



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 9, 2026 – 07:02 AM UTC

PDB ID : 5XA0 / pdb_00005xa0
Title : Crystal structure of inositol 1,4,5-trisphosphate receptor cytosolic domain
Authors : Hamada, K.; Miyatake, H.; Terauchi, A.; Mikoshiba, K.
Deposited on : 2017-03-10
Resolution : 5.81 Å(reported)

This is a Full wwPDB X-ray Structure 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/XrayValidationReportHelp>

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:

MolProbity	:	4-5-2 with Phenix2.0
Xtriage (Phenix)	:	2.0
EDS	:	3.0
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
CCP4	:	9.0.010 (Gargrove)
Density-Fitness	:	1.0.12
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.49

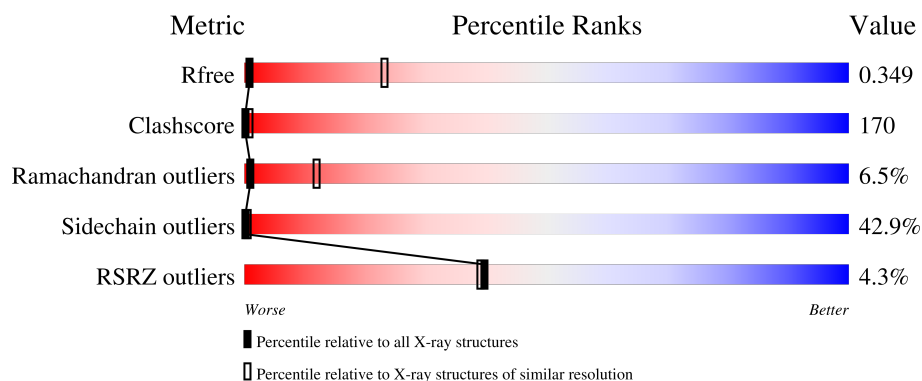
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 5.81 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)	Similar resolution (#Entries, resolution range(Å))
R_{free}	180053	1126 (7.64-4.00)
Clashscore	190562	1191 (7.64-4.00)
Ramachandran outliers	187476	1019 (7.60-4.00)
Sidechain outliers	187428	1009 (7.60-3.98)
RSRZ outliers	180081	1119 (7.64-4.00)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments of the lower 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 electron density. The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	1581	3% 35% 31% 12% • 21%
1	B	1581	4% 35% 30% 12% • 21%

2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 17919 atoms, of which 2970 are hydrogens and 0 are deuteriums.

In the tables below, the ZeroOcc column contains the number of atoms modelled with zero occupancy, 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.

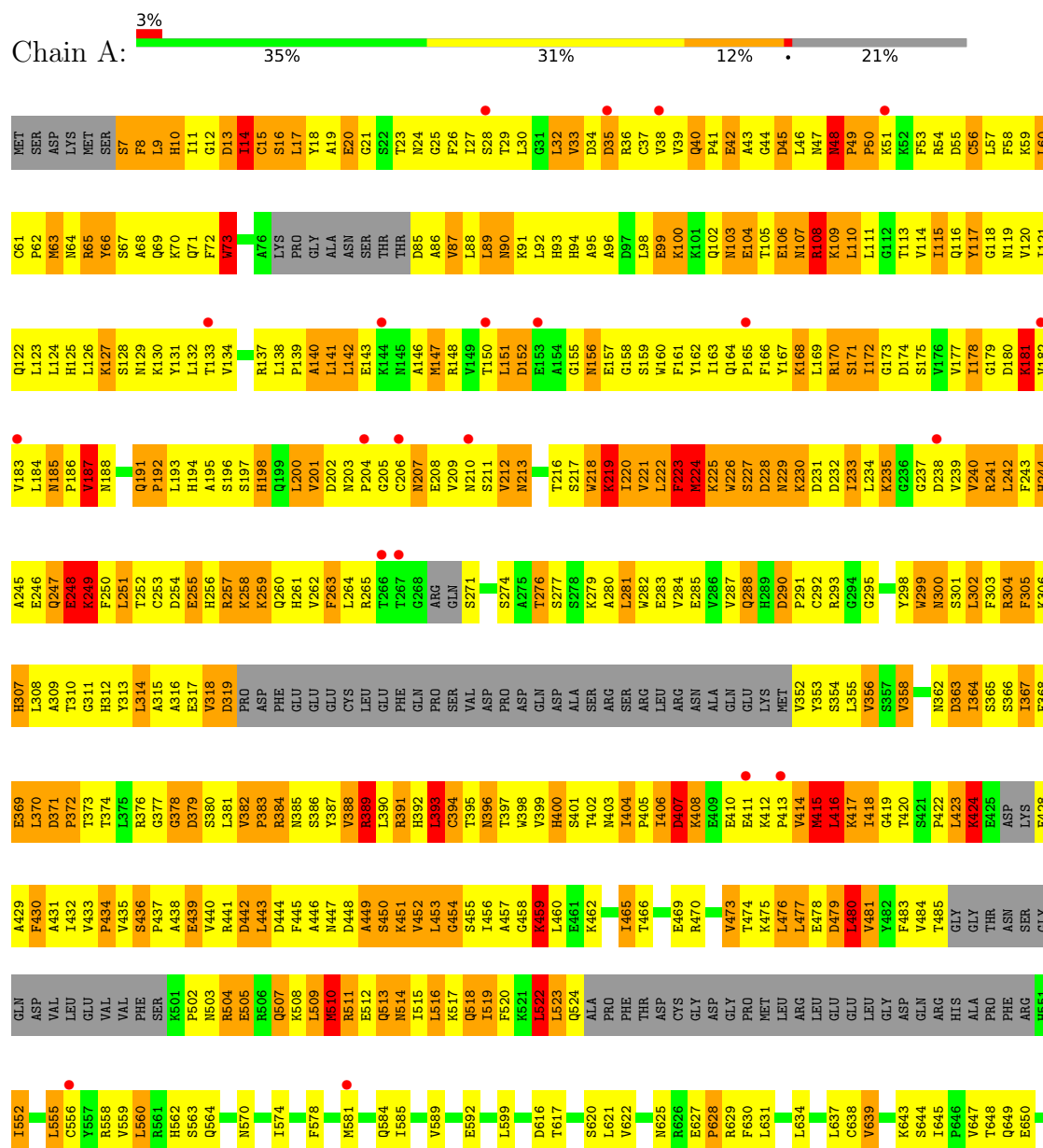
- Molecule 1 is a protein called Inositol 1,4,5-trisphosphate receptor type 1.

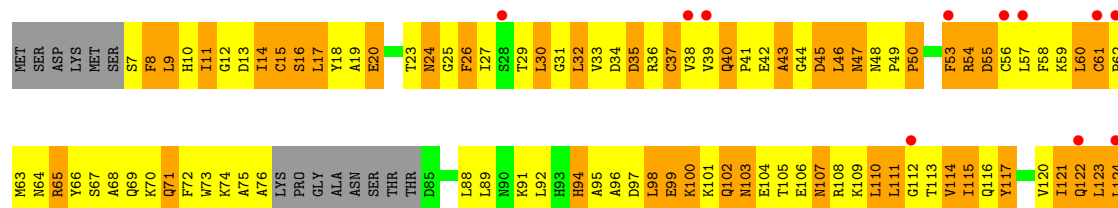
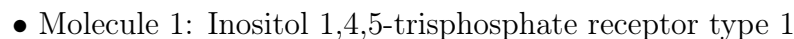
Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
1	A	1252	Total	C	H	N	O	S	0	0	0
			8949	4595	1485	1410	1444	15			
1	B	1252	Total	C	H	N	O	S	0	0	0
			8970	4610	1485	1412	1447	16			

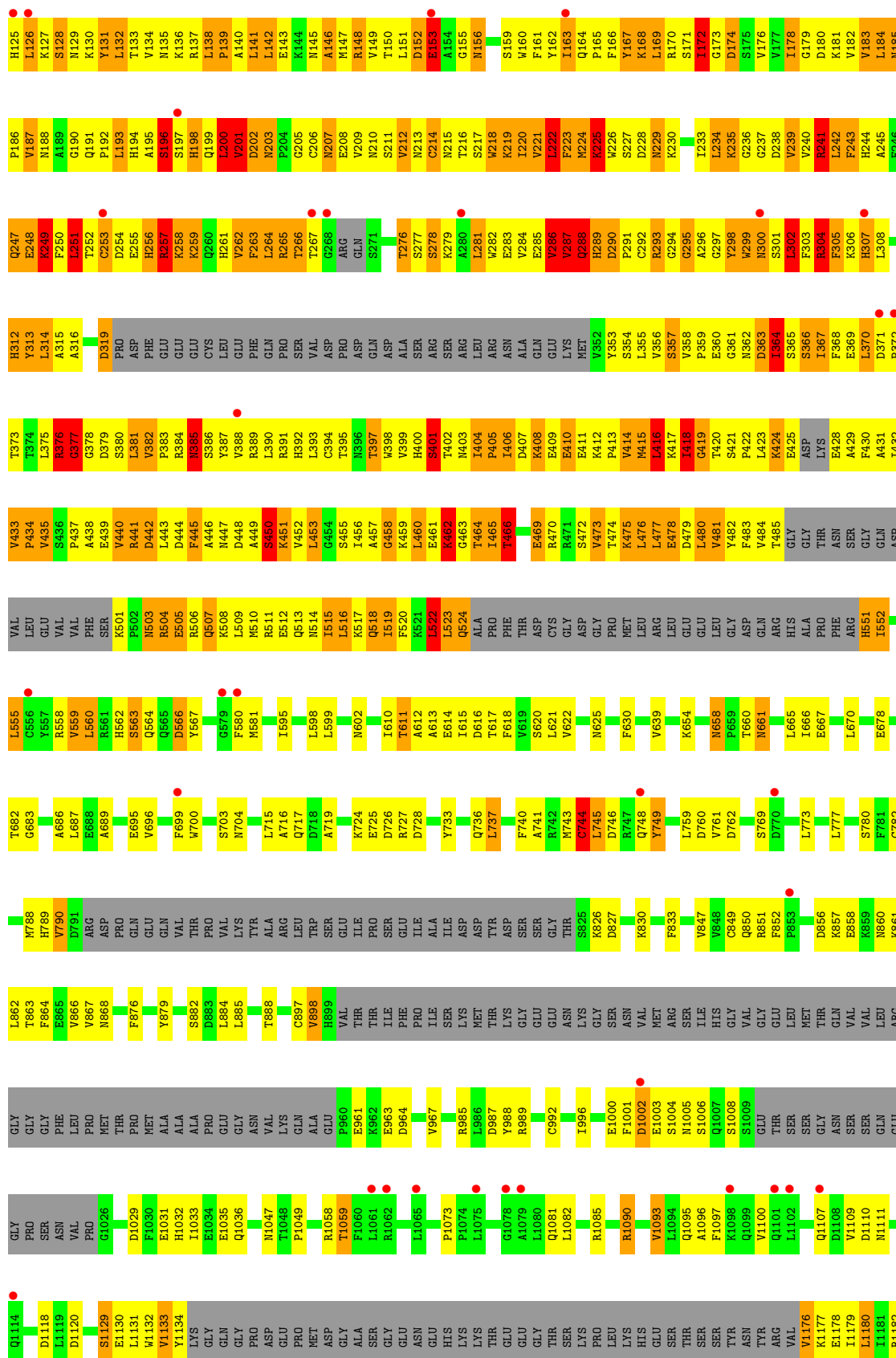
3 Residue-property plots

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 electron 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 dot above a residue indicates a poor fit to the electron density ($RSRZ > 2$). 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: Inositol 1,4,5-trisphosphate receptor type 1







SER	GLN	LYS	ALA	VAL	GLU	SER	CYS	ILE	ARG	VAL	LEU	SER	ASP	VAL	LYS	SER	ASN	THR	LEU	PHE	ASP	SER	GLN	VAL	ASN	THR	LEU	PHE	LYS	ALA	LEU	SER	HIS	ASN	ILE	VAL	GLN	LYS	THR	ALA	SER	ALA	ARG	Q1356	M1357	S1360	E1361	R1364	S1369	P1370	L1371	I1375	H1376	L1377	V1378	E1379	L1380	L1381	A1382	V1383	C1384	T1385	K1395	C1396	N1397	L1400	P1401	L1402	D1403	D1404	D1413	T1420	A1421	N1424	F1425	H1428	C1429	Y1430	V1431	D1432	T1433	E1434	V1435	K1438	E1439	I1440	Y1441	T1442	S1443	N1444	N1452								
																																												F1453	L1454	C1458	R1459	A1460	C1461	N1462	N1463	T1464	S1472	I1473	L1474	E1475	K1476	Y1477	V1478	T1481	T1488	F1489	F1490	S1491	S1492	P1493																																	
																																												L1183	S1184	C1187	V1188	S1191	A1192	S1193	V1194	R1195	K1196	S1197	R1198	R1203	L1204	L1205	R1206	N1207	N1208	G1209	A1210	H1211	A1212	V1213	V1214	L1215	E1216	L1217	L1218	Q1219	I1220	P1221	Y1222	K1223	K1224	A1225	E1226	D1227	T1228	K1229	M1230	Q1231	E1232	I1233	M1234	R1235	L1236	A1237	H1238	E1239	Q1242	N1243	F1244	C1245	A1246	G1247	N1248
																																												Q1249	Q1250	N1251	L1255	H1256	K1257	H1258	I1259	N1260	L1261	A1270	Q1274	H1275	M1278	N1279	N1280	F1281	Q1282	L1283	C1284	S1285	E1286	I1287	N1288	V1292	Q1293	T1301	I1309	I1315	V1316	K1317	A1318	E1319	I1323	M1329	V1330	E1339	D1340	V1341	L1342	V1343	F1344	Y1345	N1346	D1347	S1350	I1355									

4 Data and refinement statistics

Property	Value	Source
Space group	P 42	Depositor
Cell constants a, b, c, α , β , γ	128.16Å 128.16Å 369.49Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	48.81 – 5.81 48.81 – 5.81	Depositor EDS
% Data completeness (in resolution range)	99.9 (48.81-5.81) 99.9 (48.81-5.81)	Depositor EDS
R_{merge}	(Not available)	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.85 (at 5.73Å)	Xtriage
Refinement program	PHENIX (1.10.1_2155: ???)	Depositor
R, R_{free}	0.278 , 0.342 0.281 , 0.349	Depositor DCC
R_{free} test set	818 reflections (5.00%)	wwPDB-VP
Wilson B-factor (Å ²)	88.0	Xtriage
Anisotropy	0.101	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.24 , 389.8	EDS
L-test for twinning ²	$\langle L \rangle = 0.21$, $\langle L^2 \rangle = 0.07$	Xtriage
Estimated twinning fraction	0.368 for h,-k,-l	Xtriage
F_o, F_c correlation	0.80	EDS
Total number of atoms	17919	wwPDB-VP
Average B, all atoms (Å ²)	31.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 4.62% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

²Theoretical values of $\langle |L| \rangle$, $\langle L^2 \rangle$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

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	0.72	4/7527 (0.1%)	1.02	24/10362 (0.2%)
1	B	0.73	5/7549 (0.1%)	1.01	17/10390 (0.2%)
All	All	0.73	9/15076 (0.1%)	1.02	41/20752 (0.2%)

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	7
1	B	0	11
All	All	0	18

All (9) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	A	137	ARG	C-N	5.79	1.45	1.33
1	B	288	GLN	CG-CD	-5.74	1.37	1.52
1	A	430	PHE	CD1-CE1	5.63	1.55	1.38
1	B	462	LYS	CD-CE	5.55	1.69	1.52
1	B	385	ASN	CB-CG	-5.53	1.38	1.52
1	B	377	GLY	N-CA	5.42	1.53	1.45
1	A	415	MET	CB-CG	5.35	1.68	1.52
1	B	1176	VAL	CA-C	5.28	1.64	1.52
1	A	459	LYS	CD-CE	5.05	1.67	1.52

All (41) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	377	GLY	N-CA-C	12.29	142.31	113.18
1	B	376	ARG	CA-C-N	10.13	141.26	121.41
1	B	376	ARG	C-N-CA	10.13	141.26	121.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	108	ARG	CG-CD-NE	-9.50	91.10	112.00
1	B	243	PHE	CA-C-N	-9.39	103.20	122.07
1	B	243	PHE	C-N-CA	-9.39	103.20	122.07
1	A	480	LEU	CB-CG-CD1	-8.83	84.22	110.70
1	A	249	LYS	CG-CD-CE	-7.75	93.46	111.30
1	B	1226	GLU	N-CA-C	7.74	127.29	110.80
1	A	459	LYS	CG-CD-CE	7.19	127.83	111.30
1	A	416	LEU	CA-CB-CG	7.17	141.40	116.30
1	B	224	MET	CG-SD-CE	6.91	116.10	100.90
1	B	304	ARG	CA-CB-CG	6.68	127.46	114.10
1	A	181	LYS	CD-CE-NZ	6.66	133.22	111.90
1	A	99	GLU	CB-CG-CD	-6.59	101.40	112.60
1	B	555	LEU	CA-CB-CG	-6.40	93.92	116.30
1	A	73	TRP	CA-CB-CG	6.20	125.38	113.60
1	A	555	LEU	CA-CB-CG	-6.16	94.74	116.30
1	B	241	ARG	CG-CD-NE	-6.13	98.50	112.00
1	A	454	GLY	CA-C-N	6.02	129.46	120.31
1	A	454	GLY	C-N-CA	6.02	129.46	120.31
1	A	1243	ASN	N-CA-C	5.96	119.91	111.92
1	A	389	ARG	NE-CZ-NH1	-5.93	115.57	121.50
1	B	251	LEU	CB-CG-CD2	-5.86	93.12	110.70
1	A	48	ASN	C-N-CD	-5.80	101.22	125.00
1	A	219	LYS	CG-CD-CE	-5.75	98.07	111.30
1	A	522	LEU	CB-CG-CD2	-5.53	94.11	110.70
1	B	481	VAL	N-CA-C	-5.49	107.09	111.81
1	A	241	ARG	NE-CZ-NH1	-5.46	116.04	121.50
1	B	418	ILE	CG1-CB-CG2	-5.46	94.33	110.70
1	B	440	VAL	CA-C-N	-5.44	113.23	120.79
1	B	440	VAL	C-N-CA	-5.44	113.23	120.79
1	A	108	ARG	CA-CB-CG	5.41	124.92	114.10
1	A	21	GLY	N-CA-C	-5.29	102.83	111.78
1	A	14	ILE	CG1-CB-CG2	5.29	126.58	110.70
1	A	415	MET	CA-CB-CG	5.29	124.67	114.10
1	A	415	MET	CA-C-N	5.27	132.16	123.25
1	A	415	MET	C-N-CA	5.27	132.16	123.25
1	A	416	LEU	CD1-CG-CD2	-5.20	99.37	110.80
1	B	287	VAL	CA-CB-CG1	5.10	119.07	110.40
1	B	263	PHE	N-CA-C	5.02	116.55	108.32

There are no chirality outliers.

All (18) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	1204	LEU	Peptide
1	A	14	ILE	Peptide
1	A	213	ASN	Peptide
1	A	226	TRP	Peptide
1	A	248	GLU	Peptide
1	A	393	LEU	Peptide
1	A	416	LEU	Peptide
1	B	1090	ARG	Peptide
1	B	1225	ALA	Peptide
1	B	286	VAL	Peptide
1	B	287	VAL	Peptide
1	B	293	ARG	Peptide
1	B	295	GLY	Peptide
1	B	302	LEU	Peptide
1	B	376	ARG	Peptide
1	B	377	GLY	Peptide
1	B	414	VAL	Peptide
1	B	744	CYS	Peptide

5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	7464	1485	5141	2123	2
1	B	7485	1485	5190	2173	2
All	All	14949	2970	10331	4296	4

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 170.

All (4296) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:263:PHE:HB2	1:A:415:MET:CB	1.50	1.38
1:A:456:ILE:C	1:A:459:LYS:HD3	1.51	1.35
1:B:459:LYS:C	1:B:462:LYS:HE3	1.48	1.35
1:A:32:LEU:CD1	1:A:445:PHE:HA	1.54	1.35

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:18:TYR:O	1:B:181:LYS:NZ	1.59	1.32
1:B:462:LYS:HE2	1:B:462:LYS:N	1.44	1.30
1:B:283:GLU:O	1:B:306:LYS:HB2	1.25	1.28
1:A:20:GLU:OE1	1:A:20:GLU:N	1.66	1.28
1:B:239:VAL:HA	1:B:282:TRP:O	1.32	1.27
1:B:19:ALA:HA	1:B:218:TRP:CZ3	1.71	1.26
1:A:181:LYS:CE	1:A:219:LYS:HB3	1.66	1.24
1:B:249:LYS:CD	1:B:416:LEU:HG	1.65	1.24
1:A:194:HIS:O	1:A:210:ASN:HB2	1.35	1.24
1:A:263:PHE:CB	1:A:415:MET:HB3	1.67	1.23
1:A:454:GLY:O	1:A:459:LYS:NZ	1.73	1.22
1:B:249:LYS:HD2	1:B:416:LEU:CG	1.69	1.22
1:A:133:THR:HB	1:A:150:THR:O	1.38	1.22
1:B:104:GLU:O	1:B:108:ARG:HB2	1.07	1.21
1:A:478:GLU:OE2	1:A:555:LEU:HD21	1.38	1.20
1:A:391:ARG:CG	1:A:398:TRP:HA	1.69	1.20
1:B:617:THR:O	1:B:621:LEU:CB	1.89	1.20
1:B:11:ILE:HG21	1:B:62:PRO:HD3	1.22	1.19
1:B:447:ASN:O	1:B:451:LYS:NZ	1.75	1.19
1:A:20:GLU:HB2	1:A:217:SER:HB2	1.23	1.19
1:A:41:PRO:HD3	1:A:207:ASN:ND2	1.58	1.19
1:B:224:MET:SD	1:B:288:GLN:NE2	2.12	1.19
1:B:18:TYR:CB	1:B:46:LEU:HG	1.72	1.18
1:B:516:LEU:HA	1:B:519:ILE:HD11	1.24	1.18
1:A:15:CYS:HB2	1:A:221:VAL:O	1.44	1.17
1:A:16:SER:O	1:A:221:VAL:HB	1.41	1.17
1:B:399:VAL:HA	1:B:419:GLY:HA2	1.24	1.17
1:B:184:LEU:HB3	1:B:193:LEU:HB3	1.26	1.17
1:B:186:PRO:HD2	1:B:191:GLN:O	1.44	1.16
1:B:400:HIS:HB2	1:B:420:THR:CA	1.76	1.16
1:B:451:LYS:N	1:B:451:LYS:HZ3	1.40	1.16
1:B:477:LEU:O	1:B:480:LEU:HD22	1.41	1.16
1:B:298:TYR:HA	1:B:381:LEU:HA	1.26	1.15
1:B:400:HIS:CB	1:B:420:THR:HA	1.74	1.15
1:B:448:ASP:O	1:B:451:LYS:HE3	1.46	1.15
1:B:242:LEU:HB2	1:B:430:PHE:CD2	1.79	1.15
1:B:18:TYR:HB2	1:B:46:LEU:HG	1.17	1.15
1:A:454:GLY:N	1:A:518:GLN:OE1	1.78	1.15
1:B:302:LEU:HG	1:B:367:ILE:HG13	1.29	1.15
1:B:390:LEU:HD11	1:B:399:VAL:HB	1.26	1.14
1:B:133:THR:OG1	1:B:156:ASN:OD1	1.65	1.14

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:49:PRO:HG3	1:A:291:PRO:CA	1.76	1.14
1:B:242:LEU:HA	1:B:431:ALA:O	1.46	1.14
1:A:281:LEU:HB2	1:A:308:LEU:HD21	1.21	1.14
1:A:401:SER:HA	1:A:418:ILE:HB	1.21	1.14
1:B:39:VAL:HG11	1:B:195:ALA:HB1	1.23	1.14
1:B:234:LEU:HG	1:B:384:ARG:HD3	1.14	1.14
1:B:130:LYS:HG3	1:B:153:GLU:H	1.12	1.13
1:B:140:ALA:HB3	1:B:143:GLU:HB2	1.23	1.13
1:B:391:ARG:HH12	1:B:393:LEU:HD13	1.11	1.12
1:A:14:ILE:HG23	1:A:60:LEU:HD22	1.31	1.12
1:A:405:PRO:HB3	1:A:415:MET:O	1.47	1.12
1:A:39:VAL:O	1:A:207:ASN:N	1.82	1.12
1:A:316:ALA:HB3	1:A:368:PHE:CZ	1.85	1.12
1:B:243:PHE:HB3	1:B:431:ALA:HB3	1.30	1.12
1:A:14:ILE:CG1	1:A:59:LYS:HA	1.79	1.12
1:B:610:ILE:O	1:B:614:GLU:CB	1.98	1.12
1:A:191:GLN:HB2	1:A:192:PRO:HD2	1.12	1.11
1:A:316:ALA:HB3	1:A:368:PHE:HZ	0.96	1.11
1:B:257:ARG:HB3	1:B:259:LYS:HZ3	1.00	1.11
1:A:391:ARG:HG3	1:A:398:TRP:CA	1.80	1.11
1:A:391:ARG:HA	1:A:399:VAL:HG23	1.30	1.11
1:B:60:LEU:HD22	1:B:60:LEU:H	1.13	1.11
1:B:110:LEU:HD12	1:B:113:THR:HG22	1.28	1.11
1:A:54:ARG:HA	1:A:57:LEU:HD21	1.33	1.11
1:B:453:LEU:HD11	1:B:515:ILE:HD12	1.28	1.11
1:A:388:VAL:N	1:A:430:PHE:O	1.81	1.11
1:A:19:ALA:HB3	1:A:24:ASN:HA	1.11	1.10
1:B:138:LEU:HG	1:B:148:ARG:HB3	1.31	1.10
1:B:224:MET:HE3	1:B:295:GLY:H	1.09	1.10
1:B:19:ALA:N	1:B:25:GLY:O	1.83	1.10
1:B:106:GLU:HG3	1:B:107:ASN:OD1	1.50	1.10
1:B:111:LEU:HB2	1:B:226:TRP:HZ3	1.08	1.10
1:B:181:LYS:HB3	1:B:217:SER:HB2	1.17	1.10
1:B:392:HIS:NE2	1:B:394:CYS:SG	2.25	1.10
1:A:247:GLN:O	1:A:249:LYS:N	1.84	1.10
1:A:23:THR:HG21	1:A:195:ALA:HB3	1.30	1.10
1:B:38:VAL:HG11	1:B:206:CYS:HB3	1.31	1.10
1:B:241:ARG:HH22	1:B:433:VAL:HG23	1.01	1.10
1:A:448:ASP:HA	1:A:451:LYS:HD2	1.21	1.10
1:B:242:LEU:HB2	1:B:430:PHE:HD2	0.98	1.10
1:B:406:ILE:H	1:B:415:MET:HB2	1.05	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:462:LYS:CE	1:B:462:LYS:H	1.63	1.10
1:A:111:LEU:HD22	1:A:226:TRP:HA	1.11	1.09
1:A:450:SER:CB	1:A:514:ASN:HB3	1.80	1.09
1:B:1472:SER:O	1:B:1476:LYS:CB	2.01	1.09
1:A:27:ILE:HD11	1:A:39:VAL:HG22	1.27	1.09
1:A:57:LEU:HB3	1:A:126:LEU:HD12	1.25	1.09
1:A:63:MET:HE3	1:A:120:VAL:HG11	1.25	1.09
1:A:364:ILE:O	1:A:367:ILE:HG12	1.51	1.09
1:B:18:TYR:OH	1:B:24:ASN:HB2	1.51	1.09
1:A:14:ILE:CB	1:A:59:LYS:HA	1.82	1.09
1:B:143:GLU:HG3	1:B:210:ASN:HB3	1.15	1.09
1:B:286:VAL:HA	1:B:304:ARG:NE	1.66	1.08
1:A:56:CYS:HA	1:A:125:HIS:NE2	1.69	1.08
1:A:298:TYR:H	1:A:301:SER:HB2	1.07	1.08
1:A:387:TYR:HA	1:A:431:ALA:HA	1.31	1.08
1:A:1211:HIS:O	1:A:1215:LEU:CB	2.01	1.08
1:A:240:VAL:HA	1:A:434:PRO:HA	1.14	1.08
1:A:477:LEU:HD11	1:A:552:ILE:HD12	1.33	1.08
1:B:281:LEU:HB3	1:B:308:LEU:HD23	1.18	1.08
1:B:256:HIS:O	1:B:258:LYS:N	1.87	1.08
1:B:458:GLY:O	1:B:462:LYS:NZ	1.85	1.08
1:B:516:LEU:HA	1:B:519:ILE:CD1	1.84	1.08
1:A:63:MET:HA	1:A:160:TRP:HE1	1.05	1.07
1:A:108:ARG:HB2	1:A:108:ARG:CZ	1.68	1.07
1:A:1380:LEU:O	1:A:1384:CYS:N	1.86	1.07
1:B:224:MET:CE	1:B:295:GLY:H	1.66	1.07
1:B:437:PRO:O	1:B:441:ARG:NE	1.87	1.07
1:B:459:LYS:HA	1:B:462:LYS:HD3	1.31	1.07
1:A:58:PHE:HE2	1:A:123:LEU:HB3	1.12	1.07
1:A:70:LYS:HA	1:A:73:TRP:CE2	1.88	1.07
1:A:364:ILE:HA	1:A:367:ILE:CD1	1.84	1.07
1:A:477:LEU:HA	1:A:480:LEU:CD1	1.83	1.07
1:B:19:ALA:HB3	1:B:25:GLY:N	1.69	1.07
1:B:371:ASP:OD2	1:B:389:ARG:HB2	1.53	1.07
1:A:239:VAL:HG13	1:A:435:VAL:HB	1.32	1.07
1:B:104:GLU:O	1:B:108:ARG:CB	2.01	1.07
1:B:111:LEU:HB2	1:B:226:TRP:CZ3	1.87	1.07
1:B:181:LYS:HE2	1:B:219:LYS:H	1.16	1.07
1:B:477:LEU:HB2	1:B:555:LEU:HD13	1.08	1.07
1:A:19:ALA:HB3	1:A:24:ASN:CA	1.85	1.06
1:A:963:GLU:O	1:A:967:VAL:CB	2.02	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:20:GLU:HG3	1:A:181:LYS:HE3	1.35	1.06
1:A:242:LEU:HD11	1:A:251:LEU:HD21	1.35	1.06
1:B:355:LEU:HB2	1:B:417:LYS:HE3	1.33	1.06
1:A:41:PRO:HD3	1:A:207:ASN:HD21	0.92	1.06
1:A:443:LEU:HA	1:A:509:LEU:HD21	1.36	1.06
1:B:58:PHE:HB3	1:B:123:LEU:HD12	1.37	1.06
1:B:510:MET:HA	1:B:513:GLN:HE22	1.02	1.06
1:A:369:GLU:HB3	1:A:393:LEU:HD11	1.33	1.06
1:B:14:ILE:HG22	1:B:57:LEU:HD21	1.37	1.06
1:B:249:LYS:HG2	1:B:264:LEU:CD1	1.86	1.06
1:B:477:LEU:HB2	1:B:555:LEU:CD1	1.84	1.06
1:B:507:GLN:OE1	1:B:507:GLN:N	1.88	1.06
1:A:14:ILE:HD12	1:A:224:MET:HB3	1.37	1.06
1:A:66:TYR:HA	1:A:99:GLU:OE1	1.55	1.06
1:A:181:LYS:HG3	1:A:219:LYS:HA	1.38	1.06
1:A:181:LYS:NZ	1:A:219:LYS:HB3	1.70	1.06
1:B:194:HIS:O	1:B:209:VAL:HG13	1.55	1.06
1:B:243:PHE:HB3	1:B:431:ALA:CB	1.86	1.06
1:A:264:LEU:O	1:A:416:LEU:N	1.88	1.05
1:A:281:LEU:HB3	1:A:435:VAL:HG21	1.37	1.05
1:A:477:LEU:CA	1:A:480:LEU:HD11	1.84	1.05
1:B:180:ASP:O	1:B:219:LYS:HA	1.56	1.05
1:B:195:ALA:HA	1:B:209:VAL:HG22	1.31	1.05
1:B:465:ILE:O	1:B:469:GLU:HB3	1.56	1.05
1:A:227:SER:HA	1:A:230:LYS:HD2	1.38	1.05
1:A:446:ALA:HB2	1:A:509:LEU:HG	1.38	1.05
1:B:244:HIS:HB3	1:B:430:PHE:CE1	1.91	1.05
1:B:313:TYR:CE2	1:B:361:GLY:HA3	1.92	1.05
1:A:18:TYR:HB2	1:A:219:LYS:HE3	1.37	1.05
1:A:58:PHE:CE2	1:A:123:LEU:HB3	1.92	1.05
1:B:15:CYS:HB3	1:B:220:ILE:HD11	1.10	1.05
1:B:18:TYR:HB2	1:B:46:LEU:CG	1.85	1.05
1:B:443:LEU:HA	1:B:509:LEU:HD21	1.08	1.05
1:B:451:LYS:HE2	1:B:452:VAL:N	1.70	1.05
1:A:32:LEU:HD11	1:A:445:PHE:HA	1.38	1.05
1:A:105:THR:O	1:A:108:ARG:NH2	1.90	1.05
1:A:387:TYR:HA	1:A:431:ALA:CA	1.86	1.05
1:A:255:GLU:HA	1:A:255:GLU:OE2	1.50	1.05
1:A:516:LEU:HB2	1:A:520:PHE:CZ	1.92	1.05
1:A:166:PHE:HB2	1:A:182:VAL:CA	1.86	1.04
1:A:285:GLU:O	1:A:304:ARG:HB2	1.57	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:257:ARG:HH12	1:B:408:LYS:HG3	1.14	1.04
1:B:442:ASP:OD1	1:B:442:ASP:N	1.80	1.04
1:A:117:TYR:HE1	1:A:170:ARG:HG2	1.19	1.04
1:A:402:THR:H	1:A:418:ILE:HG22	1.13	1.04
1:A:298:TYR:H	1:A:301:SER:CB	1.71	1.04
1:A:477:LEU:HB2	1:A:555:LEU:CB	1.86	1.04
1:A:68:ALA:HB1	1:A:95:ALA:HB1	1.07	1.03
1:A:68:ALA:HB1	1:A:95:ALA:CB	1.88	1.03
1:A:219:LYS:O	1:A:219:LYS:NZ	1.90	1.03
1:B:263:PHE:CD2	1:B:415:MET:HA	1.94	1.03
1:A:68:ALA:CB	1:A:95:ALA:HB1	1.88	1.03
1:A:406:ILE:HG13	1:A:417:LYS:HD2	1.37	1.03
1:B:390:LEU:HD11	1:B:399:VAL:CB	1.87	1.03
1:B:462:LYS:N	1:B:462:LYS:CE	2.18	1.03
1:A:33:VAL:CG1	1:A:452:VAL:HG22	1.88	1.03
1:A:388:VAL:CG2	1:A:432:ILE:HG12	1.87	1.03
1:A:478:GLU:HG2	1:A:555:LEU:HD11	1.40	1.03
1:A:474:THR:HG22	1:A:552:ILE:HD11	1.03	1.03
1:A:49:PRO:CG	1:A:291:PRO:HA	1.89	1.02
1:A:139:PRO:HA	1:A:147:MET:HA	1.04	1.02
1:A:155:GLY:O	1:A:156:ASN:ND2	1.90	1.02
1:B:221:VAL:HG21	1:B:292:CYS:HB2	1.36	1.02
1:B:249:LYS:HG3	1:B:265:ARG:N	1.73	1.02
1:B:286:VAL:HG12	1:B:288:GLN:HG3	1.35	1.02
1:A:166:PHE:CB	1:A:182:VAL:HA	1.90	1.02
1:A:166:PHE:CE2	1:A:217:SER:HA	1.95	1.02
1:A:253:CYS:HB2	1:A:307:HIS:CE1	1.94	1.02
1:A:263:PHE:CG	1:A:415:MET:HB3	1.93	1.02
1:A:307:HIS:O	1:A:311:GLY:N	1.93	1.02
1:A:388:VAL:HG21	1:A:432:ILE:HG12	1.05	1.02
1:B:47:ASN:O	1:B:291:PRO:HG3	1.57	1.02
1:B:57:LEU:HD22	1:B:126:LEU:HD21	1.38	1.02
1:A:257:ARG:HB3	1:A:259:LYS:HE3	1.40	1.02
1:A:316:ALA:CB	1:A:355:LEU:HG	1.90	1.02
1:A:446:ALA:HA	1:A:515:ILE:HD11	1.03	1.02
1:A:510:MET:HA	1:A:515:ILE:HB	1.41	1.02
1:B:140:ALA:HB1	1:B:208:GLU:CD	1.84	1.02
1:A:263:PHE:CD2	1:A:415:MET:HB3	1.93	1.01
1:A:364:ILE:CA	1:A:367:ILE:HD11	1.90	1.01
1:A:448:ASP:HA	1:A:451:LYS:CD	1.89	1.01
1:A:298:TYR:HE1	1:A:381:LEU:HB3	1.18	1.01

