYR:CE1	1:I:642:PHE:HB2	2.45	0.51
1:K:491:PHE:CZ	1:K:616:PHE:HB2	2.45	0.51
1:L:348:PHE:O	1:L:350:SER:N	2.43	0.51
1:B:365:ASP:HB3	1:B:368:LYS:HB2	1.92	0.51
1:C:635:ARG:HG3	1:C:636:ASP:N	2.19	0.51
1:E:452:VAL:HG22	1:E:458:SER:O	2.10	0.51
1:E:554:ILE:HG22	1:E:591:VAL:HA	1.92	0.51
1:I:533:ASP:O	1:I:535:LEU:HD23	2.10	0.51
1:J:408:ILE:HD12	1:J:409:ILE:CA	2.40	0.51
1:K:462:SER:HB3	1:L:478:SER:O	2.11	0.51
1:B:486:ARG:HG2	1:B:499:ILE:CD1	2.40	0.51
1:E:462:SER:HB3	1:F:478:SER:O	2.11	0.51
1:E:533:ASP:O	1:E:535:LEU:HD23	2.10	0.51
1:I:342:ALA:HA	1:I:362:THR:HG21	1.93	0.51
1:I:554:ILE:HG22	1:I:591:VAL:HA	1.92	0.51
1:J:547:SER:O	1:J:548:LYS:HG2	2.10	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:567:GLU:CA	1:J:570:GLN:OE1	2.56	0.51
1:J:595:ILE:HD12	1:J:595:ILE:N	2.21	0.51
1:K:533:ASP:O	1:K:535:LEU:HD23	2.10	0.51
1:C:522:LYS:N	1:C:522:LYS:CD	2.72	0.51
1:F:628:THR:HG22	1:F:629:ASP:OD1	2.10	0.51
1:H:348:PHE:O	1:H:350:SER:N	2.43	0.51
1:H:440:ILE:HD11	1:H:653:ILE:HD13	1.91	0.51
1:H:547:SER:O	1:H:548:LYS:HG2	2.09	0.51
1:I:342:ALA:CB	1:I:362:THR:O	2.58	0.51
1:J:440:ILE:HD11	1:J:653:ILE:HD13	1.91	0.51
1:L:472:ASP:OD2	1:L:474:SER:HB3	2.10	0.51
1:B:658:ILE:CD1	1:B:658:ILE:N	2.73	0.51
1:C:428:LYS:HD2	1:C:473:HIS:HD2	1.74	0.51
1:E:353:PHE:CE2	1:E:395:TYR:CD2	2.86	0.51
1:G:462:SER:HB3	1:H:478:SER:O	2.11	0.51
1:G:492:TYR:O	1:G:606:ILE:HG13	2.11	0.51
1:G:533:ASP:O	1:G:535:LEU:HD23	2.10	0.51
1:H:443:LYS:HG3	1:H:471:ALA:HB2	1.93	0.51
1:H:476:ILE:HG22	1:H:477:GLY:N	2.18	0.51
1:K:342:ALA:HA	1:K:362:THR:HG21	1.93	0.51
1:A:448:TYR:CE1	1:A:642:PHE:HB2	2.45	0.51
1:A:457:SER:N	1:A:634:THR:HG23	2.26	0.51
1:A:462:SER:HB3	1:B:478:SER:O	2.11	0.51
1:C:492:TYR:O	1:C:606:ILE:HG13	2.11	0.51
1:C:554:ILE:HG22	1:C:591:VAL:HA	1.92	0.51
1:F:443:LYS:HG3	1:F:471:ALA:HB2	1.93	0.51
1:F:565:GLU:CB	1:F:570:GLN:NE2	2.55	0.51
1:I:452:VAL:HG22	1:I:458:SER:O	2.10	0.51
1:I:462:SER:HB3	1:J:478:SER:O	2.11	0.51
1:K:353:PHE:O	1:K:354:GLY:C	2.48	0.51
1:L:365:ASP:HB3	1:L:368:LYS:HB2	1.92	0.51
1:A:403:PRO:N	1:C:366:SER:CB	2.73	0.51
1:B:595:ILE:HD12	1:B:595:ILE:N	2.21	0.51
1:C:353:PHE:C	1:C:355:SER:N	2.65	0.51
1:C:385:THR:CB	1:C:387:VAL:CG2	2.88	0.51
1:D:348:PHE:O	1:D:350:SER:N	2.43	0.51
1:D:408:ILE:HD12	1:D:409:ILE:CA	2.40	0.51
1:D:660:GLN:HA	1:D:661:LEU:HB3	1.90	0.51
1:J:628:THR:HG22	1:J:629:ASP:OD1	2.10	0.51
1:K:385:THR:C	1:K:387:VAL:H	2.16	0.51
1:K:452:VAL:HG22	1:K:458:SER:O	2.10	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:340:VAL:O	1:A:341:THR:O	2.29	0.51
1:A:492:TYR:O	1:A:606:ILE:HG13	2.11	0.51
1:A:533:ASP:O	1:A:535:LEU:HD23	2.10	0.51
1:A:554:ILE:HG22	1:A:591:VAL:HA	1.92	0.51
1:A:651:LEU:HD23	1:A:652:THR:N	2.26	0.51
1:D:472:ASP:OD2	1:D:474:SER:HB3	2.10	0.51
1:D:492:TYR:CE1	1:D:606:ILE:HB	2.46	0.51
1:E:340:VAL:O	1:E:341:THR:O	2.29	0.51
1:F:396:LEU:HD22	1:F:406:PRO:HG2	1.90	0.51
1:H:402:ALA:H	1:H:403:PRO:CD	2.24	0.51
1:J:661:LEU:O	1:J:661:LEU:CG	2.58	0.51
1:K:448:TYR:CE1	1:K:642:PHE:HB2	2.45	0.51
1:K:492:TYR:O	1:K:606:ILE:HG13	2.11	0.51
1:K:651:LEU:HD23	1:K:652:THR:N	2.26	0.51
1:L:402:ALA:H	1:L:403:PRO:CD	2.24	0.51
1:L:523:VAL:HG11	1:L:532:GLU:HB2	1.92	0.51
1:B:366:SER:HA	1:K:403:PRO:HG3	1.89	0.51
1:C:340:VAL:O	1:C:341:THR:O	2.29	0.51
1:C:651:LEU:HD23	1:C:652:THR:N	2.26	0.51
1:F:365:ASP:HB3	1:F:368:LYS:HB2	1.93	0.51
1:H:408:ILE:HD12	1:H:409:ILE:CA	2.40	0.51
1:H:628:THR:HG22	1:H:629:ASP:OD1	2.10	0.51
1:J:361:GLN:HE22	1:J:456:ASN:HD21	1.57	0.51
1:K:353:PHE:CE2	1:K:395:TYR:CD2	2.86	0.51
1:K:651:LEU:CD2	1:K:653:ILE:HG13	2.39	0.51
1:A:386:THR:C	1:A:388:GLN:N	2.67	0.51
1:A:452:VAL:HG22	1:A:458:SER:O	2.10	0.51
1:D:402:ALA:H	1:D:403:PRO:CD	2.24	0.51
1:E:385:THR:C	1:E:387:VAL:N	2.66	0.51
1:F:396:LEU:HD13	1:F:405:THR:O	2.11	0.51
1:F:567:GLU:O	1:F:568:ASN:OD1	2.29	0.51
1:G:342:ALA:CB	1:G:362:THR:O	2.58	0.51
1:G:554:ILE:HG22	1:G:591:VAL:HA	1.92	0.51
1:G:559:ALA:O	1:G:561:GLY:CA	2.34	0.51
1:H:361:GLN:HE22	1:H:456:ASN:HD21	1.57	0.51
1:H:595:ILE:HD12	1:H:595:ILE:N	2.21	0.51
1:H:661:LEU:C	1:H:661:LEU:HD22	2.30	0.51
1:K:340:VAL:O	1:K:341:THR:O	2.29	0.51
1:L:396:LEU:HD22	1:L:406:PRO:HG2	1.90	0.51
1:L:628:THR:HG22	1:L:629:ASP:OD1	2.10	0.51
1:B:382:LEU:HD12	1:B:646:LEU:HD13	1.92	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:443:LYS:HG3	1:B:471:ALA:HB2	1.93	0.50
1:B:661:LEU:O	1:B:661:LEU:CG	2.58	0.50
1:D:361:GLN:HE22	1:D:456:ASN:HD21	1.57	0.50
1:D:628:THR:HG22	1:D:629:ASP:OD1	2.10	0.50
1:F:402:ALA:H	1:F:403:PRO:CD	2.24	0.50
1:F:607:ALA:C	1:F:609:ILE:H	2.20	0.50
1:H:492:TYR:CE1	1:H:606:ILE:HB	2.46	0.50
1:H:523:VAL:HG11	1:H:532:GLU:HB2	1.92	0.50
1:J:382:LEU:HD12	1:J:646:LEU:HD13	1.92	0.50
1:K:342:ALA:CB	1:K:362:THR:O	2.58	0.50
1:K:386:THR:C	1:K:388:GLN:N	2.67	0.50
1:L:382:LEU:HD12	1:L:646:LEU:HD13	1.92	0.50
1:L:408:ILE:O	1:L:409:ILE:HG23	2.09	0.50
1:A:387:VAL:HB	1:A:390:GLU:CD	2.37	0.50
1:A:556:GLY:HA3	1:A:589:TYR:HD1	1.74	0.50
1:B:402:ALA:H	1:B:403:PRO:CD	2.24	0.50
1:C:387:VAL:HB	1:C:390:GLU:CD	2.36	0.50
1:D:510:MET:HB3	1:D:542:SER:HB3	1.94	0.50
1:D:607:ALA:C	1:D:609:ILE:H	2.20	0.50
1:E:492:TYR:O	1:E:606:ILE:HG13	2.11	0.50
1:F:565:GLU:OE2	1:F:590:TYR:OH	2.28	0.50
1:G:340:VAL:O	1:G:341:THR:O	2.29	0.50
1:H:472:ASP:OD2	1:H:474:SER:HB3	2.10	0.50
1:I:466:THR:HB	1:J:466:THR:HB	1.94	0.50
1:I:651:LEU:HD23	1:I:652:THR:N	2.26	0.50
1:J:492:TYR:CE1	1:J:606:ILE:HB	2.46	0.50
1:K:457:SER:N	1:K:634:THR:HG23	2.26	0.50
1:L:443:LYS:HG3	1:L:471:ALA:HB2	1.93	0.50
1:L:660:GLN:HA	1:L:661:LEU:HB3	1.90	0.50
1:A:342:ALA:HB1	1:A:362:THR:HG22	1.94	0.50
1:A:495:PRO:HD2	1:A:499:ILE:HG22	1.93	0.50
1:D:365:ASP:HB3	1:D:368:LYS:HB2	1.93	0.50
1:D:396:LEU:HD13	1:D:405:THR:O	2.11	0.50
1:D:423:THR:HB	1:D:656:GLU:HB3	1.94	0.50
1:D:661:LEU:O	1:D:661:LEU:CG	2.58	0.50
1:F:361:GLN:HE22	1:F:456:ASN:HD21	1.57	0.50
1:G:342:ALA:HA	1:G:362:THR:HG21	1.93	0.50
1:H:396:LEU:HD13	1:H:405:THR:O	2.11	0.50
1:H:565:GLU:OE2	1:H:590:TYR:OH	2.28	0.50
1:H:567:GLU:O	1:H:568:ASN:OD1	2.29	0.50
1:I:403:PRO:CD	1:L:366:SER:HA	2.28	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:423:THR:HB	1:L:656:GLU:HB3	1.94	0.50
1:L:510:MET:HB3	1:L:542:SER:HB3	1.93	0.50
1:A:348:PHE:HZ	1:A:400:ASN:HD21	1.58	0.50
1:A:391:ASP:C	1:A:393:LYS:H	2.20	0.50
1:B:492:TYR:CE1	1:B:606:ILE:HB	2.46	0.50
1:B:567:GLU:O	1:B:568:ASN:OD1	2.29	0.50
1:C:340:VAL:O	1:C:341:THR:OG1	2.30	0.50
1:C:391:ASP:C	1:C:393:LYS:H	2.20	0.50
1:D:342:ALA:O	1:D:346:ASP:OD1	2.30	0.50
1:D:382:LEU:HD12	1:D:646:LEU:HD13	1.92	0.50
1:D:566:ASN:O	1:D:570:GLN:OE1	2.30	0.50
1:E:495:PRO:HD2	1:E:499:ILE:HG22	1.93	0.50
1:I:340:VAL:O	1:I:341:THR:O	2.29	0.50
1:J:566:ASN:O	1:J:568:ASN:OD1	2.30	0.50
1:K:466:THR:HB	1:L:466:THR:HB	1.94	0.50
1:L:661:LEU:O	1:L:661:LEU:CG	2.58	0.50
1:A:517:PHE:HE1	1:A:615:LYS:HD2	1.77	0.50
1:C:342:ALA:HA	1:C:362:THR:HG21	1.93	0.50
1:C:631:VAL:CG2	1:D:476:ILE:CG2	2.72	0.50
1:D:567:GLU:O	1:D:568:ASN:OD1	2.29	0.50
1:D:644:ASN:C	1:D:644:ASN:HD22	2.19	0.50
1:E:466:THR:HB	1:F:466:THR:HB	1.94	0.50
1:E:510:MET:HB3	1:E:542:SER:OG	2.12	0.50
1:F:347:THR:O	1:F:348:PHE:O	2.30	0.50
1:G:391:ASP:C	1:G:393:LYS:H	2.20	0.50
1:H:342:ALA:O	1:H:346:ASP:OD1	2.30	0.50
1:H:607:ALA:C	1:H:609:ILE:H	2.20	0.50
1:J:402:ALA:H	1:J:403:PRO:CD	2.24	0.50
1:J:523:VAL:HG11	1:J:532:GLU:HB2	1.92	0.50
1:K:385:THR:C	1:K:387:VAL:N	2.66	0.50
1:L:566:ASN:O	1:L:568:ASN:OD1	2.30	0.50
1:B:607:ALA:C	1:B:609:ILE:H	2.20	0.50
1:C:387:VAL:O	1:C:388:GLN:OE1	2.30	0.50
1:E:385:THR:CB	1:E:387:VAL:CG2	2.88	0.50
1:E:385:THR:C	1:E:387:VAL:H	2.16	0.50
1:F:523:VAL:HG11	1:F:532:GLU:HB2	1.92	0.50
1:G:651:LEU:HD23	1:G:652:THR:N	2.26	0.50
1:H:567:GLU:CA	1:H:570:GLN:OE1	2.56	0.50
1:H:644:ASN:C	1:H:644:ASN:HD22	2.20	0.50
1:J:347:THR:O	1:J:348:PHE:O	2.30	0.50
1:J:443:LYS:HG3	1:J:471:ALA:HB2	1.93	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:476:ILE:HG22	1:J:477:GLY:N	2.18	0.50
1:J:510:MET:HB3	1:J:542:SER:HB3	1.94	0.50
1:J:644:ASN:C	1:J:644:ASN:HD22	2.20	0.50
1:K:387:VAL:HB	1:K:390:GLU:CD	2.37	0.50
1:K:517:PHE:HE1	1:K:615:LYS:HD2	1.77	0.50
1:B:361:GLN:HE22	1:B:456:ASN:HD21	1.57	0.50
1:B:423:THR:HB	1:B:656:GLU:HB3	1.94	0.50
1:B:448:TYR:O	1:B:453:GLU:HG3	2.12	0.50
1:E:517:PHE:HE1	1:E:615:LYS:HD2	1.77	0.50
1:F:492:TYR:CE1	1:F:606:ILE:HB	2.46	0.50
1:F:566:ASN:O	1:F:570:GLN:CD	2.55	0.50
1:G:510:MET:HB3	1:G:542:SER:OG	2.12	0.50
1:I:517:PHE:HE1	1:I:615:LYS:HD2	1.77	0.50
1:J:345:TYR:C	1:J:345:TYR:HD1	2.18	0.50
1:J:566:ASN:O	1:J:570:GLN:CD	2.55	0.50
1:J:567:GLU:O	1:J:568:ASN:OD1	2.29	0.50
1:J:655:LEU:HD12	1:J:655:LEU:N	2.27	0.50
1:K:387:VAL:O	1:K:388:GLN:OE1	2.30	0.50
1:L:448:TYR:O	1:L:453:GLU:HG3	2.12	0.50
1:A:466:THR:HB	1:B:466:THR:HB	1.94	0.50
1:B:507:ASP:OD1	1:B:545:ARG:HG3	2.12	0.50
1:B:644:ASN:C	1:B:644:ASN:HD22	2.20	0.50
1:C:462:SER:HB3	1:D:478:SER:O	2.11	0.50
1:C:525:ASN:HB3	1:C:526:PRO:CD	2.17	0.50
1:E:342:ALA:HA	1:E:362:THR:HG21	1.93	0.50
1:H:365:ASP:HB3	1:H:368:LYS:HB2	1.92	0.50
1:I:492:TYR:O	1:I:606:ILE:HG13	2.11	0.50
1:I:510:MET:HB3	1:I:542:SER:OG	2.12	0.50
1:J:396:LEU:HD13	1:J:405:THR:O	2.11	0.50
1:K:495:PRO:HD2	1:K:499:ILE:HG22	1.93	0.50
1:B:566:ASN:O	1:B:570:GLN:OE1	2.30	0.50
1:D:655:LEU:HD12	1:D:655:LEU:N	2.27	0.50
1:E:342:ALA:HB2	1:E:362:THR:HG22	1.94	0.50
1:E:607:ALA:O	1:E:609:ILE:O	2.30	0.50
1:G:466:THR:HB	1:H:466:THR:HB	1.94	0.50
1:H:347:THR:O	1:H:348:PHE:O	2.30	0.50
1:I:582:ASN:HB2	1:I:585:GLY:C	2.37	0.50
1:I:607:ALA:O	1:I:609:ILE:O	2.30	0.50
1:J:365:ASP:HB3	1:J:368:LYS:HB2	1.93	0.50
1:K:401:LEU:HG	1:K:402:ALA:N	2.27	0.50
1:L:347:THR:O	1:L:348:PHE:O	2.30	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:492:TYR:CE1	1:L:606:ILE:HB	2.46	0.50
1:L:567:GLU:O	1:L:568:ASN:OD1	2.29	0.50
1:L:644:ASN:C	1:L:644:ASN:HD22	2.20	0.50
1:A:361:GLN:NE2	1:A:361:GLN:HA	2.27	0.49
1:C:466:THR:HB	1:D:466:THR:HB	1.94	0.49
1:C:510:MET:HB3	1:C:542:SER:OG	2.12	0.49
1:D:448:TYR:O	1:D:453:GLU:HG3	2.12	0.49
1:D:566:ASN:O	1:D:568:ASN:OD1	2.30	0.49
1:E:391:ASP:C	1:E:393:LYS:H	2.20	0.49
1:E:582:ASN:HB2	1:E:585:GLY:C	2.37	0.49
1:E:651:LEU:HD23	1:E:652:THR:N	2.26	0.49
1:F:567:GLU:CA	1:F:570:GLN:OE1	2.56	0.49
1:G:582:ASN:HB2	1:G:585:GLY:C	2.37	0.49
1:J:566:ASN:O	1:J:570:GLN:OE1	2.30	0.49
1:K:342:ALA:HB1	1:K:362:THR:HG22	1.94	0.49
1:A:342:ALA:HB2	1:A:362:THR:HG22	1.93	0.49
1:A:582:ASN:HB2	1:A:585:GLY:C	2.37	0.49
1:B:492:TYR:CE1	1:B:611:LEU:O	2.66	0.49
1:B:510:MET:HB3	1:B:542:SER:HB3	1.94	0.49
1:C:342:ALA:HB2	1:C:362:THR:HG22	1.93	0.49
1:C:353:PHE:CE2	1:C:395:TYR:CD2	2.86	0.49
1:D:443:LYS:HG3	1:D:471:ALA:HB2	1.93	0.49
1:E:387:VAL:HB	1:E:390:GLU:CD	2.37	0.49
1:F:566:ASN:O	1:F:568:ASN:OD1	2.30	0.49
1:F:644:ASN:C	1:F:644:ASN:HD22	2.20	0.49
1:G:387:VAL:O	1:G:388:GLN:OE1	2.30	0.49
1:G:387:VAL:HB	1:G:390:GLU:CD	2.37	0.49
1:H:566:ASN:O	1:H:570:GLN:CD	2.55	0.49
1:H:655:LEU:HD12	1:H:655:LEU:N	2.27	0.49
1:I:385:THR:C	1:I:387:VAL:H	2.16	0.49
1:I:391:ASP:OD1	1:I:391:ASP:N	2.46	0.49
1:I:556:GLY:HA3	1:I:589:TYR:HD1	1.74	0.49
1:J:607:ALA:C	1:J:609:ILE:H	2.20	0.49
1:L:361:GLN:HE22	1:L:456:ASN:HD21	1.57	0.49
1:A:387:VAL:O	1:A:388:GLN:OE1	2.30	0.49
1:A:510:MET:HB3	1:A:542:SER:OG	2.12	0.49
1:B:347:THR:O	1:B:348:PHE:O	2.30	0.49
1:B:566:ASN:O	1:B:568:ASN:OD1	2.30	0.49
1:B:566:ASN:CA	1:B:570:GLN:HE22	2.26	0.49
1:C:348:PHE:HZ	1:C:400:ASN:HD21	1.59	0.49
1:C:517:PHE:HE1	1:C:615:LYS:HD2	1.77	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:607:ALA:O	1:C:609:ILE:O	2.30	0.49
1:D:492:TYR:CE1	1:D:611:LEU:O	2.65	0.49
1:E:361:GLN:HA	1:E:361:GLN:NE2	2.27	0.49
1:E:457:SER:N	1:E:634:THR:HG23	2.26	0.49
1:G:361:GLN:NE2	1:G:361:GLN:HA	2.27	0.49
1:G:495:PRO:HD2	1:G:499:ILE:HG22	1.93	0.49
1:J:507:ASP:OD1	1:J:545:ARG:HG3	2.12	0.49
1:K:607:ALA:O	1:K:609:ILE:O	2.30	0.49
1:L:492:TYR:CE1	1:L:611:LEU:O	2.66	0.49
1:L:507:ASP:OD1	1:L:545:ARG:HG3	2.12	0.49
1:L:655:LEU:HD12	1:L:655:LEU:N	2.27	0.49
1:B:567:GLU:CA	1:B:570:GLN:OE1	2.56	0.49
1:F:507:ASP:OD1	1:F:545:ARG:HG3	2.12	0.49
1:H:566:ASN:O	1:H:568:ASN:OD1	2.30	0.49
1:H:566:ASN:O	1:H:570:GLN:OE1	2.30	0.49
1:I:457:SER:N	1:I:634:THR:HG23	2.26	0.49
1:J:565:GLU:CB	1:J:570:GLN:NE2	2.55	0.49
1:K:510:MET:HB3	1:K:542:SER:OG	2.12	0.49
1:A:401:LEU:HG	1:A:402:ALA:N	2.28	0.49
1:B:353:PHE:CZ	1:B:395:TYR:HE2	2.11	0.49
1:B:396:LEU:HD13	1:B:405:THR:O	2.11	0.49
1:D:566:ASN:CA	1:D:570:GLN:HE22	2.26	0.49
1:D:567:GLU:CA	1:D:570:GLN:OE1	2.56	0.49
1:E:342:ALA:HB1	1:E:362:THR:HG22	1.94	0.49
1:F:342:ALA:O	1:F:346:ASP:OD1	2.30	0.49
1:F:566:ASN:O	1:F:570:GLN:OE1	2.30	0.49
1:F:655:LEU:N	1:F:655:LEU:HD12	2.27	0.49
1:G:342:ALA:HB2	1:G:362:THR:HG22	1.93	0.49
1:I:391:ASP:C	1:I:393:LYS:H	2.20	0.49
1:J:342:ALA:O	1:J:346:ASP:OD1	2.30	0.49
1:K:361:GLN:HA	1:K:361:GLN:NE2	2.27	0.49
1:B:342:ALA:O	1:B:346:ASP:OD1	2.30	0.49
1:B:366:SER:HB3	1:K:403:PRO:N	2.17	0.49
1:B:448:TYR:CD1	1:B:642:PHE:HB2	2.48	0.49
1:C:495:PRO:HD2	1:C:499:ILE:HG22	1.93	0.49
1:C:582:ASN:HB2	1:C:585:GLY:C	2.37	0.49
1:E:522:LYS:N	1:E:522:LYS:CD	2.72	0.49
1:G:607:ALA:O	1:G:609:ILE:O	2.30	0.49
1:I:387:VAL:O	1:I:388:GLN:OE1	2.30	0.49
1:I:387:VAL:HB	1:I:390:GLU:CD	2.37	0.49
1:I:495:PRO:HD2	1:I:499:ILE:HG22	1.93	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:427:ASN:ND2	1:J:427:ASN:N	2.54	0.49
1:K:342:ALA:HB2	1:K:362:THR:HG22	1.94	0.49
1:L:396:LEU:HD13	1:L:405:THR:O	2.11	0.49
1:L:566:ASN:O	1:L:570:GLN:CD	2.55	0.49
1:L:567:GLU:CA	1:L:570:GLN:OE1	2.56	0.49
1:B:447:TYR:HB2	1:B:467:TYR:CG	2.48	0.49
1:C:587:ASP:OD2	1:C:587:ASP:O	2.30	0.49
1:D:408:ILE:HD12	1:D:409:ILE:HA	1.95	0.49
1:D:507:ASP:OD1	1:D:545:ARG:HG3	2.12	0.49
1:E:587:ASP:OD2	1:E:587:ASP:O	2.30	0.49
1:F:492:TYR:CE1	1:F:611:LEU:O	2.66	0.49
1:G:385:THR:C	1:G:387:VAL:H	2.16	0.49
1:G:401:LEU:HG	1:G:402:ALA:N	2.27	0.49
1:G:466:THR:CG2	1:H:466:THR:HB	2.43	0.49
1:H:346:ASP:HB3	1:H:360:VAL:HB	1.95	0.49
1:H:448:TYR:O	1:H:453:GLU:HG3	2.12	0.49
1:H:510:MET:HB3	1:H:542:SER:HB3	1.94	0.49
1:H:664:HIS:CD2	1:H:664:HIS:C	2.89	0.49
1:J:540:ILE:HG22	1:J:555:ILE:CG1	2.43	0.49
1:K:391:ASP:C	1:K:393:LYS:H	2.20	0.49
1:K:582:ASN:HB2	1:K:585:GLY:C	2.37	0.49
1:L:597:TYR:HB2	1:L:598:PRO:HD3	1.95	0.49
1:L:607:ALA:C	1:L:609:ILE:H	2.20	0.49
1:D:347:THR:O	1:D:348:PHE:O	2.30	0.49
1:D:566:ASN:O	1:D:570:GLN:CD	2.55	0.49
1:E:387:VAL:O	1:E:388:GLN:OE1	2.30	0.49
1:F:566:ASN:CA	1:F:570:GLN:HE22	2.26	0.49
1:J:448:TYR:CD1	1:J:642:PHE:HB2	2.48	0.49
1:A:353:PHE:O	1:A:354:GLY:C	2.48	0.49
1:A:385:THR:HG23	1:A:385:THR:O	2.12	0.49
1:A:522:LYS:N	1:A:522:LYS:CD	2.72	0.49
1:A:607:ALA:O	1:A:609:ILE:O	2.30	0.49
1:B:412:ASN:ND2	1:B:486:ARG:NH2	2.61	0.49
1:C:361:GLN:HA	1:C:361:GLN:NE2	2.27	0.49
1:C:385:THR:HG1	1:C:387:VAL:CG2	2.08	0.49
1:D:401:LEU:CG	1:D:403:PRO:HD2	2.42	0.49
1:D:448:TYR:CD1	1:D:642:PHE:HB2	2.48	0.49
1:F:346:ASP:HB3	1:F:360:VAL:HB	1.95	0.49
1:F:423:THR:HB	1:F:656:GLU:HB3	1.94	0.49
1:F:664:HIS:CD2	1:F:664:HIS:C	2.89	0.49
1:G:391:ASP:N	1:G:391:ASP:OD1	2.46	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:457:SER:N	1:G:634:THR:HG23	2.26	0.49
1:G:587:ASP:OD2	1:G:587:ASP:O	2.30	0.49
1:G:631:VAL:CG2	1:H:476:ILE:CG2	2.72	0.49
1:H:507:ASP:OD1	1:H:545:ARG:HG3	2.12	0.49
1:H:661:LEU:O	1:H:661:LEU:CG	2.58	0.49
1:I:361:GLN:NE2	1:I:361:GLN:HA	2.27	0.49
1:J:412:ASN:ND2	1:J:486:ARG:NH2	2.61	0.49
1:L:447:TYR:HB2	1:L:467:TYR:CG	2.48	0.49
1:L:448:TYR:CD1	1:L:642:PHE:HB2	2.48	0.49
1:L:566:ASN:CA	1:L:570:GLN:HE22	2.26	0.49
1:A:391:ASP:OD1	1:A:391:ASP:N	2.46	0.49
1:B:345:TYR:C	1:B:345:TYR:HD1	2.18	0.49
1:C:385:THR:HG23	1:C:385:THR:O	2.12	0.49
1:C:401:LEU:HG	1:C:402:ALA:N	2.27	0.49
1:D:662:GLU:O	1:D:662:GLU:HG2	2.13	0.49
1:F:540:ILE:HG22	1:F:555:ILE:CG1	2.43	0.49
1:H:412:ASN:ND2	1:H:486:ARG:NH2	2.61	0.49
1:H:447:TYR:HB2	1:H:467:TYR:CG	2.48	0.49
1:J:423:THR:HB	1:J:656:GLU:HB3	1.94	0.49
1:J:447:TYR:HB2	1:J:467:TYR:CG	2.48	0.49
1:J:448:TYR:O	1:J:453:GLU:HG3	2.12	0.49
1:J:662:GLU:O	1:J:662:GLU:HG2	2.13	0.49
1:K:358:GLN:HB2	1:K:379:LYS:CB	2.43	0.49
1:K:385:THR:HG23	1:K:385:THR:O	2.12	0.49
1:K:466:THR:CG2	1:L:466:THR:HB	2.43	0.49
1:L:566:ASN:O	1:L:570:GLN:OE1	2.30	0.49
1:A:536:TYR:CD1	1:A:586:ARG:CD	2.96	0.48
1:C:391:ASP:OD1	1:C:391:ASP:N	2.46	0.48
1:C:466:THR:CG2	1:D:466:THR:HB	2.43	0.48
1:C:518:ASN:OD1	1:C:533:ASP:CB	2.60	0.48
1:E:403:PRO:N	1:H:366:SER:HB3	2.16	0.48
1:F:365:ASP:O	1:F:366:SER:C	2.54	0.48
1:G:385:THR:HG23	1:G:385:THR:O	2.12	0.48
1:G:518:ASN:OD1	1:G:533:ASP:CB	2.60	0.48
1:H:448:TYR:CD1	1:H:642:PHE:HB2	2.48	0.48
1:I:385:THR:HG23	1:I:385:THR:O	2.12	0.48
1:I:466:THR:CG2	1:J:466:THR:HB	2.43	0.48
1:I:558:PHE:O	1:I:586:ARG:HB3	2.13	0.48
1:J:597:TYR:HB2	1:J:598:PRO:HD3	1.95	0.48
1:K:536:TYR:CD1	1:K:586:ARG:CD	2.96	0.48
1:L:660:GLN:CA	1:L:661:LEU:CB	2.71	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:346:ASP:HB3	1:B:360:VAL:HB	1.95	0.48
1:B:401:LEU:CG	1:B:403:PRO:HD2	2.42	0.48
1:D:414:LEU:HD23	1:D:414:LEU:HA	1.70	0.48
1:E:536:TYR:CD1	1:E:586:ARG:CD	2.96	0.48
1:F:510:MET:HB3	1:F:542:SER:HB3	1.94	0.48
1:G:607:ALA:C	1:G:609:ILE:O	2.57	0.48
1:H:492:TYR:CE1	1:H:611:LEU:O	2.66	0.48
1:H:661:LEU:C	1:H:663:HIS:N	2.61	0.48
1:A:358:GLN:HB2	1:A:379:LYS:CB	2.43	0.48
1:B:566:ASN:O	1:B:570:GLN:CD	2.55	0.48
1:C:358:GLN:HB2	1:C:379:LYS:CB	2.43	0.48
1:C:582:ASN:C	1:C:585:GLY:H	2.21	0.48
1:D:508:ARG:HG2	1:D:577:PHE:HA	1.96	0.48
1:E:358:GLN:HB2	1:E:379:LYS:CB	2.43	0.48
1:F:412:ASN:ND2	1:F:486:ARG:NH2	2.61	0.48
1:F:447:TYR:HB2	1:F:467:TYR:CG	2.48	0.48
1:G:342:ALA:HB1	1:G:362:THR:HG22	1.94	0.48
1:G:517:PHE:HE1	1:G:615:LYS:HD2	1.77	0.48
1:H:423:THR:HB	1:H:656:GLU:HB3	1.94	0.48
1:I:342:ALA:HB1	1:I:362:THR:HG22	1.94	0.48
1:I:358:GLN:HB2	1:I:379:LYS:CB	2.43	0.48
1:J:346:ASP:HB3	1:J:360:VAL:HB	1.95	0.48
1:B:597:TYR:HB2	1:B:598:PRO:HD3	1.95	0.48
1:B:655:LEU:HD12	1:B:655:LEU:N	2.27	0.48
1:C:342:ALA:HB1	1:C:362:THR:HG22	1.94	0.48
1:C:361:GLN:CG	1:C:455:PHE:CD1	2.97	0.48
1:C:424:TYR:HB3	1:C:475:VAL:H	1.78	0.48
1:D:658:ILE:CD1	1:D:658:ILE:N	2.74	0.48
1:E:385:THR:HG23	1:E:385:THR:O	2.12	0.48
1:E:466:THR:CG2	1:F:466:THR:HB	2.43	0.48
1:E:607:ALA:C	1:E:609:ILE:O	2.57	0.48
1:F:662:GLU:O	1:F:662:GLU:HG2	2.13	0.48
1:I:607:ALA:C	1:I:609:ILE:O	2.57	0.48
1:J:492:TYR:CE1	1:J:611:LEU:O	2.66	0.48
1:E:582:ASN:C	1:E:585:GLY:H	2.21	0.48
1:F:448:TYR:O	1:F:453:GLU:HG3	2.12	0.48
1:F:595:ILE:HD12	1:F:595:ILE:N	2.21	0.48
1:G:358:GLN:HB2	1:G:379:LYS:CB	2.43	0.48
1:G:545:ARG:HB3	1:G:597:TYR:CE2	2.49	0.48
1:I:342:ALA:HB2	1:I:362:THR:HG22	1.93	0.48
1:I:401:LEU:HG	1:I:402:ALA:N	2.27	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:408:ILE:HD12	1:J:409:ILE:HA	1.95	0.48
1:K:558:PHE:O	1:K:586:ARG:HB3	2.13	0.48
1:L:565:GLU:CB	1:L:570:GLN:NE2	2.55	0.48
1:A:545:ARG:HB3	1:A:597:TYR:CE2	2.48	0.48
1:B:492:TYR:HE1	1:B:611:LEU:O	1.97	0.48
1:B:662:GLU:O	1:B:662:GLU:HG2	2.13	0.48
1:D:412:ASN:ND2	1:D:486:ARG:NH2	2.61	0.48
1:D:565:GLU:OE2	1:D:590:TYR:OH	2.28	0.48
1:E:361:GLN:CG	1:E:455:PHE:CD1	2.97	0.48
1:E:418:THR:HG21	1:E:464:MET:HE1	1.96	0.48
1:E:545:ARG:HB3	1:E:597:TYR:CE2	2.49	0.48
1:G:558:PHE:O	1:G:586:ARG:HB3	2.13	0.48
1:A:466:THR:CG2	1:B:466:THR:HB	2.43	0.48
1:A:518:ASN:OD1	1:A:533:ASP:CB	2.60	0.48
1:B:664:HIS:CD2	1:B:664:HIS:C	2.90	0.48
1:C:536:TYR:CD1	1:C:586:ARG:CD	2.96	0.48
1:C:545:ARG:HB3	1:C:597:TYR:CE2	2.48	0.48
1:E:340:VAL:O	1:E:341:THR:OG1	2.30	0.48
1:E:387:VAL:CG2	1:E:390:GLU:OE1	2.55	0.48
1:F:448:TYR:C	1:F:453:GLU:HG3	2.39	0.48
1:G:418:THR:HG21	1:G:464:MET:HE1	1.96	0.48
1:J:342:ALA:O	1:J:346:ASP:OD2	2.29	0.48
1:J:401:LEU:CG	1:J:403:PRO:HD2	2.42	0.48
1:L:342:ALA:O	1:L:346:ASP:OD1	2.30	0.48
1:A:575:ASN:O	1:A:576:ASP:HB2	2.13	0.48
1:B:366:SER:C	1:K:403:PRO:HB3	2.29	0.48
1:E:401:LEU:HG	1:E:402:ALA:N	2.27	0.48
1:F:408:ILE:HD12	1:F:409:ILE:HA	1.95	0.48
1:F:448:TYR:CD1	1:F:642:PHE:HB2	2.48	0.48
1:H:540:ILE:HG22	1:H:555:ILE:CG1	2.43	0.48
1:I:349:VAL:C	1:I:351:GLU:N	2.71	0.48
1:J:508:ARG:HG2	1:J:577:PHE:HA	1.96	0.48
1:L:414:LEU:HD23	1:L:414:LEU:HA	1.70	0.48
1:L:448:TYR:C	1:L:453:GLU:HG3	2.39	0.48
1:L:508:ARG:HG2	1:L:577:PHE:HA	1.96	0.48
1:A:349:VAL:C	1:A:351:GLU:N	2.71	0.48
1:A:582:ASN:C	1:A:585:GLY:H	2.21	0.48
1:A:587:ASP:OD2	1:A:587:ASP:O	2.30	0.48
1:C:607:ALA:C	1:C:609:ILE:O	2.57	0.48
1:D:664:HIS:CD2	1:D:664:HIS:C	2.90	0.48
1:F:651:LEU:HD22	1:F:652:THR:N	2.29	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:597:TYR:HB2	1:H:598:PRO:HD3	1.95	0.48
1:I:582:ASN:C	1:I:585:GLY:H	2.21	0.48
1:J:486:ARG:HG2	1:J:499:ILE:HD11	1.96	0.48
1:J:566:ASN:CA	1:J:570:GLN:HE22	2.26	0.48
1:K:522:LYS:N	1:K:522:LYS:CD	2.72	0.48
1:K:575:ASN:O	1:K:576:ASP:HB2	2.13	0.48
1:L:492:TYR:O	1:L:606:ILE:HG12	2.14	0.48
1:L:653:ILE:HG22	1:L:655:LEU:HD12	1.96	0.48
1:L:662:GLU:O	1:L:662:GLU:HG2	2.13	0.48
1:A:361:GLN:CG	1:A:455:PHE:CD1	2.97	0.48
1:B:486:ARG:HG2	1:B:499:ILE:HD11	1.96	0.48
1:D:447:TYR:HB2	1:D:467:TYR:CG	2.48	0.48
1:D:448:TYR:C	1:D:453:GLU:HG3	2.39	0.48
1:D:492:TYR:O	1:D:606:ILE:HG12	2.14	0.48
1:D:653:ILE:HG22	1:D:655:LEU:HD12	1.96	0.48
1:F:508:ARG:HG2	1:F:577:PHE:HA	1.96	0.48
1:H:566:ASN:CA	1:H:570:GLN:HE22	2.26	0.48
1:H:651:LEU:HD22	1:H:652:THR:N	2.29	0.48
1:J:365:ASP:O	1:J:366:SER:C	2.54	0.48
1:J:651:LEU:HD22	1:J:652:THR:N	2.29	0.48
1:J:653:ILE:HG22	1:J:655:LEU:HD12	1.96	0.48
1:K:607:ALA:C	1:K:609:ILE:O	2.57	0.48
1:L:346:ASP:HB3	1:L:360:VAL:HB	1.95	0.48
1:L:408:ILE:HD12	1:L:409:ILE:HA	1.95	0.48
1:L:412:ASN:ND2	1:L:486:ARG:NH2	2.61	0.48
1:L:664:HIS:CD2	1:L:664:HIS:C	2.90	0.48
1:B:463:LYS:O	1:B:466:THR:HG23	2.14	0.47
1:B:540:ILE:HG22	1:B:555:ILE:CG1	2.43	0.47
1:B:651:LEU:HD22	1:B:652:THR:N	2.29	0.47
1:C:457:SER:N	1:C:634:THR:HG23	2.26	0.47
1:C:575:ASN:O	1:C:576:ASP:HB2	2.14	0.47
1:D:651:LEU:HD22	1:D:652:THR:N	2.29	0.47
1:E:391:ASP:N	1:E:391:ASP:OD1	2.46	0.47
1:E:558:PHE:O	1:E:586:ARG:HB3	2.13	0.47
1:F:492:TYR:HE1	1:F:611:LEU:O	1.97	0.47
1:G:424:TYR:HB3	1:G:475:VAL:H	1.79	0.47
1:G:575:ASN:O	1:G:576:ASP:HB2	2.14	0.47
1:I:418:THR:HG21	1:I:464:MET:HE1	1.96	0.47
1:K:545:ARG:HB3	1:K:597:TYR:CE2	2.49	0.47
1:L:492:TYR:HE1	1:L:611:LEU:O	1.97	0.47
1:C:340:VAL:CA	1:C:345:TYR:CZ	2.82	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:349:VAL:C	1:C:351:GLU:N	2.71	0.47
1:C:448:TYR:CG	1:C:642:PHE:HB2	2.50	0.47
1:G:579:LYS:HB3	1:G:589:TYR:OH	2.15	0.47
1:H:486:ARG:HG2	1:H:499:ILE:HD11	1.96	0.47
1:A:424:TYR:HB3	1:A:475:VAL:H	1.79	0.47
1:A:607:ALA:C	1:A:609:ILE:O	2.57	0.47
1:B:408:ILE:HD12	1:B:409:ILE:HA	1.95	0.47
1:C:387:VAL:CG2	1:C:390:GLU:OE1	2.54	0.47
1:C:579:LYS:HB3	1:C:589:TYR:OH	2.15	0.47
1:D:365:ASP:CG	1:D:367:THR:CG2	2.88	0.47
1:H:342:ALA:O	1:H:346:ASP:OD2	2.29	0.47
1:H:492:TYR:O	1:H:606:ILE:HG12	2.14	0.47
1:H:662:GLU:O	1:H:662:GLU:HG2	2.13	0.47
1:I:361:GLN:CG	1:I:455:PHE:CD1	2.97	0.47
1:I:536:TYR:CD1	1:I:586:ARG:CD	2.96	0.47
1:K:448:TYR:CG	1:K:642:PHE:HB2	2.49	0.47
1:K:518:ASN:OD1	1:K:533:ASP:CB	2.60	0.47
1:L:427:ASN:ND2	1:L:427:ASN:N	2.54	0.47
1:B:448:TYR:C	1:B:453:GLU:HG3	2.39	0.47
1:C:558:PHE:O	1:C:586:ARG:HB3	2.13	0.47
1:D:353:PHE:CE2	1:D:392:ILE:HD13	2.50	0.47
1:D:492:TYR:HE1	1:D:611:LEU:O	1.97	0.47
1:E:361:GLN:HG2	1:E:455:PHE:CD1	2.50	0.47
1:F:597:TYR:HB2	1:F:598:PRO:HD3	1.95	0.47
1:G:361:GLN:CG	1:G:455:PHE:CD1	2.97	0.47
1:H:408:ILE:HD12	1:H:409:ILE:HA	1.95	0.47
1:H:448:TYR:CE1	1:H:642:PHE:HB2	2.50	0.47
1:H:653:ILE:HG22	1:H:655:LEU:HD12	1.96	0.47
1:I:536:TYR:HD1	1:I:586:ARG:HD3	1.79	0.47
1:J:448:TYR:C	1:J:453:GLU:HG3	2.39	0.47
1:K:361:GLN:CG	1:K:455:PHE:CD1	2.97	0.47
1:K:582:ASN:C	1:K:585:GLY:H	2.21	0.47
1:L:346:ASP:OD1	1:L:346:ASP:N	2.48	0.47
1:L:353:PHE:CE2	1:L:392:ILE:HD13	2.50	0.47
1:L:651:LEU:HD22	1:L:652:THR:N	2.29	0.47
1:A:353:PHE:CE2	1:A:395:TYR:CD2	2.86	0.47
1:C:387:VAL:CB	1:C:390:GLU:CD	2.87	0.47
1:D:345:TYR:C	1:D:345:TYR:HD1	2.18	0.47
1:D:346:ASP:HB3	1:D:360:VAL:HB	1.95	0.47
1:D:385:THR:HG22	1:D:388:GLN:OE1	2.15	0.47
1:D:448:TYR:CE1	1:D:642:PHE:HB2	2.50	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:626:ASP:HA	1:E:627:PRO:HD3	1.65	0.47
1:F:486:ARG:HG2	1:F:499:ILE:HD11	1.96	0.47
1:G:388:GLN:C	1:G:389:ARG:CG	2.88	0.47
1:G:626:ASP:HA	1:G:627:PRO:HD3	1.65	0.47
1:H:365:ASP:CG	1:H:367:THR:CG2	2.88	0.47
1:I:424:TYR:HB3	1:I:475:VAL:H	1.79	0.47
1:I:545:ARG:HB3	1:I:597:TYR:CE2	2.49	0.47
1:I:596:ASN:HB3	1:I:601:VAL:HG22	1.97	0.47
1:J:463:LYS:O	1:J:466:THR:HG23	2.14	0.47
1:L:365:ASP:CG	1:L:367:THR:CG2	2.88	0.47
1:L:565:GLU:OE2	1:L:590:TYR:OH	2.28	0.47
1:B:408:ILE:O	1:B:409:ILE:HG23	2.09	0.47
1:F:401:LEU:CG	1:F:403:PRO:HD2	2.42	0.47
1:F:448:TYR:CE1	1:F:642:PHE:HB2	2.50	0.47
