



Full wwPDB X-ray Structure Validation Report ⓘ

Feb 1, 2016 – 09:29 AM GMT

PDB ID : 3IKM
Title : Crystal structure of human mitochondrial DNA polymerase holoenzyme
Authors : Lee, Y-S.; Kennedy, W.D.; Yin, Y.W.
Deposited on : 2009-08-06
Resolution : 3.24 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
<http://wwpdb.org/validation/2016/XrayValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467
Mogul : 1.7 (RC4), CSD as536be (2015)
Xtriage (Phenix) : 1.9-1692
EDS : rb-20026688
Percentile statistics : 20151230.v01 (using entries in the PDB archive December 30th 2015)
Refmac : 5.8.0135
CCP4 : 6.5.0
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : trunk26865

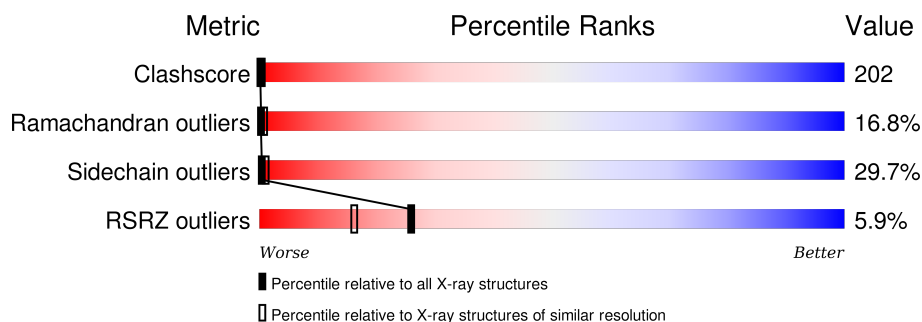
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.24 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))
Clashscore	102246	1227 (3.28-3.20)
Ramachandran outliers	100387	1204 (3.28-3.20)
Sidechain outliers	100360	1203 (3.28-3.20)
RSRZ outliers	91569	1097 (3.28-3.20)

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 on the lower bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria. 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	1172	<div> <div>5%</div> <div>5%</div> <div>45%</div> <div>34%</div> <div>9%</div> <div>7%</div> </div>
1	D	1172	<div> <div>5%</div> <div>5%</div> <div>43%</div> <div>34%</div> <div>11%</div> <div>7%</div> </div>
2	B	427	<div> <div>7%</div> <div>11%</div> <div>50%</div> <div>17%</div> <div>5%</div> <div>17%</div> </div>
2	C	427	<div> <div>5%</div> <div>6%</div> <div>54%</div> <div>25%</div> <div>8%</div> <div>7%</div> </div>
2	E	427	<div> <div>7%</div> <div>6%</div> <div>55%</div> <div>19%</div> <div>•</div> <div>17%</div> </div>
2	F	427	<div> <div>5%</div> <div>5%</div> <div>54%</div> <div>25%</div> <div>8%</div> <div>7%</div> </div>

2 Entry composition

There are 2 unique types of molecules in this entry. The entry contains 29480 atoms, of which 0 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 DNA polymerase subunit gamma-1.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	A	1093	Total	C	N	O	S	0	0	0
			8681	5505	1537	1589	50			
1	D	1094	Total	C	N	O	S	0	0	0
			8695	5515	1540	1590	50			

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	354	ASN	-	SEE REMARK 999	UNP P54098
A	355	SER	-	SEE REMARK 999	UNP P54098
D	354	ASN	-	SEE REMARK 999	UNP P54098
D	354A	SER	-	SEE REMARK 999	UNP P54098