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:72:PHE:HD2	1:A:92:LEU:HB3	1.18	1.01
1:A:856:ASP:O	1:A:860:ASN:CB	2.08	1.01
1:A:1126:VAL:O	1:A:1130:GLU:N	1.93	1.01
1:B:70:LYS:O	1:B:74:LYS:HG3	1.58	1.01
1:A:14:ILE:HG12	1:A:59:LYS:HA	1.36	1.01
1:A:191:GLN:HB2	1:A:192:PRO:CD	1.90	1.01
1:A:257:ARG:CB	1:A:259:LYS:HE3	1.91	1.01
1:A:263:PHE:HB2	1:A:415:MET:HB3	1.04	1.01
1:B:15:CYS:HB3	1:B:220:ILE:CD1	1.91	1.01
1:B:148:ARG:NH2	1:B:150:THR:HG21	1.75	1.01
1:A:15:CYS:HB3	1:A:223:PHE:HB2	1.42	1.01
1:A:69:GLN:HG3	1:A:99:GLU:OE1	1.61	1.01
1:A:127:LYS:NZ	1:A:128:SER:OG	1.92	1.01
1:A:247:GLN:NE2	1:A:403:ASN:OD1	1.93	1.01
1:B:137:ARG:HA	1:B:147:MET:HA	1.43	1.01
1:A:450:SER:HB3	1:A:514:ASN:CB	1.89	1.00
1:A:198:HIS:HB2	1:A:208:GLU:CD	1.84	1.00
1:B:55:ASP:HA	1:B:127:LYS:HG3	1.03	1.00
1:B:400:HIS:N	1:B:419:GLY:C	2.19	1.00
1:A:230:LYS:O	1:A:232:ASP:N	1.92	1.00
1:A:265:ARG:HA	1:A:416:LEU:CD1	1.90	1.00
1:B:24:ASN:O	1:B:41:PRO:HA	1.57	1.00
1:B:654:LYS:O	1:B:658:ASN:N	1.95	1.00
1:B:1473:ILE:O	1:B:1477:TYR:CB	2.09	1.00
1:A:368:PHE:C	1:A:393:LEU:HD22	1.86	1.00
1:B:241:ARG:NH2	1:B:433:VAL:HG23	1.75	1.00
1:A:177:VAL:HG12	1:A:178:ILE:HG12	1.43	1.00
1:B:516:LEU:O	1:B:519:ILE:HG12	1.61	1.00
1:A:139:PRO:HA	1:A:147:MET:CA	1.91	1.00
1:A:458:GLY:CA	1:A:459:LYS:HE3	1.91	1.00
1:A:459:LYS:CD	1:A:459:LYS:N	2.24	1.00
1:B:58:PHE:HB3	1:B:123:LEU:CD1	1.90	1.00
1:B:404:ILE:HD13	1:B:417:LYS:H	1.27	1.00
1:A:298:TYR:CE1	1:A:381:LEU:HB3	1.96	1.00
1:B:353:TYR:HD1	1:B:422:PRO:HB3	1.25	1.00
1:A:249:LYS:HA	1:A:264:LEU:HD21	1.40	0.99
1:A:410:GLU:HB3	1:A:413:PRO:HB3	1.43	0.99
1:A:447:ASN:O	1:A:450:SER:OG	1.77	0.99
1:A:402:THR:H	1:A:418:ILE:CG2	1.75	0.99
1:B:37:CYS:SG	1:B:209:VAL:HB	2.01	0.99
1:B:298:TYR:HB3	1:B:381:LEU:HD22	1.42	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:181:LYS:HG3	1:A:219:LYS:CA	1.92	0.99
1:A:453:LEU:HB2	1:A:522:LEU:HD21	1.40	0.99
1:B:400:HIS:HB2	1:B:420:THR:HA	1.03	0.99
1:B:406:ILE:O	1:B:408:LYS:HD2	1.62	0.99
1:A:13:ASP:OD1	1:A:226:TRP:N	1.96	0.99
1:B:14:ILE:CG2	1:B:57:LEU:HD21	1.92	0.99
1:A:264:LEU:CB	1:A:418:ILE:HD11	1.91	0.99
1:A:474:THR:HG22	1:A:552:ILE:CD1	1.92	0.99
1:A:477:LEU:HD12	1:A:555:LEU:HD12	1.40	0.99
1:A:315:ALA:HB2	1:A:358:VAL:HG22	1.38	0.99
1:B:462:LYS:HD2	1:B:463:GLY:N	1.78	0.99
1:A:111:LEU:CD2	1:A:226:TRP:HA	1.92	0.98
1:A:457:ALA:HB2	1:A:522:LEU:CB	1.92	0.98
1:A:518:GLN:HE22	1:A:522:LEU:HD11	1.26	0.98
1:B:251:LEU:HD11	1:B:282:TRP:HH2	1.28	0.98
1:A:249:LYS:NZ	1:A:402:THR:O	1.96	0.98
1:A:288:GLN:HA	1:A:288:GLN:HE21	1.25	0.98
1:A:1374:HIS:O	1:A:1378:VAL:N	1.95	0.98
1:B:138:LEU:HD11	1:B:148:ARG:HD2	1.43	0.98
1:B:443:LEU:CA	1:B:509:LEU:HD21	1.93	0.98
1:A:299:TRP:O	1:A:300:ASN:ND2	1.95	0.98
1:B:241:ARG:NH2	1:B:241:ARG:O	1.96	0.98
1:B:507:GLN:NE2	1:B:563:SER:HA	1.78	0.98
1:A:167:TYR:CE2	1:A:181:LYS:HB2	1.99	0.98
1:B:850:GLN:O	1:B:852:PHE:N	1.96	0.98
1:B:414:VAL:HG23	1:B:415:MET:HE1	1.45	0.98
1:A:63:MET:HA	1:A:160:TRP:NE1	1.77	0.97
1:B:403:ASN:OD1	1:B:416:LEU:HD11	1.62	0.97
1:A:111:LEU:HD22	1:A:226:TRP:CA	1.93	0.97
1:A:283:GLU:HG2	1:A:306:LYS:HB3	1.45	0.97
1:B:598:LEU:O	1:B:602:ASN:N	1.96	0.97
1:A:198:HIS:HB2	1:A:208:GLU:CG	1.93	0.97
1:A:249:LYS:HB2	1:A:430:PHE:HZ	1.27	0.97
1:A:453:LEU:O	1:A:522:LEU:HG	1.65	0.97
1:B:65:ARG:NH2	1:B:106:GLU:OE2	1.98	0.97
1:B:249:LYS:HD2	1:B:416:LEU:HG	0.97	0.97
1:B:477:LEU:CB	1:B:555:LEU:HD13	1.94	0.97
1:B:132:LEU:H	1:B:132:LEU:HD12	1.29	0.97
1:B:243:PHE:CB	1:B:431:ALA:HB3	1.92	0.97
1:B:510:MET:HG2	1:B:516:LEU:CG	1.94	0.97
1:A:110:LEU:O	1:A:110:LEU:HG	1.59	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:253:CYS:SG	1:A:307:HIS:NE2	2.38	0.97
1:A:385:ASN:HA	1:A:432:ILE:O	1.62	0.97
1:B:263:PHE:HB2	1:B:415:MET:O	1.64	0.97
1:B:400:HIS:O	1:B:418:ILE:HG22	1.64	0.97
1:B:503:ASN:HB2	1:B:506:ARG:NH1	1.79	0.97
1:A:65:ARG:HA	1:A:160:TRP:HH2	1.30	0.97
1:A:617:THR:O	1:A:621:LEU:CB	2.13	0.97
1:B:55:ASP:HA	1:B:127:LYS:CG	1.92	0.97
1:A:235:LYS:HG3	1:A:238:ASP:HB2	1.45	0.97
1:A:450:SER:HB3	1:A:514:ASN:HB3	0.99	0.97
1:A:481:VAL:O	1:A:484:VAL:HG12	1.64	0.97
1:B:369:GLU:OE2	1:B:391:ARG:NH2	1.98	0.97
1:A:14:ILE:HB	1:A:58:PHE:O	1.65	0.97
1:A:446:ALA:HA	1:A:515:ILE:CD1	1.95	0.96
1:B:147:MET:HE2	1:B:186:PRO:HG3	1.46	0.96
1:A:167:TYR:HE2	1:A:181:LYS:HB2	1.29	0.96
1:A:283:GLU:OE1	1:A:306:LYS:NZ	1.97	0.96
1:A:14:ILE:HG23	1:A:60:LEU:CD2	1.95	0.96
1:A:139:PRO:CA	1:A:147:MET:HA	1.95	0.96
1:A:263:PHE:HB2	1:A:415:MET:HB2	1.47	0.96
1:B:459:LYS:C	1:B:462:LYS:CE	2.39	0.96
1:A:19:ALA:CB	1:A:24:ASN:HA	1.95	0.96
1:A:38:VAL:HB	1:A:206:CYS:HB3	1.48	0.96
1:B:45:ASP:N	1:B:48:ASN:O	1.96	0.96
1:B:224:MET:HE2	1:B:294:GLY:HA2	1.45	0.96
1:B:313:TYR:O	1:B:358:VAL:HG22	1.62	0.96
1:B:414:VAL:HG23	1:B:415:MET:SD	2.06	0.96
1:A:250:PHE:CZ	1:A:280:ALA:HB1	2.01	0.96
1:A:302:LEU:HD11	1:A:393:LEU:HD21	1.46	0.96
1:B:14:ILE:HB	1:B:223:PHE:CD2	2.00	0.96
1:A:474:THR:CG2	1:A:552:ILE:HD11	1.96	0.96
1:B:286:VAL:CG1	1:B:288:GLN:HG3	1.96	0.96
1:A:255:GLU:OE2	1:A:260:GLN:HA	1.64	0.96
1:B:9:LEU:HD21	1:B:115:ILE:HB	1.43	0.96
1:B:18:TYR:HA	1:B:26:PHE:HD1	1.30	0.96
1:B:244:HIS:NE2	1:B:249:LYS:HB3	1.79	0.96
1:B:298:TYR:HB3	1:B:381:LEU:CD2	1.96	0.96
1:A:33:VAL:HG12	1:A:452:VAL:HG22	1.47	0.96
1:A:1210:ALA:O	1:A:1214:VAL:CB	2.13	0.96
1:A:1375:ILE:O	1:A:1380:LEU:N	1.98	0.96
1:A:132:LEU:HA	1:A:151:LEU:CD2	1.94	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:116:GLN:OE1	1:B:174:ASP:N	1.98	0.95
1:B:286:VAL:HA	1:B:304:ARG:HE	1.24	0.95
1:B:361:GLY:O	1:B:366:SER:OG	1.83	0.95
1:B:398:TRP:O	1:B:420:THR:N	1.99	0.95
1:A:446:ALA:CA	1:A:515:ILE:HD11	1.96	0.95
1:B:184:LEU:CB	1:B:193:LEU:HB3	1.96	0.95
1:B:504:ARG:HD3	1:B:504:ARG:C	1.90	0.95
1:B:1378:VAL:O	1:B:1382:ALA:N	1.99	0.95
1:A:11:ILE:O	1:A:111:LEU:HD23	1.65	0.95
1:A:456:ILE:CA	1:A:459:LYS:HD3	1.96	0.95
1:A:306:LYS:HB2	1:A:313:TYR:CE1	2.01	0.95
1:A:455:SER:O	1:A:459:LYS:HG3	1.66	0.95
1:A:457:ALA:CB	1:A:522:LEU:HA	1.96	0.95
1:B:446:ALA:HB1	1:B:513:GLN:NE2	1.80	0.95
1:A:181:LYS:HZ3	1:A:219:LYS:HB3	1.32	0.95
1:A:353:TYR:CD2	1:A:397:THR:HG21	2.01	0.95
1:A:511:ARG:HA	1:A:516:LEU:CG	1.97	0.95
1:B:56:CYS:HA	1:B:125:HIS:NE2	1.81	0.95
1:B:363:ASP:O	1:B:366:SER:HB2	1.65	0.95
1:B:510:MET:SD	1:B:563:SER:OG	2.24	0.95
1:B:197:SER:HA	1:B:207:ASN:OD1	1.65	0.95
1:B:484:VAL:HG21	1:B:562:HIS:HB3	1.48	0.95
1:A:459:LYS:N	1:A:459:LYS:HD2	1.80	0.95
1:A:249:LYS:HB2	1:A:430:PHE:CZ	2.02	0.94
1:A:857:LYS:O	1:A:861:LYS:CB	2.14	0.94
1:A:9:LEU:HD12	1:A:9:LEU:H	1.32	0.94
1:B:55:ASP:CA	1:B:127:LYS:HG3	1.95	0.94
1:B:477:LEU:HA	1:B:480:LEU:CD1	1.97	0.94
1:B:142:LEU:O	1:B:198:HIS:ND1	2.01	0.94
1:A:48:ASN:N	1:A:49:PRO:HD2	1.80	0.94
1:B:15:CYS:CB	1:B:220:ILE:HD11	1.98	0.94
1:B:1129:SER:O	1:B:1133:VAL:N	1.99	0.94
1:A:14:ILE:HG12	1:A:59:LYS:CA	1.96	0.94
1:B:262:VAL:HG11	1:B:314:LEU:HB2	1.47	0.94
1:B:302:LEU:HG	1:B:367:ILE:CG1	1.96	0.94
1:B:448:ASP:HA	1:B:451:LYS:HD3	1.45	0.94
1:A:166:PHE:HB2	1:A:182:VAL:HA	0.96	0.94
1:A:315:ALA:HB2	1:A:358:VAL:CG2	1.97	0.94
1:B:13:ASP:HA	1:B:226:TRP:HB3	1.49	0.94
1:B:63:MET:SD	1:B:64:ASN:N	2.41	0.94
1:B:257:ARG:HB3	1:B:259:LYS:NZ	1.82	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:414:VAL:HG23	1:B:415:MET:CE	1.98	0.94
1:B:449:ALA:CB	1:B:515:ILE:HD13	1.97	0.94
1:A:234:LEU:HD21	1:A:432:ILE:HG21	1.50	0.94
1:A:32:LEU:HG	1:A:448:ASP:OD2	1.66	0.94
1:A:65:ARG:HA	1:A:160:TRP:CH2	2.03	0.94
1:B:130:LYS:HE3	1:B:151:LEU:HB3	1.50	0.93
1:B:1474:LEU:O	1:B:1478:VAL:CB	2.16	0.93
1:B:504:ARG:HD3	1:B:504:ARG:O	1.68	0.93
1:A:436:SER:O	1:A:440:VAL:HG23	1.69	0.93
1:B:399:VAL:CA	1:B:419:GLY:HA2	1.99	0.93
1:B:453:LEU:HD12	1:B:518:GLN:CB	1.96	0.93
1:A:13:ASP:C	1:A:14:ILE:HG13	1.92	0.93
1:B:451:LYS:NZ	1:B:451:LYS:N	2.15	0.93
1:B:477:LEU:HA	1:B:480:LEU:HD11	1.50	0.93
1:B:510:MET:HB2	1:B:515:ILE:CG2	1.97	0.93
1:A:240:VAL:CA	1:A:434:PRO:HA	1.97	0.93
1:A:364:ILE:HA	1:A:367:ILE:HD11	0.95	0.93
1:A:401:SER:HA	1:A:418:ILE:CB	1.98	0.93
1:A:264:LEU:HD13	1:A:418:ILE:CD1	1.98	0.93
1:B:63:MET:HG3	1:B:160:TRP:CD1	2.04	0.93
1:B:132:LEU:HB2	1:B:149:VAL:CG1	1.98	0.93
1:B:257:ARG:HG3	1:B:257:ARG:O	1.69	0.93
1:A:316:ALA:HB1	1:A:355:LEU:HG	1.46	0.93
1:B:506:ARG:HB2	1:B:507:GLN:OE1	1.69	0.93
1:A:523:LEU:HD23	1:A:524:GLN:NE2	1.84	0.93
1:A:1096:ALA:O	1:A:1100:VAL:N	2.02	0.93
1:A:185:ASN:N	1:A:185:ASN:HD22	1.64	0.92
1:B:287:VAL:HG13	1:B:289:HIS:H	1.32	0.92
1:B:353:TYR:HD1	1:B:422:PRO:CB	1.82	0.92
1:B:407:ASP:HB2	1:B:413:PRO:HA	1.51	0.92
1:B:510:MET:HG2	1:B:516:LEU:HG	1.49	0.92
1:A:20:GLU:HG3	1:A:181:LYS:CE	1.98	0.92
1:A:1057:GLY:O	1:A:1059:THR:N	2.01	0.92
1:B:11:ILE:HG13	1:B:113:THR:HG23	1.51	0.92
1:B:130:LYS:HD3	1:B:153:GLU:OE1	1.70	0.92
1:B:373:THR:CG2	1:B:389:ARG:HE	1.82	0.92
1:B:400:HIS:H	1:B:419:GLY:CA	1.81	0.92
1:B:400:HIS:H	1:B:419:GLY:C	1.76	0.92
1:B:451:LYS:HE2	1:B:452:VAL:H	1.35	0.92
1:A:72:PHE:CD2	1:A:92:LEU:HB3	2.03	0.92
1:A:315:ALA:HB2	1:A:358:VAL:HG13	1.51	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:523:LEU:HD23	1:A:524:GLN:HE22	1.34	0.92
1:B:19:ALA:HA	1:B:218:TRP:HZ3	1.12	0.92
1:A:716:ALA:O	1:A:718:ASP:N	2.02	0.92
1:A:448:ASP:CA	1:A:451:LYS:HD2	2.00	0.92
1:B:94:HIS:O	1:B:98:LEU:HB2	1.70	0.92
1:B:404:ILE:HD11	1:B:418:ILE:H	1.34	0.92
1:A:23:THR:HG21	1:A:195:ALA:CB	1.99	0.92
1:A:247:GLN:O	1:A:249:LYS:HG2	1.70	0.92
1:A:446:ALA:CB	1:A:509:LEU:HG	1.99	0.92
1:B:196:SER:O	1:B:207:ASN:ND2	2.02	0.92
1:A:447:ASN:C	1:A:450:SER:HG	1.78	0.92
1:B:66:TYR:HE1	1:B:100:LYS:HA	1.32	0.92
1:B:181:LYS:HE2	1:B:219:LYS:N	1.83	0.92
1:B:484:VAL:CG2	1:B:562:HIS:HB3	2.00	0.92
1:A:368:PHE:CD1	1:A:392:HIS:HA	2.05	0.91
1:B:516:LEU:H	1:B:516:LEU:HD12	1.35	0.91
1:A:62:PRO:HD3	1:A:110:LEU:HD22	1.51	0.91
1:A:315:ALA:CB	1:A:358:VAL:HG13	2.00	0.91
1:B:235:LYS:NZ	1:B:238:ASP:OD2	2.02	0.91
1:A:367:ILE:C	1:A:393:LEU:HB2	1.94	0.91
1:A:389:ARG:HH12	1:A:428:GLU:N	1.69	0.91
1:B:69:GLN:HB3	1:B:100:LYS:NZ	1.84	0.91
1:B:1377:LEU:O	1:B:1381:LEU:N	2.03	0.91
1:A:11:ILE:HD11	1:A:115:ILE:HD11	1.52	0.91
1:A:11:ILE:CG2	1:A:110:LEU:HD21	2.01	0.91
1:B:234:LEU:HG	1:B:384:ARG:CD	1.98	0.91
1:A:16:SER:CB	1:A:53:PHE:HZ	1.83	0.91
1:A:20:GLU:OE2	1:A:181:LYS:HG2	1.70	0.91
1:A:33:VAL:HG21	1:A:451:LYS:HG3	1.50	0.91
1:A:167:TYR:HD2	1:A:180:ASP:OD1	1.54	0.91
1:A:249:LYS:O	1:A:264:LEU:HD11	1.71	0.91
1:B:195:ALA:CA	1:B:209:VAL:HG22	2.00	0.91
1:A:20:GLU:CB	1:A:217:SER:HB2	2.00	0.91
1:B:453:LEU:HD12	1:B:518:GLN:HB3	1.52	0.91
1:A:139:PRO:C	1:A:148:ARG:HD3	1.95	0.91
1:A:564:GLN:HE22	1:A:570:ASN:CB	1.83	0.91
1:B:130:LYS:HD3	1:B:153:GLU:CD	1.96	0.91
1:B:38:VAL:CG1	1:B:206:CYS:HB3	2.00	0.91
1:A:302:LEU:CD1	1:A:393:LEU:HD21	2.01	0.90
1:A:581:MET:O	1:A:585:ILE:N	2.03	0.90
1:A:1371:LEU:O	1:A:1375:ILE:N	2.04	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:314:LEU:HD11	1:A:316:ALA:HB2	1.52	0.90
1:A:447:ASN:N	1:A:513:GLN:OE1	2.03	0.90
1:B:181:LYS:HB3	1:B:217:SER:CB	2.01	0.90
1:B:484:VAL:HG13	1:B:562:HIS:ND1	1.86	0.90
1:A:454:GLY:C	1:A:459:LYS:HZ1	1.79	0.90
1:B:54:ARG:HD3	1:B:54:ARG:H	1.36	0.90
1:A:117:TYR:CE1	1:A:170:ARG:HG2	2.07	0.90
1:A:181:LYS:HG3	1:A:219:LYS:CB	2.02	0.90
1:A:239:VAL:HG21	1:A:308:LEU:HD23	1.51	0.90
1:A:243:PHE:C	1:A:430:PHE:HD1	1.78	0.90
1:A:264:LEU:HB2	1:A:418:ILE:HD11	1.51	0.90
1:B:353:TYR:CD1	1:B:422:PRO:HB3	2.06	0.90
1:B:441:ARG:NE	1:B:441:ARG:H	1.67	0.90
1:B:240:VAL:HG12	1:B:434:PRO:CA	2.02	0.90
1:B:480:LEU:HD23	1:B:559:VAL:HG22	1.53	0.90
1:B:140:ALA:HB3	1:B:143:GLU:CB	2.02	0.90
1:B:460:LEU:N	1:B:462:LYS:HE3	1.84	0.90
1:B:510:MET:HA	1:B:513:GLN:NE2	1.86	0.90
1:A:62:PRO:HG3	1:A:110:LEU:HD13	1.53	0.90
1:A:370:LEU:HA	1:A:390:LEU:HD13	1.51	0.90
1:A:511:ARG:HA	1:A:516:LEU:HG	1.51	0.90
1:B:33:VAL:HG13	1:B:476:LEU:HD12	1.54	0.90
1:A:477:LEU:O	1:A:555:LEU:HD22	1.71	0.90
1:B:857:LYS:O	1:B:861:LYS:CB	2.20	0.90
1:A:13:ASP:HA	1:A:224:MET:O	1.71	0.90
1:B:457:ALA:O	1:B:462:LYS:NZ	2.04	0.90
1:B:507:GLN:NE2	1:B:562:HIS:O	2.05	0.90
1:A:49:PRO:HG3	1:A:291:PRO:HA	0.93	0.90
1:A:49:PRO:HD3	1:A:291:PRO:HB3	1.51	0.90
1:A:32:LEU:HD13	1:A:445:PHE:HA	1.50	0.89
1:A:138:LEU:C	1:A:148:ARG:HG2	1.98	0.89
1:A:223:PHE:CE2	1:A:292:CYS:HB3	2.07	0.89
1:B:448:ASP:C	1:B:451:LYS:HZ1	1.80	0.89
1:B:133:THR:HG22	1:B:152:ASP:OD2	1.72	0.89
1:B:399:VAL:HG13	1:B:419:GLY:H	1.35	0.89
1:A:39:VAL:HG23	1:A:209:VAL:HG22	1.53	0.89
1:B:20:GLU:OE1	1:B:20:GLU:N	2.05	0.89
1:B:257:ARG:NH1	1:B:408:LYS:HG3	1.87	0.89
1:A:457:ALA:HB3	1:A:522:LEU:HD12	1.54	0.89
1:A:1202:GLN:O	1:A:1204:LEU:N	2.04	0.89
1:B:399:VAL:HA	1:B:419:GLY:CA	2.01	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:371:ASP:C	1:A:389:ARG:HG2	1.98	0.89
1:B:13:ASP:O	1:B:59:LYS:HB2	1.72	0.89
1:B:143:GLU:HG3	1:B:210:ASN:CB	2.00	0.89
1:B:474:THR:HA	1:B:477:LEU:CD1	2.03	0.89
1:A:108:ARG:NH2	1:A:109:LYS:H	1.69	0.89
1:A:249:LYS:CA	1:A:264:LEU:HD21	2.02	0.89
1:B:19:ALA:CA	1:B:218:TRP:CZ3	2.56	0.89
1:B:140:ALA:HB1	1:B:208:GLU:OE2	1.72	0.89
1:B:400:HIS:ND1	1:B:402:THR:OG1	2.05	0.89
1:B:744:CYS:HA	1:B:1085:ARG:N	1.87	0.89
1:A:219:LYS:NZ	1:A:221:VAL:HG23	1.87	0.89
1:B:355:LEU:HB2	1:B:417:LYS:CE	2.03	0.89
1:B:228:ASP:O	1:B:233:ILE:HD11	1.73	0.89
1:A:316:ALA:HA	1:A:356:VAL:H	1.37	0.89
1:A:484:VAL:HG22	1:A:562:HIS:CD2	2.09	0.89
1:B:9:LEU:HD23	1:B:176:VAL:HB	1.55	0.89
1:A:28:SER:OG	1:A:55:ASP:OD2	1.89	0.88
1:A:243:PHE:C	1:A:430:PHE:CD1	2.51	0.88
1:A:308:LEU:HD12	1:A:309:ALA:H	1.37	0.88
1:A:400:HIS:O	1:A:418:ILE:C	2.16	0.88
1:B:59:LYS:HG2	1:B:124:LEU:HD23	1.54	0.88
1:B:115:ILE:CD1	1:B:121:ILE:HG21	2.02	0.88
1:B:117:TYR:HD2	1:B:165:PRO:HD3	1.38	0.88
1:A:391:ARG:HG3	1:A:398:TRP:HA	0.91	0.88
1:A:407:ASP:CG	1:A:415:MET:HG2	1.98	0.88
1:A:1209:GLY:O	1:A:1213:VAL:CB	2.21	0.88
1:A:466:THR:O	1:A:470:ARG:N	2.06	0.88
1:B:1090:ARG:O	1:B:1176:VAL:N	2.06	0.88
1:A:166:PHE:HE2	1:A:217:SER:HA	1.37	0.88
1:A:194:HIS:O	1:A:210:ASN:CB	2.21	0.88
1:A:249:LYS:CE	1:A:416:LEU:HD13	2.04	0.88
1:A:250:PHE:C	1:A:264:LEU:HG	1.98	0.88
1:A:453:LEU:CB	1:A:522:LEU:HD21	2.03	0.88
1:A:298:TYR:N	1:A:301:SER:HB2	1.87	0.88
1:A:392:HIS:CE1	1:A:395:THR:HG23	2.09	0.88
1:A:406:ILE:CG1	1:A:417:LYS:HD2	2.03	0.88
1:B:249:LYS:HG2	1:B:264:LEU:HD12	1.52	0.88
1:A:219:LYS:HZ3	1:A:221:VAL:HG23	1.38	0.88
1:B:257:ARG:CB	1:B:259:LYS:HZ3	1.86	0.88
1:B:507:GLN:HA	1:B:510:MET:SD	2.14	0.88
1:A:395:THR:O	1:A:397:THR:N	2.06	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:240:VAL:HG12	1:B:434:PRO:CB	2.03	0.88
1:B:443:LEU:HD12	1:B:509:LEU:CD1	2.04	0.88
1:A:315:ALA:HB2	1:A:358:VAL:CG1	2.04	0.88
1:A:407:ASP:HB3	1:A:415:MET:SD	2.13	0.88
1:B:44:GLY:HA3	1:B:50:PRO:HD3	1.52	0.88
1:A:20:GLU:CD	1:A:181:LYS:HZ2	1.82	0.87
1:B:19:ALA:CA	1:B:218:TRP:HZ3	1.86	0.87
1:B:69:GLN:HB3	1:B:100:LYS:HZ2	1.37	0.87
1:B:235:LYS:HD2	1:B:238:ASP:OD1	1.74	0.87
1:A:103:ASN:O	1:A:104:GLU:C	2.15	0.87
1:A:249:LYS:HE3	1:A:416:LEU:HD13	1.56	0.87
1:B:7:SER:OG	1:B:178:ILE:HB	1.72	0.87
1:B:65:ARG:NH2	1:B:99:GLU:OE1	2.07	0.87
1:B:59:LYS:O	1:B:123:LEU:HD13	1.75	0.87
1:A:63:MET:HE3	1:A:120:VAL:CG1	2.04	0.87
1:A:449:ALA:O	1:A:452:VAL:HG23	1.72	0.87
1:A:465:ILE:O	1:A:466:THR:OG1	1.92	0.87
1:A:1421:ALA:HB2	1:A:1456:ASP:HA	1.56	0.87
1:B:58:PHE:HA	1:B:124:LEU:O	1.73	0.87
1:B:134:VAL:HG13	1:B:147:MET:HB3	1.55	0.87
1:B:290:ASP:OD2	1:B:293:ARG:N	2.08	0.87
1:B:1194:VAL:O	1:B:1196:LYS:N	2.07	0.87
1:A:37:CYS:SG	1:A:209:VAL:HB	2.12	0.87
1:A:739:LEU:O	1:A:741:ALA:N	2.06	0.87
1:B:11:ILE:HA	1:B:60:LEU:HD23	1.56	0.87
1:B:385:ASN:OD1	1:B:433:VAL:HG22	1.74	0.87
1:B:399:VAL:O	1:B:420:THR:OG1	1.92	0.87
1:B:400:HIS:CB	1:B:419:GLY:O	2.22	0.87
1:A:62:PRO:CD	1:A:110:LEU:HD22	2.05	0.87
1:A:129:ASN:OD1	1:A:441:ARG:NH2	2.08	0.87
1:A:1294:HIS:O	1:A:1298:CYS:N	2.07	0.87
1:B:222:LEU:HD22	1:B:223:PHE:N	1.90	0.87
1:B:480:LEU:CD2	1:B:559:VAL:HG22	2.05	0.87
1:A:17:LEU:HD13	1:A:18:TYR:N	1.89	0.87
1:A:33:VAL:HG21	1:A:451:LYS:CG	2.05	0.87
1:A:455:SER:O	1:A:459:LYS:CG	2.23	0.87
1:B:244:HIS:CE1	1:B:249:LYS:HB3	2.09	0.87
1:A:118:GLY:H	1:A:163:ILE:HG23	1.40	0.86
1:A:514:ASN:ND2	1:A:518:GLN:HG2	1.90	0.86
1:B:223:PHE:HA	1:B:293:ARG:HA	1.57	0.86
1:A:142:LEU:HD22	1:A:198:HIS:HB3	1.57	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:130:LYS:HB3	1:B:151:LEU:HD22	1.56	0.86
1:B:193:LEU:HD23	1:B:193:LEU:O	1.74	0.86
1:B:60:LEU:HD22	1:B:60:LEU:N	1.79	0.86
1:B:117:TYR:CE2	1:B:182:VAL:HG11	2.10	0.86
1:B:316:ALA:HB3	1:B:392:HIS:ND1	1.89	0.86
1:B:181:LYS:HE3	1:B:219:LYS:HG2	1.57	0.86
1:A:456:ILE:N	1:A:459:LYS:HZ3	1.73	0.86
1:B:14:ILE:HB	1:B:223:PHE:CE2	2.10	0.86
1:B:44:GLY:HA3	1:B:50:PRO:CD	2.05	0.86
1:B:229:ASN:HB3	1:B:235:LYS:CB	2.05	0.86
1:B:240:VAL:HG12	1:B:434:PRO:HA	1.57	0.86
1:B:387:TYR:O	1:B:388:VAL:HG13	1.75	0.86
1:A:125:HIS:O	1:A:129:ASN:HA	1.75	0.86
1:A:509:LEU:HD11	1:A:513:GLN:HG3	1.58	0.86
1:B:373:THR:HG23	1:B:389:ARG:HE	1.39	0.86
1:B:478:GLU:OE1	1:B:478:GLU:N	2.08	0.86
1:B:480:LEU:HA	1:B:483:PHE:CE1	2.11	0.86
1:B:65:ARG:HH12	1:B:106:GLU:CD	1.84	0.86
1:B:451:LYS:HE2	1:B:451:LYS:C	2.01	0.86
1:A:123:LEU:O	1:A:131:TYR:HA	1.75	0.86
1:A:264:LEU:HD23	1:A:265:ARG:N	1.91	0.86
1:B:181:LYS:CB	1:B:217:SER:HB2	2.03	0.86
1:B:507:GLN:HG3	1:B:563:SER:HA	1.58	0.85
1:B:100:LYS:O	1:B:103:ASN:ND2	2.08	0.85
1:B:251:LEU:HD11	1:B:282:TRP:CH2	2.11	0.85
1:A:388:VAL:HG21	1:A:432:ILE:CG1	2.00	0.85
1:B:66:TYR:CE1	1:B:100:LYS:HD3	2.12	0.85
1:A:117:TYR:OH	1:A:174:ASP:HB2	1.75	0.85
1:B:404:ILE:O	1:B:416:LEU:HA	1.76	0.85
1:B:555:LEU:O	1:B:559:VAL:HG23	1.75	0.85
1:A:32:LEU:HD12	1:A:448:ASP:HB2	1.58	0.85
1:B:36:ARG:HG2	1:B:37:CYS:H	1.41	0.85
1:B:203:ASN:HB3	1:B:206:CYS:HB2	1.56	0.85
1:B:249:LYS:HD2	1:B:416:LEU:CB	2.05	0.85
1:A:168:LYS:O	1:A:168:LYS:HD3	1.74	0.85
1:A:264:LEU:HD13	1:A:418:ILE:HD12	1.58	0.85
1:A:389:ARG:HH22	1:A:430:PHE:H	1.23	0.85
1:B:38:VAL:HA	1:B:207:ASN:O	1.75	0.85
1:B:263:PHE:CG	1:B:415:MET:HE3	2.11	0.85
1:B:298:TYR:HB2	1:B:380:SER:HB3	1.59	0.85
1:B:503:ASN:HB2	1:B:506:ARG:HH11	1.38	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:47:ASN:C	1:A:49:PRO:HD2	2.01	0.85
1:A:239:VAL:HG13	1:A:435:VAL:CB	2.06	0.85
1:A:410:GLU:CB	1:A:413:PRO:HB3	2.06	0.85
1:B:218:TRP:HA	1:B:218:TRP:HE3	1.41	0.85
1:A:453:LEU:HD23	1:A:518:GLN:CG	2.06	0.85
1:B:222:LEU:HD22	1:B:223:PHE:H	1.40	0.85
1:B:312:HIS:CD2	1:B:359:PRO:HA	2.12	0.85
1:A:14:ILE:HB	1:A:59:LYS:HA	1.56	0.85
1:A:1287:ILE:HA	1:A:1343:VAL:CB	2.07	0.85
1:B:11:ILE:H	1:B:113:THR:H	1.24	0.85
1:B:507:GLN:HE21	1:B:562:HIS:C	1.83	0.85
1:A:118:GLY:N	1:A:163:ILE:HG23	1.92	0.84
1:A:239:VAL:O	1:A:435:VAL:N	2.10	0.84
1:B:283:GLU:O	1:B:306:LYS:CB	2.19	0.84
1:B:446:ALA:O	1:B:450:SER:OG	1.93	0.84
1:B:744:CYS:HA	1:B:1085:ARG:CA	2.06	0.84
1:A:164:GLN:OE1	1:A:185:ASN:ND2	2.09	0.84
1:B:18:TYR:HA	1:B:26:PHE:CD1	2.12	0.84
1:B:115:ILE:HD12	1:B:121:ILE:HG21	1.59	0.84
1:B:417:LYS:C	1:B:418:ILE:HG12	2.02	0.84
1:B:16:SER:OG	1:B:292:CYS:HB3	1.76	0.84
1:B:115:ILE:O	1:B:176:VAL:HG23	1.77	0.84
1:A:259:LYS:HB3	1:A:259:LYS:HZ3	1.43	0.84
1:B:451:LYS:CE	1:B:452:VAL:H	1.89	0.84
1:A:402:THR:N	1:A:418:ILE:HG22	1.91	0.84
1:A:477:LEU:HA	1:A:480:LEU:HD11	0.92	0.84
1:B:264:LEU:HD21	1:B:430:PHE:HE1	1.43	0.84
1:B:27:ILE:HA	1:B:39:VAL:HG12	1.59	0.84
1:A:40:GLN:HG2	1:A:205:GLY:O	1.78	0.84
1:A:32:LEU:N	1:A:448:ASP:OD2	2.09	0.84
1:A:72:PHE:HB2	1:A:92:LEU:HD22	1.57	0.84
1:A:193:LEU:HA	1:A:211:SER:HA	1.59	0.84
1:A:459:LYS:CE	1:A:459:LYS:H	1.91	0.84
1:B:510:MET:CG	1:B:516:LEU:HG	2.06	0.84
1:B:514:ASN:O	1:B:518:GLN:N	2.09	0.84
1:A:184:LEU:C	1:A:185:ASN:HD22	1.84	0.84
1:A:1272:THR:O	1:A:1276:ILE:CB	2.26	0.84
1:B:304:ARG:HG3	1:B:304:ARG:NH1	1.92	0.84
1:B:372:PRO:HA	1:B:388:VAL:HG12	1.59	0.84
1:A:17:LEU:HD22	1:A:219:LYS:C	2.03	0.83
1:A:54:ARG:HA	1:A:57:LEU:CD2	2.08	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:253:CYS:HB3	1:A:262:VAL:HG12	1.57	0.83
1:A:383:PRO:HG2	1:A:386:SER:CB	2.07	0.83
1:A:391:ARG:NE	1:A:392:HIS:O	2.10	0.83
1:B:130:LYS:HG3	1:B:153:GLU:N	1.92	0.83
1:B:453:LEU:CD1	1:B:515:ILE:HD12	2.07	0.83
1:B:61:CYS:SG	1:B:124:LEU:HB3	2.17	0.83
1:B:181:LYS:HE3	1:B:219:LYS:CG	2.08	0.83
1:B:181:LYS:HA	1:B:218:TRP:O	1.79	0.83
1:B:384:ARG:HD2	1:B:434:PRO:CG	2.08	0.83
1:B:17:LEU:HA	1:B:219:LYS:O	1.78	0.83
1:B:72:PHE:HA	1:B:92:LEU:HD13	1.58	0.83
1:B:452:VAL:O	1:B:456:ILE:HG12	1.78	0.83
1:A:241:ARG:HB3	1:A:250:PHE:CZ	2.13	0.83
1:A:510:MET:SD	1:A:516:LEU:HD23	2.19	0.83
1:B:20:GLU:OE2	1:B:217:SER:OG	1.95	0.83
1:B:184:LEU:HB3	1:B:193:LEU:CB	2.08	0.83
1:A:245:ALA:N	1:A:429:ALA:O	2.10	0.83
1:A:520:PHE:HA	1:A:523:LEU:HD11	1.58	0.83
1:B:451:LYS:HE2	1:B:451:LYS:CA	2.06	0.83
1:A:455:SER:C	1:A:459:LYS:CE	2.51	0.83
1:A:518:GLN:HE21	1:A:518:GLN:HA	1.40	0.83
1:B:59:LYS:HG2	1:B:124:LEU:CD2	2.07	0.83
1:B:237:GLY:HA2	1:B:283:GLU:OE2	1.78	0.83
1:B:1129:SER:O	1:B:1132:TRP:N	2.11	0.83
1:A:14:ILE:HG21	1:A:59:LYS:C	2.03	0.83
1:A:15:CYS:CB	1:A:221:VAL:O	2.26	0.83
1:B:194:HIS:CE1	1:B:212:VAL:H	1.97	0.83
1:B:397:THR:OG1	1:B:421:SER:O	1.96	0.83
1:A:39:VAL:CG2	1:A:209:VAL:HG22	2.09	0.83
1:A:72:PHE:HA	1:A:92:LEU:HD13	1.59	0.83
1:A:453:LEU:HD23	1:A:518:GLN:CB	2.07	0.83
1:B:34:ASP:OD2	1:B:452:VAL:HG13	1.77	0.83
1:B:130:LYS:HD2	1:B:151:LEU:O	1.79	0.83
1:B:181:LYS:HD3	1:B:218:TRP:N	1.93	0.83
1:A:457:ALA:HB2	1:A:522:LEU:CA	2.07	0.83
1:B:191:GLN:HB3	1:B:212:VAL:HA	1.59	0.83
1:B:298:TYR:CA	1:B:381:LEU:HD13	2.08	0.83
1:B:406:ILE:N	1:B:415:MET:HB2	1.91	0.83
1:B:449:ALA:HB1	1:B:515:ILE:HD13	1.61	0.83
1:B:451:LYS:HZ3	1:B:451:LYS:H	1.24	0.83
1:A:306:LYS:HA	1:A:313:TYR:HA	1.58	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:9:LEU:CD2	1:B:176:VAL:HB	2.09	0.83
1:B:135:ASN:O	1:B:138:LEU:HD23	1.79	0.83
1:B:390:LEU:CD1	1:B:399:VAL:HB	2.08	0.83
1:B:400:HIS:HB3	1:B:419:GLY:O	1.77	0.83
1:B:415:MET:SD	1:B:415:MET:N	2.52	0.83
1:A:14:ILE:HG21	1:A:59:LYS:CA	2.08	0.82
1:A:315:ALA:CB	1:A:358:VAL:HG22	2.09	0.82
1:A:457:ALA:HB2	1:A:522:LEU:HA	1.61	0.82
1:A:244:HIS:CD2	1:A:430:PHE:CD1	2.67	0.82
1:B:18:TYR:HD1	1:B:26:PHE:CE1	1.97	0.82
1:B:75:ALA:O	1:B:89:LEU:HD13	1.78	0.82
1:B:303:PHE:C	1:B:367:ILE:HG21	2.04	0.82
1:B:475:LYS:HA	1:B:478:GLU:OE2	1.79	0.82
1:B:863:THR:O	1:B:867:VAL:CB	2.27	0.82
1:B:879:TYR:H	1:B:884:LEU:CB	1.92	0.82
1:A:18:TYR:CB	1:A:219:LYS:HE3	2.08	0.82
1:B:11:ILE:CG2	1:B:62:PRO:HD3	2.08	0.82
1:A:9:LEU:HD11	1:A:179:GLY:N	1.94	0.82
1:A:57:LEU:C	1:A:126:LEU:HG	2.03	0.82
1:A:263:PHE:CD2	1:A:415:MET:CB	2.62	0.82
1:A:288:GLN:HA	1:A:288:GLN:NE2	1.88	0.82
1:A:281:LEU:CB	1:A:308:LEU:HD21	2.06	0.82
1:B:33:VAL:CG1	1:B:452:VAL:HG21	2.09	0.82
1:B:57:LEU:CD2	1:B:126:LEU:HD21	2.10	0.82
1:B:286:VAL:HG11	1:B:303:PHE:CD1	2.14	0.82
1:B:507:GLN:HA	1:B:563:SER:OG	1.80	0.82
1:B:131:TYR:HB2	1:B:153:GLU:O	1.79	0.82
1:B:305:PHE:CZ	1:B:367:ILE:HA	2.15	0.82
1:A:259:LYS:O	1:A:261:HIS:CD2	2.32	0.82
1:B:117:TYR:HE2	1:B:182:VAL:HG11	1.43	0.82
1:B:420:THR:HG22	1:B:425:GLU:O	1.79	0.82
1:A:15:CYS:CB	1:A:223:PHE:HB2	2.09	0.82
1:A:589:VAL:O	1:A:592:GLU:CB	2.28	0.82
1:B:507:GLN:CG	1:B:563:SER:HA	2.08	0.82
1:A:9:LEU:HD11	1:A:179:GLY:H	1.44	0.82
1:A:65:ARG:NH1	1:A:99:GLU:O	2.13	0.82
1:A:117:TYR:HE2	1:A:173:GLY:N	1.76	0.82
1:A:207:ASN:ND2	1:A:207:ASN:H	1.77	0.82
1:A:478:GLU:HG2	1:A:555:LEU:CD1	2.09	0.82
1:B:46:LEU:HD13	1:B:221:VAL:HG13	1.62	0.82
1:B:105:THR:O	1:B:109:LYS:HE3	1.80	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:400:HIS:N	1:B:420:THR:N	2.26	0.82
1:B:437:PRO:O	1:B:441:ARG:CZ	2.27	0.82
1:A:315:ALA:HB2	1:A:358:VAL:CB	2.09	0.82
1:A:455:SER:O	1:A:459:LYS:HE2	1.79	0.82
1:B:286:VAL:HG11	1:B:303:PHE:CE1	2.15	0.82
1:B:725:GLU:CB	1:B:861:LYS:CB	2.58	0.82
1:B:963:GLU:O	1:B:967:VAL:CB	2.28	0.82
1:A:51:LYS:O	1:A:306:LYS:NZ	2.13	0.81
1:A:265:ARG:HA	1:A:416:LEU:CG	2.09	0.81
1:A:400:HIS:CE1	1:A:423:LEU:HD12	2.15	0.81
1:A:16:SER:CB	1:A:53:PHE:CZ	2.63	0.81
1:A:447:ASN:OD1	1:A:513:GLN:NE2	2.13	0.81
1:B:27:ILE:CD1	1:B:209:VAL:HG21	2.10	0.81
1:B:192:PRO:HB2	1:B:194:HIS:NE2	1.95	0.81
1:B:507:GLN:CD	1:B:563:SER:HA	2.05	0.81
1:A:63:MET:HE2	1:A:160:TRP:CG	2.15	0.81
1:A:87:VAL:HA	1:A:90:ASN:ND2	1.96	0.81
1:A:132:LEU:HA	1:A:151:LEU:HD22	1.60	0.81
1:A:450:SER:HA	1:A:518:GLN:HG3	1.60	0.81
1:B:12:GLY:HA2	1:B:59:LYS:HE3	1.63	0.81
1:B:36:ARG:NH1	1:B:200:LEU:HD21	1.95	0.81
1:B:407:ASP:HB2	1:B:413:PRO:CA	2.10	0.81
1:B:518:GLN:O	1:B:522:LEU:HD22	1.80	0.81
1:A:124:LEU:HD11	1:A:129:ASN:C	2.05	0.81
1:B:130:LYS:CE	1:B:151:LEU:HB3	2.09	0.81
1:B:224:MET:HG3	1:B:294:GLY:CA	2.09	0.81
1:B:249:LYS:HG2	1:B:264:LEU:HD13	1.62	0.81
1:B:399:VAL:HG13	1:B:419:GLY:N	1.95	0.81
1:A:367:ILE:O	1:A:393:LEU:HB2	1.78	0.81
1:A:387:TYR:O	1:A:388:VAL:HG13	1.81	0.81
1:A:453:LEU:HD23	1:A:518:GLN:HB3	1.61	0.81
1:B:403:ASN:HA	1:B:416:LEU:CD1	2.10	0.81
1:A:392:HIS:CD2	1:A:394:CYS:H	1.98	0.81
1:B:184:LEU:O	1:B:193:LEU:N	2.13	0.81
1:A:11:ILE:HG22	1:A:110:LEU:HD21	1.62	0.81
1:A:417:LYS:O	1:A:418:ILE:HG12	1.80	0.81
1:B:131:TYR:N	1:B:153:GLU:O	2.14	0.81
1:B:459:LYS:HA	1:B:462:LYS:CD	2.10	0.81
1:A:12:GLY:C	1:A:111:LEU:HD21	2.06	0.81
1:A:1472:SER:O	1:A:1476:LYS:CB	2.28	0.81
1:A:13:ASP:HB3	1:A:225:LYS:HA	1.61	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:249:LYS:HG3	1:A:401:SER:OG	1.81	0.81
1:A:256:HIS:O	1:A:259:LYS:HE2	1.80	0.81
1:A:302:LEU:HD11	1:A:393:LEU:CD2	2.10	0.81
1:A:54:ARG:O	1:A:57:LEU:HG	1.80	0.81
1:A:259:LYS:HE2	1:A:259:LYS:H	1.44	0.81
1:A:404:ILE:O	1:A:417:LYS:HB2	1.80	0.81
1:A:15:CYS:HB3	1:A:222:LEU:C	2.06	0.80
1:A:142:LEU:HD22	1:A:198:HIS:CB	2.11	0.80
1:B:27:ILE:HD12	1:B:209:VAL:HG21	1.61	0.80
1:B:103:ASN:HA	1:B:107:ASN:OD1	1.80	0.80
1:B:111:LEU:HD22	1:B:226:TRP:CH2	2.16	0.80
1:B:141:LEU:HD23	1:B:141:LEU:H	1.46	0.80
1:B:226:TRP:CD1	1:B:226:TRP:H	1.97	0.80
1:B:441:ARG:H	1:B:441:ARG:CZ	1.93	0.80
1:A:126:LEU:CD1	1:A:224:MET:HG2	2.12	0.80
1:B:132:LEU:HA	1:B:152:ASP:H	1.46	0.80
1:B:133:THR:HG22	1:B:152:ASP:CB	2.10	0.80
1:B:140:ALA:CB	1:B:143:GLU:HB2	2.10	0.80
1:A:452:VAL:HG21	1:A:476:LEU:HD22	1.61	0.80
1:B:65:ARG:HH21	1:B:99:GLU:CD	1.88	0.80
1:B:132:LEU:HB2	1:B:149:VAL:HG13	1.63	0.80
1:B:36:ARG:HH12	1:B:200:LEU:HD21	1.45	0.80
1:B:407:ASP:N	1:B:415:MET:HG2	1.97	0.80
1:B:476:LEU:O	1:B:479:ASP:HB2	1.81	0.80
1:B:286:VAL:CG1	1:B:303:PHE:CD1	2.64	0.80
1:B:301:SER:HB3	1:B:303:PHE:CE1	2.17	0.80
1:B:406:ILE:H	1:B:415:MET:CB	1.93	0.80
1:A:198:HIS:HB2	1:A:208:GLU:HG3	1.61	0.80
1:A:369:GLU:HB3	1:A:393:LEU:CD1	2.12	0.80
1:A:391:ARG:HH12	1:A:396:ASN:HA	1.47	0.80
1:A:1306:VAL:O	1:A:1310:LYS:N	2.15	0.80
1:B:33:VAL:HB	1:B:448:ASP:C	2.06	0.80
1:B:130:LYS:HE2	1:B:151:LEU:HD22	1.64	0.80
1:B:504:ARG:HB3	1:B:505:GLU:OE2	1.81	0.80
1:A:456:ILE:C	1:A:459:LYS:CD	2.46	0.80
1:B:224:MET:HG3	1:B:294:GLY:C	2.06	0.80
1:B:224:MET:HA	1:B:225:LYS:HD2	1.63	0.80
1:B:448:ASP:HA	1:B:451:LYS:CD	2.10	0.80
1:A:306:LYS:HB2	1:A:313:TYR:CD1	2.17	0.80
1:B:18:TYR:OH	1:B:24:ASN:CB	2.30	0.80
1:B:196:SER:HB3	1:B:208:GLU:O	1.82	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:287:VAL:HB	1:B:304:ARG:NH1	1.97	0.80
1:B:368:PHE:HB3	1:B:391:ARG:O	1.82	0.80
1:A:1421:ALA:O	1:A:1425:PHE:N	2.14	0.80
1:B:448:ASP:O	1:B:452:VAL:HG22	1.80	0.80
1:B:457:ALA:HA	1:B:460:LEU:CD2	2.12	0.80
1:A:181:LYS:HZ3	1:A:219:LYS:CB	1.95	0.79
1:A:249:LYS:HA	1:A:264:LEU:CD2	2.13	0.79
1:A:455:SER:O	1:A:459:LYS:CD	2.30	0.79
1:A:457:ALA:N	1:A:459:LYS:HZ2	1.80	0.79
1:A:981:ILE:O	1:A:985:ARG:CB	2.30	0.79
1:A:264:LEU:CD1	1:A:418:ILE:HD11	2.11	0.79
1:A:480:LEU:HD12	1:A:480:LEU:H	1.46	0.79
1:A:17:LEU:HD13	1:A:18:TYR:H	1.48	0.79
1:A:225:LYS:HZ1	1:A:295:GLY:HA2	1.45	0.79
1:A:447:ASN:O	1:A:451:LYS:NZ	2.14	0.79
1:A:447:ASN:HA	1:A:513:GLN:HE22	1.47	0.79
1:A:511:ARG:CG	1:A:516:LEU:HD11	2.13	0.79
1:B:223:PHE:CD1	1:B:294:GLY:N	2.50	0.79
1:B:234:LEU:CG	1:B:384:ARG:HD3	2.06	0.79
1:A:11:ILE:HD12	1:A:113:THR:OG1	1.83	0.79
1:A:20:GLU:HB2	1:A:217:SER:CB	2.09	0.79
1:A:122:GLN:HG3	1:A:160:TRP:NE1	1.97	0.79
1:A:233:ILE:HG22	1:A:382:VAL:O	1.80	0.79
1:A:239:VAL:CG1	1:A:435:VAL:HB	2.11	0.79
1:A:406:ILE:HG13	1:A:417:LYS:CD	2.11	0.79
1:A:450:SER:HA	1:A:453:LEU:CD2	2.13	0.79
1:A:477:LEU:HB2	1:A:555:LEU:HB3	1.64	0.79
1:B:102:GLN:O	1:B:106:GLU:HG2	1.83	0.79
1:B:244:HIS:ND1	1:B:264:LEU:HD11	1.98	0.79
1:B:451:LYS:NZ	1:B:451:LYS:H	1.77	0.79
1:B:224:MET:HE3	1:B:295:GLY:N	1.94	0.79
1:B:507:GLN:HE21	1:B:563:SER:HA	1.42	0.79
1:B:1178:GLU:O	1:B:1182:ARG:CB	2.30	0.79
1:A:387:TYR:CA	1:A:431:ALA:HA	2.11	0.79
1:A:455:SER:C	1:A:459:LYS:HE2	2.07	0.79
1:B:20:GLU:OE1	1:B:181:LYS:NZ	2.15	0.79
1:B:508:LYS:HG3	1:B:511:ARG:HH21	1.48	0.79
1:A:41:PRO:CD	1:A:207:ASN:HD21	1.85	0.79
1:A:247:GLN:HE22	1:A:416:LEU:HD22	1.46	0.79
1:A:298:TYR:HB2	1:A:301:SER:OG	1.82	0.79
1:A:458:GLY:HA3	1:A:459:LYS:HE3	1.64	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:511:ARG:HB3	1:A:512:GLU:OE2	1.83	0.79
1:B:242:LEU:CA	1:B:431:ALA:O	2.29	0.79
1:B:264:LEU:O	1:B:416:LEU:HB2	1.83	0.79
1:B:475:LYS:O	1:B:475:LYS:HD3	1.80	0.79
1:A:263:PHE:HD2	1:A:415:MET:CB	1.95	0.79
1:B:46:LEU:CD1	1:B:221:VAL:HG13	2.12	0.79
1:B:181:LYS:HA	1:B:218:TRP:C	2.08	0.79
1:B:458:GLY:C	1:B:462:LYS:HZ2	1.89	0.79
1:A:63:MET:HG2	1:A:160:TRP:NE1	1.97	0.79
1:A:142:LEU:CD2	1:A:198:HIS:HB3	2.12	0.79
1:A:244:HIS:HA	1:A:430:PHE:HA	1.64	0.79
1:B:218:TRP:HA	1:B:218:TRP:CE3	2.13	0.79
1:B:856:ASP:O	1:B:860:ASN:CB	2.31	0.79
1:A:16:SER:HB2	1:A:53:PHE:HZ	1.47	0.79
1:A:45:ASP:OD1	1:A:48:ASN:HB2	1.81	0.79
1:A:57:LEU:HB3	1:A:126:LEU:CD1	2.12	0.79
1:A:241:ARG:NH1	1:A:277:SER:O	2.16	0.79
1:B:11:ILE:HG21	1:B:62:PRO:CD	2.11	0.79
1:B:111:LEU:CB	1:B:226:TRP:HZ3	1.93	0.79
1:B:405:PRO:HB3	1:B:413:PRO:HB2	1.65	0.79
1:A:40:GLN:OE1	1:A:43:ALA:N	2.14	0.78
1:B:36:ARG:NH2	1:B:202:ASP:OD2	2.16	0.78
1:B:76:ALA:CB	1:B:89:LEU:HD22	2.13	0.78
1:B:250:PHE:O	1:B:265:ARG:N	2.16	0.78
1:B:264:LEU:HD21	1:B:430:PHE:CE1	2.18	0.78
1:B:459:LYS:O	1:B:462:LYS:HE3	1.83	0.78
1:B:516:LEU:HD12	1:B:516:LEU:N	1.97	0.78
1:A:14:ILE:O	1:A:223:PHE:HB3	1.82	0.78
1:A:63:MET:CE	1:A:120:VAL:HG11	2.08	0.78
1:A:72:PHE:HA	1:A:92:LEU:CD1	2.14	0.78
1:A:225:LYS:O	1:A:228:ASP:HB2	1.82	0.78
1:A:318:VAL:HA	1:A:353:TYR:HA	1.65	0.78
1:A:318:VAL:HG13	1:A:353:TYR:CG	2.18	0.78
1:A:373:THR:N	1:A:389:ARG:HD2	1.98	0.78
1:A:423:LEU:O	1:A:424:LYS:HB2	1.83	0.78
1:A:452:VAL:CG2	1:A:476:LEU:HD22	2.13	0.78
1:B:39:VAL:HG11	1:B:195:ALA:CB	2.08	0.78
1:B:1420:ILE:O	1:B:1424:ASN:N	2.16	0.78
1:A:405:PRO:HA	1:A:416:LEU:HA	1.66	0.78
1:A:505:GLU:N	1:A:505:GLU:OE1	2.16	0.78
1:B:372:PRO:CA	1:B:388:VAL:HG12	2.14	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1179:ILE:O	1:B:1183:LEU:CB	2.32	0.78
1:A:306:LYS:HB2	1:A:313:TYR:HE1	1.48	0.78
1:A:406:ILE:HD12	1:A:415:MET:HB2	1.66	0.78
1:B:305:PHE:CE1	1:B:367:ILE:HA	2.18	0.78
1:B:510:MET:HB2	1:B:515:ILE:HG23	1.64	0.78
1:B:58:PHE:CB	1:B:123:LEU:HD12	2.13	0.78
1:B:103:ASN:N	1:B:106:GLU:OE2	2.17	0.78
1:B:17:LEU:CA	1:B:46:LEU:HD11	2.13	0.78
1:B:240:VAL:HG12	1:B:434:PRO:HB3	1.66	0.78
1:B:433:VAL:HG12	1:B:434:PRO:HD2	1.65	0.78
1:A:406:ILE:CD1	1:A:415:MET:HB2	2.14	0.78
1:A:450:SER:O	1:A:518:GLN:OE1	2.02	0.78
1:B:66:TYR:CE1	1:B:100:LYS:HA	2.17	0.78
1:B:229:ASN:HB3	1:B:235:LYS:HB2	1.65	0.78
1:B:395:THR:HG22	1:B:397:THR:HG22	1.66	0.78
1:B:458:GLY:C	1:B:462:LYS:NZ	2.40	0.78
1:A:181:LYS:CE	1:A:219:LYS:CB	2.57	0.78
1:A:227:SER:HA	1:A:230:LYS:CD	2.13	0.78
1:A:508:LYS:NZ	1:A:512:GLU:HG3	1.98	0.78
1:A:511:ARG:HD3	1:A:512:GLU:OE2	1.83	0.78
1:A:132:LEU:HA	1:A:151:LEU:HD21	1.64	0.78
1:A:140:ALA:H	1:A:147:MET:C	1.91	0.78
1:A:389:ARG:NH1	1:A:389:ARG:HB3	1.98	0.78
1:B:54:ARG:HG2	1:B:283:GLU:CD	2.08	0.78
1:B:121:ILE:HD11	1:B:163:ILE:HG21	1.64	0.78
1:B:226:TRP:HA	1:B:229:ASN:OD1	1.83	0.78
1:A:306:LYS:HG2	1:A:307:HIS:H	1.46	0.77
1:A:704:ASN:HA	1:A:733:TYR:HA	1.65	0.77
1:B:102:GLN:HB3	1:B:106:GLU:OE2	1.84	0.77
1:B:181:LYS:CB	1:B:218:TRP:H	1.96	0.77
1:B:251:LEU:HD21	1:B:282:TRP:CZ2	2.17	0.77
1:A:27:ILE:CD1	1:A:39:VAL:HG22	2.13	0.77
1:A:69:GLN:CD	1:A:96:ALA:HA	2.09	0.77
1:A:244:HIS:H	1:A:248:GLU:HA	1.49	0.77
1:B:184:LEU:HD13	1:B:193:LEU:HD22	1.65	0.77
1:B:441:ARG:NE	1:B:441:ARG:N	2.31	0.77
1:B:480:LEU:HA	1:B:483:PHE:CZ	2.19	0.77
1:A:142:LEU:HD13	1:A:208:GLU:OE2	1.84	0.77
1:A:502:PRO:HG3	1:A:562:HIS:NE2	1.99	0.77
1:A:505:GLU:CD	1:A:505:GLU:H	1.89	0.77
1:B:383:PRO:O	1:B:386:SER:OG	2.00	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:461:GLU:C	1:B:462:LYS:HE2	2.08	0.77
1:B:696:VAL:O	1:B:700:TRP:CB	2.32	0.77
1:A:126:LEU:HD13	1:A:224:MET:HG2	1.64	0.77
1:A:244:HIS:CE1	1:A:428:GLU:HB3	2.19	0.77
1:B:241:ARG:N	1:B:241:ARG:HE	1.81	0.77
1:B:513:GLN:OE1	1:B:513:GLN:N	2.18	0.77
1:A:33:VAL:HG12	1:A:452:VAL:CG2	2.14	0.77
1:A:454:GLY:C	1:A:459:LYS:NZ	2.39	0.77
1:A:622:VAL:HA	1:A:628:PRO:HA	1.67	0.77
1:A:455:SER:O	1:A:459:LYS:CE	2.32	0.77
1:B:244:HIS:CG	1:B:264:LEU:HD11	2.20	0.77
1:B:298:TYR:HA	1:B:381:LEU:CA	2.10	0.77
1:B:404:ILE:HD12	1:B:404:ILE:N	2.00	0.77
1:A:358:VAL:HG11	1:A:366:SER:OG	1.84	0.77
1:B:519:ILE:O	1:B:523:LEU:HD21	1.84	0.77
1:A:315:ALA:O	1:A:356:VAL:HG23	1.84	0.77
1:A:477:LEU:CD1	1:A:552:ILE:HD12	2.13	0.77
1:B:14:ILE:HB	1:B:223:PHE:HD2	1.49	0.77
1:B:181:LYS:HG2	1:B:219:LYS:N	1.98	0.77
1:A:14:ILE:CG2	1:A:60:LEU:HD22	2.12	0.77
1:A:27:ILE:HG22	1:A:56:CYS:SG	2.25	0.77
1:A:65:ARG:HH22	1:A:103:ASN:HA	1.50	0.77
1:A:66:TYR:OH	1:A:100:LYS:HG2	1.84	0.77
1:A:259:LYS:HB3	1:A:259:LYS:NZ	1.97	0.77
1:A:509:LEU:HD12	1:A:513:GLN:HB2	1.67	0.77
1:B:102:GLN:C	1:B:106:GLU:HG2	2.10	0.77
1:B:224:MET:HB2	1:B:295:GLY:HA3	1.67	0.77
1:B:1000:GLU:CB	1:B:1006:SER:H	1.97	0.77
1:A:46:LEU:O	1:A:291:PRO:HD3	1.84	0.76
1:A:133:THR:HG23	1:A:156:ASN:OD1	1.85	0.76
1:B:299:TRP:CD1	1:B:380:SER:HG	2.02	0.76
1:B:458:GLY:O	1:B:462:LYS:CE	2.32	0.76
1:B:1090:ARG:CB	1:B:1093:VAL:HA	2.15	0.76
1:A:27:ILE:HD11	1:A:39:VAL:CG2	2.14	0.76
1:A:391:ARG:HD2	1:A:398:TRP:CD1	2.20	0.76
1:B:313:TYR:CZ	1:B:361:GLY:HA3	2.21	0.76
1:A:223:PHE:HE2	1:A:292:CYS:HB3	1.48	0.76
1:A:249:LYS:C	1:A:264:LEU:HD21	2.09	0.76
1:A:306:LYS:HG2	1:A:307:HIS:N	2.00	0.76
1:B:66:TYR:HB3	1:B:69:GLN:HG2	1.67	0.76
1:B:414:VAL:CG2	1:B:415:MET:SD	2.73	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:440:VAL:HB	1:B:441:ARG:CZ	2.16	0.76
1:A:55:ASP:HA	1:A:127:LYS:HD3	1.66	0.76
1:A:218:TRP:HA	1:A:218:TRP:CE3	2.18	0.76
1:B:251:LEU:CD1	1:B:282:TRP:HH2	1.98	0.76
1:B:473:VAL:CG2	1:B:477:LEU:HD21	2.15	0.76
1:B:864:PHE:O	1:B:868:ASN:CB	2.34	0.76
1:B:1209:GLY:O	1:B:1213:VAL:CB	2.33	0.76
1:A:19:ALA:HA	1:A:218:TRP:CZ3	2.20	0.76
1:B:185:ASN:HA	1:B:191:GLN:O	1.84	0.76
1:B:392:HIS:CD2	1:B:395:THR:H	2.03	0.76
1:A:7:SER:O	1:A:178:ILE:N	2.19	0.76
1:A:15:CYS:HA	1:A:223:PHE:CB	2.15	0.76
1:A:365:SER:HA	1:A:394:CYS:CB	2.15	0.76
1:B:132:LEU:H	1:B:132:LEU:CD1	1.97	0.76
1:B:212:VAL:HG22	1:B:213:ASN:H	1.49	0.76
1:B:443:LEU:HD12	1:B:509:LEU:HD11	1.65	0.76
1:B:18:TYR:C	1:B:218:TRP:HZ3	1.92	0.76
1:A:57:LEU:O	1:A:126:LEU:N	2.19	0.76
1:A:478:GLU:OE2	1:A:555:LEU:CD2	2.29	0.76
1:A:516:LEU:HD12	1:A:520:PHE:CE2	2.21	0.76
1:B:33:VAL:HG11	1:B:449:ALA:HA	1.66	0.76
1:B:385:ASN:CG	1:B:433:VAL:HG22	2.11	0.76
1:A:306:LYS:HD3	1:A:311:GLY:C	2.10	0.76
1:A:510:MET:CB	1:A:516:LEU:HD23	2.16	0.76
1:A:510:MET:HB3	1:A:516:LEU:HD23	1.68	0.76
1:A:555:LEU:O	1:A:559:VAL:HG23	1.86	0.76
1:B:253:CYS:HB2	1:B:307:HIS:HE1	1.51	0.76
1:B:264:LEU:HB2	1:B:418:ILE:HD11	1.68	0.76
1:A:449:ALA:O	1:A:453:LEU:HD22	1.86	0.76
1:B:224:MET:CE	1:B:295:GLY:N	2.47	0.76
1:B:405:PRO:CB	1:B:413:PRO:HB2	2.15	0.76
1:B:503:ASN:O	1:B:507:GLN:CD	2.29	0.76
1:A:233:ILE:HG21	1:A:381:LEU:CB	2.17	0.75
1:A:282:TRP:CE3	1:A:307:HIS:HA	2.22	0.75
