



Full wwPDB EM Validation Report ⓘ

Apr 29, 2026 – 06:17 PM JST

PDB ID : 9VEF / pdb_00009vef
EMDB ID : EMD-65006
Title : The cryo-EM structure of human Piezo2-MDFIC complex (composite map)
Authors : Zhang, Y.; Dai, F.; Zhou, Z.; Dai, F.; Cheng, D.; Ma, X.; Omidkhoda, S.F.;
Clarke, J.; Zhang, H.; Laden, M.; Guo, Y.; Li, J.V.; Liu, R.; Wong, E.S.;
Zhang, Y.; Cox, C.D.
Deposited on : 2025-06-09
Resolution : 3.75 Å(reported)

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

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : **FAILED**
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : **NOT EXECUTED**
MapQ : **FAILED**
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

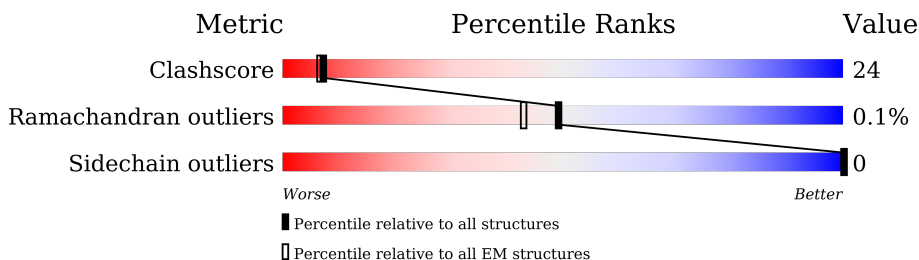
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.75 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	229148	23984
Ramachandran outliers	224038	23583
Sidechain outliers	223484	23102

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 $\leq 5\%$.

Mol	Chain	Length	Quality of chain
1	A	3020	26% 20% 54%
1	C	3020	26% 20% 54%
1	E	3020	26% 20% 54%
2	B	267	6% . 92%
2	D	267	6% . 92%
2	F	267	6% . 92%

2 Entry composition

There are 2 unique types of molecules in this entry. The entry contains 34857 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 Piezo-type mechanosensitive ion channel component 2, Green fluorescent protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	1396	Total	C	N	O	S	0	0
			11465	7568	1851	1969	77		
1	C	1396	Total	C	N	O	S	0	0
			11465	7568	1851	1969	77		
1	E	1396	Total	C	N	O	S	0	0
			11465	7568	1851	1969	77		

There are 105 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	2753	SER	-	linker	UNP Q9H5I5
A	2754	ASN	-	linker	UNP Q9H5I5
A	2755	SER	-	linker	UNP Q9H5I5
A	2756	LEU	-	linker	UNP Q9H5I5
A	2757	GLU	-	linker	UNP Q9H5I5
A	2758	VAL	-	linker	UNP Q9H5I5
A	2759	LEU	-	linker	UNP Q9H5I5
A	2760	PHE	-	linker	UNP Q9H5I5
A	2761	GLN	-	linker	UNP Q9H5I5
A	2762	GLY	-	linker	UNP Q9H5I5
A	2763	PRO	-	linker	UNP Q9H5I5
A	2764	THR	-	linker	UNP Q9H5I5
A	2765	ALA	-	linker	UNP Q9H5I5
A	2766	ALA	-	linker	UNP Q9H5I5
A	2767	ALA	-	linker	UNP Q9H5I5
A	2768	ALA	-	linker	UNP Q9H5I5
A	2769	VAL	-	linker	UNP Q9H5I5
A	2832	LEU	PHE	conflict	UNP P42212
A	2833	THR	SER	conflict	UNP P42212
A	2974	LYS	ALA	conflict	UNP P42212
A	2999	LEU	HIS	conflict	UNP P42212
A	3007	SER	-	expression tag	UNP P42212
A	3008	GLY	-	expression tag	UNP P42212

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Chain	Residue	Modelled	Actual	Comment	Reference
A	3009	GLY	-	expression tag	UNP P42212
A	3010	GLY	-	expression tag	UNP P42212
A	3011	HIS	-	expression tag	UNP P42212
A	3012	HIS	-	expression tag	UNP P42212
A	3013	HIS	-	expression tag	UNP P42212
A	3014	HIS	-	expression tag	UNP P42212
A	3015	HIS	-	expression tag	UNP P42212
A	3016	HIS	-	expression tag	UNP P42212
A	3017	HIS	-	expression tag	UNP P42212
A	3018	HIS	-	expression tag	UNP P42212
A	3019	HIS	-	expression tag	UNP P42212
A	3020	HIS	-	expression tag	UNP P42212
C	2753	SER	-	linker	UNP Q9H5I5
C	2754	ASN	-	linker	UNP Q9H5I5
C	2755	SER	-	linker	UNP Q9H5I5
C	2756	LEU	-	linker	UNP Q9H5I5
C	2757	GLU	-	linker	UNP Q9H5I5
C	2758	VAL	-	linker	UNP Q9H5I5
C	2759	LEU	-	linker	UNP Q9H5I5
C	2760	PHE	-	linker	UNP Q9H5I5
C	2761	GLN	-	linker	UNP Q9H5I5
C	2762	GLY	-	linker	UNP Q9H5I5
C	2763	PRO	-	linker	UNP Q9H5I5
C	2764	THR	-	linker	UNP Q9H5I5
C	2765	ALA	-	linker	UNP Q9H5I5
C	2766	ALA	-	linker	UNP Q9H5I5
C	2767	ALA	-	linker	UNP Q9H5I5
C	2768	ALA	-	linker	UNP Q9H5I5
C	2769	VAL	-	linker	UNP Q9H5I5
C	2832	LEU	PHE	conflict	UNP P42212
C	2833	THR	SER	conflict	UNP P42212
C	2974	LYS	ALA	conflict	UNP P42212
C	2999	LEU	HIS	conflict	UNP P42212
C	3007	SER	-	expression tag	UNP P42212
C	3008	GLY	-	expression tag	UNP P42212
C	3009	GLY	-	expression tag	UNP P42212
C	3010	GLY	-	expression tag	UNP P42212
C	3011	HIS	-	expression tag	UNP P42212
C	3012	HIS	-	expression tag	UNP P42212
C	3013	HIS	-	expression tag	UNP P42212
C	3014	HIS	-	expression tag	UNP P42212
C	3015	HIS	-	expression tag	UNP P42212

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Chain	Residue	Modelled	Actual	Comment	Reference
C	3016	HIS	-	expression tag	UNP P42212
C	3017	HIS	-	expression tag	UNP P42212
C	3018	HIS	-	expression tag	UNP P42212
C	3019	HIS	-	expression tag	UNP P42212
C	3020	HIS	-	expression tag	UNP P42212
E	2753	SER	-	linker	UNP Q9H5I5
E	2754	ASN	-	linker	UNP Q9H5I5
E	2755	SER	-	linker	UNP Q9H5I5
E	2756	LEU	-	linker	UNP Q9H5I5
E	2757	GLU	-	linker	UNP Q9H5I5
E	2758	VAL	-	linker	UNP Q9H5I5
E	2759	LEU	-	linker	UNP Q9H5I5
E	2760	PHE	-	linker	UNP Q9H5I5
E	2761	GLN	-	linker	UNP Q9H5I5
E	2762	GLY	-	linker	UNP Q9H5I5
E	2763	PRO	-	linker	UNP Q9H5I5
E	2764	THR	-	linker	UNP Q9H5I5
E	2765	ALA	-	linker	UNP Q9H5I5
E	2766	ALA	-	linker	UNP Q9H5I5
E	2767	ALA	-	linker	UNP Q9H5I5
E	2768	ALA	-	linker	UNP Q9H5I5
E	2769	VAL	-	linker	UNP Q9H5I5
E	2832	LEU	PHE	conflict	UNP P42212
E	2833	THR	SER	conflict	UNP P42212
E	2974	LYS	ALA	conflict	UNP P42212
E	2999	LEU	HIS	conflict	UNP P42212
E	3007	SER	-	expression tag	UNP P42212
E	3008	GLY	-	expression tag	UNP P42212
E	3009	GLY	-	expression tag	UNP P42212
E	3010	GLY	-	expression tag	UNP P42212
E	3011	HIS	-	expression tag	UNP P42212
E	3012	HIS	-	expression tag	UNP P42212
E	3013	HIS	-	expression tag	UNP P42212
E	3014	HIS	-	expression tag	UNP P42212
E	3015	HIS	-	expression tag	UNP P42212
E	3016	HIS	-	expression tag	UNP P42212
E	3017	HIS	-	expression tag	UNP P42212
E	3018	HIS	-	expression tag	UNP P42212
E	3019	HIS	-	expression tag	UNP P42212
E	3020	HIS	-	expression tag	UNP P42212

- Molecule 2 is a protein called MyoD family inhibitor domain-containing protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	B	22	Total 154	C 91	N 22	O 33	S 8	0	0
2	D	22	Total 154	C 91	N 22	O 33	S 8	0	0
2	F	22	Total 154	C 91	N 22	O 33	S 8	0	0

There are 63 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
B	-20	MET	-	initiating methionine	UNP Q9P1T7
B	-19	ASP	-	expression tag	UNP Q9P1T7
B	-18	TYR	-	expression tag	UNP Q9P1T7
B	-17	LYS	-	expression tag	UNP Q9P1T7
B	-16	ASP	-	expression tag	UNP Q9P1T7
B	-15	ASP	-	expression tag	UNP Q9P1T7
B	-14	ASP	-	expression tag	UNP Q9P1T7
B	-13	ASP	-	expression tag	UNP Q9P1T7
B	-12	LYS	-	expression tag	UNP Q9P1T7
B	-11	GLY	-	expression tag	UNP Q9P1T7
B	-10	LEU	-	expression tag	UNP Q9P1T7
B	-9	GLU	-	expression tag	UNP Q9P1T7
B	-8	VAL	-	expression tag	UNP Q9P1T7
B	-7	LEU	-	expression tag	UNP Q9P1T7
B	-6	PHE	-	expression tag	UNP Q9P1T7
B	-5	GLN	-	expression tag	UNP Q9P1T7
B	-4	GLY	-	expression tag	UNP Q9P1T7
B	-3	PRO	-	expression tag	UNP Q9P1T7
B	-2	GLY	-	expression tag	UNP Q9P1T7
B	-1	SER	-	expression tag	UNP Q9P1T7
B	0	SER	-	expression tag	UNP Q9P1T7
D	-20	MET	-	initiating methionine	UNP Q9P1T7
D	-19	ASP	-	expression tag	UNP Q9P1T7
D	-18	TYR	-	expression tag	UNP Q9P1T7
D	-17	LYS	-	expression tag	UNP Q9P1T7
D	-16	ASP	-	expression tag	UNP Q9P1T7
D	-15	ASP	-	expression tag	UNP Q9P1T7
D	-14	ASP	-	expression tag	UNP Q9P1T7
D	-13	ASP	-	expression tag	UNP Q9P1T7
D	-12	LYS	-	expression tag	UNP Q9P1T7
D	-11	GLY	-	expression tag	UNP Q9P1T7
D	-10	LEU	-	expression tag	UNP Q9P1T7
D	-9	GLU	-	expression tag	UNP Q9P1T7

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Chain	Residue	Modelled	Actual	Comment	Reference
D	-8	VAL	-	expression tag	UNP Q9P1T7
D	-7	LEU	-	expression tag	UNP Q9P1T7
D	-6	PHE	-	expression tag	UNP Q9P1T7
D	-5	GLN	-	expression tag	UNP Q9P1T7
D	-4	GLY	-	expression tag	UNP Q9P1T7
D	-3	PRO	-	expression tag	UNP Q9P1T7
D	-2	GLY	-	expression tag	UNP Q9P1T7
D	-1	SER	-	expression tag	UNP Q9P1T7
D	0	SER	-	expression tag	UNP Q9P1T7
F	-20	MET	-	initiating methionine	UNP Q9P1T7
F	-19	ASP	-	expression tag	UNP Q9P1T7
F	-18	TYR	-	expression tag	UNP Q9P1T7
F	-17	LYS	-	expression tag	UNP Q9P1T7
F	-16	ASP	-	expression tag	UNP Q9P1T7
F	-15	ASP	-	expression tag	UNP Q9P1T7
F	-14	ASP	-	expression tag	UNP Q9P1T7
F	-13	ASP	-	expression tag	UNP Q9P1T7
F	-12	LYS	-	expression tag	UNP Q9P1T7
F	-11	GLY	-	expression tag	UNP Q9P1T7
F	-10	LEU	-	expression tag	UNP Q9P1T7
F	-9	GLU	-	expression tag	UNP Q9P1T7
F	-8	VAL	-	expression tag	UNP Q9P1T7
F	-7	LEU	-	expression tag	UNP Q9P1T7
F	-6	PHE	-	expression tag	UNP Q9P1T7
F	-5	GLN	-	expression tag	UNP Q9P1T7
F	-4	GLY	-	expression tag	UNP Q9P1T7
F	-3	PRO	-	expression tag	UNP Q9P1T7
F	-2	GLY	-	expression tag	UNP Q9P1T7
F	-1	SER	-	expression tag	UNP Q9P1T7
F	0	SER	-	expression tag	UNP Q9P1T7

WORLDWIDE
PDB
PROTEIN DATA BANK



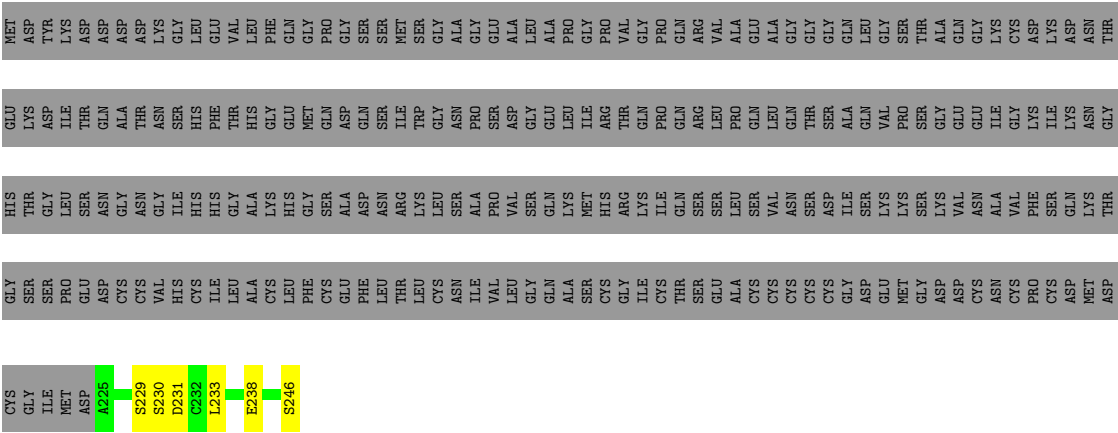






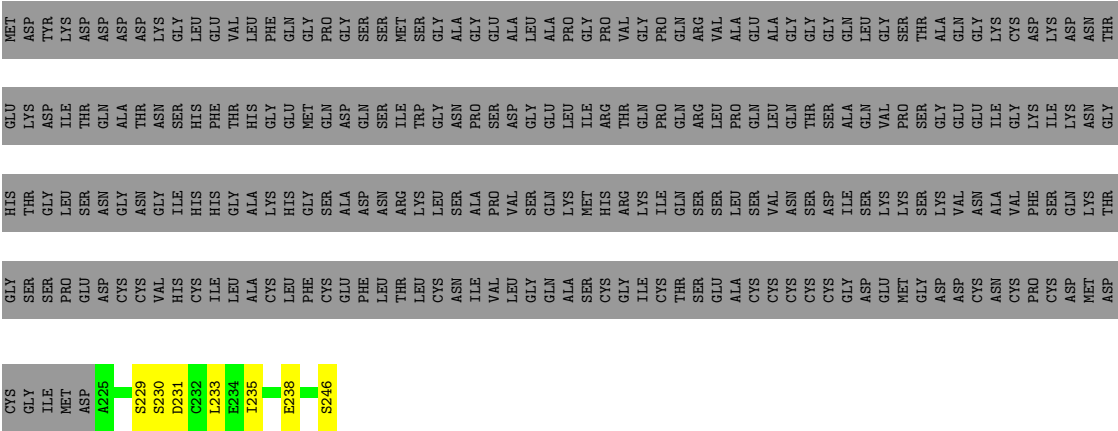






● Molecule 2: MyoD family inhibitor domain-containing protein

Chain F: 6% . 92%



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	83949	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	49.41	Depositor
Minimum defocus (nm)	1400	Depositor
Maximum defocus (nm)	2400	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	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	0.45	0/11775	0.55	0/15965
1	C	0.45	0/11775	0.55	0/15965
1	E	0.45	0/11775	0.55	0/15965
2	B	0.30	0/155	0.55	0/207
2	D	0.31	0/155	0.55	0/207
2	F	0.30	0/155	0.55	0/207
All	All	0.45	0/35790	0.55	0/48516

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	3
1	C	0	3
1	E	0	3
All	All	0	9

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (9) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	2477	SER	Peptide
1	A	2624	THR	Peptide
1	A	2653	PHE	Peptide
1	C	2477	SER	Peptide
1	C	2624	THR	Peptide
1	C	2653	PHE	Peptide

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Mol	Chain	Res	Type	Group
1	E	2477	SER	Peptide
1	E	2624	THR	Peptide
1	E	2653	PHE	Peptide

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	11465	0	11592	577	0
1	C	11465	0	11592	570	0
1	E	11465	0	11592	577	0
2	B	154	0	137	6	0
2	D	154	0	137	7	0
2	F	154	0	137	7	0
All	All	34857	0	35187	1648	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 24.