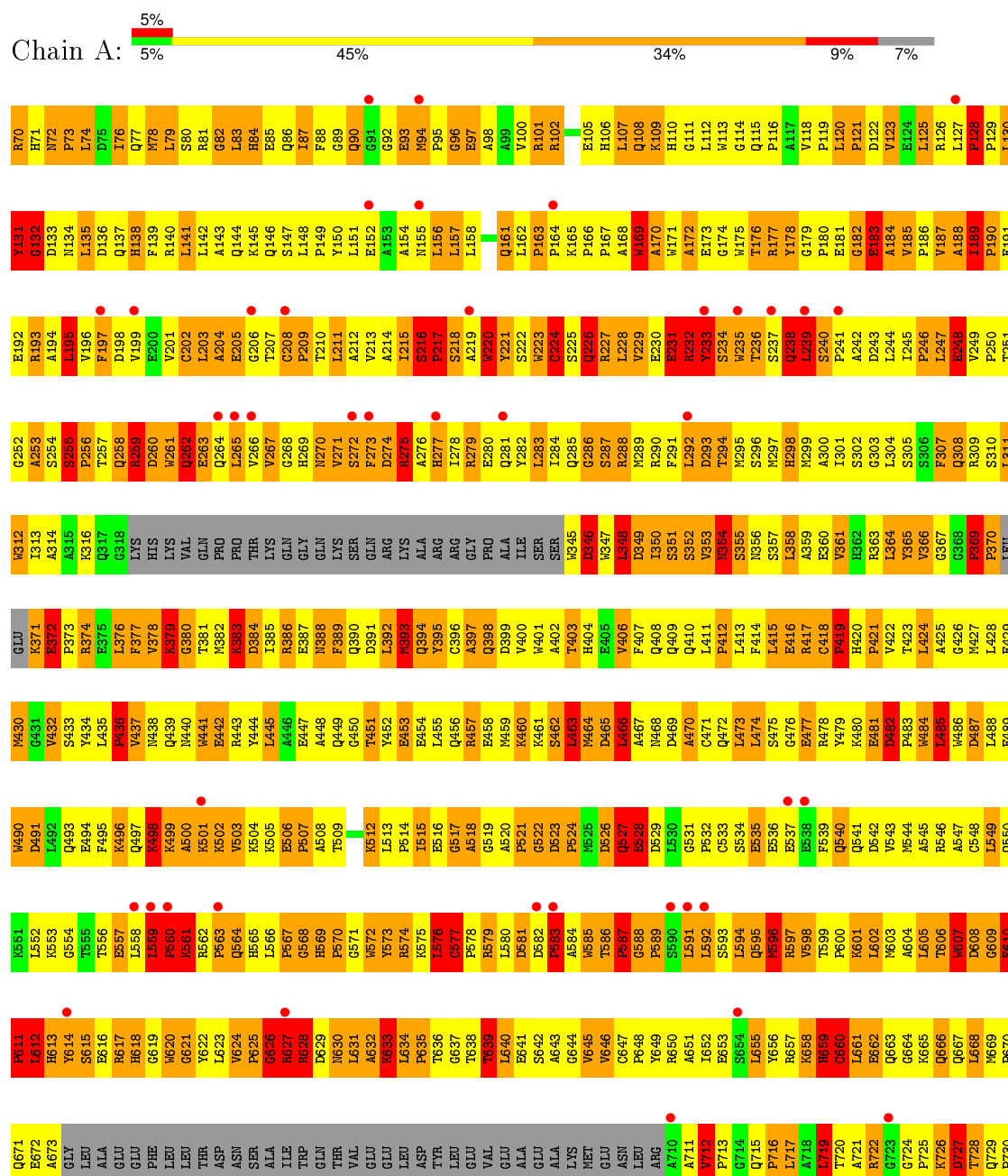
- Molecule 2 is a protein called DNA polymerase subunit gamma-2.

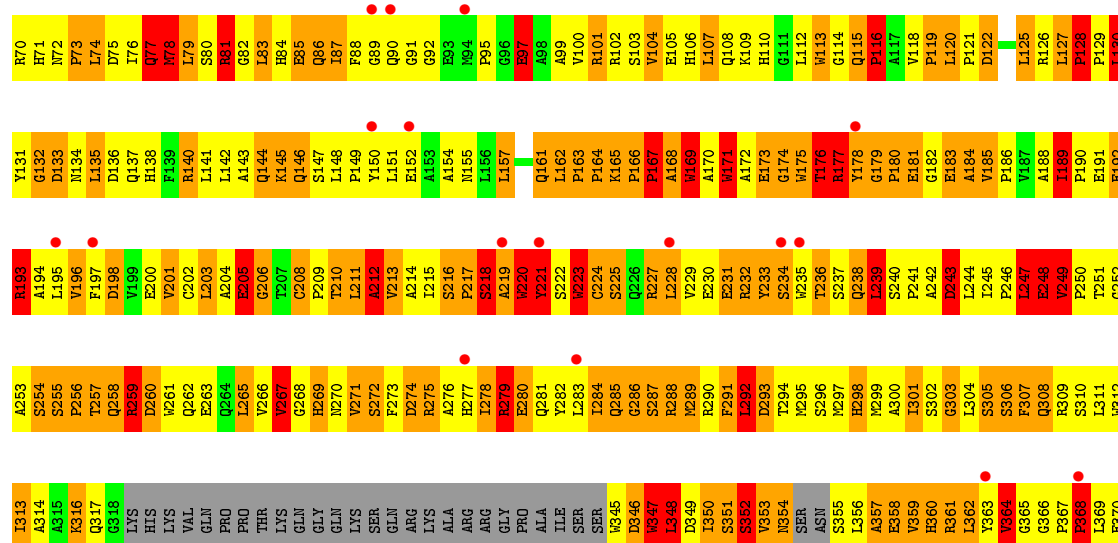
Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	B	355	Total	C	N	O	S	0	0	0
			2871	1840	504	511	16			
2	C	396	Total	C	N	O	S	0	0	0
			3181	2031	563	571	16			
2	E	355	Total	C	N	O	S	0	0	0
			2871	1840	504	511	16			
2	F	396	Total	C	N	O	S	0	0	0
			3181	2031	563	571	16			