1:F:592:ILE:O	1:F:593:GLY:O	2.32	0.47
1:F:653:ILE:HG22	1:F:655:LEU:HD12	1.96	0.47
1:G:341:THR:O	1:G:341:THR:OG1	2.33	0.47
1:G:596:ASN:HB3	1:G:601:VAL:HG22	1.97	0.47
1:J:353:PHE:CE2	1:J:392:ILE:HD13	2.50	0.47
1:A:347:THR:C	1:A:349:VAL:N	2.73	0.47
1:A:361:GLN:HG2	1:A:455:PHE:CD1	2.50	0.47
1:A:418:THR:HG21	1:A:464:MET:HE1	1.96	0.47
1:B:492:TYR:O	1:B:606:ILE:HG12	2.14	0.47
1:B:508:ARG:HG2	1:B:577:PHE:HA	1.96	0.47
1:B:653:ILE:HG22	1:B:655:LEU:HD12	1.96	0.47
1:D:495:PRO:O	1:D:496:GLU:C	2.57	0.47
1:D:592:ILE:O	1:D:593:GLY:O	2.32	0.47
1:E:424:TYR:HB3	1:E:475:VAL:H	1.78	0.47
1:E:575:ASN:O	1:E:576:ASP:HB2	2.14	0.47
1:F:365:ASP:CG	1:F:367:THR:CG2	2.87	0.47
1:G:536:TYR:CD1	1:G:586:ARG:CD	2.96	0.47
1:G:582:ASN:C	1:G:585:GLY:H	2.21	0.47
1:H:448:TYR:C	1:H:453:GLU:HG3	2.39	0.47
1:H:492:TYR:HE1	1:H:611:LEU:O	1.97	0.47
1:H:592:ILE:O	1:H:593:GLY:O	2.32	0.47
1:J:346:ASP:OD1	1:J:346:ASP:N	2.47	0.47
1:J:448:TYR:CE1	1:J:642:PHE:HB2	2.50	0.47
1:J:492:TYR:HE1	1:J:611:LEU:O	1.97	0.47
1:J:495:PRO:O	1:J:496:GLU:C	2.57	0.47
1:J:565:GLU:OE2	1:J:590:TYR:OH	2.28	0.47
1:J:660:GLN:HA	1:J:661:LEU:HB3	1.90	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:579:LYS:HB3	1:K:589:TYR:OH	2.15	0.47
1:K:587:ASP:OD2	1:K:587:ASP:O	2.30	0.47
1:L:401:LEU:CG	1:L:403:PRO:HD2	2.42	0.47
1:L:476:ILE:HG22	1:L:477:GLY:N	2.18	0.47
1:L:486:ARG:HG2	1:L:499:ILE:HD11	1.96	0.47
1:A:387:VAL:CB	1:A:390:GLU:CD	2.88	0.47
1:A:558:PHE:O	1:A:586:ARG:HB3	2.13	0.47
1:B:342:ALA:O	1:B:346:ASP:OD2	2.29	0.47
1:B:353:PHE:CE2	1:B:392:ILE:HD13	2.50	0.47
1:B:385:THR:HG22	1:B:388:GLN:OE1	2.15	0.47
1:D:597:TYR:HB2	1:D:598:PRO:HD3	1.95	0.47
1:E:579:LYS:HB3	1:E:589:TYR:OH	2.14	0.47
1:E:596:ASN:HB3	1:E:601:VAL:HG22	1.97	0.47
1:F:342:ALA:O	1:F:346:ASP:OD2	2.29	0.47
1:F:495:PRO:O	1:F:496:GLU:C	2.57	0.47
1:G:403:PRO:HD3	1:J:366:SER:HA	1.96	0.47
1:G:448:TYR:CG	1:G:642:PHE:HB2	2.50	0.47
1:I:356:ILE:O	1:I:356:ILE:CG1	2.63	0.47
1:I:387:VAL:CB	1:I:390:GLU:CD	2.88	0.47
1:I:559:ALA:O	1:I:561:GLY:CA	2.34	0.47
1:I:579:LYS:HB3	1:I:589:TYR:OH	2.15	0.47
1:J:592:ILE:O	1:J:593:GLY:O	2.32	0.47
1:K:431:GLU:HB2	1:K:435:TRP:CE3	2.50	0.47
1:A:448:TYR:CG	1:A:642:PHE:HB2	2.50	0.47
1:B:365:ASP:CG	1:B:367:THR:CG2	2.88	0.47
1:D:664:HIS:O	1:D:665:HIS:CB	2.63	0.47
1:E:347:THR:C	1:E:349:VAL:N	2.73	0.47
1:H:345:TYR:C	1:H:345:TYR:HD1	2.18	0.47
1:J:492:TYR:O	1:J:606:ILE:HG12	2.14	0.47
1:K:420:LEU:HD12	1:K:479:SER:O	2.15	0.47
1:L:345:TYR:C	1:L:345:TYR:HD1	2.18	0.47
1:L:463:LYS:O	1:L:466:THR:HG23	2.14	0.47
1:C:388:GLN:C	1:C:389:ARG:CG	2.88	0.47
1:C:596:ASN:HB3	1:C:601:VAL:HG22	1.97	0.47
1:D:455:PHE:O	1:D:456:ASN:HB2	2.16	0.47
1:F:463:LYS:O	1:F:466:THR:HG23	2.14	0.47
1:F:522:LYS:HG2	1:F:524:VAL:HG23	1.97	0.47
1:G:347:THR:C	1:G:349:VAL:N	2.73	0.47
1:H:353:PHE:CE2	1:H:392:ILE:HD13	2.50	0.47
1:H:463:LYS:O	1:H:466:THR:HG23	2.14	0.47
1:H:656:GLU:OE1	1:H:664:HIS:CG	2.68	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:431:GLU:HB2	1:I:435:TRP:CE3	2.50	0.47
1:K:387:VAL:CB	1:K:390:GLU:CD	2.88	0.47
1:L:455:PHE:O	1:L:456:ASN:HB2	2.15	0.47
1:A:420:LEU:HD12	1:A:479:SER:O	2.15	0.46
1:C:386:THR:C	1:C:388:GLN:N	2.67	0.46
1:C:420:LEU:HD12	1:C:479:SER:O	2.15	0.46
1:D:398:ASP:CB	1:E:663:HIS:HB3	2.44	0.46
1:D:463:LYS:O	1:D:466:THR:HG23	2.14	0.46
1:E:448:TYR:CG	1:E:642:PHE:HB2	2.49	0.46
1:F:492:TYR:O	1:F:606:ILE:HG12	2.14	0.46
1:H:385:THR:HG22	1:H:388:GLN:OE1	2.15	0.46
1:H:664:HIS:O	1:H:665:HIS:CB	2.63	0.46
1:I:448:TYR:CG	1:I:642:PHE:HB2	2.50	0.46
1:J:365:ASP:CG	1:J:367:THR:CG2	2.88	0.46
1:K:418:THR:HG21	1:K:464:MET:HE1	1.96	0.46
1:L:540:ILE:HG22	1:L:555:ILE:CG1	2.43	0.46
1:A:431:GLU:HB2	1:A:435:TRP:CE3	2.50	0.46
1:B:365:ASP:HB3	1:B:367:THR:HG23	1.97	0.46
1:B:656:GLU:OE1	1:B:664:HIS:CG	2.68	0.46
1:C:418:THR:HG21	1:C:464:MET:HE1	1.96	0.46
1:D:399:TYR:CA	1:E:664:HIS:HB2	2.24	0.46
1:D:540:ILE:HG22	1:D:555:ILE:CG1	2.43	0.46
1:E:387:VAL:CB	1:E:390:GLU:CD	2.88	0.46
1:E:431:GLU:HB2	1:E:435:TRP:CE3	2.50	0.46
1:H:495:PRO:O	1:H:496:GLU:C	2.57	0.46
1:I:587:ASP:O	1:I:587:ASP:OD2	2.30	0.46
1:J:522:LYS:HG2	1:J:524:VAL:HG23	1.97	0.46
1:K:596:ASN:HB3	1:K:601:VAL:HG22	1.97	0.46
1:L:656:GLU:OE1	1:L:664:HIS:CG	2.68	0.46
1:B:366:SER:N	1:B:367:THR:HG22	2.31	0.46
1:C:536:TYR:HD1	1:C:586:ARG:HD3	1.79	0.46
1:C:559:ALA:O	1:C:561:GLY:CA	2.34	0.46
1:D:365:ASP:HB3	1:D:367:THR:HG23	1.97	0.46
1:D:486:ARG:HG2	1:D:499:ILE:HD11	1.96	0.46
1:D:656:GLU:OE1	1:D:664:HIS:CG	2.68	0.46
1:E:536:TYR:HD1	1:E:586:ARG:HD3	1.79	0.46
1:F:353:PHE:CE2	1:F:392:ILE:HD13	2.50	0.46
1:G:431:GLU:HB2	1:G:435:TRP:CE3	2.50	0.46
1:I:347:THR:C	1:I:349:VAL:N	2.73	0.46
1:J:656:GLU:OE1	1:J:664:HIS:CG	2.68	0.46
1:J:664:HIS:O	1:J:665:HIS:CB	2.63	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:424:TYR:HB3	1:K:475:VAL:H	1.79	0.46
1:L:366:SER:N	1:L:367:THR:HG22	2.31	0.46
1:L:448:TYR:CE1	1:L:642:PHE:HB2	2.50	0.46
1:A:401:LEU:CD2	1:A:402:ALA:CB	2.69	0.46
1:B:448:TYR:CE1	1:B:642:PHE:HB2	2.50	0.46
1:C:340:VAL:O	1:C:341:THR:CB	2.63	0.46
1:C:431:GLU:HB2	1:C:435:TRP:CE3	2.50	0.46
1:D:408:ILE:O	1:D:409:ILE:HG23	2.09	0.46
1:E:519:SER:O	1:E:533:ASP:HB2	2.16	0.46
1:F:401:LEU:HD12	1:F:401:LEU:HA	1.63	0.46
1:G:361:GLN:HG2	1:G:455:PHE:CD1	2.49	0.46
1:G:386:THR:C	1:G:388:GLN:N	2.67	0.46
1:G:519:SER:O	1:G:533:ASP:HB2	2.16	0.46
1:G:536:TYR:HD1	1:G:586:ARG:HD3	1.79	0.46
1:G:651:LEU:HD23	1:G:651:LEU:C	2.40	0.46
1:H:455:PHE:O	1:H:456:ASN:HB2	2.15	0.46
1:J:596:ASN:ND2	1:J:598:PRO:HD2	2.31	0.46
1:K:361:GLN:HG2	1:K:455:PHE:CD1	2.49	0.46
1:L:592:ILE:O	1:L:593:GLY:O	2.32	0.46
1:A:396:LEU:O	1:A:399:TYR:CE1	2.69	0.46
1:A:579:LYS:HB3	1:A:589:TYR:OH	2.15	0.46
1:B:495:PRO:O	1:B:496:GLU:C	2.57	0.46
1:B:592:ILE:O	1:B:593:GLY:O	2.32	0.46
1:B:596:ASN:ND2	1:B:598:PRO:HD2	2.31	0.46
1:C:519:SER:O	1:C:533:ASP:HB2	2.16	0.46
1:C:651:LEU:HD23	1:C:651:LEU:C	2.40	0.46
1:D:366:SER:N	1:D:367:THR:HG22	2.31	0.46
1:D:401:LEU:HD12	1:D:401:LEU:HA	1.63	0.46
1:D:565:GLU:CB	1:D:570:GLN:NE2	2.55	0.46
1:E:386:THR:O	1:E:388:GLN:HG3	2.16	0.46
1:F:455:PHE:O	1:F:456:ASN:HB2	2.16	0.46
1:F:541:VAL:HG12	1:F:542:SER:N	2.31	0.46
1:G:387:VAL:CB	1:G:390:GLU:CD	2.88	0.46
1:G:420:LEU:HD12	1:G:479:SER:O	2.15	0.46
1:I:396:LEU:O	1:I:399:TYR:CE1	2.69	0.46
1:I:401:LEU:CD2	1:I:402:ALA:CB	2.69	0.46
1:J:500:LYS:HB3	1:J:500:LYS:HE2	1.81	0.46
1:A:340:VAL:O	1:A:341:THR:OG1	2.30	0.46
1:A:596:ASN:HB3	1:A:601:VAL:HG22	1.97	0.46
1:A:651:LEU:HD23	1:A:651:LEU:C	2.40	0.46
1:B:405:THR:OG1	1:B:406:PRO:HD2	2.15	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:386:THR:O	1:C:388:GLN:HG3	2.16	0.46
1:E:356:ILE:O	1:E:356:ILE:CG1	2.63	0.46
1:F:385:THR:HG22	1:F:388:GLN:OE1	2.15	0.46
1:F:386:THR:O	1:F:390:GLU:HG3	2.16	0.46
1:F:656:GLU:OE1	1:F:664:HIS:CG	2.68	0.46
1:G:386:THR:O	1:G:388:GLN:HG3	2.16	0.46
1:G:522:LYS:N	1:G:522:LYS:CD	2.72	0.46
1:H:365:ASP:O	1:H:366:SER:C	2.54	0.46
1:H:508:ARG:HG2	1:H:577:PHE:HA	1.96	0.46
1:I:522:LYS:N	1:I:522:LYS:CD	2.72	0.46
1:I:575:ASN:O	1:I:576:ASP:HB2	2.13	0.46
1:I:651:LEU:HD23	1:I:651:LEU:C	2.40	0.46
1:K:536:TYR:HD1	1:K:586:ARG:HD3	1.79	0.46
1:L:365:ASP:O	1:L:366:SER:C	2.54	0.46
1:A:386:THR:O	1:A:388:GLN:HG3	2.16	0.46
1:C:485:VAL:HG22	1:C:624:TYR:CD2	2.51	0.46
1:E:420:LEU:HD12	1:E:479:SER:O	2.15	0.46
1:E:651:LEU:HD23	1:E:651:LEU:C	2.40	0.46
1:F:596:ASN:ND2	1:F:598:PRO:HD2	2.31	0.46
1:L:522:LYS:HG2	1:L:524:VAL:HG23	1.97	0.46
1:A:341:THR:O	1:A:341:THR:OG1	2.33	0.46
1:A:485:VAL:HG22	1:A:624:TYR:CD2	2.51	0.46
1:A:536:TYR:HD1	1:A:586:ARG:HD3	1.79	0.46
1:A:559:ALA:O	1:A:561:GLY:CA	2.34	0.46
1:B:660:GLN:HA	1:B:661:LEU:HB3	1.90	0.46
1:D:596:ASN:ND2	1:D:598:PRO:HD2	2.31	0.46
1:E:396:LEU:O	1:E:399:TYR:CE1	2.69	0.46
1:G:526:PRO:O	1:G:527:ASP:C	2.59	0.46
1:H:522:LYS:HG2	1:H:524:VAL:HG23	1.97	0.46
1:I:361:GLN:HG2	1:I:455:PHE:CD1	2.50	0.46
1:I:386:THR:C	1:I:388:GLN:N	2.67	0.46
1:I:626:ASP:HA	1:I:627:PRO:HD3	1.65	0.46
1:J:481:THR:O	1:J:481:THR:HG22	2.16	0.46
1:J:541:VAL:HG12	1:J:542:SER:N	2.31	0.46
1:L:402:ALA:O	1:L:403:PRO:C	2.59	0.46
1:L:541:VAL:HG12	1:L:542:SER:N	2.30	0.46
1:L:596:ASN:ND2	1:L:598:PRO:HD2	2.31	0.46
1:A:492:TYR:HA	1:A:606:ILE:HD12	1.98	0.46
1:B:455:PHE:O	1:B:456:ASN:HB2	2.15	0.46
1:B:522:LYS:HG2	1:B:524:VAL:HG23	1.97	0.46
1:B:565:GLU:OE2	1:B:590:TYR:OH	2.28	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:559:ALA:O	1:E:561:GLY:CA	2.34	0.46
1:H:401:LEU:CD1	1:H:403:PRO:HD2	2.46	0.46
1:H:541:VAL:HG12	1:H:542:SER:N	2.31	0.46
1:I:582:ASN:C	1:I:584:ASP:N	2.74	0.46
1:J:366:SER:N	1:J:367:THR:HG22	2.31	0.46
1:J:385:THR:HG22	1:J:388:GLN:OE1	2.15	0.46
1:K:347:THR:C	1:K:349:VAL:N	2.73	0.46
1:K:356:ILE:O	1:K:356:ILE:CG1	2.63	0.46
1:K:651:LEU:HD23	1:K:651:LEU:C	2.40	0.46
1:L:365:ASP:HB3	1:L:367:THR:HG23	1.97	0.46
1:B:386:THR:O	1:B:390:GLU:HG3	2.16	0.46
1:D:476:ILE:HG22	1:D:477:GLY:N	2.18	0.46
1:E:349:VAL:C	1:E:351:GLU:N	2.71	0.46
1:E:403:PRO:CD	1:H:366:SER:HA	2.45	0.46
1:H:546:ASP:CG	1:H:547:SER:O	2.59	0.46
1:I:386:THR:O	1:I:388:GLN:HG3	2.16	0.46
1:I:526:PRO:O	1:I:527:ASP:C	2.59	0.46
1:K:385:THR:HG1	1:K:388:GLN:N	2.14	0.46
1:K:582:ASN:C	1:K:584:ASP:N	2.74	0.46
1:A:385:THR:HG1	1:A:388:GLN:N	2.14	0.45
1:B:565:GLU:CB	1:B:570:GLN:NE2	2.55	0.45
1:D:481:THR:O	1:D:481:THR:HG22	2.16	0.45
1:D:522:LYS:HG2	1:D:524:VAL:HG23	1.97	0.45
1:F:401:LEU:CD1	1:F:403:PRO:HD2	2.46	0.45
1:F:546:ASP:CG	1:F:547:SER:O	2.59	0.45
1:F:660:GLN:HA	1:F:661:LEU:HB3	1.90	0.45
1:H:386:THR:O	1:H:390:GLU:HG3	2.16	0.45
1:K:391:ASP:OD1	1:K:391:ASP:N	2.46	0.45
1:B:401:LEU:CD1	1:B:403:PRO:HD2	2.46	0.45
1:B:545:ARG:NH1	1:B:597:TYR:HB3	2.32	0.45
1:C:385:THR:HG1	1:C:388:GLN:N	2.14	0.45
1:C:396:LEU:O	1:C:399:TYR:CE1	2.69	0.45
1:D:346:ASP:OD1	1:D:346:ASP:N	2.47	0.45
1:D:546:ASP:CG	1:D:547:SER:O	2.59	0.45
1:E:492:TYR:HA	1:E:606:ILE:HD12	1.98	0.45
1:E:631:VAL:CG2	1:F:476:ILE:CG2	2.72	0.45
1:F:365:ASP:HB3	1:F:367:THR:HG23	1.97	0.45
1:F:500:LYS:HB3	1:F:500:LYS:HE2	1.80	0.45
1:G:629:ASP:HB2	1:G:632:ILE:HD13	1.98	0.45
1:I:518:ASN:OD1	1:I:533:ASP:CB	2.60	0.45
1:J:545:ARG:NH1	1:J:597:TYR:HB3	2.32	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:546:ASP:CG	1:J:547:SER:O	2.59	0.45
1:K:396:LEU:O	1:K:399:TYR:CE1	2.69	0.45
1:K:485:VAL:HG22	1:K:624:TYR:CD2	2.51	0.45
1:L:481:THR:O	1:L:481:THR:HG22	2.16	0.45
1:L:495:PRO:O	1:L:496:GLU:C	2.57	0.45
1:A:502:ASN:ND2	1:A:635:ARG:CB	2.59	0.45
1:D:401:LEU:CD1	1:D:403:PRO:HD2	2.46	0.45
1:D:492:TYR:OH	1:D:611:LEU:O	2.34	0.45
1:D:545:ARG:NH1	1:D:597:TYR:HB3	2.32	0.45
1:E:629:ASP:HB2	1:E:632:ILE:HD13	1.99	0.45
1:F:664:HIS:O	1:F:665:HIS:CB	2.63	0.45
1:G:358:GLN:HB2	1:G:379:LYS:HB2	1.99	0.45
1:G:440:ILE:CG1	1:G:441:ILE:N	2.79	0.45
1:H:481:THR:O	1:H:481:THR:HG22	2.16	0.45
1:J:455:PHE:O	1:J:456:ASN:HB2	2.15	0.45
1:A:629:ASP:HB2	1:A:632:ILE:HD13	1.99	0.45
1:B:365:ASP:O	1:B:366:SER:C	2.54	0.45
1:B:541:VAL:HG12	1:B:542:SER:N	2.30	0.45
1:C:526:PRO:O	1:C:527:ASP:C	2.59	0.45
1:D:484:MET:O	1:D:624:TYR:HA	2.17	0.45
1:D:541:VAL:HG12	1:D:542:SER:N	2.31	0.45
1:F:402:ALA:O	1:F:403:PRO:C	2.59	0.45
1:H:427:ASN:ND2	1:H:427:ASN:N	2.54	0.45
1:J:484:MET:O	1:J:624:TYR:HA	2.17	0.45
1:J:599:ALA:O	1:J:600:ASP:C	2.59	0.45
1:J:664:HIS:CD2	1:J:664:HIS:C	2.90	0.45
1:K:340:VAL:O	1:K:341:THR:OG1	2.30	0.45
1:L:385:THR:HG22	1:L:388:GLN:OE1	2.15	0.45
1:L:664:HIS:O	1:L:665:HIS:CB	2.63	0.45
1:B:366:SER:HA	1:K:403:PRO:CD	2.47	0.45
1:B:441:ILE:HD11	1:B:644:ASN:CG	2.42	0.45
1:C:346:ASP:OD1	1:C:362:THR:HB	2.17	0.45
1:C:347:THR:C	1:C:349:VAL:N	2.73	0.45
1:C:356:ILE:O	1:C:356:ILE:CG1	2.63	0.45
1:C:440:ILE:CG1	1:C:441:ILE:N	2.79	0.45
1:C:629:ASP:HB2	1:C:632:ILE:HD13	1.99	0.45
1:F:508:ARG:HA	1:F:542:SER:O	2.17	0.45
1:G:340:VAL:O	1:G:341:THR:CB	2.63	0.45
1:G:396:LEU:O	1:G:399:TYR:CE1	2.69	0.45
1:G:660:GLN:HG3	1:G:661:LEU:H	1.77	0.45
1:I:485:VAL:HG22	1:I:624:TYR:CD2	2.51	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:629:ASP:HB2	1:I:632:ILE:HD13	1.99	0.45
1:J:401:LEU:CD1	1:J:403:PRO:HD2	2.46	0.45
1:K:629:ASP:HB2	1:K:632:ILE:HD13	1.99	0.45
1:L:599:ALA:O	1:L:600:ASP:C	2.59	0.45
1:A:519:SER:O	1:A:533:ASP:HB2	2.16	0.45
1:B:648:PRO:HD2	1:B:649:GLN:H	1.81	0.45
1:E:460:ALA:HB2	1:E:631:VAL:HG22	1.99	0.45
1:F:441:ILE:HD11	1:F:644:ASN:CG	2.42	0.45
1:F:484:MET:O	1:F:624:TYR:HA	2.17	0.45
1:G:460:ALA:HB2	1:G:631:VAL:HG22	1.99	0.45
1:G:485:VAL:HG22	1:G:624:TYR:CD2	2.51	0.45
1:H:366:SER:N	1:H:367:THR:HG22	2.31	0.45
1:H:596:ASN:ND2	1:H:598:PRO:HD2	2.31	0.45
1:I:420:LEU:HD12	1:I:479:SER:O	2.15	0.45
1:J:605:ASN:OD1	1:J:607:ALA:HB3	2.17	0.45
1:J:661:LEU:C	1:J:661:LEU:CD2	2.89	0.45
1:K:386:THR:O	1:K:388:GLN:HG3	2.16	0.45
1:K:626:ASP:HA	1:K:627:PRO:HD3	1.65	0.45
1:L:405:THR:OG1	1:L:406:PRO:HD2	2.15	0.45
1:L:484:MET:O	1:L:624:TYR:HA	2.17	0.45
1:A:440:ILE:CG1	1:A:441:ILE:N	2.79	0.45
1:B:599:ALA:O	1:B:600:ASP:C	2.59	0.45
1:B:605:ASN:OD1	1:B:607:ALA:HB3	2.17	0.45
1:B:623:LEU:HD12	1:B:623:LEU:HA	1.86	0.45
1:B:664:HIS:O	1:B:665:HIS:CB	2.63	0.45
1:D:441:ILE:HD11	1:D:644:ASN:CG	2.42	0.45
1:D:658:ILE:CG2	1:D:659:SER:N	2.79	0.45
1:F:353:PHE:CZ	1:F:395:TYR:HE2	2.11	0.45
1:H:401:LEU:CG	1:H:403:PRO:HD2	2.42	0.45
1:H:402:ALA:O	1:H:403:PRO:C	2.59	0.45
1:H:605:ASN:OD1	1:H:607:ALA:HB3	2.17	0.45
1:H:660:GLN:HA	1:H:661:LEU:HB3	1.90	0.45
1:I:358:GLN:HB2	1:I:379:LYS:HB2	1.99	0.45
1:K:559:ALA:O	1:K:561:GLY:CA	2.34	0.45
1:L:545:ARG:NH1	1:L:597:TYR:HB3	2.32	0.45
1:L:648:PRO:HD2	1:L:649:GLN:H	1.82	0.45
1:B:572:TYR:HB2	1:B:591:VAL:HG23	1.99	0.45
1:D:389:ARG:HA	1:D:392:ILE:HG22	1.99	0.45
1:D:508:ARG:HA	1:D:542:SER:O	2.17	0.45
1:E:358:GLN:HB2	1:E:379:LYS:HB2	1.99	0.45
1:E:485:VAL:HG22	1:E:624:TYR:CD2	2.51	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:518:ASN:OD1	1:E:533:ASP:CB	2.60	0.45
1:F:366:SER:N	1:F:367:THR:HG22	2.31	0.45
1:F:599:ALA:O	1:F:600:ASP:C	2.59	0.45
1:F:605:ASN:OD1	1:F:607:ALA:HB3	2.17	0.45
1:F:648:PRO:HD2	1:F:649:GLN:H	1.82	0.45
1:G:346:ASP:OD1	1:G:362:THR:HB	2.17	0.45
1:H:545:ARG:NH1	1:H:597:TYR:HB3	2.32	0.45
1:I:353:PHE:C	1:I:355:SER:N	2.65	0.45
1:J:508:ARG:HA	1:J:542:SER:O	2.17	0.45
1:K:492:TYR:HA	1:K:606:ILE:HD12	1.98	0.45
1:L:605:ASN:OD1	1:L:607:ALA:HB3	2.17	0.45
1:A:437:GLU:C	1:A:439:GLN:N	2.75	0.45
1:A:554:ILE:HD11	1:A:577:PHE:CD2	2.52	0.45
1:C:361:GLN:HG2	1:C:455:PHE:CD1	2.50	0.45
1:D:572:TYR:HB2	1:D:591:VAL:HG23	1.99	0.45
1:E:340:VAL:O	1:E:341:THR:CB	2.63	0.45
1:G:658:ILE:HA	1:G:658:ILE:HD13	1.52	0.45
1:I:595:ILE:HG23	1:I:602:ILE:HG12	1.99	0.45
1:J:658:ILE:CD1	1:J:658:ILE:N	2.74	0.45
1:K:437:GLU:C	1:K:439:GLN:N	2.75	0.45
1:L:658:ILE:CD1	1:L:658:ILE:N	2.74	0.45
1:B:402:ALA:O	1:B:403:PRO:C	2.59	0.45
1:B:438:GLY:HA2	1:B:441:ILE:CG2	2.47	0.45
1:B:511:GLU:O	1:B:511:GLU:HG3	2.17	0.45
1:G:387:VAL:CG2	1:G:390:GLU:OE1	2.55	0.45
1:G:517:PHE:CE1	1:G:615:LYS:HB3	2.52	0.45
1:H:346:ASP:OD1	1:H:346:ASP:N	2.48	0.45
1:H:365:ASP:HB3	1:H:367:THR:HG23	1.97	0.45
1:H:441:ILE:HD11	1:H:644:ASN:CG	2.42	0.45
1:I:530:LEU:HD23	1:I:530:LEU:HA	1.82	0.45
1:J:441:ILE:HD11	1:J:644:ASN:CG	2.42	0.45
1:J:648:PRO:HD2	1:J:649:GLN:H	1.82	0.45
1:K:341:THR:O	1:K:341:THR:OG1	2.33	0.45
1:L:508:ARG:HA	1:L:542:SER:O	2.17	0.45
1:L:546:ASP:CG	1:L:547:SER:O	2.59	0.45
1:A:517:PHE:CE1	1:A:615:LYS:HB3	2.52	0.44
1:B:658:ILE:CG2	1:B:659:SER:N	2.79	0.44
1:C:441:ILE:O	1:C:441:ILE:HG13	2.17	0.44
1:D:462:SER:O	1:D:466:THR:CG2	2.66	0.44
1:E:595:ILE:HG23	1:E:602:ILE:HG12	1.99	0.44
1:F:438:GLY:HA2	1:F:441:ILE:CG2	2.48	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:658:ILE:CG2	1:F:659:SER:N	2.79	0.44
1:H:353:PHE:CZ	1:H:395:TYR:HE2	2.11	0.44
1:H:572:TYR:HB2	1:H:591:VAL:HG23	1.99	0.44
1:H:599:ALA:O	1:H:600:ASP:C	2.59	0.44
1:I:340:VAL:O	1:I:341:THR:CB	2.63	0.44
1:I:346:ASP:OD1	1:I:362:THR:HB	2.17	0.44
1:I:387:VAL:CG2	1:I:390:GLU:HB2	2.47	0.44
1:I:492:TYR:HA	1:I:606:ILE:HD12	1.98	0.44
1:J:365:ASP:HB3	1:J:367:THR:HG23	1.97	0.44
1:J:401:LEU:HD12	1:J:401:LEU:HA	1.63	0.44
1:K:519:SER:O	1:K:533:ASP:HB2	2.16	0.44
1:L:358:GLN:HB3	1:L:379:LYS:HA	1.99	0.44
1:L:386:THR:O	1:L:390:GLU:HG3	2.16	0.44
1:B:377:LYS:HB2	1:B:411:PRO:HG2	1.99	0.44
1:B:484:MET:O	1:B:624:TYR:HA	2.17	0.44
1:B:508:ARG:HA	1:B:542:SER:O	2.17	0.44
1:B:546:ASP:CG	1:B:547:SER:O	2.59	0.44
1:C:492:TYR:HA	1:C:606:ILE:HD12	1.98	0.44
1:D:353:PHE:CZ	1:D:395:TYR:HE2	2.11	0.44
1:E:385:THR:HG1	1:E:388:GLN:N	2.14	0.44
1:E:524:VAL:HG22	1:E:529:GLY:C	2.42	0.44
1:F:377:LYS:HB2	1:F:411:PRO:HG2	1.99	0.44
1:G:510:MET:SD	1:G:602:ILE:HG21	2.58	0.44
1:I:441:ILE:O	1:I:441:ILE:HG13	2.17	0.44
1:J:402:ALA:O	1:J:403:PRO:C	2.59	0.44
1:K:340:VAL:O	1:K:341:THR:CG2	2.64	0.44
1:L:389:ARG:HA	1:L:392:ILE:HG22	1.99	0.44
1:A:358:GLN:HB2	1:A:379:LYS:HB2	1.99	0.44
1:A:460:ALA:HB2	1:A:631:VAL:HG22	1.99	0.44
1:A:556:GLY:CA	1:A:589:TYR:CD1	3.01	0.44
1:D:342:ALA:O	1:D:346:ASP:OD2	2.29	0.44
1:D:605:ASN:OD1	1:D:607:ALA:HB3	2.17	0.44
1:D:648:PRO:HD2	1:D:649:GLN:H	1.81	0.44
1:E:510:MET:SD	1:E:602:ILE:HG21	2.57	0.44
1:E:534:VAL:C	1:E:535:LEU:HD23	2.43	0.44
1:F:389:ARG:HA	1:F:392:ILE:HG22	1.99	0.44
1:F:427:ASN:ND2	1:F:427:ASN:N	2.54	0.44
1:F:572:TYR:HB2	1:F:591:VAL:HG23	1.99	0.44
1:G:349:VAL:C	1:G:351:GLU:N	2.71	0.44
1:G:492:TYR:HA	1:G:606:ILE:HD12	1.98	0.44
1:I:377:LYS:HA	1:I:378:PRO:HD3	1.84	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:I:437:GLU:C	1:I:439:GLN:N	2.75	0.44
1:I:510:MET:SD	1:I:602:ILE:HG21	2.57	0.44
1:I:519:SER:O	1:I:533:ASP:HB2	2.16	0.44
1:J:358:GLN:HB3	1:J:379:LYS:HA	1.99	0.44
1:K:346:ASP:OD1	1:K:362:THR:HB	2.17	0.44
1:K:440:ILE:CG1	1:K:441:ILE:N	2.79	0.44
1:L:438:GLY:HA2	1:L:441:ILE:CG2	2.47	0.44
1:B:481:THR:O	1:B:481:THR:HG22	2.16	0.44
1:B:607:ALA:C	1:B:609:ILE:N	2.76	0.44
1:C:460:ALA:HB2	1:C:631:VAL:HG22	1.99	0.44
1:E:387:VAL:CG2	1:E:390:GLU:HB2	2.48	0.44
1:E:554:ILE:HD11	1:E:577:PHE:CD2	2.52	0.44
1:F:511:GLU:O	1:F:511:GLU:HG3	2.17	0.44
1:H:448:TYR:CG	1:H:642:PHE:HB2	2.53	0.44
1:H:458:SER:HB2	1:H:632:ILE:O	2.17	0.44
1:H:462:SER:O	1:H:466:THR:CG2	2.66	0.44
1:I:502:ASN:ND2	1:I:635:ARG:CB	2.59	0.44
1:I:534:VAL:C	1:I:535:LEU:HD23	2.43	0.44
1:K:510:MET:SD	1:K:602:ILE:HG21	2.57	0.44
1:L:658:ILE:CG2	1:L:659:SER:N	2.79	0.44
1:A:346:ASP:OD1	1:A:362:THR:HB	2.17	0.44
1:B:389:ARG:HA	1:B:392:ILE:HG22	1.99	0.44
1:C:510:MET:SD	1:C:602:ILE:HG21	2.58	0.44
1:C:517:PHE:CE1	1:C:615:LYS:HB3	2.52	0.44
1:D:511:GLU:HG3	1:D:511:GLU:O	2.17	0.44
1:E:401:LEU:HD21	1:E:402:ALA:HB2	1.96	0.44
1:E:441:ILE:HG13	1:E:441:ILE:O	2.17	0.44
1:F:458:SER:HB2	1:F:632:ILE:O	2.17	0.44
1:F:545:ARG:NH1	1:F:597:TYR:HB3	2.32	0.44
1:G:484:MET:HG3	1:G:627:PRO:HG3	2.00	0.44
1:H:405:THR:OG1	1:H:406:PRO:HD2	2.16	0.44
1:H:508:ARG:HA	1:H:542:SER:O	2.17	0.44
1:J:353:PHE:CZ	1:J:395:TYR:HE2	2.11	0.44
1:J:458:SER:HB2	1:J:632:ILE:O	2.17	0.44
1:J:524:VAL:O	1:J:524:VAL:HG12	2.17	0.44
1:L:365:ASP:C	1:L:367:THR:CG2	2.90	0.44
1:L:462:SER:O	1:L:466:THR:CG2	2.66	0.44
1:L:567:GLU:C	1:L:568:ASN:CG	2.86	0.44
1:L:607:ALA:C	1:L:609:ILE:N	2.76	0.44
1:B:365:ASP:C	1:B:367:THR:CG2	2.90	0.44
1:C:556:GLY:CA	1:C:589:TYR:CD1	3.01	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:377:LYS:HB2	1:D:411:PRO:HG2	1.99	0.44
1:D:599:ALA:O	1:D:600:ASP:C	2.59	0.44
1:E:437:GLU:C	1:E:439:GLN:N	2.75	0.44
1:E:484:MET:HG3	1:E:627:PRO:HG3	2.00	0.44
1:F:481:THR:O	1:F:481:THR:HG22	2.16	0.44
1:F:560:SER:HB2	1:F:588:LYS:NZ	2.33	0.44
1:G:437:GLU:C	1:G:439:GLN:N	2.75	0.44
1:I:484:MET:HG3	1:I:627:PRO:HG3	2.00	0.44
1:I:517:PHE:CE1	1:I:615:LYS:HB3	2.53	0.44
1:I:524:VAL:HG22	1:I:529:GLY:C	2.42	0.44
1:J:386:THR:O	1:J:390:GLU:HG3	2.16	0.44
1:K:460:ALA:HB2	1:K:631:VAL:HG22	1.99	0.44
1:K:554:ILE:HD11	1:K:577:PHE:CD2	2.52	0.44
1:L:377:LYS:HB2	1:L:411:PRO:HG2	1.99	0.44
1:L:401:LEU:CD1	1:L:403:PRO:HD2	2.46	0.44
1:L:441:ILE:HD11	1:L:644:ASN:CG	2.42	0.44
1:L:484:MET:HG3	1:L:627:PRO:HG3	2.00	0.44
1:L:511:GLU:O	1:L:511:GLU:HG3	2.17	0.44
1:A:387:VAL:CG2	1:A:390:GLU:HB2	2.48	0.44
1:A:510:MET:SD	1:A:602:ILE:HG21	2.57	0.44
1:B:461:LYS:O	1:B:461:LYS:HG2	2.18	0.44
1:C:534:VAL:C	1:C:535:LEU:HD23	2.43	0.44
1:E:582:ASN:C	1:E:584:ASP:N	2.74	0.44
1:E:650:TYR:O	1:E:651:LEU:HB2	2.18	0.44
1:E:658:ILE:HA	1:E:658:ILE:HD13	1.52	0.44
1:F:462:SER:O	1:F:466:THR:CG2	2.66	0.44
1:H:377:LYS:HB2	1:H:411:PRO:HG2	1.99	0.44
1:H:524:VAL:HG12	1:H:524:VAL:O	2.17	0.44
1:H:648:PRO:HD2	1:H:649:GLN:H	1.81	0.44
1:I:554:ILE:HD11	1:I:577:PHE:CD2	2.52	0.44
1:J:389:ARG:HA	1:J:392:ILE:HG22	1.99	0.44
1:J:438:GLY:HA2	1:J:441:ILE:CG2	2.47	0.44
1:J:567:GLU:C	1:J:568:ASN:CG	2.86	0.44
1:K:530:LEU:HD23	1:K:530:LEU:HA	1.82	0.44
1:L:500:LYS:HB3	1:L:500:LYS:HE2	1.80	0.44
1:L:524:VAL:O	1:L:524:VAL:HG12	2.17	0.44
1:A:534:VAL:C	1:A:535:LEU:HD23	2.43	0.44
1:B:462:SER:O	1:B:466:THR:CG2	2.66	0.44
1:B:484:MET:HG3	1:B:627:PRO:HG3	2.00	0.44
1:B:500:LYS:HB3	1:B:500:LYS:HE2	1.80	0.44
1:C:358:GLN:HB2	1:C:379:LYS:HB2	1.99	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:660:GLN:HG3	1:C:661:LEU:H	1.77	0.44
1:D:367:THR:CG2	1:D:368:LYS:N	2.45	0.44
1:D:405:THR:OG1	1:D:406:PRO:HD2	2.15	0.44
1:D:524:VAL:O	1:D:524:VAL:HG12	2.17	0.44
1:D:560:SER:HB2	1:D:588:LYS:NZ	2.33	0.44
1:E:517:PHE:CE1	1:E:615:LYS:HB3	2.52	0.44
1:F:658:ILE:HD12	1:F:658:ILE:HA	1.67	0.44
1:G:375:ALA:HA	1:G:409:ILE:HD12	2.00	0.44
1:G:441:ILE:HG13	1:G:441:ILE:O	2.17	0.44
1:I:341:THR:O	1:I:341:THR:OG1	2.33	0.44
1:I:385:THR:HG1	1:I:388:GLN:N	2.15	0.44
1:I:403:PRO:HD3	1:L:366:SER:HA	2.00	0.44
1:K:379:LYS:O	1:K:379:LYS:CG	2.66	0.44
1:K:534:VAL:C	1:K:535:LEU:HD23	2.43	0.44
1:B:458:SER:HB2	1:B:632:ILE:O	2.17	0.44
1:C:595:ILE:HG23	1:C:602:ILE:HG12	1.99	0.44
1:D:358:GLN:HB3	1:D:379:LYS:HA	1.99	0.44
1:E:379:LYS:O	1:E:379:LYS:CG	2.66	0.44
1:F:346:ASP:OD1	1:F:346:ASP:N	2.48	0.44
1:G:365:ASP:OD2	1:G:367:THR:HB	2.18	0.44
1:H:438:GLY:HA2	1:H:441:ILE:CG2	2.47	0.44
1:I:340:VAL:O	1:I:341:THR:OG1	2.30	0.44
1:I:379:LYS:O	1:I:379:LYS:CG	2.66	0.44
1:I:635:ARG:HE	1:I:635:ARG:HB2	1.66	0.44
1:J:405:THR:OG1	1:J:406:PRO:HD2	2.15	0.44
1:J:658:ILE:CG2	1:J:659:SER:N	2.79	0.44
1:K:517:PHE:CE1	1:K:615:LYS:HB3	2.52	0.44
1:K:556:GLY:CA	1:K:589:TYR:CD1	3.01	0.44
1:L:458:SER:HB2	1:L:632:ILE:O	2.17	0.44
1:A:516:SER:HB2	1:A:617:GLU:OE1	2.18	0.43
1:B:567:GLU:C	1:B:568:ASN:CG	2.86	0.43
1:D:386:THR:O	1:D:390:GLU:HG3	2.16	0.43
1:D:567:GLU:C	1:D:568:ASN:CG	2.86	0.43
1:E:530:LEU:HD23	1:E:530:LEU:HA	1.82	0.43
1:F:464:MET:O	1:F:468:VAL:HG23	2.18	0.43
1:G:385:THR:HG1	1:G:388:GLN:N	2.15	0.43
1:G:650:TYR:O	1:G:651:LEU:HB2	2.18	0.43
1:H:484:MET:O	1:H:624:TYR:HA	2.17	0.43
1:H:560:SER:HB2	1:H:588:LYS:NZ	2.33	0.43
1:I:460:ALA:HB2	1:I:631:VAL:HG22	1.99	0.43
1:J:492:TYR:OH	1:J:611:LEU:O	2.34	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:358:GLN:HB2	1:K:379:LYS:HB2	1.99	0.43
1:K:436:LEU:C	1:K:436:LEU:CD2	2.91	0.43
1:K:484:MET:HG3	1:K:627:PRO:HG3	2.00	0.43
1:L:461:LYS:O	1:L:461:LYS:HG2	2.18	0.43
1:L:557:PRO:CG	1:L:580:LEU:HD11	2.37	0.43
1:A:650:TYR:O	1:A:651:LEU:HB2	2.18	0.43
1:B:560:SER:HB2	1:B:588:LYS:NZ	2.33	0.43
1:C:650:TYR:O	1:C:651:LEU:HB2	2.18	0.43
1:D:438:GLY:HA2	1:D:441:ILE:CG2	2.47	0.43
1:E:408:ILE:H	1:E:408:ILE:CD1	2.30	0.43
1:E:635:ARG:HE	1:E:635:ARG:HB2	1.66	0.43
1:F:461:LYS:O	1:F:461:LYS:HG2	2.18	0.43
1:F:484:MET:HG3	1:F:627:PRO:HG3	2.00	0.43
1:G:541:VAL:CG1	1:G:542:SER:N	2.82	0.43
1:G:554:ILE:HD11	1:G:577:PHE:CD2	2.52	0.43
1:H:648:PRO:CD	1:H:649:GLN:H	2.32	0.43
1:H:658:ILE:CG2	1:H:659:SER:N	2.79	0.43
1:I:340:VAL:O	1:I:341:THR:CG2	2.64	0.43
1:I:375:ALA:HA	1:I:409:ILE:HD12	2.00	0.43
1:J:462:SER:O	1:J:466:THR:CG2	2.66	0.43
1:K:387:VAL:CG2	1:K:390:GLU:HB2	2.47	0.43
1:K:595:ILE:HG23	1:K:602:ILE:HG12	1.99	0.43
1:A:356:ILE:O	1:A:356:ILE:CG1	2.63	0.43
1:A:361:GLN:HA	1:A:361:GLN:HE21	1.84	0.43
1:A:379:LYS:O	1:A:379:LYS:CG	2.66	0.43
1:A:541:VAL:CG1	1:A:542:SER:N	2.81	0.43
1:A:664:HIS:HD2	1:K:348:PHE:CZ	2.17	0.43
1:B:377:LYS:CE	1:B:641:VAL:HG21	2.48	0.43
1:C:365:ASP:OD2	1:C:367:THR:HB	2.18	0.43
1:C:401:LEU:HD21	1:C:402:ALA:HB2	1.96	0.43
1:C:437:GLU:C	1:C:439:GLN:N	2.75	0.43
1:C:541:VAL:CG1	1:C:542:SER:N	2.82	0.43
1:C:635:ARG:HE	1:C:635:ARG:HB2	1.66	0.43
1:D:365:ASP:C	1:D:367:THR:CG2	2.90	0.43
1:D:607:ALA:C	1:D:609:ILE:N	2.76	0.43
1:D:623:LEU:HD12	1:D:623:LEU:HA	1.86	0.43
1:E:375:ALA:HA	1:E:409:ILE:HD12	2.00	0.43
1:E:656:GLU:HA	1:E:657:PRO:HD3	1.82	0.43
1:H:464:MET:O	1:H:468:VAL:HG23	2.18	0.43
1:H:557:PRO:CG	1:H:580:LEU:HD11	2.37	0.43
1:I:385:THR:OG1	1:I:387:VAL:HG22	2.11	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:661:LEU:C	1:J:663:HIS:N	2.61	0.43
1:K:516:SER:HB2	1:K:617:GLU:OE1	2.18	0.43