All (1648) 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:E:2430:ALA:N	1:E:2658:SER:HG	1.44	1.15
1:C:2430:ALA:N	1:C:2658:SER:HG	1.43	1.15
1:A:2430:ALA:N	1:A:2658:SER:HG	1.45	1.14
1:E:2468:SER:HB2	1:E:2472:PHE:HB3	1.46	0.95
1:A:2468:SER:HB2	1:A:2472:PHE:HB3	1.46	0.94
1:C:2468:SER:HB2	1:C:2472:PHE:HB3	1.46	0.93
1:A:2392:ARG:HE	1:E:1542:HIS:HA	1.34	0.91
1:A:1542:HIS:HA	1:C:2392:ARG:HE	1.34	0.91
1:C:1542:HIS:HA	1:E:2392:ARG:HE	1.34	0.91
1:C:1950:PHE:HZ	1:C:2300:CYS:HG	1.19	0.88
1:E:2457:GLN:HB2	1:E:2460:GLN:HB2	1.56	0.88
1:C:2457:GLN:HB2	1:C:2460:GLN:HB2	1.56	0.88
1:A:2413:ILE:O	1:A:2417:TRP:HB3	1.74	0.88
1:E:2413:ILE:O	1:E:2417:TRP:HB3	1.74	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1449:SER:OG	1:E:1700:ARG:NH1	2.08	0.86
1:A:1449:SER:OG	1:A:1700:ARG:NH1	2.08	0.86
1:C:1449:SER:OG	1:C:1700:ARG:NH1	2.08	0.86
1:A:2457:GLN:HB2	1:A:2460:GLN:HB2	1.56	0.86
1:C:2413:ILE:O	1:C:2417:TRP:HB3	1.74	0.85
1:E:1950:PHE:HZ	1:E:2300:CYS:HG	1.25	0.85
1:C:2698:PHE:HD2	1:E:2694:HIS:HD1	1.19	0.85
1:A:2694:HIS:HD1	1:E:2698:PHE:HD2	1.21	0.84
1:C:2471:LYS:O	1:C:2475:ALA:N	2.11	0.84
1:A:2471:LYS:O	1:A:2475:ALA:N	2.11	0.84
1:A:2698:PHE:HD2	1:C:2694:HIS:HD1	1.21	0.84
1:E:2471:LYS:O	1:E:2475:ALA:N	2.11	0.83
1:A:1950:PHE:HZ	1:A:2300:CYS:HG	1.25	0.82
1:E:2486:LEU:O	1:E:2488:ASN:ND2	2.13	0.82
1:A:2486:LEU:O	1:A:2488:ASN:ND2	2.13	0.81
1:A:2469:PHE:HA	1:A:2473:ILE:HD12	1.63	0.81
1:C:2486:LEU:O	1:C:2488:ASN:ND2	2.13	0.81
1:E:2469:PHE:HA	1:E:2473:ILE:HD12	1.63	0.81
1:C:2323:LEU:O	1:C:2331:ASN:ND2	2.15	0.80
1:E:1374:LEU:HG	1:E:1378:SER:HB2	1.63	0.80
1:E:2482:ALA:HB1	1:E:2604:GLN:HB3	1.63	0.80
1:C:2469:PHE:HA	1:C:2473:ILE:HD12	1.62	0.80
1:A:2482:ALA:HB1	1:A:2604:GLN:HB3	1.63	0.80
1:A:2600:LYS:NZ	1:A:2601:PRO:O	2.14	0.80
1:C:1374:LEU:HG	1:C:1378:SER:HB2	1.63	0.80
1:C:2482:ALA:HB1	1:C:2604:GLN:HB3	1.63	0.80
1:A:1374:LEU:HG	1:A:1378:SER:HB2	1.63	0.80
1:E:2323:LEU:O	1:E:2331:ASN:ND2	2.15	0.79
1:A:2323:LEU:O	1:A:2331:ASN:ND2	2.15	0.79
1:E:2600:LYS:NZ	1:E:2601:PRO:O	2.14	0.79
1:C:2259:LYS:NZ	1:C:2306:SER:OG	2.16	0.79
1:C:2558:LYS:HD2	1:C:2559:ASN:H	1.48	0.79
1:A:2558:LYS:HD2	1:A:2559:ASN:H	1.48	0.78
1:A:2748:ARG:NH1	1:E:1550:ASP:OD1	2.17	0.78
1:A:2582:THR:HA	1:A:2615:THR:HG22	1.67	0.77
1:E:2582:THR:HA	1:E:2615:THR:HG22	1.67	0.77
1:A:2466:GLN:HA	1:A:2491:LYS:HD3	1.66	0.77
1:E:2259:LYS:NZ	1:E:2306:SER:OG	2.16	0.77
1:C:2600:LYS:NZ	1:C:2601:PRO:O	2.14	0.77
1:A:1012:VAL:HB	1:A:1039:ILE:HB	1.67	0.77
1:E:2558:LYS:HD2	1:E:2559:ASN:H	1.48	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1012:VAL:HB	1:E:1039:ILE:HB	1.67	0.77
1:C:2582:THR:HA	1:C:2615:THR:HG22	1.67	0.76
1:E:2466:GLN:HA	1:E:2491:LYS:HD3	1.66	0.76
1:A:2259:LYS:NZ	1:A:2306:SER:OG	2.15	0.76
1:A:2630:TRP:CH2	1:C:2510:SER:HB3	2.21	0.76
1:A:2468:SER:HA	1:A:2471:LYS:HG2	1.68	0.76
1:C:2466:GLN:HA	1:C:2491:LYS:HD3	1.66	0.75
1:E:2468:SER:HA	1:E:2471:LYS:HG2	1.67	0.75
1:C:1012:VAL:HB	1:C:1039:ILE:HB	1.67	0.75
1:A:1550:ASP:OD1	1:C:2748:ARG:NH1	2.18	0.75
1:C:1114:PHE:O	1:C:1118:PHE:N	2.20	0.75
1:C:1209:ILE:HG23	1:C:1214:PHE:HB2	1.68	0.75
1:E:1710:HIS:HA	1:E:1713:ASN:HD22	1.52	0.75
1:A:1227:VAL:HG12	1:A:1231:MET:HE1	1.68	0.75
1:A:1114:PHE:O	1:A:1118:PHE:N	2.20	0.74
1:A:2485:PHE:HD2	1:A:2486:LEU:HD22	1.52	0.74
1:E:1227:VAL:HG12	1:E:1231:MET:HE1	1.68	0.74
1:A:1710:HIS:HA	1:A:1713:ASN:HD22	1.51	0.74
1:C:2485:PHE:HD2	1:C:2486:LEU:HD22	1.53	0.74
1:A:2467:GLN:HB2	1:A:2470:ASN:HB2	1.69	0.74
1:C:2207:GLY:O	1:C:2211:PHE:N	2.20	0.74
1:E:2485:PHE:HD2	1:E:2486:LEU:HD22	1.53	0.74
1:C:1227:VAL:HG12	1:C:1231:MET:HE1	1.68	0.74
1:E:2207:GLY:O	1:E:2211:PHE:N	2.20	0.73
1:E:1114:PHE:O	1:E:1118:PHE:N	2.20	0.73
1:C:1710:HIS:HA	1:C:1713:ASN:HD22	1.51	0.73
1:C:2468:SER:HA	1:C:2471:LYS:HG2	1.67	0.73
1:C:2630:TRP:CH2	1:E:2510:SER:HB3	2.23	0.73
1:E:1209:ILE:HG23	1:E:1214:PHE:HB2	1.68	0.73
1:A:1263:MET:HE3	1:A:1340:LYS:HA	1.71	0.73
1:C:2467:GLN:HB2	1:C:2470:ASN:HB2	1.69	0.73
1:A:1209:ILE:HG23	1:A:1214:PHE:HB2	1.68	0.73
1:C:1023:ASN:ND2	1:C:1199:PHE:O	2.22	0.73
1:A:1023:ASN:ND2	1:A:1199:PHE:O	2.22	0.73
1:E:1099:THR:H	1:E:1102:HIS:HB2	1.53	0.73
1:E:2480:THR:O	1:E:2484:GLN:N	2.22	0.73
1:A:1228:TYR:HA	1:A:1231:MET:HE2	1.71	0.72
1:A:2510:SER:HB3	1:E:2630:TRP:CH2	2.23	0.72
1:C:979:ARG:NH1	1:C:983:SER:OG	2.21	0.72
1:E:1023:ASN:ND2	1:E:1199:PHE:O	2.22	0.72
1:E:979:ARG:NH1	1:E:983:SER:OG	2.21	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1263:MET:HE3	1:E:1340:LYS:HA	1.71	0.72
1:E:2467:GLN:HB2	1:E:2470:ASN:HB2	1.70	0.72
1:A:2207:GLY:O	1:A:2211:PHE:N	2.20	0.72
1:E:1228:TYR:HA	1:E:1231:MET:HE2	1.71	0.72
1:A:1099:THR:H	1:A:1102:HIS:HB2	1.53	0.72
1:A:979:ARG:NH1	1:A:983:SER:OG	2.21	0.72
1:C:1550:ASP:OD1	1:E:2748:ARG:NH1	2.17	0.72
1:C:1099:THR:H	1:C:1102:HIS:HB2	1.53	0.72
1:C:1228:TYR:HA	1:C:1231:MET:HE2	1.71	0.72
1:E:1154:LEU:O	1:E:1157:ARG:NH2	2.23	0.71
1:A:1154:LEU:O	1:A:1157:ARG:NH2	2.23	0.71
1:A:2359:THR:HG23	1:A:2723:LEU:HD23	1.72	0.71
1:C:1154:LEU:O	1:C:1157:ARG:NH2	2.23	0.71
1:A:1368:CYS:O	1:A:1394:TYR:OH	2.09	0.71
1:C:2345:THR:OG1	1:E:2686:ARG:NH1	2.24	0.71
1:E:1487:ARG:NH2	1:E:1557:ASP:OD1	2.24	0.71
1:A:1487:ARG:NH2	1:A:1557:ASP:OD1	2.24	0.71
1:C:1487:ARG:NH2	1:C:1557:ASP:OD1	2.24	0.71
1:C:2359:THR:HG23	1:C:2723:LEU:HD23	1.72	0.70
1:C:1263:MET:HE3	1:C:1340:LYS:HA	1.71	0.70
1:A:2594:PRO:HD2	1:A:2599:SER:HA	1.73	0.70
1:A:1542:HIS:HA	1:C:2392:ARG:NE	2.06	0.70
1:A:2686:ARG:NH1	1:E:2345:THR:OG1	2.25	0.70
1:C:2480:THR:O	1:C:2484:GLN:N	2.22	0.70
1:E:2594:PRO:HD2	1:E:2599:SER:HA	1.73	0.70
1:C:1368:CYS:O	1:C:1394:TYR:OH	2.09	0.70
1:E:2359:THR:HG23	1:E:2723:LEU:HD23	1.72	0.70
1:C:1198:ARG:HH22	1:C:1209:ILE:HD13	1.57	0.70
1:C:2467:GLN:OE1	1:C:2467:GLN:N	2.25	0.70
1:E:1368:CYS:O	1:E:1394:TYR:OH	2.09	0.69
1:E:1198:ARG:HH22	1:E:1209:ILE:HD13	1.57	0.69
1:A:1198:ARG:HH22	1:A:1209:ILE:HD13	1.57	0.69
1:A:2467:GLN:OE1	1:A:2467:GLN:N	2.25	0.69
1:E:1384:ALA:HA	1:E:1984:PHE:HE2	1.57	0.69
1:A:1713:ASN:HA	1:A:1717:GLU:HB2	1.75	0.69
1:C:2458:GLN:NE2	1:E:2527:SER:O	2.26	0.69
1:C:2698:PHE:HD2	1:E:2694:HIS:ND1	1.91	0.69
1:E:1061:LEU:HD22	1:E:1064:LEU:HD12	1.74	0.69
1:C:1061:LEU:HD22	1:C:1064:LEU:HD12	1.74	0.69
1:C:1713:ASN:HA	1:C:1717:GLU:HB2	1.75	0.69
1:A:1282:CYS:SG	1:A:1288:MET:HB3	2.33	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1452:ALA:HB2	1:A:2037:GLY:HA3	1.75	0.69
1:C:2594:PRO:HD2	1:C:2599:SER:HA	1.73	0.69
1:C:2738:ARG:HG3	1:E:2697:MET:HA	1.74	0.69
1:E:1699:THR:N	1:E:1702:SER:OG	2.23	0.69
1:E:1388:ALA:HB2	1:E:1406:PRO:HG2	1.75	0.69
1:E:1962:ARG:HB2	1:E:2042:ASP:HB3	1.75	0.69
1:A:1384:ALA:HA	1:A:1984:PHE:HE2	1.57	0.69
1:C:2234:LEU:HD21	1:E:2421:LEU:HA	1.75	0.69
1:A:2345:THR:OG1	1:C:2686:ARG:NH1	2.26	0.68
1:C:1282:CYS:SG	1:C:1288:MET:HB3	2.33	0.68
1:C:1542:HIS:HA	1:E:2392:ARG:NE	2.07	0.68
1:A:1554:PHE:HE1	1:A:2732:LYS:HG3	1.58	0.68
1:A:1061:LEU:HD22	1:A:1064:LEU:HD12	1.74	0.68
1:A:2458:GLN:NE2	1:C:2527:SER:O	2.25	0.68
1:A:1134:VAL:HG22	1:A:1228:TYR:HB3	1.76	0.68
1:E:1554:PHE:HE1	1:E:2732:LYS:HG3	1.58	0.68
1:E:2467:GLN:N	1:E:2467:GLN:OE1	2.25	0.68
1:A:1025:PRO:HD2	1:A:1028:GLU:HG3	1.76	0.68
1:C:1384:ALA:HA	1:C:1984:PHE:HE2	1.57	0.68
1:A:1962:ARG:HB2	1:A:2042:ASP:HB3	1.75	0.68
1:C:1554:PHE:HE1	1:C:2732:LYS:HG3	1.58	0.68
1:C:2371:ILE:HG12	1:C:2737:TYR:HE2	1.58	0.68
1:E:1316:PHE:O	1:E:1352:TYR:OH	2.12	0.68
1:E:1713:ASN:HA	1:E:1717:GLU:HB2	1.75	0.68
1:A:1316:PHE:O	1:A:1352:TYR:OH	2.12	0.68
1:C:1962:ARG:HB2	1:C:2042:ASP:HB3	1.75	0.68
1:C:2591:VAL:HA	1:C:2601:PRO:HA	1.76	0.68
1:E:1025:PRO:HD2	1:E:1028:GLU:HG3	1.76	0.68
1:E:1282:CYS:SG	1:E:1288:MET:HB3	2.33	0.68
1:A:2454:MET:HE2	1:A:2500:GLU:H	1.59	0.67
1:A:2591:VAL:HA	1:A:2601:PRO:HA	1.76	0.67
1:C:1452:ALA:HB2	1:C:2037:GLY:HA3	1.75	0.67
1:A:2527:SER:O	1:E:2458:GLN:NE2	2.27	0.67
1:A:2371:ILE:HG12	1:A:2737:TYR:HE2	1.58	0.67
1:A:2738:ARG:HG3	1:C:2697:MET:HA	1.75	0.67
1:E:2326:SER:OG	1:E:2331:ASN:OD1	2.11	0.67
1:E:2591:VAL:HA	1:E:2601:PRO:HA	1.76	0.67
1:A:1272:GLN:NE2	1:A:1693:LYS:O	2.28	0.67
1:A:1388:ALA:HB2	1:A:1406:PRO:HG2	1.75	0.67
1:A:2697:MET:HA	1:E:2738:ARG:HG3	1.75	0.67
1:A:2698:PHE:HD2	1:C:2694:HIS:ND1	1.93	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1388:ALA:HB2	1:C:1406:PRO:HG2	1.75	0.67
1:A:2608:GLU:HA	1:A:2611:PHE:HB2	1.75	0.67
1:A:2480:THR:O	1:A:2484:GLN:N	2.22	0.66
1:A:2694:HIS:ND1	1:E:2698:PHE:HD2	1.92	0.66
1:C:1134:VAL:HG22	1:C:1228:TYR:HB3	1.76	0.66
1:C:1272:GLN:NE2	1:C:1693:LYS:O	2.28	0.66
1:E:1100:ARG:HB2	1:E:1255:ALA:HA	1.78	0.66
1:E:2566:ALA:HA	1:E:2569:ILE:HD12	1.78	0.66
1:E:2608:GLU:HA	1:E:2611:PHE:HB2	1.76	0.66
1:C:1316:PHE:O	1:C:1352:TYR:OH	2.12	0.66
1:C:2015:TYR:HD2	1:C:2016:VAL:HG13	1.60	0.66
1:C:2454:MET:HE2	1:C:2500:GLU:H	1.59	0.66
1:E:2371:ILE:HG12	1:E:2737:TYR:HE2	1.58	0.66
1:A:2566:ALA:HA	1:A:2569:ILE:HD12	1.77	0.66
1:E:1452:ALA:HB2	1:E:2037:GLY:HA3	1.75	0.66
1:C:1025:PRO:HD2	1:C:1028:GLU:HG3	1.76	0.66
1:C:2566:ALA:HA	1:C:2569:ILE:HD12	1.77	0.66
1:E:2454:MET:HE2	1:E:2500:GLU:H	1.59	0.66
1:E:1134:VAL:HG22	1:E:1228:TYR:HB3	1.76	0.66
1:A:1100:ARG:HB2	1:A:1255:ALA:HA	1.77	0.66
1:C:955:VAL:HG13	1:C:1223:PRO:HB2	1.78	0.66
1:C:1699:THR:N	1:C:1702:SER:OG	2.23	0.66
1:E:2015:TYR:HD2	1:E:2016:VAL:HG13	1.60	0.66
1:E:2558:LYS:HD2	1:E:2559:ASN:N	2.11	0.66
1:A:2392:ARG:NE	1:E:1542:HIS:HA	2.07	0.66
1:E:1476:GLN:OE1	1:E:1479:ARG:NH1	2.29	0.66
1:A:2421:LEU:HA	1:E:2234:LEU:HD21	1.76	0.65
1:C:2608:GLU:HA	1:C:2611:PHE:HB2	1.76	0.65
1:A:1876:HIS:ND1	1:A:1877:GLU:OE1	2.26	0.65
1:C:1934:HIS:ND1	1:C:1942:THR:O	2.30	0.65
1:C:2465:ASP:OD1	1:C:2471:LYS:NZ	2.25	0.65
1:A:1934:HIS:ND1	1:A:1942:THR:O	2.30	0.65
1:C:2558:LYS:HD2	1:C:2559:ASN:N	2.11	0.65
1:E:1272:GLN:NE2	1:E:1693:LYS:O	2.28	0.65
1:C:1450:ARG:NH1	1:C:1453:GLU:OE2	2.30	0.65
1:A:1476:GLN:OE1	1:A:1479:ARG:NH1	2.29	0.65
1:A:2015:TYR:HD2	1:A:2016:VAL:HG13	1.60	0.65
1:C:1100:ARG:HB2	1:C:1255:ALA:HA	1.77	0.65
1:A:1699:THR:N	1:A:1702:SER:OG	2.23	0.65
1:C:1888:MET:HG3	1:C:1889:PHE:H	1.62	0.65
1:E:955:VAL:HG13	1:E:1223:PRO:HB2	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1876:HIS:ND1	1:C:1877:GLU:OE1	2.26	0.65
1:C:2532:PHE:O	1:C:2553:LEU:N	2.29	0.65
1:C:1476:GLN:OE1	1:C:1479:ARG:NH1	2.29	0.65
1:E:2443:ILE:HG13	1:E:2532:PHE:HD1	1.62	0.65
1:A:955:VAL:HG13	1:A:1223:PRO:HB2	1.78	0.64
1:A:2326:SER:OG	1:A:2331:ASN:OD1	2.11	0.64
1:C:1477:LEU:HB2	1:C:2317:ARG:HH22	1.62	0.64
1:A:2234:LEU:HD21	1:C:2421:LEU:HA	1.77	0.64
1:A:2480:THR:HB	1:A:2484:GLN:HG2	1.79	0.64
1:E:1934:HIS:ND1	1:E:1942:THR:O	2.30	0.64
1:E:2480:THR:HB	1:E:2484:GLN:HG2	1.79	0.64
1:A:1888:MET:HG3	1:A:1889:PHE:H	1.62	0.64
1:C:2488:ASN:HB2	1:C:2489:TYR:CZ	2.33	0.64
1:E:2442:THR:OG1	1:E:2533:SER:OG	2.16	0.64
1:A:2442:THR:OG1	1:A:2533:SER:OG	2.16	0.64
1:A:2443:ILE:HG13	1:A:2532:PHE:HD1	1.62	0.64
1:E:1450:ARG:NH1	1:E:1453:GLU:OE2	2.30	0.64
1:E:2415:ILE:HG23	1:E:2416:VAL:HG23	1.80	0.64
1:A:1477:LEU:HB2	1:A:2317:ARG:HH22	1.62	0.64
1:C:2442:THR:OG1	1:C:2533:SER:OG	2.16	0.64
1:C:2491:LYS:NZ	1:C:2492:GLU:OE2	2.31	0.64
1:E:1888:MET:HG3	1:E:1889:PHE:H	1.62	0.64
1:A:1450:ARG:NH1	1:A:1453:GLU:OE2	2.30	0.64
1:E:2491:LYS:NZ	1:E:2492:GLU:OE2	2.31	0.64
1:A:1278:ASP:O	1:A:1283:ARG:NH2	2.31	0.64
1:A:2558:LYS:HD2	1:A:2559:ASN:N	2.11	0.64
1:E:1278:ASP:O	1:E:1283:ARG:NH2	2.31	0.64
1:E:1282:CYS:HG	1:E:1900:PHE:HD2	1.46	0.64
1:E:1477:LEU:HB2	1:E:2317:ARG:HH22	1.62	0.64
1:C:1540:VAL:O	1:E:2392:ARG:NH2	2.31	0.64
1:A:2510:SER:O	1:A:2514:LYS:N	2.31	0.64
1:C:2510:SER:O	1:C:2514:LYS:N	2.31	0.63
1:E:1007:PRO:O	1:E:1011:SER:OG	2.12	0.63
1:A:1540:VAL:HB	1:A:1542:HIS:HB2	1.80	0.63
1:C:2326:SER:OG	1:C:2331:ASN:OD1	2.11	0.63
1:A:1540:VAL:O	1:C:2392:ARG:NH2	2.31	0.63
1:A:2630:TRP:HZ2	1:C:2511:PRO:HD2	1.63	0.63
1:C:1278:ASP:O	1:C:1283:ARG:NH2	2.31	0.63
1:C:2523:ASP:HB2	1:C:2526:SER:HB3	1.81	0.63
1:E:2523:ASP:HB2	1:E:2526:SER:HB3	1.81	0.63
1:A:2491:LYS:NZ	1:A:2492:GLU:OE2	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2539:ASN:OD1	1:E:2540:LEU:N	2.31	0.63
1:A:1370:TYR:HB3	1:A:1374:LEU:HD13	1.81	0.63
1:A:2488:ASN:HB2	1:A:2489:TYR:CZ	2.33	0.63
1:A:2539:ASN:OD1	1:A:2540:LEU:N	2.31	0.63
2:D:238:GLU:HG2	1:E:2404:MET:HE1	1.79	0.63
1:E:1875:THR:HA	1:E:1878:LEU:HB2	1.81	0.63
1:E:2488:ASN:HB2	1:E:2489:TYR:CZ	2.33	0.63
1:A:2392:ARG:NH2	1:E:1540:VAL:O	2.32	0.63
1:C:2480:THR:HB	1:C:2484:GLN:HG2	1.79	0.63
1:C:2415:ILE:HG23	1:C:2416:VAL:HG23	1.80	0.63
1:C:2443:ILE:HG13	1:C:2532:PHE:HD1	1.62	0.63
1:E:1466:ARG:HE	1:E:2185:ALA:HB1	1.63	0.63
1:E:1876:HIS:ND1	1:E:1877:GLU:OE1	2.26	0.63