3 Residue-property plots

These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of errors 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.

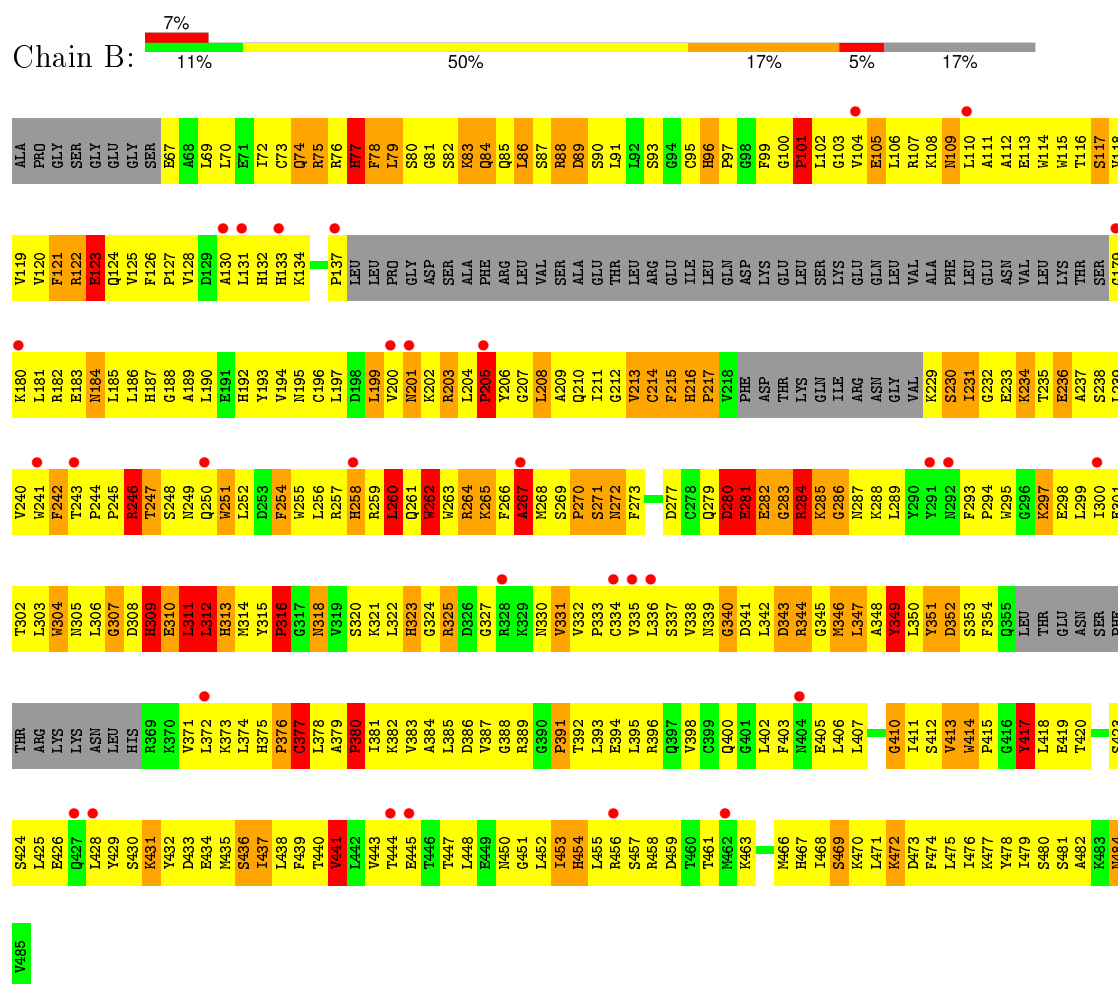
- Molecule 1: DNA polymerase subunit gamma-1