1:A:480:LEU:HD12	1:A:480:LEU:N	2.00	0.75
1:A:511:ARG:HA	1:A:516:LEU:CD1	2.15	0.75
1:B:20:GLU:CD	1:B:181:LYS:HZ3	1.93	0.75
1:B:39:VAL:CG1	1:B:195:ALA:HB1	2.10	0.75
1:B:510:MET:HB2	1:B:515:ILE:HG21	1.69	0.75
1:B:133:THR:C	1:B:149:VAL:HG22	2.11	0.75
1:A:244:HIS:CD2	1:A:429:ALA:H	2.04	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:502:PRO:HD2	1:A:507:GLN:NE2	2.01	0.75
1:B:385:ASN:N	1:B:432:ILE:HB	2.01	0.75
1:B:400:HIS:CA	1:B:420:THR:HA	2.17	0.75
1:A:298:TYR:O	1:A:301:SER:N	2.20	0.75
1:A:304:ARG:HD3	1:A:367:ILE:CG2	2.17	0.75
1:B:148:ARG:HG3	1:B:148:ARG:HH11	1.50	0.75
1:B:513:GLN:NE2	1:B:515:ILE:HG22	2.00	0.75
1:A:30:LEU:HB2	1:A:34:ASP:O	1.86	0.75
1:A:39:VAL:HG23	1:A:209:VAL:CG2	2.15	0.75
1:A:265:ARG:HA	1:A:416:LEU:HD12	1.68	0.75
1:A:416:LEU:O	1:A:418:ILE:HG23	1.86	0.75
1:B:44:GLY:HA3	1:B:50:PRO:N	2.02	0.75
1:B:473:VAL:HG23	1:B:477:LEU:HD21	1.69	0.75
1:A:129:ASN:CG	1:A:441:ARG:HH12	1.94	0.75
1:B:54:ARG:HH11	1:B:54:ARG:N	1.83	0.75
1:B:130:LYS:CD	1:B:151:LEU:HB3	2.17	0.75
1:B:138:LEU:O	1:B:148:ARG:N	2.19	0.75
1:B:407:ASP:CB	1:B:413:PRO:HA	2.17	0.75
1:B:448:ASP:C	1:B:451:LYS:HE3	2.10	0.75
1:A:143:GLU:CB	1:A:210:ASN:HD21	2.00	0.75
1:B:32:LEU:HD23	1:B:32:LEU:H	1.51	0.75
1:A:45:ASP:OD1	1:A:45:ASP:N	2.20	0.75
1:A:124:LEU:HB2	1:A:131:TYR:HE1	1.52	0.75
1:A:263:PHE:CD2	1:A:415:MET:SD	2.80	0.75
1:A:303:PHE:HB3	1:A:305:PHE:CE1	2.21	0.75
1:A:457:ALA:HB2	1:A:522:LEU:CG	2.17	0.75
1:B:300:ASN:O	1:B:301:SER:HB2	1.86	0.75
1:B:510:MET:C	1:B:516:LEU:HD11	2.12	0.75
1:A:124:LEU:HD12	1:A:125:HIS:H	1.52	0.75
1:A:230:LYS:HA	1:A:230:LYS:CE	2.17	0.75
1:A:287:VAL:CG2	1:A:302:LEU:HB3	2.16	0.75
1:A:448:ASP:O	1:A:449:ALA:C	2.30	0.75
1:B:130:LYS:HA	1:B:153:GLU:HA	1.67	0.75
1:B:457:ALA:HA	1:B:460:LEU:HD21	1.69	0.75
1:A:249:LYS:C	1:A:264:LEU:HD11	2.12	0.74
1:B:49:PRO:HG2	1:B:53:PHE:CZ	2.21	0.74
1:B:277:SER:HA	1:B:508:LYS:NZ	2.01	0.74
1:A:61:CYS:SG	1:A:62:PRO:HD2	2.27	0.74
1:A:85:ASP:N	1:A:88:LEU:HD13	2.02	0.74
1:A:373:THR:HG23	1:A:389:ARG:CD	2.16	0.74
1:A:57:LEU:HB2	1:A:126:LEU:HB2	1.70	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:59:LYS:HB3	1:A:126:LEU:CD2	2.16	0.74
1:A:304:ARG:NH2	1:A:363:ASP:O	2.20	0.74
1:B:223:PHE:HD1	1:B:293:ARG:HA	1.52	0.74
1:B:224:MET:CE	1:B:294:GLY:HA2	2.16	0.74
1:B:450:SER:C	1:B:451:LYS:HZ3	1.95	0.74
1:B:515:ILE:O	1:B:518:GLN:HB2	1.87	0.74
1:A:122:GLN:HG3	1:A:160:TRP:CD1	2.22	0.74
1:A:244:HIS:CD2	1:A:430:PHE:N	2.55	0.74
1:A:303:PHE:O	1:A:367:ILE:HG22	1.87	0.74
1:A:315:ALA:C	1:A:356:VAL:O	2.29	0.74
1:B:277:SER:HA	1:B:508:LYS:HZ3	1.52	0.74
1:A:20:GLU:OE2	1:A:218:TRP:N	2.21	0.74
1:A:166:PHE:CZ	1:A:217:SER:HA	2.22	0.74
1:A:233:ILE:HG21	1:A:381:LEU:HB3	1.69	0.74
1:B:54:ARG:HD2	1:B:283:GLU:HG2	1.69	0.74
1:B:249:LYS:CB	1:B:266:THR:HA	2.17	0.74
1:B:1129:SER:C	1:B:1133:VAL:H	1.94	0.74
1:A:244:HIS:HD2	1:A:430:PHE:N	1.86	0.74
1:B:71:GLN:HA	1:B:74:LYS:HD3	1.68	0.74
1:B:181:LYS:HD3	1:B:217:SER:C	2.12	0.74
1:B:250:PHE:HA	1:B:430:PHE:HZ	1.53	0.74
1:B:400:HIS:O	1:B:402:THR:N	2.20	0.74
1:B:473:VAL:O	1:B:477:LEU:HG	1.88	0.74
1:A:477:LEU:CD1	1:A:552:ILE:HA	2.18	0.74
1:B:54:ARG:HE	1:B:308:LEU:CD1	1.99	0.74
1:B:142:LEU:HB3	1:B:198:HIS:ND1	2.03	0.74
1:B:406:ILE:HB	1:B:415:MET:CG	2.16	0.74
1:A:12:GLY:O	1:A:111:LEU:HD21	1.88	0.74
1:A:32:LEU:CD1	1:A:445:PHE:CA	2.51	0.74
1:A:62:PRO:HG3	1:A:110:LEU:CD1	2.18	0.74
1:B:400:HIS:H	1:B:419:GLY:N	1.86	0.74
1:A:510:MET:HB3	1:A:516:LEU:CD2	2.18	0.74
1:B:191:GLN:HB3	1:B:212:VAL:CA	2.16	0.74
1:A:392:HIS:HE1	1:A:395:THR:HG23	1.52	0.74
1:B:391:ARG:HD3	1:B:398:TRP:NE1	2.02	0.74
1:B:446:ALA:HB1	1:B:513:GLN:HE21	1.53	0.74
1:B:448:ASP:O	1:B:451:LYS:CE	2.33	0.74
1:A:69:GLN:O	1:A:72:PHE:N	2.19	0.73
1:A:198:HIS:CB	1:A:208:GLU:HG3	2.17	0.73
1:A:371:ASP:HB3	1:A:389:ARG:HG3	1.68	0.73
1:B:58:PHE:HE1	1:B:125:HIS:CG	2.05	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:102:GLN:NE2	1:B:106:GLU:OE1	2.21	0.73
1:B:298:TYR:CB	1:B:381:LEU:HD22	2.15	0.73
1:B:304:ARG:NE	1:B:304:ARG:H	1.85	0.73
1:A:372:PRO:HA	1:A:389:ARG:CD	2.18	0.73
1:A:510:MET:SD	1:A:516:LEU:N	2.61	0.73
1:B:66:TYR:HE1	1:B:100:LYS:CA	2.00	0.73
1:B:511:ARG:HA	1:B:516:LEU:CD1	2.18	0.73
1:A:33:VAL:H	1:A:448:ASP:HB3	1.53	0.73
1:A:129:ASN:HB2	1:A:441:ARG:HH12	1.53	0.73
1:A:138:LEU:O	1:A:148:ARG:HG2	1.89	0.73
1:A:459:LYS:CD	1:A:459:LYS:H	1.93	0.73
1:A:484:VAL:O	1:A:562:HIS:NE2	2.21	0.73
1:B:249:LYS:HD3	1:B:266:THR:N	2.03	0.73
1:B:286:VAL:HG13	1:B:303:PHE:HA	1.70	0.73
1:B:439:GLU:HA	1:B:442:ASP:OD2	1.88	0.73
1:A:62:PRO:CG	1:A:110:LEU:HD22	2.18	0.73
1:A:305:PHE:HD2	1:A:314:LEU:HD23	1.53	0.73
1:A:511:ARG:HA	1:A:516:LEU:HD11	1.71	0.73
1:B:243:PHE:CD1	1:B:250:PHE:HE1	2.06	0.73
1:B:264:LEU:HB3	1:B:416:LEU:HB3	1.71	0.73
1:B:407:ASP:C	1:B:408:LYS:HZ2	1.97	0.73
1:A:282:TRP:CE3	1:A:282:TRP:HA	2.23	0.73
1:A:287:VAL:HG21	1:A:302:LEU:HG	1.69	0.73
1:A:299:TRP:HZ3	1:A:370:LEU:HB2	1.54	0.73
1:A:511:ARG:HG2	1:A:516:LEU:HD11	1.71	0.73
1:A:599:LEU:CB	1:A:637:LEU:CB	2.67	0.73
1:A:738:ASN:O	1:A:740:PHE:N	2.22	0.73
1:B:461:GLU:N	1:B:462:LYS:NZ	2.37	0.73
1:B:1400:LEU:HA	1:B:1442:THR:HA	1.71	0.73
1:A:67:SER:O	1:A:70:LYS:CB	2.37	0.73
1:A:239:VAL:HG21	1:A:308:LEU:CD2	2.18	0.73
1:A:477:LEU:CD1	1:A:552:ILE:HG23	2.19	0.73
1:A:510:MET:SD	1:A:515:ILE:HB	2.29	0.73
1:B:131:TYR:H	1:B:153:GLU:C	1.95	0.73
1:B:186:PRO:CD	1:B:191:GLN:O	2.32	0.73
1:B:371:ASP:OD2	1:B:389:ARG:CB	2.35	0.73
1:A:391:ARG:NH2	1:A:395:THR:O	2.21	0.73
1:B:11:ILE:HA	1:B:60:LEU:CD2	2.19	0.73
1:A:181:LYS:CG	1:A:219:LYS:HA	2.17	0.73
1:B:137:ARG:CA	1:B:147:MET:HA	2.18	0.73
1:B:507:GLN:HB3	1:B:563:SER:O	1.88	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:156:ASN:HD21	1:A:159:SER:H	1.35	0.73
1:A:234:LEU:HD21	1:A:432:ILE:CG2	2.18	0.73
1:B:308:LEU:HD22	1:B:308:LEU:N	2.03	0.73
1:B:399:VAL:C	1:B:420:THR:N	2.47	0.73
1:B:462:LYS:HD2	1:B:463:GLY:H	1.52	0.73
1:A:235:LYS:HE3	1:A:384:ARG:HH11	1.53	0.72
1:A:368:PHE:CE1	1:A:392:HIS:HA	2.24	0.72
1:B:32:LEU:HD12	1:B:445:PHE:CD1	2.24	0.72
1:A:16:SER:HB2	1:A:53:PHE:CZ	2.23	0.72
1:A:17:LEU:HD11	1:A:218:TRP:CE3	2.24	0.72
1:A:128:SER:O	1:A:130:LYS:HG3	1.88	0.72
1:A:372:PRO:C	1:A:389:ARG:HD2	2.14	0.72
1:A:405:PRO:CA	1:A:416:LEU:HA	2.19	0.72
1:A:406:ILE:HD12	1:A:406:ILE:H	1.54	0.72
1:B:196:SER:C	1:B:207:ASN:HD21	1.97	0.72
1:B:404:ILE:CD1	1:B:417:LYS:H	2.02	0.72
1:A:202:ASP:C	1:A:204:PRO:HD3	2.15	0.72
1:B:11:ILE:HG23	1:B:61:CYS:N	2.04	0.72
1:B:133:THR:H	1:B:152:ASP:CG	1.98	0.72
1:B:221:VAL:HG21	1:B:292:CYS:CB	2.17	0.72
1:B:303:PHE:C	1:B:367:ILE:CG2	2.61	0.72
1:B:507:GLN:HG3	1:B:563:SER:C	2.13	0.72
1:A:14:ILE:CG2	1:A:59:LYS:HA	2.18	0.72
1:A:63:MET:HE2	1:A:160:TRP:CD2	2.25	0.72
1:A:161:PHE:CB	1:A:184:LEU:HD21	2.20	0.72
1:A:243:PHE:O	1:A:430:PHE:HD1	1.71	0.72
1:A:306:LYS:HD3	1:A:311:GLY:CA	2.19	0.72
1:B:316:ALA:HA	1:B:355:LEU:HG	1.71	0.72
1:A:11:ILE:CD1	1:A:115:ILE:HD11	2.18	0.72
1:A:243:PHE:O	1:A:430:PHE:CD1	2.42	0.72
1:A:373:THR:HG21	1:A:428:GLU:O	1.88	0.72
1:B:226:TRP:H	1:B:226:TRP:HD1	1.36	0.72
1:B:281:LEU:HB3	1:B:308:LEU:CD2	2.09	0.72
1:B:286:VAL:CG1	1:B:288:GLN:CG	2.68	0.72
1:B:314:LEU:HD23	1:B:314:LEU:O	1.89	0.72
1:A:306:LYS:HD3	1:A:311:GLY:HA2	1.72	0.72
1:A:316:ALA:HB1	1:A:355:LEU:CG	2.20	0.72
1:A:457:ALA:HB1	1:A:522:LEU:HA	1.72	0.72
1:A:748:GLN:CB	1:A:1074:PRO:HA	2.20	0.72
1:B:66:TYR:CB	1:B:69:GLN:HG2	2.18	0.72
1:B:96:ALA:O	1:B:99:GLU:N	2.21	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:166:PHE:HE1	1:B:216:THR:O	1.72	0.72
1:A:59:LYS:HB3	1:A:126:LEU:HD23	1.71	0.72
1:A:259:LYS:HE2	1:A:259:LYS:N	2.04	0.72
1:A:298:TYR:CE1	1:A:381:LEU:HD22	2.24	0.72
1:B:404:ILE:HD13	1:B:417:LYS:N	2.03	0.72
1:A:163:ILE:HD12	1:A:183:VAL:O	1.88	0.72
1:A:264:LEU:CD1	1:A:418:ILE:CD1	2.67	0.72
1:A:304:ARG:HD3	1:A:367:ILE:HG23	1.71	0.72
1:A:459:LYS:N	1:A:459:LYS:CE	2.50	0.72
1:A:465:ILE:O	1:A:469:GLU:HB3	1.89	0.72
1:B:130:LYS:HE3	1:B:151:LEU:HD13	1.72	0.72
1:B:137:ARG:HA	1:B:147:MET:CA	2.19	0.72
1:B:224:MET:SD	1:B:288:GLN:CD	2.73	0.72
1:B:244:HIS:HB3	1:B:430:PHE:HE1	1.47	0.72
1:B:249:LYS:HD3	1:B:266:THR:H	1.55	0.72
1:A:458:GLY:N	1:A:459:LYS:CE	2.52	0.72
1:B:142:LEU:CD1	1:B:200:LEU:HA	2.20	0.72
1:A:17:LEU:HD22	1:A:219:LYS:O	1.89	0.72
1:A:306:LYS:CB	1:A:313:TYR:CD1	2.72	0.72
1:A:454:GLY:HA2	1:A:522:LEU:HD11	1.71	0.72
1:B:148:ARG:HG3	1:B:148:ARG:NH1	2.03	0.72
1:B:222:LEU:HD13	1:B:223:PHE:O	1.90	0.72
1:A:265:ARG:HA	1:A:416:LEU:HG	1.71	0.71
1:B:111:LEU:HD13	1:B:226:TRP:HH2	1.54	0.71
1:B:239:VAL:CG2	1:B:281:LEU:HD22	2.19	0.71
1:B:511:ARG:HA	1:B:516:LEU:HD11	1.72	0.71
1:A:14:ILE:N	1:A:224:MET:O	2.22	0.71
1:B:222:LEU:O	1:B:293:ARG:HA	1.90	0.71
1:B:224:MET:CA	1:B:225:LYS:HD2	2.21	0.71
1:B:241:ARG:H	1:B:241:ARG:CD	2.03	0.71
1:B:299:TRP:N	1:B:381:LEU:H	1.87	0.71
1:B:856:ASP:HA	1:B:860:ASN:H	1.55	0.71
1:A:118:GLY:HA2	1:A:163:ILE:O	1.91	0.71
1:B:181:LYS:HG2	1:B:219:LYS:HB3	1.72	0.71
1:B:250:PHE:HA	1:B:430:PHE:CZ	2.24	0.71
1:B:371:ASP:OD2	1:B:398:TRP:HZ3	1.72	0.71
1:A:247:GLN:C	1:A:249:LYS:H	1.98	0.71
1:A:363:ASP:OD1	1:A:366:SER:HB2	1.90	0.71
1:A:407:ASP:H	1:A:415:MET:HG2	1.56	0.71
1:B:32:LEU:HD23	1:B:32:LEU:N	2.04	0.71
1:A:138:LEU:HB3	1:A:148:ARG:NE	2.05	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:218:TRP:HA	1:A:218:TRP:HE3	1.53	0.71
1:A:391:ARG:NH2	1:A:392:HIS:O	2.23	0.71
1:A:401:SER:CA	1:A:418:ILE:HB	2.12	0.71
1:A:1372:MET:C	1:A:1375:ILE:H	1.99	0.71
1:B:55:ASP:O	1:B:125:HIS:NE2	2.22	0.71
1:B:522:LEU:N	1:B:522:LEU:HD23	2.04	0.71
1:B:522:LEU:C	1:B:523:LEU:HD13	2.14	0.71
1:A:485:THR:HG22	1:A:562:HIS:HE1	1.55	0.71
1:B:32:LEU:HD12	1:B:445:PHE:HD1	1.54	0.71
1:B:440:VAL:HB	1:B:441:ARG:NH2	2.06	0.71
1:B:725:GLU:HA	1:B:862:LEU:HA	1.72	0.71
1:A:32:LEU:HD12	1:A:445:PHE:HA	1.69	0.71
1:A:160:TRP:C	1:A:187:VAL:HG23	2.14	0.71
1:A:283:GLU:CG	1:A:306:LYS:HB3	2.19	0.71
1:A:363:ASP:OD2	1:A:366:SER:OG	2.08	0.71
1:B:198:HIS:C	1:B:200:LEU:H	1.99	0.71
1:B:450:SER:O	1:B:453:LEU:HG	1.91	0.71
1:A:9:LEU:CD1	1:A:179:GLY:H	2.03	0.71
1:A:519:ILE:O	1:A:523:LEU:HD13	1.90	0.71
1:A:769:SER:O	1:A:773:LEU:CB	2.38	0.71
1:B:65:ARG:NH1	1:B:106:GLU:CD	2.49	0.71
1:B:130:LYS:HD3	1:B:153:GLU:OE2	1.90	0.71
1:A:163:ILE:HD12	1:A:164:GLN:H	1.55	0.71
1:A:456:ILE:O	1:A:459:LYS:HD3	1.89	0.71
1:B:241:ARG:HD3	1:B:435:VAL:HG22	1.73	0.71
1:B:257:ARG:HB2	1:B:257:ARG:HH11	1.55	0.71
1:B:509:LEU:HD12	1:B:509:LEU:O	1.91	0.71
1:A:125:HIS:O	1:A:129:ASN:CA	2.39	0.71
1:A:166:PHE:HB3	1:A:181:LYS:O	1.91	0.71
1:B:420:THR:HG22	1:B:425:GLU:C	2.16	0.71
1:A:65:ARG:HH12	1:A:103:ASN:N	1.89	0.70
1:A:484:VAL:HG13	1:A:562:HIS:ND1	2.06	0.70
1:B:54:ARG:HE	1:B:308:LEU:HD12	1.55	0.70
1:B:133:THR:HG22	1:B:152:ASP:CG	2.16	0.70
1:B:507:GLN:HG3	1:B:563:SER:O	1.90	0.70
1:A:70:LYS:HA	1:A:73:TRP:CD2	2.26	0.70
1:A:263:PHE:CB	1:A:415:MET:CB	2.39	0.70
1:A:264:LEU:HB3	1:A:418:ILE:HD11	1.70	0.70
1:A:477:LEU:HD11	1:A:552:ILE:CD1	2.17	0.70
1:B:110:LEU:CD1	1:B:113:THR:HG22	2.14	0.70
1:B:219:LYS:HZ1	1:B:220:ILE:N	1.89	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:99:GLU:O	1:A:103:ASN:ND2	2.24	0.70
1:A:518:GLN:HE22	1:A:522:LEU:CD1	2.03	0.70
1:B:224:MET:HG3	1:B:294:GLY:N	2.05	0.70
1:B:304:ARG:HG3	1:B:304:ARG:HH11	1.53	0.70
1:A:55:ASP:OD1	1:A:127:LYS:HD3	1.90	0.70
1:A:94:HIS:O	1:A:98:LEU:HG	1.90	0.70
1:A:235:LYS:HE3	1:A:384:ARG:NH1	2.06	0.70
1:A:263:PHE:CE2	1:A:414:VAL:HB	2.27	0.70
1:A:305:PHE:CZ	1:A:368:PHE:HB2	2.26	0.70
1:A:510:MET:CE	1:A:515:ILE:HG22	2.21	0.70
1:B:11:ILE:HD12	1:B:62:PRO:HG3	1.73	0.70
1:B:147:MET:CE	1:B:186:PRO:HG3	2.22	0.70
1:B:181:LYS:CE	1:B:219:LYS:CG	2.69	0.70
1:B:264:LEU:HB3	1:B:416:LEU:CB	2.21	0.70
1:A:15:CYS:HA	1:A:223:PHE:HB2	1.73	0.70
1:A:124:LEU:HB2	1:A:131:TYR:CE1	2.25	0.70
1:A:143:GLU:OE1	1:A:198:HIS:NE2	2.23	0.70
1:A:253:CYS:CB	1:A:307:HIS:CE1	2.73	0.70
1:B:181:LYS:HB3	1:B:218:TRP:H	1.56	0.70
1:B:302:LEU:HG	1:B:367:ILE:CD1	2.21	0.70
1:B:503:ASN:HB3	1:B:506:ARG:HD2	1.73	0.70
1:A:143:GLU:HB3	1:A:210:ASN:HD21	1.55	0.70
1:A:304:ARG:CZ	1:A:367:ILE:HD13	2.22	0.70
1:A:477:LEU:HD13	1:A:552:ILE:HG23	1.73	0.70
1:A:1058:ARG:O	1:A:1059:THR:CB	2.39	0.70
1:B:57:LEU:HD22	1:B:126:LEU:CD2	2.20	0.70
1:B:308:LEU:HD22	1:B:308:LEU:H	1.55	0.70
1:B:398:TRP:C	1:B:420:THR:H	1.98	0.70
1:A:62:PRO:HD3	1:A:110:LEU:CD2	2.21	0.70
1:A:385:ASN:OD1	1:A:434:PRO:HD2	1.91	0.70
1:B:133:THR:HG22	1:B:152:ASP:HB3	1.73	0.70
1:B:244:HIS:CE1	1:B:247:GLN:CB	2.75	0.70
1:B:283:GLU:HB2	1:B:308:LEU:HD11	1.73	0.70
1:B:369:GLU:H	1:B:393:LEU:HD21	1.57	0.70
1:A:457:ALA:HB2	1:A:522:LEU:HB3	1.74	0.70
1:A:478:GLU:N	1:A:555:LEU:HD13	2.06	0.70
1:B:313:TYR:CD2	1:B:361:GLY:HA3	2.26	0.70
1:A:310:THR:HB	1:A:312:HIS:CE1	2.27	0.70
1:B:111:LEU:N	1:B:111:LEU:HD12	2.06	0.70
1:B:162:TYR:HB2	1:B:185:ASN:O	1.92	0.70
1:B:181:LYS:HG2	1:B:219:LYS:CB	2.22	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:461:GLU:N	1:B:462:LYS:HZ3	1.90	0.70
1:A:62:PRO:CG	1:A:110:LEU:HD13	2.22	0.69
1:A:316:ALA:HB1	1:A:355:LEU:HA	1.74	0.69
1:A:372:PRO:CA	1:A:389:ARG:HD2	2.22	0.69
1:A:510:MET:HE2	1:A:515:ILE:HG22	1.74	0.69
1:B:60:LEU:N	1:B:60:LEU:CD2	2.55	0.69
1:B:132:LEU:CB	1:B:149:VAL:CG1	2.70	0.69
1:B:449:ALA:HB3	1:B:515:ILE:HD13	1.74	0.69
1:A:58:PHE:CD1	1:A:125:HIS:HA	2.27	0.69
1:A:166:PHE:HE2	1:A:217:SER:CA	2.05	0.69
1:A:207:ASN:ND2	1:A:207:ASN:N	2.39	0.69
1:A:353:TYR:CD2	1:A:397:THR:CG2	2.73	0.69
1:A:1421:ALA:CB	1:A:1456:ASP:HA	2.22	0.69
1:B:14:ILE:HD12	1:B:14:ILE:H	1.57	0.69
1:B:32:LEU:HD11	1:B:444:ASP:CB	2.22	0.69
1:B:115:ILE:HD11	1:B:121:ILE:HG21	1.74	0.69
1:B:235:LYS:HA	1:B:296:ALA:CB	2.21	0.69
1:B:239:VAL:HG22	1:B:281:LEU:HD22	1.74	0.69
1:B:514:ASN:HB3	1:B:518:GLN:NE2	2.06	0.69
1:A:56:CYS:CA	1:A:125:HIS:NE2	2.52	0.69
1:A:65:ARG:NH2	1:A:103:ASN:HA	2.06	0.69
1:A:129:ASN:CB	1:A:441:ARG:HH12	2.05	0.69
1:A:255:GLU:OE2	1:A:255:GLU:CA	2.32	0.69
1:A:478:GLU:HA	1:A:555:LEU:CD2	2.22	0.69
1:B:299:TRP:HA	1:B:299:TRP:CE3	2.27	0.69
1:B:507:GLN:HG3	1:B:563:SER:CA	2.21	0.69
1:B:1281:PHE:O	1:B:1285:SER:N	2.25	0.69
1:A:166:PHE:CE2	1:A:217:SER:CA	2.74	0.69
1:A:250:PHE:O	1:A:264:LEU:HG	1.91	0.69
1:A:707:ILE:CB	1:A:737:LEU:CB	2.70	0.69
1:B:287:VAL:HG12	1:B:304:ARG:HH22	1.56	0.69
1:B:470:ARG:O	1:B:474:THR:HG23	1.91	0.69
1:A:1372:MET:HA	1:A:1375:ILE:CB	2.22	0.69
1:B:117:TYR:HD2	1:B:165:PRO:CD	2.03	0.69
1:B:132:LEU:CB	1:B:149:VAL:HG13	2.23	0.69
1:B:142:LEU:HD11	1:B:200:LEU:HA	1.74	0.69
1:B:391:ARG:HD3	1:B:398:TRP:HE1	1.56	0.69
1:B:397:THR:HB	1:B:422:PRO:O	1.91	0.69
1:B:456:ILE:O	1:B:460:LEU:HD13	1.92	0.69
1:B:678:GLU:HA	1:B:686:ALA:HA	1.74	0.69
1:A:108:ARG:HE	1:A:108:ARG:H	1.41	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:181:LYS:HE2	1:A:219:LYS:HB3	1.69	0.69
1:A:306:LYS:CB	1:A:313:TYR:HD1	2.06	0.69
1:A:315:ALA:O	1:A:356:VAL:O	2.10	0.69
1:A:457:ALA:CB	1:A:522:LEU:HD12	2.23	0.69
1:A:458:GLY:N	1:A:459:LYS:HE3	2.07	0.69
1:B:194:HIS:CD2	1:B:214:CYS:HB2	2.27	0.69
1:B:401:SER:HA	1:B:418:ILE:HG22	1.73	0.69
1:A:19:ALA:C	1:A:20:GLU:OE1	2.35	0.69
1:A:133:THR:CB	1:A:150:THR:O	2.29	0.69
1:A:134:VAL:HG21	1:A:186:PRO:HG2	1.75	0.69
1:A:446:ALA:HB1	1:A:513:GLN:HB3	1.74	0.69
1:A:453:LEU:CB	1:A:522:LEU:CD2	2.71	0.69
1:A:453:LEU:HD22	1:A:453:LEU:H	1.56	0.69
1:A:516:LEU:CD1	1:A:520:PHE:CZ	2.75	0.69
1:B:265:ARG:HG2	1:B:414:VAL:HB	1.74	0.69
1:B:303:PHE:N	1:B:367:ILE:HG21	2.08	0.69
1:A:16:SER:O	1:A:221:VAL:CB	2.30	0.69
1:A:19:ALA:CA	1:A:218:TRP:CZ3	2.76	0.69
1:A:60:LEU:HD13	1:A:123:LEU:CD2	2.22	0.69
1:A:86:ALA:O	1:A:90:ASN:ND2	2.25	0.69
1:A:181:LYS:CG	1:A:219:LYS:CA	2.70	0.69
1:A:242:LEU:HD23	1:A:250:PHE:HE1	1.58	0.69
1:A:246:GLU:CB	1:A:429:ALA:HB3	2.23	0.69
1:A:264:LEU:HB3	1:A:418:ILE:CD1	2.23	0.69
1:A:287:VAL:HG21	1:A:302:LEU:CG	2.22	0.69
1:A:389:ARG:HH12	1:A:428:GLU:CA	2.05	0.69
1:A:446:ALA:HB1	1:A:515:ILE:HG12	1.73	0.69
1:A:520:PHE:HA	1:A:523:LEU:CD1	2.22	0.69
1:B:11:ILE:H	1:B:113:THR:N	1.90	0.69
1:B:12:GLY:O	1:B:59:LYS:HD2	1.92	0.69
1:B:17:LEU:O	1:B:46:LEU:HD21	1.93	0.69
1:B:181:LYS:CG	1:B:219:LYS:HB3	2.23	0.69
1:B:224:MET:HE3	1:B:295:GLY:O	1.92	0.69
1:B:253:CYS:CB	1:B:307:HIS:HE1	2.05	0.69
1:B:441:ARG:CZ	1:B:441:ARG:N	2.54	0.69
1:B:453:LEU:HD11	1:B:515:ILE:CD1	2.15	0.69
1:B:1454:LEU:O	1:B:1458:CYS:N	2.25	0.69
1:A:253:CYS:HB2	1:A:307:HIS:HE1	1.55	0.69
1:B:32:LEU:HB2	1:B:445:PHE:CD1	2.28	0.69
1:B:167:TYR:N	1:B:167:TYR:CD1	2.60	0.69
1:B:249:LYS:HB2	1:B:266:THR:HA	1.73	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:251:LEU:H	1:B:430:PHE:HZ	1.37	0.69
1:B:302:LEU:O	1:B:304:ARG:NH2	2.26	0.69
1:A:305:PHE:CD2	1:A:314:LEU:HD23	2.28	0.69
1:A:458:GLY:N	1:A:459:LYS:NZ	2.41	0.69
1:A:477:LEU:HB2	1:A:555:LEU:HB2	1.70	0.69
1:B:130:LYS:HB3	1:B:151:LEU:CD2	2.23	0.69
1:B:249:LYS:NZ	1:B:266:THR:OG1	2.23	0.69
1:B:361:GLY:O	1:B:366:SER:CB	2.41	0.69
1:A:117:TYR:CE2	1:A:173:GLY:N	2.59	0.68
1:A:200:LEU:HD22	1:A:203:ASN:HB2	1.72	0.68
1:A:256:HIS:H	1:A:261:HIS:CD2	2.12	0.68
1:A:705:LYS:HA	1:A:736:GLN:CB	2.22	0.68
1:B:264:LEU:HG	1:B:418:ILE:CD1	2.23	0.68
1:B:302:LEU:CG	1:B:367:ILE:HG13	2.17	0.68
1:B:476:LEU:HA	1:B:479:ASP:OD2	1.93	0.68
1:B:1133:VAL:CB	1:B:1180:LEU:CB	2.71	0.68
1:A:60:LEU:HD13	1:A:123:LEU:HD23	1.74	0.68
1:A:69:GLN:NE2	1:A:96:ALA:O	2.26	0.68
1:A:193:LEU:HD12	1:A:218:TRP:HE1	1.56	0.68
1:B:440:VAL:HB	1:B:441:ARG:NH1	2.08	0.68
1:B:448:ASP:C	1:B:451:LYS:NZ	2.51	0.68
1:B:769:SER:CB	1:B:782:CYS:CB	2.72	0.68
1:A:105:THR:C	1:A:108:ARG:NH2	2.50	0.68
1:A:125:HIS:HB3	1:A:130:LYS:H	1.59	0.68
1:A:142:LEU:HD13	1:A:208:GLU:CD	2.19	0.68
1:A:285:GLU:C	1:A:304:ARG:HB2	2.17	0.68
1:A:368:PHE:HA	1:A:393:LEU:N	2.08	0.68
1:B:33:VAL:HG11	1:B:452:VAL:HG21	1.74	0.68
1:B:257:ARG:NH1	1:B:257:ARG:HB2	2.07	0.68
1:B:438:ALA:O	1:B:442:ASP:OD1	2.10	0.68
1:B:477:LEU:C	1:B:555:LEU:HD22	2.18	0.68
1:A:26:PHE:CD2	1:A:53:PHE:HE2	2.11	0.68
1:A:523:LEU:HD23	1:A:524:GLN:CD	2.19	0.68
1:B:19:ALA:HB3	1:B:24:ASN:C	2.17	0.68
1:B:281:LEU:CB	1:B:308:LEU:HD23	2.10	0.68
1:A:249:LYS:NZ	1:A:416:LEU:HD13	2.09	0.68
1:A:507:GLN:HE22	1:A:562:HIS:HD2	1.41	0.68
1:B:106:GLU:CG	1:B:107:ASN:OD1	2.37	0.68
1:A:244:HIS:CD2	1:A:430:PHE:CG	2.81	0.68
1:A:298:TYR:CE1	1:A:381:LEU:CB	2.75	0.68
1:A:443:LEU:CA	1:A:509:LEU:HD21	2.19	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:458:GLY:N	1:A:459:LYS:HZ2	1.91	0.68
1:B:71:GLN:HB2	1:B:92:LEU:CD2	2.24	0.68
1:B:263:PHE:CD1	1:B:415:MET:HE3	2.28	0.68
1:A:18:TYR:HB2	1:A:219:LYS:CE	2.22	0.68
1:A:66:TYR:HA	1:A:99:GLU:CD	2.18	0.68
1:A:167:TYR:CE2	1:A:181:LYS:CB	2.75	0.68
1:A:249:LYS:CB	1:A:264:LEU:CD1	2.72	0.68
1:A:459:LYS:N	1:A:459:LYS:HE3	2.09	0.68
1:A:10:HIS:CE1	1:A:114:VAL:HG22	2.28	0.68
1:A:13:ASP:CA	1:A:224:MET:O	2.41	0.68
1:A:400:HIS:O	1:A:418:ILE:HB	1.93	0.68
1:A:407:ASP:N	1:A:407:ASP:OD1	2.24	0.68
1:A:457:ALA:N	1:A:459:LYS:NZ	2.42	0.68
1:A:1287:ILE:CA	1:A:1343:VAL:CB	2.72	0.68
1:A:17:LEU:HA	1:A:219:LYS:NZ	2.09	0.68
1:A:108:ARG:NH2	1:A:109:LYS:N	2.42	0.68
1:A:138:LEU:HD22	1:A:148:ARG:NH2	2.07	0.68
1:A:367:ILE:O	1:A:393:LEU:N	2.26	0.68
1:B:11:ILE:HG13	1:B:113:THR:H	1.58	0.68
1:B:241:ARG:H	1:B:241:ARG:HD3	1.59	0.68
1:A:70:LYS:HA	1:A:73:TRP:CZ2	2.28	0.68
1:A:469:GLU:O	1:A:473:VAL:HG12	1.94	0.68
1:B:111:LEU:HD13	1:B:112:GLY:H	1.59	0.68
1:B:139:PRO:HB3	1:B:146:ALA:N	2.08	0.68
1:B:404:ILE:HD12	1:B:404:ILE:H	1.58	0.68
1:A:19:ALA:HA	1:A:218:TRP:CE3	2.29	0.67
1:A:181:LYS:CD	1:A:219:LYS:HB3	2.23	0.67
1:A:453:LEU:HB3	1:A:522:LEU:CD2	2.24	0.67
1:B:257:ARG:C	1:B:259:LYS:HE2	2.19	0.67
1:A:251:LEU:H	1:A:251:LEU:HD13	1.59	0.67
1:B:395:THR:CG2	1:B:397:THR:HG22	2.25	0.67
1:B:400:HIS:HB2	1:B:419:GLY:O	1.92	0.67
1:B:475:LYS:NZ	1:B:478:GLU:OE2	2.25	0.67
1:A:26:PHE:CE1	1:A:44:GLY:HA3	2.29	0.67
1:A:230:LYS:HA	1:A:230:LYS:HE2	1.76	0.67
1:A:384:ARG:HG3	1:A:385:ASN:CG	2.19	0.67
1:A:407:ASP:HA	1:A:415:MET:HE2	1.75	0.67
1:B:244:HIS:CE1	1:B:247:GLN:HB3	2.29	0.67
1:A:18:TYR:HE2	1:A:26:PHE:CZ	2.12	0.67
1:A:263:PHE:HE2	1:A:414:VAL:HG23	1.59	0.67
1:A:298:TYR:O	1:A:301:SER:HB2	1.95	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:391:ARG:CZ	1:A:392:HIS:O	2.42	0.67
1:A:513:GLN:HE21	1:A:514:ASN:H	1.42	0.67
1:B:29:THR:CG2	1:B:125:HIS:HE1	2.07	0.67
1:B:477:LEU:O	1:B:480:LEU:N	2.26	0.67
1:B:1434:GLU:N	1:B:1493:PRO:O	2.27	0.67
1:A:19:ALA:CA	1:A:218:TRP:HZ3	2.07	0.67
1:A:65:ARG:HH22	1:A:103:ASN:CA	2.06	0.67
1:A:400:HIS:O	1:A:418:ILE:O	2.12	0.67
1:A:457:ALA:O	1:A:460:LEU:N	2.27	0.67
1:B:102:GLN:O	1:B:103:ASN:C	2.35	0.67
1:B:243:PHE:CE1	1:B:250:PHE:HE1	2.11	0.67
1:B:297:GLY:HA2	1:B:303:PHE:CZ	2.30	0.67
1:B:1210:ALA:O	1:B:1214:VAL:CB	2.43	0.67
1:A:243:PHE:O	1:A:430:PHE:HA	1.94	0.67
1:A:369:GLU:N	1:A:393:LEU:HD13	2.09	0.67
1:B:63:MET:CG	1:B:160:TRP:CD1	2.78	0.67
1:B:107:ASN:OD1	1:B:107:ASN:N	2.28	0.67
1:B:194:HIS:HB2	1:B:210:ASN:O	1.95	0.67
1:B:373:THR:HG21	1:B:389:ARG:HE	1.58	0.67
1:B:447:ASN:C	1:B:451:LYS:HZ2	1.95	0.67
1:A:69:GLN:HE22	1:A:100:LYS:HG3	1.60	0.67
1:A:166:PHE:H	1:A:182:VAL:HG12	1.60	0.67
1:A:259:LYS:O	1:A:261:HIS:NE2	2.28	0.67
1:A:264:LEU:CB	1:A:418:ILE:CD1	2.72	0.67
1:A:389:ARG:NH1	1:A:428:GLU:C	2.53	0.67
1:A:1240:PHE:O	1:A:1242:GLN:N	2.28	0.67
1:B:264:LEU:HG	1:B:418:ILE:HD11	1.77	0.67
1:B:394:CYS:SG	1:B:395:THR:N	2.67	0.67
1:B:476:LEU:HD23	1:B:477:LEU:HD23	1.77	0.67
1:A:222:LEU:C	1:A:222:LEU:HD22	2.20	0.67
1:A:406:ILE:HG13	1:A:417:LYS:HA	1.76	0.67
1:A:478:GLU:HA	1:A:555:LEU:HD22	1.77	0.67
1:B:181:LYS:CE	1:B:219:LYS:HG3	2.25	0.67
1:B:242:LEU:HD12	1:B:430:PHE:CD2	2.28	0.67
1:B:384:ARG:HD2	1:B:434:PRO:CD	2.23	0.67
1:B:440:VAL:N	1:B:441:ARG:NH2	2.43	0.67
1:B:508:LYS:HG3	1:B:511:ARG:NH2	2.09	0.67
1:A:15:CYS:SG	1:A:223:PHE:N	2.67	0.67
1:A:49:PRO:O	1:A:53:PHE:HB2	1.93	0.67
1:A:65:ARG:HH12	1:A:103:ASN:H	1.43	0.67
1:A:200:LEU:HD23	1:A:203:ASN:H	1.60	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:254:ASP:HB2	1:A:263:PHE:CZ	2.30	0.67
1:A:373:THR:HG23	1:A:389:ARG:NE	2.09	0.67
1:B:142:LEU:HB3	1:B:198:HIS:HD1	1.59	0.67
1:B:148:ARG:HH12	1:B:150:THR:HB	1.59	0.67
1:B:230:LYS:O	1:B:233:ILE:O	2.12	0.67
1:B:458:GLY:O	1:B:462:LYS:CD	2.43	0.67
1:A:478:GLU:CG	1:A:555:LEU:HD11	2.23	0.67
1:B:386:SER:O	1:B:387:TYR:HB2	1.95	0.67
1:B:407:ASP:HB2	1:B:413:PRO:CB	2.24	0.67
1:B:552:ILE:HD13	1:B:552:ILE:H	1.60	0.67
1:A:49:PRO:CD	1:A:291:PRO:HB3	2.24	0.66
1:A:305:PHE:CZ	1:A:390:LEU:HD21	2.30	0.66
1:B:228:ASP:OD1	1:B:381:LEU:HD21	1.94	0.66
1:B:385:ASN:HA	1:B:432:ILE:H	1.59	0.66
1:B:511:ARG:N	1:B:516:LEU:HD11	2.11	0.66
1:A:9:LEU:HD12	1:A:9:LEU:N	2.08	0.66
1:A:30:LEU:HD21	1:A:38:VAL:HG11	1.76	0.66
1:A:161:PHE:CB	1:A:186:PRO:HA	2.24	0.66
1:A:240:VAL:CA	1:A:433:VAL:O	2.43	0.66
1:A:244:HIS:H	1:A:248:GLU:CA	2.08	0.66
1:A:390:LEU:O	1:A:399:VAL:HB	1.95	0.66
1:B:23:THR:HG21	1:B:41:PRO:HG3	1.77	0.66
1:B:27:ILE:CD1	1:B:209:VAL:HG11	2.25	0.66
1:B:46:LEU:HD12	1:B:219:LYS:CE	2.26	0.66
1:B:243:PHE:CD1	1:B:250:PHE:CE1	2.83	0.66
1:B:365:SER:HB2	1:B:392:HIS:NE2	2.10	0.66
1:B:403:ASN:HA	1:B:416:LEU:HD13	1.78	0.66
1:B:474:THR:HA	1:B:477:LEU:HD11	1.77	0.66
1:A:138:LEU:O	1:A:148:ARG:N	2.27	0.66
1:A:181:LYS:CG	1:A:219:LYS:CB	2.73	0.66
1:A:281:LEU:HD22	1:A:435:VAL:HG13	1.75	0.66
1:A:303:PHE:HB2	1:A:368:PHE:O	1.94	0.66
1:A:560:LEU:HD21	1:A:564:GLN:OE1	1.95	0.66
1:B:130:LYS:CB	1:B:151:LEU:HD22	2.25	0.66
1:B:297:GLY:CA	1:B:303:PHE:CE2	2.78	0.66
1:B:448:ASP:HA	1:B:451:LYS:CE	2.25	0.66
1:B:477:LEU:O	1:B:480:LEU:CD2	2.33	0.66
1:A:372:PRO:HA	1:A:389:ARG:HD2	1.77	0.66
1:A:389:ARG:HH22	1:A:430:PHE:N	1.94	0.66
1:A:507:GLN:HE22	1:A:562:HIS:CD2	2.12	0.66
1:B:148:ARG:NH1	1:B:150:THR:HB	2.10	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:193:LEU:HD23	1:B:193:LEU:C	2.20	0.66
1:B:235:LYS:HD2	1:B:238:ASP:CG	2.19	0.66
1:B:504:ARG:HH22	1:B:567:TYR:CB	2.09	0.66
1:B:746:ASP:CB	1:B:988:TYR:HA	2.25	0.66
1:A:249:LYS:N	1:A:430:PHE:HE1	1.94	0.66
1:A:249:LYS:HB3	1:A:264:LEU:HD13	1.78	0.66
1:A:403:ASN:O	1:A:405:PRO:HD3	1.95	0.66
1:B:23:THR:CG2	1:B:41:PRO:HG3	2.26	0.66
1:B:249:LYS:CE	1:B:416:LEU:HG	2.25	0.66
1:B:1248:ASN:O	1:B:1250:GLN:N	2.28	0.66
1:A:65:ARG:HH22	1:A:103:ASN:N	1.92	0.66
1:A:523:LEU:H	1:A:523:LEU:HD22	1.60	0.66
1:B:116:GLN:O	1:B:117:TYR:HB2	1.96	0.66
1:B:316:ALA:O	1:B:392:HIS:HE1	1.78	0.66
1:B:371:ASP:OD2	1:B:398:TRP:CZ3	2.49	0.66
1:B:474:THR:HA	1:B:477:LEU:HD12	1.76	0.66
1:B:504:ARG:HA	1:B:504:ARG:HE	1.60	0.66
1:B:595:ILE:O	1:B:599:LEU:N	2.21	0.66
1:A:247:GLN:OE1	1:A:249:LYS:HE3	1.95	0.66
1:A:477:LEU:HD12	1:A:555:LEU:CD1	2.23	0.66
1:A:516:LEU:HB2	1:A:520:PHE:CE1	2.30	0.66
1:B:181:LYS:CA	1:B:218:TRP:H	2.09	0.66
1:B:290:ASP:CG	1:B:293:ARG:H	2.02	0.66
1:B:298:TYR:HA	1:B:381:LEU:HD13	1.75	0.66
1:B:448:ASP:C	1:B:451:LYS:CE	2.68	0.66
1:B:477:LEU:HB2	1:B:555:LEU:CG	2.25	0.66
1:B:1130:GLU:O	1:B:1217:LEU:CB	2.44	0.66
1:A:63:MET:HG3	1:A:120:VAL:HG12	1.78	0.66
1:A:299:TRP:O	1:A:300:ASN:CG	2.38	0.66
1:A:400:HIS:O	1:A:418:ILE:CA	2.43	0.66
1:B:36:ARG:HG2	1:B:37:CYS:N	2.11	0.66
1:B:243:PHE:HB3	1:B:431:ALA:HB2	1.76	0.66
1:A:9:LEU:HD21	1:A:220:ILE:HG22	1.77	0.66
1:A:12:GLY:HA3	1:A:111:LEU:CD2	2.26	0.66
1:A:32:LEU:HD11	1:A:445:PHE:CA	2.21	0.66
1:A:985:ARG:HA	1:A:989:ARG:CB	2.26	0.66
1:B:65:ARG:O	1:B:65:ARG:NE	2.26	0.66
1:B:249:LYS:CG	1:B:264:LEU:CD1	2.71	0.66
1:B:287:VAL:HA	1:B:290:ASP:O	1.96	0.66
1:B:456:ILE:HD12	1:B:469:GLU:CD	2.21	0.66
1:A:179:GLY:C	1:A:220:ILE:HB	2.21	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:287:VAL:O	1:A:288:GLN:NE2	2.29	0.66
1:B:130:LYS:CE	1:B:151:LEU:HD22	2.25	0.66
1:B:194:HIS:ND1	1:B:210:ASN:O	2.28	0.66
1:B:305:PHE:O	1:B:313:TYR:CD2	2.48	0.66
1:B:481:VAL:HG12	1:B:558:ARG:CB	2.25	0.66
1:B:504:ARG:HA	1:B:504:ARG:NE	2.10	0.66
1:A:19:ALA:HB3	1:A:24:ASN:N	2.11	0.65
1:A:219:LYS:O	1:A:219:LYS:HD3	1.96	0.65
1:A:243:PHE:HB2	1:A:248:GLU:O	1.95	0.65
1:A:281:LEU:HD22	1:A:435:VAL:CG1	2.25	0.65
1:A:518:GLN:HA	1:A:518:GLN:NE2	2.11	0.65
1:B:181:LYS:HG2	1:B:219:LYS:CA	2.25	0.65
1:B:233:ILE:C	1:B:233:ILE:HD12	2.20	0.65
1:A:244:HIS:CA	1:A:430:PHE:HD1	2.10	0.65
1:A:249:LYS:HZ3	1:A:418:ILE:CG2	2.10	0.65
1:A:305:PHE:CE2	1:A:368:PHE:HB2	2.30	0.65
1:B:63:MET:HG3	1:B:160:TRP:NE1	2.11	0.65
1:B:132:LEU:HB2	1:B:149:VAL:HG12	1.77	0.65
1:B:255:GLU:OE2	1:B:258:LYS:HD3	1.96	0.65
1:B:264:LEU:CB	1:B:418:ILE:HD11	2.26	0.65
1:B:278:SER:OG	1:B:443:LEU:HD11	1.96	0.65
1:B:1251:ASN:CB	1:B:1283:LEU:O	2.44	0.65
1:B:1439:GLU:HA	1:B:1444:ASN:HA	1.78	0.65
1:A:156:ASN:HD21	1:A:159:SER:CB	2.09	0.65
1:A:256:HIS:N	1:A:261:HIS:CD2	2.64	0.65
1:A:769:SER:O	1:A:773:LEU:N	2.29	0.65
1:B:111:LEU:CB	1:B:226:TRP:CZ3	2.74	0.65
1:B:461:GLU:H	1:B:462:LYS:NZ	1.93	0.65
1:B:511:ARG:CA	1:B:516:LEU:HD11	2.27	0.65
1:B:1095:GLN:CB	1:B:1178:GLU:CB	2.73	0.65
1:A:14:ILE:HG12	1:A:59:LYS:HB2	1.78	0.65
1:A:307:HIS:HB2	1:A:312:HIS:O	1.97	0.65
1:A:460:LEU:C	1:A:460:LEU:HD13	2.21	0.65
1:B:130:LYS:HD2	1:B:151:LEU:C	2.20	0.65
1:B:148:ARG:CZ	1:B:150:THR:HG21	2.25	0.65
1:B:1215:LEU:HA	1:B:1218:LEU:CB	2.27	0.65
1:A:55:ASP:HA	1:A:127:LYS:CD	2.27	0.65
1:A:244:HIS:HD2	1:A:430:PHE:CG	2.15	0.65
1:A:314:LEU:CD1	1:A:316:ALA:HB2	2.26	0.65
1:A:391:ARG:HA	1:A:399:VAL:CG2	2.18	0.65
1:A:450:SER:CA	1:A:518:GLN:HG3	2.27	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:15:CYS:SG	1:B:223:PHE:N	2.70	0.65
1:B:183:VAL:HG12	1:B:192:PRO:HB3	1.78	0.65
1:B:372:PRO:HB3	1:B:388:VAL:HG12	1.78	0.65
1:B:1096:ALA:O	1:B:1100:VAL:N	2.28	0.65
1:A:14:ILE:CD1	1:A:224:MET:HB3	2.22	0.65
1:A:55:ASP:OD1	1:A:127:LYS:CD	2.45	0.65
1:A:125:HIS:HB2	1:A:130:LYS:O	1.95	0.65
1:A:242:LEU:CD1	1:A:251:LEU:HD21	2.21	0.65
1:B:141:LEU:HD23	1:B:141:LEU:N	2.12	0.65
1:B:150:THR:HG23	1:B:150:THR:O	1.95	0.65
1:B:399:VAL:CA	1:B:420:THR:N	2.59	0.65
1:A:251:LEU:O	1:A:251:LEU:HD22	1.96	0.65
1:B:203:ASN:ND2	1:B:206:CYS:SG	2.70	0.65
1:B:392:HIS:HE2	1:B:394:CYS:HG	0.66	0.65
1:B:399:VAL:CA	1:B:420:THR:H	2.09	0.65
1:B:1218:LEU:O	1:B:1230:MET:C	2.39	0.65
1:A:72:PHE:CB	1:A:92:LEU:HD22	2.26	0.65
1:A:111:LEU:HD13	1:A:226:TRP:O	1.96	0.65
1:A:219:LYS:CE	1:A:221:VAL:HG23	2.26	0.65
1:B:11:ILE:HG23	1:B:60:LEU:C	2.22	0.65
1:B:61:CYS:SG	1:B:124:LEU:N	2.70	0.65
1:A:7:SER:O	1:A:178:ILE:HD13	1.96	0.65
1:A:279:LYS:O	1:A:282:TRP:CZ2	2.49	0.65
1:B:16:SER:O	1:B:220:ILE:HD13	1.96	0.65
1:B:19:ALA:HB2	1:B:218:TRP:HH2	1.62	0.65
1:B:184:LEU:O	1:B:192:PRO:HA	1.96	0.65
1:B:473:VAL:HG13	1:B:474:THR:H	1.62	0.65
1:B:503:ASN:CG	1:B:506:ARG:HG3	2.22	0.65
1:B:560:LEU:HD22	1:B:564:GLN:CG	2.26	0.65
1:A:14:ILE:HG12	1:A:59:LYS:CB	2.27	0.65
1:A:58:PHE:HE2	1:A:123:LEU:CB	2.01	0.65
1:A:89:LEU:O	1:A:93:HIS:HB2	1.97	0.65
1:A:307:HIS:N	1:A:312:HIS:O	2.30	0.65
1:A:455:SER:C	1:A:459:LYS:HZ3	2.05	0.65
1:B:504:ARG:HH12	1:B:567:TYR:HA	1.62	0.65
1:B:518:GLN:N	1:B:518:GLN:HE21	1.94	0.65
1:B:616:ASP:O	1:B:620:SER:CB	2.45	0.65
1:A:56:CYS:HA	1:A:125:HIS:CE1	2.31	0.64
1:A:239:VAL:HG13	1:A:435:VAL:CG2	2.27	0.64
1:A:390:LEU:HD12	1:A:391:ARG:H	1.62	0.64
1:A:1460:ALA:CB	1:A:1466:ASP:HA	2.26	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1400:LEU:HA	1:B:1442:THR:CA	2.26	0.64
1:A:34:ASP:OD1	1:A:36:ARG:NH2	2.29	0.64
1:A:225:LYS:NZ	1:A:295:GLY:HA2	2.11	0.64
1:A:502:PRO:HD3	1:A:562:HIS:NE2	2.11	0.64
1:A:518:GLN:O	1:A:522:LEU:HD22	1.96	0.64
1:B:17:LEU:HA	1:B:46:LEU:HD11	1.77	0.64
1:B:130:LYS:HD2	1:B:151:LEU:HB3	1.79	0.64
1:B:302:LEU:HD23	1:B:304:ARG:NH2	2.12	0.64
1:B:390:LEU:HD21	1:B:399:VAL:HB	1.78	0.64
1:B:682:THR:O	1:B:686:ALA:HB3	1.97	0.64
1:A:106:GLU:HA	1:A:109:LYS:HZ1	1.62	0.64
1:A:1250:GLN:CB	1:A:1254:LEU:CB	2.75	0.64
1:B:33:VAL:HG11	1:B:452:VAL:CG2	2.27	0.64
1:B:67:SER:O	1:B:71:GLN:HG2	1.96	0.64
1:B:241:ARG:HE	1:B:241:ARG:C	2.05	0.64
1:B:287:VAL:HG12	1:B:304:ARG:NH2	2.11	0.64
1:B:305:PHE:N	1:B:305:PHE:CD1	2.63	0.64
1:A:250:PHE:N	1:A:264:LEU:HD21	2.13	0.64
1:A:449:ALA:O	1:A:452:VAL:CG2	2.45	0.64
1:A:507:GLN:CD	1:A:563:SER:HA	2.22	0.64
1:B:15:CYS:SG	1:B:223:PHE:HB2	2.38	0.64
1:B:117:TYR:CD2	1:B:165:PRO:HD3	2.27	0.64
1:B:453:LEU:HD12	1:B:518:GLN:HB2	1.80	0.64
1:A:11:ILE:HD11	1:A:115:ILE:CD1	2.24	0.64
1:A:108:ARG:CZ	1:A:108:ARG:CB	2.56	0.64
1:A:138:LEU:HB3	1:A:148:ARG:CG	2.28	0.64
1:A:247:GLN:HE22	1:A:416:LEU:CD2	2.10	0.64
1:A:391:ARG:HG2	1:A:392:HIS:N	2.13	0.64
1:A:1059:THR:CB	1:A:1098:LYS:HA	2.28	0.64
1:B:134:VAL:N	1:B:149:VAL:HG22	2.12	0.64
1:B:385:ASN:OD1	1:B:385:ASN:N	2.26	0.64
1:A:20:GLU:CD	1:A:181:LYS:NZ	2.55	0.64
1:A:317:GLU:HG3	1:A:395:THR:HG21	1.79	0.64
1:A:387:TYR:HA	1:A:431:ALA:CB	2.27	0.64
1:A:744:CYS:HA	1:A:1077:SER:CB	2.26	0.64
1:B:18:TYR:HB3	1:B:46:LEU:HG	1.78	0.64
1:B:249:LYS:HG3	1:B:265:ARG:CA	2.26	0.64
1:B:363:ASP:O	1:B:366:SER:N	2.30	0.64
1:A:38:VAL:CB	1:A:206:CYS:HB3	2.27	0.64
1:A:69:GLN:CG	1:A:99:GLU:OE1	2.44	0.64
1:A:224:MET:HE3	1:A:229:ASN:OD1	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:452:VAL:CB	1:A:476:LEU:HD22	2.27	0.64
1:B:11:ILE:HG13	1:B:113:THR:N	2.13	0.64
1:B:26:PHE:CE1	1:B:45:ASP:HA	2.32	0.64
1:B:241:ARG:O	1:B:241:ARG:NE	2.31	0.64
1:B:476:LEU:O	1:B:480:LEU:HD13	1.97	0.64
1:A:389:ARG:NH1	1:A:428:GLU:N	2.46	0.64
1:A:460:LEU:HD13	1:A:460:LEU:O	1.98	0.64
1:A:1274:GLN:HA	1:A:1330:VAL:HA	1.80	0.64
1:B:244:HIS:NE2	1:B:249:LYS:CB	2.56	0.64
1:B:298:TYR:HB3	1:B:381:LEU:CD1	2.28	0.64
1:B:443:LEU:HD12	1:B:509:LEU:CD2	2.27	0.64
1:B:477:LEU:HD12	1:B:555:LEU:HD13	1.80	0.64
1:B:482:TYR:O	1:B:482:TYR:HD1	1.81	0.64
1:A:207:ASN:N	1:A:207:ASN:HD22	1.95	0.64
1:A:256:HIS:C	1:A:258:LYS:H	2.04	0.64
1:A:516:LEU:HD13	1:A:520:PHE:HZ	1.62	0.64
1:B:72:PHE:HA	1:B:92:LEU:HB3	1.79	0.64
1:B:134:VAL:HA	1:B:149:VAL:HG23	1.80	0.64
1:B:234:LEU:N	1:B:234:LEU:HD23	2.12	0.64
1:B:478:GLU:O	1:B:481:VAL:HG22	1.98	0.64
1:A:109:LYS:O	1:A:109:LYS:HG2	1.97	0.64
1:A:216:THR:HG23	1:A:217:SER:N	2.11	0.64
1:A:230:LYS:HE2	1:A:230:LYS:CA	2.28	0.64
1:A:369:GLU:CB	1:A:393:LEU:HD11	2.18	0.64
1:A:1375:ILE:HA	1:A:1379:GLU:H	1.63	0.64
1:B:18:TYR:C	1:B:218:TRP:CZ3	2.76	0.64
1:B:194:HIS:HE1	1:B:212:VAL:H	1.45	0.64
1:B:248:GLU:O	1:B:250:PHE:CE1	2.51	0.64
1:A:15:CYS:CA	1:A:223:PHE:HB2	2.28	0.63
1:A:90:ASN:ND2	1:A:90:ASN:H	1.95	0.63
1:A:103:ASN:O	1:A:104:GLU:O	2.15	0.63
1:A:520:PHE:O	1:A:523:LEU:HD21	1.96	0.63
1:B:133:THR:O	1:B:135:ASN:OD1	2.15	0.63
1:B:135:ASN:HB2	1:B:138:LEU:HD21	1.81	0.63
1:B:249:LYS:CA	1:B:266:THR:HA	2.28	0.63
1:B:372:PRO:CB	1:B:388:VAL:HG12	2.27	0.63
1:A:20:GLU:HG2	1:A:217:SER:HB3	1.79	0.63
1:A:140:ALA:N	1:A:148:ARG:HD3	2.13	0.63
1:B:37:CYS:SG	1:B:209:VAL:CB	2.84	0.63
1:B:224:MET:C	1:B:225:LYS:HD2	2.22	0.63
1:B:400:HIS:CB	1:B:419:GLY:C	2.71	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:264:LEU:HD22	1:A:416:LEU:CD1	2.29	0.63
1:A:983:ASN:O	1:A:987:ASP:CB	2.47	0.63
1:B:242:LEU:HG	1:B:251:LEU:CD2	2.28	0.63
1:B:264:LEU:HB2	1:B:416:LEU:O	1.99	0.63
1:B:385:ASN:H	1:B:432:ILE:HB	1.62	0.63
1:B:476:LEU:CD2	1:B:477:LEU:HD23	2.28	0.63
1:A:64:ASN:HB3	1:A:122:GLN:CD	2.24	0.63
1:A:249:LYS:CB	1:A:430:PHE:CZ	2.80	0.63
1:A:314:LEU:C	1:A:358:VAL:HG22	2.24	0.63
1:B:31:GLY:HA3	1:B:448:ASP:OD2	1.98	0.63
1:B:240:VAL:HA	1:B:434:PRO:HA	1.79	0.63
1:B:399:VAL:HA	1:B:420:THR:N	2.12	0.63
1:B:1090:ARG:CB	1:B:1177:LYS:N	2.61	0.63
1:A:96:ALA:O	1:A:100:LYS:HG3	1.98	0.63
1:A:223:PHE:CD2	1:A:292:CYS:HB3	2.33	0.63
1:A:281:LEU:HD22	1:A:435:VAL:HG22	1.81	0.63
1:A:380:SER:O	1:A:380:SER:OG	2.17	0.63
1:B:184:LEU:O	1:B:192:PRO:C	2.42	0.63
1:B:223:PHE:HD1	1:B:293:ARG:CA	2.11	0.63
1:B:251:LEU:N	1:B:430:PHE:HZ	1.95	0.63
1:B:384:ARG:HD2	1:B:434:PRO:CB	2.28	0.63
1:B:392:HIS:CD2	1:B:394:CYS:HG	2.05	0.63
1:A:33:VAL:HG12	1:A:452:VAL:HG13	1.80	0.63
1:A:198:HIS:O	1:A:208:GLU:HG2	1.98	0.63
1:A:249:LYS:CA	1:A:264:LEU:CD2	2.73	0.63
1:A:1224:LYS:HA	1:A:1271:VAL:CB	2.29	0.63
1:A:1335:VAL:CB	1:A:1385:THR:CB	2.76	0.63
1:B:34:ASP:HB3	1:B:452:VAL:CG1	2.29	0.63
1:B:390:LEU:HD11	1:B:399:VAL:CG2	2.29	0.63
1:B:449:ALA:HB1	1:B:515:ILE:CD1	2.26	0.63
1:B:522:LEU:H	1:B:522:LEU:CD2	2.12	0.63
1:A:124:LEU:HD13	1:A:131:TYR:CD1	2.34	0.63
1:A:1059:THR:CB	1:A:1098:LYS:CA	2.76	0.63
1:B:142:LEU:C	1:B:198:HIS:HD1	2.06	0.63
1:B:522:LEU:N	1:B:522:LEU:CD2	2.61	0.63
1:A:39:VAL:HG12	1:A:207:ASN:OD1	1.99	0.63
1:A:125:HIS:O	1:A:129:ASN:N	2.31	0.63
1:A:1061:LEU:O	1:A:1065:LEU:N	2.23	0.63
1:B:15:CYS:HB3	1:B:220:ILE:CG1	2.29	0.63
1:B:33:VAL:CG1	1:B:476:LEU:HD12	2.27	0.63
1:B:241:ARG:NH2	1:B:433:VAL:CG2	2.57	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:313:TYR:N	1:B:313:TYR:CD1	2.66	0.63
1:B:391:ARG:NH1	1:B:393:LEU:HD13	1.97	0.63
1:B:759:LEU:CB	1:B:762:ASP:CB	2.77	0.63
1:B:759:LEU:O	1:B:762:ASP:N	2.32	0.63
1:B:1288:ASN:H	1:B:1343:VAL:CB	2.11	0.63
1:A:33:VAL:HG13	1:A:448:ASP:C	2.24	0.62
1:A:59:LYS:HG2	1:A:124:LEU:HB3	1.79	0.62
1:A:124:LEU:HA	1:A:131:TYR:HD1	1.63	0.62
1:A:769:SER:CB	1:A:779:ALA:HA	2.29	0.62
1:B:37:CYS:C	1:B:38:VAL:HG13	2.24	0.62
1:B:72:PHE:N	1:B:92:LEU:HD22	2.13	0.62
1:B:301:SER:HB3	1:B:303:PHE:CD1	2.34	0.62
1:B:401:SER:HA	1:B:418:ILE:CG2	2.28	0.62
1:A:39:VAL:N	1:A:207:ASN:O	2.25	0.62
1:A:306:LYS:HE3	1:A:313:TYR:CE1	2.34	0.62
1:A:435:VAL:HG12	1:A:440:VAL:HG22	1.79	0.62
1:B:58:PHE:CE1	1:B:125:HIS:CG	2.87	0.62
1:B:130:LYS:HG2	1:B:151:LEU:CA	2.29	0.62
1:B:138:LEU:CD1	1:B:148:ARG:HD2	2.24	0.62
1:B:241:ARG:HE	1:B:241:ARG:CA	2.11	0.62
1:B:286:VAL:HG12	1:B:288:GLN:CG	2.22	0.62
1:B:1286:GLU:O	1:B:1344:PHE:CB	2.47	0.62
1:A:20:GLU:CG	1:A:181:LYS:CE	2.76	0.62
1:A:20:GLU:CG	1:A:217:SER:CB	2.76	0.62
1:A:156:ASN:ND2	1:A:159:SER:H	1.96	0.62
1:A:160:TRP:O	1:A:187:VAL:HG23	1.98	0.62
1:A:166:PHE:CB	1:A:181:LYS:O	2.47	0.62
1:A:183:VAL:HG12	1:A:192:PRO:HA	1.82	0.62