1:A:2465:ASP:C	1:A:2491:LYS:HE2	2.24	0.62
1:C:2465:ASP:C	1:C:2491:LYS:HE2	2.24	0.62
1:A:1709:ASN:O	1:A:1713:ASN:ND2	2.32	0.62
1:C:2630:TRP:HZ2	1:E:2511:PRO:HD2	1.65	0.62
1:A:2234:LEU:HG	1:C:2417:TRP:CZ3	2.34	0.62
1:A:2415:ILE:HG23	1:A:2416:VAL:HG23	1.80	0.62
1:A:2523:ASP:HB2	1:A:2526:SER:HB3	1.81	0.62
1:A:2568:MET:HE1	1:A:2631:TRP:HH2	1.65	0.62
1:C:1540:VAL:HB	1:C:1542:HIS:HB2	1.81	0.62
1:C:2539:ASN:OD1	1:C:2540:LEU:N	2.31	0.62
1:A:1466:ARG:HE	1:A:2185:ALA:HB1	1.63	0.62
1:E:1709:ASN:O	1:E:1713:ASN:ND2	2.33	0.62
1:E:2465:ASP:C	1:E:2491:LYS:HE2	2.24	0.62
1:C:2015:TYR:CD2	1:C:2016:VAL:HG13	2.35	0.62
1:A:2529:SER:HA	1:A:2556:PRO:HA	1.82	0.62
1:C:978:LEU:HD22	1:C:981:LEU:HD21	1.82	0.62
1:C:2568:MET:HE1	1:C:2631:TRP:HH2	1.65	0.62
1:E:978:LEU:HD22	1:E:981:LEU:HD21	1.82	0.62
1:E:1540:VAL:HB	1:E:1542:HIS:HB2	1.80	0.62
1:E:2568:MET:HE1	1:E:2631:TRP:HH2	1.65	0.62
1:C:1930:ILE:HG12	1:C:2022:GLN:NE2	2.15	0.62
1:E:1930:ILE:HG12	1:E:2022:GLN:NE2	2.15	0.62
1:A:1350:ILE:HD11	1:A:1422:LEU:HB3	1.82	0.61
1:E:2634:ASN:ND2	1:E:2645:SER:OG	2.33	0.61
1:A:978:LEU:HD22	1:A:981:LEU:HD21	1.82	0.61
1:C:1709:ASN:O	1:C:1713:ASN:ND2	2.32	0.61
1:A:2015:TYR:CD2	1:A:2016:VAL:HG13	2.35	0.61
1:A:2458:GLN:NE2	1:C:2527:SER:OG	2.33	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1875:THR:HA	1:C:1878:LEU:HB2	1.81	0.61
1:C:2370:ASP:OD2	1:E:2690:SER:HA	2.01	0.61
1:E:1498:MET:HE2	1:E:1498:MET:H	1.65	0.61
1:A:1950:PHE:HZ	1:A:2300:CYS:SG	2.23	0.61
1:A:2477:SER:C	1:A:2479:ASP:H	2.09	0.61
1:A:2417:TRP:CZ3	1:E:2234:LEU:HG	2.36	0.61
1:C:1350:ILE:HD11	1:C:1422:LEU:HB3	1.82	0.61
1:C:1370:TYR:HB3	1:C:1374:LEU:HD13	1.81	0.61
1:C:2634:ASN:ND2	1:C:2645:SER:OG	2.33	0.61
1:E:1370:TYR:HB3	1:E:1374:LEU:HD13	1.81	0.61
1:E:2015:TYR:CD2	1:E:2016:VAL:HG13	2.34	0.61
1:A:2511:PRO:HD2	1:E:2630:TRP:HZ2	1.66	0.61
1:C:1713:ASN:O	1:C:1718:SER:N	2.34	0.61
2:B:245:PRO:HG2	1:C:2374:HIS:ND1	2.15	0.61
1:A:1875:THR:HA	1:A:1878:LEU:HB2	1.81	0.61
1:C:1466:ARG:HE	1:C:2185:ALA:HB1	1.63	0.61
1:E:982:ALA:HA	1:E:985:VAL:HG12	1.83	0.61
1:C:982:ALA:HA	1:C:985:VAL:HG12	1.83	0.61
1:C:1107:LEU:HA	1:C:1110:CYS:HB2	1.83	0.61
1:A:1498:MET:HE2	1:A:1498:MET:H	1.65	0.60
1:A:1930:ILE:HG12	1:A:2022:GLN:NE2	2.15	0.60
1:C:2529:SER:HA	1:C:2556:PRO:HA	1.82	0.60
1:E:2529:SER:HA	1:E:2556:PRO:HA	1.82	0.60
1:A:1140:ASP:OD1	1:A:1143:ALA:N	2.34	0.60
1:C:1498:MET:HE2	1:C:1498:MET:H	1.65	0.60
1:E:2465:ASP:OD1	1:E:2471:LYS:NZ	2.25	0.60
1:A:1354:VAL:O	1:A:1358:THR:OG1	2.18	0.60
1:C:1950:PHE:HZ	1:C:2300:CYS:SG	2.23	0.60
1:E:1350:ILE:HD11	1:E:1422:LEU:HB3	1.82	0.60
1:A:1107:LEU:HA	1:A:1110:CYS:HB2	1.83	0.60
1:E:950:ILE:HD11	1:E:965:PHE:HB3	1.83	0.60
1:E:1950:PHE:HZ	1:E:2300:CYS:SG	2.23	0.60
1:A:2634:ASN:ND2	1:A:2645:SER:OG	2.33	0.60
1:E:2532:PHE:O	1:E:2553:LEU:N	2.29	0.60
1:A:1098:ILE:HG23	1:A:1102:HIS:HB3	1.84	0.60
1:E:1140:ASP:OD1	1:E:1143:ALA:N	2.34	0.60
1:A:1713:ASN:O	1:A:1718:SER:N	2.34	0.60
1:E:1098:ILE:HG23	1:E:1102:HIS:HB3	1.84	0.60
1:E:2477:SER:C	1:E:2479:ASP:H	2.09	0.60
1:A:2488:ASN:HB2	1:A:2489:TYR:CE1	2.37	0.60
1:C:2477:SER:C	1:C:2479:ASP:H	2.09	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1252:ARG:HG3	1:C:1257:ASP:HB2	1.84	0.59
1:C:2290:LEU:O	1:C:2293:GLN:N	2.36	0.59
1:C:1098:ILE:HG23	1:C:1102:HIS:HB3	1.84	0.59
1:C:2488:ASN:HB2	1:C:2489:TYR:CE1	2.37	0.59
1:A:967:ILE:HG12	1:A:1178:THR:HG22	1.85	0.59
1:C:1477:LEU:HB2	1:C:2317:ARG:NH2	2.18	0.59
1:E:1107:LEU:HA	1:E:1110:CYS:HB2	1.83	0.59
1:A:2690:SER:HA	1:E:2370:ASP:OD2	2.01	0.59
1:E:1713:ASN:O	1:E:1718:SER:N	2.33	0.59
1:A:2234:LEU:HG	1:C:2417:TRP:CH2	2.38	0.59
1:A:2527:SER:OG	1:E:2458:GLN:NE2	2.34	0.59
1:C:950:ILE:HD11	1:C:965:PHE:HB3	1.83	0.59
1:C:1140:ASP:OD1	1:C:1143:ALA:N	2.34	0.59
1:C:2234:LEU:HG	1:E:2417:TRP:CZ3	2.37	0.59
1:E:2290:LEU:O	1:E:2293:GLN:N	2.36	0.59
1:E:2510:SER:O	1:E:2514:LYS:N	2.31	0.59
1:A:950:ILE:HD11	1:A:965:PHE:HB3	1.83	0.59
1:C:2468:SER:HB2	1:C:2472:PHE:CB	2.28	0.59
1:A:982:ALA:HA	1:A:985:VAL:HG12	1.83	0.59
1:C:2458:GLN:NE2	1:E:2527:SER:OG	2.35	0.59
1:E:2179:ILE:HG22	1:E:2180:HIS:CD2	2.38	0.59
1:E:2488:ASN:HB2	1:E:2489:TYR:CE1	2.37	0.59
1:A:2532:PHE:O	1:A:2553:LEU:N	2.29	0.59
1:C:2568:MET:HE1	1:C:2631:TRP:CH2	2.38	0.59
1:A:2630:TRP:CZ2	1:C:2511:PRO:HD2	2.38	0.58
1:C:961:PHE:HB2	1:C:993:ILE:HG23	1.85	0.58
1:A:2290:LEU:O	1:A:2293:GLN:N	2.36	0.58
1:A:2568:MET:HE1	1:A:2631:TRP:CH2	2.38	0.58
1:C:967:ILE:HG12	1:C:1178:THR:HG22	1.85	0.58
1:E:967:ILE:HG12	1:E:1178:THR:HG22	1.85	0.58
1:A:1363:LEU:HB3	1:A:1387:LEU:HD11	1.85	0.58
1:A:2370:ASP:OD2	1:C:2690:SER:HA	2.02	0.58
1:A:2510:SER:OG	1:E:2505:SER:O	2.21	0.58
1:C:2179:ILE:HG22	1:C:2180:HIS:CD2	2.38	0.58
1:A:1477:LEU:HB2	1:A:2317:ARG:NH2	2.18	0.58
1:A:1699:THR:H	1:A:1702:SER:HG	1.50	0.58
1:A:2179:ILE:HG22	1:A:2180:HIS:CD2	2.38	0.58
1:A:2505:SER:O	1:C:2510:SER:OG	2.20	0.58
1:C:2497:ALA:HB3	1:C:2650:LEU:HB3	1.86	0.58
1:A:1252:ARG:HG3	1:A:1257:ASP:HB2	1.84	0.58
1:A:1666:TRP:O	1:A:1670:ILE:HD12	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2497:ALA:HB3	1:A:2650:LEU:HB3	1.86	0.58
1:E:964:VAL:O	1:E:968:SER:N	2.30	0.58
1:E:1666:TRP:O	1:E:1670:ILE:HD12	2.04	0.58
1:E:2497:ALA:HB3	1:E:2650:LEU:HB3	1.86	0.58
1:C:2558:LYS:NZ	1:C:2560:ILE:HB	2.18	0.58
1:E:992:VAL:HA	1:E:995:VAL:HG22	1.86	0.58
1:E:1699:THR:H	1:E:1702:SER:HG	1.48	0.58
1:C:2618:LEU:HD12	1:C:2630:TRP:O	2.04	0.58
1:A:2348:ARG:NH1	1:A:2352:ASP:OD2	2.37	0.58
1:C:1666:TRP:O	1:C:1670:ILE:HD12	2.04	0.58
1:E:1252:ARG:HG3	1:E:1257:ASP:HB2	1.84	0.58
1:E:1477:LEU:HB2	1:E:2317:ARG:NH2	2.18	0.58
1:A:1455:PHE:HE2	1:A:1959:ARG:HH22	1.52	0.58
1:A:2558:LYS:NZ	1:A:2560:ILE:HB	2.18	0.58
1:A:2618:LEU:HD12	1:A:2630:TRP:O	2.04	0.58
1:C:2505:SER:O	1:E:2510:SER:OG	2.22	0.58
1:E:2558:LYS:NZ	1:E:2560:ILE:HB	2.18	0.58
1:E:2618:LEU:HD12	1:E:2630:TRP:O	2.04	0.58
1:E:2568:MET:HE1	1:E:2631:TRP:CH2	2.38	0.57
1:C:990:THR:HG21	1:C:1063:MET:HE1	1.87	0.57
1:C:1363:LEU:HB3	1:C:1387:LEU:HD11	1.85	0.57
1:C:992:VAL:HA	1:C:995:VAL:HG22	1.86	0.57
1:E:1363:LEU:HB3	1:E:1387:LEU:HD11	1.85	0.57
1:C:2630:TRP:CZ2	1:E:2511:PRO:HD2	2.39	0.57
1:C:2717:VAL:HG11	1:C:2725:LEU:HB2	1.87	0.57
1:C:1699:THR:H	1:C:1702:SER:HG	1.51	0.57
1:C:2576:SER:C	1:C:2579:THR:H	2.13	0.57
1:E:1653:TRP:HA	1:E:1656:PHE:HD2	1.70	0.57
1:A:2537:GLN:HA	1:A:2548:ILE:HG12	1.86	0.57
1:C:959:SER:OG	1:C:962:ASN:OD1	2.22	0.57
1:E:2371:ILE:HG12	1:E:2737:TYR:CE2	2.39	0.57
1:C:2348:ARG:NH1	1:C:2352:ASP:OD2	2.37	0.57
1:E:2504:ASN:OD1	1:E:2505:SER:N	2.38	0.57
1:A:2504:ASN:OD1	1:A:2505:SER:N	2.38	0.57
1:A:2511:PRO:HD2	1:E:2630:TRP:CZ2	2.40	0.57
1:C:2537:GLN:HA	1:C:2548:ILE:HG12	1.86	0.57
1:A:1213:TYR:OH	1:A:1222:ASN:O	2.21	0.57
1:E:1310:THR:HB	1:E:2016:VAL:HG21	1.87	0.57
1:A:950:ILE:HD13	1:A:969:TRP:CD1	2.40	0.57
1:A:1966:MET:HE3	1:A:1966:MET:HA	1.87	0.57
1:E:2348:ARG:NH1	1:E:2352:ASP:OD2	2.37	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:961:PHE:HB2	1:A:993:ILE:HG23	1.85	0.56
1:C:964:VAL:O	1:C:968:SER:N	2.30	0.56
1:C:2234:LEU:HG	1:E:2417:TRP:CH2	2.39	0.56
1:E:990:THR:HG21	1:E:1063:MET:HE1	1.86	0.56
1:E:2700:GLU:N	1:E:2700:GLU:OE1	2.38	0.56
1:A:992:VAL:HA	1:A:995:VAL:HG22	1.86	0.56
1:A:2417:TRP:CH2	1:E:2234:LEU:HG	2.39	0.56
1:C:1310:THR:HB	1:C:2016:VAL:HG21	1.87	0.56
1:C:1653:TRP:HA	1:C:1656:PHE:HD2	1.70	0.56
1:C:1712:MET:C	1:C:1717:GLU:HG2	2.30	0.56
1:C:2179:ILE:HD13	1:C:2257:LEU:HD12	1.87	0.56
1:C:2434:ASN:HB2	1:C:2657:VAL:HG13	1.87	0.56
1:E:950:ILE:HD13	1:E:969:TRP:CD1	2.40	0.56
1:A:1653:TRP:HA	1:A:1656:PHE:HD2	1.70	0.56
1:A:2534:TRP:CZ2	1:A:2551:ASP:HB3	2.40	0.56
1:C:1098:ILE:HG13	1:C:1109:ASN:HB3	1.87	0.56
1:C:1213:TYR:OH	1:C:1222:ASN:O	2.21	0.56
1:C:1966:MET:HE3	1:C:1966:MET:HA	1.87	0.56
1:C:2371:ILE:HG12	1:C:2737:TYR:CE2	2.39	0.56
1:E:2179:ILE:HD13	1:E:2257:LEU:HD12	1.87	0.56
1:E:2534:TRP:CZ2	1:E:2551:ASP:HB3	2.40	0.56
1:E:2576:SER:C	1:E:2579:THR:H	2.13	0.56
1:E:2717:VAL:HG11	1:E:2725:LEU:HB2	1.87	0.56
1:C:2504:ASN:OD1	1:C:2505:SER:N	2.38	0.56
1:E:2583:ILE:O	1:E:2613:ASP:HB3	2.06	0.56
1:A:1931:ILE:O	1:A:1935:MET:HG3	2.06	0.56
1:E:2477:SER:C	1:E:2479:ASP:N	2.62	0.56
1:A:1712:MET:C	1:A:1717:GLU:HG2	2.30	0.56
1:A:2430:ALA:N	1:A:2658:SER:OG	2.27	0.56
1:C:1041:PRO:O	1:C:1045:VAL:HG22	2.06	0.56
1:E:961:PHE:HB2	1:E:993:ILE:HG23	1.85	0.56
1:E:1098:ILE:HG13	1:E:1109:ASN:HB3	1.88	0.56
1:A:1709:ASN:HA	1:A:1712:MET:HE2	1.88	0.56
1:A:2179:ILE:HD13	1:A:2257:LEU:HD12	1.87	0.56
1:E:1712:MET:C	1:E:1717:GLU:HG2	2.30	0.56
1:E:2468:SER:HB2	1:E:2472:PHE:CB	2.28	0.56
1:A:1310:THR:HB	1:A:2016:VAL:HG21	1.87	0.56
1:A:1384:ALA:HA	1:A:1984:PHE:CE2	2.40	0.56
1:C:1931:ILE:O	1:C:1935:MET:HG3	2.06	0.56
1:C:2207:GLY:HA3	1:C:2291:VAL:HG11	1.88	0.56
1:E:1709:ASN:HA	1:E:1712:MET:HE2	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1098:ILE:HG13	1:A:1109:ASN:HB3	1.88	0.56
1:C:950:ILE:HD13	1:C:969:TRP:CD1	2.40	0.56
1:C:1455:PHE:HE2	1:C:1959:ARG:HH22	1.52	0.56
1:C:2339:ARG:HG3	1:C:2345:THR:HG21	1.88	0.56
1:E:1931:ILE:O	1:E:1935:MET:HG3	2.06	0.56
1:A:1041:PRO:O	1:A:1045:VAL:HG22	2.06	0.56
1:A:2437:LEU:HD23	1:A:2437:LEU:H	1.70	0.56
1:A:2583:ILE:O	1:A:2613:ASP:HB3	2.06	0.56
1:C:1709:ASN:HA	1:C:1712:MET:HE2	1.88	0.56
1:C:2700:GLU:N	1:C:2700:GLU:OE1	2.38	0.56
1:E:1966:MET:HE3	1:E:1966:MET:HA	1.87	0.56
1:E:2207:GLY:HA3	1:E:2291:VAL:HG11	1.88	0.56
1:A:2717:VAL:HG11	1:A:2725:LEU:HB2	1.87	0.55
1:C:1044:TRP:CH2	1:C:1188:PRO:HG3	2.41	0.55
1:E:1455:PHE:HE2	1:E:1959:ARG:HH22	1.52	0.55
1:A:990:THR:HG21	1:A:1063:MET:HE1	1.86	0.55
1:A:2468:SER:HB2	1:A:2472:PHE:CB	2.28	0.55
1:A:2576:SER:HA	1:A:2579:THR:HB	1.89	0.55
1:E:1041:PRO:O	1:E:1045:VAL:HG22	2.06	0.55
1:E:2537:GLN:HA	1:E:2548:ILE:HG12	1.86	0.55
1:E:2587:TYR:O	1:E:2612:MET:HB3	2.06	0.55
1:E:2437:LEU:HD23	1:E:2437:LEU:H	1.70	0.55
1:E:2585:LYS:NZ	1:E:2609:ASN:O	2.23	0.55
1:E:2646:GLN:HG3	1:E:2647:ALA:N	2.22	0.55
1:C:2587:TYR:O	1:C:2612:MET:HB3	2.06	0.55
1:E:2339:ARG:HG3	1:E:2345:THR:HG21	1.88	0.55
1:A:959:SER:OG	1:A:962:ASN:OD1	2.22	0.55
1:E:1044:TRP:CH2	1:E:1188:PRO:HG3	2.41	0.55
1:E:2434:ASN:HB2	1:E:2657:VAL:HG13	1.87	0.55
1:E:2480:THR:HB	1:E:2484:GLN:CG	2.36	0.55
1:A:1044:TRP:CH2	1:A:1188:PRO:HG3	2.41	0.55
1:A:2319:LEU:HD12	1:A:2719:GLU:HB2	1.89	0.55
1:A:2587:TYR:O	1:A:2612:MET:HB3	2.06	0.55
1:C:1280:ILE:HG13	1:C:1436:HIS:CE1	2.42	0.55
1:C:2319:LEU:HD12	1:C:2719:GLU:HB2	1.89	0.55
1:E:1319:GLY:HA3	1:E:1352:TYR:CZ	2.42	0.55
1:A:2465:ASP:OD1	1:A:2471:LYS:NZ	2.25	0.55
1:A:2510:SER:HB3	1:E:2630:TRP:HH2	1.72	0.55
1:C:2437:LEU:HD23	1:C:2437:LEU:H	1.70	0.55
1:C:2534:TRP:CZ2	1:C:2551:ASP:HB3	2.40	0.55
1:A:2576:SER:C	1:A:2579:THR:H	2.13	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1100:ARG:HB2	1:C:1256:GLY:H	1.71	0.55
1:A:2207:GLY:HA3	1:A:2291:VAL:HG11	1.88	0.55
1:C:1319:GLY:HA3	1:C:1352:TYR:CZ	2.42	0.55
1:C:2464:MET:HE2	1:C:2465:ASP:O	2.07	0.55
1:E:1001:GLN:HG3	1:E:1051:SER:HB3	1.89	0.55
1:E:2576:SER:HA	1:E:2579:THR:HB	1.89	0.55
1:E:2592:LYS:N	1:E:2600:LYS:O	2.34	0.55
1:A:964:VAL:HA	1:A:967:ILE:HB	1.89	0.55
1:A:1007:PRO:O	1:A:1011:SER:OG	2.12	0.55
1:A:1156:ARG:HG3	1:A:1165:ILE:HD13	1.89	0.55
1:A:2434:ASN:HB2	1:A:2657:VAL:HG13	1.87	0.55
1:A:2630:TRP:HH2	1:C:2510:SER:HB3	1.70	0.55
1:C:2477:SER:C	1:C:2479:ASP:N	2.62	0.55
1:E:2350:VAL:HG12	1:E:2351:MET:HE2	1.89	0.55
1:A:978:LEU:O	1:A:982:ALA:N	2.38	0.54
1:C:2533:SER:HB2	1:C:2551:ASP:O	2.07	0.54
1:E:1403:CYS:SG	1:E:1404:THR:N	2.80	0.54
1:A:1029:LEU:O	1:A:1036:SER:HB3	2.07	0.54
1:A:1280:ILE:HG13	1:A:1436:HIS:CE1	2.42	0.54
1:A:2339:ARG:HG3	1:A:2345:THR:HG21	1.88	0.54
1:A:2480:THR:HB	1:A:2484:GLN:CG	2.36	0.54
1:A:2700:GLU:N	1:A:2700:GLU:OE1	2.38	0.54
1:E:1029:LEU:O	1:E:1036:SER:HB3	2.07	0.54
1:E:1100:ARG:HB2	1:E:1256:GLY:H	1.72	0.54
1:E:1180:GLN:HB3	1:E:1226:LEU:HD21	1.89	0.54
1:A:1319:GLY:HA3	1:A:1352:TYR:CZ	2.42	0.54
1:A:1699:THR:O	1:A:1703:ILE:N	2.37	0.54
1:C:1156:ARG:HG3	1:C:1165:ILE:HD13	1.89	0.54
1:C:2480:THR:HB	1:C:2484:GLN:CG	2.36	0.54
1:C:2718:ARG:NH2	1:C:2726:GLU:OE2	2.40	0.54
1:E:1156:ARG:HG3	1:E:1165:ILE:HD13	1.89	0.54
1:E:2490:GLU:N	1:E:2493:ASP:OD2	2.41	0.54
1:A:2507:TRP:HB3	1:A:2629:GLU:HB3	1.90	0.54
1:A:2533:SER:HB2	1:A:2551:ASP:O	2.07	0.54
1:A:2592:LYS:O	1:A:2599:SER:OG	2.21	0.54
1:C:1029:LEU:O	1:C:1036:SER:HB3	2.07	0.54
1:C:1180:GLN:HB3	1:C:1226:LEU:HD21	1.89	0.54
1:C:1209:ILE:O	1:C:1213:TYR:N	2.41	0.54
1:C:2583:ILE:O	1:C:2613:ASP:HB3	2.06	0.54
1:E:1384:ALA:HA	1:E:1984:PHE:CE2	2.40	0.54
1:A:2464:MET:HE2	1:A:2465:ASP:O	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2490:GLU:N	1:C:2493:ASP:OD2	2.41	0.54