1:K:541:VAL:CG1	1:K:542:SER:N	2.82	0.43
1:C:484:MET:HG3	1:C:627:PRO:HG3	2.00	0.43
1:D:458:SER:HB2	1:D:632:ILE:O	2.17	0.43
1:E:346:ASP:OD1	1:E:362:THR:HB	2.17	0.43
1:E:365:ASP:OD2	1:E:367:THR:HB	2.18	0.43
1:E:661:LEU:HD13	1:E:662:GLU:CB	2.42	0.43
1:F:524:VAL:O	1:F:524:VAL:HG12	2.17	0.43
1:H:389:ARG:HA	1:H:392:ILE:HG22	1.99	0.43
1:H:600:ASP:O	1:H:600:ASP:CG	2.62	0.43
1:I:541:VAL:CG1	1:I:542:SER:N	2.82	0.43
1:J:377:LYS:CE	1:J:641:VAL:HG21	2.48	0.43
1:J:448:TYR:CG	1:J:642:PHE:HB2	2.53	0.43
1:K:375:ALA:HA	1:K:409:ILE:HD12	2.00	0.43
1:L:342:ALA:O	1:L:346:ASP:OD2	2.29	0.43
1:L:353:PHE:CZ	1:L:395:TYR:CD2	3.05	0.43
1:A:365:ASP:OD2	1:A:367:THR:HB	2.18	0.43
1:B:358:GLN:HB3	1:B:379:LYS:HA	1.99	0.43
1:B:524:VAL:O	1:B:524:VAL:HG12	2.17	0.43
1:D:464:MET:O	1:D:468:VAL:HG23	2.18	0.43
1:F:375:ALA:HA	1:F:409:ILE:O	2.19	0.43
1:F:648:PRO:CD	1:F:649:GLN:H	2.32	0.43
1:G:516:SER:HB2	1:G:617:GLU:OE1	2.18	0.43
1:G:524:VAL:HG22	1:G:529:GLY:C	2.42	0.43
1:H:375:ALA:HA	1:H:409:ILE:O	2.19	0.43
1:I:658:ILE:HD13	1:I:658:ILE:HA	1.52	0.43
1:J:353:PHE:CZ	1:J:395:TYR:CD2	3.05	0.43
1:J:511:GLU:O	1:J:511:GLU:HG3	2.17	0.43
1:K:661:LEU:HA	1:K:661:LEU:HD22	1.75	0.43
1:L:560:SER:HB2	1:L:588:LYS:NZ	2.33	0.43
1:A:484:MET:HG3	1:A:627:PRO:HG3	2.00	0.43
1:B:447:TYR:HB2	1:B:467:TYR:CD1	2.54	0.43
1:C:387:VAL:CG2	1:C:390:GLU:HB2	2.47	0.43
1:E:541:VAL:CG1	1:E:542:SER:N	2.82	0.43
1:G:595:ILE:HG23	1:G:602:ILE:HG12	1.99	0.43
1:H:377:LYS:CE	1:H:641:VAL:HG21	2.48	0.43
1:H:461:LYS:O	1:H:461:LYS:HG2	2.18	0.43
1:H:492:TYR:OH	1:H:611:LEU:O	2.34	0.43
1:I:361:GLN:HA	1:I:361:GLN:HE21	1.84	0.43
1:I:593:GLY:HA3	1:I:603:TYR:O	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:572:TYR:HB2	1:J:591:VAL:CG2	2.49	0.43
1:J:572:TYR:HB2	1:J:591:VAL:HG23	1.99	0.43
1:K:401:LEU:CD2	1:K:402:ALA:CB	2.68	0.43
1:L:572:TYR:HB2	1:L:591:VAL:CG2	2.49	0.43
1:A:375:ALA:HA	1:A:409:ILE:HD12	2.00	0.43
1:A:533:ASP:O	1:A:535:LEU:CD2	2.67	0.43
1:B:448:TYR:CG	1:B:642:PHE:HB2	2.53	0.43
1:B:653:ILE:HG22	1:B:655:LEU:CD1	2.49	0.43
1:C:375:ALA:HA	1:C:409:ILE:HD12	2.00	0.43
1:C:379:LYS:O	1:C:379:LYS:CG	2.66	0.43
1:C:661:LEU:HA	1:C:661:LEU:HD22	1.75	0.43
1:D:600:ASP:O	1:D:600:ASP:CG	2.62	0.43
1:E:361:GLN:HA	1:E:361:GLN:HE21	1.84	0.43
1:E:469:ASP:HA	1:E:475:VAL:HG11	2.01	0.43
1:F:358:GLN:HB3	1:F:379:LYS:HA	1.99	0.43
1:F:448:TYR:CG	1:F:642:PHE:HB2	2.53	0.43
1:F:607:ALA:C	1:F:609:ILE:N	2.76	0.43
1:G:593:GLY:HA3	1:G:603:TYR:O	2.19	0.43
1:I:408:ILE:H	1:I:408:ILE:CD1	2.30	0.43
1:J:377:LYS:HB2	1:J:411:PRO:HG2	1.99	0.43
1:K:441:ILE:O	1:K:441:ILE:HG13	2.17	0.43
1:K:469:ASP:HA	1:K:475:VAL:HG11	2.00	0.43
1:K:623:LEU:HD12	1:K:623:LEU:HA	1.79	0.43
1:L:377:LYS:CE	1:L:641:VAL:HG21	2.48	0.43
1:L:494:THR:HG22	1:L:499:ILE:HG22	2.01	0.43
1:L:572:TYR:HB2	1:L:591:VAL:HG23	1.99	0.43
1:L:592:ILE:CD1	1:L:606:ILE:HD13	2.49	0.43
1:L:653:ILE:HG22	1:L:655:LEU:CD1	2.49	0.43
1:A:582:ASN:C	1:A:584:ASP:N	2.74	0.43
1:A:626:ASP:HA	1:A:627:PRO:HD3	1.65	0.43
1:B:410:SER:HA	1:B:411:PRO:HD3	1.81	0.43
1:B:494:THR:HG22	1:B:499:ILE:HG22	2.01	0.43
1:C:516:SER:HB2	1:C:617:GLU:OE1	2.18	0.43
1:C:554:ILE:HD11	1:C:577:PHE:CD2	2.52	0.43
1:D:377:LYS:CE	1:D:641:VAL:HG21	2.48	0.43
1:D:448:TYR:CG	1:D:642:PHE:HB2	2.53	0.43
1:E:492:TYR:CE1	1:E:606:ILE:HB	2.54	0.43
1:E:533:ASP:O	1:E:535:LEU:CD2	2.67	0.43
1:F:377:LYS:CE	1:F:641:VAL:HG21	2.48	0.43
1:F:592:ILE:CD1	1:F:606:ILE:HD13	2.49	0.43
1:G:379:LYS:O	1:G:379:LYS:CG	2.66	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:387:VAL:CG2	1:G:390:GLU:HB2	2.48	0.43
1:H:388:GLN:O	1:H:392:ILE:HG22	2.19	0.43
1:H:484:MET:HG3	1:H:627:PRO:HG3	2.00	0.43
1:K:533:ASP:O	1:K:535:LEU:CD2	2.67	0.43
1:K:593:GLY:HA3	1:K:603:TYR:O	2.19	0.43
1:L:353:PHE:CZ	1:L:395:TYR:HE2	2.11	0.43
1:L:375:ALA:HA	1:L:409:ILE:O	2.19	0.43
1:A:441:ILE:O	1:A:441:ILE:HG13	2.17	0.43
1:B:492:TYR:OH	1:B:611:LEU:O	2.34	0.43
1:C:408:ILE:H	1:C:408:ILE:CD1	2.30	0.43
1:C:593:GLY:HA3	1:C:603:TYR:O	2.19	0.43
1:D:353:PHE:CZ	1:D:395:TYR:CD2	3.05	0.43
1:F:465:LEU:HD21	1:F:480:ALA:CB	2.49	0.43
1:G:530:LEU:HD23	1:G:530:LEU:HA	1.82	0.43
1:G:534:VAL:C	1:G:535:LEU:HD23	2.43	0.43
1:H:401:LEU:HD12	1:H:401:LEU:HA	1.62	0.43
1:H:572:TYR:HB2	1:H:591:VAL:CG2	2.49	0.43
1:H:653:ILE:HG22	1:H:655:LEU:CD1	2.49	0.43
1:I:492:TYR:CE1	1:I:606:ILE:HB	2.54	0.43
1:I:533:ASP:O	1:I:535:LEU:CD2	2.67	0.43
1:I:557:PRO:HD3	1:I:580:LEU:HD11	2.01	0.43
1:K:557:PRO:HD3	1:K:580:LEU:HD11	2.01	0.43
1:A:408:ILE:H	1:A:408:ILE:CD1	2.30	0.43
1:B:464:MET:O	1:B:468:VAL:HG23	2.19	0.43
1:B:592:ILE:CD1	1:B:606:ILE:HD13	2.49	0.43
1:D:365:ASP:O	1:D:366:SER:C	2.54	0.43
1:D:422:VAL:HG11	1:D:475:VAL:HG13	2.01	0.43
1:D:447:TYR:HB2	1:D:467:TYR:CD1	2.54	0.43
1:D:484:MET:HG3	1:D:627:PRO:HG3	2.00	0.43
1:D:557:PRO:CG	1:D:580:LEU:HD11	2.37	0.43
1:D:592:ILE:CD1	1:D:606:ILE:HD13	2.49	0.43
1:E:440:ILE:CG1	1:E:441:ILE:N	2.79	0.43
1:F:388:GLN:O	1:F:392:ILE:HG22	2.19	0.43
1:G:533:ASP:O	1:G:535:LEU:CD2	2.67	0.43
1:J:388:GLN:O	1:J:392:ILE:HG22	2.19	0.43
1:J:648:PRO:CD	1:J:649:GLN:H	2.32	0.43
1:K:340:VAL:O	1:K:341:THR:CB	2.63	0.43
1:K:365:ASP:OD2	1:K:367:THR:HB	2.18	0.43
1:K:377:LYS:HA	1:K:378:PRO:HD3	1.84	0.43
1:K:524:VAL:HG22	1:K:529:GLY:C	2.42	0.43
1:L:388:GLN:O	1:L:392:ILE:HG22	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:600:ASP:O	1:L:600:ASP:CG	2.62	0.43
1:B:401:LEU:HA	1:B:401:LEU:HD12	1.62	0.42
1:B:465:LEU:HD21	1:B:480:ALA:CB	2.49	0.42
1:B:572:TYR:HB2	1:B:591:VAL:CG2	2.49	0.42
1:C:458:SER:HB2	1:C:632:ILE:O	2.19	0.42
1:D:465:LEU:HD21	1:D:480:ALA:CB	2.49	0.42
1:D:607:ALA:O	1:D:609:ILE:N	2.52	0.42
1:E:340:VAL:O	1:E:341:THR:CG2	2.64	0.42
1:E:516:SER:HB2	1:E:617:GLU:OE1	2.18	0.42
1:F:447:TYR:HB2	1:F:467:TYR:CD1	2.54	0.42
1:G:582:ASN:C	1:G:584:ASP:N	2.74	0.42
1:H:358:GLN:HB3	1:H:379:LYS:HA	1.99	0.42
1:H:511:GLU:O	1:H:511:GLU:HG3	2.17	0.42
1:H:542:SER:HB2	1:H:595:ILE:CD1	2.49	0.42
1:H:607:ALA:C	1:H:609:ILE:N	2.76	0.42
1:H:607:ALA:O	1:H:609:ILE:N	2.52	0.42
1:I:650:TYR:O	1:I:651:LEU:HB2	2.18	0.42
1:J:375:ALA:HA	1:J:409:ILE:O	2.19	0.42
1:J:414:LEU:HD23	1:J:414:LEU:HA	1.70	0.42
1:L:448:TYR:CG	1:L:642:PHE:HB2	2.53	0.42
1:C:469:ASP:HA	1:C:475:VAL:HG11	2.00	0.42
1:C:533:ASP:O	1:C:535:LEU:CD2	2.67	0.42
1:C:582:ASN:C	1:C:584:ASP:N	2.74	0.42
1:C:647:ARG:N	1:C:648:PRO:HD3	2.34	0.42
1:D:375:ALA:HA	1:D:409:ILE:O	2.19	0.42
1:F:653:ILE:HG22	1:F:655:LEU:CD1	2.49	0.42
1:G:623:LEU:HD12	1:G:623:LEU:HA	1.79	0.42
1:H:422:VAL:HG11	1:H:475:VAL:HG13	2.01	0.42
1:I:340:VAL:C	1:I:341:THR:CG2	2.62	0.42
1:I:454:ILE:HG23	1:I:455:PHE:N	2.34	0.42
1:I:660:GLN:HG3	1:I:661:LEU:H	1.77	0.42
1:J:464:MET:O	1:J:468:VAL:HG23	2.18	0.42
1:J:592:ILE:CD1	1:J:606:ILE:HD13	2.49	0.42
1:L:607:ALA:O	1:L:609:ILE:N	2.52	0.42
1:A:469:ASP:HA	1:A:475:VAL:HG11	2.01	0.42
1:A:595:ILE:HG23	1:A:602:ILE:HG12	1.99	0.42
1:A:661:LEU:CD1	1:A:662:GLU:CA	2.85	0.42
1:D:572:TYR:HB2	1:D:591:VAL:CG2	2.49	0.42
1:E:436:LEU:C	1:E:436:LEU:CD2	2.91	0.42
1:E:458:SER:HB2	1:E:632:ILE:O	2.19	0.42
1:E:661:LEU:HD22	1:E:661:LEU:HA	1.75	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:340:VAL:CA	1:G:345:TYR:CZ	2.82	0.42
1:G:454:ILE:HG23	1:G:455:PHE:N	2.34	0.42
1:I:346:ASP:OD1	1:I:362:THR:CB	2.67	0.42
1:I:365:ASP:OD2	1:I:367:THR:HB	2.18	0.42
1:J:461:LYS:O	1:J:461:LYS:HG2	2.18	0.42
1:J:484:MET:HG3	1:J:627:PRO:HG3	1.99	0.42
1:J:560:SER:HB2	1:J:588:LYS:NZ	2.33	0.42
1:K:647:ARG:N	1:K:648:PRO:HD3	2.34	0.42
1:L:464:MET:O	1:L:468:VAL:HG23	2.18	0.42
1:L:542:SER:HB2	1:L:595:ILE:CD1	2.50	0.42
1:B:600:ASP:CG	1:B:600:ASP:O	2.62	0.42
1:C:492:TYR:CE1	1:C:606:ILE:HB	2.54	0.42
1:D:658:ILE:HD12	1:D:658:ILE:HA	1.67	0.42
1:E:556:GLY:CA	1:E:589:TYR:CD1	3.01	0.42
1:F:557:PRO:CG	1:F:580:LEU:HD11	2.37	0.42
1:G:408:ILE:H	1:G:408:ILE:CD1	2.30	0.42
1:G:469:ASP:HA	1:G:475:VAL:HG11	2.00	0.42
1:G:661:LEU:HD22	1:G:661:LEU:HA	1.75	0.42
1:H:592:ILE:CD1	1:H:606:ILE:HD13	2.49	0.42
1:I:469:ASP:HA	1:I:475:VAL:HG11	2.01	0.42
1:I:516:SER:HB2	1:I:617:GLU:OE1	2.18	0.42
1:J:607:ALA:C	1:J:609:ILE:N	2.76	0.42
1:A:492:TYR:CE1	1:A:606:ILE:HB	2.54	0.42
1:A:526:PRO:O	1:A:527:ASP:C	2.59	0.42
1:C:454:ILE:HG23	1:C:455:PHE:N	2.34	0.42
1:D:388:GLN:O	1:D:392:ILE:HG22	2.19	0.42
1:D:400:ASN:HD22	1:E:662:GLU:HG2	1.02	0.42
1:E:593:GLY:HA3	1:E:603:TYR:O	2.19	0.42
1:F:414:LEU:HD23	1:F:414:LEU:HA	1.70	0.42
1:F:567:GLU:C	1:F:568:ASN:CG	2.86	0.42
1:G:492:TYR:CE1	1:G:606:ILE:HB	2.54	0.42
1:I:448:TYR:CZ	1:I:642:PHE:HB2	2.55	0.42
1:I:521:ARG:HE	1:I:521:ARG:HB2	1.71	0.42
1:J:494:THR:HG22	1:J:499:ILE:HG22	2.01	0.42
1:J:600:ASP:O	1:J:600:ASP:CG	2.62	0.42
1:J:607:ALA:O	1:J:609:ILE:N	2.52	0.42
1:A:593:GLY:HA3	1:A:603:TYR:O	2.19	0.42
1:C:524:VAL:HG22	1:C:529:GLY:C	2.42	0.42
1:D:402:ALA:O	1:D:403:PRO:C	2.59	0.42
1:D:545:ARG:CZ	1:D:597:TYR:HB3	2.50	0.42
1:E:636:ASP:C	1:E:638:SER:H	2.27	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:607:ALA:O	1:F:609:ILE:N	2.52	0.42
1:F:623:LEU:HD12	1:F:623:LEU:HA	1.86	0.42
1:G:448:TYR:CZ	1:G:642:PHE:HB2	2.55	0.42
1:K:650:TYR:O	1:K:651:LEU:HB2	2.18	0.42
1:L:465:LEU:HD21	1:L:480:ALA:CB	2.49	0.42
1:L:545:ARG:CZ	1:L:597:TYR:HB3	2.50	0.42
1:B:542:SER:HB2	1:B:595:ILE:CD1	2.49	0.42
1:B:607:ALA:O	1:B:609:ILE:N	2.52	0.42
1:D:648:PRO:CD	1:D:649:GLN:H	2.32	0.42
1:D:661:LEU:C	1:D:663:HIS:N	2.61	0.42
1:E:556:GLY:CA	1:E:589:TYR:HD1	2.32	0.42
1:G:346:ASP:OD1	1:G:362:THR:CB	2.67	0.42
1:H:408:ILE:O	1:H:409:ILE:HG23	2.09	0.42
1:I:556:GLY:CA	1:I:589:TYR:CD1	3.01	0.42
1:I:661:LEU:HA	1:I:661:LEU:HD22	1.75	0.42
1:J:436:LEU:O	1:J:437:GLU:C	2.63	0.42
1:J:447:TYR:HB2	1:J:467:TYR:CD1	2.54	0.42
1:J:542:SER:HB2	1:J:595:ILE:CD1	2.50	0.42
1:K:361:GLN:HA	1:K:361:GLN:HE21	1.84	0.42
1:K:526:PRO:O	1:K:527:ASP:C	2.59	0.42
1:C:385:THR:C	1:C:387:VAL:HG22	2.45	0.42
1:D:461:LYS:O	1:D:461:LYS:HG2	2.18	0.42
1:E:385:THR:C	1:E:387:VAL:HG22	2.45	0.42
1:E:448:TYR:HA	1:E:452:VAL:HB	2.02	0.42
1:E:448:TYR:CZ	1:E:642:PHE:HB2	2.55	0.42
1:F:422:VAL:HG11	1:F:475:VAL:HG13	2.01	0.42
1:F:436:LEU:O	1:F:437:GLU:C	2.63	0.42
1:F:600:ASP:CG	1:F:600:ASP:O	2.62	0.42
1:G:385:THR:C	1:G:387:VAL:HG22	2.45	0.42
1:I:385:THR:C	1:I:387:VAL:HG22	2.45	0.42
1:I:647:ARG:N	1:I:648:PRO:HD3	2.34	0.42
1:J:405:THR:HB	1:J:406:PRO:HD3	1.43	0.42
1:A:458:SER:HB2	1:A:632:ILE:O	2.19	0.42
1:A:595:ILE:HG23	1:A:602:ILE:HG23	2.02	0.42
1:B:367:THR:C	1:B:369:PRO:HD3	2.34	0.42
1:B:388:GLN:O	1:B:392:ILE:HG22	2.19	0.42
1:B:436:LEU:O	1:B:437:GLU:C	2.63	0.42
1:B:648:PRO:CD	1:B:649:GLN:H	2.32	0.42
1:C:346:ASP:OD1	1:C:362:THR:CB	2.67	0.42
1:C:502:ASN:ND2	1:C:635:ARG:CB	2.59	0.42
1:D:507:ASP:C	1:D:509:SER:H	2.28	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:346:ASP:OD1	1:E:362:THR:CB	2.67	0.42
1:E:647:ARG:N	1:E:648:PRO:HD3	2.34	0.42
1:F:545:ARG:CZ	1:F:597:TYR:HB3	2.50	0.42
1:F:572:TYR:HB2	1:F:591:VAL:CG2	2.49	0.42
1:G:377:LYS:HA	1:G:378:PRO:HD3	1.84	0.42
1:H:545:ARG:CZ	1:H:597:TYR:HB3	2.50	0.42
1:K:346:ASP:OD1	1:K:362:THR:CB	2.67	0.42
1:K:454:ILE:HG23	1:K:455:PHE:N	2.34	0.42
1:K:458:SER:HB2	1:K:632:ILE:O	2.19	0.42
1:K:492:TYR:CE1	1:K:606:ILE:HB	2.54	0.42
1:L:448:TYR:CZ	1:L:642:PHE:HB2	2.55	0.42
1:A:346:ASP:OD1	1:A:362:THR:CB	2.67	0.42
1:B:346:ASP:OD1	1:B:346:ASP:N	2.47	0.42
1:B:658:ILE:HD12	1:B:658:ILE:HA	1.67	0.42
1:C:424:TYR:O	1:C:658:ILE:HB	2.20	0.42
1:D:358:GLN:HE21	1:D:377:LYS:HZ2	1.67	0.42
1:E:557:PRO:HD3	1:E:580:LEU:HD11	2.01	0.42
1:E:660:GLN:HG3	1:E:661:LEU:H	1.77	0.42
1:G:556:GLY:CA	1:G:589:TYR:HD1	2.32	0.42
1:H:567:GLU:C	1:H:568:ASN:CG	2.86	0.42
1:H:658:ILE:CD1	1:H:658:ILE:N	2.74	0.42
1:I:636:ASP:C	1:I:638:SER:H	2.27	0.42
1:J:653:ILE:HG22	1:J:655:LEU:CD1	2.49	0.42
1:K:465:LEU:HD23	1:K:465:LEU:HA	1.84	0.42
1:L:422:VAL:HG11	1:L:475:VAL:HG13	2.01	0.42
1:L:507:ASP:C	1:L:509:SER:H	2.28	0.42
1:A:340:VAL:O	1:A:341:THR:CB	2.63	0.41
1:A:424:TYR:O	1:A:658:ILE:HB	2.20	0.41
1:A:524:VAL:HG22	1:A:529:GLY:C	2.42	0.41
1:A:557:PRO:HD3	1:A:580:LEU:HD11	2.01	0.41
1:A:660:GLN:HG3	1:A:661:LEU:H	1.77	0.41
1:B:375:ALA:HA	1:B:409:ILE:O	2.19	0.41
1:B:448:TYR:CZ	1:B:642:PHE:HB2	2.55	0.41
1:B:545:ARG:CZ	1:B:597:TYR:HB3	2.50	0.41
1:C:557:PRO:HD3	1:C:580:LEU:HD11	2.01	0.41
1:C:629:ASP:CG	1:D:664:HIS:NE2	2.78	0.41
1:D:494:THR:HG22	1:D:499:ILE:HG22	2.01	0.41
1:E:526:PRO:O	1:E:527:ASP:C	2.59	0.41
1:F:448:TYR:CZ	1:F:642:PHE:HB2	2.55	0.41
1:I:525:ASN:OD1	1:I:526:PRO:N	2.53	0.41
1:J:545:ARG:CZ	1:J:597:TYR:HB3	2.50	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:408:ILE:H	1:K:408:ILE:CD1	2.30	0.41
1:K:448:TYR:CZ	1:K:642:PHE:HB2	2.55	0.41
1:A:387:VAL:CG2	1:A:390:GLU:OE1	2.55	0.41
1:A:629:ASP:CG	1:B:664:HIS:NE2	2.78	0.41
1:C:448:TYR:HA	1:C:452:VAL:HB	2.02	0.41
1:C:556:GLY:CA	1:C:589:TYR:HD1	2.32	0.41
1:D:653:ILE:HG22	1:D:655:LEU:CD1	2.49	0.41
1:F:353:PHE:CZ	1:F:395:TYR:CD2	3.05	0.41
1:F:494:THR:HG22	1:F:499:ILE:HG22	2.01	0.41
1:F:642:PHE:CZ	1:F:644:ASN:HB2	2.55	0.41
1:G:525:ASN:OD1	1:G:526:PRO:N	2.53	0.41
1:I:465:LEU:HD23	1:I:465:LEU:HA	1.84	0.41
1:I:493:LYS:HE3	1:I:493:LYS:HB2	1.91	0.41
1:I:629:ASP:CG	1:J:664:HIS:NE2	2.78	0.41
1:J:496:GLU:N	1:J:496:GLU:OE1	2.30	0.41
1:J:507:ASP:C	1:J:509:SER:H	2.28	0.41
1:L:401:LEU:HD12	1:L:401:LEU:HA	1.63	0.41
1:A:387:VAL:C	1:A:388:GLN:HG3	2.44	0.41
1:A:510:MET:HB3	1:A:542:SER:HG	1.85	0.41
1:A:550:ILE:HG12	1:A:596:ASN:HA	2.02	0.41
1:C:361:GLN:HA	1:C:361:GLN:HE21	1.84	0.41
1:C:595:ILE:HG23	1:C:602:ILE:HG23	2.02	0.41
1:E:377:LYS:HA	1:E:378:PRO:HD3	1.84	0.41
1:E:525:ASN:OD1	1:E:526:PRO:N	2.53	0.41
1:F:408:ILE:O	1:F:409:ILE:HG23	2.09	0.41
1:F:492:TYR:OH	1:F:611:LEU:O	2.34	0.41
1:G:361:GLN:HG2	1:G:455:PHE:CB	2.50	0.41
1:G:502:ASN:ND2	1:G:635:ARG:CB	2.59	0.41
1:G:557:PRO:HD3	1:G:580:LEU:HD11	2.01	0.41
1:H:494:THR:HG22	1:H:499:ILE:HG22	2.01	0.41
1:H:642:PHE:CZ	1:H:644:ASN:HB2	2.55	0.41
1:I:401:LEU:HD21	1:I:402:ALA:HB2	1.96	0.41
1:I:458:SER:HB2	1:I:632:ILE:O	2.19	0.41
1:I:550:ILE:HG12	1:I:596:ASN:HA	2.02	0.41
1:I:556:GLY:CA	1:I:589:TYR:HD1	2.32	0.41
1:J:408:ILE:O	1:J:409:ILE:HG23	2.09	0.41
1:K:387:VAL:C	1:K:388:GLN:HG3	2.44	0.41
1:K:502:ASN:ND2	1:K:635:ARG:CB	2.59	0.41
1:L:436:LEU:O	1:L:437:GLU:C	2.63	0.41
1:A:382:LEU:CD1	1:A:647:ARG:HH21	2.33	0.41
1:C:369:PRO:C	1:C:371:TYR:H	2.29	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:384:LEU:HD23	1:C:384:LEU:HA	1.85	0.41
1:C:436:LEU:C	1:C:436:LEU:CD2	2.91	0.41
1:C:448:TYR:CZ	1:C:642:PHE:HB2	2.55	0.41
1:E:454:ILE:HG23	1:E:455:PHE:N	2.34	0.41
1:G:448:TYR:HA	1:G:452:VAL:HB	2.02	0.41
1:G:458:SER:HB2	1:G:632:ILE:O	2.19	0.41
1:G:556:GLY:CA	1:G:589:TYR:CD1	3.01	0.41
1:H:447:TYR:HB2	1:H:467:TYR:CD1	2.54	0.41
1:I:387:VAL:CG2	1:I:390:GLU:OE1	2.55	0.41
1:K:385:THR:C	1:K:387:VAL:HG22	2.45	0.41
1:K:525:ASN:OD1	1:K:526:PRO:N	2.53	0.41
1:K:629:ASP:CG	1:L:664:HIS:NE2	2.78	0.41
1:A:453:GLU:O	1:A:454:ILE:HG13	2.21	0.41
1:C:382:LEU:CD1	1:C:647:ARG:HH21	2.33	0.41
1:D:542:SER:HB2	1:D:595:ILE:CD1	2.50	0.41
1:E:453:GLU:O	1:E:454:ILE:HG13	2.21	0.41
1:G:629:ASP:CG	1:H:664:HIS:NE2	2.78	0.41
1:G:647:ARG:N	1:G:648:PRO:HD3	2.34	0.41
1:G:656:GLU:HA	1:G:657:PRO:HD3	1.82	0.41
1:J:422:VAL:HG11	1:J:475:VAL:HG13	2.01	0.41
1:K:550:ILE:HG12	1:K:596:ASN:HA	2.02	0.41
1:K:556:GLY:CA	1:K:589:TYR:HD1	2.32	0.41
1:K:595:ILE:HG23	1:K:602:ILE:HG23	2.02	0.41
1:L:642:PHE:CZ	1:L:644:ASN:HB2	2.55	0.41
1:A:448:TYR:CZ	1:A:642:PHE:HB2	2.55	0.41
1:B:396:LEU:HD13	1:B:405:THR:OG1	2.21	0.41
1:B:642:PHE:CZ	1:B:644:ASN:HB2	2.55	0.41
1:C:384:LEU:O	1:C:385:THR:C	2.64	0.41
1:C:636:ASP:C	1:C:638:SER:H	2.27	0.41
1:E:629:ASP:CG	1:F:664:HIS:NE2	2.78	0.41
1:G:595:ILE:HG23	1:G:602:ILE:HG23	2.02	0.41
1:H:375:ALA:HA	1:H:409:ILE:CD1	2.51	0.41
1:H:396:LEU:HD13	1:H:405:THR:OG1	2.21	0.41
1:H:448:TYR:CZ	1:H:642:PHE:HB2	2.55	0.41
1:I:440:ILE:CG1	1:I:441:ILE:N	2.79	0.41
1:J:448:TYR:CZ	1:J:642:PHE:HB2	2.55	0.41
1:L:367:THR:CG2	1:L:368:LYS:N	2.44	0.41
1:L:447:TYR:HB2	1:L:467:TYR:CD1	2.54	0.41
1:A:340:VAL:O	1:A:341:THR:CG2	2.64	0.41
1:A:361:GLN:HG2	1:A:455:PHE:CB	2.50	0.41
1:A:647:ARG:N	1:A:648:PRO:HD3	2.34	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:408:ILE:O	1:B:408:ILE:CG1	2.64	0.41
1:B:455:PHE:O	1:B:456:ASN:CB	2.69	0.41
1:B:647:ARG:HD3	1:B:649:GLN:CG	2.51	0.41
1:C:377:LYS:HA	1:C:378:PRO:HD3	1.84	0.41
1:C:521:ARG:HE	1:C:521:ARG:HB2	1.71	0.41
1:E:361:GLN:HG2	1:E:455:PHE:CB	2.50	0.41
1:F:507:ASP:C	1:F:509:SER:H	2.28	0.41
1:G:636:ASP:C	1:G:638:SER:H	2.27	0.41
1:H:358:GLN:HE21	1:H:377:LYS:NZ	2.19	0.41
1:I:424:TYR:O	1:I:658:ILE:HB	2.20	0.41
1:I:424:TYR:HB2	1:I:475:VAL:H	1.85	0.41
1:I:595:ILE:HG23	1:I:602:ILE:HG23	2.02	0.41
1:J:642:PHE:CZ	1:J:644:ASN:HB2	2.55	0.41
1:K:361:GLN:HG2	1:K:455:PHE:CB	2.50	0.41
1:K:382:LEU:CD1	1:K:647:ARG:HH21	2.33	0.41
1:K:424:TYR:HB2	1:K:475:VAL:H	1.85	0.41
1:L:375:ALA:HA	1:L:409:ILE:CD1	2.51	0.41
1:L:455:PHE:O	1:L:456:ASN:CB	2.69	0.41
1:L:663:HIS:HE1	1:L:665:HIS:O	2.04	0.41
1:A:369:PRO:C	1:A:371:TYR:H	2.29	0.41
1:A:385:THR:C	1:A:387:VAL:HG22	2.45	0.41
1:B:375:ALA:HA	1:B:409:ILE:CD1	2.51	0.41
1:B:663:HIS:HE1	1:B:665:HIS:O	2.04	0.41
1:C:361:GLN:HG2	1:C:455:PHE:CB	2.50	0.41
1:C:387:VAL:C	1:C:388:GLN:HG3	2.44	0.41
1:E:369:PRO:C	1:E:371:TYR:H	2.29	0.41
1:E:384:LEU:O	1:E:385:THR:C	2.64	0.41
1:E:523:VAL:HG12	1:E:534:VAL:HG22	2.03	0.41
1:F:437:GLU:O	1:F:438:GLY:C	2.64	0.41
1:F:542:SER:HB2	1:F:595:ILE:CD1	2.50	0.41
1:G:369:PRO:C	1:G:371:TYR:H	2.29	0.41
1:G:453:GLU:O	1:G:454:ILE:HG13	2.21	0.41
1:G:595:ILE:CG2	1:G:602:ILE:HG12	2.51	0.41
1:K:448:TYR:HA	1:K:452:VAL:HB	2.02	0.41
1:A:414:LEU:HD23	1:A:414:LEU:HA	1.91	0.41
1:A:448:TYR:HA	1:A:452:VAL:HB	2.02	0.41
1:A:455:PHE:O	1:A:456:ASN:C	2.64	0.41
1:A:523:VAL:HG22	1:A:524:VAL:H	1.86	0.41
1:B:448:TYR:CD2	1:B:642:PHE:HB2	2.56	0.41
1:B:507:ASP:C	1:B:509:SER:H	2.28	0.41
1:C:420:LEU:HG	1:C:422:VAL:HG23	2.03	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:523:VAL:HG12	1:C:534:VAL:HG22	2.03	0.41
1:D:396:LEU:HD13	1:D:405:THR:OG1	2.21	0.41
1:D:448:TYR:CZ	1:D:642:PHE:HB2	2.55	0.41
1:D:455:PHE:O	1:D:456:ASN:CB	2.69	0.41
1:D:642:PHE:CZ	1:D:644:ASN:HB2	2.55	0.41
1:E:353:PHE:O	1:E:355:SER:HB3	2.21	0.41
1:E:523:VAL:HG22	1:E:524:VAL:H	1.86	0.41
1:F:358:GLN:HE21	1:F:377:LYS:NZ	2.19	0.41
1:F:405:THR:OG1	1:F:406:PRO:HD2	2.15	0.41
1:G:361:GLN:HA	1:G:361:GLN:HE21	1.84	0.41
1:G:382:LEU:CD1	1:G:647:ARG:HH21	2.33	0.41
1:G:523:VAL:HG12	1:G:534:VAL:HG22	2.03	0.41
1:H:365:ASP:C	1:H:367:THR:CG2	2.90	0.41
1:H:448:TYR:CD2	1:H:642:PHE:HB2	2.56	0.41
1:I:369:PRO:C	1:I:371:TYR:H	2.29	0.41
1:J:455:PHE:O	1:J:456:ASN:CB	2.69	0.41
1:K:455:PHE:O	1:K:456:ASN:C	2.64	0.41
1:K:660:GLN:HG3	1:K:661:LEU:H	1.77	0.41
1:L:651:LEU:HD23	1:L:651:LEU:HA	1.83	0.41
1:A:454:ILE:HG23	1:A:455:PHE:N	2.34	0.41
1:B:342:ALA:C	1:B:346:ASP:CG	2.85	0.41
1:B:353:PHE:CZ	1:B:395:TYR:CD2	3.05	0.41
1:C:353:PHE:O	1:C:355:SER:HB3	2.21	0.41
1:C:647:ARG:CG	1:C:650:TYR:CD1	3.04	0.41
1:D:385:THR:HG23	1:D:387:VAL:HG13	2.03	0.41
1:E:424:TYR:O	1:E:658:ILE:HB	2.20	0.41
1:E:571:PRO:O	1:E:572:TYR:C	2.63	0.41
1:H:410:SER:HA	1:H:411:PRO:HD3	1.81	0.41
1:J:541:VAL:HG12	1:J:542:SER:H	1.86	0.41
1:K:523:VAL:HG12	1:K:534:VAL:HG22	2.03	0.41
1:L:396:LEU:HD13	1:L:405:THR:OG1	2.21	0.41
1:L:410:SER:HA	1:L:411:PRO:HD3	1.81	0.41
1:A:636:ASP:C	1:A:638:SER:H	2.27	0.40
1:C:391:ASP:C	1:C:393:LYS:N	2.79	0.40
1:C:455:PHE:O	1:C:456:ASN:C	2.64	0.40
1:C:516:SER:H	1:C:617:GLU:CD	2.29	0.40
1:C:525:ASN:OD1	1:C:526:PRO:N	2.53	0.40
1:D:663:HIS:HE1	1:D:665:HIS:O	2.04	0.40
1:F:358:GLN:HE21	1:F:377:LYS:HZ2	1.69	0.40
1:F:396:LEU:HD13	1:F:405:THR:OG1	2.21	0.40
1:H:437:GLU:O	1:H:438:GLY:C	2.64	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:455:PHE:O	1:H:456:ASN:CB	2.69	0.40
1:H:658:ILE:HD12	1:H:658:ILE:HA	1.67	0.40
1:I:391:ASP:C	1:I:393:LYS:N	2.79	0.40
1:I:420:LEU:HG	1:I:422:VAL:HG23	2.03	0.40
1:I:516:SER:H	1:I:617:GLU:CD	2.29	0.40
1:K:391:ASP:C	1:K:393:LYS:N	2.79	0.40
1:K:424:TYR:O	1:K:658:ILE:HB	2.20	0.40
1:K:523:VAL:HG22	1:K:524:VAL:H	1.86	0.40
1:K:636:ASP:C	1:K:638:SER:H	2.27	0.40
1:L:556:GLY:HA2	1:L:557:PRO:HD3	1.90	0.40
1:A:355:SER:OG	1:A:356:ILE:N	2.55	0.40
1:A:623:LEU:HD12	1:A:623:LEU:HA	1.79	0.40
1:B:533:ASP:O	1:B:535:LEU:HG	2.22	0.40
1:B:557:PRO:CG	1:B:580:LEU:HD11	2.37	0.40
1:D:358:GLN:HE21	1:D:377:LYS:NZ	2.19	0.40
1:D:372:ALA:O	1:D:406:PRO:HA	2.21	0.40
1:E:391:ASP:C	1:E:393:LYS:N	2.79	0.40
1:F:372:ALA:O	1:F:406:PRO:HA	2.21	0.40
1:G:496:GLU:CD	1:G:496:GLU:H	2.30	0.40
1:G:550:ILE:HG12	1:G:596:ASN:HA	2.02	0.40
1:H:647:ARG:HD3	1:H:649:GLN:CG	2.51	0.40
1:I:453:GLU:O	1:I:454:ILE:HG13	2.21	0.40
1:J:375:ALA:HA	1:J:409:ILE:CD1	2.51	0.40
1:J:448:TYR:CD2	1:J:642:PHE:HB2	2.56	0.40
1:K:353:PHE:O	1:K:355:SER:HB3	2.21	0.40
1:K:420:LEU:HG	1:K:422:VAL:HG23	2.03	0.40
1:K:453:GLU:O	1:K:454:ILE:HG13	2.21	0.40
1:K:483:GLN:HG2	1:K:626:ASP:HB3	2.04	0.40
1:L:385:THR:HG23	1:L:387:VAL:HG13	2.04	0.40
1:L:623:LEU:HA	1:L:623:LEU:HD12	1.86	0.40
1:A:424:TYR:HB2	1:A:475:VAL:H	1.85	0.40
1:A:595:ILE:CG2	1:A:602:ILE:HG12	2.51	0.40
1:B:372:ALA:O	1:B:406:PRO:HA	2.21	0.40
1:B:422:VAL:HG11	1:B:475:VAL:HG13	2.01	0.40
1:C:626:ASP:HA	1:C:627:PRO:HD3	1.65	0.40
1:D:375:ALA:HA	1:D:409:ILE:CD1	2.51	0.40
1:D:448:TYR:CD2	1:D:642:PHE:HB2	2.56	0.40
1:D:647:ARG:HD3	1:D:649:GLN:CG	2.51	0.40
1:E:550:ILE:HG12	1:E:596:ASN:HA	2.02	0.40
1:E:647:ARG:CG	1:E:650:TYR:CD1	3.04	0.40
1:F:385:THR:HG23	1:F:387:VAL:HG13	2.04	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:448:TYR:CD2	1:F:642:PHE:HB2	2.56	0.40
1:F:663:HIS:HE1	1:F:665:HIS:O	2.04	0.40
1:G:424:TYR:O	1:G:658:ILE:HB	2.20	0.40
1:G:465:LEU:HA	1:G:465:LEU:HD23	1.84	0.40
1:H:348:PHE:C	1:H:350:SER:N	2.80	0.40
1:I:629:ASP:OD1	1:I:629:ASP:N	2.55	0.40
1:J:358:GLN:HE21	1:J:377:LYS:NZ	2.19	0.40
1:J:396:LEU:HD13	1:J:405:THR:OG1	2.21	0.40
1:J:453:GLU:C	1:J:454:ILE:HG13	2.47	0.40
1:J:570:GLN:O	1:J:571:PRO:C	2.65	0.40
1:L:448:TYR:CD2	1:L:642:PHE:HB2	2.56	0.40
1:L:648:PRO:CD	1:L:649:GLN:H	2.32	0.40
1:L:659:SER:HB3	1:L:660:GLN:H	1.78	0.40
1:A:518:ASN:OD1	1:A:533:ASP:CG	2.65	0.40
1:A:523:VAL:HG12	1:A:534:VAL:HG22	2.03	0.40
1:A:525:ASN:OD1	1:A:526:PRO:N	2.53	0.40
1:A:556:GLY:CA	1:A:589:TYR:HD1	2.32	0.40
1:B:348:PHE:C	1:B:350:SER:N	2.80	0.40
1:B:486:ARG:HH22	1:B:638:SER:HB3	1.86	0.40
1:C:340:VAL:O	1:C:341:THR:CG2	2.64	0.40
1:C:629:ASP:OD1	1:C:629:ASP:N	2.55	0.40
1:D:408:ILE:O	1:D:408:ILE:CG1	2.64	0.40
1:D:533:ASP:O	1:D:535:LEU:HG	2.22	0.40
1:F:356:ILE:O	1:F:379:LYS:N	2.48	0.40
1:G:340:VAL:O	1:G:341:THR:CG2	2.64	0.40
1:G:455:PHE:O	1:G:456:ASN:C	2.64	0.40
1:H:414:LEU:HD23	1:H:414:LEU:HA	1.70	0.40
1:H:486:ARG:HH22	1:H:638:SER:HB3	1.86	0.40
1:H:507:ASP:C	1:H:509:SER:H	2.28	0.40
1:H:663:HIS:HE1	1:H:665:HIS:O	2.04	0.40
1:I:453:GLU:HG2	1:I:641:VAL:HA	2.04	0.40
1:I:523:VAL:HG22	1:I:524:VAL:H	1.86	0.40
1:K:369:PRO:C	1:K:371:TYR:H	2.29	0.40
1:K:629:ASP:OD1	1:K:629:ASP:N	2.55	0.40
1:A:459:PHE:CD2	1:A:459:PHE:C	3.00	0.40
1:B:437:GLU:O	1:B:438:GLY:C	2.64	0.40
1:C:424:TYR:HB2	1:C:475:VAL:H	1.85	0.40
1:C:518:ASN:OD1	1:C:533:ASP:CG	2.65	0.40
1:C:523:VAL:HG22	1:C:524:VAL:H	1.86	0.40
1:C:595:ILE:CG2	1:C:602:ILE:HG12	2.51	0.40
1:C:658:ILE:HD13	1:C:658:ILE:HA	1.52	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:453:GLU:C	1:D:454:ILE:HG13	2.47	0.40
1:E:595:ILE:HG23	1:E:602:ILE:HG23	2.02	0.40
1:F:375:ALA:HA	1:F:409:ILE:CD1	2.51	0.40
1:F:455:PHE:O	1:F:456:ASN:CB	2.69	0.40
1:G:424:TYR:HB2	1:G:475:VAL:H	1.85	0.40
1:G:464:MET:CG	1:G:465:LEU:N	2.82	0.40
1:H:465:LEU:HD21	1:H:480:ALA:CB	2.49	0.40
1:I:459:PHE:CD2	1:I:459:PHE:C	3.00	0.40
1:I:464:MET:CG	1:I:465:LEU:N	2.82	0.40
1:J:465:LEU:HD21	1:J:480:ALA:CB	2.49	0.40
1:J:507:ASP:C	1:J:509:SER:N	2.80	0.40
1:J:658:ILE:HD12	1:J:658:ILE:HA	1.67	0.40
1:K:349:VAL:C	1:K:351:GLU:N	2.71	0.40
1:K:355:SER:OG	1:K:356:ILE:N	2.55	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	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15
1	B	322/335 (96%)	261 (81%)	42 (13%)	19 (6%)	1	13
1	C	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15
1	D	322/335 (96%)	262 (81%)	41 (13%)	19 (6%)	1	13
1	E	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15
1	F	322/335 (96%)	262 (81%)	41 (13%)	19 (6%)	1	13
1	G	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15
1	H	322/335 (96%)	262 (81%)	41 (13%)	19 (6%)	1	13
1	I	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	J	322/335 (96%)	261 (81%)	42 (13%)	19 (6%)	1	13
1	K	324/335 (97%)	260 (80%)	47 (14%)	17 (5%)	1	15
1	L	322/335 (96%)	261 (81%)	42 (13%)	19 (6%)	1	13
All	All	3876/4020 (96%)	3129 (81%)	531 (14%)	216 (6%)	2	14