1:A:282:TRP:CZ3	1:A:308:LEU:HD12	2.33	0.62
1:A:457:ALA:CA	1:A:459:LYS:HZ2	2.12	0.62
1:A:508:LYS:HZ2	1:A:512:GLU:HG3	1.62	0.62
1:B:135:ASN:CG	1:B:148:ARG:HG2	2.24	0.62
1:B:449:ALA:O	1:B:452:VAL:HG22	1.99	0.62
1:B:459:LYS:CA	1:B:462:LYS:HD3	2.20	0.62
1:B:507:GLN:HE21	1:B:563:SER:CA	2.13	0.62
1:A:13:ASP:C	1:A:14:ILE:CG1	2.70	0.62
1:A:264:LEU:C	1:A:416:LEU:HD12	2.24	0.62
1:A:281:LEU:HD22	1:A:435:VAL:CG2	2.28	0.62
1:A:484:VAL:O	1:A:502:PRO:HG3	1.99	0.62
1:A:485:THR:CG2	1:A:562:HIS:HE1	2.12	0.62
1:B:191:GLN:CB	1:B:212:VAL:HA	2.28	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:384:ARG:HH11	1:B:434:PRO:HD3	1.64	0.62
1:B:388:VAL:O	1:B:429:ALA:HA	2.00	0.62
1:A:166:PHE:H	1:A:182:VAL:CG1	2.13	0.62
1:A:512:GLU:OE2	1:A:512:GLU:N	2.32	0.62
1:B:65:ARG:CZ	1:B:106:GLU:OE2	2.47	0.62
1:B:147:MET:SD	1:B:186:PRO:HB3	2.40	0.62
1:A:65:ARG:HH22	1:A:102:GLN:C	2.07	0.62
1:A:124:LEU:HA	1:A:131:TYR:CD1	2.34	0.62
1:A:133:THR:OG1	1:A:152:ASP:N	2.31	0.62
1:A:453:LEU:HD13	1:A:476:LEU:HD21	1.82	0.62
1:A:509:LEU:CD1	1:A:513:GLN:HB2	2.29	0.62
1:B:504:ARG:HH12	1:B:567:TYR:CA	2.11	0.62
1:B:830:LYS:O	1:B:833:PHE:N	2.32	0.62
1:B:1090:ARG:CB	1:B:1177:LYS:H	2.12	0.62
1:A:14:ILE:HG21	1:A:59:LYS:HA	1.79	0.62
1:A:167:TYR:CD2	1:A:180:ASP:OD1	2.45	0.62
1:A:263:PHE:HE2	1:A:414:VAL:CG2	2.13	0.62
1:A:281:LEU:O	1:A:282:TRP:CD2	2.51	0.62
1:B:12:GLY:HA2	1:B:59:LYS:CE	2.29	0.62
1:B:299:TRP:CZ2	1:B:379:ASP:N	2.67	0.62
1:B:406:ILE:HB	1:B:415:MET:HG3	1.80	0.62
1:B:63:MET:SD	1:B:63:MET:C	2.83	0.62
1:B:384:ARG:HD2	1:B:434:PRO:HG3	1.81	0.62
1:B:459:LYS:CA	1:B:462:LYS:CE	2.78	0.62
1:B:475:LYS:HD3	1:B:475:LYS:C	2.22	0.62
1:B:485:THR:H	1:B:506:ARG:NH1	1.96	0.62
1:B:1421:ALA:O	1:B:1425:PHE:CB	2.48	0.62
1:A:481:VAL:CG2	1:A:555:LEU:CD2	2.77	0.62
1:A:1381:LEU:HA	1:A:1385:THR:CB	2.29	0.62
1:B:461:GLU:N	1:B:462:LYS:CE	2.63	0.62
1:A:243:PHE:CA	1:A:248:GLU:O	2.48	0.62
1:B:241:ARG:N	1:B:241:ARG:NE	2.48	0.62
1:B:257:ARG:HH12	1:B:408:LYS:CG	2.01	0.62
1:B:476:LEU:HD23	1:B:480:LEU:HD11	1.81	0.62
1:A:281:LEU:O	1:A:282:TRP:CE3	2.53	0.61
1:A:303:PHE:CE1	1:A:370:LEU:HD21	2.35	0.61
1:A:389:ARG:HH12	1:A:428:GLU:C	2.08	0.61
1:B:167:TYR:CE1	1:B:169:LEU:HD13	2.35	0.61
1:B:196:SER:HB3	1:B:208:GLU:C	2.24	0.61
1:B:240:VAL:HG23	1:B:240:VAL:O	1.99	0.61
1:B:462:LYS:CD	1:B:463:GLY:N	2.60	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:477:LEU:CB	1:B:555:LEU:HD22	2.30	0.61
1:A:32:LEU:CD1	1:A:448:ASP:HB2	2.29	0.61
1:A:61:CYS:SG	1:A:62:PRO:CD	2.89	0.61
1:A:316:ALA:HB2	1:A:355:LEU:HG	1.77	0.61
1:A:395:THR:O	1:A:397:THR:HG23	2.00	0.61
1:B:16:SER:CB	1:B:292:CYS:HB3	2.29	0.61
1:B:241:ARG:CZ	1:B:433:VAL:HB	2.30	0.61
1:B:353:TYR:CE2	1:B:395:THR:HG23	2.35	0.61
1:A:9:LEU:H	1:A:9:LEU:CD1	2.09	0.61
1:A:18:TYR:HB3	1:A:219:LYS:HD2	1.81	0.61
1:A:235:LYS:CE	1:A:384:ARG:HH11	2.12	0.61
1:A:391:ARG:HH22	1:A:395:THR:C	2.07	0.61
1:A:1481:ILE:O	1:A:1485:ILE:CB	2.48	0.61
1:B:242:LEU:HD21	1:B:282:TRP:CD2	2.35	0.61
1:B:456:ILE:HG22	1:B:460:LEU:CD1	2.30	0.61
1:A:10:HIS:NE2	1:A:114:VAL:HG22	2.15	0.61
1:A:181:LYS:HG3	1:A:219:LYS:HB2	1.82	0.61
1:A:244:HIS:CD2	1:A:429:ALA:C	2.78	0.61
1:A:358:VAL:CG1	1:A:366:SER:OG	2.48	0.61
1:A:417:LYS:O	1:A:418:ILE:CG1	2.48	0.61
1:B:207:ASN:ND2	1:B:208:GLU:O	2.34	0.61
1:B:253:CYS:HB2	1:B:307:HIS:CE1	2.35	0.61
1:B:384:ARG:HD2	1:B:434:PRO:HB3	1.82	0.61
1:B:477:LEU:CD1	1:B:555:LEU:HD13	2.30	0.61
1:B:759:LEU:O	1:B:761:VAL:N	2.34	0.61
1:A:317:GLU:O	1:A:353:TYR:HB3	2.00	0.61
1:A:518:GLN:HE21	1:A:518:GLN:CA	2.13	0.61
1:A:1381:LEU:O	1:A:1385:THR:O	2.18	0.61
1:B:26:PHE:CE1	1:B:46:LEU:HD23	2.35	0.61
1:B:29:THR:HG21	1:B:125:HIS:CE1	2.35	0.61
1:B:263:PHE:CE2	1:B:414:VAL:HB	2.35	0.61
1:B:286:VAL:HA	1:B:304:ARG:CD	2.30	0.61
1:B:298:TYR:CB	1:B:381:LEU:HD13	2.30	0.61
1:B:459:LYS:CA	1:B:462:LYS:HE3	2.27	0.61
1:B:503:ASN:CB	1:B:506:ARG:HD2	2.30	0.61
1:A:11:ILE:HG21	1:A:110:LEU:HD21	1.79	0.61
1:A:14:ILE:O	1:A:224:MET:HB2	2.00	0.61
1:A:166:PHE:N	1:A:182:VAL:HG12	2.16	0.61
1:A:175:SER:O	1:A:177:VAL:HG23	2.00	0.61
1:A:220:ILE:HD12	1:A:220:ILE:N	2.16	0.61
1:A:242:LEU:HD23	1:A:250:PHE:CE1	2.35	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:257:ARG:HB2	1:A:259:LYS:HE3	1.78	0.61
1:A:264:LEU:HD23	1:A:264:LEU:C	2.24	0.61
1:B:19:ALA:HA	1:B:218:TRP:CH2	2.35	0.61
1:B:71:GLN:HB2	1:B:92:LEU:HD21	1.83	0.61
1:B:290:ASP:OD2	1:B:293:ARG:CA	2.47	0.61
1:B:390:LEU:CD1	1:B:399:VAL:H	2.13	0.61
1:B:416:LEU:H	1:B:416:LEU:CD2	2.14	0.61
1:B:478:GLU:OE1	1:B:478:GLU:CA	2.46	0.61
1:A:108:ARG:HH21	1:A:109:LYS:H	1.47	0.61
1:B:11:ILE:HG13	1:B:113:THR:CG2	2.27	0.61
1:B:32:LEU:HB2	1:B:445:PHE:CE1	2.35	0.61
1:B:54:ARG:H	1:B:54:ARG:CD	2.06	0.61
1:B:100:LYS:HA	1:B:103:ASN:OD1	2.00	0.61
1:B:251:LEU:N	1:B:430:PHE:CZ	2.67	0.61
1:B:19:ALA:N	1:B:218:TRP:HZ3	1.98	0.61
1:B:42:GLU:O	1:B:43:ALA:C	2.43	0.61
1:B:121:ILE:HD11	1:B:163:ILE:CG2	2.31	0.61
1:B:195:ALA:HA	1:B:209:VAL:CG2	2.19	0.61
1:B:241:ARG:O	1:B:241:ARG:CZ	2.48	0.61
1:B:447:ASN:C	1:B:451:LYS:NZ	2.57	0.61
1:B:462:LYS:CD	1:B:463:GLY:H	2.13	0.61
1:A:244:HIS:H	1:A:248:GLU:C	2.08	0.61
1:A:244:HIS:N	1:A:430:PHE:HD1	1.99	0.61
1:A:509:LEU:O	1:A:513:GLN:HB2	2.01	0.61
1:A:1332:ALA:HB2	1:A:1376:HIS:CB	2.31	0.61
1:B:10:HIS:CE1	1:B:114:VAL:HG23	2.36	0.61
1:B:69:GLN:O	1:B:73:TRP:CB	2.49	0.61
1:B:399:VAL:CG1	1:B:419:GLY:H	2.11	0.61
1:B:449:ALA:O	1:B:452:VAL:CG2	2.49	0.61
1:A:19:ALA:HB2	1:A:218:TRP:CZ3	2.36	0.61
1:A:235:LYS:CG	1:A:238:ASP:HB2	2.26	0.61
1:A:259:LYS:O	1:A:259:LYS:HG2	1.99	0.61
1:B:38:VAL:HB	1:B:207:ASN:H	1.66	0.61
1:B:241:ARG:NH1	1:B:433:VAL:HB	2.16	0.61
1:B:289:HIS:CE1	1:B:290:ASP:HB3	2.35	0.61
1:A:117:TYR:CZ	1:A:171:SER:O	2.54	0.60
1:B:18:TYR:HD1	1:B:26:PHE:HE1	1.46	0.60
1:A:443:LEU:HD13	1:A:509:LEU:HD11	1.83	0.60
1:B:181:LYS:HE3	1:B:219:LYS:HG3	1.82	0.60
1:B:184:LEU:HD22	1:B:193:LEU:CB	2.31	0.60
1:B:236:GLY:N	1:B:296:ALA:HA	2.16	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:399:VAL:HG13	1:B:419:GLY:CA	2.31	0.60
1:A:168:LYS:HE2	1:A:168:LYS:HA	1.84	0.60
1:A:370:LEU:HA	1:A:390:LEU:CD1	2.27	0.60
1:B:225:LYS:HD2	1:B:225:LYS:N	2.16	0.60
1:B:249:LYS:HG3	1:B:265:ARG:C	2.26	0.60
1:B:302:LEU:HD23	1:B:304:ARG:HH22	1.65	0.60
1:B:312:HIS:NE2	1:B:359:PRO:HA	2.15	0.60
1:B:384:ARG:HD2	1:B:434:PRO:HD3	1.83	0.60
1:B:456:ILE:O	1:B:459:LYS:HB2	2.02	0.60
1:B:510:MET:HG2	1:B:516:LEU:CD1	2.31	0.60
1:A:138:LEU:HD22	1:A:148:ARG:HH21	1.65	0.60
1:A:243:PHE:HA	1:A:248:GLU:O	2.01	0.60
1:A:477:LEU:O	1:A:480:LEU:CD1	2.49	0.60
1:A:477:LEU:CB	1:A:555:LEU:HB3	2.30	0.60
1:A:520:PHE:O	1:A:523:LEU:CD2	2.50	0.60
1:B:131:TYR:H	1:B:153:GLU:CA	2.14	0.60
1:B:264:LEU:O	1:B:414:VAL:O	2.18	0.60
1:B:1029:ASP:O	1:B:1033:ILE:CB	2.48	0.60
1:A:106:GLU:CD	1:A:109:LYS:HZ1	2.10	0.60
1:A:123:LEU:O	1:A:131:TYR:CA	2.49	0.60
1:B:33:VAL:H	1:B:448:ASP:CG	2.10	0.60
1:B:1218:LEU:CB	1:B:1234:MET:CB	2.79	0.60
1:A:20:GLU:OE1	1:A:20:GLU:CA	2.47	0.60
1:A:166:PHE:H	1:A:182:VAL:CB	2.15	0.60
1:A:234:LEU:CD2	1:A:432:ILE:HG21	2.28	0.60
1:A:315:ALA:O	1:A:317:GLU:OE1	2.20	0.60
1:A:318:VAL:HG22	1:A:353:TYR:CE2	2.37	0.60
1:A:513:GLN:NE2	1:A:514:ASN:H	1.98	0.60
1:B:10:HIS:O	1:B:60:LEU:HD21	2.02	0.60
1:B:46:LEU:HD12	1:B:219:LYS:HE3	1.83	0.60
1:B:55:ASP:N	1:B:127:LYS:HZ1	2.00	0.60
1:B:57:LEU:HD22	1:B:126:LEU:HD11	1.82	0.60
1:B:287:VAL:HG13	1:B:289:HIS:N	2.11	0.60
1:B:287:VAL:H	1:B:304:ARG:CZ	2.15	0.60
1:B:368:PHE:HA	1:B:393:LEU:CD2	2.32	0.60
1:B:385:ASN:OD1	1:B:433:VAL:CG2	2.48	0.60
1:A:14:ILE:HD13	1:A:126:LEU:HD21	1.82	0.60
1:A:193:LEU:HD12	1:A:218:TRP:NE1	2.17	0.60
1:A:318:VAL:HG13	1:A:353:TYR:CD1	2.36	0.60
1:A:510:MET:CG	1:A:516:LEU:HD23	2.31	0.60
1:A:1125:ILE:O	1:A:1129:SER:CB	2.50	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:10:HIS:HB3	1:B:112:GLY:HA3	1.83	0.60
1:B:19:ALA:CB	1:B:25:GLY:N	2.58	0.60
1:B:224:MET:O	1:B:225:LYS:O	2.20	0.60
1:B:245:ALA:HB2	1:B:387:TYR:CE2	2.36	0.60
1:B:249:LYS:CG	1:B:264:LEU:HD13	2.31	0.60
1:B:520:PHE:CD1	1:B:520:PHE:O	2.55	0.60
1:A:183:VAL:CG1	1:A:192:PRO:HA	2.31	0.60
1:A:200:LEU:CD2	1:A:203:ASN:HB2	2.32	0.60
1:A:255:GLU:OE2	1:A:260:GLN:CA	2.44	0.60
1:A:287:VAL:HG21	1:A:302:LEU:HB3	1.82	0.60
1:A:298:TYR:N	1:A:301:SER:CB	2.54	0.60
1:A:1243:ASN:N	1:A:1244:PHE:HA	2.17	0.60
1:B:58:PHE:CD2	1:B:123:LEU:HD12	2.35	0.60
1:B:167:TYR:HD1	1:B:167:TYR:H	1.50	0.60
1:B:194:HIS:CE1	1:B:212:VAL:N	2.68	0.60
1:B:384:ARG:CD	1:B:434:PRO:HB3	2.32	0.60
1:B:451:LYS:CE	1:B:451:LYS:CA	2.78	0.60
1:B:475:LYS:O	1:B:478:GLU:OE1	2.19	0.60
1:A:20:GLU:CG	1:A:217:SER:HB2	2.31	0.60
1:A:163:ILE:HD13	1:A:184:LEU:HB2	1.82	0.60
1:A:229:ASN:OD1	1:A:229:ASN:N	2.30	0.60
1:A:400:HIS:CE1	1:A:423:LEU:CD1	2.84	0.60
1:A:477:LEU:C	1:A:555:LEU:HD22	2.26	0.60
1:B:143:GLU:HG2	1:B:208:GLU:OE1	2.01	0.60
1:B:480:LEU:HD21	1:B:555:LEU:HD22	1.83	0.60
1:B:1377:LEU:O	1:B:1381:LEU:CB	2.49	0.60
1:A:242:LEU:O	1:A:250:PHE:HA	2.02	0.60
1:A:308:LEU:HD12	1:A:309:ALA:N	2.13	0.60
1:A:516:LEU:HD12	1:A:520:PHE:CZ	2.35	0.60
1:A:871:ARG:HA	1:A:980:PHE:HA	1.83	0.60
1:B:18:TYR:HD1	1:B:26:PHE:CD1	2.19	0.60
1:B:133:THR:CG2	1:B:152:ASP:OD2	2.48	0.60
1:B:384:ARG:HA	1:B:432:ILE:HB	1.84	0.60
1:B:385:ASN:OD1	1:B:433:VAL:N	2.35	0.60
1:B:403:ASN:CG	1:B:416:LEU:HD11	2.26	0.60
1:A:132:LEU:O	1:A:159:SER:HA	2.01	0.59
1:A:198:HIS:CG	1:A:208:GLU:HG3	2.36	0.59
1:B:304:ARG:HH11	1:B:304:ARG:CG	2.15	0.59
1:A:367:ILE:HB	1:A:393:LEU:HD23	1.85	0.59
1:A:1377:LEU:O	1:A:1381:LEU:N	2.35	0.59
1:B:385:ASN:OD1	1:B:433:VAL:CA	2.49	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:20:GLU:CG	1:A:217:SER:HB3	2.33	0.59
1:A:141:LEU:N	1:A:141:LEU:HD23	2.18	0.59
1:A:239:VAL:HG22	1:A:435:VAL:CG2	2.33	0.59
1:A:240:VAL:CB	1:A:433:VAL:O	2.50	0.59
1:A:247:GLN:HB3	1:A:249:LYS:HG2	1.84	0.59
1:A:253:CYS:CB	1:A:262:VAL:HG12	2.32	0.59
1:A:372:PRO:N	1:A:389:ARG:HG2	2.16	0.59
1:A:406:ILE:HD13	1:A:415:MET:HG3	1.84	0.59
1:A:564:GLN:NE2	1:A:570:ASN:CB	2.62	0.59
1:A:1459:ARG:CB	1:A:1463:ASN:O	2.50	0.59
1:B:183:VAL:HG11	1:B:192:PRO:HG3	1.84	0.59
1:B:242:LEU:CD1	1:B:251:LEU:HD22	2.32	0.59
1:B:244:HIS:NE2	1:B:249:LYS:N	2.50	0.59
1:A:29:THR:OG1	1:A:125:HIS:HE1	1.84	0.59
1:A:254:ASP:H	1:A:263:PHE:HE1	1.50	0.59
1:A:306:LYS:HG3	1:A:313:TYR:CD1	2.37	0.59
1:A:365:SER:HA	1:A:394:CYS:HB3	1.84	0.59
1:A:383:PRO:HG2	1:A:386:SER:HB2	1.85	0.59
1:A:477:LEU:CB	1:A:555:LEU:CB	2.73	0.59
1:A:1127:GLU:O	1:A:1131:LEU:CB	2.50	0.59
1:B:54:ARG:NH2	1:B:127:LYS:HZ3	1.99	0.59
1:B:130:LYS:HE3	1:B:151:LEU:CB	2.28	0.59
1:B:141:LEU:C	1:B:142:LEU:HD23	2.28	0.59
1:B:148:ARG:HH11	1:B:148:ARG:CG	2.16	0.59
1:B:191:GLN:HB3	1:B:212:VAL:C	2.28	0.59
1:B:196:SER:HB2	1:B:209:VAL:HA	1.84	0.59
1:B:252:THR:OG1	1:B:265:ARG:HG3	2.02	0.59
1:B:400:HIS:C	1:B:418:ILE:HG22	2.28	0.59
1:B:510:MET:CA	1:B:513:GLN:HE22	1.95	0.59
1:A:26:PHE:CZ	1:A:45:ASP:O	2.56	0.59
1:A:111:LEU:HD22	1:A:226:TRP:CB	2.31	0.59
1:A:227:SER:O	1:A:230:LYS:CE	2.50	0.59
1:B:63:MET:CE	1:B:64:ASN:C	2.76	0.59
1:B:116:GLN:HE22	1:B:173:GLY:CA	2.16	0.59
1:B:240:VAL:CG1	1:B:434:PRO:HB3	2.33	0.59
1:B:249:LYS:CG	1:B:264:LEU:HD12	2.30	0.59
1:B:263:PHE:CZ	1:B:415:MET:HE1	2.37	0.59
1:B:278:SER:CB	1:B:443:LEU:HD11	2.32	0.59
1:B:285:GLU:C	1:B:286:VAL:HG22	2.27	0.59
1:B:400:HIS:O	1:B:418:ILE:CG2	2.47	0.59
1:B:724:LYS:CB	1:B:858:GLU:HA	2.32	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:19:ALA:HB3	1:A:23:THR:C	2.28	0.59
1:A:57:LEU:O	1:A:126:LEU:HG	2.02	0.59
1:A:198:HIS:CB	1:A:208:GLU:CG	2.74	0.59
1:A:288:GLN:HG3	1:A:293:ARG:CB	2.32	0.59
1:A:303:PHE:CZ	1:A:370:LEU:HD21	2.37	0.59
1:A:459:LYS:HE3	1:A:459:LYS:H	1.68	0.59
1:A:481:VAL:HG13	1:A:558:ARG:CB	2.33	0.59
1:A:484:VAL:HG13	1:A:562:HIS:CE1	2.38	0.59
1:B:96:ALA:C	1:B:100:LYS:HE3	2.28	0.59
1:B:1031:GLU:O	1:B:1035:GLU:CB	2.50	0.59
1:A:453:LEU:HD23	1:A:518:GLN:CD	2.27	0.59
1:B:135:ASN:CB	1:B:138:LEU:HD21	2.33	0.59
1:B:198:HIS:O	1:B:200:LEU:N	2.36	0.59
1:B:229:ASN:HB3	1:B:235:LYS:HB3	1.84	0.59
1:A:124:LEU:HD12	1:A:125:HIS:N	2.17	0.59
1:A:391:ARG:HG3	1:A:398:TRP:CG	2.37	0.59
1:B:16:SER:N	1:B:220:ILE:HD11	2.18	0.59
1:B:184:LEU:O	1:B:191:GLN:O	2.20	0.59
1:B:447:ASN:O	1:B:451:LYS:HG3	2.01	0.59
1:A:38:VAL:HB	1:A:206:CYS:CB	2.27	0.59
1:A:139:PRO:O	1:A:148:ARG:HD3	2.02	0.59
1:A:224:MET:HG3	1:A:228:ASP:OD1	2.03	0.59
1:A:282:TRP:CZ2	1:A:307:HIS:CE1	2.91	0.59
1:A:1488:THR:O	1:A:1492:SER:N	2.36	0.59
1:B:54:ARG:CD	1:B:283:GLU:HG2	2.32	0.59
1:A:98:LEU:O	1:A:102:GLN:HB2	2.02	0.59
1:A:249:LYS:CB	1:A:264:LEU:HD13	2.33	0.59
1:A:303:PHE:HB3	1:A:305:PHE:HE1	1.64	0.59
1:A:391:ARG:HG2	1:A:392:HIS:H	1.65	0.59
1:B:99:GLU:O	1:B:103:ASN:OD1	2.21	0.59
1:B:244:HIS:CD2	1:B:250:PHE:H	2.20	0.59
1:B:421:SER:HB2	1:B:425:GLU:C	2.27	0.59
1:A:106:GLU:HA	1:A:109:LYS:CE	2.32	0.58
1:A:395:THR:C	1:A:397:THR:HG23	2.28	0.58
1:B:12:GLY:N	1:B:60:LEU:HD23	2.18	0.58
1:B:250:PHE:O	1:B:264:LEU:HA	2.02	0.58
1:B:381:LEU:O	1:B:382:VAL:HG13	2.02	0.58
1:B:404:ILE:O	1:B:416:LEU:HD22	2.03	0.58
1:B:406:ILE:C	1:B:415:MET:HG2	2.27	0.58
1:B:458:GLY:O	1:B:462:LYS:HG3	2.02	0.58
1:B:484:VAL:HG22	1:B:562:HIS:HB3	1.82	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:15:CYS:HA	1:A:223:PHE:HB3	1.84	0.58
1:A:117:TYR:OH	1:A:174:ASP:N	2.35	0.58
1:A:226:TRP:CD1	1:A:226:TRP:H	2.20	0.58
1:A:516:LEU:CD1	1:A:520:PHE:HZ	2.14	0.58
1:B:18:TYR:CD1	1:B:26:PHE:CE1	2.87	0.58
1:B:195:ALA:C	1:B:209:VAL:HG22	2.27	0.58
1:B:234:LEU:HD13	1:B:238:ASP:OD2	2.03	0.58
1:A:58:PHE:CZ	1:A:123:LEU:C	2.81	0.58
1:A:772:ASN:O	1:A:773:LEU:C	2.42	0.58
1:A:1451:GLU:O	1:A:1455:VAL:N	2.35	0.58
1:B:11:ILE:CG1	1:B:113:THR:HG23	2.30	0.58
1:B:242:LEU:CB	1:B:430:PHE:CD2	2.72	0.58
1:B:302:LEU:O	1:B:302:LEU:HD23	2.02	0.58
1:A:453:LEU:HD23	1:A:518:GLN:HG3	1.84	0.58
1:B:11:ILE:N	1:B:112:GLY:HA2	2.17	0.58
1:B:54:ARG:CG	1:B:283:GLU:HG2	2.33	0.58
1:B:249:LYS:HA	1:B:266:THR:HA	1.84	0.58
1:B:390:LEU:HD11	1:B:399:VAL:H	1.68	0.58
1:B:510:MET:HG2	1:B:516:LEU:CD2	2.32	0.58
1:A:124:LEU:CA	1:A:131:TYR:HD1	2.16	0.58
1:A:287:VAL:HG11	1:A:302:LEU:HD23	1.84	0.58
1:A:452:VAL:HB	1:A:476:LEU:HD22	1.85	0.58
1:B:14:ILE:O	1:B:223:PHE:HB2	2.04	0.58
1:B:111:LEU:HD12	1:B:111:LEU:H	1.68	0.58
1:B:166:PHE:CE1	1:B:183:VAL:HG22	2.38	0.58
1:B:263:PHE:CZ	1:B:265:ARG:HG3	2.39	0.58
1:B:506:ARG:C	1:B:507:GLN:OE1	2.46	0.58
1:B:654:LYS:O	1:B:658:ASN:CB	2.51	0.58
1:B:1207:ASN:O	1:B:1211:HIS:CB	2.51	0.58
1:A:49:PRO:CG	1:A:291:PRO:CA	2.66	0.58
1:A:185:ASN:N	1:A:185:ASN:ND2	2.36	0.58
1:A:355:LEU:HD11	1:A:399:VAL:HG22	1.85	0.58
1:A:453:LEU:CD2	1:A:518:GLN:HB3	2.31	0.58
1:B:33:VAL:CG1	1:B:452:VAL:CG2	2.81	0.58
1:B:287:VAL:HG22	1:B:289:HIS:C	2.29	0.58
1:B:439:GLU:O	1:B:442:ASP:OD1	2.21	0.58
1:B:509:LEU:HD12	1:B:509:LEU:C	2.28	0.58
1:A:11:ILE:HB	1:A:113:THR:OG1	2.02	0.58
1:A:453:LEU:HB3	1:A:522:LEU:HD23	1.86	0.58
1:A:643:LYS:O	1:A:647:VAL:N	2.28	0.58
1:B:239:VAL:HG23	1:B:308:LEU:HD21	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:450:SER:OG	1:B:515:ILE:HB	2.03	0.58
1:B:506:ARG:CB	1:B:507:GLN:OE1	2.50	0.58
1:B:719:ALA:HB1	1:B:727:ARG:CB	2.34	0.58
1:B:1270:ALA:HB3	1:B:1319:GLU:CB	2.34	0.58
1:A:63:MET:HG2	1:A:160:TRP:CD1	2.39	0.58
1:A:281:LEU:HB3	1:A:435:VAL:CG2	2.22	0.58
1:A:315:ALA:CA	1:A:358:VAL:HG22	2.34	0.58
1:A:401:SER:OG	1:A:430:PHE:CZ	2.54	0.58
1:A:407:ASP:OD2	1:A:413:PRO:HB2	2.04	0.58
1:A:709:SER:O	1:A:713:ARG:CB	2.52	0.58
1:A:16:SER:HG	1:A:53:PHE:HZ	1.48	0.58
1:A:121:ILE:HD12	1:A:122:GLN:O	2.04	0.58
1:A:316:ALA:CB	1:A:355:LEU:HA	2.33	0.58
1:A:502:PRO:CG	1:A:562:HIS:NE2	2.67	0.58
1:A:1102:LEU:O	1:A:1103:LEU:C	2.46	0.58
1:B:11:ILE:HG23	1:B:61:CYS:CA	2.33	0.58
1:B:38:VAL:CB	1:B:206:CYS:HB3	2.34	0.58
1:B:54:ARG:HG2	1:B:283:GLU:CG	2.33	0.58
1:B:200:LEU:HD12	1:B:208:GLU:N	2.19	0.58
1:A:20:GLU:OE1	1:A:217:SER:C	2.47	0.58
1:A:227:SER:O	1:A:230:LYS:HE3	2.04	0.58
1:A:233:ILE:HD12	1:A:298:TYR:CE1	2.38	0.58
1:A:416:LEU:O	1:A:418:ILE:HD13	2.04	0.58
1:B:139:PRO:HB3	1:B:145:ASN:C	2.29	0.58
1:B:241:ARG:NE	1:B:241:ARG:CA	2.67	0.58
1:B:404:ILE:CD1	1:B:417:LYS:N	2.65	0.58
1:B:414:VAL:CG2	1:B:415:MET:HE1	2.29	0.58
1:B:744:CYS:C	1:B:745:LEU:O	2.47	0.58
1:A:249:LYS:HD3	1:A:402:THR:N	2.18	0.57
1:A:725:GLU:CB	1:A:862:LEU:HA	2.34	0.57
1:A:1248:ASN:O	1:A:1250:GLN:N	2.37	0.57
1:B:19:ALA:HB1	1:B:23:THR:O	2.04	0.57
1:B:132:LEU:HA	1:B:152:ASP:N	2.16	0.57
1:B:132:LEU:C	1:B:149:VAL:HG13	2.29	0.57
1:B:222:LEU:O	1:B:293:ARG:CB	2.51	0.57
1:B:513:GLN:NE2	1:B:515:ILE:CG2	2.67	0.57
1:A:14:ILE:CG1	1:A:59:LYS:CA	2.65	0.57
1:A:17:LEU:CD1	1:A:218:TRP:CE3	2.86	0.57
1:A:41:PRO:HD3	1:A:207:ASN:CG	2.25	0.57
1:A:104:GLU:O	1:A:108:ARG:NE	2.37	0.57
1:A:122:GLN:NE2	1:A:131:TYR:CD2	2.72	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:219:LYS:NZ	1:A:221:VAL:CG2	2.66	0.57
1:A:237:GLY:HA2	1:A:284:VAL:O	2.05	0.57
1:B:9:LEU:HD11	1:B:115:ILE:HG12	1.86	0.57
1:B:57:LEU:HD13	1:B:126:LEU:HD11	1.85	0.57
1:B:223:PHE:HA	1:B:293:ARG:CA	2.32	0.57
1:B:263:PHE:HE2	1:B:414:VAL:HB	1.69	0.57
1:B:746:ASP:CB	1:B:987:ASP:O	2.52	0.57
1:A:13:ASP:CB	1:A:225:LYS:HA	2.33	0.57
1:A:14:ILE:O	1:A:223:PHE:C	2.47	0.57
1:A:304:ARG:CD	1:A:367:ILE:HG23	2.33	0.57
1:A:1316:VAL:CB	1:A:1326:CYS:CB	2.82	0.57
1:B:11:ILE:CG2	1:B:61:CYS:HA	2.34	0.57
1:A:18:TYR:HE2	1:A:26:PHE:CE1	2.22	0.57
1:A:32:LEU:HD11	1:A:444:ASP:C	2.30	0.57
1:A:256:HIS:O	1:A:259:LYS:CE	2.50	0.57
1:A:369:GLU:H	1:A:393:LEU:HD13	1.68	0.57
1:A:481:VAL:HG21	1:A:555:LEU:CD2	2.35	0.57
1:B:11:ILE:CA	1:B:60:LEU:HD23	2.32	0.57
1:B:167:TYR:CD1	1:B:169:LEU:HD13	2.39	0.57
1:B:453:LEU:HD23	1:B:453:LEU:H	1.68	0.57
1:B:469:GLU:O	1:B:473:VAL:HG12	2.03	0.57
1:B:516:LEU:HD23	1:B:563:SER:HB3	1.85	0.57
1:A:59:LYS:HE2	1:A:124:LEU:HD23	1.87	0.57
1:A:264:LEU:O	1:A:416:LEU:HD12	2.04	0.57
1:B:184:LEU:O	1:B:192:PRO:CA	2.53	0.57
1:B:283:GLU:C	1:B:306:LYS:HB2	2.22	0.57
1:B:399:VAL:HG13	1:B:419:GLY:HA2	1.86	0.57
1:B:449:ALA:N	1:B:451:LYS:HZ1	2.02	0.57
1:B:847:VAL:CB	1:B:898:VAL:N	2.67	0.57
1:A:306:LYS:CA	1:A:313:TYR:HD1	2.18	0.57
1:A:318:VAL:HG12	1:A:352:VAL:O	2.04	0.57
1:A:389:ARG:NH1	1:A:428:GLU:CA	2.68	0.57
1:A:1285:SER:HA	1:A:1292:VAL:CB	2.34	0.57
1:B:66:TYR:O	1:B:69:GLN:HG2	2.05	0.57
1:B:222:LEU:O	1:B:293:ARG:CA	2.52	0.57
1:B:297:GLY:HA3	1:B:303:PHE:CE2	2.40	0.57
1:B:400:HIS:N	1:B:419:GLY:CA	2.55	0.57
1:A:368:PHE:CA	1:A:393:LEU:HD22	2.33	0.57
1:A:1322:PHE:O	1:A:1323:ILE:CB	2.53	0.57
1:B:221:VAL:CG2	1:B:292:CYS:HB2	2.23	0.57
1:B:249:LYS:CG	1:B:265:ARG:N	2.59	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:286:VAL:HG21	1:B:303:PHE:CZ	2.39	0.57
1:B:1090:ARG:CB	1:B:1177:LYS:CB	2.82	0.57
1:A:288:GLN:CG	1:A:293:ARG:CB	2.83	0.57
1:A:386:SER:O	1:A:388:VAL:HG22	2.04	0.57
1:A:406:ILE:N	1:A:417:LYS:HG2	2.19	0.57
1:B:11:ILE:O	1:B:111:LEU:O	2.23	0.57
1:B:57:LEU:HB3	1:B:126:LEU:HG	1.87	0.57
1:B:142:LEU:CB	1:B:198:HIS:HD1	2.17	0.57
1:B:242:LEU:N	1:B:242:LEU:HD23	2.19	0.57
1:B:249:LYS:HA	1:B:266:THR:CA	2.35	0.57
1:B:404:ILE:H	1:B:416:LEU:HD13	1.70	0.57
1:B:477:LEU:C	1:B:555:LEU:CD2	2.77	0.57
1:B:503:ASN:HB3	1:B:506:ARG:CD	2.34	0.57
1:A:117:TYR:CE2	1:A:171:SER:O	2.58	0.57
1:A:181:LYS:O	1:A:182:VAL:CG1	2.52	0.57
1:A:364:ILE:O	1:A:394:CYS:SG	2.50	0.57
1:A:364:ILE:HD12	1:A:367:ILE:HD11	1.86	0.57
1:A:447:ASN:HA	1:A:450:SER:OG	2.05	0.57
1:B:224:MET:HE2	1:B:294:GLY:CA	2.26	0.57
1:B:241:ARG:C	1:B:241:ARG:NE	2.62	0.57
1:B:249:LYS:HG2	1:B:264:LEU:HB3	1.85	0.57
1:B:301:SER:CB	1:B:303:PHE:CE1	2.88	0.57
1:A:41:PRO:HG3	1:A:207:ASN:OD1	2.05	0.57
1:A:126:LEU:HD11	1:A:224:MET:HG2	1.86	0.57
1:A:168:LYS:HA	1:A:168:LYS:CE	2.34	0.57
1:A:251:LEU:N	1:A:264:LEU:HG	2.20	0.57
1:A:368:PHE:CG	1:A:392:HIS:HA	2.39	0.57
1:A:1313:GLN:O	1:A:1323:ILE:HA	2.05	0.57
1:B:15:CYS:SG	1:B:222:LEU:C	2.88	0.57
1:B:16:SER:O	1:B:220:ILE:HA	2.05	0.57
1:B:220:ILE:HD12	1:B:221:VAL:N	2.20	0.57
1:B:250:PHE:C	1:B:264:LEU:HD22	2.30	0.57
1:B:399:VAL:HA	1:B:420:THR:H	1.68	0.57
1:B:461:GLU:H	1:B:462:LYS:HZ3	1.51	0.57
1:A:146:ALA:HB3	1:A:210:ASN:HD22	1.69	0.56
1:A:158:GLY:O	1:A:186:PRO:HB3	2.05	0.56
1:A:244:HIS:CA	1:A:430:PHE:CD1	2.87	0.56
1:A:302:LEU:CD1	1:A:393:LEU:CD2	2.78	0.56
1:A:407:ASP:CA	1:A:415:MET:HE2	2.35	0.56
1:A:458:GLY:C	1:A:459:LYS:HE3	2.29	0.56
1:A:1331:MET:CB	1:A:1377:LEU:CB	2.83	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1460:ALA:HB3	1:A:1466:ASP:HA	1.87	0.56
1:B:185:ASN:ND2	1:B:192:PRO:HD3	2.20	0.56
1:B:241:ARG:H	1:B:241:ARG:NE	2.03	0.56
1:B:245:ALA:HB2	1:B:387:TYR:HE2	1.70	0.56
1:B:503:ASN:ND2	1:B:506:ARG:HG3	2.20	0.56
1:B:519:ILE:HA	1:B:522:LEU:CD2	2.35	0.56
1:A:63:MET:HG2	1:A:160:TRP:CE2	2.40	0.56
1:A:156:ASN:HD21	1:A:159:SER:HB2	1.69	0.56
1:A:699:PHE:O	1:A:702:ASP:CB	2.52	0.56
1:B:264:LEU:CB	1:B:416:LEU:HB3	2.34	0.56
1:B:298:TYR:HB3	1:B:381:LEU:HD13	1.87	0.56
1:B:398:TRP:C	1:B:420:THR:HB	2.31	0.56
1:B:451:LYS:N	1:B:451:LYS:CE	2.68	0.56
1:A:26:PHE:CD2	1:A:53:PHE:CE2	2.93	0.56
1:A:72:PHE:N	1:A:92:LEU:HD21	2.20	0.56
1:A:282:TRP:HE3	1:A:308:LEU:H	1.53	0.56
1:A:291:PRO:HB2	1:A:292:CYS:SG	2.45	0.56
1:A:365:SER:O	1:A:392:HIS:NE2	2.36	0.56
1:A:368:PHE:HA	1:A:393:LEU:H	1.71	0.56
1:A:417:LYS:C	1:A:418:ILE:HD13	2.30	0.56
1:A:417:LYS:NZ	1:A:419:GLY:HA3	2.20	0.56
1:A:443:LEU:CD1	1:A:513:GLN:HG3	2.35	0.56
1:A:457:ALA:N	1:A:459:LYS:HD3	2.16	0.56
1:B:135:ASN:HB2	1:B:138:LEU:CD2	2.36	0.56
1:B:141:LEU:H	1:B:141:LEU:CD2	2.16	0.56
1:B:180:ASP:C	1:B:219:LYS:HA	2.29	0.56
1:B:363:ASP:O	1:B:366:SER:CB	2.46	0.56
1:B:523:LEU:HD13	1:B:523:LEU:N	2.21	0.56
1:A:259:LYS:HD2	1:A:261:HIS:CE1	2.40	0.56
1:A:282:TRP:HA	1:A:282:TRP:HE3	1.71	0.56
1:A:367:ILE:O	1:A:393:LEU:CB	2.52	0.56
1:A:373:THR:CG2	1:A:389:ARG:HD3	2.36	0.56
1:A:374:THR:O	1:A:374:THR:OG1	2.24	0.56
1:A:666:ILE:CB	1:A:667:GLU:C	2.78	0.56
1:A:1300:GLU:CB	1:A:1304:ARG:CB	2.83	0.56
1:B:391:ARG:HB2	1:B:398:TRP:CE2	2.40	0.56
1:A:184:LEU:C	1:A:185:ASN:ND2	2.60	0.56
1:B:14:ILE:HG22	1:B:57:LEU:CD2	2.25	0.56
1:B:60:LEU:HD23	1:B:60:LEU:O	2.06	0.56
1:B:138:LEU:HD11	1:B:148:ARG:CD	2.27	0.56
1:B:181:LYS:HB3	1:B:218:TRP:N	2.20	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:249:LYS:HA	1:B:265:ARG:C	2.30	0.56
1:B:256:HIS:HE1	1:B:407:ASP:CG	2.13	0.56
1:B:297:GLY:CA	1:B:303:PHE:CZ	2.88	0.56
1:B:368:PHE:HA	1:B:393:LEU:HD23	1.87	0.56
1:B:1198:ARG:CB	1:B:1239:GLU:H	2.19	0.56
1:A:33:VAL:HG12	1:A:452:VAL:CG1	2.35	0.56
1:A:181:LYS:O	1:A:182:VAL:HG13	2.06	0.56
1:A:453:LEU:O	1:A:522:LEU:CG	2.47	0.56
1:B:72:PHE:HB2	1:B:92:LEU:O	2.06	0.56
1:B:238:ASP:O	1:B:283:GLU:HA	2.05	0.56
1:B:355:LEU:N	1:B:417:LYS:HZ1	2.04	0.56
1:A:65:ARG:CA	1:A:160:TRP:HH2	2.13	0.56
1:A:87:VAL:C	1:A:90:ASN:HD22	2.14	0.56
1:A:254:ASP:O	1:A:261:HIS:HB2	2.05	0.56
1:A:477:LEU:HB2	1:A:555:LEU:CD1	2.35	0.56
1:B:238:ASP:H	1:B:284:VAL:HG23	1.71	0.56
1:B:363:ASP:H	1:B:366:SER:CB	2.19	0.56
1:B:1347:ASP:H	1:B:1350:SER:CB	2.19	0.56
1:A:9:LEU:HD11	1:A:179:GLY:CA	2.36	0.56
1:A:14:ILE:HG21	1:A:60:LEU:N	2.21	0.56
1:A:235:LYS:NZ	1:A:235:LYS:H	2.04	0.56
1:A:455:SER:C	1:A:459:LYS:NZ	2.63	0.56
1:B:55:ASP:O	1:B:125:HIS:CD2	2.59	0.56
1:B:220:ILE:HD12	1:B:220:ILE:C	2.31	0.56
1:B:249:LYS:HA	1:B:265:ARG:O	2.05	0.56
1:B:443:LEU:CD1	1:B:509:LEU:CD1	2.81	0.56
1:B:514:ASN:OD1	1:B:517:LYS:CB	2.53	0.56
1:A:20:GLU:CD	1:A:217:SER:CB	2.79	0.56
1:A:26:PHE:HZ	1:A:45:ASP:O	1.89	0.56
1:A:131:TYR:HB3	1:A:159:SER:OG	2.04	0.56
1:A:239:VAL:HG22	1:A:435:VAL:HG21	1.88	0.56
1:A:263:PHE:CB	1:A:406:ILE:HD13	2.36	0.56
1:A:406:ILE:HG13	1:A:417:LYS:CG	2.35	0.56
1:A:1371:LEU:CB	1:A:1374:HIS:CB	2.84	0.56
1:A:1372:MET:O	1:A:1375:ILE:N	2.38	0.56
1:B:42:GLU:O	1:B:44:GLY:N	2.39	0.56
1:B:54:ARG:C	1:B:127:LYS:HZ1	2.14	0.56
1:B:101:LYS:O	1:B:102:GLN:C	2.49	0.56
1:B:166:PHE:CE1	1:B:216:THR:O	2.55	0.56
1:B:181:LYS:HE2	1:B:219:LYS:HG3	1.87	0.56
1:B:392:HIS:CD2	1:B:394:CYS:SG	2.98	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:518:GLN:NE2	1:B:518:GLN:CA	2.69	0.56
1:A:15:CYS:SG	1:A:222:LEU:HA	2.46	0.56
1:A:20:GLU:CD	1:A:217:SER:C	2.73	0.56
1:A:87:VAL:HG12	1:A:91:LYS:CE	2.36	0.56
1:B:11:ILE:HB	1:B:112:GLY:HA2	1.87	0.56
1:B:233:ILE:C	1:B:234:LEU:HD23	2.31	0.56
1:B:242:LEU:HD21	1:B:282:TRP:CE2	2.40	0.56
1:B:302:LEU:O	1:B:367:ILE:HD12	2.06	0.56
1:B:481:VAL:O	1:B:484:VAL:O	2.23	0.56
1:A:249:LYS:HB3	1:A:264:LEU:CD1	2.35	0.55
1:A:404:ILE:O	1:A:416:LEU:C	2.49	0.55
1:B:14:ILE:HB	1:B:223:PHE:HE2	1.70	0.55
1:B:314:LEU:HD23	1:B:314:LEU:C	2.31	0.55
1:B:404:ILE:O	1:B:416:LEU:CA	2.51	0.55
1:B:424:LYS:HG3	1:B:425:GLU:N	2.21	0.55
1:B:1000:GLU:CB	1:B:1006:SER:CB	2.84	0.55
1:B:1270:ALA:HB2	1:B:1315:ILE:C	2.31	0.55
1:A:123:LEU:C	1:A:131:TYR:HD1	2.14	0.55
1:A:203:ASN:N	1:A:204:PRO:HD3	2.21	0.55
1:A:285:GLU:O	1:A:304:ARG:CB	2.44	0.55
1:A:290:ASP:CB	1:A:291:PRO:HD2	2.36	0.55
1:B:71:GLN:CB	1:B:92:LEU:HD21	2.35	0.55
1:B:233:ILE:HD12	1:B:234:LEU:O	2.06	0.55
1:B:298:TYR:HB3	1:B:381:LEU:CG	2.35	0.55
1:A:18:TYR:HE2	1:A:26:PHE:CE2	2.24	0.55
1:A:32:LEU:H	1:A:448:ASP:CG	2.08	0.55
1:A:40:GLN:OE1	1:A:43:ALA:HB2	2.06	0.55
1:A:48:ASN:N	1:A:49:PRO:CD	2.64	0.55
1:A:219:LYS:O	1:A:219:LYS:CD	2.55	0.55
1:B:259:LYS:NZ	1:B:261:HIS:HE1	2.03	0.55
1:A:220:ILE:N	1:A:220:ILE:CD1	2.69	0.55
1:A:391:ARG:HE	1:A:392:HIS:C	2.14	0.55
1:A:1317:LYS:HA	1:A:1322:PHE:O	2.06	0.55
1:B:18:TYR:CD1	1:B:26:PHE:HE1	2.23	0.55
1:B:46:LEU:HD12	1:B:219:LYS:HE2	1.87	0.55
1:B:65:ARG:NH1	1:B:106:GLU:OE2	2.40	0.55
1:B:200:LEU:HD12	1:B:207:ASN:HA	1.89	0.55
1:B:504:ARG:NH2	1:B:567:TYR:CB	2.69	0.55
1:A:30:LEU:HD12	1:A:36:ARG:HB2	1.88	0.55
1:A:161:PHE:HA	1:A:187:VAL:N	2.22	0.55
1:A:241:ARG:CB	1:A:250:PHE:CZ	2.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:457:ALA:C	1:A:459:LYS:HZ2	2.14	0.55
1:B:192:PRO:O	1:B:212:VAL:N	2.39	0.55
1:B:194:HIS:HE1	1:B:212:VAL:C	2.14	0.55
1:B:299:TRP:CG	1:B:380:SER:HA	2.42	0.55
1:B:458:GLY:O	1:B:462:LYS:CG	2.55	0.55
1:A:310:THR:HB	1:A:312:HIS:NE2	2.22	0.55
1:B:35:ASP:HB2	1:B:151:LEU:HD12	1.87	0.55
1:B:220:ILE:HG23	1:B:220:ILE:O	2.06	0.55
1:B:390:LEU:C	1:B:390:LEU:HD12	2.31	0.55
1:B:1441:TYR:O	1:B:1442:THR:CB	2.54	0.55
1:A:11:ILE:O	1:A:13:ASP:OD2	2.23	0.55
1:A:30:LEU:HB3	1:A:34:ASP:HB2	1.88	0.55
1:A:33:VAL:N	1:A:448:ASP:HB3	2.19	0.55
1:A:249:LYS:CB	1:A:264:LEU:HD11	2.37	0.55
1:A:256:HIS:C	1:A:258:LYS:N	2.63	0.55
1:A:290:ASP:OD1	1:A:291:PRO:HD2	2.07	0.55
1:A:318:VAL:HG13	1:A:353:TYR:CD2	2.41	0.55
1:A:511:ARG:CA	1:A:516:LEU:HG	2.31	0.55
1:B:71:GLN:O	1:B:74:LYS:N	2.27	0.55
1:B:150:THR:C	1:B:152:ASP:OD1	2.50	0.55
1:B:194:HIS:HE1	1:B:212:VAL:N	2.02	0.55
1:B:259:LYS:HG2	1:B:261:HIS:CE1	2.41	0.55
1:B:299:TRP:H	1:B:381:LEU:H	1.55	0.55
1:B:312:HIS:HE2	1:B:359:PRO:HB3	1.72	0.55
1:B:405:PRO:HB2	1:B:413:PRO:HB2	1.87	0.55
1:B:1000:GLU:CB	1:B:1006:SER:N	2.68	0.55
1:A:18:TYR:CE1	1:A:25:GLY:CA	2.90	0.55
1:A:58:PHE:CE2	1:A:123:LEU:CB	2.81	0.55
1:A:405:PRO:HA	1:A:416:LEU:CA	2.35	0.55
1:A:879:TYR:H	1:A:884:LEU:CB	2.20	0.55
1:B:40:GLN:NE2	1:B:205:GLY:O	2.40	0.55
1:B:200:LEU:HD12	1:B:207:ASN:CA	2.36	0.55
1:B:256:HIS:CE1	1:B:407:ASP:OD1	2.60	0.55
1:B:1292:VAL:CB	1:B:1344:PHE:CB	2.85	0.55
1:A:11:ILE:HG22	1:A:12:GLY:N	2.20	0.55
1:A:256:HIS:O	1:A:258:LYS:N	2.40	0.55
1:A:458:GLY:CA	1:A:459:LYS:CE	2.78	0.55
1:B:17:LEU:N	1:B:46:LEU:HD11	2.22	0.55
1:B:72:PHE:CA	1:B:92:LEU:HD13	2.35	0.55
1:B:134:VAL:HA	1:B:149:VAL:CG2	2.36	0.55
1:B:299:TRP:CD2	1:B:380:SER:HA	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:373:THR:HG23	1:B:389:ARG:NE	2.17	0.55
1:B:397:THR:CB	1:B:422:PRO:HA	2.37	0.55
1:B:435:VAL:CG1	1:B:439:GLU:OE2	2.55	0.55
1:B:456:ILE:HG23	1:B:469:GLU:OE1	2.07	0.55
1:A:11:ILE:HG21	1:A:110:LEU:HD11	1.88	0.55
1:A:27:ILE:O	1:A:56:CYS:HB2	2.07	0.55
1:A:106:GLU:HA	1:A:109:LYS:NZ	2.21	0.55
1:A:263:PHE:CD2	1:A:414:VAL:O	2.60	0.55
1:A:522:LEU:HD13	1:A:522:LEU:N	2.20	0.55
1:A:1382:ALA:HB1	1:A:1423:ILE:O	2.07	0.55
1:B:130:LYS:HG2	1:B:151:LEU:HA	1.89	0.55
1:B:138:LEU:CG	1:B:148:ARG:HB3	2.22	0.55
1:B:242:LEU:HD21	1:B:282:TRP:CE3	2.41	0.55
1:B:313:TYR:CZ	1:B:361:GLY:CA	2.88	0.55
1:B:473:VAL:HG22	1:B:477:LEU:HD21	1.87	0.55
1:B:1129:SER:O	1:B:1130:GLU:C	2.47	0.55
1:A:17:LEU:HA	1:A:219:LYS:HZ2	1.71	0.54
1:A:301:SER:O	1:A:302:LEU:CB	2.55	0.54
1:A:407:ASP:H	1:A:415:MET:CG	2.20	0.54
1:A:437:PRO:O	1:A:441:ARG:HG3	2.07	0.54
1:B:145:ASN:O	1:B:146:ALA:CB	2.55	0.54
1:B:1130:GLU:C	1:B:1217:LEU:CB	2.81	0.54
1:B:1206:ARG:O	1:B:1210:ALA:HB3	2.07	0.54
1:A:256:HIS:HB3	1:A:261:HIS:NE2	2.22	0.54
1:A:365:SER:HA	1:A:394:CYS:SG	2.47	0.54
1:A:446:ALA:HB3	1:A:513:GLN:OE1	2.06	0.54
1:A:1248:ASN:O	1:A:1249:GLN:C	2.49	0.54
1:B:54:ARG:HD2	1:B:306:LYS:NZ	2.22	0.54
1:B:133:THR:N	1:B:152:ASP:HB2	2.23	0.54
1:B:371:ASP:HB3	1:B:389:ARG:C	2.32	0.54
1:B:387:TYR:O	1:B:388:VAL:CG1	2.51	0.54
1:A:18:TYR:H	1:A:219:LYS:HD3	1.72	0.54
1:A:33:VAL:CG1	1:A:452:VAL:HG13	2.36	0.54
1:A:59:LYS:CG	1:A:124:LEU:HB3	2.37	0.54
1:A:302:LEU:HD12	1:A:393:LEU:HD21	1.86	0.54
1:A:1416:PRO:CB	1:A:1462:ASN:CB	2.86	0.54
1:B:20:GLU:CG	1:B:181:LYS:HZ3	2.21	0.54
1:B:29:THR:CG2	1:B:125:HIS:CE1	2.88	0.54
1:B:54:ARG:H	1:B:54:ARG:HH11	1.52	0.54
1:B:235:LYS:HA	1:B:296:ALA:HB2	1.90	0.54
1:B:296:ALA:O	1:B:303:PHE:HE2	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:507:GLN:NE2	1:B:562:HIS:C	2.59	0.54
1:A:33:VAL:HG21	1:A:451:LYS:HG2	1.88	0.54
1:A:166:PHE:H	1:A:182:VAL:HB	1.71	0.54
1:A:253:CYS:HG	1:A:307:HIS:CD2	2.24	0.54
1:A:367:ILE:C	1:A:368:PHE:HD1	2.15	0.54
1:A:481:VAL:HG21	1:A:555:LEU:HD23	1.88	0.54
1:B:7:SER:HG	1:B:178:ILE:HB	1.70	0.54
1:B:399:VAL:C	1:B:420:THR:OG1	2.50	0.54
1:B:505:GLU:OE2	1:B:505:GLU:N	2.39	0.54
1:B:1218:LEU:CB	1:B:1234:MET:CA	2.86	0.54
1:B:1315:ILE:O	1:B:1316:VAL:C	2.50	0.54
1:B:1369:SER:O	1:B:1370:PRO:CB	2.54	0.54
1:A:146:ALA:CB	1:A:210:ASN:HD22	2.21	0.54
1:A:163:ILE:HD12	1:A:164:GLN:N	2.22	0.54
1:A:248:GLU:O	1:A:249:LYS:C	2.49	0.54
1:A:249:LYS:HB2	1:A:264:LEU:HD11	1.89	0.54
1:A:407:ASP:O	1:A:408:LYS:HD3	2.08	0.54
1:A:450:SER:O	1:A:451:LYS:C	2.49	0.54
1:A:1328:ASP:CB	1:A:1374:HIS:N	2.71	0.54
1:B:34:ASP:N	1:B:448:ASP:OD1	2.41	0.54
1:B:111:LEU:C	1:B:226:TRP:CZ3	2.85	0.54
1:B:132:LEU:CB	1:B:149:VAL:HG12	2.35	0.54
1:B:254:ASP:OD1	1:B:279:LYS:NZ	2.40	0.54
1:B:263:PHE:CE2	1:B:265:ARG:HG2	2.43	0.54
1:B:1090:ARG:H	1:B:1177:LYS:H	1.54	0.54
1:A:49:PRO:O	1:A:50:PRO:O	2.26	0.54
1:A:264:LEU:HD22	1:A:416:LEU:HD12	1.89	0.54
1:A:453:LEU:HD22	1:A:453:LEU:N	2.22	0.54
1:A:621:LEU:CB	1:A:630:PHE:CB	2.85	0.54
1:B:106:GLU:HG3	1:B:107:ASN:N	2.23	0.54
1:B:244:HIS:HB3	1:B:430:PHE:CD1	2.41	0.54
1:B:298:TYR:N	1:B:381:LEU:HD13	2.21	0.54
1:B:451:LYS:CE	1:B:452:VAL:N	2.51	0.54
1:B:1248:ASN:CB	1:B:1283:LEU:O	2.55	0.54
1:A:18:TYR:CB	1:A:219:LYS:CE	2.83	0.54
1:B:249:LYS:HD3	1:B:266:THR:OG1	2.08	0.54
1:B:390:LEU:CD2	1:B:399:VAL:HB	2.38	0.54
1:B:476:LEU:O	1:B:479:ASP:CB	2.53	0.54
1:B:507:GLN:NE2	1:B:563:SER:CA	2.63	0.54
1:A:164:GLN:OE1	1:A:164:GLN:N	2.40	0.54
1:A:165:PRO:HB2	1:A:170:ARG:O	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:166:PHE:CE2	1:A:217:SER:CB	2.91	0.54
1:A:281:LEU:O	1:A:308:LEU:HG	2.07	0.54
1:A:287:VAL:HG21	1:A:302:LEU:CB	2.38	0.54
1:A:456:ILE:HG12	1:A:473:VAL:HB	1.90	0.54
1:B:33:VAL:HG12	1:B:452:VAL:HG21	1.86	0.54
1:B:196:SER:HB2	1:B:210:ASN:N	2.23	0.54
1:B:238:ASP:O	1:B:284:VAL:HG22	2.08	0.54
1:A:124:LEU:HD12	1:A:130:LYS:C	2.33	0.54
1:A:263:PHE:HE2	1:A:414:VAL:CB	2.21	0.54
1:A:456:ILE:CA	1:A:459:LYS:CD	2.79	0.54
1:A:518:GLN:NE2	1:A:518:GLN:CA	2.66	0.54
1:A:1274:GLN:O	1:A:1333:GLU:CB	2.56	0.54
1:B:203:ASN:HB3	1:B:206:CYS:CB	2.34	0.54
1:B:256:HIS:HE1	1:B:407:ASP:OD1	1.91	0.54
1:B:298:TYR:CA	1:B:381:LEU:HA	2.17	0.54
1:B:390:LEU:HD11	1:B:399:VAL:N	2.22	0.54
1:A:46:LEU:O	1:A:291:PRO:CD	2.54	0.54
1:A:134:VAL:HG21	1:A:186:PRO:CG	2.38	0.54
1:A:181:LYS:NZ	1:A:219:LYS:CB	2.53	0.54
1:A:244:HIS:N	1:A:430:PHE:CD1	2.76	0.54
1:A:307:HIS:O	1:A:311:GLY:CA	2.56	0.54
1:A:1376:HIS:HA	1:A:1380:LEU:CB	2.37	0.54
1:A:1442:THR:O	1:A:1445:HIS:N	2.41	0.54
1:B:14:ILE:C	1:B:223:PHE:CD2	2.86	0.54
1:B:26:PHE:CZ	1:B:45:ASP:C	2.85	0.54
1:B:193:LEU:HA	1:B:211:SER:CB	2.38	0.54
1:B:194:HIS:CB	1:B:210:ASN:O	2.56	0.54
1:B:240:VAL:HB	1:B:433:VAL:O	2.08	0.54
1:B:316:ALA:O	1:B:392:HIS:CE1	2.61	0.54
1:B:384:ARG:HB3	1:B:433:VAL:HA	1.90	0.54
1:B:515:ILE:HA	1:B:518:GLN:HB2	1.88	0.54
1:A:139:PRO:HA	1:A:147:MET:C	2.33	0.53
1:A:314:LEU:O	1:A:358:VAL:CG2	2.56	0.53
1:A:371:ASP:O	1:A:389:ARG:HG2	2.09	0.53
1:A:502:PRO:CD	1:A:562:HIS:NE2	2.71	0.53
1:B:237:GLY:CA	1:B:283:GLU:OE2	2.53	0.53
1:B:506:ARG:O	1:B:510:MET:SD	2.66	0.53
1:A:9:LEU:CD2	1:A:220:ILE:CG2	2.87	0.53
1:A:87:VAL:HG12	1:A:91:LYS:HE3	1.90	0.53
1:A:399:VAL:HA	1:A:420:THR:CB	2.38	0.53
1:A:1341:VAL:C	1:A:1343:VAL:H	2.16	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:20:GLU:CG	1:B:181:LYS:HD2	2.39	0.53
1:B:1032:HIS:O	1:B:1036:GLN:N	2.42	0.53
1:B:1191:SER:HA	1:B:1237:ALA:HA	1.90	0.53
1:B:1198:ARG:CB	1:B:1239:GLU:N	2.71	0.53
1:A:8:PHE:N	1:A:8:PHE:CD1	2.76	0.53
1:A:14:ILE:CG2	1:A:60:LEU:N	2.72	0.53
1:A:124:LEU:HD13	1:A:131:TYR:CE1	2.43	0.53
1:A:389:ARG:HB3	1:A:389:ARG:CZ	2.38	0.53
1:A:450:SER:CA	1:A:453:LEU:CD2	2.86	0.53
1:B:181:LYS:CD	1:B:218:TRP:N	2.70	0.53
1:B:263:PHE:CD2	1:B:415:MET:HE3	2.44	0.53
1:B:481:VAL:CG1	1:B:558:ARG:CB	2.86	0.53
1:B:885:LEU:O	1:B:888:THR:N	2.37	0.53
1:A:57:LEU:CB	1:A:126:LEU:HB2	2.37	0.53
1:A:184:LEU:C	1:A:184:LEU:HD23	2.33	0.53
1:A:244:HIS:HA	1:A:430:PHE:HD1	1.72	0.53
1:A:373:THR:CG2	1:A:389:ARG:CD	2.85	0.53
1:A:449:ALA:O	1:A:453:LEU:CD2	2.55	0.53
1:A:477:LEU:O	1:A:480:LEU:HD12	2.08	0.53
1:A:562:HIS:CD2	1:A:562:HIS:C	2.87	0.53
1:A:1232:GLU:HA	1:A:1236:LEU:H	1.73	0.53
1:B:33:VAL:HG21	1:B:449:ALA:HB2	1.91	0.53
1:B:162:TYR:CD1	1:B:187:VAL:HA	2.44	0.53
1:B:316:ALA:CA	1:B:355:LEU:HG	2.37	0.53
1:B:462:LYS:CE	1:B:463:GLY:H	2.21	0.53
1:B:476:LEU:CD2	1:B:480:LEU:HD11	2.38	0.53
1:B:477:LEU:HD12	1:B:555:LEU:CD1	2.37	0.53
1:B:744:CYS:O	1:B:745:LEU:CB	2.55	0.53
1:A:304:ARG:C	1:A:305:PHE:CD1	2.86	0.53
1:A:373:THR:HG23	1:A:389:ARG:HD3	1.91	0.53
1:A:406:ILE:CD1	1:A:417:LYS:HA	2.39	0.53
1:A:1332:ALA:HB2	1:A:1376:HIS:C	2.33	0.53
1:B:148:ARG:HG2	1:B:148:ARG:O	2.09	0.53
1:B:244:HIS:CB	1:B:430:PHE:CE1	2.81	0.53
1:B:440:VAL:HB	1:B:441:ARG:HH22	1.72	0.53
1:B:1001:PHE:O	1:B:1002:ASP:C	2.52	0.53
1:A:63:MET:CG	1:A:160:TRP:CE2	2.92	0.53
1:A:117:TYR:OH	1:A:174:ASP:CB	2.53	0.53
1:A:143:GLU:CD	1:A:198:HIS:HE2	2.16	0.53
1:A:249:LYS:HB2	1:A:264:LEU:CD1	2.39	0.53
1:A:255:GLU:OE2	1:A:259:LYS:O	2.26	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:383:PRO:HG2	1:A:386:SER:OG	2.08	0.53
1:B:111:LEU:HD22	1:B:226:TRP:CZ3	2.43	0.53
1:B:166:PHE:CD1	1:B:183:VAL:HG22	2.43	0.53
1:B:285:GLU:C	1:B:286:VAL:CG2	2.81	0.53
1:A:26:PHE:CE1	1:A:50:PRO:HD2	2.43	0.53
1:A:63:MET:HE2	1:A:160:TRP:CB	2.39	0.53
1:A:63:MET:CG	1:A:120:VAL:HG12	2.38	0.53
1:A:100:LYS:HA	1:A:103:ASN:HD22	1.74	0.53
1:A:139:PRO:C	1:A:148:ARG:CD	2.77	0.53
1:A:367:ILE:O	1:A:392:HIS:HD2	1.91	0.53
1:A:443:LEU:O	1:A:513:GLN:OE1	2.26	0.53
1:B:34:ASP:H	1:B:448:ASP:CG	2.17	0.53
1:B:54:ARG:HG2	1:B:283:GLU:HG2	1.90	0.53
1:B:63:MET:SD	1:B:64:ASN:C	2.92	0.53
1:B:147:MET:CE	1:B:186:PRO:HB3	2.37	0.53
1:B:250:PHE:CA	1:B:430:PHE:HZ	2.19	0.53
1:B:253:CYS:CB	1:B:307:HIS:CE1	2.91	0.53
1:A:87:VAL:CA	1:A:90:ASN:ND2	2.68	0.53
1:A:224:MET:CG	1:A:228:ASP:OD1	2.56	0.53
1:A:233:ILE:HD12	1:A:298:TYR:HE1	1.72	0.53
1:A:391:ARG:CZ	1:A:397:THR:H	2.21	0.53
1:A:481:VAL:CG2	1:A:555:LEU:HD23	2.38	0.53
1:A:1218:LEU:O	1:A:1221:PRO:O	2.26	0.53
1:B:54:ARG:HG3	1:B:283:GLU:OE2	2.08	0.53
1:B:201:VAL:HG23	1:B:202:ASP:H	1.73	0.53