1:A:2490:GLU:N	1:A:2493:ASP:OD2	2.41	0.54
1:C:1007:PRO:O	1:C:1011:SER:OG	2.12	0.54
1:C:1935:MET:HB3	1:C:2293:GLN:OE1	2.08	0.54
1:E:959:SER:OG	1:E:962:ASN:OD1	2.22	0.54
1:E:2464:MET:HE2	1:E:2465:ASP:O	2.07	0.54
1:A:1360:LYS:HD2	1:A:1410:ALA:HB3	1.90	0.54
1:A:2477:SER:C	1:A:2479:ASP:N	2.62	0.54
1:C:2576:SER:HA	1:C:2579:THR:HB	1.89	0.54
1:C:2646:GLN:HG3	1:C:2647:ALA:N	2.22	0.54
1:E:2319:LEU:HD12	1:E:2719:GLU:HB2	1.89	0.54
1:E:2533:SER:HB2	1:E:2551:ASP:O	2.07	0.54
1:A:1100:ARG:HB2	1:A:1256:GLY:H	1.71	0.54
1:A:1209:ILE:O	1:A:1213:TYR:N	2.41	0.54
1:A:2460:GLN:HB3	1:A:2498:GLU:OE1	2.08	0.54
1:A:2646:GLN:HG3	1:A:2647:ALA:N	2.22	0.54
1:A:2653:PHE:HB3	1:A:2654:ASN:CG	2.33	0.54
1:C:1360:LYS:HD2	1:C:1410:ALA:HB3	1.90	0.54
1:C:2653:PHE:HB3	1:C:2654:ASN:CG	2.33	0.54
1:E:2460:GLN:HB3	1:E:2498:GLU:OE1	2.08	0.54
1:A:2454:MET:SD	1:A:2455:SER:N	2.81	0.54
1:E:2575:GLU:O	1:E:2579:THR:N	2.41	0.54
1:C:964:VAL:HA	1:C:967:ILE:HB	1.89	0.54
1:C:1267:ALA:HB1	1:C:1439:ALA:HB2	1.90	0.54
1:C:2460:GLN:HB3	1:C:2498:GLU:OE1	2.08	0.54
1:E:1209:ILE:O	1:E:1213:TYR:N	2.41	0.54
1:E:1280:ILE:HG13	1:E:1436:HIS:CE1	2.42	0.54
1:E:2444:THR:HG22	1:E:2450:PRO:HB3	1.90	0.54
1:E:2454:MET:SD	1:E:2455:SER:N	2.81	0.54
1:C:2575:GLU:O	1:C:2579:THR:N	2.41	0.53
1:E:964:VAL:HA	1:E:967:ILE:HB	1.89	0.53
1:E:1306:PHE:HB2	1:E:1324:CYS:SG	2.48	0.53
1:E:1360:LYS:HD2	1:E:1410:ALA:HB3	1.90	0.53
1:E:2592:LYS:O	1:E:2599:SER:OG	2.21	0.53
1:A:2371:ILE:HG12	1:A:2737:TYR:CE2	2.39	0.53
1:A:2580:PRO:HA	1:A:2617:ILE:HB	1.90	0.53
1:C:1306:PHE:HB2	1:C:1324:CYS:SG	2.48	0.53
1:E:2653:PHE:HB3	1:E:2654:ASN:CG	2.32	0.53
1:A:1267:ALA:HB1	1:A:1439:ALA:HB2	1.90	0.53
1:A:2433:ILE:HG21	1:A:2492:GLU:O	2.08	0.53
1:C:1354:VAL:O	1:C:1358:THR:OG1	2.18	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1384:ALA:HA	1:C:1984:PHE:CE2	2.40	0.53
1:A:1346:TRP:CZ2	1:A:1422:LEU:HD23	2.43	0.53
1:A:1476:GLN:HA	1:A:1479:ARG:CZ	2.39	0.53
1:C:1346:TRP:CZ2	1:C:1422:LEU:HD23	2.43	0.53
1:C:2464:MET:O	1:C:2491:LYS:NZ	2.38	0.53
1:E:2621:ASP:OD1	1:E:2623:THR:N	2.41	0.53
1:A:1001:GLN:HG3	1:A:1051:SER:HB3	1.89	0.53
1:C:1476:GLN:HA	1:C:1479:ARG:CZ	2.39	0.53
1:E:1310:THR:HG23	1:E:1321:LEU:HD21	1.91	0.53
1:E:1476:GLN:HA	1:E:1479:ARG:CZ	2.39	0.53
1:A:964:VAL:O	1:A:968:SER:N	2.30	0.53
1:A:1180:GLN:HB3	1:A:1226:LEU:HD21	1.89	0.53
1:A:1403:CYS:SG	1:A:1404:THR:N	2.80	0.53
1:A:1487:ARG:NH2	1:A:1491:TYR:OH	2.42	0.53
1:A:2575:GLU:O	1:A:2579:THR:N	2.41	0.53
1:C:1001:GLN:HG3	1:C:1051:SER:HB3	1.89	0.53
1:C:1699:THR:O	1:C:1703:ILE:N	2.37	0.53
1:E:2718:ARG:NH2	1:E:2726:GLU:OE2	2.40	0.53
1:A:1935:MET:HB3	1:A:2293:GLN:OE1	2.08	0.53
1:C:1403:CYS:SG	1:C:1404:THR:N	2.80	0.53
1:C:1487:ARG:NH2	1:C:1491:TYR:OH	2.41	0.53
1:C:1493:LYS:HG2	1:C:1497:ARG:HE	1.74	0.53
1:E:1493:LYS:HG2	1:E:1497:ARG:HE	1.73	0.53
1:E:1935:MET:HB3	1:E:2293:GLN:OE1	2.08	0.53
1:E:1941:ILE:O	1:E:1971:TYR:OH	2.21	0.53
1:E:2507:TRP:HB3	1:E:2629:GLU:HB3	1.90	0.53
1:A:1493:LYS:HG2	1:A:1497:ARG:HE	1.73	0.53
1:A:2621:ASP:OD1	1:A:2623:THR:N	2.41	0.53
1:C:1934:HIS:HE1	1:C:1943:LEU:HB2	1.73	0.53
1:E:1267:ALA:HB1	1:E:1439:ALA:HB2	1.91	0.53
1:E:1923:GLU:HG2	1:E:1927:TYR:CE2	2.44	0.53
1:C:2433:ILE:HG21	1:C:2492:GLU:O	2.08	0.53
1:C:2454:MET:SD	1:C:2455:SER:N	2.81	0.53
1:C:2507:TRP:HB3	1:C:2629:GLU:HB3	1.90	0.53
1:E:1487:ARG:NH2	1:E:1491:TYR:OH	2.42	0.53
1:E:2477:SER:OG	1:E:2478:ARG:N	2.42	0.53
1:A:2412:LEU:HD13	1:E:2347:LEU:HD21	1.91	0.53
1:A:2530:VAL:HG11	1:A:2586:ILE:HD13	1.91	0.53
1:C:2350:VAL:HG12	1:C:2351:MET:HE2	1.89	0.53
1:E:2433:ILE:HG21	1:E:2492:GLU:O	2.08	0.53
1:A:1710:HIS:ND1	1:A:1888:MET:HE1	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2477:SER:OG	1:A:2478:ARG:N	2.42	0.52
1:C:950:ILE:HG21	1:C:969:TRP:CD1	2.44	0.52
1:C:1923:GLU:HG2	1:C:1927:TYR:CE2	2.44	0.52
1:C:2638:ASN:C	1:C:2639:ARG:HD3	2.34	0.52
1:E:1346:TRP:CZ2	1:E:1422:LEU:HD23	2.43	0.52
1:E:1711:ILE:HG13	1:E:1886:LYS:HE2	1.91	0.52
1:A:1923:GLU:HG2	1:A:1927:TYR:CE2	2.44	0.52
1:A:2638:ASN:C	1:A:2639:ARG:HD3	2.34	0.52
1:C:1710:HIS:ND1	1:C:1888:MET:HE1	2.24	0.52
1:E:1213:TYR:OH	1:E:1222:ASN:O	2.21	0.52
1:E:1477:LEU:HD13	1:E:2317:ARG:HH12	1.74	0.52
1:C:2347:LEU:HD21	1:E:2412:LEU:HD13	1.90	0.52
1:E:950:ILE:HG21	1:E:969:TRP:CD1	2.44	0.52
1:E:2477:SER:HB3	1:E:2479:ASP:HB2	1.92	0.52
1:E:2583:ILE:HD11	1:E:2616:ILE:HG12	1.91	0.52
1:A:950:ILE:HG21	1:A:969:TRP:CD1	2.44	0.52
1:A:1314:SER:O	1:A:1980:TYR:OH	2.27	0.52
1:A:2718:ARG:NH2	1:A:2726:GLU:OE2	2.40	0.52
1:C:1098:ILE:HD12	1:C:1102:HIS:HB3	1.92	0.52
1:C:1314:SER:O	1:C:1980:TYR:OH	2.27	0.52
1:E:1354:VAL:O	1:E:1358:THR:OG1	2.18	0.52
1:E:1710:HIS:ND1	1:E:1888:MET:HE1	2.24	0.52
1:A:1098:ILE:HD12	1:A:1102:HIS:HB3	1.92	0.52
1:A:2350:VAL:HG12	1:A:2351:MET:HE2	1.89	0.52
1:C:2560:ILE:O	1:C:2563:LYS:N	2.41	0.52
1:A:1091:SER:HB3	1:A:1251:VAL:HG22	1.92	0.52
1:A:2523:ASP:OD1	1:A:2523:ASP:N	2.41	0.52
1:A:2592:LYS:HG3	1:A:2594:PRO:HD3	1.92	0.52
1:A:2653:PHE:HB3	1:A:2654:ASN:ND2	2.25	0.52
1:C:1878:LEU:HG	1:C:1882:GLU:HB2	1.92	0.52
1:C:2444:THR:HG22	1:C:2450:PRO:HB3	1.90	0.52
1:C:2477:SER:HB3	1:C:2479:ASP:HB2	1.92	0.52
1:E:1162:ILE:HA	1:E:1165:ILE:HG12	1.92	0.52
1:E:1878:LEU:HG	1:E:1882:GLU:HB2	1.92	0.52
1:E:1934:HIS:CE1	1:E:1943:LEU:HB2	2.45	0.52
1:A:1888:MET:HG3	1:A:1889:PHE:N	2.25	0.52
1:A:1934:HIS:HE1	1:A:1943:LEU:HB2	1.74	0.52
1:E:2560:ILE:O	1:E:2563:LYS:N	2.41	0.52
1:A:1306:PHE:HB2	1:A:1324:CYS:SG	2.48	0.52
1:A:1477:LEU:HD13	1:A:2317:ARG:HH12	1.74	0.52
1:C:978:LEU:O	1:C:982:ALA:N	2.38	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1310:THR:HG23	1:C:1321:LEU:HD21	1.91	0.52
1:C:2464:MET:HG3	1:C:2491:LYS:HD2	1.92	0.52
1:E:1091:SER:HB3	1:E:1251:VAL:HG22	1.92	0.52
1:E:2530:VAL:HG11	1:E:2586:ILE:HD13	1.90	0.52
1:E:2653:PHE:HB3	1:E:2654:ASN:ND2	2.25	0.52
1:C:2530:VAL:HG11	1:C:2586:ILE:HD13	1.90	0.52
1:C:2585:LYS:NZ	1:C:2609:ASN:O	2.23	0.52
1:E:1888:MET:HG3	1:E:1889:PHE:N	2.25	0.52
1:C:950:ILE:HG21	1:C:969:TRP:NE1	2.25	0.52
1:C:1687:MET:CE	1:C:1889:PHE:HB2	2.40	0.52
1:C:2580:PRO:HA	1:C:2617:ILE:HB	1.91	0.52
1:C:2583:ILE:HD11	1:C:2616:ILE:HG12	1.91	0.52
1:E:1687:MET:CE	1:E:1889:PHE:HB2	2.40	0.52
1:E:1934:HIS:HE1	1:E:1943:LEU:HB2	1.73	0.52
1:E:2580:PRO:HA	1:E:2617:ILE:HB	1.91	0.52
1:A:1711:ILE:HG13	1:A:1886:LYS:HE2	1.91	0.51
1:A:2347:LEU:HD21	1:C:2412:LEU:HD13	1.92	0.51
1:A:2365:TRP:CZ3	1:A:2718:ARG:CZ	2.93	0.51
1:A:2444:THR:HG22	1:A:2450:PRO:HB3	1.90	0.51
1:C:1091:SER:HB3	1:C:1251:VAL:HG22	1.92	0.51
1:C:2477:SER:OG	1:C:2478:ARG:N	2.42	0.51
1:C:2464:MET:HE1	1:C:2468:SER:OG	2.11	0.51
1:E:2638:ASN:C	1:E:2639:ARG:HD3	2.34	0.51
1:C:2558:LYS:CD	1:C:2559:ASN:H	2.22	0.51
1:A:2464:MET:HE1	1:A:2468:SER:OG	2.11	0.51
1:A:2583:ILE:HD11	1:A:2616:ILE:HG12	1.92	0.51
1:C:1711:ILE:HG13	1:C:1886:LYS:HE2	1.91	0.51
1:C:2523:ASP:OD1	1:C:2523:ASP:N	2.41	0.51
1:E:1098:ILE:HD12	1:E:1102:HIS:HB3	1.92	0.51
1:E:2365:TRP:CZ3	1:E:2718:ARG:CZ	2.93	0.51
1:A:1310:THR:HG23	1:A:1321:LEU:HD21	1.91	0.51
1:A:2684:PHE:O	1:A:2687:GLU:HB3	2.11	0.51
1:C:2630:TRP:HH2	1:E:2510:SER:HB3	1.71	0.51
1:E:950:ILE:HG21	1:E:969:TRP:NE1	2.25	0.51
1:E:1711:ILE:HG23	1:E:1716:ARG:HH22	1.75	0.51
1:A:1878:LEU:HG	1:A:1882:GLU:HB2	1.92	0.51
1:A:1934:HIS:CE1	1:A:1943:LEU:HB2	2.45	0.51
1:A:2404:MET:HE1	2:F:238:GLU:HB3	1.93	0.51
1:C:2003:PRO:N	1:C:2004:PRO:HD2	2.26	0.51
1:C:2621:ASP:OD1	1:C:2623:THR:N	2.41	0.51
1:E:2482:ALA:O	1:E:2485:PHE:HB3	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2479:ASP:O	1:A:2480:THR:C	2.54	0.51
1:C:1948:LEU:HD23	1:C:1967:MET:HE2	1.93	0.51
1:E:935:TRP:HE3	1:E:1077:GLN:HG3	1.75	0.51
1:E:2003:PRO:N	1:E:2004:PRO:HD2	2.26	0.51
1:A:935:TRP:HE3	1:A:1077:GLN:HG3	1.74	0.51
1:A:950:ILE:HG21	1:A:969:TRP:NE1	2.25	0.51
1:A:1162:ILE:HA	1:A:1165:ILE:HG12	1.92	0.51
1:A:1192:CYS:SG	1:A:1193:ARG:N	2.82	0.51
1:A:1498:MET:HE2	1:A:1498:MET:N	2.26	0.51
1:A:2440:SER:OG	1:A:2535:SER:HB3	2.11	0.51
1:C:991:CYS:O	1:C:995:VAL:HG13	2.11	0.51
1:C:1477:LEU:HD13	1:C:2317:ARG:HH12	1.74	0.51
1:C:1934:HIS:CE1	1:C:1943:LEU:HB2	2.45	0.51
1:C:2365:TRP:CZ3	1:C:2718:ARG:CZ	2.93	0.51
1:E:1138:ARG:HA	1:E:1225:PHE:CZ	2.46	0.51
1:E:2592:LYS:HG3	1:E:2594:PRO:HD3	1.92	0.51
1:A:1948:LEU:HD23	1:A:1967:MET:HE2	1.93	0.51
1:A:2464:MET:O	1:A:2491:LYS:NZ	2.38	0.51
1:A:2627:ASN:OD1	1:A:2628:SER:N	2.40	0.51
1:C:935:TRP:HE3	1:C:1077:GLN:HG3	1.75	0.51
1:C:1008:GLU:HG3	1:C:1009:ASN:H	1.76	0.51
1:C:2653:PHE:HB3	1:C:2654:ASN:ND2	2.25	0.51
1:C:2684:PHE:O	1:C:2687:GLU:HB3	2.11	0.51
1:E:978:LEU:O	1:E:982:ALA:N	2.38	0.51
1:E:1314:SER:O	1:E:1980:TYR:OH	2.27	0.51
1:A:991:CYS:O	1:A:995:VAL:HG13	2.11	0.51
1:A:2477:SER:HB3	1:A:2479:ASP:HB2	1.92	0.51
1:C:1138:ARG:HA	1:C:1225:PHE:CZ	2.46	0.51
1:C:1708:GLN:O	1:C:1711:ILE:HG22	2.11	0.51
1:C:2592:LYS:HG3	1:C:2594:PRO:HD3	1.92	0.51
1:E:2684:PHE:O	1:E:2687:GLU:HB3	2.11	0.51
1:A:1285:TYR:HA	1:A:1288:MET:HG2	1.93	0.50
1:A:1711:ILE:HG23	1:A:1716:ARG:HH22	1.75	0.50
1:A:2003:PRO:N	1:A:2004:PRO:HD2	2.26	0.50
1:A:2486:LEU:O	1:A:2488:ASN:N	2.45	0.50
1:E:1008:GLU:HG3	1:E:1009:ASN:H	1.76	0.50
1:E:1708:GLN:O	1:E:1711:ILE:HG22	2.11	0.50
1:C:1498:MET:HE2	1:C:1498:MET:N	2.26	0.50
1:C:1674:HIS:CD2	1:C:1920:ALA:HA	2.46	0.50
1:C:1888:MET:HG3	1:C:1889:PHE:N	2.25	0.50
1:C:2473:ILE:HG12	1:C:2483:MET:HE2	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:991:CYS:O	1:E:995:VAL:HG13	2.11	0.50
1:E:1100:ARG:O	1:E:1103:LEU:HB3	2.11	0.50
1:E:2473:ILE:HG12	1:E:2483:MET:HE2	1.93	0.50
1:A:1138:ARG:HA	1:A:1225:PHE:CZ	2.46	0.50
1:C:1162:ILE:HA	1:C:1165:ILE:HG12	1.92	0.50
1:C:2592:LYS:O	1:C:2599:SER:OG	2.21	0.50
1:E:1708:GLN:HG2	1:E:1712:MET:HE1	1.94	0.50
1:E:2627:ASN:OD1	1:E:2628:SER:N	2.40	0.50
1:A:1708:GLN:HG2	1:A:1712:MET:HE1	1.93	0.50
1:A:2482:ALA:O	1:A:2485:PHE:HB3	2.11	0.50
1:E:944:LYS:HZ1	1:E:1092:ARG:HH12	1.59	0.50
1:E:1216:ASP:OD1	1:E:1217:PHE:N	2.44	0.50
1:E:2464:MET:HG3	1:E:2491:LYS:HD2	1.92	0.50
1:E:2486:LEU:O	1:E:2488:ASN:N	2.45	0.50
1:A:2515:GLN:HA	1:A:2518:ILE:HD12	1.94	0.50
1:C:1711:ILE:HG23	1:C:1716:ARG:HH22	1.75	0.50
1:C:2440:SER:OG	1:C:2535:SER:HB3	2.11	0.50
1:C:2482:ALA:O	1:C:2485:PHE:HB3	2.11	0.50
1:A:1076:HIS:HA	1:A:1079:TYR:CE1	2.47	0.50
1:A:1216:ASP:OD1	1:A:1217:PHE:N	2.44	0.50
1:C:1100:ARG:O	1:C:1103:LEU:HB3	2.11	0.50
1:C:2486:LEU:O	1:C:2488:ASN:N	2.45	0.50
1:C:2495:THR:O	1:C:2651:VAL:HA	2.12	0.50
1:C:2592:LYS:N	1:C:2600:LYS:O	2.34	0.50
1:E:2479:ASP:O	1:E:2480:THR:C	2.54	0.50
1:E:2523:ASP:OD1	1:E:2523:ASP:N	2.41	0.50
1:A:1008:GLU:HG3	1:A:1009:ASN:H	1.76	0.50
1:A:1674:HIS:CD2	1:A:1920:ALA:HA	2.47	0.50
1:A:1687:MET:CE	1:A:1889:PHE:HB2	2.40	0.50
1:A:1700:ARG:HA	1:A:1703:ILE:HD12	1.93	0.50
1:A:2464:MET:HG3	1:A:2491:LYS:HD2	1.92	0.50
1:C:1216:ASP:OD1	1:C:1217:PHE:N	2.44	0.50
1:C:1465:ALA:O	1:C:1468:GLU:HG3	2.12	0.50
1:C:1700:ARG:HA	1:C:1703:ILE:HD12	1.93	0.50
1:C:1941:ILE:O	1:C:1971:TYR:OH	2.21	0.50
1:C:2353:TRP:CE3	1:C:2362:LEU:HD12	2.47	0.50
1:E:1465:ALA:O	1:E:1468:GLU:HG3	2.12	0.50
1:E:1948:LEU:HD23	1:E:1967:MET:HE2	1.93	0.50
1:A:1100:ARG:O	1:A:1103:LEU:HB3	2.11	0.50
1:A:1135:ILE:HG23	1:A:1143:ALA:HB1	1.94	0.50
1:A:2585:LYS:HB3	1:A:2611:PHE:HD2	1.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1708:GLN:HG2	1:C:1712:MET:HE1	1.94	0.50
1:C:2585:LYS:HB3	1:C:2611:PHE:HD2	1.77	0.50
1:C:2627:ASN:OD1	1:C:2628:SER:N	2.40	0.50
1:E:920:PHE:HA	1:E:923:PHE:CE1	2.47	0.50
1:E:2365:TRP:CD1	1:E:2365:TRP:C	2.90	0.50
1:E:2440:SER:OG	1:E:2535:SER:HB3	2.11	0.50
1:E:2580:PRO:HB3	1:E:2617:ILE:HB	1.94	0.50
1:A:1983:GLN:HB2	1:A:2011:LYS:HB2	1.94	0.50
1:A:2353:TRP:CE3	1:A:2362:LEU:HD12	2.47	0.50
1:A:2473:ILE:HG12	1:A:2483:MET:HE2	1.93	0.50
1:C:1192:CYS:SG	1:C:1193:ARG:N	2.82	0.50
1:E:2495:THR:O	1:E:2651:VAL:HA	2.12	0.50
1:A:2563:LYS:O	1:A:2567:LYS:HG3	2.12	0.49
1:E:1498:MET:HE2	1:E:1498:MET:N	2.26	0.49
1:A:2365:TRP:CD1	1:A:2365:TRP:C	2.90	0.49
1:E:1124:LEU:HD22	1:E:1162:ILE:HD12	1.94	0.49
1:E:2353:TRP:CE3	1:E:2362:LEU:HD12	2.47	0.49
1:E:2485:PHE:CD2	1:E:2486:LEU:HD22	2.41	0.49
1:A:1708:GLN:O	1:A:1711:ILE:HG22	2.11	0.49
1:A:2463:VAL:O	1:A:2464:MET:HB3	2.12	0.49
1:A:2495:THR:O	1:A:2651:VAL:HA	2.12	0.49
1:A:2560:ILE:O	1:A:2563:LYS:N	2.41	0.49
1:A:2585:LYS:NZ	1:A:2609:ASN:O	2.23	0.49
1:A:2722:GLU:C	1:A:2723:LEU:HD12	2.38	0.49
1:C:920:PHE:HA	1:C:923:PHE:CE1	2.47	0.49
1:E:1674:HIS:CD2	1:E:1920:ALA:HA	2.46	0.49
1:A:1124:LEU:HD22	1:A:1162:ILE:HD12	1.94	0.49
1:C:1210:LYS:NZ	1:C:1393:GLY:O	2.45	0.49
1:E:1076:HIS:HA	1:E:1079:TYR:CE1	2.47	0.49
1:C:1480:GLN:O	1:C:1484:ILE:HD12	2.13	0.49