All (216) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	348	PHE
1	A	354	GLY
1	A	389	ARG
1	A	390	GLU
1	A	526	PRO
1	A	576	ASP
1	A	635	ARG
1	A	662	GLU
1	B	368	LYS
1	B	406	PRO
1	B	556	GLY
1	B	568	ASN
1	B	593	GLY
1	B	660	GLN
1	B	661	LEU
1	B	662	GLU
1	C	348	PHE
1	C	354	GLY
1	C	389	ARG
1	C	390	GLU
1	C	526	PRO
1	C	576	ASP
1	C	635	ARG
1	C	662	GLU
1	D	368	LYS
1	D	406	PRO
1	D	556	GLY
1	D	568	ASN
1	D	593	GLY
1	D	660	GLN
1	D	661	LEU
1	D	662	GLU
1	E	348	PHE

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	E	354	GLY
1	E	389	ARG
1	E	390	GLU
1	E	526	PRO
1	E	576	ASP
1	E	635	ARG
1	E	662	GLU
1	F	368	LYS
1	F	406	PRO
1	F	556	GLY
1	F	568	ASN
1	F	593	GLY
1	F	660	GLN
1	F	661	LEU
1	F	662	GLU
1	G	348	PHE
1	G	354	GLY
1	G	389	ARG
1	G	390	GLU
1	G	526	PRO
1	G	576	ASP
1	G	635	ARG
1	G	662	GLU
1	H	368	LYS
1	H	406	PRO
1	H	556	GLY
1	H	568	ASN
1	H	593	GLY
1	H	660	GLN
1	H	661	LEU
1	H	662	GLU
1	I	348	PHE
1	I	354	GLY
1	I	389	ARG
1	I	390	GLU
1	I	526	PRO
1	I	576	ASP
1	I	635	ARG
1	I	662	GLU
1	J	368	LYS
1	J	406	PRO
1	J	556	GLY