1:B:242:LEU:HG	1:B:251:LEU:HD21	1.91	0.53
1:B:304:ARG:N	1:B:367:ILE:HG21	2.22	0.53
1:B:364:ILE:O	1:B:365:SER:OG	2.17	0.53
1:B:741:ALA:O	1:B:876:PHE:HA	2.09	0.53
1:B:1218:LEU:CB	1:B:1234:MET:HA	2.39	0.53
1:A:152:ASP:OD2	1:A:156:ASN:HB3	2.09	0.53
1:A:372:PRO:CA	1:A:389:ARG:HG2	2.39	0.53
1:A:383:PRO:HB2	1:A:386:SER:OG	2.08	0.53
1:A:400:HIS:O	1:A:418:ILE:CB	2.57	0.53
1:B:54:ARG:CG	1:B:283:GLU:CD	2.82	0.53
1:B:65:ARG:NH1	1:B:106:GLU:OE1	2.42	0.53
1:B:75:ALA:O	1:B:89:LEU:CD1	2.54	0.53
1:B:243:PHE:HB2	1:B:431:ALA:HB3	1.86	0.53
1:B:264:LEU:CG	1:B:418:ILE:HD11	2.38	0.53
1:B:287:VAL:H	1:B:304:ARG:NE	2.07	0.53
1:B:474:THR:O	1:B:478:GLU:CD	2.52	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:484:VAL:O	1:B:562:HIS:CE1	2.62	0.53
1:B:1428:HIS:CB	1:B:1452:ASN:CB	2.86	0.53
1:A:14:ILE:CG2	1:A:59:LYS:CA	2.78	0.53
1:A:15:CYS:CB	1:A:223:PHE:CB	2.86	0.53
1:A:456:ILE:HA	1:A:459:LYS:HG2	1.91	0.53
1:B:303:PHE:O	1:B:304:ARG:C	2.52	0.53
1:B:743:MET:O	1:B:745:LEU:N	2.38	0.53
1:A:197:SER:C	1:A:198:HIS:ND1	2.67	0.52
1:A:370:LEU:HD23	1:A:390:LEU:HD13	1.89	0.52
1:A:387:TYR:C	1:A:388:VAL:HG22	2.34	0.52
1:B:242:LEU:HB2	1:B:430:PHE:CE2	2.39	0.52
1:B:365:SER:CB	1:B:392:HIS:HE2	2.22	0.52
1:B:400:HIS:HB2	1:B:419:GLY:C	2.33	0.52
1:B:480:LEU:O	1:B:484:VAL:HG12	2.09	0.52
1:A:30:LEU:HD21	1:A:38:VAL:CG1	2.39	0.52
1:A:63:MET:HE2	1:A:160:TRP:HB3	1.91	0.52
1:A:243:PHE:O	1:A:430:PHE:CA	2.57	0.52
1:A:406:ILE:CG1	1:A:417:LYS:HA	2.39	0.52
1:A:450:SER:HA	1:A:453:LEU:HD21	1.90	0.52
1:B:16:SER:C	1:B:46:LEU:HD11	2.34	0.52
1:B:290:ASP:OD2	1:B:293:ARG:O	2.27	0.52
1:B:380:SER:CB	1:B:381:LEU:HD22	2.39	0.52
1:B:1281:PHE:O	1:B:1285:SER:CB	2.57	0.52
1:A:28:SER:HA	1:A:56:CYS:HB2	1.91	0.52
1:A:235:LYS:HD2	1:A:238:ASP:OD2	2.10	0.52
1:A:1057:GLY:C	1:A:1059:THR:H	2.09	0.52
1:B:138:LEU:O	1:B:148:ARG:CB	2.58	0.52
1:B:194:HIS:CE1	1:B:214:CYS:SG	3.02	0.52
1:B:290:ASP:OD2	1:B:293:ARG:CB	2.57	0.52
1:B:448:ASP:CA	1:B:451:LYS:CE	2.87	0.52
1:B:507:GLN:CA	1:B:510:MET:SD	2.92	0.52
1:B:1316:VAL:O	1:B:1323:ILE:CB	2.58	0.52
1:A:13:ASP:O	1:A:14:ILE:HG13	2.08	0.52
1:A:124:LEU:CA	1:A:131:TYR:CD1	2.92	0.52
1:A:235:LYS:HD2	1:A:238:ASP:CG	2.35	0.52
1:A:508:LYS:CE	1:A:512:GLU:HG3	2.39	0.52
1:B:16:SER:N	1:B:220:ILE:CD1	2.73	0.52
1:B:1309:ILE:CB	1:B:1361:GLU:O	2.58	0.52
1:B:1460:ALA:HB1	1:B:1464:THR:HA	1.90	0.52
1:A:33:VAL:CG1	1:A:448:ASP:O	2.58	0.52
1:A:306:LYS:CD	1:A:311:GLY:HA2	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:466:THR:OG1	1:A:469:GLU:CB	2.57	0.52
1:A:485:THR:HG22	1:A:562:HIS:CE1	2.42	0.52
1:A:581:MET:HA	1:A:584:GLN:CB	2.39	0.52
1:B:19:ALA:HB2	1:B:218:TRP:CH2	2.43	0.52
1:B:46:LEU:HD12	1:B:221:VAL:HG13	1.89	0.52
1:B:54:ARG:CG	1:B:283:GLU:OE2	2.57	0.52
1:B:68:ALA:O	1:B:96:ALA:HB2	2.09	0.52
1:B:276:THR:HG23	1:B:277:SER:O	2.09	0.52
1:B:286:VAL:HG11	1:B:288:GLN:CG	2.38	0.52
1:B:440:VAL:HB	1:B:441:ARG:HH12	1.74	0.52
1:B:1400:LEU:CA	1:B:1442:THR:HA	2.39	0.52
1:B:1434:GLU:HA	1:B:1493:PRO:HA	1.92	0.52
1:A:16:SER:OG	1:A:53:PHE:HZ	1.91	0.52
1:A:315:ALA:N	1:A:358:VAL:HG22	2.25	0.52
1:A:507:GLN:OE1	1:A:563:SER:HA	2.10	0.52
1:B:111:LEU:CD1	1:B:112:GLY:H	2.22	0.52
1:B:130:LYS:HG2	1:B:151:LEU:CD2	2.39	0.52
1:B:195:ALA:O	1:B:208:GLU:O	2.28	0.52
1:B:353:TYR:CE2	1:B:395:THR:CG2	2.93	0.52
1:B:380:SER:O	1:B:381:LEU:HB2	2.09	0.52
1:A:26:PHE:HE1	1:A:44:GLY:HA3	1.74	0.52
1:A:58:PHE:HZ	1:A:123:LEU:O	1.93	0.52
1:A:243:PHE:CG	1:A:244:HIS:N	2.74	0.52
1:A:249:LYS:NZ	1:A:416:LEU:O	2.41	0.52
1:A:299:TRP:HA	1:A:299:TRP:CE3	2.45	0.52
1:A:457:ALA:C	1:A:459:LYS:HD2	2.35	0.52
1:A:710:LYS:O	1:A:714:GLU:CB	2.58	0.52
1:B:1177:LYS:O	1:B:1180:LEU:CB	2.57	0.52
1:B:1355:ILE:C	1:B:1357:MET:N	2.64	0.52
1:A:15:CYS:CA	1:A:223:PHE:CB	2.85	0.52
1:A:29:THR:OG1	1:A:125:HIS:CE1	2.62	0.52
1:A:300:ASN:C	1:A:300:ASN:HD22	2.17	0.52
1:A:313:TYR:O	1:A:358:VAL:O	2.27	0.52
1:A:364:ILE:O	1:A:367:ILE:CG1	2.42	0.52
1:A:384:ARG:NE	1:A:385:ASN:HD21	2.08	0.52
1:A:484:VAL:HG13	1:A:562:HIS:CG	2.45	0.52
1:A:1442:THR:CB	1:A:1445:HIS:CB	2.87	0.52
1:B:24:ASN:ND2	1:B:41:PRO:O	2.38	0.52
1:B:110:LEU:HD23	1:B:110:LEU:H	1.75	0.52
1:B:249:LYS:CD	1:B:266:THR:N	2.72	0.52
1:B:440:VAL:O	1:B:441:ARG:C	2.51	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1223:GLU:CB	1:B:1227:ASP:O	2.58	0.52
1:A:242:LEU:H	1:A:250:PHE:HE1	1.48	0.52
1:A:262:VAL:O	1:A:263:PHE:HB3	2.10	0.52
1:A:282:TRP:CE3	1:A:308:LEU:HG	2.45	0.52
1:A:769:SER:O	1:A:773:LEU:CA	2.57	0.52
1:A:1076:VAL:O	1:A:1079:ALA:HB3	2.09	0.52
1:B:9:LEU:CD2	1:B:115:ILE:HB	2.30	0.52
1:B:287:VAL:HA	1:B:290:ASP:C	2.35	0.52
1:B:287:VAL:N	1:B:304:ARG:CZ	2.73	0.52
1:B:480:LEU:CD2	1:B:555:LEU:HD22	2.40	0.52
1:A:33:VAL:HG22	1:A:448:ASP:CB	2.39	0.52
1:A:244:HIS:CG	1:A:430:PHE:CD1	2.97	0.52
1:A:1460:ALA:HB1	1:A:1466:ASP:HA	1.92	0.52
1:A:1482:VAL:O	1:A:1486:VAL:CB	2.58	0.52
1:B:249:LYS:HD2	1:B:416:LEU:HB2	1.90	0.52
1:B:299:TRP:HZ2	1:B:379:ASP:N	2.05	0.52
1:B:417:LYS:O	1:B:418:ILE:HG12	2.10	0.52
1:B:450:SER:HA	1:B:453:LEU:HD11	1.92	0.52
1:B:507:GLN:CB	1:B:563:SER:O	2.58	0.52
1:A:39:VAL:O	1:A:206:CYS:C	2.51	0.51
1:A:142:LEU:HD13	1:A:208:GLU:OE1	2.10	0.51
1:A:166:PHE:CZ	1:A:217:SER:OG	2.63	0.51
1:A:166:PHE:CZ	1:A:217:SER:CB	2.92	0.51
1:A:191:GLN:CB	1:A:192:PRO:CD	2.74	0.51
1:A:235:LYS:HB2	1:A:238:ASP:OD1	2.11	0.51
1:B:20:GLU:CD	1:B:181:LYS:CD	2.83	0.51
1:B:55:ASP:N	1:B:127:LYS:NZ	2.57	0.51
1:B:219:LYS:NZ	1:B:219:LYS:HB2	2.24	0.51
1:B:408:LYS:HA	1:B:408:LYS:NZ	2.24	0.51
1:B:437:PRO:C	1:B:441:ARG:NH2	2.67	0.51
1:B:510:MET:HG3	1:B:516:LEU:HG	1.88	0.51
1:A:7:SER:CB	1:A:178:ILE:HD13	2.39	0.51
1:A:18:TYR:CE1	1:A:25:GLY:C	2.89	0.51
1:A:133:THR:OG1	1:A:152:ASP:HB3	2.10	0.51
1:A:477:LEU:C	1:A:480:LEU:CD1	2.83	0.51
1:A:1057:GLY:C	1:A:1059:THR:N	2.67	0.51
1:A:1341:VAL:C	1:A:1343:VAL:N	2.65	0.51
1:B:132:LEU:HD12	1:B:132:LEU:N	2.12	0.51
1:B:162:TYR:HB2	1:B:185:ASN:HB3	1.92	0.51
1:B:243:PHE:HD1	1:B:250:PHE:CE1	2.27	0.51
1:B:244:HIS:ND1	1:B:264:LEU:CD1	2.72	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:459:LYS:CA	1:B:462:LYS:CD	2.84	0.51
1:A:69:GLN:NE2	1:A:96:ALA:HA	2.26	0.51
1:A:196:SER:HB2	1:A:198:HIS:CE1	2.46	0.51
1:A:212:VAL:HG13	1:A:213:ASN:O	2.10	0.51
1:A:290:ASP:CG	1:A:291:PRO:HD2	2.35	0.51
1:A:304:ARG:NH2	1:A:367:ILE:HD13	2.26	0.51
1:A:455:SER:C	1:A:459:LYS:CD	2.83	0.51
1:A:478:GLU:HA	1:A:555:LEU:HD21	1.93	0.51
1:A:514:ASN:ND2	1:A:518:GLN:CG	2.70	0.51
1:A:1421:ALA:HB2	1:A:1456:ASP:CA	2.35	0.51
1:B:20:GLU:N	1:B:20:GLU:CD	2.68	0.51
1:B:54:ARG:NH1	1:B:55:ASP:OD1	2.28	0.51
1:B:76:ALA:HB3	1:B:89:LEU:HD22	1.90	0.51
1:B:167:TYR:N	1:B:167:TYR:HD1	2.04	0.51
1:B:263:PHE:CE1	1:B:415:MET:HE1	2.46	0.51
1:B:313:TYR:CD2	1:B:361:GLY:CA	2.94	0.51
1:B:397:THR:HB	1:B:422:PRO:HA	1.93	0.51
1:B:744:CYS:HA	1:B:1085:ARG:CB	2.39	0.51
1:A:11:ILE:HG22	1:A:110:LEU:CD2	2.37	0.51
1:A:263:PHE:CD2	1:A:415:MET:CG	2.93	0.51
1:A:303:PHE:CZ	1:A:370:LEU:HD11	2.45	0.51
1:A:435:VAL:HG12	1:A:440:VAL:CG2	2.40	0.51
1:B:76:ALA:HB2	1:B:89:LEU:HD22	1.91	0.51
1:B:184:LEU:HB3	1:B:193:LEU:H	1.75	0.51
1:B:233:ILE:CD1	1:B:234:LEU:O	2.59	0.51
1:B:300:ASN:O	1:B:301:SER:CB	2.54	0.51
1:B:399:VAL:CB	1:B:419:GLY:HA2	2.40	0.51
1:B:443:LEU:HD12	1:B:509:LEU:HD21	1.91	0.51
1:B:504:ARG:HH22	1:B:567:TYR:N	2.08	0.51
1:A:14:ILE:C	1:A:15:CYS:SG	2.93	0.51
1:A:26:PHE:HE1	1:A:45:ASP:N	2.09	0.51
1:A:239:VAL:C	1:A:435:VAL:HG23	2.36	0.51
1:A:454:GLY:O	1:A:459:LYS:CE	2.54	0.51
1:A:964:ASP:O	1:A:967:VAL:N	2.44	0.51
1:A:1225:ALA:O	1:A:1320:GLY:HA3	2.10	0.51
1:B:13:ASP:CA	1:B:226:TRP:HB3	2.32	0.51
1:B:33:VAL:HG21	1:B:449:ALA:CA	2.40	0.51
1:B:285:GLU:O	1:B:286:VAL:HG22	2.10	0.51
1:B:286:VAL:HG13	1:B:303:PHE:CD1	2.43	0.51
1:B:296:ALA:C	1:B:303:PHE:HE2	2.18	0.51
1:B:312:HIS:NE2	1:B:359:PRO:CA	2.74	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:399:VAL:HA	1:B:419:GLY:C	2.35	0.51
1:B:461:GLU:CB	1:B:462:LYS:HZ3	2.24	0.51
1:B:728:ASP:CB	1:B:866:VAL:N	2.73	0.51
1:A:72:PHE:N	1:A:92:LEU:CD2	2.74	0.51
1:A:457:ALA:CB	1:A:522:LEU:CG	2.88	0.51
1:B:181:LYS:HG2	1:B:218:TRP:C	2.36	0.51
1:B:239:VAL:CG2	1:B:308:LEU:HD21	2.41	0.51
1:B:384:ARG:CB	1:B:434:PRO:HD3	2.41	0.51
1:B:445:PHE:O	1:B:449:ALA:HB2	2.10	0.51
1:A:19:ALA:CB	1:A:218:TRP:CZ3	2.94	0.51
1:A:57:LEU:O	1:A:125:HIS:CD2	2.63	0.51
1:A:58:PHE:N	1:A:126:LEU:HG	2.26	0.51
1:A:140:ALA:CA	1:A:148:ARG:HD3	2.40	0.51
1:A:476:LEU:HA	1:A:479:ASP:OD1	2.10	0.51
1:A:1380:LEU:CB	1:A:1384:CYS:CB	2.89	0.51
1:B:16:SER:O	1:B:220:ILE:CD1	2.58	0.51
1:B:33:VAL:HB	1:B:448:ASP:CB	2.41	0.51
1:B:111:LEU:HD13	1:B:112:GLY:N	2.25	0.51
1:B:116:GLN:HE22	1:B:173:GLY:HA2	1.75	0.51
1:B:126:LEU:HD23	1:B:126:LEU:N	2.25	0.51
1:B:221:VAL:CG2	1:B:292:CYS:CB	2.88	0.51
1:B:287:VAL:CB	1:B:304:ARG:NH1	2.72	0.51
1:B:473:VAL:HG13	1:B:474:THR:N	2.26	0.51
1:B:612:ALA:O	1:B:613:ALA:C	2.53	0.51
1:B:666:ILE:CB	1:B:670:LEU:H	2.24	0.51
1:B:1339:GLU:CB	1:B:1385:THR:N	2.74	0.51
1:A:12:GLY:CA	1:A:111:LEU:CD2	2.89	0.51
1:A:243:PHE:CB	1:A:248:GLU:O	2.59	0.51
1:A:680:VAL:CB	1:A:685:ASN:CB	2.89	0.51
1:A:707:ILE:O	1:A:708:ARG:CB	2.58	0.51
1:B:168:LYS:HB2	1:B:168:LYS:NZ	2.26	0.51
1:B:183:VAL:CG1	1:B:192:PRO:HG3	2.41	0.51
1:B:244:HIS:NE2	1:B:247:GLN:C	2.69	0.51
1:B:300:ASN:HD22	1:B:369:GLU:HG3	1.75	0.51
1:B:462:LYS:N	1:B:462:LYS:CD	2.72	0.51
1:A:98:LEU:O	1:A:102:GLN:CD	2.54	0.51
1:A:511:ARG:CA	1:A:516:LEU:CG	2.81	0.51
1:A:879:TYR:CB	1:A:883:ASP:CB	2.88	0.51
1:B:30:LEU:HD13	1:B:36:ARG:HB3	1.93	0.51
1:B:66:TYR:HB2	1:B:69:GLN:HG2	1.92	0.51
1:B:299:TRP:HA	1:B:299:TRP:HE3	1.71	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:513:GLN:OE1	1:B:513:GLN:CA	2.58	0.51
1:B:1224:LYS:O	1:B:1225:ALA:C	2.53	0.51
1:A:240:VAL:HA	1:A:433:VAL:O	2.09	0.51
1:A:552:ILE:HD13	1:A:552:ILE:N	2.25	0.51
1:B:11:ILE:CA	1:B:112:GLY:HA2	2.41	0.51
1:B:185:ASN:HA	1:B:191:GLN:C	2.36	0.51
1:B:252:THR:OG1	1:B:265:ARG:HB2	2.10	0.51
1:B:406:ILE:HB	1:B:415:MET:CB	2.41	0.51
1:B:482:TYR:CD1	1:B:482:TYR:C	2.89	0.51
1:A:65:ARG:HD2	1:A:65:ARG:O	2.11	0.50
1:A:233:ILE:HG21	1:A:381:LEU:HB2	1.94	0.50
1:A:249:LYS:HZ3	1:A:418:ILE:HG21	1.75	0.50
1:A:298:TYR:OH	1:A:381:LEU:HD13	2.12	0.50
1:A:317:GLU:OE1	1:A:356:VAL:HG23	2.11	0.50
1:A:456:ILE:HG21	1:A:473:VAL:HG21	1.94	0.50
1:A:879:TYR:O	1:A:884:LEU:CB	2.59	0.50
1:A:1061:LEU:HA	1:A:1101:GLN:CA	2.41	0.50
1:A:1270:ALA:HB2	1:A:1315:ILE:O	2.11	0.50
1:B:8:PHE:HE2	1:B:114:VAL:HG21	1.76	0.50
1:B:63:MET:SD	1:B:160:TRP:NE1	2.82	0.50
1:B:193:LEU:C	1:B:193:LEU:CD2	2.81	0.50
1:B:475:LYS:HA	1:B:478:GLU:CD	2.36	0.50
1:B:477:LEU:C	1:B:480:LEU:HD22	2.29	0.50
1:A:20:GLU:OE1	1:A:217:SER:O	2.30	0.50
1:A:163:ILE:CD1	1:A:183:VAL:O	2.58	0.50
1:A:166:PHE:HZ	1:A:217:SER:HG	1.53	0.50
1:A:181:LYS:CG	1:A:219:LYS:HB3	2.39	0.50
1:A:198:HIS:ND1	1:A:198:HIS:N	2.59	0.50
1:A:225:LYS:O	1:A:228:ASP:CB	2.57	0.50
1:A:230:LYS:CE	1:A:230:LYS:CA	2.86	0.50
1:A:477:LEU:C	1:A:555:LEU:HD13	2.36	0.50
1:A:622:VAL:HA	1:A:628:PRO:CA	2.39	0.50
1:A:830:LYS:O	1:A:833:PHE:N	2.45	0.50
1:A:1284:CYS:CB	1:A:1295:PHE:CB	2.89	0.50
1:B:192:PRO:O	1:B:211:SER:HA	2.11	0.50
1:B:264:LEU:HB3	1:B:416:LEU:HB2	1.94	0.50
1:B:281:LEU:C	1:B:308:LEU:HD23	2.36	0.50
1:B:308:LEU:CD2	1:B:308:LEU:H	2.21	0.50
1:B:385:ASN:HD21	1:B:433:VAL:HG13	1.76	0.50
1:B:424:LYS:HZ2	1:B:424:LYS:C	2.11	0.50
1:B:503:ASN:CB	1:B:506:ARG:CD	2.89	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:885:LEU:O	1:B:888:THR:CB	2.58	0.50
1:A:222:LEU:H	1:A:222:LEU:CD1	2.25	0.50
1:A:384:ARG:NE	1:A:385:ASN:OD1	2.43	0.50
1:B:11:ILE:CG1	1:B:113:THR:H	2.22	0.50
1:B:26:PHE:CZ	1:B:45:ASP:CA	2.94	0.50
1:B:440:VAL:CG1	1:B:441:ARG:NH1	2.75	0.50
1:A:40:GLN:OE1	1:A:43:ALA:CB	2.58	0.50
1:A:124:LEU:HD11	1:A:129:ASN:O	2.11	0.50
1:A:133:THR:HG23	1:A:156:ASN:CG	2.35	0.50
1:A:263:PHE:CE2	1:A:414:VAL:CB	2.95	0.50
1:A:407:ASP:OD2	1:A:413:PRO:HG2	2.10	0.50
1:A:451:LYS:NZ	1:A:451:LYS:H	2.09	0.50
1:A:481:VAL:HG23	1:A:555:LEU:HD22	1.93	0.50
1:A:507:GLN:NE2	1:A:562:HIS:CD2	2.80	0.50
1:A:522:LEU:CD1	1:A:522:LEU:N	2.73	0.50
1:A:1371:LEU:O	1:A:1374:HIS:C	2.55	0.50
1:B:100:LYS:H	1:B:100:LYS:HE2	1.76	0.50
1:B:457:ALA:HA	1:B:460:LEU:HD22	1.93	0.50
1:B:519:ILE:O	1:B:522:LEU:HD21	2.11	0.50
1:B:1090:ARG:C	1:B:1176:VAL:N	2.70	0.50
1:A:239:VAL:HG23	1:A:282:TRP:C	2.36	0.50
1:A:371:ASP:HB3	1:A:389:ARG:CG	2.37	0.50
1:A:391:ARG:HH21	1:A:392:HIS:HB3	1.76	0.50
1:A:516:LEU:HD12	1:A:520:PHE:HE2	1.72	0.50
1:B:256:HIS:C	1:B:258:LYS:N	2.67	0.50
1:B:286:VAL:HG21	1:B:303:PHE:CE1	2.46	0.50
1:B:369:GLU:N	1:B:393:LEU:HD21	2.26	0.50
1:B:407:ASP:HB2	1:B:413:PRO:HB3	1.94	0.50
1:B:503:ASN:HB3	1:B:506:ARG:CG	2.42	0.50
1:A:103:ASN:C	1:A:107:ASN:HD21	2.20	0.50
1:A:143:GLU:HG2	1:A:210:ASN:ND2	2.26	0.50
1:A:363:ASP:OD1	1:A:366:SER:CB	2.58	0.50
1:A:524:GLN:OE1	1:A:524:GLN:N	2.45	0.50
1:A:1332:ALA:HA	1:A:1376:HIS:O	2.11	0.50
1:B:365:SER:CB	1:B:392:HIS:NE2	2.74	0.50
1:B:1090:ARG:N	1:B:1176:VAL:N	2.59	0.50
1:A:8:PHE:CD2	1:A:177:VAL:HG13	2.46	0.50
1:A:69:GLN:HE21	1:A:99:GLU:CB	2.23	0.50
1:A:263:PHE:CD2	1:A:414:VAL:C	2.90	0.50
1:A:303:PHE:CE2	1:A:370:LEU:HD11	2.47	0.50
1:A:389:ARG:NH1	1:A:428:GLU:O	2.44	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:442:ASP:N	1:A:442:ASP:OD1	2.44	0.50
1:A:445:PHE:HD2	1:A:483:PHE:CD2	2.30	0.50
1:A:1034:GLU:O	1:A:1037:ALA:HB3	2.11	0.50
1:B:117:TYR:CZ	1:B:182:VAL:HG11	2.46	0.50
1:B:148:ARG:CZ	1:B:150:THR:CG2	2.90	0.50
1:B:475:LYS:O	1:B:479:ASP:CG	2.54	0.50
1:A:103:ASN:C	1:A:107:ASN:ND2	2.70	0.50
1:A:126:LEU:O	1:A:129:ASN:ND2	2.45	0.50
1:A:263:PHE:HD2	1:A:415:MET:CG	2.25	0.50
1:A:477:LEU:HB2	1:A:555:LEU:CG	2.42	0.50
1:A:516:LEU:O	1:A:517:LYS:C	2.55	0.50
1:A:747:ARG:CB	1:A:752:ILE:CB	2.89	0.50
1:A:1059:THR:CB	1:A:1098:LYS:N	2.75	0.50
1:A:1417:GLU:CB	1:A:1458:CYS:CB	2.90	0.50
1:B:14:ILE:H	1:B:14:ILE:CD1	2.25	0.50
1:B:461:GLU:CA	1:B:462:LYS:HZ3	2.25	0.50
1:B:510:MET:HG2	1:B:516:LEU:HD21	1.94	0.50
1:B:1356:GLN:HA	1:B:1397:ASN:CB	2.42	0.50
1:A:104:GLU:O	1:A:108:ARG:CZ	2.60	0.50
1:A:281:LEU:CD2	1:A:435:VAL:HG22	2.41	0.50
1:A:306:LYS:CG	1:A:313:TYR:CD1	2.95	0.50
1:A:373:THR:CG2	1:A:428:GLU:O	2.57	0.50
1:A:445:PHE:O	1:A:445:PHE:CD1	2.65	0.50
1:A:449:ALA:O	1:A:450:SER:C	2.55	0.50
1:A:511:ARG:N	1:A:516:LEU:HD21	2.27	0.50
1:A:1416:PRO:CB	1:A:1462:ASN:N	2.75	0.50
1:A:1417:GLU:CB	1:A:1461:CYS:O	2.60	0.50
1:B:11:ILE:HA	1:B:60:LEU:CG	2.41	0.50
1:B:27:ILE:HD11	1:B:209:VAL:HG11	1.94	0.50
1:B:159:SER:O	1:B:161:PHE:HD1	1.95	0.50
1:B:263:PHE:CD2	1:B:415:MET:CA	2.82	0.50
1:B:453:LEU:HD21	1:B:515:ILE:HD12	1.93	0.50
1:A:140:ALA:N	1:A:147:MET:C	2.67	0.49
1:A:140:ALA:HB2	1:A:147:MET:O	2.12	0.49
1:A:235:LYS:H	1:A:235:LYS:HZ3	1.59	0.49
1:A:256:HIS:HB3	1:A:261:HIS:CE1	2.47	0.49
1:A:282:TRP:HZ3	1:A:309:ALA:H	1.58	0.49
1:A:391:ARG:HE	1:A:392:HIS:N	2.10	0.49
1:B:46:LEU:O	1:B:292:CYS:SG	2.70	0.49
1:B:68:ALA:HB1	1:B:95:ALA:C	2.36	0.49
1:B:155:GLY:HA3	1:B:159:SER:OG	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:184:LEU:HD22	1:B:193:LEU:HB2	1.94	0.49
1:B:355:LEU:CD1	1:B:417:LYS:HZ1	2.25	0.49
1:B:625:ASN:CB	1:B:630:PHE:CB	2.89	0.49
1:B:1107:GLN:CB	1:B:1111:ASN:CB	2.90	0.49
1:B:1360:SER:O	1:B:1364:ARG:CB	2.60	0.49
1:A:139:PRO:O	1:A:140:ALA:O	2.30	0.49
1:A:168:LYS:HD3	1:A:168:LYS:C	2.31	0.49
1:A:387:TYR:CA	1:A:431:ALA:CB	2.90	0.49
1:A:518:GLN:O	1:A:519:ILE:C	2.55	0.49
1:A:1000:GLU:CB	1:A:1005:ASN:HA	2.42	0.49
1:B:19:ALA:N	1:B:218:TRP:CZ3	2.79	0.49
1:B:63:MET:HE1	1:B:64:ASN:C	2.37	0.49
1:B:147:MET:HE1	1:B:161:PHE:CZ	2.46	0.49
1:B:240:VAL:CG1	1:B:434:PRO:HA	2.37	0.49
1:B:424:LYS:O	1:B:424:LYS:NZ	2.24	0.49
1:B:598:LEU:O	1:B:599:LEU:C	2.54	0.49
1:B:654:LYS:C	1:B:658:ASN:CB	2.85	0.49
1:A:20:GLU:OE2	1:A:218:TRP:C	2.55	0.49
1:A:179:GLY:CA	1:A:220:ILE:HB	2.41	0.49
1:A:242:LEU:N	1:A:250:PHE:CE1	2.67	0.49
1:B:130:LYS:CG	1:B:151:LEU:HD22	2.42	0.49
1:B:226:TRP:CD1	1:B:226:TRP:N	2.71	0.49
1:A:14:ILE:HG23	1:A:60:LEU:HD23	1.91	0.49
1:A:26:PHE:CE2	1:A:53:PHE:HE2	2.29	0.49
1:A:33:VAL:HG11	1:A:451:LYS:HG2	1.94	0.49
1:A:105:THR:HA	1:A:108:ARG:CZ	2.42	0.49
1:A:138:LEU:CB	1:A:148:ARG:CG	2.89	0.49
1:A:193:LEU:HD23	1:A:211:SER:OG	2.13	0.49
1:A:306:LYS:HA	1:A:313:TYR:HD1	1.76	0.49
1:A:435:VAL:CG1	1:A:440:VAL:HG22	2.42	0.49
1:A:510:MET:SD	1:A:515:ILE:CB	3.00	0.49
1:A:1421:ALA:O	1:A:1426:LEU:N	2.42	0.49
1:B:57:LEU:HD23	1:B:57:LEU:C	2.38	0.49
1:B:740:PHE:CB	1:B:1047:ASN:CB	2.90	0.49
1:B:788:MET:O	1:B:790:VAL:N	2.45	0.49
1:B:1371:LEU:C	1:B:1375:ILE:CB	2.85	0.49
1:A:26:PHE:CD1	1:A:44:GLY:HA3	2.47	0.49
1:A:314:LEU:C	1:A:358:VAL:CG2	2.85	0.49
1:A:460:LEU:C	1:A:460:LEU:CD1	2.85	0.49
1:A:616:ASP:O	1:A:620:SER:CB	2.60	0.49
1:B:63:MET:HE1	1:B:65:ARG:HA	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:181:LYS:HG3	1:B:219:LYS:HB3	1.95	0.49
1:B:453:LEU:H	1:B:453:LEU:CD2	2.23	0.49
1:A:20:GLU:HG3	1:A:181:LYS:CD	2.42	0.49
1:A:395:THR:O	1:A:397:THR:OG1	2.28	0.49
1:A:446:ALA:HB3	1:A:513:GLN:CD	2.37	0.49
1:A:1240:PHE:O	1:A:1241:LEU:C	2.55	0.49
1:B:7:SER:OG	1:B:7:SER:O	2.28	0.49
1:B:18:TYR:CE2	1:B:19:ALA:O	2.66	0.49
1:B:262:VAL:CG2	1:B:357:SER:N	2.76	0.49
1:B:453:LEU:HD23	1:B:453:LEU:N	2.27	0.49
1:A:304:ARG:CZ	1:A:367:ILE:HG23	2.42	0.49
1:A:1270:ALA:CB	1:A:1316:VAL:HA	2.43	0.49
1:B:33:VAL:HB	1:B:448:ASP:OD1	2.12	0.49
1:B:438:ALA:C	1:B:441:ARG:HH21	2.20	0.49
1:B:448:ASP:CA	1:B:451:LYS:NZ	2.75	0.49
1:B:456:ILE:HG23	1:B:469:GLU:CD	2.38	0.49
1:A:67:SER:HA	1:A:70:LYS:CB	2.42	0.49
1:A:372:PRO:HA	1:A:389:ARG:HG2	1.93	0.49
1:B:10:HIS:HA	1:B:113:THR:O	2.12	0.49
1:B:122:GLN:NE2	1:B:131:TYR:CZ	2.81	0.49
1:B:130:LYS:CG	1:B:151:LEU:C	2.86	0.49
1:B:251:LEU:CD1	1:B:282:TRP:CH2	2.84	0.49
1:B:366:SER:O	1:B:367:ILE:HG23	2.12	0.49
1:B:477:LEU:HB2	1:B:555:LEU:CD2	2.42	0.49
1:B:510:MET:HG2	1:B:516:LEU:HD11	1.94	0.49
1:B:618:PHE:O	1:B:622:VAL:CB	2.61	0.49
1:A:72:PHE:CA	1:A:92:LEU:HD22	2.43	0.49
1:A:265:ARG:HA	1:A:416:LEU:HD11	1.90	0.49
1:A:265:ARG:CA	1:A:416:LEU:HD12	2.42	0.49
1:A:283:GLU:CD	1:A:306:LYS:HB3	2.37	0.49
1:A:385:ASN:OD1	1:A:434:PRO:CD	2.59	0.49
1:A:407:ASP:OD2	1:A:415:MET:O	2.30	0.49
1:B:58:PHE:HD2	1:B:123:LEU:HD12	1.75	0.49
1:B:184:LEU:HD22	1:B:193:LEU:HB3	1.95	0.49
1:B:249:LYS:O	1:B:250:PHE:CB	2.61	0.49
1:B:460:LEU:HD13	1:B:460:LEU:H	1.76	0.49
1:B:484:VAL:HG13	1:B:484:VAL:O	2.13	0.49
1:B:1029:ASP:O	1:B:1033:ILE:N	2.35	0.49
1:B:1198:ARG:HA	1:B:1238:HIS:HA	1.93	0.49
1:A:169:LEU:HD23	1:A:170:ARG:NH2	2.26	0.49
1:A:263:PHE:HB2	1:A:406:ILE:CD1	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:390:LEU:HD12	1:A:391:ARG:N	2.27	0.49
1:B:137:ARG:C	1:B:147:MET:HA	2.38	0.49
1:B:219:LYS:O	1:B:219:LYS:HE3	2.13	0.49
1:B:264:LEU:CD2	1:B:430:PHE:CE1	2.94	0.49
1:B:1130:GLU:O	1:B:1131:LEU:C	2.55	0.49
1:A:9:LEU:HD21	1:A:220:ILE:CG2	2.42	0.48
1:A:196:SER:OG	1:A:198:HIS:CE1	2.66	0.48
1:A:299:TRP:HA	1:A:299:TRP:HE3	1.78	0.48
1:A:299:TRP:CZ3	1:A:370:LEU:HD12	2.48	0.48
1:A:964:ASP:C	1:A:967:VAL:H	2.21	0.48
1:B:131:TYR:CB	1:B:153:GLU:O	2.58	0.48
1:B:398:TRP:O	1:B:420:THR:HB	2.13	0.48
1:B:443:LEU:CD1	1:B:509:LEU:HD13	2.42	0.48
1:A:17:LEU:CD1	1:A:18:TYR:N	2.69	0.48
1:A:39:VAL:O	1:A:206:CYS:HA	2.13	0.48
1:A:160:TRP:O	1:A:187:VAL:N	2.31	0.48
1:A:166:PHE:HZ	1:A:217:SER:OG	1.96	0.48
1:A:181:LYS:CD	1:A:219:LYS:CB	2.91	0.48
1:A:391:ARG:NH2	1:A:397:THR:H	2.11	0.48
1:A:450:SER:HA	1:A:453:LEU:HD23	1.91	0.48
1:A:475:LYS:O	1:A:479:ASP:OD1	2.31	0.48
1:B:19:ALA:HB3	1:B:25:GLY:H	1.66	0.48
1:B:38:VAL:CB	1:B:207:ASN:H	2.27	0.48
1:B:71:GLN:HB2	1:B:92:LEU:HD22	1.94	0.48
1:B:102:GLN:HB3	1:B:106:GLU:CD	2.38	0.48
1:B:179:GLY:HA2	1:B:220:ILE:HG22	1.94	0.48
1:B:181:LYS:CB	1:B:218:TRP:N	2.70	0.48
1:B:249:LYS:O	1:B:265:ARG:O	2.31	0.48
1:B:477:LEU:CG	1:B:555:LEU:HD13	2.42	0.48
1:B:748:GLN:O	1:B:749:TYR:CB	2.62	0.48
1:B:1293:GLN:HA	1:B:1345:TYR:O	2.13	0.48
1:A:8:PHE:HD2	1:A:177:VAL:HG22	1.77	0.48
1:A:11:ILE:HG13	1:A:113:THR:O	2.14	0.48
1:A:263:PHE:HB2	1:A:415:MET:CG	2.37	0.48
1:A:353:TYR:CG	1:A:397:THR:CG2	2.96	0.48
1:A:1214:VAL:O	1:A:1218:LEU:CB	2.61	0.48
1:B:46:LEU:HB3	1:B:221:VAL:HG11	1.95	0.48
1:B:400:HIS:HB2	1:B:420:THR:C	2.36	0.48
1:B:482:TYR:HD1	1:B:482:TYR:C	2.21	0.48
1:B:612:ALA:O	1:B:615:ILE:N	2.46	0.48
1:B:985:ARG:HA	1:B:989:ARG:CB	2.43	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1274:GLN:CB	1:B:1330:VAL:HA	2.43	0.48
1:A:15:CYS:HB3	1:A:223:PHE:CB	2.29	0.48
1:A:33:VAL:HG22	1:A:448:ASP:HB3	1.95	0.48
1:A:138:LEU:HB3	1:A:148:ARG:HG3	1.96	0.48
1:A:156:ASN:HD21	1:A:159:SER:N	2.08	0.48
1:A:244:HIS:HB2	1:A:430:PHE:CE1	2.49	0.48
1:A:249:LYS:C	1:A:264:LEU:CD2	2.84	0.48
1:A:301:SER:O	1:A:302:LEU:HB2	2.13	0.48
1:A:303:PHE:CE2	1:A:370:LEU:HD21	2.49	0.48
1:A:391:ARG:HD2	1:A:398:TRP:NE1	2.29	0.48
1:A:392:HIS:NE2	1:A:394:CYS:HB2	2.28	0.48
1:A:1073:PRO:O	1:A:1074:PRO:CB	2.59	0.48
1:A:1381:LEU:O	1:A:1382:ALA:C	2.55	0.48
1:B:49:PRO:HG2	1:B:53:PHE:CE1	2.48	0.48
1:B:251:LEU:HD21	1:B:282:TRP:CH2	2.49	0.48
1:B:312:HIS:NE2	1:B:359:PRO:HB3	2.27	0.48
1:B:380:SER:HB3	1:B:381:LEU:HD22	1.94	0.48
1:B:598:LEU:O	1:B:602:ASN:CA	2.60	0.48
1:A:134:VAL:CG2	1:A:186:PRO:HG2	2.42	0.48
1:A:163:ILE:HB	1:A:184:LEU:HD12	1.96	0.48
1:A:249:LYS:HE2	1:A:264:LEU:HD22	1.96	0.48
1:A:739:LEU:O	1:A:740:PHE:C	2.56	0.48
1:A:1071:ASP:O	1:A:1073:PRO:N	2.46	0.48
1:A:1203:ARG:O	1:A:1205:LEU:O	2.31	0.48
1:B:32:LEU:HD21	1:B:444:ASP:CB	2.43	0.48
1:B:57:LEU:HD23	1:B:58:PHE:N	2.27	0.48
1:B:75:ALA:C	1:B:89:LEU:HD13	2.38	0.48
1:B:128:SER:HB3	1:B:130:LYS:HB3	1.95	0.48
1:B:319:ASP:CG	1:B:354:SER:OG	2.56	0.48
1:B:388:VAL:O	1:B:428:GLU:O	2.32	0.48
1:B:504:ARG:HH12	1:B:566:ASP:C	2.22	0.48
1:B:1130:GLU:CA	1:B:1217:LEU:CB	2.92	0.48
1:A:239:VAL:CG2	1:A:308:LEU:HD23	2.33	0.48
1:A:249:LYS:CB	1:A:430:PHE:HZ	2.09	0.48
1:A:372:PRO:CA	1:A:389:ARG:CD	2.87	0.48
1:A:481:VAL:CG1	1:A:558:ARG:CB	2.91	0.48
1:A:502:PRO:CD	1:A:507:GLN:NE2	2.72	0.48
1:A:514:ASN:O	1:A:518:GLN:HB2	2.14	0.48
1:B:66:TYR:HD1	1:B:99:GLU:HB3	1.78	0.48
1:B:130:LYS:HE3	1:B:151:LEU:CD1	2.43	0.48
1:B:160:TRP:O	1:B:161:PHE:CD1	2.66	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:303:PHE:O	1:B:367:ILE:HG22	2.13	0.48
1:B:462:LYS:H	1:B:462:LYS:NZ	2.10	0.48
1:A:304:ARG:CZ	1:A:367:ILE:CD1	2.90	0.48
1:A:447:ASN:O	1:A:451:LYS:HD2	2.13	0.48
1:A:1050:LEU:O	1:A:1051:ASP:C	2.57	0.48
1:B:11:ILE:CB	1:B:112:GLY:HA2	2.44	0.48
1:B:16:SER:OG	1:B:292:CYS:CB	2.57	0.48
1:B:18:TYR:HB2	1:B:46:LEU:CB	2.40	0.48
1:B:26:PHE:CE1	1:B:45:ASP:C	2.92	0.48
1:B:313:TYR:CE1	1:B:361:GLY:N	2.82	0.48
1:B:415:MET:O	1:B:416:LEU:C	2.57	0.48
1:B:522:LEU:CD2	1:B:523:LEU:HD13	2.43	0.48
1:B:1400:LEU:HA	1:B:1442:THR:CB	2.44	0.48
1:A:15:CYS:CB	1:A:222:LEU:C	2.85	0.48
1:A:26:PHE:HD2	1:A:53:PHE:HE2	1.62	0.48
1:A:387:TYR:CA	1:A:431:ALA:HB2	2.43	0.48
1:A:436:SER:OG	1:A:439:GLU:HB2	2.12	0.48
1:A:466:THR:OG1	1:A:469:GLU:HB2	2.12	0.48
1:B:13:ASP:OD2	1:B:60:LEU:HD11	2.13	0.48
1:B:283:GLU:HB3	1:B:306:LYS:HB3	1.96	0.48
1:A:122:GLN:CG	1:A:160:TRP:CD1	2.95	0.48
1:A:140:ALA:C	1:A:141:LEU:HD23	2.38	0.48
1:A:241:ARG:CZ	1:A:276:THR:HB	2.43	0.48
1:A:263:PHE:HB3	1:A:406:ILE:HD13	1.94	0.48
1:A:393:LEU:HA	1:A:393:LEU:HD12	1.18	0.48
1:A:448:ASP:O	1:A:451:LYS:HG2	2.14	0.48
1:B:100:LYS:O	1:B:103:ASN:CG	2.56	0.48
1:B:142:LEU:HD12	1:B:200:LEU:HA	1.96	0.48
1:B:241:ARG:NH2	1:B:433:VAL:N	2.62	0.48
1:B:244:HIS:CE1	1:B:247:GLN:HB2	2.48	0.48
1:B:263:PHE:CZ	1:B:265:ARG:CG	2.96	0.48
1:B:385:ASN:H	1:B:432:ILE:CB	2.27	0.48
1:B:457:ALA:O	1:B:460:LEU:HD22	2.14	0.48
1:B:1355:ILE:O	1:B:1356:GLN:C	2.55	0.48
1:A:35:ASP:OD1	1:A:35:ASP:N	2.45	0.48
1:A:1215:LEU:CB	1:A:1259:ILE:CB	2.92	0.48
1:B:58:PHE:HE1	1:B:125:HIS:CD2	2.32	0.48
1:B:130:LYS:CD	1:B:153:GLU:OE2	2.62	0.48
1:B:224:MET:SD	1:B:294:GLY:HA2	2.54	0.48
1:B:263:PHE:CD1	1:B:415:MET:CE	2.95	0.48
1:B:353:TYR:CD1	1:B:422:PRO:CB	2.74	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:384:ARG:HH11	1:B:434:PRO:CD	2.25	0.48
1:B:400:HIS:CA	1:B:420:THR:CA	2.91	0.48
1:B:461:GLU:N	1:B:462:LYS:HE2	2.29	0.48
1:B:484:VAL:HG23	1:B:506:ARG:HD2	1.96	0.48
1:A:32:LEU:HD13	1:A:445:PHE:CA	2.34	0.47
1:A:117:TYR:CE2	1:A:174:ASP:N	2.78	0.47
1:A:139:PRO:O	1:A:148:ARG:CD	2.63	0.47
1:A:389:ARG:NH2	1:A:430:PHE:N	2.61	0.47
1:A:418:ILE:HD12	1:A:418:ILE:HG21	1.48	0.47
1:A:442:ASP:HA	1:A:483:PHE:HZ	1.79	0.47
1:B:18:TYR:HB2	1:B:46:LEU:CD2	2.44	0.47
1:B:66:TYR:HB2	1:B:69:GLN:CD	2.38	0.47
1:B:185:ASN:HD21	1:B:192:PRO:HD3	1.79	0.47
1:B:298:TYR:CB	1:B:380:SER:HB3	2.38	0.47
1:B:441:ARG:H	1:B:441:ARG:HE	1.55	0.47
1:A:14:ILE:HB	1:A:58:PHE:C	2.38	0.47
1:A:139:PRO:HD2	1:A:148:ARG:HE	1.80	0.47
1:A:201:VAL:O	1:A:204:PRO:HD3	2.14	0.47
1:A:243:PHE:O	1:A:430:PHE:CB	2.63	0.47
1:A:254:ASP:HB2	1:A:263:PHE:CE1	2.49	0.47
1:A:303:PHE:HB3	1:A:305:PHE:CZ	2.49	0.47
1:B:20:GLU:CD	1:B:20:GLU:C	2.82	0.47
1:B:53:PHE:HE2	1:B:292:CYS:HA	1.79	0.47
1:B:102:GLN:CD	1:B:106:GLU:HB3	2.39	0.47
1:B:304:ARG:HE	1:B:304:ARG:H	1.60	0.47
1:B:450:SER:N	1:B:515:ILE:CD1	2.76	0.47
1:B:996:ILE:HA	1:B:1008:SER:CB	2.43	0.47
1:A:8:PHE:HA	1:A:177:VAL:HA	1.96	0.47
1:A:33:VAL:HG22	1:A:448:ASP:CG	2.40	0.47
1:A:181:LYS:HA	1:A:219:LYS:HA	1.96	0.47
1:A:264:LEU:CD2	1:A:264:LEU:C	2.88	0.47
1:A:316:ALA:HA	1:A:356:VAL:N	2.17	0.47
1:A:445:PHE:O	1:A:446:ALA:C	2.55	0.47
1:B:32:LEU:HD11	1:B:444:ASP:C	2.39	0.47
1:B:138:LEU:O	1:B:148:ARG:HB3	2.15	0.47
1:B:257:ARG:CZ	1:B:408:LYS:HG3	2.42	0.47
1:B:290:ASP:OD2	1:B:290:ASP:O	2.31	0.47
1:B:316:ALA:CB	1:B:355:LEU:HG	2.45	0.47
1:B:355:LEU:HB2	1:B:417:LYS:NZ	2.28	0.47
1:B:437:PRO:O	1:B:441:ARG:NH2	2.47	0.47
1:B:522:LEU:HD23	1:B:523:LEU:HD13	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:384:ARG:HE	1:A:385:ASN:CG	2.23	0.47
1:A:523:LEU:HD23	1:A:524:GLN:OE1	2.13	0.47
1:B:47:ASN:OD1	1:B:47:ASN:N	2.47	0.47
1:B:134:VAL:HG11	1:B:147:MET:SD	2.54	0.47
1:B:249:LYS:CG	1:B:266:THR:N	2.78	0.47
1:B:263:PHE:CE1	1:B:415:MET:CE	2.98	0.47
1:B:522:LEU:HD22	1:B:522:LEU:H	1.79	0.47
1:B:666:ILE:N	1:B:670:LEU:CB	2.78	0.47
1:B:847:VAL:CB	1:B:897:CYS:CB	2.92	0.47
1:A:317:GLU:H	1:A:356:VAL:HG22	1.78	0.47
1:A:644:SER:O	1:A:645:ILE:C	2.58	0.47
1:B:37:CYS:C	1:B:38:VAL:CG1	2.88	0.47
1:B:399:VAL:O	1:B:428:GLU:N	2.47	0.47
1:B:507:GLN:OE1	1:B:507:GLN:CA	2.61	0.47
1:B:666:ILE:N	1:B:667:GLU:CA	2.77	0.47
1:A:69:GLN:O	1:A:70:LYS:C	2.57	0.47
1:A:465:ILE:O	1:A:469:GLU:OE1	2.32	0.47
1:B:191:GLN:OE1	1:B:213:ASN:N	2.47	0.47
1:B:212:VAL:HG22	1:B:213:ASN:N	2.22	0.47
1:B:283:GLU:HA	1:B:283:GLU:OE1	2.15	0.47
1:B:440:VAL:CB	1:B:441:ARG:NH1	2.78	0.47
1:B:473:VAL:HG22	1:B:477:LEU:HD11	1.96	0.47
1:B:510:MET:O	1:B:515:ILE:HG22	2.15	0.47
1:B:519:ILE:HA	1:B:522:LEU:HD22	1.96	0.47
1:A:12:GLY:CA	1:A:111:LEU:HD21	2.45	0.47
1:A:108:ARG:HH21	1:A:108:ARG:N	2.13	0.47
1:A:222:LEU:CD1	1:A:222:LEU:N	2.78	0.47
1:A:249:LYS:HB3	1:A:264:LEU:HD22	1.97	0.47
1:A:251:LEU:HA	1:A:263:PHE:O	2.14	0.47
1:A:298:TYR:CD1	1:A:381:LEU:HD22	2.49	0.47
1:A:485:THR:HA	1:A:562:HIS:CE1	2.49	0.47
1:A:621:LEU:O	1:A:625:ASN:CB	2.62	0.47
1:A:761:VAL:O	1:A:762:ASP:CB	2.62	0.47
1:A:1419:LYS:O	1:A:1423:ILE:N	2.30	0.47
1:A:1434:GLU:O	1:A:1435:VAL:CB	2.63	0.47
1:B:11:ILE:N	1:B:112:GLY:CA	2.77	0.47
1:B:13:ASP:OD1	1:B:60:LEU:CD2	2.63	0.47
1:B:15:CYS:C	1:B:220:ILE:HD11	2.39	0.47
1:B:65:ARG:O	1:B:99:GLU:OE2	2.31	0.47
1:B:139:PRO:HB3	1:B:146:ALA:CA	2.44	0.47
1:B:166:PHE:CE1	1:B:216:THR:C	2.92	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:249:LYS:HG3	1:B:265:ARG:H	1.74	0.47
1:B:263:PHE:CE2	1:B:265:ARG:CG	2.98	0.47
1:B:303:PHE:CA	1:B:367:ILE:HG21	2.43	0.47
1:B:407:ASP:OD2	1:B:414:VAL:HG22	2.14	0.47
1:B:408:LYS:HA	1:B:408:LYS:CE	2.45	0.47
1:B:477:LEU:HB2	1:B:555:LEU:HD22	1.96	0.47
1:B:1315:ILE:O	1:B:1318:ALA:N	2.48	0.47
1:A:18:TYR:CE1	1:A:24:ASN:OD1	2.68	0.47
1:A:58:PHE:CZ	1:A:124:LEU:C	2.93	0.47
1:A:391:ARG:HG3	1:A:398:TRP:CB	2.40	0.47
1:A:1134:TYR:H	1:A:1180:LEU:CB	2.28	0.47
1:B:196:SER:HB2	1:B:209:VAL:CA	2.44	0.47
1:B:303:PHE:C	1:B:367:ILE:HG22	2.38	0.47
1:B:391:ARG:HB2	1:B:398:TRP:CD2	2.50	0.47
1:B:478:GLU:O	1:B:481:VAL:CG2	2.62	0.47
1:B:1000:GLU:H	1:B:1006:SER:H	1.63	0.47
1:B:1370:PRO:N	1:B:1413:ASP:CB	2.78	0.47
1:A:7:SER:HB2	1:A:178:ILE:CD1	2.44	0.47
1:A:138:LEU:CB	1:A:148:ARG:HG2	2.45	0.47
1:A:222:LEU:O	1:A:222:LEU:HD13	2.14	0.47
1:A:589:VAL:CB	1:A:629:ARG:CB	2.93	0.47
1:A:1375:ILE:HA	1:A:1379:GLU:CB	2.45	0.47
1:B:8:PHE:CE2	1:B:114:VAL:HG21	2.50	0.47
1:B:20:GLU:CD	1:B:181:LYS:NZ	2.69	0.47
1:B:33:VAL:CG1	1:B:476:LEU:CD1	2.92	0.47
1:B:150:THR:N	1:B:152:ASP:OD1	2.47	0.47
1:B:371:ASP:HB3	1:B:389:ARG:O	2.14	0.47
1:B:401:SER:CA	1:B:418:ILE:HG22	2.41	0.47
1:B:456:ILE:O	1:B:459:LYS:N	2.43	0.47
1:B:460:LEU:N	1:B:460:LEU:HD13	2.30	0.47
1:B:879:TYR:O	1:B:884:LEU:CB	2.63	0.47
1:B:1355:ILE:O	1:B:1357:MET:N	2.48	0.47
1:A:32:LEU:HD11	1:A:444:ASP:O	2.14	0.47
1:A:33:VAL:HG11	1:A:452:VAL:HG22	1.86	0.47
1:A:108:ARG:HE	1:A:108:ARG:N	2.10	0.47
1:A:305:PHE:HZ	1:A:368:PHE:HB2	1.75	0.47
1:A:372:PRO:HA	1:A:389:ARG:CG	2.45	0.47
1:A:379:ASP:N	1:A:379:ASP:OD1	2.46	0.47
1:A:447:ASN:C	1:A:450:SER:OG	2.44	0.47
1:A:1126:VAL:O	1:A:1130:GLU:CA	2.61	0.47
1:A:1331:MET:O	1:A:1335:VAL:CB	2.63	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:19:ALA:CA	1:B:218:TRP:CH2	2.95	0.47
1:B:63:MET:CE	1:B:64:ASN:O	2.63	0.47
1:B:133:THR:CG2	1:B:152:ASP:CB	2.87	0.47
1:B:303:PHE:N	1:B:367:ILE:CG2	2.78	0.47
1:B:439:GLU:C	1:B:442:ASP:OD1	2.58	0.47
1:B:1223:GLU:CB	1:B:1228:THR:HA	2.45	0.47
1:B:1260:ASN:O	1:B:1261:LEU:CB	2.63	0.47
1:A:12:GLY:HA3	1:A:111:LEU:HG	1.96	0.46
1:A:57:LEU:O	1:A:126:LEU:CB	2.63	0.46
1:A:227:SER:C	1:A:230:LYS:HG2	2.40	0.46
1:A:485:THR:CG2	1:A:562:HIS:CE1	2.95	0.46
1:A:1227:ASP:O	1:A:1228:THR:C	2.58	0.46
1:A:1364:ARG:O	1:A:1365:MET:CB	2.63	0.46
1:B:49:PRO:HG2	1:B:53:PHE:CE2	2.50	0.46
1:B:223:PHE:CD2	1:B:294:GLY:O	2.67	0.46
1:B:416:LEU:CD2	1:B:416:LEU:N	2.75	0.46
1:B:457:ALA:C	1:B:459:LYS:N	2.73	0.46
1:B:507:GLN:CG	1:B:563:SER:O	2.59	0.46
1:B:725:GLU:HA	1:B:862:LEU:CA	2.44	0.46
1:A:11:ILE:C	1:A:13:ASP:OD2	2.57	0.46
1:A:178:ILE:HG22	1:A:178:ILE:O	2.15	0.46
1:A:235:LYS:CD	1:A:238:ASP:CG	2.88	0.46
1:A:256:HIS:O	1:A:261:HIS:NE2	2.33	0.46
1:A:257:ARG:O	1:A:258:LYS:HB2	2.15	0.46
1:A:466:THR:OG1	1:A:469:GLU:HB3	2.15	0.46
1:A:1287:ILE:CB	1:A:1343:VAL:CB	2.93	0.46
1:B:299:TRP:CZ2	1:B:379:ASP:C	2.93	0.46
1:A:105:THR:O	1:A:109:LYS:NZ	2.48	0.46
1:A:264:LEU:CD2	1:A:416:LEU:HD12	2.44	0.46
1:A:315:ALA:HB1	1:A:358:VAL:HG13	1.91	0.46
1:A:368:PHE:N	1:A:393:LEU:HD22	2.30	0.46
1:A:507:GLN:NE2	1:A:563:SER:HA	2.30	0.46
1:B:29:THR:HG22	1:B:125:HIS:HE1	1.78	0.46
1:B:251:LEU:HA	1:B:264:LEU:HA	1.96	0.46
1:B:287:VAL:CG1	1:B:288:GLN:N	2.78	0.46
1:B:290:ASP:CG	1:B:293:ARG:CB	2.89	0.46
1:B:390:LEU:HD11	1:B:399:VAL:CA	2.44	0.46
1:B:519:ILE:HA	1:B:522:LEU:HD21	1.96	0.46
1:B:1232:GLU:HA	1:B:1236:LEU:CB	2.46	0.46
1:A:37:CYS:SG	1:A:209:VAL:CB	2.97	0.46
1:A:240:VAL:C	1:A:433:VAL:O	2.59	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:448:ASP:OD1	1:A:451:LYS:HD3	2.15	0.46
1:A:457:ALA:CB	1:A:522:LEU:CD1	2.93	0.46
1:A:508:LYS:HZ2	1:A:512:GLU:CG	2.27	0.46
1:A:1317:LYS:HA	1:A:1322:PHE:C	2.40	0.46
1:B:242:LEU:HD11	1:B:251:LEU:HD22	1.97	0.46
1:B:313:TYR:CE2	1:B:361:GLY:CA	2.82	0.46
1:B:366:SER:C	1:B:367:ILE:HG23	2.41	0.46
1:B:407:ASP:H	1:B:415:MET:HG2	1.77	0.46
1:B:470:ARG:O	1:B:473:VAL:HG13	2.15	0.46
1:A:125:HIS:CB	1:A:130:LYS:O	2.64	0.46
1:A:453:LEU:CD1	1:A:476:LEU:HD21	2.45	0.46
1:A:503:ASN:ND2	1:A:505:GLU:OE2	2.48	0.46
1:A:512:GLU:N	1:A:512:GLU:CD	2.70	0.46
1:B:16:SER:HG	1:B:53:PHE:HZ	1.62	0.46
1:B:30:LEU:CD1	1:B:36:ARG:HB3	2.46	0.46
1:B:264:LEU:C	1:B:416:LEU:HB2	2.40	0.46
1:B:665:LEU:C	1:B:667:GLU:HA	2.40	0.46
1:A:17:LEU:HD11	1:A:218:TRP:CD2	2.50	0.46
1:A:124:LEU:HD11	1:A:130:LYS:N	2.30	0.46
1:A:1132:TRP:O	1:A:1133:VAL:C	2.59	0.46
1:B:33:VAL:CB	1:B:448:ASP:C	2.84	0.46
1:B:102:GLN:O	1:B:105:THR:N	2.48	0.46
1:B:135:ASN:O	1:B:137:ARG:N	2.48	0.46
1:B:143:GLU:OE1	1:B:196:SER:OG	2.34	0.46
1:B:180:ASP:O	1:B:218:TRP:O	2.33	0.46
1:B:235:LYS:CE	1:B:238:ASP:OD2	2.63	0.46
1:B:290:ASP:CG	1:B:293:ARG:N	2.69	0.46
1:B:1031:GLU:O	1:B:1035:GLU:N	2.37	0.46
1:B:1370:PRO:CA	1:B:1413:ASP:CB	2.94	0.46
1:A:18:TYR:CE2	1:A:26:PHE:CD2	3.04	0.46
1:A:57:LEU:O	1:A:125:HIS:HD2	1.97	0.46
1:B:111:LEU:N	1:B:111:LEU:CD1	2.74	0.46
1:B:242:LEU:HD21	1:B:282:TRP:CZ3	2.51	0.46
1:B:244:HIS:NE2	1:B:247:GLN:HB3	2.30	0.46
1:B:319:ASP:CG	1:B:354:SER:HG	2.24	0.46
1:B:416:LEU:H	1:B:416:LEU:HD22	1.79	0.46
1:B:462:LYS:HD2	1:B:462:LYS:C	2.33	0.46
1:B:666:ILE:N	1:B:667:GLU:HA	2.31	0.46
1:B:725:GLU:CB	1:B:861:LYS:C	2.88	0.46
1:B:1378:VAL:O	1:B:1382:ALA:CB	2.64	0.46
1:A:164:GLN:N	1:A:164:GLN:CD	2.74	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:447:ASN:CA	1:A:450:SER:OG	2.64	0.46
1:A:451:LYS:CD	1:A:451:LYS:H	2.29	0.46
1:A:638:CYS:O	1:A:639:VAL:CB	2.63	0.46
1:A:1300:GLU:O	1:A:1302:HIS:N	2.47	0.46
1:B:239:VAL:O	1:B:239:VAL:CG1	2.64	0.46
1:B:314:LEU:HG	1:B:315:ALA:N	2.31	0.46
1:B:440:VAL:CB	1:B:441:ARG:CZ	2.90	0.46
1:B:477:LEU:HB3	1:B:555:LEU:HB3	1.98	0.46
1:B:511:ARG:HD2	1:B:511:ARG:C	2.41	0.46
1:B:612:ALA:C	1:B:614:GLU:N	2.72	0.46
1:B:1193:SER:O	1:B:1194:VAL:C	2.59	0.46
1:A:9:LEU:HD22	1:A:220:ILE:HG21	1.97	0.46
1:A:32:LEU:CG	1:A:448:ASP:HB2	2.46	0.46
1:A:129:ASN:HB2	1:A:441:ARG:NH1	2.27	0.46
1:A:255:GLU:OE2	1:A:259:LYS:C	2.59	0.46
1:A:302:LEU:O	1:A:303:PHE:CD2	2.69	0.46
1:A:654:LYS:O	1:A:658:ASN:N	2.37	0.46
1:B:13:ASP:O	1:B:59:LYS:CB	2.54	0.46
1:B:37:CYS:O	1:B:38:VAL:CG1	2.64	0.46
1:B:130:LYS:HG2	1:B:151:LEU:C	2.41	0.46
1:B:134:VAL:CG1	1:B:147:MET:HB3	2.38	0.46
1:B:139:PRO:O	1:B:140:ALA:HB2	2.16	0.46
1:B:224:MET:CE	1:B:288:GLN:NE2	2.78	0.46
1:B:290:ASP:OD2	1:B:290:ASP:C	2.58	0.46
1:B:385:ASN:HD21	1:B:433:VAL:CG1	2.29	0.46
1:B:389:ARG:HH11	1:B:428:GLU:HG2	1.80	0.46
1:B:453:LEU:HD21	1:B:515:ILE:CD1	2.46	0.46
1:B:611:THR:O	1:B:614:GLU:CB	2.64	0.46
1:A:19:ALA:N	1:A:218:TRP:HZ3	2.13	0.46
1:A:98:LEU:O	1:A:102:GLN:CB	2.63	0.46
1:A:138:LEU:HB3	1:A:148:ARG:CZ	2.46	0.46
1:A:181:LYS:HZ3	1:A:219:LYS:N	2.14	0.46
1:A:283:GLU:OE1	1:A:306:LYS:CB	2.64	0.46
1:A:389:ARG:HB3	1:A:389:ARG:HH11	1.80	0.46
1:A:411:GLU:O	1:A:412:LYS:C	2.58	0.46
1:A:445:PHE:HD2	1:A:483:PHE:CG	2.34	0.46
1:A:1374:HIS:O	1:A:1378:VAL:CA	2.64	0.46
1:B:11:ILE:HG23	1:B:61:CYS:HA	1.96	0.46
1:B:20:GLU:HG3	1:B:181:LYS:HZ3	1.81	0.46
1:B:63:MET:CG	1:B:160:TRP:HD1	2.29	0.46
1:B:355:LEU:H	1:B:417:LYS:NZ	2.14	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:363:ASP:H	1:B:366:SER:HB2	1.80	0.46
1:A:20:GLU:HG2	1:A:217:SER:CB	2.44	0.45
1:A:46:LEU:O	1:A:291:PRO:HB3	2.16	0.45
1:A:100:LYS:O	1:A:103:ASN:HB2	2.17	0.45
1:A:180:ASP:O	1:A:220:ILE:HD13	2.16	0.45
1:A:254:ASP:HB2	1:A:263:PHE:HZ	1.76	0.45
1:A:452:VAL:CG2	1:A:476:LEU:CD2	2.90	0.45
1:A:484:VAL:HG23	1:A:563:SER:OG	2.16	0.45
1:A:744:CYS:O	1:A:745:LEU:CB	2.64	0.45
1:A:767:CYS:O	1:A:771:GLU:CB	2.64	0.45
1:B:13:ASP:OD1	1:B:60:LEU:HD21	2.16	0.45
1:B:33:VAL:HG21	1:B:449:ALA:N	2.30	0.45
1:B:256:HIS:NE2	1:B:415:MET:HE2	2.31	0.45
1:B:399:VAL:CG1	1:B:419:GLY:HA2	2.46	0.45
1:B:456:ILE:HG22	1:B:460:LEU:HD11	1.98	0.45
1:B:476:LEU:HD23	1:B:477:LEU:N	2.31	0.45
1:B:826:LYS:CB	1:B:830:LYS:HA	2.46	0.45
1:B:885:LEU:C	1:B:888:THR:H	2.22	0.45
1:A:18:TYR:CE2	1:A:26:PHE:CE1	3.04	0.45
1:A:40:GLN:CG	1:A:205:GLY:O	2.58	0.45
1:A:142:LEU:HD21	1:A:198:HIS:HB3	1.96	0.45
1:A:185:ASN:OD1	1:A:191:GLN:HB3	2.16	0.45
1:A:249:LYS:NZ	1:A:418:ILE:HG21	2.31	0.45
1:B:133:THR:CG2	1:B:152:ASP:HB3	2.42	0.45