1:C:1710:HIS:CG	1:C:1888:MET:HE1	2.48	0.49
1:C:2365:TRP:CD1	1:C:2365:TRP:C	2.90	0.49
1:C:2414:CYS:HA	1:C:2418:PHE:CD2	2.48	0.49
1:C:2479:ASP:O	1:C:2480:THR:C	2.54	0.49
1:C:2722:GLU:C	1:C:2723:LEU:HD12	2.38	0.49
1:E:1207:ASN:ND2	1:E:1394:TYR:OH	2.46	0.49
1:E:1210:LYS:NZ	1:E:1393:GLY:O	2.46	0.49
1:E:2463:VAL:O	1:E:2464:MET:HB3	2.12	0.49
1:A:1210:LYS:NZ	1:A:1393:GLY:O	2.46	0.49
1:A:1465:ALA:O	1:A:1468:GLU:HG3	2.12	0.49
1:A:2558:LYS:CD	1:A:2559:ASN:H	2.22	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1285:TYR:HA	1:E:1288:MET:HG2	1.93	0.49
1:E:1360:LYS:HE3	1:E:1412:ILE:HG12	1.94	0.49
1:E:2464:MET:HE1	1:E:2468:SER:OG	2.11	0.49
1:A:1307:ILE:O	1:A:1311:THR:OG1	2.28	0.49
1:A:1360:LYS:HE3	1:A:1412:ILE:HG12	1.94	0.49
1:C:2580:PRO:HB3	1:C:2617:ILE:HB	1.94	0.49
1:E:1700:ARG:HA	1:E:1703:ILE:HD12	1.93	0.49
1:E:2585:LYS:HB3	1:E:2611:PHE:HD2	1.77	0.49
1:A:920:PHE:HA	1:A:923:PHE:CE1	2.47	0.49
1:A:2414:CYS:HA	1:A:2418:PHE:CD2	2.48	0.49
1:A:2580:PRO:HB3	1:A:2617:ILE:HB	1.94	0.49
1:C:1207:ASN:ND2	1:C:1394:TYR:OH	2.46	0.49
1:C:1285:TYR:HA	1:C:1288:MET:HG2	1.93	0.49
1:E:1149:TRP:HE3	1:E:1169:TYR:HD1	1.61	0.49
1:E:1263:MET:HE1	1:E:1343:LEU:HB2	1.95	0.49
1:A:924:LEU:HA	1:A:927:PHE:CE1	2.48	0.49
1:A:1710:HIS:CG	1:A:1888:MET:HE1	2.48	0.49
1:C:979:ARG:HA	1:C:982:ALA:HB3	1.95	0.49
1:C:1135:ILE:HG23	1:C:1143:ALA:HB1	1.94	0.49
1:C:1149:TRP:HE3	1:C:1169:TYR:HD1	1.61	0.49
1:C:2563:LYS:O	1:C:2567:LYS:HG3	2.12	0.49
1:E:1320:TYR:OH	1:E:1353:ASN:OD1	2.31	0.49
1:E:2179:ILE:HG22	1:E:2180:HIS:HD2	1.77	0.49
1:A:1941:ILE:O	1:A:1971:TYR:OH	2.21	0.49
1:E:1135:ILE:HG23	1:E:1143:ALA:HB1	1.94	0.49
1:E:2563:LYS:O	1:E:2567:LYS:HG3	2.12	0.49
1:E:2722:GLU:C	1:E:2723:LEU:HD12	2.38	0.49
1:E:1480:GLN:O	1:E:1484:ILE:HD12	2.13	0.48
1:E:1699:THR:O	1:E:1703:ILE:N	2.37	0.48
1:A:1320:TYR:OH	1:A:1353:ASN:OD1	2.31	0.48
1:A:1454:LEU:HD21	1:A:1880:ALA:HB2	1.95	0.48
1:C:1983:GLN:HB2	1:C:2011:LYS:HB2	1.94	0.48
1:E:2515:GLN:HA	1:E:2518:ILE:HD12	1.94	0.48
1:C:1076:HIS:HA	1:C:1079:TYR:CE1	2.47	0.48
1:C:1360:LYS:HE3	1:C:1412:ILE:HG12	1.94	0.48
1:C:1454:LEU:HD21	1:C:1880:ALA:HB2	1.95	0.48
1:C:2438:ASP:OD1	1:C:2456:ALA:HB3	2.13	0.48
1:A:1207:ASN:ND2	1:A:1394:TYR:OH	2.46	0.48
1:A:1263:MET:HE1	1:A:1343:LEU:HB2	1.95	0.48
1:A:2438:ASP:OD1	1:A:2456:ALA:HB3	2.13	0.48
1:A:2536:ILE:N	1:A:2549:ALA:O	2.46	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:924:LEU:HA	1:E:927:PHE:CE1	2.48	0.48
1:E:2438:ASP:OD1	1:E:2456:ALA:HB3	2.13	0.48
1:C:1270:PHE:CZ	1:C:1275:PRO:HD3	2.49	0.48
1:C:2515:GLN:HA	1:C:2518:ILE:HD12	1.94	0.48
1:E:1710:HIS:CG	1:E:1888:MET:HE1	2.48	0.48
1:E:2414:CYS:HA	1:E:2418:PHE:CD2	2.48	0.48
1:E:2536:ILE:HG22	1:E:2538:ARG:HG3	1.96	0.48
1:E:2536:ILE:N	1:E:2549:ALA:O	2.46	0.48
1:A:1181:TYR:HD1	1:A:1226:LEU:HD13	1.78	0.48
1:C:1934:HIS:CE1	1:C:1943:LEU:HD13	2.49	0.48
1:C:2463:VAL:O	1:C:2464:MET:HB3	2.12	0.48
1:C:2536:ILE:N	1:C:2549:ALA:O	2.46	0.48
1:A:1006:LYS:HG3	1:A:1008:GLU:HG2	1.95	0.48
1:A:1934:HIS:CE1	1:A:1943:LEU:HD13	2.49	0.48
1:A:2568:MET:SD	1:A:2618:LEU:HB2	2.54	0.48
1:C:1124:LEU:HD22	1:C:1162:ILE:HD12	1.94	0.48
1:C:1553:LEU:O	1:C:1555:GLU:N	2.46	0.48
1:C:2568:MET:SD	1:C:2618:LEU:HB2	2.54	0.48
1:E:1983:GLN:HB2	1:E:2011:LYS:HB2	1.94	0.48
1:A:1270:PHE:CZ	1:A:1275:PRO:HD3	2.49	0.48
1:C:2317:ARG:HG3	1:C:2719:GLU:OE1	2.14	0.48
1:E:1553:LEU:O	1:E:1555:GLU:N	2.46	0.48
1:E:2476:PHE:CD1	1:E:2605:LEU:HD22	2.49	0.48
1:A:2476:PHE:CD1	1:A:2605:LEU:HD22	2.49	0.48
1:C:924:LEU:HA	1:C:927:PHE:CE1	2.48	0.48
1:C:2485:PHE:CD2	1:C:2486:LEU:HD22	2.41	0.48
1:A:979:ARG:HA	1:A:982:ALA:HB3	1.95	0.48
1:A:1042:THR:O	1:A:1046:GLY:N	2.43	0.48
1:C:1263:MET:HE1	1:C:1343:LEU:HB2	1.95	0.48
1:C:1313:ILE:HG23	1:C:1980:TYR:CE1	2.49	0.48
1:C:1669:SER:HA	1:C:1672:ARG:HE	1.79	0.48
1:E:1454:LEU:HD21	1:E:1880:ALA:HB2	1.95	0.48
1:A:1480:GLN:O	1:A:1484:ILE:HD12	2.13	0.47
1:A:1543:ALA:HB2	1:C:2390:GLN:O	2.14	0.47
1:A:1684:GLU:HA	1:A:1707:TYR:OH	2.14	0.47
1:A:2536:ILE:HG22	1:A:2538:ARG:HG3	1.96	0.47
1:A:2621:ASP:C	1:A:2623:THR:N	2.72	0.47
1:C:1014:CYS:HB3	1:C:1035:TYR:HA	1.96	0.47
1:C:1320:TYR:OH	1:C:1353:ASN:OD1	2.31	0.47
1:C:1006:LYS:HG3	1:C:1008:GLU:HG2	1.95	0.47
1:C:1683:ILE:HA	1:C:1894:LEU:HD21	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2476:PHE:CD1	1:C:2605:LEU:HD22	2.49	0.47
1:C:2561:THR:O	1:C:2564:ASN:N	2.47	0.47
1:E:1934:HIS:CE1	1:E:1943:LEU:HD13	2.49	0.47
1:E:2568:MET:SD	1:E:2618:LEU:HB2	2.54	0.47
1:A:1149:TRP:HE3	1:A:1169:TYR:HD1	1.61	0.47
1:A:2561:THR:O	1:A:2564:ASN:N	2.47	0.47
1:C:936:TRP:NE1	1:C:1093:THR:O	2.46	0.47
1:E:1006:LYS:HG3	1:E:1008:GLU:HG2	1.95	0.47
1:A:2616:ILE:HG22	1:A:2633:LEU:HG	1.97	0.47
1:A:2620:ARG:HA	1:A:2628:SER:O	2.15	0.47
1:C:960:LEU:HB3	1:C:1045:VAL:O	2.14	0.47
1:C:2616:ILE:HG22	1:C:2633:LEU:HG	1.97	0.47
1:E:936:TRP:NE1	1:E:1093:THR:O	2.46	0.47
1:E:1313:ILE:HG23	1:E:1980:TYR:CE1	2.49	0.47
1:E:2317:ARG:HG3	1:E:2719:GLU:OE1	2.14	0.47
1:A:1141:PHE:O	1:A:1144:MET:HB2	2.15	0.47
1:A:2037:GLY:O	1:A:2038:LEU:HG	2.14	0.47
1:A:2463:VAL:HG12	1:A:2464:MET:HG2	1.96	0.47
1:A:2472:PHE:HZ	1:A:2590:TYR:HE1	1.62	0.47
1:A:2509:ILE:HG12	1:A:2510:SER:H	1.79	0.47
1:C:1181:TYR:HD1	1:C:1226:LEU:HD13	1.78	0.47
1:E:979:ARG:HA	1:E:982:ALA:HB3	1.95	0.47
1:E:1696:ASN:O	1:E:1698:PRO:HD3	2.15	0.47
1:E:2495:THR:HB	1:E:2652:VAL:HB	1.97	0.47
1:E:2561:THR:O	1:E:2564:ASN:N	2.47	0.47
1:C:1368:CYS:C	1:C:1394:TYR:HH	2.19	0.47
1:C:2584:GLU:OE1	1:C:2584:GLU:N	2.48	0.47
1:E:1270:PHE:CZ	1:E:1275:PRO:HD3	2.49	0.47
1:A:1696:ASN:O	1:A:1698:PRO:HD3	2.15	0.47
1:A:2486:LEU:HA	1:A:2489:TYR:HE2	1.80	0.47
1:C:1371:ILE:O	1:C:1375:VAL:HG23	2.15	0.47
1:C:1696:ASN:O	1:C:1698:PRO:HD3	2.15	0.47
1:C:2353:TRP:CZ3	1:C:2362:LEU:HD12	2.50	0.47
1:C:2536:ILE:HG22	1:C:2538:ARG:HG3	1.96	0.47
1:C:2698:PHE:CE2	1:E:2694:HIS:HA	2.50	0.47
1:E:2586:ILE:C	1:E:2611:PHE:HE2	2.23	0.47
1:E:2620:ARG:HA	1:E:2628:SER:O	2.15	0.47
1:A:950:ILE:HD13	1:A:969:TRP:NE1	2.30	0.47
1:A:960:LEU:HB3	1:A:1045:VAL:O	2.14	0.47
1:A:1371:ILE:O	1:A:1375:VAL:HG23	2.15	0.47
1:A:1669:SER:HA	1:A:1672:ARG:HE	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2179:ILE:HG22	1:A:2180:HIS:HD2	1.78	0.47
1:C:1706:TYR:O	1:C:1710:HIS:ND1	2.48	0.47
1:C:2441:VAL:HA	1:C:2534:TRP:HA	1.97	0.47
1:C:2586:ILE:C	1:C:2611:PHE:HE2	2.23	0.47
1:E:1181:TYR:HD1	1:E:1226:LEU:HD13	1.78	0.47
1:E:1192:CYS:SG	1:E:1193:ARG:N	2.82	0.47
1:E:1305:ILE:HA	1:E:1418:CYS:SG	2.55	0.47
1:E:1684:GLU:HA	1:E:1707:TYR:OH	2.14	0.47
1:E:2353:TRP:CZ3	1:E:2362:LEU:HD12	2.50	0.47
1:A:1305:ILE:HA	1:A:1418:CYS:SG	2.55	0.47
1:A:1706:TYR:O	1:A:1710:HIS:ND1	2.48	0.47
1:C:1699:THR:O	1:C:1703:ILE:HG13	2.15	0.47
1:E:1141:PHE:O	1:E:1144:MET:HB2	2.15	0.47
1:E:1699:THR:O	1:E:1703:ILE:HG13	2.15	0.47
1:E:2441:VAL:HA	1:E:2534:TRP:HA	1.97	0.47
1:A:2404:MET:HE1	2:F:238:GLU:HG2	1.97	0.47
1:C:2495:THR:HB	1:C:2652:VAL:HB	1.97	0.47
1:E:1014:CYS:HB3	1:E:1035:TYR:HA	1.96	0.47
1:E:1706:TYR:O	1:E:1710:HIS:ND1	2.48	0.47
1:E:2037:GLY:O	1:E:2038:LEU:HG	2.14	0.47
1:E:2623:THR:HG23	1:E:2626:TYR:O	2.15	0.47
1:A:2317:ARG:HG3	1:A:2719:GLU:OE1	2.14	0.46
1:A:2394:GLN:HG3	1:A:2395:LYS:O	2.15	0.46
1:A:2510:SER:HB2	1:A:2513:SER:OG	2.16	0.46
1:C:1684:GLU:HA	1:C:1707:TYR:OH	2.14	0.46
1:C:2486:LEU:HA	1:C:2489:TYR:HE2	1.80	0.46
1:E:2472:PHE:HZ	1:E:2590:TYR:HE1	1.62	0.46
1:A:1043:GLU:HA	1:A:1047:LEU:C	2.40	0.46
1:A:2584:GLU:OE1	1:A:2584:GLU:N	2.48	0.46
1:A:2586:ILE:C	1:A:2611:PHE:HE2	2.23	0.46
1:C:1043:GLU:HA	1:C:1047:LEU:C	2.40	0.46
1:C:1543:ALA:HB2	1:E:2390:GLN:O	2.14	0.46
1:C:2394:GLN:HG3	1:C:2395:LYS:O	2.15	0.46
1:C:2509:ILE:HG12	1:C:2510:SER:H	1.79	0.46
1:C:2510:SER:HB2	1:C:2513:SER:OG	2.16	0.46
1:C:2730:TYR:CZ	1:C:2734:ILE:HG13	2.51	0.46
2:D:246:SER:O	2:D:246:SER:OG	2.23	0.46
1:E:2509:ILE:HG12	1:E:2510:SER:H	1.79	0.46
1:A:1014:CYS:HB3	1:A:1035:TYR:HA	1.96	0.46
1:C:1305:ILE:HA	1:C:1418:CYS:SG	2.55	0.46
1:C:1682:ARG:HH12	1:C:1901:TYR:HE2	1.64	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2466:GLN:N	1:C:2467:GLN:OE1	2.48	0.46
1:C:2620:ARG:HA	1:C:2628:SER:O	2.14	0.46
1:A:2390:GLN:O	1:E:1543:ALA:HB2	2.14	0.46
1:A:2466:GLN:N	1:A:2467:GLN:OE1	2.49	0.46
1:A:2639:ARG:HG2	1:A:2649:GLU:HG3	1.98	0.46
1:A:2730:TYR:CZ	1:A:2734:ILE:HG13	2.50	0.46
1:C:2463:VAL:HG12	1:C:2464:MET:HG2	1.96	0.46
1:E:960:LEU:HB3	1:E:1045:VAL:O	2.14	0.46
1:E:1682:ARG:HH12	1:E:1901:TYR:HE2	1.64	0.46
1:E:2274:MET:HB2	1:E:2274:MET:HE3	1.60	0.46
1:E:2616:ILE:HG22	1:E:2633:LEU:HG	1.97	0.46
1:A:1682:ARG:HH12	1:A:1901:TYR:HE2	1.64	0.46
1:A:2274:MET:HB2	1:A:2274:MET:HE3	1.60	0.46
1:A:2495:THR:HB	1:A:2652:VAL:HB	1.97	0.46
1:A:2623:THR:HG23	1:A:2626:TYR:O	2.16	0.46
1:C:2037:GLY:O	1:C:2038:LEU:HG	2.14	0.46
1:C:2179:ILE:HG22	1:C:2180:HIS:HD2	1.78	0.46
1:E:1453:GLU:HB2	1:E:1700:ARG:NH2	2.31	0.46
1:E:1669:SER:HA	1:E:1672:ARG:HE	1.79	0.46
1:E:2486:LEU:HA	1:E:2489:TYR:HE2	1.80	0.46
1:A:1699:THR:O	1:A:1703:ILE:HG13	2.15	0.46
1:A:2353:TRP:CZ3	1:A:2362:LEU:HD12	2.50	0.46
1:A:2391:PRO:HB2	1:A:2394:GLN:HB2	1.98	0.46
1:A:2413:ILE:HD12	1:A:2413:ILE:HA	1.84	0.46
1:C:950:ILE:HD13	1:C:969:TRP:NE1	2.30	0.46
1:C:2415:ILE:HD13	1:C:2681:ILE:HG21	1.98	0.46
1:C:2589:TYR:O	1:C:2590:TYR:CG	2.69	0.46
1:C:2623:THR:HG23	1:C:2626:TYR:O	2.15	0.46
1:E:1683:ILE:HA	1:E:1894:LEU:HD21	1.97	0.46
1:E:2510:SER:HB2	1:E:2513:SER:OG	2.15	0.46
1:A:2441:VAL:HA	1:A:2534:TRP:HA	1.97	0.46
1:C:1453:GLU:HB2	1:C:1700:ARG:NH2	2.31	0.46
1:E:2410:VAL:HA	1:E:2413:ILE:HG22	1.98	0.46
1:E:2466:GLN:N	1:E:2467:GLN:OE1	2.49	0.46
1:E:2472:PHE:HA	1:E:2475:ALA:CB	2.46	0.46
1:A:1313:ILE:HG23	1:A:1980:TYR:CE1	2.49	0.46
1:A:1321:LEU:HD23	1:A:1321:LEU:HA	1.78	0.46
1:A:1683:ILE:HA	1:A:1894:LEU:HD21	1.96	0.46
1:A:2410:VAL:HA	1:A:2413:ILE:HG22	1.98	0.46
1:A:2415:ILE:HD13	1:A:2681:ILE:HG21	1.98	0.46
1:C:2621:ASP:C	1:C:2623:THR:N	2.72	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:950:ILE:HD13	1:E:969:TRP:NE1	2.30	0.46
1:E:1371:ILE:O	1:E:1375:VAL:HG23	2.14	0.46
1:E:2394:GLN:HG3	1:E:2395:LYS:O	2.15	0.46
1:E:2584:GLU:N	1:E:2584:GLU:OE1	2.48	0.46
1:E:2589:TYR:O	1:E:2590:TYR:CG	2.69	0.46
1:A:1453:GLU:HB2	1:A:1700:ARG:NH2	2.31	0.46
1:C:2410:VAL:HA	1:C:2413:ILE:HG22	1.98	0.46
1:C:2618:LEU:HD23	1:C:2620:ARG:NH2	2.31	0.46
1:E:2415:ILE:HD13	1:E:2681:ILE:HG21	1.98	0.46
1:A:2016:VAL:O	1:A:2020:LEU:N	2.45	0.46
1:A:2434:ASN:ND2	1:A:2538:ARG:HD2	2.31	0.46
1:C:994:ILE:HB	1:C:1061:LEU:HD21	1.98	0.46
1:C:1141:PHE:O	1:C:1144:MET:HB2	2.15	0.46
1:C:2179:ILE:C	1:C:2181:PRO:HD3	2.41	0.46
1:C:2274:MET:HB2	1:C:2274:MET:HE3	1.60	0.46
1:E:2179:ILE:C	1:E:2181:PRO:HD3	2.41	0.46
1:A:936:TRP:NE1	1:A:1093:THR:O	2.46	0.45
1:A:2448:TYR:CE2	1:A:2517:MET:HG2	2.51	0.45
1:A:2472:PHE:HA	1:A:2475:ALA:CB	2.46	0.45
1:A:2485:PHE:CD2	1:A:2486:LEU:HD22	2.40	0.45
1:A:2589:TYR:O	1:A:2590:TYR:CG	2.69	0.45
1:C:2434:ASN:ND2	1:C:2538:ARG:HD2	2.31	0.45
1:C:2461:LEU:HD23	1:C:2497:ALA:HA	1.98	0.45
1:E:1043:GLU:HA	1:E:1047:LEU:C	2.40	0.45
1:E:2553:LEU:HD23	1:E:2553:LEU:HA	1.78	0.45
1:E:2743:MET:HE2	1:E:2743:MET:HB3	1.55	0.45
1:A:2365:TRP:HZ3	1:A:2718:ARG:CZ	2.30	0.45
1:A:2694:HIS:HA	1:E:2698:PHE:CE2	2.51	0.45
1:C:1923:GLU:HG2	1:C:1927:TYR:HE2	1.80	0.45
1:E:2448:TYR:CE2	1:E:2517:MET:HG2	2.51	0.45
1:A:2202:ILE:HA	1:A:2205:VAL:HG12	1.98	0.45
1:A:2743:MET:HE2	1:A:2743:MET:HB3	1.55	0.45
1:C:2202:ILE:HA	1:C:2205:VAL:HG12	1.98	0.45
1:C:2472:PHE:HA	1:C:2475:ALA:CB	2.46	0.45
1:C:2472:PHE:HZ	1:C:2590:TYR:HE1	1.62	0.45
1:C:2639:ARG:HG2	1:C:2649:GLU:HG3	1.98	0.45
1:E:2608:GLU:CA	1:E:2611:PHE:HB2	2.46	0.45
1:E:2621:ASP:C	1:E:2623:THR:N	2.72	0.45
1:E:2730:TYR:CZ	1:E:2734:ILE:HG13	2.51	0.45
1:A:2583:ILE:HD13	1:A:2616:ILE:HG23	1.98	0.45
1:A:2592:LYS:N	1:A:2600:LYS:O	2.34	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:989:TRP:O	1:C:993:ILE:HG12	2.17	0.45
1:C:1982:PHE:HB2	1:C:2009:VAL:CG2	2.47	0.45
1:C:2430:ALA:N	1:C:2659:PRO:HD2	2.32	0.45
1:E:1982:PHE:HB2	1:E:2009:VAL:CG2	2.47	0.45
1:E:2202:ILE:HA	1:E:2205:VAL:HG12	1.98	0.45
1:E:2396:LYS:HB2	1:E:2401:LYS:HD3	1.99	0.45
1:E:2618:LEU:HD23	1:E:2620:ARG:NH2	2.31	0.45
1:C:2396:LYS:HB2	1:C:2401:LYS:HD3	1.99	0.45
1:E:1923:GLU:HG2	1:E:1927:TYR:HE2	1.80	0.45
1:E:2440:SER:HA	1:E:2455:SER:HA	1.98	0.45
1:E:2463:VAL:HG12	1:E:2464:MET:HG2	1.97	0.45
1:E:2464:MET:O	1:E:2491:LYS:NZ	2.38	0.45
1:E:2472:PHE:CE2	1:E:2476:PHE:HE2	2.35	0.45
1:A:1280:ILE:CD1	1:A:1440:ASP:HB2	2.47	0.45
1:A:2467:GLN:C	1:A:2471:LYS:HZ2	2.25	0.45
1:A:2472:PHE:HA	1:A:2475:ALA:HB3	1.99	0.45
1:A:2512:PRO:HG2	1:E:2630:TRP:CH2	2.51	0.45
1:C:2334:LEU:HD23	1:C:2334:LEU:HA	1.74	0.45