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	J	568	ASN
1	J	593	GLY
1	J	660	GLN
1	J	661	LEU
1	J	662	GLU
1	K	348	PHE
1	K	354	GLY
1	K	389	ARG
1	K	390	GLU
1	K	526	PRO
1	K	576	ASP
1	K	635	ARG
1	K	662	GLU
1	L	368	LYS
1	L	406	PRO
1	L	556	GLY
1	L	568	ASN
1	L	593	GLY
1	L	660	GLN
1	L	661	LEU
1	L	662	GLU
1	A	402	ALA
1	A	474	SER
1	A	490	ASN
1	A	664	HIS
1	B	367	THR
1	B	560	SER
1	B	600	ASP
1	C	402	ALA
1	C	474	SER
1	C	490	ASN
1	C	664	HIS
1	D	367	THR
1	D	560	SER
1	D	600	ASP
1	E	402	ALA
1	E	474	SER
1	E	490	ASN
1	E	664	HIS
1	F	367	THR
1	F	560	SER
1	F	600	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	G	402	ALA
1	G	474	SER
1	G	490	ASN
1	G	664	HIS
1	H	367	THR
1	H	560	SER
1	H	600	ASP
1	I	402	ALA
1	I	474	SER
1	I	490	ASN
1	I	664	HIS
1	J	367	THR
1	J	560	SER
1	J	600	ASP
1	K	402	ALA
1	K	474	SER
1	K	490	ASN
1	K	664	HIS
1	L	367	THR
1	L	560	SER
1	L	600	ASP
1	A	341	THR
1	A	587	ASP
1	A	651	LEU
1	B	348	PHE
1	B	574	GLY
1	B	608	LYS
1	B	663	HIS
1	C	341	THR
1	C	587	ASP
1	C	651	LEU
1	D	348	PHE
1	D	574	GLY
1	D	608	LYS
1	D	663	HIS
1	E	341	THR
1	E	587	ASP
1	E	651	LEU
1	F	348	PHE
1	F	574	GLY
1	F	608	LYS
1	F	663	HIS