1:B:134:VAL:HG13	1:B:147:MET:CB	2.36	0.45
1:B:183:VAL:HG12	1:B:192:PRO:CB	2.43	0.45
1:B:450:SER:HA	1:B:453:LEU:CD1	2.46	0.45
1:B:897:CYS:O	1:B:898:VAL:CB	2.64	0.45
1:B:1004:SER:O	1:B:1005:ASN:C	2.57	0.45
1:A:14:ILE:O	1:A:224:MET:CB	2.64	0.45
1:A:25:GLY:HA2	1:A:40:GLN:O	2.15	0.45
1:A:65:ARG:NH1	1:A:103:ASN:N	2.61	0.45
1:A:219:LYS:HE2	1:A:221:VAL:HG23	1.96	0.45
1:A:227:SER:O	1:A:228:ASP:CB	2.62	0.45
1:A:246:GLU:CB	1:A:428:GLU:OE2	2.63	0.45
1:A:317:GLU:O	1:A:353:TYR:HD2	2.00	0.45
1:A:452:VAL:HB	1:A:476:LEU:CD2	2.47	0.45
1:A:1330:VAL:O	1:A:1334:LEU:CB	2.64	0.45
1:B:59:LYS:HG2	1:B:124:LEU:HD22	1.94	0.45
1:B:66:TYR:HB3	1:B:100:LYS:HZ3	1.81	0.45
1:B:240:VAL:O	1:B:282:TRP:HB2	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:385:ASN:ND2	1:B:433:VAL:HG13	2.31	0.45
1:B:477:LEU:HA	1:B:480:LEU:HD13	1.90	0.45
1:B:518:GLN:NE2	1:B:518:GLN:HA	2.30	0.45
1:B:580:PHE:O	1:B:581:MET:C	2.57	0.45
1:A:161:PHE:HA	1:A:187:VAL:H	1.81	0.45
1:A:314:LEU:CD1	1:A:355:LEU:HD23	2.46	0.45
1:A:417:LYS:HZ3	1:A:419:GLY:HA3	1.81	0.45
1:A:748:GLN:O	1:A:1074:PRO:HA	2.17	0.45
1:A:1341:VAL:O	1:A:1344:PHE:N	2.49	0.45
1:B:33:VAL:HB	1:B:448:ASP:CG	2.41	0.45
1:B:46:LEU:HD23	1:B:46:LEU:N	2.30	0.45
1:B:68:ALA:HB1	1:B:96:ALA:N	2.32	0.45
1:B:135:ASN:O	1:B:136:LYS:C	2.58	0.45
1:B:138:LEU:O	1:B:148:ARG:CA	2.65	0.45
1:B:194:HIS:CE1	1:B:212:VAL:O	2.69	0.45
1:B:212:VAL:HG13	1:B:213:ASN:N	2.32	0.45
1:B:418:ILE:HD12	1:B:418:ILE:HG23	1.43	0.45
1:B:474:THR:O	1:B:478:GLU:OE1	2.34	0.45
1:B:503:ASN:HB3	1:B:506:ARG:HG3	1.96	0.45
1:B:724:LYS:CB	1:B:858:GLU:CA	2.93	0.45
1:B:743:MET:C	1:B:745:LEU:N	2.74	0.45
1:A:123:LEU:O	1:A:131:TYR:CB	2.65	0.45
1:A:402:THR:OG1	1:A:404:ILE:HD12	2.16	0.45
1:A:1061:LEU:HA	1:A:1101:GLN:HA	1.98	0.45
1:A:1326:CYS:O	1:A:1330:VAL:CB	2.65	0.45
1:A:1421:ALA:O	1:A:1425:PHE:CA	2.64	0.45
1:B:35:ASP:HB2	1:B:151:LEU:CD1	2.46	0.45
1:B:55:ASP:CA	1:B:127:LYS:HZ2	2.30	0.45
1:B:152:ASP:N	1:B:152:ASP:OD1	2.49	0.45
1:B:192:PRO:O	1:B:211:SER:CA	2.65	0.45
1:B:223:PHE:CG	1:B:294:GLY:O	2.70	0.45
1:B:224:MET:HB2	1:B:295:GLY:CA	2.43	0.45
1:B:371:ASP:CG	1:B:371:ASP:O	2.59	0.45
1:B:391:ARG:HG2	1:B:392:HIS:N	2.31	0.45
1:B:450:SER:N	1:B:515:ILE:HD13	2.32	0.45
1:B:466:THR:O	1:B:470:ARG:CB	2.64	0.45
1:B:1400:LEU:CB	1:B:1442:THR:HA	2.47	0.45
1:A:19:ALA:HB2	1:A:218:TRP:HZ3	1.80	0.45
1:A:66:TYR:O	1:A:73:TRP:CZ2	2.70	0.45
1:A:69:GLN:HE21	1:A:99:GLU:HB2	1.82	0.45
1:A:128:SER:O	1:A:130:LYS:CG	2.63	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:166:PHE:CZ	1:A:217:SER:CA	2.95	0.45
1:A:304:ARG:NE	1:A:367:ILE:HG23	2.32	0.45
1:A:455:SER:CA	1:A:459:LYS:HE2	2.46	0.45
1:A:458:GLY:H	1:A:459:LYS:NZ	2.12	0.45
1:A:511:ARG:N	1:A:516:LEU:CD2	2.80	0.45
1:A:666:ILE:CB	1:A:670:LEU:H	2.30	0.45
1:A:870:ALA:O	1:A:871:ARG:C	2.59	0.45
1:B:384:ARG:HB2	1:B:434:PRO:HD3	1.98	0.45
1:B:518:GLN:N	1:B:518:GLN:NE2	2.64	0.45
1:B:1246:ALA:HA	1:B:1286:GLU:HA	1.98	0.45
1:A:67:SER:N	1:A:99:GLU:OE2	2.50	0.45
1:A:179:GLY:O	1:A:220:ILE:HB	2.17	0.45
1:A:184:LEU:CA	1:A:185:ASN:HD22	2.28	0.45
1:A:230:LYS:HE2	1:A:230:LYS:N	2.31	0.45
1:A:242:LEU:O	1:A:250:PHE:CD1	2.70	0.45
1:A:244:HIS:CB	1:A:430:PHE:CD1	3.00	0.45
1:A:249:LYS:C	1:A:264:LEU:CD1	2.87	0.45
1:A:281:LEU:O	1:A:308:LEU:CG	2.65	0.45
1:A:298:TYR:CD1	1:A:381:LEU:HA	2.52	0.45
1:A:384:ARG:HG3	1:A:385:ASN:ND2	2.31	0.45
1:A:402:THR:N	1:A:418:ILE:CG2	2.59	0.45
1:A:406:ILE:CD1	1:A:415:MET:HG3	2.47	0.45
1:B:44:GLY:CA	1:B:50:PRO:HD3	2.35	0.45
1:B:184:LEU:CG	1:B:193:LEU:HB3	2.46	0.45
1:B:249:LYS:HG2	1:B:264:LEU:CB	2.46	0.45
1:B:249:LYS:HB3	1:B:264:LEU:CD1	2.46	0.45
1:B:298:TYR:O	1:B:298:TYR:HD1	1.99	0.45
1:B:304:ARG:N	1:B:367:ILE:CG2	2.80	0.45
1:B:353:TYR:CZ	1:B:395:THR:HG23	2.52	0.45
1:B:417:LYS:HA	1:B:417:LYS:HD2	1.80	0.45
1:B:1203:ARG:O	1:B:1207:ASN:N	2.50	0.45
1:A:10:HIS:CD2	1:A:113:THR:C	2.95	0.45
1:A:18:TYR:CB	1:A:219:LYS:CD	2.95	0.45
1:A:71:GLN:C	1:A:92:LEU:HD21	2.41	0.45
1:A:162:TYR:CE1	1:A:187:VAL:CG1	3.00	0.45
1:A:196:SER:O	1:A:198:HIS:N	2.50	0.45
1:A:393:LEU:HB3	1:A:394:CYS:SG	2.57	0.45
1:A:454:GLY:N	1:A:518:GLN:CD	2.66	0.45
1:B:27:ILE:HD11	1:B:37:CYS:SG	2.57	0.45
1:B:131:TYR:HD1	1:B:131:TYR:HA	1.44	0.45
1:B:133:THR:O	1:B:152:ASP:OD2	2.34	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:200:LEU:HD12	1:B:207:ASN:C	2.41	0.45
1:B:248:GLU:O	1:B:250:PHE:CZ	2.69	0.45
1:B:299:TRP:H	1:B:381:LEU:N	2.14	0.45
1:B:744:CYS:HA	1:B:1085:ARG:HA	1.92	0.45
1:A:223:PHE:HE2	1:A:292:CYS:CB	2.25	0.45
1:A:242:LEU:HD21	1:A:251:LEU:HD22	1.97	0.45
1:A:249:LYS:NZ	1:A:418:ILE:CG2	2.77	0.45
1:A:257:ARG:O	1:A:258:LYS:CB	2.65	0.45
1:A:317:GLU:OE1	1:A:317:GLU:N	2.50	0.45
1:A:407:ASP:OD2	1:A:413:PRO:CG	2.65	0.45
1:A:477:LEU:HD11	1:A:552:ILE:HG23	1.94	0.45
1:A:1442:THR:O	1:A:1445:HIS:CB	2.65	0.45
1:B:26:PHE:CE1	1:B:45:ASP:CA	2.99	0.45
1:B:135:ASN:ND2	1:B:148:ARG:HD3	2.32	0.45
1:B:299:TRP:CD1	1:B:380:SER:HA	2.51	0.45
1:B:300:ASN:ND2	1:B:369:GLU:HG3	2.32	0.45
1:B:360:GLU:O	1:B:363:ASP:HB2	2.17	0.45
1:B:477:LEU:CA	1:B:480:LEU:CD1	2.84	0.45
1:B:827:ASP:O	1:B:830:LYS:CB	2.65	0.45
1:B:1341:VAL:O	1:B:1344:PHE:C	2.60	0.45
1:B:1433:THR:CB	1:B:1492:SER:CB	2.95	0.45
1:A:7:SER:C	1:A:178:ILE:HD13	2.41	0.45
1:A:161:PHE:CB	1:A:185:ASN:O	2.65	0.45
1:A:244:HIS:CD2	1:A:429:ALA:N	2.80	0.45
1:A:451:LYS:NZ	1:A:451:LYS:N	2.65	0.45
1:A:1126:VAL:C	1:A:1130:GLU:H	2.23	0.45
1:B:247:GLN:OE1	1:B:403:ASN:ND2	2.48	0.45
1:B:259:LYS:CE	1:B:261:HIS:CE1	3.00	0.45
1:B:464:THR:OG1	1:B:465:ILE:N	2.49	0.45
1:B:473:VAL:O	1:B:477:LEU:CG	2.60	0.45
1:B:1244:PHE:CB	1:B:1248:ASN:CB	2.95	0.45
1:B:1432:ASP:O	1:B:1433:THR:CB	2.64	0.45
1:A:10:HIS:NE2	1:A:114:VAL:CG2	2.80	0.44
1:A:30:LEU:HG	1:A:36:ARG:O	2.17	0.44
1:A:508:LYS:HE3	1:A:512:GLU:HG3	2.00	0.44
1:B:88:LEU:O	1:B:92:LEU:HB2	2.17	0.44
1:B:130:LYS:CE	1:B:151:LEU:HD13	2.45	0.44
1:B:308:LEU:N	1:B:308:LEU:CD2	2.75	0.44
1:B:440:VAL:H	1:B:441:ARG:NH2	2.12	0.44
1:A:18:TYR:CE2	1:A:26:PHE:CG	3.06	0.44
1:A:20:GLU:CD	1:A:217:SER:HB3	2.42	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:30:LEU:HD13	1:A:34:ASP:HB3	1.99	0.44
1:A:33:VAL:HG11	1:A:448:ASP:O	2.16	0.44
1:A:129:ASN:CG	1:A:441:ARG:NH1	2.70	0.44
1:A:161:PHE:N	1:A:187:VAL:HG23	2.32	0.44
1:A:244:HIS:HB3	1:A:247:GLN:N	2.31	0.44
1:A:244:HIS:NE2	1:A:428:GLU:HB3	2.32	0.44
1:A:370:LEU:CA	1:A:390:LEU:HD13	2.34	0.44
1:A:417:LYS:HB3	1:A:418:ILE:H	1.31	0.44
1:A:456:ILE:HA	1:A:459:LYS:CG	2.47	0.44
1:A:978:LEU:O	1:A:982:LEU:CB	2.65	0.44
1:B:14:ILE:HD12	1:B:14:ILE:N	2.29	0.44
1:B:16:SER:C	1:B:220:ILE:HD13	2.42	0.44
1:B:181:LYS:HA	1:B:218:TRP:H	1.82	0.44
1:B:241:ARG:CZ	1:B:433:VAL:HG23	2.45	0.44
1:B:504:ARG:O	1:B:507:GLN:HB2	2.17	0.44
1:B:524:GLN:O	1:B:524:GLN:CD	2.60	0.44
1:B:847:VAL:C	1:B:849:CYS:N	2.74	0.44
1:B:1371:LEU:O	1:B:1375:ILE:CB	2.65	0.44
1:A:18:TYR:CE2	1:A:26:PHE:CD1	3.05	0.44
1:A:143:GLU:CG	1:A:210:ASN:HD21	2.30	0.44
1:A:256:HIS:ND1	1:A:257:ARG:HG2	2.32	0.44
1:A:302:LEU:HD12	1:A:302:LEU:HA	1.23	0.44
1:A:1005:ASN:O	1:A:1006:SER:CB	2.65	0.44
1:B:11:ILE:HG23	1:B:60:LEU:O	2.18	0.44
1:B:286:VAL:HG11	1:B:288:GLN:HG2	1.98	0.44
1:B:465:ILE:O	1:B:466:THR:CB	2.65	0.44
1:B:1219:GLN:HA	1:B:1230:MET:CB	2.47	0.44
1:A:65:ARG:NH2	1:A:103:ASN:N	2.64	0.44
1:A:87:VAL:CG1	1:A:91:LYS:HE3	2.48	0.44
1:A:163:ILE:C	1:A:164:GLN:CD	2.86	0.44
1:A:201:VAL:O	1:A:204:PRO:CG	2.65	0.44
1:A:263:PHE:CB	1:A:406:ILE:CD1	2.94	0.44
1:A:298:TYR:CZ	1:A:381:LEU:HD22	2.52	0.44
1:A:299:TRP:CZ3	1:A:370:LEU:HB2	2.43	0.44
1:A:451:LYS:HE3	1:A:451:LYS:HB3	1.25	0.44
1:A:1382:ALA:HB1	1:A:1423:ILE:HA	1.98	0.44
1:B:17:LEU:C	1:B:46:LEU:HD21	2.42	0.44
1:B:167:TYR:CD1	1:B:169:LEU:CD1	3.00	0.44
1:B:220:ILE:O	1:B:220:ILE:CG2	2.65	0.44
1:B:287:VAL:H	1:B:304:ARG:CD	2.31	0.44
1:B:319:ASP:OD2	1:B:354:SER:CB	2.65	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:439:GLU:HA	1:B:442:ASP:CG	2.42	0.44
1:B:484:VAL:O	1:B:562:HIS:ND1	2.50	0.44
1:B:485:THR:N	1:B:506:ARG:NH1	2.64	0.44
1:B:598:LEU:HA	1:B:602:ASN:CB	2.48	0.44
1:A:166:PHE:CE2	1:A:217:SER:HB3	2.53	0.44
1:A:230:LYS:HA	1:A:230:LYS:NZ	2.33	0.44
1:A:235:LYS:CE	1:A:384:ARG:NH1	2.75	0.44
1:A:244:HIS:HA	1:A:430:PHE:CD1	2.52	0.44
1:A:304:ARG:HD2	1:A:304:ARG:HA	1.80	0.44
1:A:438:ALA:O	1:A:442:ASP:CG	2.60	0.44
1:A:518:GLN:NE2	1:A:522:LEU:HD11	2.10	0.44
1:B:14:ILE:O	1:B:223:PHE:CB	2.66	0.44
1:B:63:MET:HE1	1:B:64:ASN:O	2.18	0.44
1:B:69:GLN:HB3	1:B:100:LYS:HZ1	1.79	0.44
1:B:145:ASN:O	1:B:146:ALA:HB2	2.18	0.44
1:B:263:PHE:CD2	1:B:263:PHE:C	2.94	0.44
1:B:285:GLU:C	1:B:304:ARG:HD2	2.43	0.44
1:B:299:TRP:H	1:B:381:LEU:CA	2.31	0.44
1:B:560:LEU:HD21	1:B:564:GLN:NE2	2.32	0.44
1:B:666:ILE:N	1:B:667:GLU:C	2.75	0.44
1:B:1292:VAL:CB	1:B:1344:PHE:O	2.65	0.44
1:A:9:LEU:HD22	1:A:220:ILE:CG2	2.48	0.44
1:A:19:ALA:CB	1:A:218:TRP:HZ3	2.29	0.44
1:A:122:GLN:O	1:A:123:LEU:HG	2.18	0.44
1:A:196:SER:CB	1:A:198:HIS:CE1	3.00	0.44
1:A:256:HIS:CE1	1:A:257:ARG:HG2	2.53	0.44
1:A:511:ARG:CA	1:A:516:LEU:HD11	2.45	0.44
1:A:574:ILE:O	1:A:578:PHE:CB	2.66	0.44
1:A:1363:ASP:O	1:A:1364:ARG:C	2.58	0.44
1:B:16:SER:O	1:B:46:LEU:HD11	2.17	0.44
1:B:33:VAL:CG2	1:B:445:PHE:CD2	3.01	0.44
1:B:47:ASN:C	1:B:291:PRO:HG3	2.36	0.44
1:B:115:ILE:HD11	1:B:121:ILE:CG2	2.46	0.44
1:B:184:LEU:HD13	1:B:193:LEU:HD13	1.98	0.44
1:B:266:THR:HG22	1:B:266:THR:O	2.18	0.44
1:B:307:HIS:HB2	1:B:312:HIS:H	1.83	0.44
1:B:392:HIS:CG	1:B:395:THR:HB	2.53	0.44
1:B:503:ASN:CB	1:B:506:ARG:HG3	2.48	0.44
1:B:510:MET:CB	1:B:515:ILE:CG2	2.84	0.44
1:B:726:ASP:O	1:B:727:ARG:C	2.60	0.44
1:B:1218:LEU:O	1:B:1230:MET:O	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:26:PHE:CZ	1:A:49:PRO:HA	2.52	0.44
1:A:170:ARG:HH11	1:A:170:ARG:HG3	1.83	0.44
1:A:263:PHE:N	1:A:263:PHE:CD1	2.84	0.44
1:A:304:ARG:HD3	1:A:367:ILE:HG22	1.95	0.44
1:A:314:LEU:O	1:A:358:VAL:HG22	2.18	0.44
1:A:452:VAL:C	1:A:456:ILE:HD12	2.43	0.44
1:B:16:SER:O	1:B:46:LEU:CD1	2.66	0.44
1:B:16:SER:HB2	1:B:292:CYS:HB3	1.98	0.44
1:B:110:LEU:HD23	1:B:110:LEU:N	2.33	0.44
1:B:223:PHE:CD1	1:B:293:ARG:HA	2.42	0.44
1:B:244:HIS:NE2	1:B:248:GLU:N	2.65	0.44
1:B:385:ASN:OD1	1:B:433:VAL:CB	2.65	0.44
1:B:404:ILE:O	1:B:416:LEU:N	2.51	0.44
1:B:683:GLY:O	1:B:687:LEU:CB	2.65	0.44
1:A:106:GLU:O	1:A:109:LYS:HE2	2.17	0.44
1:A:110:LEU:O	1:A:110:LEU:CG	2.47	0.44
1:A:127:LYS:HZ2	1:A:127:LYS:HB3	1.83	0.44
1:A:142:LEU:HD22	1:A:198:HIS:CG	2.53	0.44
1:A:256:HIS:HE1	1:A:257:ARG:HE	1.66	0.44
1:A:282:TRP:HA	1:A:308:LEU:HG	1.99	0.44
1:A:283:GLU:HG2	1:A:306:LYS:CB	2.32	0.44
1:A:683:GLY:O	1:A:684:GLU:C	2.61	0.44
1:A:1313:GLN:O	1:A:1314:THR:C	2.61	0.44
1:B:26:PHE:HZ	1:B:45:ASP:C	2.25	0.44
1:B:75:ALA:HB2	1:B:92:LEU:HD12	1.99	0.44
1:B:169:LEU:O	1:B:170:ARG:HG2	2.17	0.44
1:B:477:LEU:O	1:B:478:GLU:C	2.57	0.44
1:B:1109:VAL:O	1:B:1110:ASP:C	2.61	0.44
1:A:62:PRO:HG3	1:A:110:LEU:CD2	2.48	0.44
1:A:139:PRO:O	1:A:148:ARG:NE	2.51	0.44
1:A:140:ALA:HA	1:A:148:ARG:HD3	1.99	0.44
1:A:193:LEU:CD1	1:A:218:TRP:HE1	2.27	0.44
1:A:353:TYR:CG	1:A:397:THR:HG22	2.53	0.44
1:A:384:ARG:HE	1:A:385:ASN:HD21	1.66	0.44
1:A:696:VAL:O	1:A:700:TRP:CB	2.66	0.44
1:A:1231:GLN:O	1:A:1235:ARG:CB	2.65	0.44
1:B:389:ARG:HA	1:B:428:GLU:O	2.18	0.44
1:B:449:ALA:C	1:B:453:LEU:HD21	2.43	0.44
1:B:472:SER:O	1:B:473:VAL:C	2.60	0.44
1:B:478:GLU:OE1	1:B:479:ASP:N	2.50	0.44
1:B:515:ILE:O	1:B:519:ILE:HD13	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:139:PRO:CD	1:A:148:ARG:HE	2.31	0.43
1:A:170:ARG:CB	1:A:180:ASP:OD2	2.66	0.43
1:A:239:VAL:HG22	1:A:435:VAL:HG23	2.00	0.43
1:A:288:GLN:NE2	1:A:288:GLN:CA	2.73	0.43
1:A:303:PHE:CD1	1:A:370:LEU:HD21	2.52	0.43
1:A:387:TYR:HA	1:A:430:PHE:C	2.42	0.43
1:A:511:ARG:CB	1:A:512:GLU:OE2	2.62	0.43
1:A:1064:LEU:CB	1:A:1101:GLN:CB	2.96	0.43
1:B:11:ILE:CG2	1:B:61:CYS:CA	2.95	0.43
1:B:72:PHE:HA	1:B:92:LEU:CD1	2.40	0.43
1:B:128:SER:OG	1:B:151:LEU:CD2	2.66	0.43
1:B:249:LYS:NZ	1:B:416:LEU:HG	2.33	0.43
1:B:287:VAL:H	1:B:304:ARG:HD3	1.83	0.43
1:A:162:TYR:CE1	1:A:187:VAL:HG13	2.53	0.43
1:A:452:VAL:O	1:A:453:LEU:C	2.59	0.43
1:A:515:ILE:HG23	1:A:515:ILE:HD12	1.32	0.43
1:B:18:TYR:CA	1:B:46:LEU:HG	2.40	0.43
1:B:63:MET:HE1	1:B:65:ARG:CA	2.47	0.43
1:B:102:GLN:NE2	1:B:106:GLU:CD	2.77	0.43
1:B:115:ILE:HG22	1:B:176:VAL:CG2	2.48	0.43
1:B:196:SER:CB	1:B:209:VAL:HA	2.47	0.43
1:B:263:PHE:HE2	1:B:265:ARG:HG2	1.83	0.43
1:B:384:ARG:HD3	1:B:434:PRO:HB3	2.00	0.43
1:B:399:VAL:C	1:B:420:THR:HG1	2.23	0.43
1:B:435:VAL:HG12	1:B:439:GLU:OE2	2.16	0.43
1:B:476:LEU:O	1:B:479:ASP:N	2.51	0.43
1:A:15:CYS:HB3	1:A:223:PHE:N	2.34	0.43
1:A:58:PHE:HZ	1:A:123:LEU:C	2.27	0.43
1:A:265:ARG:CB	1:A:414:VAL:C	2.91	0.43
1:A:399:VAL:HA	1:A:420:THR:HA	2.00	0.43
1:A:443:LEU:O	1:A:444:ASP:C	2.60	0.43
1:A:456:ILE:HG21	1:A:473:VAL:CG2	2.48	0.43
1:A:507:GLN:NE2	1:A:562:HIS:HD2	2.10	0.43
1:A:769:SER:O	1:A:770:ASP:C	2.57	0.43
1:A:1202:GLN:C	1:A:1204:LEU:N	2.74	0.43
1:B:9:LEU:O	1:B:10:HIS:CG	2.71	0.43
1:B:65:ARG:O	1:B:65:ARG:HG2	2.18	0.43
1:B:96:ALA:O	1:B:97:ASP:C	2.60	0.43
1:B:109:LYS:HE2	1:B:109:LYS:HA	2.00	0.43
1:B:207:ASN:ND2	1:B:207:ASN:C	2.76	0.43
1:B:265:ARG:O	1:B:267:THR:N	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:365:SER:HB2	1:B:368:PHE:HE1	1.83	0.43
1:B:1379:GLU:O	1:B:1383:VAL:N	2.28	0.43
1:A:7:SER:O	1:A:178:ILE:CA	2.66	0.43
1:A:11:ILE:HD11	1:A:115:ILE:CG1	2.48	0.43
1:A:18:TYR:CB	1:A:219:LYS:HD2	2.48	0.43
1:A:39:VAL:HB	1:A:207:ASN:C	2.43	0.43
1:A:40:GLN:CD	1:A:43:ALA:HB2	2.43	0.43
1:A:234:LEU:HD13	1:A:384:ARG:HA	1.99	0.43
1:A:384:ARG:HE	1:A:385:ASN:ND2	2.16	0.43
1:B:46:LEU:HB3	1:B:221:VAL:CG1	2.49	0.43
1:B:130:LYS:HB2	1:B:153:GLU:OE2	2.19	0.43
1:B:287:VAL:O	1:B:288:GLN:OE1	2.36	0.43
1:B:355:LEU:HD13	1:B:417:LYS:NZ	2.33	0.43
1:B:385:ASN:H	1:B:432:ILE:C	2.26	0.43
1:B:403:ASN:HA	1:B:416:LEU:HD11	1.96	0.43
1:B:485:THR:C	1:B:501:LYS:O	2.62	0.43
1:A:451:LYS:O	1:A:452:VAL:C	2.62	0.43
1:A:510:MET:HB3	1:A:516:LEU:HD21	1.95	0.43
1:A:1129:SER:O	1:A:1133:VAL:N	2.52	0.43
1:A:1332:ALA:CB	1:A:1376:HIS:CB	2.95	0.43
1:B:116:GLN:CD	1:B:174:ASP:N	2.74	0.43
1:B:287:VAL:N	1:B:304:ARG:NE	2.66	0.43
1:B:456:ILE:HD12	1:B:469:GLU:OE2	2.17	0.43
1:B:1090:ARG:CA	1:B:1176:VAL:N	2.81	0.43
1:B:1250:GLN:O	1:B:1251:ASN:C	2.61	0.43
1:A:17:LEU:HD21	1:A:218:TRP:HB3	2.01	0.43
1:A:18:TYR:HD1	1:A:24:ASN:HB2	1.84	0.43
1:A:26:PHE:HE1	1:A:44:GLY:C	2.26	0.43
1:A:41:PRO:HD2	1:A:42:GLU:OE2	2.17	0.43
1:A:98:LEU:HB3	1:A:102:GLN:NE2	2.34	0.43
1:A:117:TYR:CD2	1:A:172:ILE:HA	2.54	0.43
1:A:138:LEU:HD13	1:A:148:ARG:CZ	2.49	0.43
1:A:157:GLU:CB	1:A:188:ASN:HD21	2.31	0.43
1:A:222:LEU:N	1:A:222:LEU:HD13	2.33	0.43
1:A:406:ILE:CD1	1:A:415:MET:CB	2.91	0.43
1:A:406:ILE:HD11	1:A:417:LYS:HA	2.01	0.43
1:A:446:ALA:CB	1:A:515:ILE:HG12	2.46	0.43
1:A:475:LYS:O	1:A:476:LEU:C	2.61	0.43
1:A:1422:TYR:HA	1:A:1426:LEU:CB	2.49	0.43
1:B:57:LEU:O	1:B:58:PHE:HD1	2.01	0.43
1:B:63:MET:O	1:B:107:ASN:ND2	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:242:LEU:HD21	1:B:282:TRP:CZ2	2.54	0.43
1:B:319:ASP:HB2	1:B:354:SER:OG	2.19	0.43
1:B:390:LEU:HD11	1:B:399:VAL:HG23	2.01	0.43
1:B:430:PHE:O	1:B:432:ILE:HD11	2.19	0.43
1:B:703:SER:O	1:B:733:TYR:O	2.36	0.43
1:B:1218:LEU:O	1:B:1230:MET:CB	2.66	0.43
1:A:27:ILE:HD13	1:A:27:ILE:HA	1.90	0.43
1:A:40:GLN:HA	1:A:207:ASN:ND2	2.34	0.43
1:A:139:PRO:N	1:A:148:ARG:HE	2.16	0.43
1:A:445:PHE:CD2	1:A:483:PHE:CG	3.06	0.43
1:A:465:ILE:C	1:A:466:THR:OG1	2.61	0.43
1:B:59:LYS:O	1:B:61:CYS:SG	2.76	0.43
1:B:63:MET:HG3	1:B:160:TRP:HE1	1.84	0.43
1:B:141:LEU:HD23	1:B:208:GLU:OE2	2.18	0.43
1:B:286:VAL:HG12	1:B:288:GLN:N	2.34	0.43
1:B:385:ASN:ND2	1:B:433:VAL:HG22	2.32	0.43
1:B:457:ALA:O	1:B:458:GLY:C	2.61	0.43
1:B:475:LYS:O	1:B:479:ASP:OD2	2.36	0.43
1:A:18:TYR:CE2	1:A:26:PHE:CE2	3.05	0.43
1:A:66:TYR:HH	1:A:100:LYS:HG2	1.80	0.43
1:A:119:ASN:O	1:A:163:ILE:HG22	2.19	0.43
1:A:123:LEU:O	1:A:131:TYR:HD1	2.02	0.43
1:A:183:VAL:HG11	1:A:192:PRO:CB	2.49	0.43
1:A:379:ASP:C	1:A:381:LEU:H	2.27	0.43
1:A:1316:VAL:CB	1:A:1323:ILE:O	2.67	0.43
1:B:66:TYR:CD1	1:B:100:LYS:CE	3.02	0.43
1:B:241:ARG:NE	1:B:433:VAL:O	2.52	0.43
1:B:304:ARG:NH2	1:B:367:ILE:HD12	2.34	0.43
1:B:305:PHE:CE1	1:B:367:ILE:HG22	2.53	0.43
1:B:743:MET:C	1:B:745:LEU:H	2.27	0.43
1:B:1242:GLN:O	1:B:1243:ASN:CB	2.67	0.43
1:A:87:VAL:HA	1:A:90:ASN:HD22	1.76	0.43
1:A:166:PHE:HE2	1:A:218:TRP:H	1.66	0.43
1:A:282:TRP:CE3	1:A:308:LEU:N	2.86	0.43
1:A:363:ASP:CG	1:A:366:SER:OG	2.61	0.43
1:A:364:ILE:HA	1:A:367:ILE:CG1	2.46	0.43
1:A:391:ARG:HD2	1:A:398:TRP:CG	2.54	0.43
1:A:457:ALA:C	1:A:459:LYS:CD	2.92	0.43
1:A:658:ASN:O	1:A:659:PRO:C	2.61	0.43
1:B:135:ASN:OD1	1:B:148:ARG:O	2.36	0.43
1:B:168:LYS:NZ	1:B:168:LYS:CB	2.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:242:LEU:C	1:B:430:PHE:CE2	2.97	0.43
1:B:389:ARG:N	1:B:389:ARG:HD3	2.34	0.43
1:B:397:THR:HB	1:B:422:PRO:C	2.44	0.43
1:B:448:ASP:O	1:B:448:ASP:OD1	2.37	0.43
1:B:504:ARG:NH1	1:B:566:ASP:C	2.76	0.43
1:B:560:LEU:HD22	1:B:564:GLN:HG3	1.99	0.43
1:B:1134:TYR:C	1:B:1233:ILE:CB	2.92	0.43
1:A:11:ILE:HG22	1:A:12:GLY:H	1.82	0.43
1:A:30:LEU:CB	1:A:34:ASP:O	2.63	0.43
1:A:181:LYS:C	1:A:182:VAL:HG13	2.44	0.43
1:A:259:LYS:N	1:A:259:LYS:CE	2.79	0.43
1:A:368:PHE:C	1:A:393:LEU:CD2	2.76	0.43
1:A:555:LEU:HD23	1:A:555:LEU:HA	1.22	0.43
1:B:13:ASP:OD1	1:B:13:ASP:C	2.62	0.43
1:B:32:LEU:N	1:B:32:LEU:CD2	2.73	0.43
1:B:54:ARG:NH1	1:B:55:ASP:H	2.17	0.43
1:B:66:TYR:CE1	1:B:100:LYS:CA	2.89	0.43
1:B:96:ALA:O	1:B:100:LYS:CE	2.66	0.43
1:B:385:ASN:H	1:B:432:ILE:CA	2.32	0.43
1:B:421:SER:N	1:B:425:GLU:O	2.52	0.43
1:B:484:VAL:HG23	1:B:506:ARG:CD	2.48	0.43
1:A:234:LEU:HD12	1:A:234:LEU:HA	1.88	0.42
1:A:243:PHE:HE1	1:A:431:ALA:HB3	1.84	0.42
1:A:244:HIS:HD2	1:A:430:PHE:CA	2.31	0.42
1:A:280:ALA:O	1:A:282:TRP:CD1	2.72	0.42
1:A:290:ASP:HB3	1:A:291:PRO:HD2	2.00	0.42
1:A:459:LYS:HD2	1:A:460:LEU:H	1.84	0.42
1:A:886:ARG:C	1:A:888:THR:N	2.75	0.42
1:A:1427:ASN:O	1:A:1431:VAL:CB	2.67	0.42
1:B:13:ASP:C	1:B:14:ILE:HG13	2.39	0.42
1:B:94:HIS:CE1	1:B:98:LEU:CD1	3.02	0.42
1:B:181:LYS:NZ	1:B:218:TRP:CE3	2.87	0.42
1:B:195:ALA:C	1:B:209:VAL:HA	2.43	0.42
1:B:235:LYS:O	1:B:238:ASP:HB2	2.19	0.42
1:B:460:LEU:C	1:B:462:LYS:CE	2.92	0.42
1:B:523:LEU:N	1:B:523:LEU:HD22	2.34	0.42
1:B:856:ASP:CA	1:B:860:ASN:H	2.29	0.42
1:A:304:ARG:NH1	1:A:367:ILE:HD13	2.34	0.42
1:A:316:ALA:HB1	1:A:355:LEU:CA	2.47	0.42
1:A:353:TYR:CB	1:A:420:THR:O	2.67	0.42
1:A:371:ASP:CG	1:A:398:TRP:CZ3	2.97	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:504:ARG:HB3	1:A:505:GLU:OE1	2.20	0.42
1:B:71:GLN:HA	1:B:71:GLN:HE21	1.84	0.42
1:B:138:LEU:O	1:B:147:MET:C	2.62	0.42
1:B:171:SER:C	1:B:172:ILE:HG13	2.42	0.42
1:B:179:GLY:HA2	1:B:220:ILE:O	2.19	0.42
1:B:230:LYS:HD3	1:B:230:LYS:HA	1.85	0.42
1:B:420:THR:CG2	1:B:425:GLU:O	2.58	0.42
1:B:555:LEU:HA	1:B:555:LEU:HD23	1.70	0.42
1:B:1438:LYS:O	1:B:1444:ASN:HA	2.20	0.42
1:A:20:GLU:CD	1:A:181:LYS:CE	2.92	0.42
1:A:72:PHE:O	1:A:72:PHE:CD1	2.73	0.42
1:A:140:ALA:CA	1:A:148:ARG:HA	2.49	0.42
1:A:162:TYR:CD1	1:A:187:VAL:HG13	2.55	0.42
1:A:186:PRO:O	1:A:188:ASN:N	2.46	0.42
1:A:222:LEU:C	1:A:222:LEU:CD2	2.89	0.42
1:A:250:PHE:HZ	1:A:280:ALA:HB1	1.71	0.42
1:A:255:GLU:OE1	1:A:259:LYS:N	2.52	0.42
1:A:418:ILE:HG23	1:A:418:ILE:HD13	1.43	0.42
1:A:454:GLY:C	1:A:459:LYS:HZ3	2.25	0.42
1:A:456:ILE:O	1:A:459:LYS:CD	2.61	0.42
1:A:631:LEU:O	1:A:634:LEU:CB	2.67	0.42
1:B:111:LEU:HD13	1:B:226:TRP:CH2	2.44	0.42
1:B:122:GLN:HE21	1:B:122:GLN:HB3	1.65	0.42
1:B:241:ARG:CZ	1:B:433:VAL:CB	2.97	0.42
1:B:249:LYS:HG3	1:B:266:THR:N	2.34	0.42
1:B:1191:SER:CB	1:B:1236:LEU:O	2.67	0.42
1:B:1401:PRO:N	1:B:1442:THR:CB	2.82	0.42
1:A:63:MET:CE	1:A:120:VAL:CG1	2.83	0.42
1:A:437:PRO:O	1:A:441:ARG:CG	2.67	0.42
1:A:473:VAL:O	1:A:477:LEU:HG	2.19	0.42
1:A:510:MET:SD	1:A:516:LEU:HB3	2.60	0.42
1:A:1062:ARG:O	1:A:1066:HIS:N	2.45	0.42
1:A:1225:ALA:N	1:A:1271:VAL:CB	2.82	0.42
1:B:26:PHE:CZ	1:B:45:ASP:HA	2.53	0.42
1:B:137:ARG:HA	1:B:147:MET:CB	2.49	0.42
1:B:168:LYS:HB2	1:B:168:LYS:HZ2	1.84	0.42
1:B:287:VAL:HG22	1:B:290:ASP:N	2.33	0.42
1:B:407:ASP:CG	1:B:413:PRO:HA	2.45	0.42
1:B:515:ILE:C	1:B:518:GLN:HB2	2.42	0.42
1:A:11:ILE:CG2	1:A:12:GLY:N	2.82	0.42
1:A:46:LEU:O	1:A:291:PRO:CG	2.67	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:58:PHE:CE2	1:A:123:LEU:C	2.98	0.42
1:A:235:LYS:HZ3	1:A:235:LYS:HG2	1.58	0.42
1:A:250:PHE:CZ	1:A:280:ALA:CB	2.88	0.42
1:A:368:PHE:O	1:A:393:LEU:HD22	2.14	0.42
1:A:389:ARG:CZ	1:A:389:ARG:CB	2.97	0.42
1:A:447:ASN:O	1:A:450:SER:CB	2.64	0.42
1:A:449:ALA:HB3	1:A:515:ILE:CD1	2.50	0.42
1:B:14:ILE:CB	1:B:223:PHE:HD2	2.24	0.42
1:B:37:CYS:O	1:B:38:VAL:HG13	2.18	0.42
1:B:66:TYR:OH	1:B:103:ASN:ND2	2.52	0.42
1:B:68:ALA:HA	1:B:71:GLN:HG2	2.02	0.42
1:B:111:LEU:C	1:B:226:TRP:HZ3	2.27	0.42
1:B:130:LYS:CB	1:B:153:GLU:OE2	2.68	0.42
1:B:192:PRO:CD	1:B:212:VAL:O	2.67	0.42
1:B:196:SER:HB2	1:B:210:ASN:H	1.83	0.42
1:B:277:SER:HA	1:B:508:LYS:CE	2.49	0.42
1:B:355:LEU:N	1:B:417:LYS:NZ	2.67	0.42
1:B:372:PRO:HB3	1:B:388:VAL:CG1	2.48	0.42
1:B:395:THR:CG2	1:B:397:THR:CG2	2.95	0.42
1:B:430:PHE:O	1:B:432:ILE:CD1	2.67	0.42
1:B:441:ARG:H	1:B:441:ARG:NH2	2.15	0.42
1:A:87:VAL:CA	1:A:90:ASN:HD22	2.30	0.42
1:A:185:ASN:OD1	1:A:191:GLN:HA	2.20	0.42
1:A:192:PRO:O	1:A:211:SER:O	2.37	0.42
1:A:226:TRP:H	1:A:226:TRP:HD1	1.65	0.42
1:A:242:LEU:O	1:A:250:PHE:HD1	2.01	0.42
1:A:246:GLU:CB	1:A:429:ALA:CB	2.94	0.42
1:A:319:ASP:O	1:A:319:ASP:CG	2.63	0.42
1:A:387:TYR:HA	1:A:430:PHE:O	2.20	0.42
1:A:391:ARG:NH2	1:A:395:THR:C	2.72	0.42
1:A:450:SER:HG	1:A:451:LYS:HZ2	1.59	0.42
1:B:263:PHE:CZ	1:B:414:VAL:CG2	3.02	0.42
1:B:384:ARG:HB3	1:B:432:ILE:O	2.20	0.42
1:B:724:LYS:O	1:B:725:GLU:CB	2.68	0.42
1:A:10:HIS:C	1:A:13:ASP:OD2	2.63	0.42
1:A:11:ILE:HB	1:A:113:THR:HG1	1.83	0.42
1:A:41:PRO:HD2	1:A:42:GLU:H	1.83	0.42
1:A:371:ASP:CG	1:A:398:TRP:CH2	2.98	0.42
1:A:649:GLN:O	1:A:650:GLU:C	2.61	0.42
1:A:1270:ALA:C	1:A:1272:THR:N	2.76	0.42
1:B:234:LEU:HD22	1:B:234:LEU:HA	1.45	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:238:ASP:H	1:B:284:VAL:CG2	2.31	0.42
1:B:410:GLU:HB3	1:B:411:GLU:H	1.49	0.42
1:B:518:GLN:HE21	1:B:518:GLN:CA	2.32	0.42
1:B:1058:ARG:O	1:B:1059:THR:CB	2.66	0.42
1:A:11:ILE:N	1:A:13:ASP:OD2	2.52	0.42
1:A:98:LEU:O	1:A:102:GLN:CG	2.68	0.42
1:A:118:GLY:CA	1:A:163:ILE:O	2.66	0.42
1:A:181:LYS:HA	1:A:220:ILE:HD13	2.02	0.42
1:A:303:PHE:CE2	1:A:370:LEU:CG	3.02	0.42
1:A:319:ASP:N	1:A:354:SER:OG	2.53	0.42
1:B:197:SER:HA	1:B:207:ASN:CG	2.37	0.42
1:B:256:HIS:HB3	1:B:259:LYS:HZ1	1.84	0.42
1:B:387:TYR:C	1:B:388:VAL:HG13	2.42	0.42
1:B:477:LEU:CA	1:B:555:LEU:HD22	2.49	0.42
1:B:480:LEU:HB3	1:B:483:PHE:HZ	1.85	0.42
1:B:509:LEU:O	1:B:513:GLN:OE1	2.38	0.42
1:B:660:THR:O	1:B:661:ASN:CB	2.68	0.42
1:B:964:ASP:C	1:B:967:VAL:H	2.26	0.42
1:B:1251:ASN:O	1:B:1255:LEU:O	2.38	0.42
1:A:14:ILE:HG12	1:A:60:LEU:H	1.84	0.42
1:A:33:VAL:HG13	1:A:448:ASP:O	2.19	0.42
1:A:63:MET:CE	1:A:160:TRP:CD2	3.00	0.42
1:A:65:ARG:NH2	1:A:103:ASN:CA	2.75	0.42
1:A:298:TYR:CE1	1:A:381:LEU:CD2	2.99	0.42
1:A:305:PHE:CE1	1:A:390:LEU:HD21	2.54	0.42
1:A:367:ILE:O	1:A:368:PHE:HD1	2.03	0.42
1:A:465:ILE:C	1:A:466:THR:HG1	2.24	0.42
1:A:476:LEU:CA	1:A:479:ASP:OD1	2.68	0.42
1:A:477:LEU:HD12	1:A:552:ILE:HA	1.99	0.42
1:A:559:VAL:O	1:A:563:SER:HB2	2.20	0.42
1:B:46:LEU:HD22	1:B:46:LEU:HA	1.54	0.42
1:B:58:PHE:CG	1:B:123:LEU:HD12	2.55	0.42
1:B:75:ALA:O	1:B:76:ALA:HB3	2.20	0.42
1:B:184:LEU:HD13	1:B:193:LEU:CD2	2.44	0.42
1:B:223:PHE:HA	1:B:294:GLY:H	1.84	0.42
1:B:263:PHE:HD2	1:B:264:LEU:O	2.03	0.42
1:B:264:LEU:N	1:B:416:LEU:O	2.49	0.42
1:B:281:LEU:HD11	1:B:440:VAL:HG22	2.02	0.42
1:B:435:VAL:HG11	1:B:439:GLU:OE2	2.19	0.42
1:B:449:ALA:O	1:B:452:VAL:HG23	2.19	0.42
1:A:11:ILE:CG1	1:A:115:ILE:HG13	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:24:ASN:OD1	1:A:24:ASN:O	2.38	0.42
1:A:124:LEU:CD1	1:A:125:HIS:H	2.28	0.42
1:A:308:LEU:CD1	1:A:309:ALA:H	2.21	0.42
1:A:406:ILE:CA	1:A:417:LYS:HG2	2.50	0.42
1:A:450:SER:HB2	1:A:451:LYS:HZ1	1.85	0.42
1:A:480:LEU:O	1:A:481:VAL:C	2.61	0.42
1:A:628:PRO:O	1:A:629:ARG:C	2.63	0.42
1:A:784:LEU:O	1:A:785:MET:C	2.61	0.42
1:A:1405:ILE:O	1:A:1408:VAL:N	2.51	0.42
1:B:10:HIS:HB3	1:B:112:GLY:CA	2.50	0.42
1:B:49:PRO:O	1:B:50:PRO:C	2.63	0.42
1:B:68:ALA:HB3	1:B:96:ALA:HA	2.02	0.42
1:B:192:PRO:N	1:B:212:VAL:O	2.53	0.42
1:B:262:VAL:HG23	1:B:357:SER:N	2.35	0.42
1:B:393:LEU:HD22	1:B:393:LEU:N	2.35	0.42
1:B:407:ASP:OD2	1:B:413:PRO:HA	2.19	0.42
1:A:87:VAL:HG12	1:A:91:LYS:HE2	2.01	0.41
1:A:473:VAL:HG23	1:A:477:LEU:HD21	2.02	0.41
1:A:1225:ALA:HB1	1:A:1226:GLU:HA	2.02	0.41
1:B:102:GLN:C	1:B:106:GLU:OE2	2.63	0.41
1:B:130:LYS:CG	1:B:151:LEU:CD2	2.98	0.41
1:B:224:MET:SD	1:B:288:GLN:OE1	2.78	0.41
1:B:243:PHE:H	1:B:431:ALA:H	1.67	0.41
1:B:259:LYS:NZ	1:B:261:HIS:CE1	2.87	0.41
1:B:263:PHE:CG	1:B:415:MET:HA	2.50	0.41
1:B:418:ILE:HB	1:B:419:GLY:H	1.70	0.41
1:B:507:GLN:CG	1:B:563:SER:CA	2.89	0.41
1:B:847:VAL:O	1:B:849:CYS:O	2.38	0.41
1:B:1003:GLU:O	1:B:1004:SER:C	2.62	0.41
1:B:1488:THR:O	1:B:1489:PHE:CB	2.67	0.41
1:A:61:CYS:O	1:A:122:GLN:O	2.38	0.41
1:A:69:GLN:HB2	1:A:73:TRP:HE1	1.84	0.41
1:A:90:ASN:H	1:A:90:ASN:HD22	1.68	0.41
1:A:252:THR:O	1:A:263:PHE:CD1	2.74	0.41
1:A:263:PHE:N	1:A:263:PHE:HD1	2.18	0.41
1:A:314:LEU:HD13	1:A:355:LEU:HD23	2.02	0.41
1:A:410:GLU:HB2	1:A:413:PRO:HB3	1.96	0.41
1:A:422:PRO:O	1:A:423:LEU:C	2.63	0.41
1:A:643:LYS:O	1:A:647:VAL:CB	2.68	0.41
1:B:20:GLU:CD	1:B:20:GLU:O	2.63	0.41
1:B:66:TYR:CB	1:B:69:GLN:CG	2.94	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:68:ALA:CB	1:B:96:ALA:HA	2.50	0.41
1:B:103:ASN:HA	1:B:106:GLU:CG	2.50	0.41
1:B:133:THR:HB	1:B:152:ASP:HB2	2.02	0.41
1:B:244:HIS:CE1	1:B:249:LYS:CB	2.94	0.41
1:B:353:TYR:CD2	1:B:395:THR:HG21	2.54	0.41
1:B:485:THR:H	1:B:506:ARG:CZ	2.32	0.41
1:A:8:PHE:CD2	1:A:177:VAL:HG22	2.55	0.41
1:A:28:SER:HG	1:A:55:ASP:CG	2.07	0.41
1:A:316:ALA:CA	1:A:355:LEU:HA	2.50	0.41
1:A:407:ASP:CB	1:A:415:MET:HG2	2.49	0.41
1:B:148:ARG:NH2	1:B:150:THR:CG2	2.66	0.41
1:B:200:LEU:CD1	1:B:208:GLU:N	2.83	0.41
1:B:245:ALA:CB	1:B:387:TYR:HE2	2.32	0.41
1:B:452:VAL:O	1:B:453:LEU:C	2.63	0.41
1:B:519:ILE:C	1:B:522:LEU:CD2	2.93	0.41
1:A:185:ASN:OD1	1:A:191:GLN:CB	2.67	0.41
1:A:241:ARG:HH11	1:A:280:ALA:HB3	1.85	0.41
1:A:244:HIS:N	1:A:248:GLU:C	2.75	0.41
1:A:287:VAL:CB	1:A:302:LEU:HB3	2.50	0.41
1:A:412:LYS:N	1:A:413:PRO:HD3	2.35	0.41
1:A:466:THR:O	1:A:470:ARG:CB	2.68	0.41
1:A:511:ARG:CA	1:A:516:LEU:HD21	2.51	0.41
1:B:26:PHE:O	1:B:39:VAL:HB	2.18	0.41
1:B:63:MET:CG	1:B:160:TRP:NE1	2.79	0.41
1:B:63:MET:CG	1:B:160:TRP:HE1	2.33	0.41
1:B:116:GLN:OE1	1:B:173:GLY:N	2.53	0.41
1:B:195:ALA:O	1:B:209:VAL:HG22	2.19	0.41
1:B:248:GLU:N	1:B:248:GLU:CD	2.78	0.41
1:B:263:PHE:CZ	1:B:414:VAL:HG23	2.56	0.41
1:B:391:ARG:HG3	1:B:398:TRP:CD1	2.56	0.41
1:B:716:ALA:O	1:B:717:GLN:C	2.63	0.41
1:B:728:ASP:CB	1:B:866:VAL:CA	2.98	0.41
1:B:1000:GLU:H	1:B:1006:SER:N	2.17	0.41
1:A:7:SER:CA	1:A:178:ILE:HD13	2.51	0.41
1:A:14:ILE:CB	1:A:58:PHE:O	2.53	0.41
1:A:62:PRO:CG	1:A:110:LEU:CD2	2.93	0.41
1:A:244:HIS:CE1	1:A:428:GLU:OE2	2.73	0.41
1:A:248:GLU:O	1:A:250:PHE:N	2.53	0.41
1:A:391:ARG:NH2	1:A:392:HIS:HB3	2.35	0.41
1:A:1094:LEU:O	1:A:1095:GLN:C	2.62	0.41
1:A:1270:ALA:HB1	1:A:1316:VAL:HA	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:101:LYS:O	1:B:102:GLN:O	2.38	0.41
1:B:194:HIS:CG	1:B:210:ASN:O	2.73	0.41
1:B:249:LYS:HZ3	1:B:266:THR:HG1	1.56	0.41
1:B:281:LEU:HD11	1:B:440:VAL:CG2	2.50	0.41
1:B:286:VAL:CA	1:B:304:ARG:NE	2.59	0.41
1:B:383:PRO:HG2	1:B:386:SER:OG	2.20	0.41
1:B:595:ILE:O	1:B:598:LEU:CB	2.69	0.41
1:B:1096:ALA:O	1:B:1097:PHE:C	2.64	0.41
1:A:166:PHE:HB2	1:A:181:LYS:O	2.21	0.41
1:A:407:ASP:OD2	1:A:413:PRO:CB	2.68	0.41
1:A:1000:GLU:CB	1:A:1005:ASN:CA	2.99	0.41
1:B:166:PHE:HZ	1:B:216:THR:N	2.18	0.41
1:B:406:ILE:O	1:B:408:LYS:CD	2.51	0.41
1:A:20:GLU:OE2	1:A:218:TRP:CA	2.68	0.41
1:A:27:ILE:O	1:A:56:CYS:CB	2.68	0.41
1:A:64:ASN:C	1:A:160:TRP:CZ2	2.99	0.41
1:A:117:TYR:HE2	1:A:174:ASP:H	1.62	0.41
1:A:129:ASN:N	1:A:129:ASN:HD22	2.17	0.41
1:A:151:LEU:N	1:A:151:LEU:HD23	2.31	0.41
1:A:368:PHE:CE1	1:A:392:HIS:CD2	3.09	0.41
1:A:1092:GLU:O	1:A:1093:VAL:CB	2.68	0.41
1:A:1418:VAL:O	1:A:1422:TYR:CB	2.69	0.41
1:B:60:LEU:HD23	1:B:60:LEU:C	2.45	0.41
1:B:219:LYS:HE2	1:B:221:VAL:HG12	2.02	0.41
1:B:263:PHE:CZ	1:B:415:MET:CE	3.03	0.41
1:B:305:PHE:O	1:B:313:TYR:CE2	2.73	0.41
1:B:371:ASP:OD2	1:B:389:ARG:CG	2.69	0.41
1:B:390:LEU:O	1:B:398:TRP:HB3	2.20	0.41
1:B:398:TRP:O	1:B:420:THR:CA	2.66	0.41
1:B:406:ILE:HG13	1:B:415:MET:HB3	2.03	0.41
1:B:479:ASP:C	1:B:481:VAL:H	2.28	0.41
1:B:516:LEU:HD23	1:B:563:SER:CB	2.50	0.41
1:B:551:HIS:CG	1:B:552:ILE:HD13	2.56	0.41
1:B:1370:PRO:HA	1:B:1413:ASP:C	2.46	0.41
1:A:53:PHE:CD1	1:A:53:PHE:O	2.73	0.41
1:A:124:LEU:CD1	1:A:129:ASN:C	2.87	0.41
1:A:124:LEU:N	1:A:131:TYR:HD1	2.19	0.41
1:A:138:LEU:CB	1:A:148:ARG:HG3	2.51	0.41
1:A:169:LEU:HD23	1:A:170:ARG:HH22	1.85	0.41
1:A:249:LYS:HB3	1:A:264:LEU:CD2	2.50	0.41
1:A:391:ARG:HE	1:A:392:HIS:CA	2.34	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:417:LYS:O	1:A:418:ILE:CD1	2.69	0.41
1:A:445:PHE:CD1	1:A:445:PHE:C	2.94	0.41
1:A:518:GLN:C	1:A:522:LEU:HD22	2.45	0.41
1:A:861:LYS:O	1:A:862:LEU:C	2.62	0.41
1:A:1061:LEU:O	1:A:1062:ARG:C	2.62	0.41
1:B:122:GLN:HB3	1:B:131:TYR:OH	2.21	0.41
1:B:208:GLU:HG2	1:B:209:VAL:N	2.36	0.41
1:B:257:ARG:O	1:B:257:ARG:CG	2.49	0.41
1:B:515:ILE:CA	1:B:518:GLN:HB2	2.51	0.41
1:B:1204:LEU:C	1:B:1206:ARG:HA	2.46	0.41
1:B:1355:ILE:O	1:B:1357:MET:CB	2.69	0.41
1:A:12:GLY:C	1:A:111:LEU:CD2	2.85	0.41
1:A:41:PRO:HD2	1:A:42:GLU:CD	2.46	0.41
1:A:57:LEU:N	1:A:57:LEU:HD23	2.36	0.41
1:A:58:PHE:CE1	1:A:124:LEU:C	2.99	0.41
1:A:60:LEU:CD1	1:A:123:LEU:CD2	2.97	0.41
1:A:108:ARG:HB2	1:A:108:ARG:NH1	2.25	0.41
1:A:124:LEU:CD1	1:A:130:LYS:N	2.83	0.41
1:A:142:LEU:HD22	1:A:198:HIS:CD2	2.56	0.41
1:A:151:LEU:HD22	1:A:151:LEU:HA	1.56	0.41
1:A:238:ASP:O	1:A:284:VAL:HG23	2.21	0.41
1:A:251:LEU:O	1:A:251:LEU:CD2	2.67	0.41
1:A:263:PHE:CE2	1:A:414:VAL:HG23	2.48	0.41
1:A:387:TYR:HA	1:A:431:ALA:HB2	1.98	0.41
1:A:647:VAL:O	1:A:648:THR:C	2.63	0.41
1:A:1225:ALA:HA	1:A:1226:GLU:C	2.46	0.41
1:B:12:GLY:H	1:B:60:LEU:HD23	1.83	0.41
1:B:15:CYS:SG	1:B:221:VAL:O	2.75	0.41
1:B:32:LEU:HD11	1:B:445:PHE:N	2.36	0.41
1:B:34:ASP:CG	1:B:448:ASP:OD1	2.64	0.41
1:B:54:ARG:N	1:B:54:ARG:NH1	2.59	0.41
1:B:56:CYS:CA	1:B:125:HIS:NE2	2.69	0.41
1:B:72:PHE:CE1	1:B:89:LEU:O	2.73	0.41
1:B:116:GLN:OE1	1:B:174:ASP:O	2.39	0.41
1:B:130:LYS:CD	1:B:151:LEU:C	2.91	0.41
1:B:164:GLN:O	1:B:182:VAL:HB	2.21	0.41
1:B:193:LEU:HA	1:B:211:SER:HB2	2.02	0.41
1:B:219:LYS:HE3	1:B:219:LYS:C	2.46	0.41
1:B:224:MET:HG3	1:B:295:GLY:N	2.35	0.41
1:B:239:VAL:HG23	1:B:282:TRP:C	2.46	0.41
1:B:281:LEU:HB2	1:B:308:LEU:HB2	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:286:VAL:CG1	1:B:303:PHE:HD1	2.27	0.41
1:B:379:ASP:O	1:B:379:ASP:OD2	2.39	0.41
1:B:400:HIS:CG	1:B:402:THR:OG1	2.73	0.41
1:B:460:LEU:N	1:B:462:LYS:CE	2.70	0.41
1:B:507:GLN:N	1:B:507:GLN:CD	2.63	0.41
1:B:598:LEU:CA	1:B:602:ASN:CB	2.98	0.41
1:B:654:LYS:O	1:B:658:ASN:CA	2.65	0.41
1:B:1130:GLU:HA	1:B:1217:LEU:CB	2.50	0.41
1:B:1134:TYR:CA	1:B:1233:ILE:CB	2.98	0.41
1:B:1270:ALA:HA	1:B:1315:ILE:CB	2.51	0.41
1:B:1491:SER:O	1:B:1492:SER:O	2.38	0.41
1:A:12:GLY:HA3	1:A:111:LEU:CG	2.51	0.41
1:A:26:PHE:HE1	1:A:44:GLY:CA	2.33	0.41
1:A:30:LEU:CD1	1:A:36:ARG:HB2	2.51	0.41
1:A:58:PHE:CE2	1:A:124:LEU:O	2.73	0.41
1:A:256:HIS:ND1	1:A:257:ARG:N	2.69	0.41
1:A:263:PHE:HD2	1:A:415:MET:SD	2.40	0.41
1:A:271:SER:CB	1:A:276:THR:HG23	2.51	0.41
1:A:317:GLU:O	1:A:353:TYR:CD2	2.74	0.41
1:A:384:ARG:CD	1:A:385:ASN:OD1	2.69	0.41
1:A:1276:ILE:O	1:A:1279:ASN:N	2.54	0.41
1:B:19:ALA:O	1:B:20:GLU:HB3	2.20	0.41
1:B:19:ALA:C	1:B:20:GLU:OE1	2.63	0.41
1:B:364:ILE:O	1:B:365:SER:CB	2.68	0.41
1:B:478:GLU:HA	1:B:555:LEU:HD21	2.03	0.41
1:A:20:GLU:HG3	1:A:181:LYS:HD3	2.03	0.40
1:A:20:GLU:CD	1:A:218:TRP:N	2.79	0.40
1:A:100:LYS:HG3	1:A:100:LYS:H	1.55	0.40
1:A:373:THR:HG22	1:A:389:ARG:HD3	2.03	0.40
1:A:458:GLY:O	1:A:462:LYS:HG3	2.20	0.40
1:A:705:LYS:CA	1:A:736:GLN:CB	2.97	0.40
1:A:1126:VAL:HA	1:A:1130:GLU:H	1.86	0.40
1:A:1285:SER:HA	1:A:1344:PHE:CB	2.50	0.40
1:B:17:LEU:C	1:B:46:LEU:HD11	2.46	0.40
1:B:259:LYS:CE	1:B:261:HIS:HE1	2.33	0.40
1:B:408:LYS:CE	1:B:408:LYS:CA	2.97	0.40
1:A:14:ILE:HG12	1:A:60:LEU:N	2.36	0.40
1:A:63:MET:C	1:A:160:TRP:HZ2	2.28	0.40
1:A:139:PRO:C	1:A:140:ALA:O	2.63	0.40
1:A:167:TYR:OH	1:A:181:LYS:HD3	2.21	0.40
1:A:181:LYS:HG2	1:A:218:TRP:C	2.46	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:181:LYS:HA	1:A:218:TRP:O	2.21	0.40
1:A:235:LYS:H	1:A:235:LYS:HG2	1.09	0.40
1:A:241:ARG:NE	1:A:276:THR:OG1	2.53	0.40
1:A:257:ARG:C	1:A:259:LYS:CE	2.94	0.40
1:A:317:GLU:C	1:A:353:TYR:HD2	2.29	0.40
1:A:352:VAL:HG12	1:A:353:TYR:HD1	1.87	0.40
1:A:456:ILE:N	1:A:459:LYS:NZ	2.56	0.40
1:A:1234:MET:O	1:A:1238:HIS:CB	2.69	0.40
1:B:115:ILE:CD1	1:B:121:ILE:CG2	2.88	0.40
1:B:241:ARG:NH2	1:B:433:VAL:CB	2.85	0.40
1:B:244:HIS:CD2	1:B:247:GLN:C	3.00	0.40
1:B:283:GLU:HB2	1:B:308:LEU:HD21	2.03	0.40
1:B:299:TRP:O	1:B:300:ASN:CB	2.69	0.40
1:B:370:LEU:HD23	1:B:370:LEU:N	2.36	0.40
1:B:371:ASP:CG	1:B:398:TRP:CZ3	2.99	0.40
1:B:777:LEU:O	1:B:780:SER:N	2.55	0.40
1:B:1081:GLN:O	1:B:1082:LEU:C	2.65	0.40
1:A:72:PHE:CA	1:A:92:LEU:CD2	2.99	0.40
1:A:391:ARG:HG2	1:A:398:TRP:HA	1.82	0.40
1:A:453:LEU:CG	1:A:518:GLN:HB3	2.50	0.40
1:A:514:ASN:HD22	1:A:518:GLN:N	2.19	0.40
1:A:1029:ASP:O	1:A:1030:PHE:C	2.65	0.40
1:A:1134:TYR:CB	1:A:1177:LYS:O	2.69	0.40
1:B:36:ARG:NH1	1:B:200:LEU:CD2	2.76	0.40
1:B:130:LYS:HG2	1:B:151:LEU:HD23	2.02	0.40
1:B:160:TRP:HB2	1:B:187:VAL:HG11	2.03	0.40
1:B:288:GLN:O	1:B:289:HIS:HB3	2.21	0.40
1:B:439:GLU:N	1:B:441:ARG:HH21	2.20	0.40
1:B:736:GLN:O	1:B:737:LEU:C	2.64	0.40
1:B:1274:GLN:CB	1:B:1329:MET:C	2.94	0.40
1:A:170:ARG:HB3	1:A:180:ASP:OD2	2.20	0.40
1:A:369:GLU:N	1:A:393:LEU:CD1	2.82	0.40
1:A:377:GLY:O	1:A:378:GLY:C	2.64	0.40
1:A:465:ILE:H	1:A:465:ILE:HG13	1.54	0.40
1:A:510:MET:HE2	1:A:515:ILE:CG2	2.47	0.40
1:A:744:CYS:CA	1:A:1077:SER:CB	2.96	0.40
1:B:19:ALA:CB	1:B:218:TRP:CH2	3.04	0.40
1:B:65:ARG:O	1:B:65:ARG:CG	2.70	0.40
1:B:181:LYS:CE	1:B:219:LYS:HG2	2.34	0.40
1:B:198:HIS:C	1:B:200:LEU:N	2.64	0.40
1:B:219:LYS:NZ	1:B:220:ILE:C	2.80	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:226:TRP:O	1:B:229:ASN:ND2	2.55	0.40
1:B:240:VAL:O	1:B:282:TRP:CB	2.70	0.40
1:B:242:LEU:HD21	1:B:282:TRP:CH2	2.55	0.40
1:B:286:VAL:HG12	1:B:288:GLN:H	1.85	0.40
1:B:286:VAL:HG21	1:B:303:PHE:CE2	2.56	0.40
1:B:666:ILE:N	1:B:667:GLU:O	2.55	0.40
1:B:695:GLU:O	1:B:699:PHE:CB	2.70	0.40
1:B:1118:ASP:O	1:B:1120:ASP:N	2.53	0.40
1:A:26:PHE:CE2	1:A:53:PHE:CE2	3.09	0.40
1:A:60:LEU:CD1	1:A:123:LEU:HD21	2.51	0.40
1:A:71:GLN:O	1:A:92:LEU:HD11	2.21	0.40
1:A:249:LYS:CE	1:A:264:LEU:HD22	2.51	0.40
1:A:313:TYR:C	1:A:358:VAL:HG23	2.46	0.40
1:A:407:ASP:OD1	1:A:415:MET:HG2	2.20	0.40
1:A:453:LEU:HD23	1:A:518:GLN:OE1	2.21	0.40
1:A:457:ALA:O	1:A:458:GLY:C	2.64	0.40
1:A:509:LEU:O	1:A:510:MET:C	2.63	0.40
1:A:510:MET:SD	1:A:516:LEU:CA	3.10	0.40
1:A:992:CYS:O	1:A:996:ILE:N	2.37	0.40
1:B:74:LYS:HB3	1:B:74:LYS:HE2	1.85	0.40
1:B:111:LEU:CD2	1:B:226:TRP:CH2	2.98	0.40
1:B:135:ASN:HB3	1:B:138:LEU:HD21	2.04	0.40
1:B:186:PRO:O	1:B:190:GLY:HA2	2.21	0.40
1:B:241:ARG:HA	1:B:282:TRP:CD1	2.57	0.40
1:B:256:HIS:CE1	1:B:407:ASP:CG	2.98	0.40
1:B:466:THR:O	1:B:470:ARG:N	2.47	0.40
1:B:682:THR:N	1:B:686:ALA:CB	2.85	0.40
1:B:1198:ARG:CB	1:B:1239:GLU:C	2.94	0.40