1:E:2430:ALA:N	1:E:2659:PRO:HD2	2.32	0.45
2:F:230:SER:O	2:F:233:LEU:HB3	2.17	0.45
1:A:1554:PHE:CE1	1:A:2732:LYS:HG3	2.46	0.45
2:B:230:SER:O	2:B:233:LEU:HB3	2.17	0.45
1:C:2305:LEU:HA	1:C:2305:LEU:HD23	1.68	0.45
1:C:2448:TYR:CE2	1:C:2517:MET:HG2	2.51	0.45
1:C:2472:PHE:HA	1:C:2475:ALA:HB3	1.99	0.45
1:C:2583:ILE:HD13	1:C:2616:ILE:HG23	1.98	0.45
1:E:2167:ILE:C	1:E:2170:PRO:HD2	2.42	0.45
1:E:2558:LYS:CD	1:E:2559:ASN:H	2.22	0.45
1:A:2340:LEU:HA	1:A:2340:LEU:HD23	1.74	0.45
1:C:1156:ARG:HB2	1:C:1162:ILE:HG12	1.99	0.45
1:E:2391:PRO:HB2	1:E:2394:GLN:HB2	1.98	0.45
1:E:2434:ASN:ND2	1:E:2538:ARG:HD2	2.31	0.45
1:E:2583:ILE:HD13	1:E:2616:ILE:HG23	1.98	0.45
1:E:2639:ARG:HG2	1:E:2649:GLU:HG3	1.98	0.45
1:A:2167:ILE:C	1:A:2170:PRO:HD2	2.42	0.45
1:A:2299:LYS:HA	1:A:2299:LYS:HD3	1.73	0.45
1:A:2585:LYS:HB3	1:A:2611:PHE:CD2	2.52	0.45
1:A:2708:LEU:HD12	1:A:2708:LEU:HA	1.68	0.45
2:D:238:GLU:CG	1:E:2404:MET:HE1	2.44	0.45
1:E:931:GLN:O	1:E:935:TRP:HD1	2.00	0.45
1:E:989:TRP:O	1:E:993:ILE:HG12	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1140:ASP:OD2	1:E:1142:TYR:HB2	2.17	0.45
1:E:1548:SER:O	1:E:1548:SER:OG	2.29	0.45
1:E:2472:PHE:HA	1:E:2475:ALA:HB3	1.99	0.45
1:E:2557:LEU:HD13	1:E:2562:ARG:NE	2.32	0.45
1:A:2179:ILE:C	1:A:2181:PRO:HD3	2.41	0.45
1:A:2396:LYS:HB2	1:A:2401:LYS:HD3	1.99	0.45
1:A:2440:SER:HA	1:A:2455:SER:HA	1.98	0.45
1:A:2461:LEU:HD23	1:A:2497:ALA:HA	1.98	0.45
1:C:1140:ASP:OD2	1:C:1142:TYR:HB2	2.17	0.45
1:C:1280:ILE:CD1	1:C:1440:ASP:HB2	2.47	0.45
1:C:2365:TRP:HZ3	1:C:2718:ARG:CZ	2.29	0.45
2:D:230:SER:O	2:D:233:LEU:HB3	2.17	0.45
1:A:2557:LEU:HD13	1:A:2562:ARG:NE	2.32	0.44
1:A:2630:TRP:CH2	1:C:2512:PRO:HG2	2.52	0.44
1:C:2167:ILE:C	1:C:2170:PRO:HD2	2.42	0.44
1:C:2378:LEU:HD23	1:C:2378:LEU:HA	1.82	0.44
1:C:2391:PRO:HB2	1:C:2394:GLN:HB2	1.98	0.44
1:C:2585:LYS:HB3	1:C:2611:PHE:CD2	2.52	0.44
1:E:2458:GLN:NE2	1:E:2459:SER:OG	2.51	0.44
1:A:994:ILE:HG12	1:A:1060:ASN:O	2.18	0.44
1:A:1542:HIS:HA	1:C:2392:ARG:HH21	1.82	0.44
1:A:2553:LEU:HB3	1:A:2555:PHE:CE2	2.52	0.44
1:A:2698:PHE:CE2	1:C:2694:HIS:HA	2.51	0.44
1:C:974:PRO:HB3	1:C:1170:CYS:HB3	1.99	0.44
1:C:1488:GLN:C	1:C:1492:LYS:HE2	2.42	0.44
1:C:1554:PHE:CE1	1:C:2732:LYS:HG3	2.46	0.44
1:E:1156:ARG:HB2	1:E:1162:ILE:HG12	1.99	0.44
1:E:1477:LEU:HD21	1:E:2720:THR:HG22	1.99	0.44
1:E:2397:LYS:O	1:E:2400:VAL:N	2.48	0.44
1:E:2461:LEU:HD23	1:E:2497:ALA:HA	1.98	0.44
1:A:1712:MET:HG3	1:A:1716:ARG:HH21	1.83	0.44
1:A:1982:PHE:HB2	1:A:2009:VAL:CG2	2.47	0.44
1:A:2560:ILE:HG23	1:A:2563:LYS:HE2	2.00	0.44
1:C:931:GLN:O	1:C:935:TRP:HD1	2.01	0.44
1:C:1149:TRP:CE3	1:C:1169:TYR:HD1	2.35	0.44
1:C:1545:MET:SD	1:E:2392:ARG:HD2	2.57	0.44
1:C:2511:PRO:O	1:C:2514:LYS:HB2	2.18	0.44
1:E:1394:TYR:CG	1:E:1395:GLN:N	2.85	0.44
1:E:2454:MET:CE	1:E:2499:LEU:HA	2.47	0.44
1:E:2621:ASP:OD1	1:E:2622:ASN:N	2.51	0.44
1:A:1140:ASP:OD2	1:A:1142:TYR:HB2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1284:SER:H	1:A:1287:ASP:HB3	1.83	0.44
1:A:2404:MET:HE1	2:F:238:GLU:CB	2.47	0.44
1:A:2618:LEU:HD23	1:A:2620:ARG:NH2	2.31	0.44
1:A:2703:ASN:OD1	1:A:2749:GLU:HA	2.18	0.44
1:C:994:ILE:HA	1:C:997:LYS:HD3	2.00	0.44
1:C:1548:SER:O	1:C:1548:SER:OG	2.29	0.44
1:C:1712:MET:HG3	1:C:1716:ARG:HH21	1.83	0.44
1:E:1280:ILE:CD1	1:E:1440:ASP:HB2	2.47	0.44
1:E:1488:GLN:C	1:E:1492:LYS:HE2	2.42	0.44
1:A:974:PRO:HB3	1:A:1170:CYS:HB3	1.99	0.44
1:A:994:ILE:HB	1:A:1061:LEU:HD21	1.99	0.44
1:A:1923:GLU:HG2	1:A:1927:TYR:HE2	1.80	0.44
1:A:2430:ALA:N	1:A:2659:PRO:HD2	2.32	0.44
1:C:1308:THR:HG23	1:C:1414:TRP:HB3	2.00	0.44
1:C:2454:MET:CE	1:C:2499:LEU:HA	2.47	0.44
1:C:2458:GLN:NE2	1:C:2459:SER:OG	2.51	0.44
1:C:2472:PHE:CE2	1:C:2476:PHE:HE2	2.35	0.44
1:E:994:ILE:HG12	1:E:1060:ASN:O	2.18	0.44
1:E:1098:ILE:HG23	1:E:1102:HIS:CB	2.47	0.44
1:E:1149:TRP:CE3	1:E:1169:TYR:HD1	2.35	0.44
1:E:1321:LEU:HA	1:E:1321:LEU:HD23	1.78	0.44
1:E:1712:MET:HG3	1:E:1716:ARG:HH21	1.83	0.44
1:E:2305:LEU:HD23	1:E:2305:LEU:HA	1.68	0.44
1:A:989:TRP:O	1:A:993:ILE:HG12	2.17	0.44
1:A:1477:LEU:HD21	1:A:2720:THR:HG22	1.99	0.44
1:C:925:GLU:HG3	1:C:926:TYR:N	2.33	0.44
1:C:1279:PHE:HA	1:C:1283:ARG:HH22	1.83	0.44
1:C:2440:SER:HA	1:C:2455:SER:HA	1.98	0.44
1:C:2557:LEU:HD13	1:C:2562:ARG:NE	2.32	0.44
1:E:994:ILE:HB	1:E:1061:LEU:HD21	1.99	0.44
1:E:1712:MET:SD	1:E:1716:ARG:NH2	2.90	0.44
1:E:2454:MET:HE3	1:E:2454:MET:HB3	1.93	0.44
1:E:2553:LEU:HB3	1:E:2555:PHE:CE2	2.52	0.44
1:E:2585:LYS:HB3	1:E:2611:PHE:CD2	2.52	0.44
1:A:2458:GLN:NE2	1:A:2459:SER:OG	2.51	0.44
1:C:994:ILE:HG12	1:C:1060:ASN:O	2.18	0.44
1:C:1098:ILE:HG23	1:C:1102:HIS:CB	2.47	0.44
1:C:1674:HIS:CE1	1:C:1919:VAL:HG12	2.53	0.44
1:E:1284:SER:H	1:E:1287:ASP:HB3	1.83	0.44
1:A:1279:PHE:HA	1:A:1283:ARG:HH22	1.83	0.44
1:A:2334:LEU:HA	1:A:2334:LEU:HD23	1.74	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2396:LYS:HD2	1:C:2396:LYS:HA	1.75	0.44
1:C:2553:LEU:HB3	1:C:2555:PHE:CE2	2.52	0.44
1:E:922:LYS:HE2	1:E:926:TYR:HE2	1.83	0.44
1:E:1124:LEU:HD13	1:E:1128:PHE:CE2	2.53	0.44
1:E:1310:THR:CB	1:E:2016:VAL:HG21	2.48	0.44
1:A:925:GLU:HG3	1:A:926:TYR:N	2.33	0.44
1:A:1488:GLN:C	1:A:1492:LYS:HE2	2.42	0.44
1:A:1666:TRP:CE2	1:A:1670:ILE:HD11	2.53	0.44
1:A:1711:ILE:HD13	1:A:1875:THR:HB	2.00	0.44
1:A:2407:MET:HA	1:A:2407:MET:HE2	2.00	0.44
1:A:2445:LEU:HD13	1:A:2530:VAL:HG22	2.00	0.44
1:A:2472:PHE:CE2	1:A:2476:PHE:HE2	2.35	0.44
1:C:2420:LEU:HD23	1:C:2420:LEU:HA	1.88	0.44
1:C:2433:ILE:HA	1:C:2656:LYS:HA	2.00	0.44
1:C:2560:ILE:HG23	1:C:2563:LYS:HE2	2.00	0.44
1:C:2709:LYS:HE2	1:C:2709:LYS:HB3	1.66	0.44
1:E:974:PRO:HB3	1:E:1170:CYS:HB3	1.99	0.44
1:E:2266:LEU:HD12	1:E:2266:LEU:HA	1.69	0.44
1:A:931:GLN:O	1:A:935:TRP:HD1	2.00	0.43
1:A:1545:MET:SD	1:C:2392:ARG:HD2	2.58	0.43
1:A:2270:ILE:HG21	1:A:2299:LYS:HG2	2.00	0.43
1:A:2709:LYS:HB3	1:A:2709:LYS:HE2	1.66	0.43
1:C:1024:ILE:HG23	1:C:1028:GLU:H	1.83	0.43
1:C:1447:LEU:HD12	1:C:1684:GLU:OE1	2.18	0.43
1:C:1688:LEU:HD13	1:C:1703:ILE:HG12	2.00	0.43
1:E:2299:LYS:HA	1:E:2299:LYS:HD3	1.73	0.43
1:E:2375:ILE:HD13	1:E:2375:ILE:HA	1.77	0.43
1:E:2703:ASN:OD1	1:E:2749:GLU:HA	2.18	0.43
1:A:1024:ILE:HG23	1:A:1028:GLU:H	1.83	0.43
1:A:1447:LEU:HD12	1:A:1684:GLU:OE1	2.18	0.43
1:A:1674:HIS:CE1	1:A:1919:VAL:HG12	2.53	0.43
1:A:2433:ILE:HA	1:A:2656:LYS:HA	2.00	0.43
1:A:2454:MET:CE	1:A:2499:LEU:HA	2.47	0.43
1:A:2585:LYS:HD3	1:A:2611:PHE:HB3	2.01	0.43
1:C:1076:HIS:HA	1:C:1079:TYR:CD1	2.54	0.43
1:C:1284:SER:H	1:C:1287:ASP:HB3	1.83	0.43
1:C:2621:ASP:OD1	1:C:2622:ASN:N	2.51	0.43
1:E:1024:ILE:HG23	1:E:1028:GLU:H	1.83	0.43
1:E:1308:THR:HG23	1:E:1414:TRP:HB3	2.00	0.43
1:E:1449:SER:HG	1:E:1700:ARG:NH1	2.14	0.43
1:E:2365:TRP:HZ3	1:E:2718:ARG:CZ	2.29	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1098:ILE:HG23	1:A:1102:HIS:CB	2.47	0.43
1:A:1553:LEU:O	1:A:1555:GLU:N	2.46	0.43
1:A:2516:LYS:O	1:A:2520:GLU:HB2	2.18	0.43
1:C:1394:TYR:CG	1:C:1395:GLN:N	2.85	0.43
1:C:2659:PRO:HB2	1:C:2661:SER:OG	2.18	0.43
1:E:1076:HIS:HA	1:E:1079:TYR:CD1	2.54	0.43
1:E:1447:LEU:HD12	1:E:1684:GLU:OE1	2.18	0.43
1:E:1674:HIS:CE1	1:E:1919:VAL:HG12	2.53	0.43
1:A:1149:TRP:CE3	1:A:1169:TYR:HD1	2.35	0.43
1:A:1156:ARG:HB2	1:A:1162:ILE:HG12	1.99	0.43
1:A:2621:ASP:OD1	1:A:2622:ASN:N	2.51	0.43
1:C:1321:LEU:HA	1:C:1321:LEU:HD23	1.78	0.43
1:C:2407:MET:HE2	1:C:2407:MET:HA	2.00	0.43
1:C:2445:LEU:HD13	1:C:2530:VAL:HG22	2.00	0.43
1:E:925:GLU:HG3	1:E:926:TYR:N	2.33	0.43
1:E:994:ILE:HA	1:E:997:LYS:HD3	2.00	0.43
1:A:2388:TYR:O	1:A:2390:GLN:NE2	2.51	0.43
1:C:2516:LYS:O	1:C:2520:GLU:HB2	2.19	0.43
1:C:2585:LYS:HD3	1:C:2611:PHE:HB3	2.01	0.43
1:E:2511:PRO:O	1:E:2514:LYS:HB2	2.18	0.43
1:E:2617:ILE:HG13	1:E:2618:LEU:O	2.19	0.43
1:E:2659:PRO:HB2	1:E:2661:SER:OG	2.18	0.43
1:A:1308:THR:HG23	1:A:1414:TRP:HB3	2.00	0.43
1:C:1362:ILE:O	1:C:1365:ILE:HG22	2.19	0.43
1:C:1477:LEU:HD21	1:C:2720:THR:HG22	1.99	0.43
1:C:2478:ARG:O	1:C:2480:THR:N	2.48	0.43
1:C:2479:ASP:C	1:C:2481:GLY:N	2.74	0.43
1:C:2703:ASN:OD1	1:C:2749:GLU:HA	2.18	0.43
1:E:1042:THR:O	1:E:1046:GLY:N	2.43	0.43
1:E:1211:TRP:NE1	1:E:1364:SER:OG	2.48	0.43
1:E:1279:PHE:HA	1:E:1283:ARG:HH22	1.83	0.43
1:E:1688:LEU:HD13	1:E:1703:ILE:HG12	2.00	0.43
1:E:2478:ARG:O	1:E:2480:THR:N	2.49	0.43
1:E:2516:LYS:O	1:E:2520:GLU:HB2	2.19	0.43
1:E:2587:TYR:OH	1:E:2649:GLU:O	2.28	0.43
1:A:2297:PHE:O	1:A:2300:CYS:HB3	2.19	0.43
1:A:2392:ARG:HD2	1:E:1545:MET:SD	2.58	0.43
1:A:2521:LEU:HD23	1:A:2521:LEU:HA	1.91	0.43
1:A:2557:LEU:HD13	1:A:2562:ARG:HB3	2.01	0.43
1:C:1112:LYS:O	1:C:1116:ASN:ND2	2.52	0.43
1:C:2297:PHE:O	1:C:2300:CYS:HB3	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2407:MET:HE2	1:E:2407:MET:HA	2.01	0.43
1:A:1362:ILE:O	1:A:1365:ILE:HG22	2.19	0.43
1:A:2392:ARG:HH21	1:E:1542:HIS:HA	1.83	0.43
1:A:2511:PRO:O	1:A:2514:LYS:HB2	2.18	0.43
1:A:2635:LEU:HD12	1:A:2635:LEU:HA	1.82	0.43
1:E:2388:TYR:O	1:E:2390:GLN:NE2	2.52	0.43
1:E:2585:LYS:HD3	1:E:2611:PHE:HB3	2.01	0.43
1:E:2635:LEU:HD12	1:E:2635:LEU:HA	1.82	0.43
1:A:994:ILE:HA	1:A:997:LYS:HD3	2.00	0.43
1:A:1153:VAL:HG13	1:A:1165:ILE:HD11	2.01	0.43
1:A:1394:TYR:CG	1:A:1395:GLN:N	2.86	0.43
1:C:938:LEU:O	1:C:942:ILE:HB	2.19	0.43
1:C:1042:THR:O	1:C:1046:GLY:N	2.43	0.43
1:C:1124:LEU:HD13	1:C:1128:PHE:CE2	2.53	0.43
1:C:2617:ILE:HG13	1:C:2618:LEU:O	2.19	0.43
1:E:1666:TRP:CE2	1:E:1670:ILE:HD11	2.53	0.43
1:E:1711:ILE:HD13	1:E:1875:THR:HB	2.00	0.43
1:E:2560:ILE:HG23	1:E:2563:LYS:HE2	2.00	0.43
1:E:2593:ALA:HB3	1:E:2655:ASP:OD1	2.19	0.43
1:A:932:VAL:O	1:A:935:TRP:HB2	2.19	0.43
1:A:1110:CYS:SG	1:A:1286:LEU:HD13	2.59	0.43
1:A:1339:ILE:HG22	1:A:1343:LEU:HG	2.01	0.43
1:A:1688:LEU:HD13	1:A:1703:ILE:HG12	2.00	0.43
2:B:237:MET:HE1	1:C:2329:TYR:CE1	2.54	0.43
1:C:1679:THR:O	1:C:1683:ILE:HD12	2.19	0.43
1:C:2016:VAL:O	1:C:2020:LEU:N	2.44	0.43
1:C:2270:ILE:HG21	1:C:2299:LYS:HG2	2.00	0.43
1:C:2388:TYR:O	1:C:2390:GLN:NE2	2.52	0.43
1:C:2576:SER:O	1:C:2579:THR:N	2.48	0.43
1:C:2585:LYS:HA	1:C:2612:MET:O	2.19	0.43
1:C:2593:ALA:HB3	1:C:2655:ASP:OD1	2.19	0.43
1:C:2630:TRP:CH2	1:E:2512:PRO:HG2	2.54	0.43
1:E:1110:CYS:SG	1:E:1286:LEU:HD13	2.59	0.43
1:E:1672:ARG:HA	1:E:1672:ARG:HD3	1.71	0.43
1:A:1112:LYS:O	1:A:1116:ASN:ND2	2.52	0.42
1:A:2659:PRO:HB2	1:A:2661:SER:OG	2.18	0.42
1:C:1666:TRP:CE2	1:C:1670:ILE:HD11	2.53	0.42
1:C:1711:ILE:HD13	1:C:1875:THR:HB	2.00	0.42
1:C:2483:MET:SD	1:C:2486:LEU:HD23	2.59	0.42
1:C:2738:ARG:HH11	1:C:2738:ARG:HD3	1.69	0.42
1:E:1112:LYS:O	1:E:1116:ASN:ND2	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2334:LEU:HD23	1:E:2334:LEU:HA	1.74	0.42
1:E:2433:ILE:HA	1:E:2656:LYS:HA	2.00	0.42
1:E:2625:LYS:HD2	1:E:2625:LYS:HA	1.86	0.42
1:A:1679:THR:O	1:A:1683:ILE:HD12	2.20	0.42
1:A:2397:LYS:O	1:A:2400:VAL:N	2.48	0.42
1:A:2619:SER:N	1:A:2630:TRP:HB2	2.34	0.42
1:A:2744:ILE:HD11	1:E:2738:ARG:HG2	2.01	0.42
1:C:932:VAL:O	1:C:935:TRP:HB2	2.19	0.42
1:C:1339:ILE:HG22	1:C:1343:LEU:HG	2.01	0.42
1:C:2397:LYS:O	1:C:2400:VAL:N	2.48	0.42
1:E:1679:THR:O	1:E:1683:ILE:HD12	2.20	0.42
1:E:2558:LYS:HZ2	1:E:2560:ILE:HB	1.83	0.42
1:A:1131:SER:HB3	1:A:1232:LEU:HD13	2.02	0.42
1:C:1110:CYS:SG	1:C:1286:LEU:HD13	2.59	0.42
1:C:1899:LYS:HA	1:C:1899:LYS:HD3	1.89	0.42
1:C:2619:SER:N	1:C:2630:TRP:HB2	2.34	0.42
1:E:997:LYS:HA	1:E:1047:LEU:HD22	2.01	0.42
1:E:1339:ILE:HG22	1:E:1343:LEU:HG	2.01	0.42
1:A:1061:LEU:O	1:A:1064:LEU:N	2.53	0.42
1:A:1076:HIS:HA	1:A:1079:TYR:CD1	2.54	0.42
1:A:1934:HIS:NE2	1:A:1943:LEU:HD13	2.34	0.42
1:A:2738:ARG:HG2	1:C:2744:ILE:HD11	2.01	0.42
1:C:1153:VAL:HG13	1:C:1165:ILE:HD11	2.01	0.42
1:C:1310:THR:CB	1:C:2016:VAL:HG21	2.48	0.42
1:C:1874:LEU:HG	1:C:1878:LEU:HD13	2.02	0.42
1:C:2413:ILE:HD12	1:C:2413:ILE:HA	1.84	0.42
1:C:2504:ASN:HA	1:E:2513:SER:OG	2.19	0.42
1:E:938:LEU:O	1:E:942:ILE:HB	2.19	0.42
1:E:1153:VAL:HG13	1:E:1165:ILE:HD11	2.01	0.42
1:E:1364:SER:HA	1:E:1367:ALA:HB3	2.01	0.42
1:E:2270:ILE:HG21	1:E:2299:LYS:HG2	2.00	0.42
1:E:2445:LEU:HD13	1:E:2530:VAL:HG22	2.00	0.42
1:A:997:LYS:HA	1:A:1047:LEU:HD22	2.02	0.42
1:A:1190:ALA:HB3	1:A:1191:PRO:HD3	2.02	0.42
1:A:1364:SER:HA	1:A:1367:ALA:HB3	2.01	0.42
1:A:2400:VAL:O	2:F:235:ILE:HD11	2.19	0.42
1:A:2479:ASP:C	1:A:2481:GLY:N	2.74	0.42
1:A:2480:THR:HB	1:A:2484:GLN:CD	2.45	0.42
1:A:2504:ASN:HA	1:C:2513:SER:OG	2.20	0.42
1:A:2521:LEU:HB3	1:A:2562:ARG:O	2.19	0.42
1:A:2593:ALA:HB3	1:A:2655:ASP:OD1	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2617:ILE:HG13	1:A:2618:LEU:O	2.19	0.42
1:A:2696:ILE:HD12	1:E:2737:TYR:HD2	1.85	0.42
1:C:2350:VAL:HG22	1:C:2362:LEU:HD11	2.02	0.42
1:C:2564:ASN:HA	1:C:2567:LYS:HE2	2.02	0.42
1:C:2627:ASN:HB2	1:C:2629:GLU:OE2	2.19	0.42