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	G	341	THR
1	G	587	ASP
1	G	651	LEU
1	H	348	PHE
1	H	574	GLY
1	H	608	LYS
1	H	663	HIS
1	I	341	THR
1	I	587	ASP
1	I	651	LEU
1	J	348	PHE
1	J	574	GLY
1	J	608	LYS
1	J	663	HIS
1	K	341	THR
1	K	587	ASP
1	K	651	LEU
1	L	348	PHE
1	L	574	GLY
1	L	608	LYS
1	L	663	HIS
1	B	349	VAL
1	B	625	SER
1	D	349	VAL
1	D	625	SER
1	F	349	VAL
1	F	625	SER
1	H	349	VAL
1	H	625	SER
1	J	349	VAL
1	J	625	SER
1	L	349	VAL
1	L	625	SER
1	B	496	GLU
1	D	496	GLU
1	J	496	GLU
1	L	496	GLU
1	A	475	VAL
1	C	475	VAL
1	E	475	VAL
1	F	496	GLU
1	G	475	VAL

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	H	496	GLU
1	I	475	VAL
1	K	475	VAL
1	B	524	VAL
1	D	524	VAL
1	F	524	VAL
1	H	524	VAL
1	J	524	VAL
1	L	524	VAL
1	A	392	ILE
1	C	392	ILE
1	E	392	ILE
1	G	392	ILE
1	I	392	ILE
1	K	392	ILE