All (4) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:376:ARG:O	1:B:378:GLY:N[2_655]	1.78	0.42
1:A:374:THR:N	1:A:376:ARG:O[2_655]	1.96	0.24
1:B:377:GLY:N	1:B:379:ASP:OD1[2_655]	2.15	0.05
1:A:373:THR:OG1	1:A:376:ARG:O[2_655]	2.16	0.04

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 X-ray entries followed by that with respect to entries of similar resolution.

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	1230/1581 (78%)	1000 (81%)	155 (13%)	75 (6%)	1	12
1	B	1230/1581 (78%)	980 (80%)	165 (13%)	85 (7%)	1	11
All	All	2460/3162 (78%)	1980 (80%)	320 (13%)	160 (6%)	1	12

All (160) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	49	PRO
1	A	50	PRO
1	A	104	GLU
1	A	140	ALA
1	A	192	PRO
1	A	227	SER
1	A	231	ASP
1	A	248	GLU
1	A	249	LYS
1	A	372	PRO
1	A	396	ASN
1	A	423	LEU
1	A	424	LYS
1	A	628	PRO
1	A	639	VAL
1	A	659	PRO
1	A	666	ILE
1	A	717	GLN
1	A	739	LEU
1	A	740	PHE
1	A	774	PRO
1	A	1006	SER
1	A	1049	PRO
1	A	1051	ASP
1	A	1058	ARG