1:E:1874:LEU:HG	1:E:1878:LEU:HD13	2.02	0.42
1:E:2620:ARG:HA	1:E:2630:TRP:HD1	1.85	0.42
1:E:2627:ASN:HB2	1:E:2629:GLU:OE2	2.19	0.42
1:A:1008:GLU:HG3	1:A:1009:ASN:N	2.35	0.42
1:A:1712:MET:SD	1:A:1716:ARG:NH2	2.90	0.42
1:A:2489:TYR:CG	1:A:2654:ASN:OD1	2.73	0.42
1:A:2608:GLU:CA	1:A:2611:PHE:HB2	2.46	0.42
1:C:944:LYS:NZ	1:C:1092:ARG:HH12	2.17	0.42
1:C:2467:GLN:HG2	1:C:2471:LYS:HE3	2.02	0.42
1:E:1103:LEU:HD13	1:E:1113:TYR:CD1	2.55	0.42
1:E:1361:ASN:O	1:E:1364:SER:OG	2.36	0.42
1:E:1983:GLN:HB2	1:E:2011:LYS:CB	2.50	0.42
1:E:2467:GLN:C	1:E:2471:LYS:HZ2	2.27	0.42
1:A:944:LYS:HZ2	1:A:1092:ARG:HH22	1.68	0.42
1:A:1211:TRP:NE1	1:A:1364:SER:OG	2.48	0.42
1:A:1310:THR:CB	1:A:2016:VAL:HG21	2.48	0.42
1:A:2701:LEU:HA	1:A:2701:LEU:HD23	1.85	0.42
1:C:1712:MET:SD	1:C:1716:ARG:NH2	2.90	0.42
1:C:2375:ILE:HA	1:C:2375:ILE:HD13	1.77	0.42
1:C:2679:LEU:HD23	1:C:2679:LEU:HA	1.74	0.42
1:E:1362:ILE:O	1:E:1365:ILE:HG22	2.19	0.42
1:E:1708:GLN:NE2	1:E:1873:PRO:HG2	2.35	0.42
1:E:2483:MET:SD	1:E:2486:LEU:HD23	2.60	0.42
1:E:2604:GLN:CD	1:E:2604:GLN:H	2.28	0.42
1:E:2619:SER:N	1:E:2630:TRP:HB2	2.34	0.42
1:E:2708:LEU:HD12	1:E:2708:LEU:HA	1.67	0.42
1:A:1047:LEU:H	1:A:1047:LEU:HD23	1.85	0.42
1:A:1078:GLU:HA	1:A:1081:ARG:HG2	2.02	0.42
1:A:1124:LEU:HD13	1:A:1128:PHE:CE2	2.53	0.42
1:A:1983:GLN:HB2	1:A:2011:LYS:CB	2.50	0.42
1:A:2585:LYS:HA	1:A:2612:MET:O	2.19	0.42
1:C:1542:HIS:HA	1:E:2392:ARG:HH21	1.84	0.42
1:C:1672:ARG:HD3	1:C:1672:ARG:HA	1.71	0.42
1:C:1983:GLN:HB2	1:C:2011:LYS:CB	2.50	0.42
1:C:2033:LEU:HD23	1:C:2033:LEU:HA	1.79	0.42
1:C:2469:PHE:O	1:C:2474:GLN:NE2	2.48	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:932:VAL:O	1:E:935:TRP:HB2	2.19	0.42
1:E:1131:SER:HB3	1:E:1232:LEU:HD13	2.02	0.42
1:E:2521:LEU:HD23	1:E:2521:LEU:HA	1.91	0.42
1:E:2701:LEU:HA	1:E:2701:LEU:HD23	1.85	0.42
1:A:938:LEU:O	1:A:942:ILE:HB	2.19	0.42
1:A:2445:LEU:HD12	1:A:2529:SER:O	2.20	0.42
1:A:2630:TRP:CZ2	1:C:2512:PRO:HD2	2.55	0.42
1:C:997:LYS:HA	1:C:1047:LEU:HD22	2.01	0.42
1:C:1008:GLU:HG3	1:C:1009:ASN:N	2.35	0.42
1:C:1447:LEU:CD1	1:C:1684:GLU:HB3	2.50	0.42
1:C:1708:GLN:NE2	1:C:1873:PRO:HG2	2.35	0.42
1:C:2436:PRO:HB3	1:C:2538:ARG:HA	2.02	0.42
1:C:2476:PHE:CE1	1:C:2605:LEU:HD22	2.55	0.42
1:C:2489:TYR:CG	1:C:2654:ASN:OD1	2.73	0.42
1:C:2557:LEU:HD13	1:C:2562:ARG:HB3	2.01	0.42
1:E:1047:LEU:HD23	1:E:1047:LEU:H	1.85	0.42
1:E:1061:LEU:O	1:E:1064:LEU:N	2.53	0.42
1:E:1266:ASP:OD2	1:E:1338:PRO:HG3	2.20	0.42
1:E:2451:ILE:HG22	1:E:2507:TRP:NE1	2.35	0.42
1:E:2557:LEU:HD13	1:E:2562:ARG:HB3	2.01	0.42
1:A:931:GLN:O	1:A:935:TRP:CD1	2.73	0.42
1:A:1447:LEU:CD1	1:A:1684:GLU:HB3	2.50	0.42
1:A:2604:GLN:H	1:A:2604:GLN:CD	2.28	0.42
1:C:922:LYS:HE2	1:C:926:TYR:HE2	1.84	0.42
1:C:2451:ILE:HG22	1:C:2507:TRP:NE1	2.35	0.42
1:C:2560:ILE:O	1:C:2561:THR:C	2.62	0.42
1:C:2737:TYR:HD2	1:E:2696:ILE:HD12	1.84	0.42
1:E:931:GLN:O	1:E:935:TRP:CD1	2.73	0.42
1:E:2297:PHE:O	1:E:2300:CYS:HB3	2.19	0.42
1:E:2436:PRO:HB3	1:E:2538:ARG:HA	2.02	0.42
1:A:1266:ASP:OD2	1:A:1338:PRO:HG3	2.20	0.41
1:A:2266:LEU:HD12	1:A:2266:LEU:HA	1.69	0.41
1:A:2451:ILE:HG22	1:A:2507:TRP:NE1	2.35	0.41
1:A:2483:MET:C	1:A:2485:PHE:N	2.76	0.41
1:A:2627:ASN:HB2	1:A:2629:GLU:OE2	2.19	0.41
1:C:1103:LEU:HD13	1:C:1113:TYR:CD1	2.55	0.41
1:C:2257:LEU:O	1:C:2261:ILE:HD12	2.20	0.41
1:C:2445:LEU:HD12	1:C:2529:SER:O	2.20	0.41
1:C:2480:THR:HB	1:C:2484:GLN:CD	2.45	0.41
1:C:2551:ASP:HB2	1:C:2599:SER:O	2.20	0.41
1:E:1190:ALA:HB3	1:E:1191:PRO:HD3	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1411:GLY:O	1:E:1415:ASP:HB2	2.20	0.41
1:E:1934:HIS:NE2	1:E:1943:LEU:HD13	2.34	0.41
1:A:2620:ARG:HA	1:A:2630:TRP:HD1	1.85	0.41
1:C:1047:LEU:HD23	1:C:1047:LEU:H	1.85	0.41
1:C:1190:ALA:HB3	1:C:1191:PRO:HD3	2.02	0.41
1:C:2344:LEU:HD12	1:C:2344:LEU:HA	1.82	0.41
1:C:2521:LEU:HB3	1:C:2562:ARG:O	2.19	0.41
2:D:238:GLU:CB	1:E:2404:MET:HE1	2.51	0.41
1:E:1370:TYR:HB3	1:E:1374:LEU:CD1	2.49	0.41
1:E:2489:TYR:CG	1:E:2654:ASN:OD1	2.73	0.41
1:E:2551:ASP:HB2	1:E:2599:SER:O	2.20	0.41
1:E:2585:LYS:HA	1:E:2612:MET:O	2.19	0.41
1:A:1024:ILE:HG23	1:A:1028:GLU:HB2	2.01	0.41
1:A:1708:GLN:NE2	1:A:1873:PRO:HG2	2.35	0.41
1:A:2257:LEU:O	1:A:2261:ILE:HD12	2.20	0.41
1:A:2410:VAL:O	1:A:2413:ILE:HG22	2.21	0.41
1:A:2560:ILE:O	1:A:2561:THR:C	2.62	0.41
1:C:1688:LEU:HD12	1:C:1688:LEU:HA	1.94	0.41
1:C:1934:HIS:NE2	1:C:1943:LEU:HD13	2.34	0.41
1:C:2483:MET:C	1:C:2485:PHE:N	2.76	0.41
1:C:2604:GLN:CD	1:C:2604:GLN:H	2.28	0.41
1:C:2659:PRO:C	1:C:2661:SER:H	2.28	0.41
1:E:1892:ASP:OD1	1:E:1893:GLU:N	2.54	0.41
1:E:2480:THR:HB	1:E:2484:GLN:CD	2.45	0.41
1:E:2560:ILE:O	1:E:2561:THR:C	2.62	0.41
1:E:2659:PRO:C	1:E:2661:SER:H	2.28	0.41
1:A:2202:ILE:HA	1:A:2202:ILE:HD13	1.92	0.41
1:A:2483:MET:SD	1:A:2486:LEU:HD23	2.60	0.41
1:A:2625:LYS:HD2	1:A:2625:LYS:HA	1.86	0.41
1:E:1008:GLU:HG3	1:E:1009:ASN:N	2.35	0.41
1:E:1078:GLU:HA	1:E:1081:ARG:HG2	2.02	0.41
1:E:2476:PHE:CE1	1:E:2605:LEU:HD22	2.55	0.41
1:E:2479:ASP:C	1:E:2481:GLY:N	2.74	0.41
1:E:2564:ASN:HA	1:E:2567:LYS:HE2	2.02	0.41
2:F:246:SER:O	2:F:246:SER:OG	2.23	0.41
1:A:1205:ASN:O	1:A:1208:ILE:HG22	2.21	0.41
1:A:1350:ILE:CD1	1:A:1422:LEU:HB3	2.50	0.41
1:A:1948:LEU:HD23	1:A:1967:MET:CE	2.51	0.41
1:A:2551:ASP:OD1	1:A:2552:LYS:N	2.52	0.41
1:A:2551:ASP:HB2	1:A:2599:SER:O	2.20	0.41
1:A:2564:ASN:HA	1:A:2567:LYS:HE2	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:944:LYS:HZ1	1:C:1092:ARG:HH12	1.67	0.41
1:C:966:LEU:HD23	1:C:1181:TYR:CG	2.56	0.41
1:C:1689:THR:HA	1:C:1692:ILE:HG22	2.03	0.41
1:E:2257:LEU:O	1:E:2261:ILE:HD12	2.20	0.41
1:E:2350:VAL:HG22	1:E:2362:LEU:HD11	2.02	0.41
1:E:2396:LYS:HA	1:E:2396:LYS:HD2	1.75	0.41
1:E:2521:LEU:HB3	1:E:2562:ARG:O	2.19	0.41
1:A:922:LYS:HE2	1:A:926:TYR:HE2	1.83	0.41
1:A:944:LYS:NZ	1:A:1092:ARG:HH12	2.17	0.41
1:A:2339:ARG:HA	1:A:2345:THR:CG2	2.51	0.41
1:A:2551:ASP:CG	1:A:2552:LYS:H	2.27	0.41
1:A:2608:GLU:HA	1:A:2611:PHE:CD1	2.56	0.41
1:A:2737:TYR:HD1	1:A:2737:TYR:HA	1.69	0.41
1:C:931:GLN:O	1:C:935:TRP:CD1	2.73	0.41
1:C:1024:ILE:HG23	1:C:1028:GLU:HB2	2.01	0.41
1:C:1980:TYR:O	1:C:1983:GLN:HG2	2.21	0.41
1:C:2620:ARG:HA	1:C:2630:TRP:HD1	1.85	0.41
1:E:931:GLN:NE2	1:E:932:VAL:HG23	2.36	0.41
1:E:944:LYS:NZ	1:E:1092:ARG:HH12	2.17	0.41
1:E:1447:LEU:CD1	1:E:1684:GLU:HB3	2.50	0.41
1:E:2339:ARG:HA	1:E:2345:THR:CG2	2.51	0.41
1:E:2408:ILE:HD13	1:E:2408:ILE:HA	1.93	0.41
1:E:2578:LYS:HA	1:E:2578:LYS:HD3	1.81	0.41
1:A:1305:ILE:HG22	1:A:1324:CYS:HB2	2.03	0.41
1:A:2026:LEU:HD23	1:A:2026:LEU:HA	1.86	0.41
1:A:2467:GLN:HG2	1:A:2471:LYS:HE3	2.02	0.41
1:A:2471:LYS:HG2	1:A:2471:LYS:HZ2	1.76	0.41
1:C:1131:SER:HB3	1:C:1232:LEU:HD13	2.02	0.41
1:C:1290:LYS:HD3	1:C:1290:LYS:HA	1.93	0.41
1:C:1304:ILE:HG22	1:C:1418:CYS:SG	2.61	0.41
1:C:1411:GLY:O	1:C:1415:ASP:HB2	2.20	0.41
1:C:1445:GLN:O	1:C:1448:ALA:HB3	2.21	0.41
1:C:2558:LYS:HG3	1:C:2561:THR:HG23	2.02	0.41
1:C:2608:GLU:HA	1:C:2611:PHE:CD1	2.56	0.41
1:E:1689:THR:HA	1:E:1692:ILE:HG22	2.03	0.41
1:E:2558:LYS:HG3	1:E:2561:THR:HG23	2.02	0.41
1:A:2476:PHE:CE1	1:A:2605:LEU:HD22	2.55	0.41
1:A:2737:TYR:HD2	1:C:2696:ILE:HD12	1.85	0.41
1:C:1061:LEU:O	1:C:1064:LEU:N	2.53	0.41
1:C:1220:ARG:HD3	1:C:1220:ARG:HA	1.96	0.41
1:C:1306:PHE:CD1	1:C:2020:LEU:HD13	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2375:ILE:HG23	1:C:2704:VAL:HB	2.03	0.41
1:C:2743:MET:HE2	1:C:2743:MET:HB3	1.55	0.41
1:E:2729:LEU:HA	1:E:2729:LEU:HD23	1.74	0.41
1:A:1103:LEU:HD13	1:A:1113:TYR:CD1	2.55	0.41
1:A:1558:SER:OG	1:A:1560:GLU:HG2	2.21	0.41
1:A:1874:LEU:HG	1:A:1878:LEU:HD13	2.02	0.41
1:A:1980:TYR:O	1:A:1983:GLN:HG2	2.21	0.41
1:A:2253:ARG:O	1:A:2254:LYS:C	2.64	0.41
1:A:2302:TYR:CD2	1:A:2302:TYR:C	2.99	0.41
1:A:2468:SER:HA	1:A:2472:PHE:H	1.86	0.41
1:A:2578:LYS:HD3	1:A:2578:LYS:HA	1.81	0.41
2:B:229:SER:C	2:B:231:ASP:N	2.79	0.41
1:C:1063:MET:HE3	1:C:1063:MET:HB3	1.86	0.41
1:C:1078:GLU:HA	1:C:1081:ARG:HG2	2.02	0.41
1:C:1214:PHE:HD1	1:C:1215:PRO:HD2	1.86	0.41
1:C:1266:ASP:OD2	1:C:1338:PRO:HG3	2.20	0.41
1:C:1364:SER:HA	1:C:1367:ALA:HB3	2.01	0.41
1:C:1948:LEU:HD23	1:C:1967:MET:CE	2.51	0.41
1:C:2366:ILE:HD13	1:C:2366:ILE:HA	1.81	0.41
1:C:2410:VAL:O	1:C:2413:ILE:HG22	2.20	0.41
1:C:2551:ASP:CG	1:C:2552:LYS:H	2.28	0.41
1:C:2578:LYS:HA	1:C:2578:LYS:HD3	1.81	0.41
1:C:2589:TYR:H	1:C:2606:LEU:HD23	1.86	0.41
1:C:2708:LEU:HD12	1:C:2708:LEU:HA	1.67	0.41
1:C:2738:ARG:HG2	1:E:2744:ILE:HD11	2.02	0.41
2:D:229:SER:C	2:D:231:ASP:H	2.29	0.41
1:E:973:LEU:HB2	1:E:974:PRO:HD3	2.03	0.41
1:E:1024:ILE:HG23	1:E:1028:GLU:HB2	2.02	0.41
1:E:1304:ILE:HG22	1:E:1418:CYS:SG	2.61	0.41
1:E:1558:SER:OG	1:E:1560:GLU:HG2	2.20	0.41
1:E:2016:VAL:O	1:E:2020:LEU:N	2.45	0.41
1:E:2366:ILE:HD13	1:E:2366:ILE:HA	1.80	0.41
1:E:2375:ILE:HG23	1:E:2704:VAL:HB	2.03	0.41
1:E:2445:LEU:HD12	1:E:2529:SER:O	2.20	0.41
1:E:2468:SER:HA	1:E:2472:PHE:H	1.86	0.41
1:E:2483:MET:C	1:E:2485:PHE:N	2.76	0.41
1:E:2551:ASP:CG	1:E:2552:LYS:H	2.27	0.41
1:E:2576:SER:O	1:E:2579:THR:N	2.48	0.41
1:A:2512:PRO:HD2	1:E:2630:TRP:CZ2	2.55	0.41
1:C:923:PHE:O	1:C:927:PHE:HD1	2.04	0.41
1:C:2340:LEU:HA	1:C:2340:LEU:HD23	1.74	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1267:ALA:HB3	1:E:1442:LYS:HZ1	1.86	0.41
1:E:1980:TYR:O	1:E:1983:GLN:HG2	2.21	0.41
1:A:1302:LEU:HD23	1:A:1302:LEU:HA	1.86	0.40
1:A:1306:PHE:CD1	1:A:2020:LEU:HD13	2.56	0.40
1:A:1712:MET:O	1:A:1717:GLU:N	2.36	0.40
1:A:1892:ASP:OD1	1:A:1893:GLU:N	2.54	0.40
1:A:2350:VAL:HG22	1:A:2362:LEU:HD11	2.02	0.40
1:A:2513:SER:OG	1:E:2504:ASN:HA	2.20	0.40
1:A:2565:ILE:HD12	1:A:2565:ILE:H	1.86	0.40
1:A:2588:PRO:HB3	1:A:2611:PHE:CE1	2.57	0.40
2:B:229:SER:C	2:B:231:ASP:H	2.29	0.40
1:C:1558:SER:OG	1:C:1560:GLU:HG2	2.21	0.40
2:D:229:SER:C	2:D:231:ASP:N	2.79	0.40
1:E:1158:ARG:O	1:E:1162:ILE:HG13	2.22	0.40
1:E:2467:GLN:HG2	1:E:2471:LYS:HE3	2.02	0.40
2:F:229:SER:C	2:F:231:ASP:N	2.79	0.40
1:A:1027:ASN:ND2	1:A:1030:ASN:OD1	2.54	0.40
1:A:1140:ASP:HA	1:A:1211:TRP:CD1	2.56	0.40
1:A:1304:ILE:HG22	1:A:1418:CYS:SG	2.61	0.40
1:A:1669:SER:HA	1:A:1672:ARG:HG2	2.03	0.40
1:A:2490:GLU:O	1:A:2491:LYS:C	2.64	0.40
1:C:931:GLN:NE2	1:C:932:VAL:HG23	2.35	0.40
1:C:1158:ARG:O	1:C:1162:ILE:HG13	2.22	0.40
1:C:1205:ASN:O	1:C:1208:ILE:HG22	2.21	0.40
1:C:1433:TYR:O	1:C:1437:VAL:HG23	2.21	0.40
1:C:1892:ASP:OD1	1:C:1893:GLU:N	2.54	0.40
1:C:2339:ARG:HA	1:C:2345:THR:CG2	2.51	0.40
1:C:2408:ILE:HD13	1:C:2408:ILE:HA	1.93	0.40
1:C:2608:GLU:CA	1:C:2611:PHE:HB2	2.46	0.40
1:E:1157:ARG:HB2	1:E:1158:ARG:HH11	1.86	0.40
1:E:1433:TYR:O	1:E:1437:VAL:HG23	2.21	0.40
1:E:1945:LEU:HB3	1:E:1946:PRO:HD3	2.03	0.40
1:E:2565:ILE:HD12	1:E:2565:ILE:H	1.86	0.40
1:A:1433:TYR:O	1:A:1437:VAL:HG23	2.21	0.40
1:A:1689:THR:HA	1:A:1692:ILE:HG22	2.03	0.40
1:A:2263:GLN:HE22	1:A:2307:ALA:HB2	1.87	0.40
1:C:2438:ASP:O	1:C:2537:GLN:HB3	2.22	0.40
1:C:2565:ILE:HG22	1:C:2569:ILE:HD11	2.03	0.40
1:E:1899:LYS:HA	1:E:1899:LYS:HD3	1.89	0.40
1:E:2454:MET:CE	1:E:2500:GLU:H	2.30	0.40
1:E:2490:GLU:O	1:E:2491:LYS:C	2.65	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2588:PRO:HB3	1:E:2611:PHE:CE1	2.57	0.40
1:A:1267:ALA:HB3	1:A:1442:LYS:HZ1	1.86	0.40
1:A:1315:ILE:HD12	1:A:1315:ILE:HA	1.90	0.40
1:A:1411:GLY:O	1:A:1415:ASP:HB2	2.20	0.40
1:A:1873:PRO:O	1:A:1876:HIS:HB3	2.22	0.40
1:A:2436:PRO:HB3	1:A:2538:ARG:HA	2.02	0.40
1:A:2448:TYR:CD1	1:A:2509:ILE:HD11	2.57	0.40
1:A:2490:GLU:HB3	1:A:2491:LYS:H	1.80	0.40
1:A:2716:LEU:O	1:A:2720:THR:HG23	2.22	0.40
2:B:238:GLU:HG2	1:C:2404:MET:HE1	2.04	0.40
1:C:1873:PRO:O	1:C:1876:HIS:HB3	2.22	0.40
1:E:966:LEU:HD23	1:E:1181:TYR:CG	2.56	0.40
1:E:1027:ASN:ND2	1:E:1030:ASN:OD1	2.55	0.40
1:E:1350:ILE:CD1	1:E:1422:LEU:HB3	2.50	0.40
1:E:1703:ILE:O	1:E:1707:TYR:HD1	2.05	0.40
1:E:1945:LEU:HD12	1:E:1945:LEU:HA	1.86	0.40
1:E:2438:ASP:O	1:E:2537:GLN:HB3	2.22	0.40
1:E:2448:TYR:CD1	1:E:2509:ILE:HD11	2.57	0.40
1:E:2589:TYR:H	1:E:2606:LEU:HD23	1.86	0.40
1:A:1099:THR:N	1:A:1102:HIS:HB2	2.30	0.40
1:A:1152:ALA:O	1:A:1155:TYR:HB3	2.22	0.40
1:A:1158:ARG:O	1:A:1162:ILE:HG13	2.21	0.40
1:A:1214:PHE:HD1	1:A:1215:PRO:HD2	1.86	0.40
1:A:1445:GLN:O	1:A:1448:ALA:HB3	2.21	0.40
1:A:2409:ILE:HD13	1:A:2409:ILE:HA	1.91	0.40
1:A:2553:LEU:HA	1:A:2553:LEU:HD23	1.78	0.40
1:A:2589:TYR:H	1:A:2606:LEU:HD23	1.86	0.40
1:C:1712:MET:O	1:C:1717:GLU:HG2	2.22	0.40
1:C:2476:PHE:CZ	1:C:2605:LEU:HD13	2.57	0.40
1:C:2490:GLU:O	1:C:2491:LYS:C	2.65	0.40
1:E:923:PHE:O	1:E:927:PHE:HD1	2.05	0.40
1:E:1140:ASP:HA	1:E:1211:TRP:CD1	2.56	0.40
1:E:1214:PHE:HD1	1:E:1215:PRO:HD2	1.86	0.40
1:E:1948:LEU:HD23	1:E:1967:MET:CE	2.51	0.40
1:E:2253:ARG:O	1:E:2254:LYS:C	2.64	0.40