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	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	B	293/304 (96%)	228 (78%)	65 (22%)	1	6
1	C	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	D	293/304 (96%)	228 (78%)	65 (22%)	1	6
1	E	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	F	293/304 (96%)	227 (78%)	66 (22%)	1	6
1	G	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	H	293/304 (96%)	228 (78%)	65 (22%)	1	6
1	I	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	J	293/304 (96%)	228 (78%)	65 (22%)	1	6
1	K	295/304 (97%)	236 (80%)	59 (20%)	1	7
1	L	293/304 (96%)	227 (78%)	66 (22%)	1	6

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
All	All	3528/3648 (97%)	2782 (79%)	746 (21%)	3 6

All (746) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	340	VAL
1	A	341	THR
1	A	343	THR
1	A	345	TYR
1	A	348	PHE
1	A	355	SER
1	A	356	ILE
1	A	357	ILE
1	A	360	VAL
1	A	361	GLN
1	A	364	THR
1	A	367	THR
1	A	374	ILE
1	A	388	GLN
1	A	391	ASP
1	A	396	LEU
1	A	401	LEU
1	A	405	THR
1	A	414	LEU
1	A	416	ILE
1	A	423	THR
1	A	427	ASN
1	A	429	LEU
1	A	441	ILE
1	A	464	MET
1	A	466	THR
1	A	474	SER
1	A	475	VAL
1	A	486	ARG
1	A	489	GLN
1	A	496	GLU
1	A	499	ILE
1	A	500	LYS
1	A	522	LYS
1	A	528	THR
1	A	532	GLU
1	A	533	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	A	536	TYR
1	A	538	VAL
1	A	547	SER
1	A	580	LEU
1	A	587	ASP
1	A	595	ILE
1	A	612	THR
1	A	613	SER
1	A	619	GLN
1	A	622	GLU
1	A	626	ASP
1	A	628	THR
1	A	634	THR
1	A	635	ARG
1	A	649	GLN
1	A	652	THR
1	A	658	ILE
1	A	659	SER
1	A	660	GLN
1	A	661	LEU
1	A	662	GLU
1	A	663	HIS
1	B	344	ASP
1	B	346	ASP
1	B	356	ILE
1	B	357	ILE
1	B	360	VAL
1	B	361	GLN
1	B	364	THR
1	B	366	SER
1	B	367	THR
1	B	374	ILE
1	B	379	LYS
1	B	380	SER
1	B	386	THR
1	B	387	VAL
1	B	404	ILE
1	B	407	SER
1	B	408	ILE
1	B	409	ILE
1	B	414	LEU
1	B	417	LYS

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	B	419	ASN
1	B	423	THR
1	B	427	ASN
1	B	429	LEU
1	B	436	LEU
1	B	446	ARG
1	B	457	SER
1	B	462	SER
1	B	464	MET
1	B	466	THR
1	B	474	SER
1	B	476	ILE
1	B	481	THR
1	B	485	VAL
1	B	486	ARG
1	B	493	LYS
1	B	496	GLU
1	B	499	ILE
1	B	500	LYS
1	B	516	SER
1	B	522	LYS
1	B	535	LEU
1	B	538	VAL
1	B	540	ILE
1	B	547	SER
1	B	548	LYS
1	B	554	ILE
1	B	566	ASN
1	B	567	GLU
1	B	580	LEU
1	B	595	ILE
1	B	596	ASN
1	B	601	VAL
1	B	612	THR
1	B	613	SER
1	B	614	GLU
1	B	632	ILE
1	B	646	LEU
1	B	649	GLN
1	B	651	LEU
1	B	652	THR
1	B	658	ILE

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	B	659	SER
1	B	661	LEU
1	B	664	HIS
1	C	340	VAL
1	C	341	THR
1	C	343	THR
1	C	345	TYR
1	C	348	PHE
1	C	355	SER
1	C	356	ILE
1	C	357	ILE
1	C	360	VAL
1	C	361	GLN
1	C	364	THR
1	C	367	THR
1	C	374	ILE
1	C	388	GLN
1	C	391	ASP
1	C	396	LEU
1	C	401	LEU
1	C	405	THR
1	C	414	LEU
1	C	416	ILE
1	C	423	THR
1	C	427	ASN
1	C	429	LEU
1	C	441	ILE
1	C	464	MET
1	C	466	THR
1	C	474	SER
1	C	475	VAL
1	C	486	ARG
1	C	489	GLN
1	C	496	GLU
1	C	499	ILE
1	C	500	LYS
1	C	522	LYS
1	C	528	THR
1	C	532	GLU
1	C	533	ASP
1	C	536	TYR
1	C	538	VAL

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	C	547	SER
1	C	580	LEU
1	C	587	ASP
1	C	595	ILE
1	C	612	THR
1	C	613	SER
1	C	619	GLN
1	C	622	GLU
1	C	626	ASP
1	C	628	THR
1	C	634	THR
1	C	635	ARG
1	C	649	GLN
1	C	652	THR
1	C	658	ILE
1	C	659	SER
1	C	660	GLN
1	C	661	LEU
1	C	662	GLU
1	C	663	HIS
1	D	344	ASP
1	D	346	ASP
1	D	356	ILE
1	D	357	ILE
1	D	360	VAL
1	D	361	GLN
1	D	364	THR
1	D	366	SER
1	D	367	THR
1	D	374	ILE
1	D	379	LYS
1	D	380	SER
1	D	386	THR
1	D	387	VAL
1	D	404	ILE
1	D	407	SER
1	D	408	ILE
1	D	409	ILE
1	D	414	LEU
1	D	417	LYS
1	D	419	ASN
1	D	423	THR

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	427	ASN
1	D	429	LEU
1	D	436	LEU
1	D	446	ARG
1	D	457	SER
1	D	462	SER
1	D	464	MET
1	D	466	THR
1	D	474	SER
1	D	476	ILE
1	D	481	THR
1	D	485	VAL
1	D	486	ARG
1	D	493	LYS
1	D	496	GLU
1	D	499	ILE
1	D	500	LYS
1	D	516	SER
1	D	522	LYS
1	D	535	LEU
1	D	538	VAL
1	D	540	ILE
1	D	547	SER
1	D	548	LYS
1	D	554	ILE
1	D	566	ASN
1	D	567	GLU
1	D	580	LEU
1	D	595	ILE
1	D	596	ASN
1	D	601	VAL
1	D	612	THR
1	D	613	SER
1	D	614	GLU
1	D	632	ILE
1	D	646	LEU
1	D	649	GLN
1	D	651	LEU
1	D	652	THR
1	D	658	ILE
1	D	659	SER
1	D	661	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	664	HIS
1	E	340	VAL
1	E	341	THR
1	E	343	THR
1	E	345	TYR
1	E	348	PHE
1	E	355	SER
1	E	356	ILE
1	E	357	ILE
1	E	360	VAL
1	E	361	GLN
1	E	364	THR
1	E	367	THR
1	E	374	ILE
1	E	388	GLN
1	E	391	ASP
1	E	396	LEU
1	E	401	LEU
1	E	405	THR
1	E	414	LEU
1	E	416	ILE
1	E	423	THR
1	E	427	ASN
1	E	429	LEU
1	E	441	ILE
1	E	464	MET
1	E	466	THR
1	E	474	SER
1	E	475	VAL
1	E	486	ARG
1	E	489	GLN
1	E	496	GLU
1	E	499	ILE
1	E	500	LYS
1	E	522	LYS
1	E	528	THR
1	E	532	GLU
1	E	533	ASP
1	E	536	TYR
1	E	538	VAL
1	E	547	SER
1	E	580	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	E	587	ASP
1	E	595	ILE
1	E	612	THR
1	E	613	SER
1	E	619	GLN
1	E	622	GLU
1	E	626	ASP
1	E	628	THR
1	E	634	THR
1	E	635	ARG
1	E	649	GLN
1	E	652	THR
1	E	658	ILE
1	E	659	SER
1	E	660	GLN
1	E	661	LEU
1	E	662	GLU
1	E	663	HIS
1	F	344	ASP
1	F	346	ASP
1	F	356	ILE
1	F	357	ILE
1	F	360	VAL
1	F	361	GLN
1	F	364	THR
1	F	366	SER
1	F	367	THR
1	F	374	ILE
1	F	379	LYS
1	F	380	SER
1	F	386	THR
1	F	387	VAL
1	F	404	ILE
1	F	407	SER
1	F	408	ILE
1	F	409	ILE
1	F	414	LEU
1	F	417	LYS
1	F	419	ASN
1	F	423	THR
1	F	427	ASN
1	F	429	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	F	436	LEU
1	F	446	ARG
1	F	457	SER
1	F	462	SER
1	F	464	MET
1	F	466	THR
1	F	474	SER
1	F	476	ILE
1	F	481	THR
1	F	485	VAL
1	F	486	ARG
1	F	493	LYS
1	F	496	GLU
1	F	499	ILE
1	F	500	LYS
1	F	516	SER
1	F	522	LYS
1	F	535	LEU
1	F	538	VAL
1	F	540	ILE
1	F	547	SER
1	F	548	LYS
1	F	554	ILE
1	F	566	ASN
1	F	567	GLU
1	F	575	ASN
1	F	580	LEU
1	F	595	ILE
1	F	596	ASN
1	F	601	VAL
1	F	612	THR
1	F	613	SER
1	F	614	GLU
1	F	632	ILE
1	F	646	LEU
1	F	649	GLN
1	F	651	LEU
1	F	652	THR
1	F	658	ILE
1	F	659	SER
1	F	661	LEU
1	F	664	HIS

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	G	340	VAL
1	G	341	THR
1	G	343	THR
1	G	345	TYR
1	G	348	PHE
1	G	355	SER
1	G	356	ILE
1	G	357	ILE
1	G	360	VAL
1	G	361	GLN
1	G	364	THR
1	G	367	THR
1	G	374	ILE
1	G	388	GLN
1	G	391	ASP
1	G	396	LEU
1	G	401	LEU
1	G	405	THR
1	G	414	LEU
1	G	416	ILE
1	G	423	THR
1	G	427	ASN
1	G	429	LEU
1	G	441	ILE
1	G	464	MET
1	G	466	THR
1	G	474	SER
1	G	475	VAL
1	G	486	ARG
1	G	489	GLN
1	G	496	GLU
1	G	499	ILE
1	G	500	LYS
1	G	522	LYS
1	G	528	THR
1	G	532	GLU
1	G	533	ASP
1	G	536	TYR
1	G	538	VAL
1	G	547	SER
1	G	580	LEU
1	G	587	ASP

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	G	595	ILE
1	G	612	THR
1	G	613	SER
1	G	619	GLN
1	G	622	GLU
1	G	626	ASP
1	G	628	THR
1	G	634	THR
1	G	635	ARG
1	G	649	GLN
1	G	652	THR
1	G	658	ILE
1	G	659	SER
1	G	660	GLN
1	G	661	LEU
1	G	662	GLU
1	G	663	HIS
1	H	344	ASP
1	H	346	ASP
1	H	356	ILE
1	H	357	ILE
1	H	360	VAL
1	H	361	GLN
1	H	364	THR
1	H	366	SER
1	H	367	THR
1	H	374	ILE
1	H	379	LYS
1	H	380	SER
1	H	386	THR
1	H	387	VAL
1	H	404	ILE
1	H	407	SER
1	H	408	ILE
1	H	409	ILE
1	H	414	LEU
1	H	417	LYS
1	H	419	ASN
1	H	423	THR
1	H	427	ASN
1	H	429	LEU
1	H	436	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	H	446	ARG
1	H	457	SER
1	H	462	SER
1	H	464	MET
1	H	466	THR
1	H	474	SER
1	H	476	ILE
1	H	481	THR
1	H	485	VAL
1	H	486	ARG
1	H	493	LYS
1	H	496	GLU
1	H	499	ILE
1	H	500	LYS
1	H	516	SER
1	H	522	LYS
1	H	535	LEU
1	H	538	VAL
1	H	540	ILE
1	H	547	SER
1	H	548	LYS
1	H	554	ILE
1	H	566	ASN
1	H	567	GLU
1	H	580	LEU
1	H	595	ILE
1	H	596	ASN
1	H	601	VAL
1	H	612	THR
1	H	613	SER
1	H	614	GLU
1	H	632	ILE
1	H	646	LEU
1	H	649	GLN
1	H	651	LEU
1	H	652	THR
1	H	658	ILE
1	H	659	SER
1	H	661	LEU
1	H	664	HIS
1	I	340	VAL
1	I	341	THR

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	I	343	THR
1	I	345	TYR
1	I	348	PHE
1	I	355	SER
1	I	356	ILE
1	I	357	ILE
1	I	360	VAL
1	I	361	GLN
1	I	364	THR
1	I	367	THR
1	I	374	ILE
1	I	388	GLN
1	I	391	ASP
1	I	396	LEU
1	I	401	LEU
1	I	405	THR
1	I	414	LEU
1	I	416	ILE
1	I	423	THR
1	I	427	ASN
1	I	429	LEU
1	I	441	ILE
1	I	464	MET
1	I	466	THR
1	I	474	SER
1	I	475	VAL
1	I	486	ARG
1	I	489	GLN
1	I	496	GLU
1	I	499	ILE
1	I	500	LYS
1	I	522	LYS
1	I	528	THR
1	I	532	GLU
1	I	533	ASP
1	I	536	TYR
1	I	538	VAL
1	I	547	SER
1	I	580	LEU
1	I	587	ASP
1	I	595	ILE
1	I	612	THR

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	I	613	SER
1	I	619	GLN
1	I	622	GLU
1	I	626	ASP
1	I	628	THR
1	I	634	THR
1	I	635	ARG
1	I	649	GLN
1	I	652	THR
1	I	658	ILE
1	I	659	SER
1	I	660	GLN
1	I	661	LEU
1	I	662	GLU
1	I	663	HIS
1	J	344	ASP
1	J	346	ASP
1	J	356	ILE
1	J	357	ILE
1	J	360	VAL
1	J	361	GLN
1	J	364	THR
1	J	366	SER
1	J	367	THR
1	J	374	ILE
1	J	379	LYS
1	J	380	SER
1	J	386	THR
1	J	387	VAL
1	J	404	ILE
1	J	407	SER
1	J	408	ILE
1	J	409	ILE
1	J	414	LEU
1	J	417	LYS
1	J	419	ASN
1	J	423	THR
1	J	427	ASN
1	J	429	LEU
1	J	436	LEU
1	J	446	ARG
1	J	457	SER

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	J	462	SER
1	J	464	MET
1	J	466	THR
1	J	474	SER
1	J	476	ILE
1	J	481	THR
1	J	485	VAL
1	J	486	ARG
1	J	493	LYS
1	J	496	GLU
1	J	499	ILE
1	J	500	LYS
1	J	516	SER
1	J	522	LYS
1	J	535	LEU
1	J	538	VAL
1	J	540	ILE
1	J	547	SER
1	J	548	LYS
1	J	554	ILE
1	J	566	ASN
1	J	567	GLU
1	J	580	LEU
1	J	595	ILE
1	J	596	ASN
1	J	601	VAL
1	J	612	THR
1	J	613	SER
1	J	614	GLU
1	J	632	ILE
1	J	646	LEU
1	J	649	GLN
1	J	651	LEU
1	J	652	THR
1	J	658	ILE
1	J	659	SER
1	J	661	LEU
1	J	664	HIS
1	K	340	VAL
1	K	341	THR
1	K	343	THR
1	K	345	TYR

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	K	348	PHE
1	K	355	SER
1	K	356	ILE
1	K	357	ILE
1	K	360	VAL
1	K	361	GLN
1	K	364	THR
1	K	367	THR
1	K	374	ILE
1	K	388	GLN
1	K	391	ASP
1	K	396	LEU
1	K	401	LEU
1	K	405	THR
1	K	414	LEU
1	K	416	ILE
1	K	423	THR
1	K	427	ASN
1	K	429	LEU
1	K	441	ILE
1	K	464	MET
1	K	466	THR
1	K	474	SER
1	K	475	VAL
1	K	486	ARG
1	K	489	GLN
1	K	496	GLU
1	K	499	ILE
1	K	500	LYS
1	K	522	LYS
1	K	528	THR
1	K	532	GLU
1	K	533	ASP
1	K	536	TYR
1	K	538	VAL
1	K	547	SER
1	K	580	LEU
1	K	587	ASP
1	K	595	ILE
1	K	612	THR
1	K	613	SER
1	K	619	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	K	622	GLU
1	K	626	ASP
1	K	628	THR
1	K	634	THR
1	K	635	ARG
1	K	649	GLN
1	K	652	THR
1	K	658	ILE
1	K	659	SER
1	K	660	GLN
1	K	661	LEU
1	K	662	GLU
1	K	663	HIS
1	L	344	ASP
1	L	346	ASP
1	L	356	ILE
1	L	357	ILE
1	L	360	VAL
1	L	361	GLN
1	L	364	THR
1	L	366	SER
1	L	367	THR
1	L	374	ILE
1	L	379	LYS
1	L	380	SER
1	L	386	THR
1	L	387	VAL
1	L	404	ILE
1	L	407	SER
1	L	408	ILE
1	L	409	ILE
1	L	414	LEU
1	L	417	LYS
1	L	419	ASN
1	L	423	THR
1	L	427	ASN
1	L	429	LEU
1	L	436	LEU
1	L	446	ARG
1	L	457	SER
1	L	462	SER
1	L	464	MET

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	L	466	THR
1	L	474	SER
1	L	476	ILE
1	L	481	THR
1	L	485	VAL
1	L	486	ARG
1	L	493	LYS
1	L	496	GLU
1	L	499	ILE
1	L	500	LYS
1	L	516	SER
1	L	522	LYS
1	L	535	LEU
1	L	538	VAL
1	L	540	ILE
1	L	547	SER
1	L	548	LYS
1	L	554	ILE
1	L	566	ASN
1	L	567	GLU
1	L	575	ASN
1	L	580	LEU
1	L	595	ILE
1	L	596	ASN
1	L	601	VAL
1	L	612	THR
1	L	613	SER
1	L	614	GLU
1	L	632	ILE
1	L	646	LEU
1	L	649	GLN
1	L	651	LEU
1	L	652	THR
1	L	658	ILE
1	L	659	SER
1	L	661	LEU
1	L	664	HIS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (154) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	361	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	A	427	ASN
1	A	456	ASN
1	A	473	HIS
1	A	490	ASN
1	A	504	GLN
1	A	596	ASN
1	A	605	ASN
1	A	610	ASN
1	A	619	GLN
1	A	644	ASN
1	B	358	GLN
1	B	361	GLN
1	B	394	ASN
1	B	412	ASN
1	B	427	ASN
1	B	489	GLN
1	B	490	ASN
1	B	503	ASN
1	B	504	GLN
1	B	570	GLN
1	B	582	ASN
1	B	596	ASN
1	B	644	ASN
1	B	663	HIS
1	C	361	GLN
1	C	427	ASN
1	C	456	ASN
1	C	473	HIS
1	C	490	ASN
1	C	504	GLN
1	C	596	ASN
1	C	605	ASN
1	C	610	ASN
1	C	619	GLN
1	C	644	ASN
1	D	358	GLN
1	D	361	GLN
1	D	394	ASN
1	D	400	ASN
1	D	412	ASN
1	D	427	ASN
1	D	489	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	490	ASN
1	D	502	ASN
1	D	503	ASN
1	D	504	GLN
1	D	570	GLN
1	D	582	ASN
1	D	596	ASN
1	D	644	ASN
1	D	663	HIS
1	E	361	GLN
1	E	427	ASN
1	E	456	ASN
1	E	473	HIS
1	E	490	ASN
1	E	504	GLN
1	E	596	ASN
1	E	605	ASN
1	E	610	ASN
1	E	619	GLN
1	E	644	ASN
1	E	663	HIS
1	F	358	GLN
1	F	361	GLN
1	F	394	ASN
1	F	412	ASN
1	F	427	ASN
1	F	489	GLN
1	F	490	ASN
1	F	502	ASN
1	F	503	ASN
1	F	504	GLN
1	F	570	GLN
1	F	582	ASN
1	F	596	ASN
1	F	644	ASN
1	F	663	HIS
1	G	361	GLN
1	G	427	ASN
1	G	456	ASN
1	G	473	HIS
1	G	490	ASN
1	G	504	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	G	596	ASN
1	G	605	ASN
1	G	610	ASN
1	G	619	GLN
1	G	644	ASN
1	H	358	GLN
1	H	361	GLN
1	H	394	ASN
1	H	412	ASN
1	H	427	ASN
1	H	489	GLN
1	H	490	ASN
1	H	502	ASN
1	H	503	ASN
1	H	570	GLN
1	H	582	ASN
1	H	596	ASN
1	H	644	ASN
1	H	663	HIS
1	I	361	GLN
1	I	427	ASN
1	I	456	ASN
1	I	473	HIS
1	I	490	ASN
1	I	504	GLN
1	I	605	ASN
1	I	610	ASN
1	I	619	GLN
1	I	644	ASN
1	J	358	GLN
1	J	361	GLN
1	J	394	ASN
1	J	412	ASN
1	J	427	ASN
1	J	489	GLN
1	J	490	ASN
1	J	502	ASN
1	J	503	ASN
1	J	504	GLN
1	J	570	GLN
1	J	582	ASN
1	J	596	ASN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	J	644	ASN
1	J	663	HIS
1	K	361	GLN
1	K	427	ASN
1	K	456	ASN
1	K	473	HIS
1	K	490	ASN
1	K	504	GLN
1	K	596	ASN
1	K	605	ASN
1	K	610	ASN
1	K	619	GLN
1	K	644	ASN
1	L	358	GLN
1	L	361	GLN
1	L	394	ASN
1	L	412	ASN
1	L	427	ASN
1	L	489	GLN
1	L	490	ASN
1	L	503	ASN
1	L	504	GLN
1	L	570	GLN
1	L	582	ASN
1	L	596	ASN
1	L	644	ASN
1	L	663	HIS

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 [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

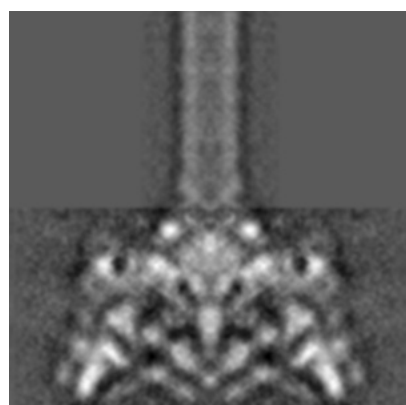
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-1048. These allow visual inspection of the internal detail of the map and identification of artifacts.