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Mol	Chain	Res	Type
1	A	1074	PRO
1	A	1093	VAL
1	A	1103	LEU
1	A	1133	VAL
1	A	1194	VAL
1	A	1203	ARG
1	A	1241	LEU
1	A	1249	GLN
1	A	1264	ASN
1	A	1323	ILE
1	A	1370	PRO
1	A	1400	LEU
1	A	1492	SER
1	B	43	ALA
1	B	50	PRO
1	B	146	ALA
1	B	172	ILE
1	B	200	LEU
1	B	212	VAL
1	B	225	LYS
1	B	257	ARG
1	B	300	ASN
1	B	410	GLU
1	B	450	SER
1	B	611	THR
1	B	639	VAL
1	B	737	LEU
1	B	749	TYR
1	B	760	ASP
1	B	773	LEU
1	B	790	VAL
1	B	851	ARG
1	B	898	VAL
1	B	1002	ASP
1	B	1049	PRO
1	B	1195	ARG
1	B	1243	ASN
1	B	1249	GLN
1	B	1370	PRO
1	B	1385	THR
1	B	1401	PRO
1	B	1433	THR

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Mol	Chain	Res	Type
1	B	1442	THR
1	B	1461	CYS
1	B	1492	SER
1	A	147	MET
1	A	224	MET
1	A	407	ASP
1	A	510	MET
1	A	664	ILE
1	A	702	ASP
1	A	708	ARG
1	A	762	ASP
1	A	854	PHE
1	A	1005	ASN
1	A	1059	THR
1	A	1347	ASP
1	A	1466	ASP
1	B	196	SER
1	B	199	GLN
1	B	362	ASN
1	B	381	LEU
1	B	401	SER
1	B	416	LEU
1	B	418	ILE
1	B	458	GLY
1	B	661	ASN
1	B	744	CYS
1	B	745	LEU
1	B	789	HIS
1	B	882	SER
1	B	1129	SER
1	B	1261	LEU
1	B	1315	ILE
1	A	258	LYS
1	A	274	SER
1	A	302	LEU
1	A	761	VAL
1	A	1322	PHE
1	A	1365	MET
1	A	1413	ASP
1	A	1459	ARG
1	B	153	GLU
1	B	222	LEU

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Mol	Chain	Res	Type
1	B	249	LYS
1	B	364	ILE
1	B	423	LEU
1	B	704	ASN
1	B	961	GLU
1	B	992	CYS
1	B	1059	THR
1	B	1221	PRO
1	A	103	ASN
1	A	383	PRO
1	A	853	PRO
1	A	1002	ASP
1	B	278	SER
1	B	412	LYS
1	B	689	ALA
1	B	1180	LEU
1	B	1395	LYS
1	B	1435	VAL
1	B	1488	THR
1	A	187	VAL
1	A	223	PHE
1	A	378	GLY
1	A	449	ALA
1	A	1271	VAL
1	A	1345	TYR
1	B	139	PRO
1	B	287	VAL
1	B	377	GLY
1	B	405	PRO
1	B	466	THR
1	B	522	LEU
1	B	566	ASP
1	B	1093	VAL
1	B	1340	ASP
1	B	1356	GLN
1	A	1242	GLN
1	B	117	TYR
1	B	266	THR
1	B	658	ASN
1	B	715	LEU
1	B	201	VAL
1	B	1133	VAL

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Mol	Chain	Res	Type
1	A	178	ILE
1	A	627	GLU
1	A	240	VAL
1	A	1435	VAL
1	B	419	GLY
1	B	1378	VAL
1	A	434	PRO
1	B	473	VAL
1	B	1073	PRO

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 X-ray entries followed by that with respect to entries of similar resolution.

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	376/1427 (26%)	227 (60%)	149 (40%)	0	1
1	B	384/1427 (27%)	207 (54%)	177 (46%)	0	0
All	All	760/2854 (27%)	434 (57%)	326 (43%)	0	0

All (326) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	7	SER
1	A	8	PHE
1	A	9	LEU
1	A	10	HIS
1	A	13	ASP
1	A	14	ILE
1	A	15	CYS
1	A	16	SER
1	A	17	LEU
1	A	20	GLU
1	A	32	LEU
1	A	33	VAL
1	A	35	ASP
1	A	40	GLN

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Mol	Chain	Res	Type
1	A	42	GLU
1	A	45	ASP
1	A	48	ASN
1	A	56	CYS
1	A	60	LEU
1	A	63	MET
1	A	65	ARG
1	A	66	TYR
1	A	73	TRP
1	A	87	VAL
1	A	89	LEU
1	A	90	ASN
1	A	100	LYS
1	A	106	GLU
1	A	107	ASN
1	A	108	ARG
1	A	109	LYS
1	A	110	LEU
1	A	115	ILE
1	A	116	GLN
1	A	117	TYR
1	A	127	LYS
1	A	141	LEU
1	A	142	LEU
1	A	151	LEU
1	A	152	ASP
1	A	156	ASN
1	A	168	LYS
1	A	170	ARG
1	A	171	SER
1	A	172	ILE
1	A	181	LYS
1	A	185	ASN
1	A	187	VAL
1	A	191	GLN
1	A	198	HIS
1	A	200	LEU
1	A	201	VAL
1	A	207	ASN
1	A	212	VAL
1	A	218	TRP
1	A	219	LYS

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Mol	Chain	Res	Type
1	A	220	ILE
1	A	221	VAL
1	A	222	LEU
1	A	223	PHE
1	A	224	MET
1	A	225	LYS
1	A	228	ASP
1	A	229	ASN
1	A	230	LYS
1	A	233	ILE
1	A	235	LYS
1	A	242	LEU
1	A	244	HIS
1	A	247	GLN
1	A	248	GLU
1	A	249	LYS
1	A	251	LEU
1	A	255	GLU
1	A	257	ARG
1	A	259	LYS
1	A	263	PHE
1	A	276	THR
1	A	281	LEU
1	A	288	GLN
1	A	290	ASP
1	A	299	TRP
1	A	300	ASN
1	A	304	ARG
1	A	305	PHE
1	A	307	HIS
1	A	314	LEU
1	A	318	VAL
1	A	319	ASP
1	A	356	VAL
1	A	358	VAL
1	A	362	ASN
1	A	363	ASP
1	A	364	ILE
1	A	367	ILE
1	A	369	GLU
1	A	370	LEU
1	A	371	ASP

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Mol	Chain	Res	Type
1	A	379	ASP
1	A	382	VAL
1	A	384	ARG
1	A	388	VAL
1	A	389	ARG
1	A	391	ARG
1	A	393	LEU
1	A	394	CYS
1	A	400	HIS
1	A	404	ILE
1	A	406	ILE
1	A	407	ASP
1	A	408	LYS
1	A	414	VAL
1	A	415	MET
1	A	416	LEU
1	A	417	LYS
1	A	418	ILE
1	A	424	LYS
1	A	436	SER
1	A	439	GLU
1	A	442	ASP
1	A	443	LEU
1	A	450	SER
1	A	451	LYS
1	A	452	VAL
1	A	453	LEU
1	A	459	LYS
1	A	465	ILE
1	A	473	VAL
1	A	476	LEU
1	A	477	LEU
1	A	479	ASP
1	A	480	LEU
1	A	481	VAL
1	A	504	ARG
1	A	505	GLU
1	A	507	GLN
1	A	509	LEU
1	A	510	MET
1	A	511	ARG
1	A	513	GLN

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Mol	Chain	Res	Type
1	A	514	ASN
1	A	516	LEU
1	A	518	GLN
1	A	519	ILE
1	A	522	LEU
1	A	523	LEU
1	A	552	ILE
1	A	556	CYS
1	A	560	LEU
1	B	8	PHE
1	B	9	LEU
1	B	11	ILE
1	B	14	ILE
1	B	15	CYS
1	B	16	SER
1	B	17	LEU
1	B	20	GLU
1	B	24	ASN
1	B	26	PHE
1	B	30	LEU
1	B	32	LEU
1	B	35	ASP
1	B	37	CYS
1	B	40	GLN
1	B	45	ASP
1	B	46	LEU
1	B	47	ASN
1	B	53	PHE
1	B	54	ARG
1	B	55	ASP
1	B	60	LEU
1	B	61	CYS
1	B	65	ARG
1	B	71	GLN
1	B	91	LYS
1	B	94	HIS
1	B	98	LEU
1	B	99	GLU
1	B	100	LYS
1	B	102	GLN
1	B	103	ASN
1	B	107	ASN

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Mol	Chain	Res	Type
1	B	110	LEU
1	B	111	LEU
1	B	114	VAL
1	B	115	ILE
1	B	120	VAL
1	B	121	ILE
1	B	122	GLN
1	B	123	LEU
1	B	124	LEU
1	B	126	LEU
1	B	128	SER
1	B	129	ASN
1	B	131	TYR
1	B	132	LEU
1	B	138	LEU
1	B	141	LEU
1	B	142	LEU
1	B	148	ARG
1	B	152	ASP
1	B	153	GLU
1	B	156	ASN
1	B	163	ILE
1	B	167	TYR
1	B	168	LYS
1	B	169	LEU
1	B	172	ILE
1	B	174	ASP
1	B	178	ILE
1	B	183	VAL
1	B	184	LEU
1	B	185	ASN
1	B	187	VAL
1	B	188	ASN
1	B	193	LEU
1	B	196	SER
1	B	198	HIS
1	B	200	LEU
1	B	201	VAL
1	B	202	ASP
1	B	203	ASN
1	B	207	ASN
1	B	214	CYS

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Mol	Chain	Res	Type
1	B	215	ASN
1	B	218	TRP
1	B	219	LYS
1	B	220	ILE
1	B	221	VAL
1	B	222	LEU
1	B	223	PHE
1	B	225	LYS
1	B	227	SER
1	B	229	ASN
1	B	234	LEU
1	B	235	LYS
1	B	239	VAL
1	B	241	ARG
1	B	242	LEU
1	B	247	GLN
1	B	248	GLU
1	B	249	LYS
1	B	251	LEU
1	B	253	CYS
1	B	256	HIS
1	B	257	ARG
1	B	258	LYS
1	B	259	LYS
1	B	262	VAL
1	B	264	LEU
1	B	265	ARG
1	B	276	THR
1	B	281	LEU
1	B	286	VAL
1	B	287	VAL
1	B	288	GLN
1	B	289	HIS
1	B	290	ASP
1	B	298	TYR
1	B	299	TRP
1	B	302	LEU
1	B	304	ARG
1	B	305	PHE
1	B	307	HIS
1	B	312	HIS
1	B	313	TYR

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Mol	Chain	Res	Type
1	B	314	LEU
1	B	319	ASP
1	B	356	VAL
1	B	357	SER
1	B	363	ASP
1	B	364	ILE
1	B	366	SER
1	B	367	ILE
1	B	370	LEU
1	B	375	LEU
1	B	382	VAL
1	B	385	ASN
1	B	397	THR
1	B	401	SER
1	B	404	ILE
1	B	406	ILE
1	B	408	LYS
1	B	409	GLU
1	B	415	MET
1	B	416	LEU
1	B	418	ILE
1	B	424	LYS
1	B	433	VAL
1	B	434	PRO
1	B	435	VAL
1	B	441	ARG
1	B	442	ASP
1	B	445	PHE
1	B	450	SER
1	B	451	LYS
1	B	453	LEU
1	B	455	SER
1	B	460	LEU
1	B	462	LYS
1	B	464	THR
1	B	465	ILE
1	B	466	THR
1	B	469	GLU
1	B	475	LYS
1	B	476	LEU
1	B	477	LEU
1	B	478	GLU

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Mol	Chain	Res	Type
1	B	480	LEU
1	B	503	ASN
1	B	504	ARG
1	B	505	GLU
1	B	507	GLN
1	B	512	GLU
1	B	515	ILE
1	B	516	LEU
1	B	518	GLN
1	B	519	ILE
1	B	522	LEU
1	B	523	LEU
1	B	524	GLN
1	B	551	HIS
1	B	552	ILE
1	B	559	VAL
1	B	560	LEU
1	B	563	SER

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (35) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	10	HIS
1	A	69	GLN
1	A	90	ASN
1	A	102	GLN
1	A	103	ASN
1	A	107	ASN
1	A	122	GLN
1	A	125	HIS
1	A	185	ASN
1	A	188	ASN
1	A	207	ASN
1	A	210	ASN
1	A	247	GLN
1	A	260	GLN
1	A	288	GLN
1	A	300	ASN
1	A	400	HIS
1	A	507	GLN
1	A	514	ASN
1	A	562	HIS

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Mol	Chain	Res	Type
1	A	564	GLN
1	B	40	GLN
1	B	71	GLN
1	B	102	GLN
1	B	185	ASN
1	B	194	HIS
1	B	203	ASN
1	B	210	ASN
1	B	215	ASN
1	B	256	HIS
1	B	261	HIS
1	B	288	GLN
1	B	307	HIS
1	B	518	GLN
1	B	564	GLN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates ⓘ

There are no oligosaccharides in this entry.

5.6 Ligand geometry ⓘ

There are no ligands in this entry.

5.7 Other polymers ⓘ

There are no such residues in this entry.

5.8 Polymer linkage issues ⓘ

There are no chain breaks in this entry.

6 Fit of model and data

6.1 Protein, DNA and RNA chains

In the following table, the column labelled ‘#RSRZ> 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q< 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2			OWAB(Å ²)	Q<0.9
1	A	1252/1581 (79%)	-0.25	43 (3%)	48	43	9, 28, 68, 157	0
1	B	1252/1581 (79%)	-0.24	64 (5%)	33	34	9, 28, 69, 127	0
All	All	2504/3162 (79%)	-0.24	107 (4%)	40	39	9, 28, 69, 157	0

All (107) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
1	B	39	VAL	7.1
1	B	1062	ARG	7.1
1	A	165	PRO	6.8
1	B	1404	ASP	6.6
1	B	1301	THR	6.1
1	A	411	GLU	5.4
1	B	1460	ALA	5.2
1	A	1387	GLY	4.9
1	B	1403	ASP	4.8
1	B	1481	ILE	4.8
1	A	1232	GLU	4.7
1	B	253	CYS	4.7
1	B	1461	CYS	4.6
1	B	268	GLY	4.6
1	A	1479	THR	4.5
1	B	197	SER	4.5
1	A	150	THR	4.5
1	A	267	THR	4.5
1	A	153	GLU	4.4
1	B	1279	ASN	4.3
1	B	1459	ARG	4.1
1	A	38	VAL	4.1
1	A	1134	TYR	4.0
1	A	238	ASP	4.0

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Mol	Chain	Res	Type	RSRZ
1	B	1188	VAL	4.0
1	A	1346	ASN	3.9
1	B	1258	HIS	3.9
1	A	144	LYS	3.9
1	B	153	GLU	3.8
1	A	206	CYS	3.8
1	B	56	CYS	3.7
1	A	413	PRO	3.7
1	B	770	ASP	3.7
1	B	1078	GLY	3.7
1	A	133	THR	3.6
1	B	28	SER	3.6
1	B	1429	CYS	3.5
1	A	1411	HIS	3.4
1	A	1349	ALA	3.4
1	B	280	ALA	3.4
1	A	1493	PRO	3.4
1	A	1260	ASN	3.4
1	B	1061	LEU	3.4
1	B	1098	LYS	3.3
1	A	266	THR	3.3
1	B	580	PHE	3.3
1	B	388	VAL	3.2
1	A	1386	GLU	3.1
1	A	782	CYS	3.1
1	A	182	VAL	3.1
1	B	1101	GLN	3.1
1	B	1075	LEU	3.0
1	A	556	CYS	3.0
1	B	1275	HIS	3.0
1	B	1102	LEU	3.0
1	A	1412	GLU	3.0
1	B	53	PHE	2.9
1	A	1388	LYS	2.9
1	B	853	PRO	2.9
1	B	307	HIS	2.9
1	B	122	GLN	2.9
1	B	1065	LEU	2.8
1	A	786	LEU	2.8
1	B	1114	GLN	2.7
1	A	28	SER	2.7
1	B	579	GLY	2.7

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Mol	Chain	Res	Type	RSRZ
1	A	1233	ILE	2.7
1	B	62	PRO	2.7
1	B	125	HIS	2.7
1	A	204	PRO	2.6
1	B	1493	PRO	2.6
1	B	699	PHE	2.6
1	A	210	ASN	2.6
1	B	371	ASP	2.6
1	B	112	GLY	2.5
1	B	748	GLN	2.5
1	A	1449	LEU	2.5
1	B	1002	ASP	2.5
1	B	57	LEU	2.5
1	B	163	ILE	2.4
1	A	51	LYS	2.4
1	A	581	MET	2.4
1	A	1279	ASN	2.4
1	B	1187	CYS	2.4
1	A	183	VAL	2.4
1	B	1079	ALA	2.3
1	B	1346	ASN	2.3
1	B	372	PRO	2.3
1	B	126	LEU	2.3
1	B	124	LEU	2.3
1	B	300	ASN	2.3
1	A	35	ASP	2.3
1	A	1132	TRP	2.2
1	B	556	CYS	2.2
1	A	1234	MET	2.2
1	B	1256	HIS	2.2
1	B	1184	SER	2.2
1	B	38	VAL	2.2
1	B	1107	GLN	2.1
1	A	961	GLU	2.1
1	B	61	CYS	2.1
1	A	789	HIS	2.1
1	B	1463	ASN	2.1
1	B	1430	TYR	2.1
1	B	267	THR	2.1
1	B	1278	MET	2.0
1	A	1492	SER	2.0

6.2 Non-standard residues in protein, DNA, RNA chains [i](#)

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

6.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

6.4 Ligands [i](#)

There are no ligands in this entry.

6.5 Other polymers [i](#)

There are no such residues in this entry.