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

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	1378/3020 (46%)	1187 (86%)	190 (14%)	1 (0%)	48	79
1	C	1378/3020 (46%)	1187 (86%)	190 (14%)	1 (0%)	48	79
1	E	1378/3020 (46%)	1187 (86%)	190 (14%)	1 (0%)	48	79
2	B	20/267 (8%)	17 (85%)	3 (15%)	0	100	100
2	D	20/267 (8%)	17 (85%)	3 (15%)	0	100	100
2	F	20/267 (8%)	17 (85%)	3 (15%)	0	100	100
All	All	4194/9861 (42%)	3612 (86%)	579 (14%)	3 (0%)	49	79

All (3) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	2487	GLU
1	C	2487	GLU
1	E	2487	GLU

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	1268/2707 (47%)	1268 (100%)	0	100	100
1	C	1268/2707 (47%)	1268 (100%)	0	100	100
1	E	1268/2707 (47%)	1268 (100%)	0	100	100
2	B	20/221 (9%)	20 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	D	20/221 (9%)	20 (100%)	0	100	100
2	F	20/221 (9%)	20 (100%)	0	100	100
All	All	3864/8784 (44%)	3864 (100%)	0	100	100

There are no protein residues with a non-rotameric sidechain to report.

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (21) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	1027	ASN
1	A	1060	ASN
1	A	1425	GLN
1	A	1674	HIS
1	A	1713	ASN
1	A	2263	GLN
1	A	2458	GLN
1	C	1027	ASN
1	C	1060	ASN
1	C	1425	GLN
1	C	1674	HIS
1	C	1713	ASN
1	C	2263	GLN
1	C	2458	GLN
1	E	1027	ASN
1	E	1060	ASN
1	E	1674	HIS
1	E	1713	ASN
1	E	2263	GLN
1	E	2458	GLN
1	E	2488	ASN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.