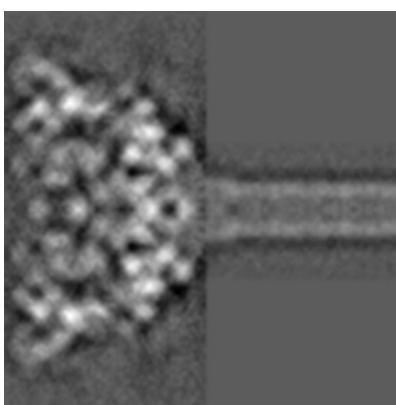
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

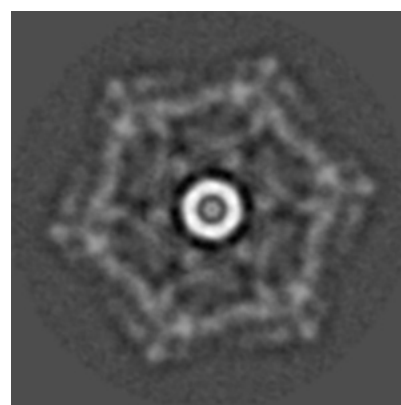
6.1.1 Primary map



X



Y

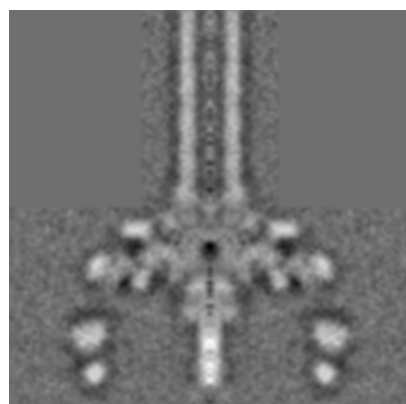


Z

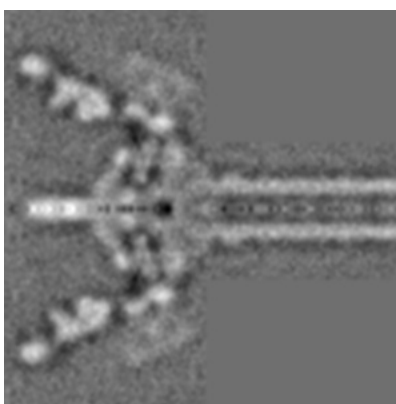
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

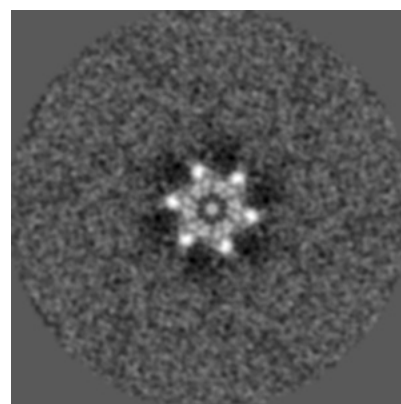
6.2.1 Primary map



X Index: 98



Y Index: 98

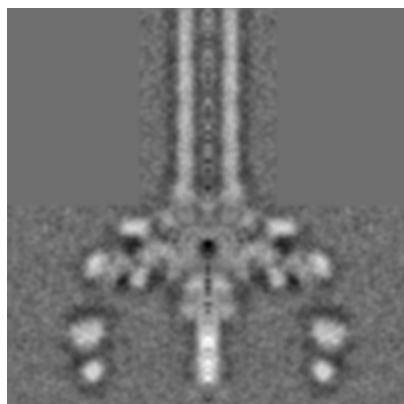


Z Index: 98

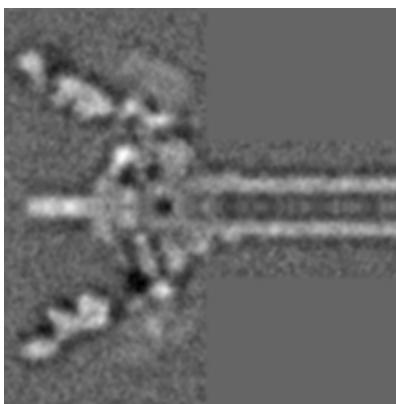
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

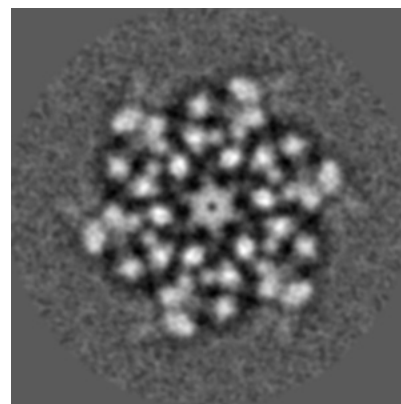
6.3.1 Primary map



X Index: 98



Y Index: 100

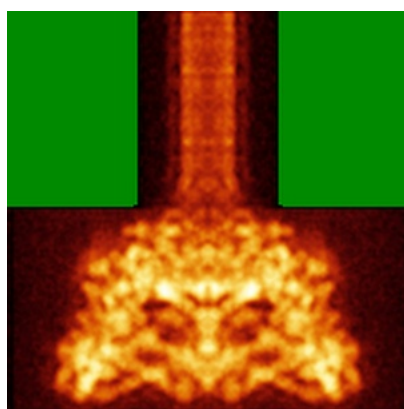


Z Index: 60

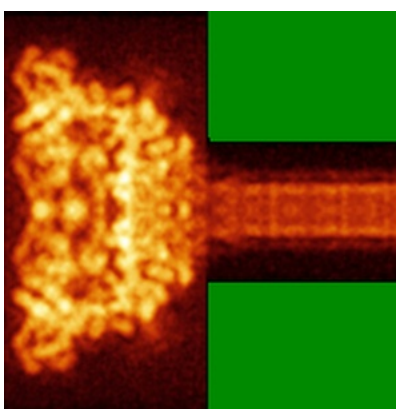
The images above show the largest variance slices of the map in three orthogonal directions.

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

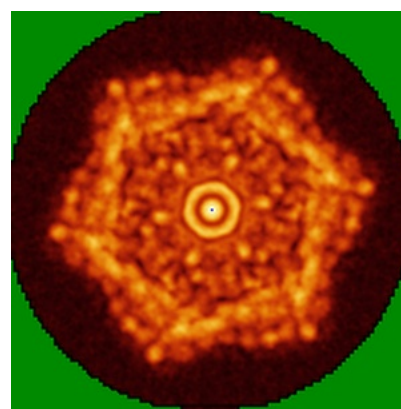
6.4.1 Primary map



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 1.45. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

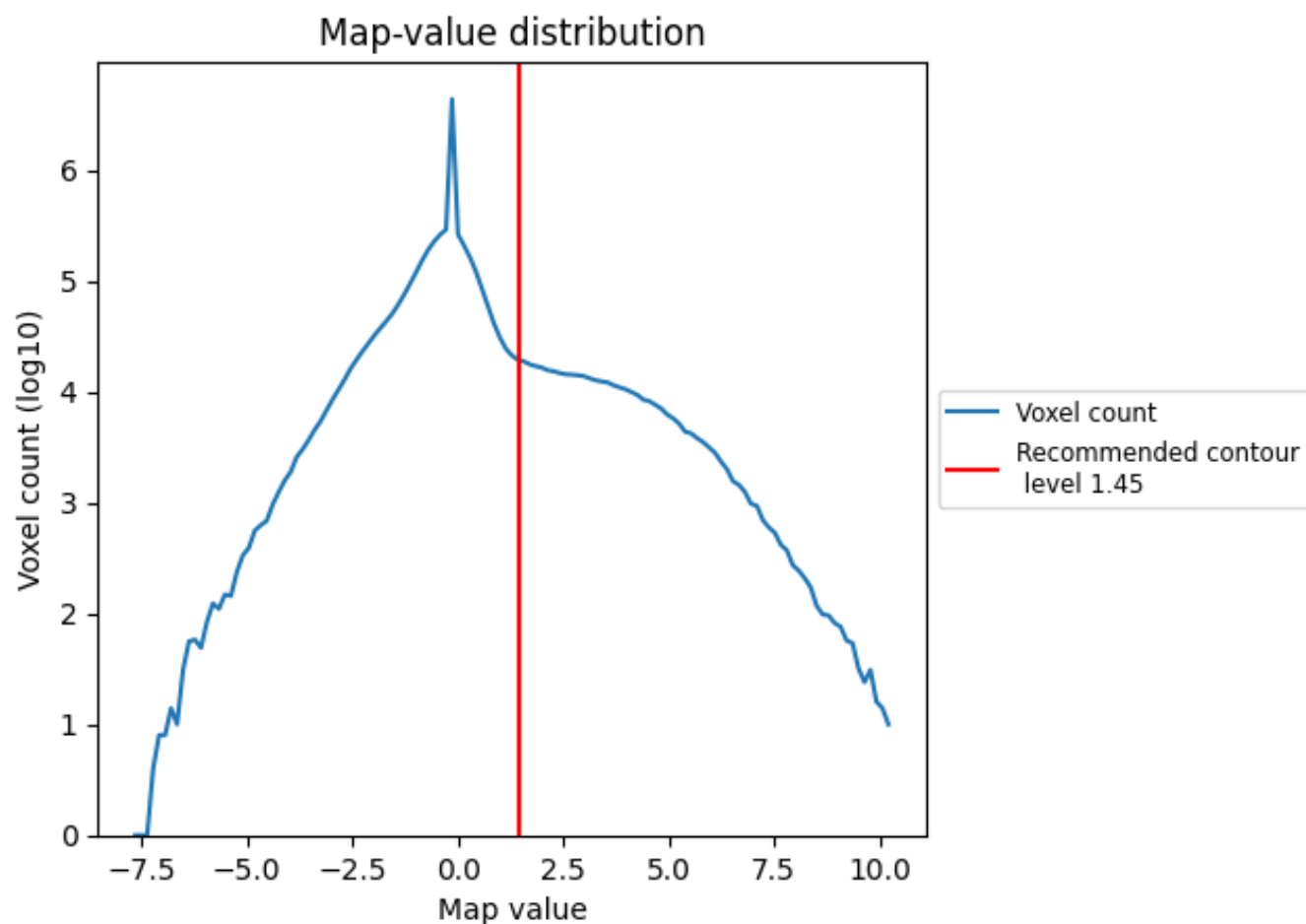
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

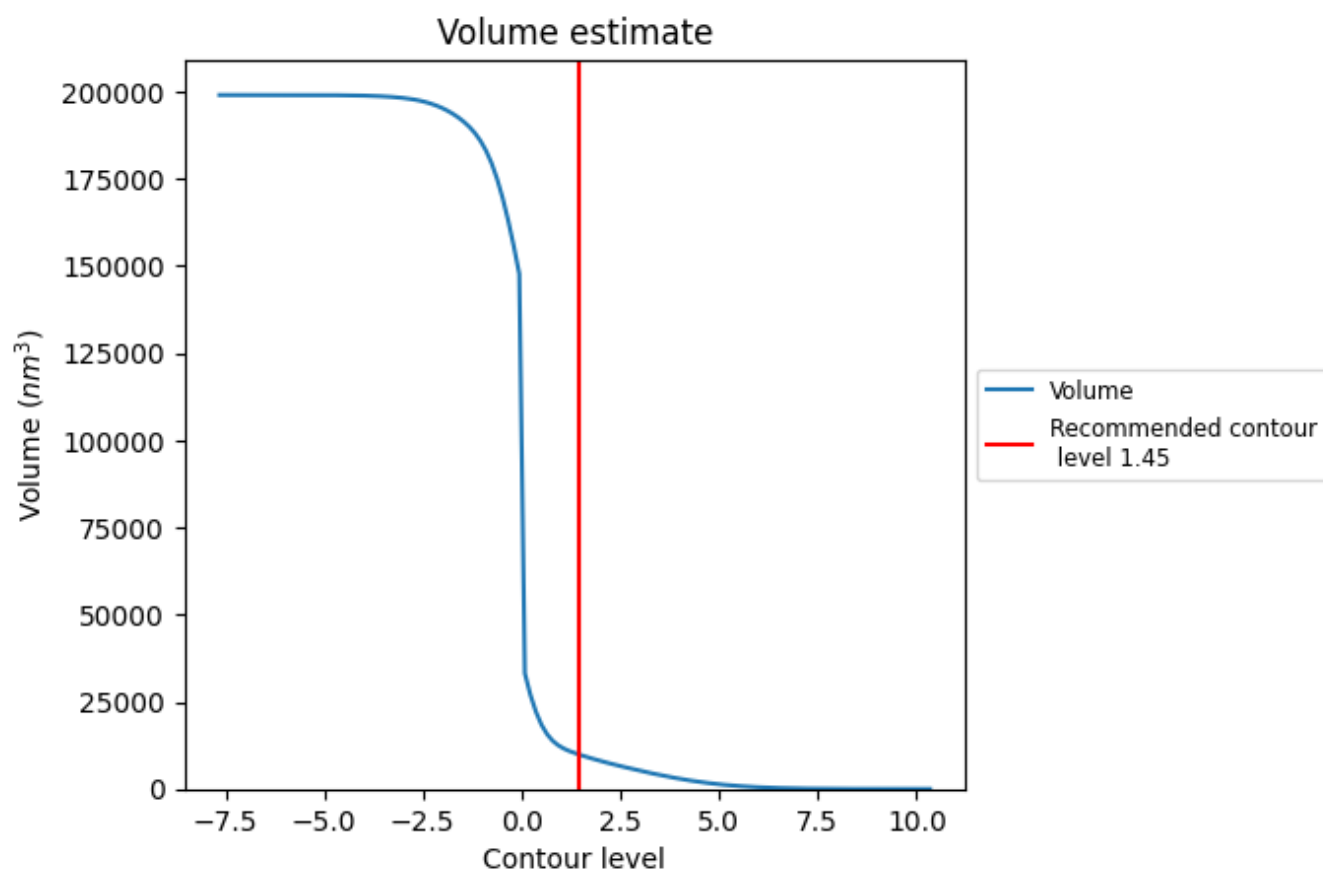
This section contains the results of statistical analysis of the map.

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.

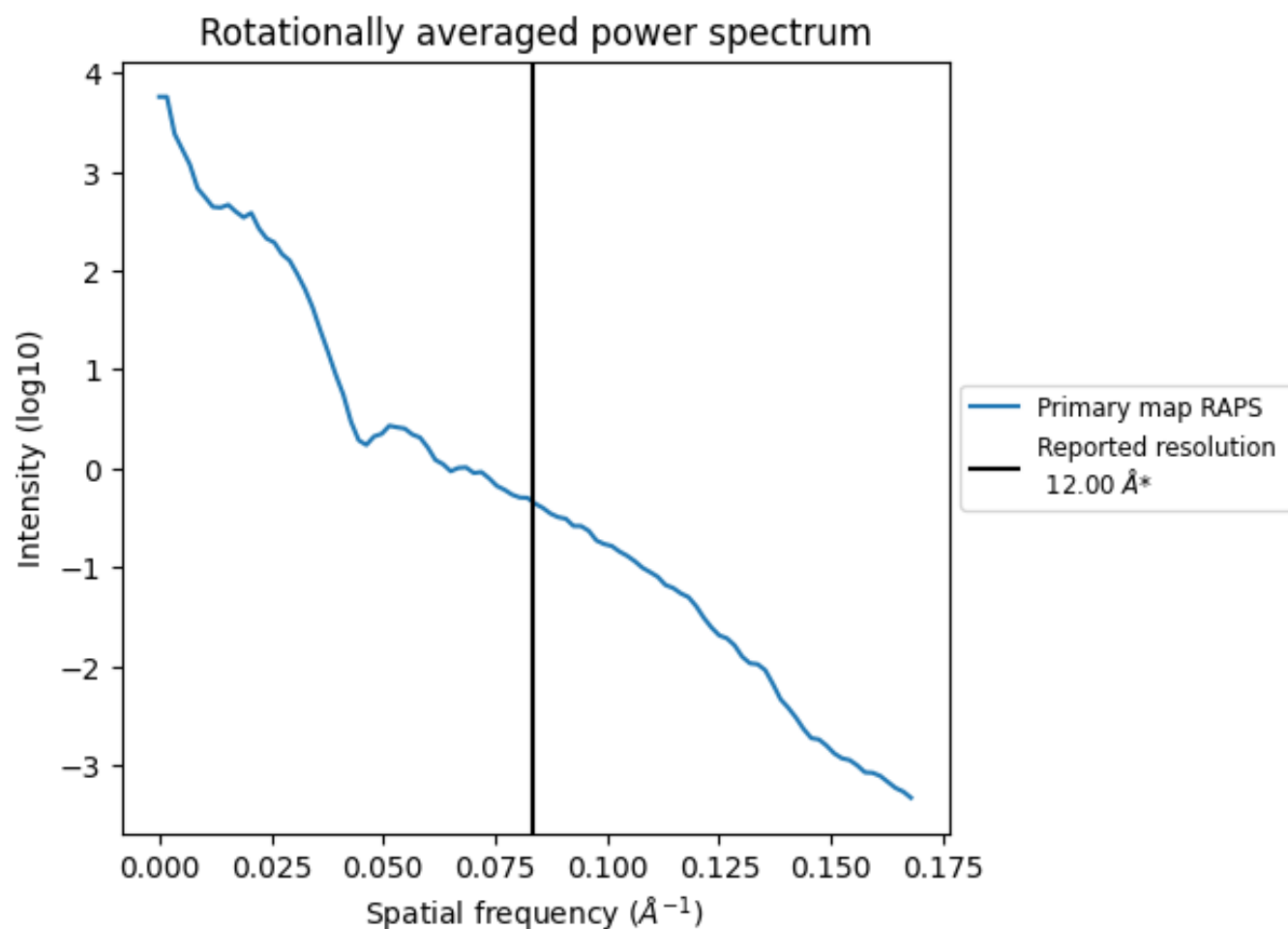
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 9896 nm³; this corresponds to an approximate mass of 8939 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 0.083 Å⁻¹

8 Fourier-Shell correlation

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

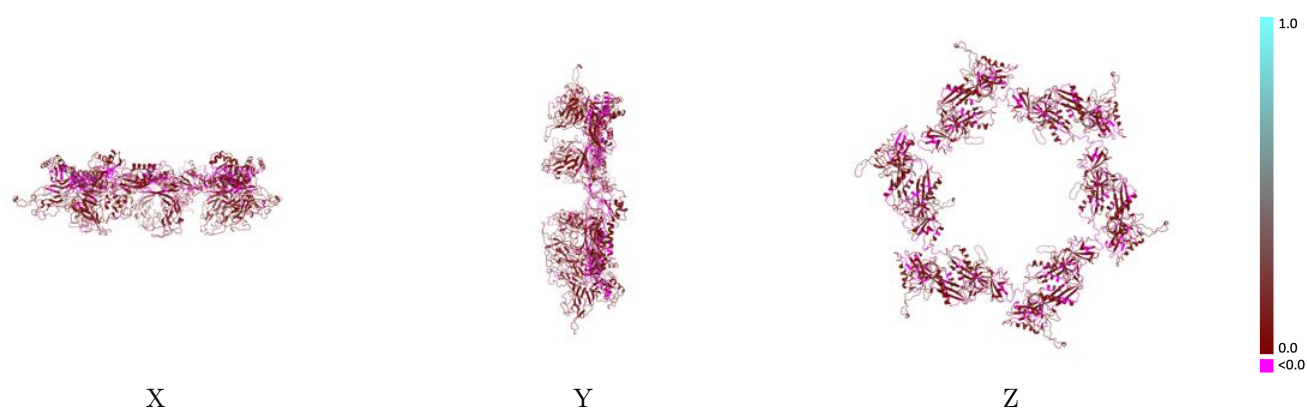
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-1048 and PDB model 3H3W. Per-residue inclusion information can be found in section 3 on page 8.

9.1 Map-model overlay [i](#)

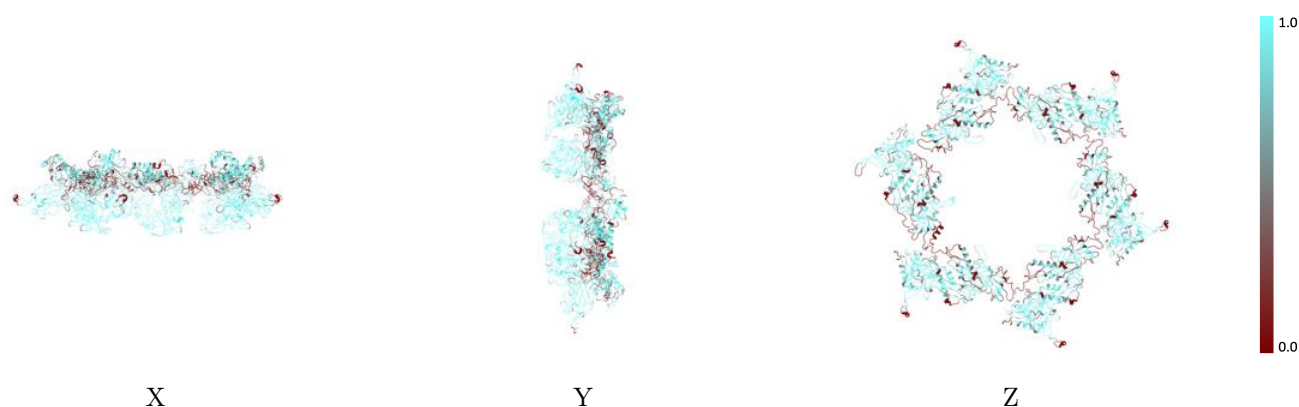
This section was not generated.

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



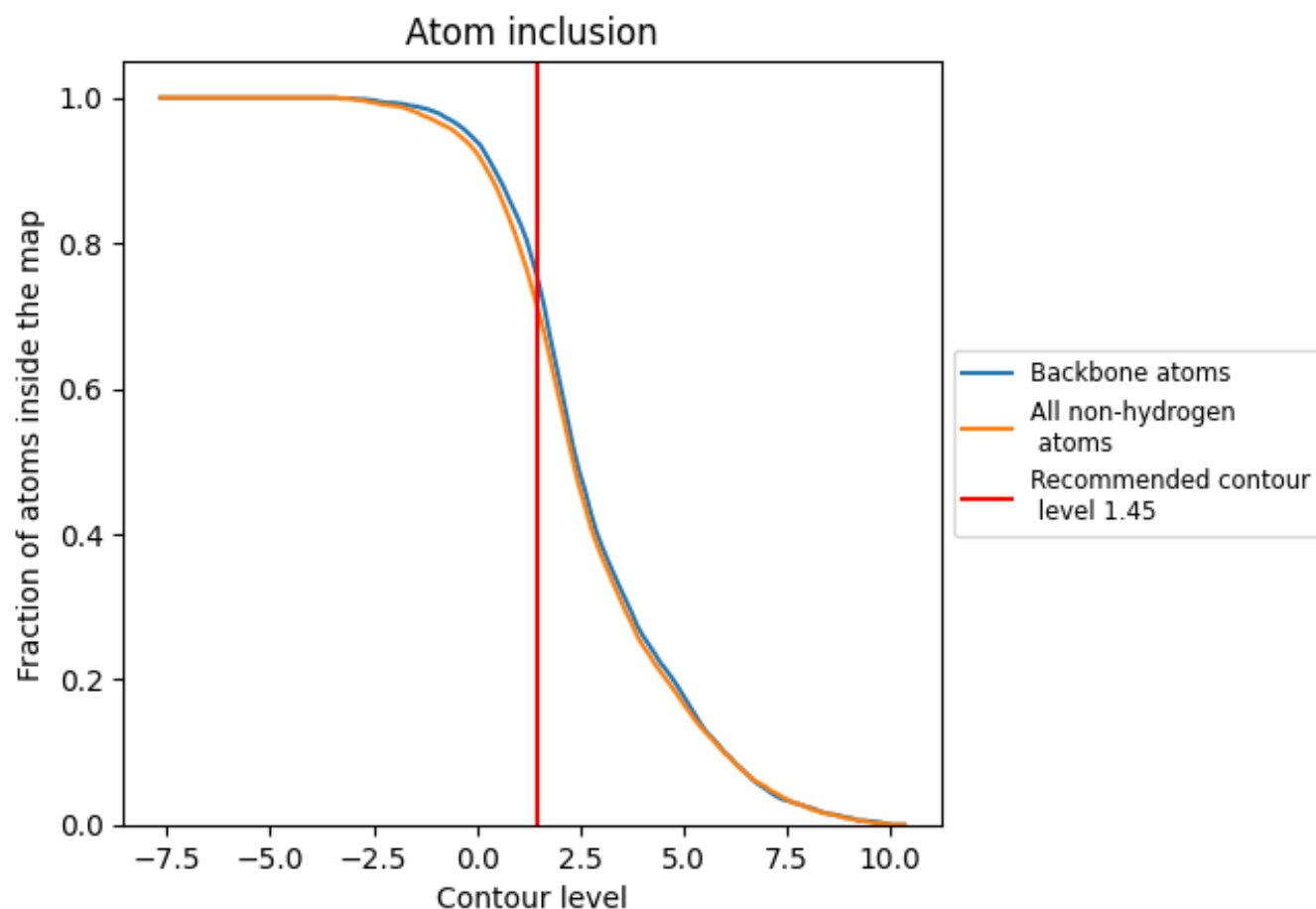
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (1.45).

9.4 Atom inclusion [i](#)



At the recommended contour level, 76% of all backbone atoms, 72% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (1.45) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.7150	0.0680
A	0.6560	0.0630
B	0.7730	0.0750
C	0.7700	0.0760
D	0.6550	0.0630
E	0.6540	0.0630
F	0.7800	0.0730
G	0.6550	0.0600
H	0.7730	0.0740
I	0.6540	0.0630
J	0.7740	0.0720
K	0.6540	0.0640
L	0.7800	0.0750