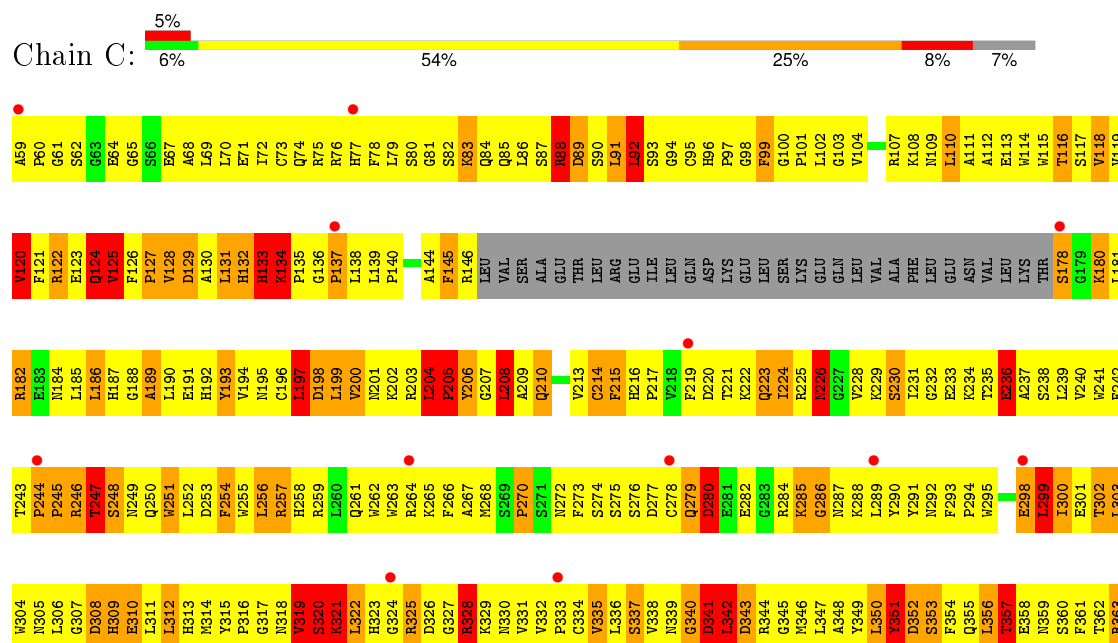


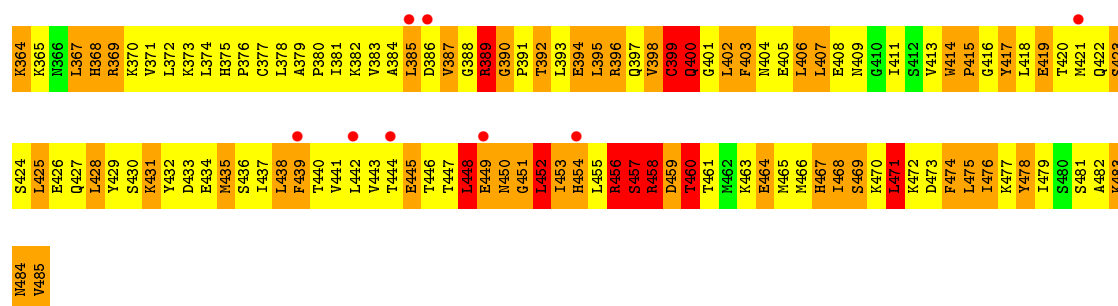
Q1214	Q1154	T1094	R1034	T974	T914	R853	E793	H733	A673	H613	K553	D491	G431	K371
I1155	I1155	S1095	K1035	Q75	A915	A854	I794	H734	GLY	Y614	G554	D492	V432	E372
T1156	T1156	R1096	S1036	Q76	F916	V855	K795	G735	LEU	S615	T595	Q493	S433	E373
M1157	M1157	V1097	Q1037	R977	G917	E856	K796	H736	ALA	R617	T556	E494	E434	R374
L1218	L1158	M1098	K1038	A978	N918	P857	I797	G737	GLU	R618	E557	F495	L435	E375
L1219	L1159	V1099	K1039	A979	N919	T858	I798	P738	GLU	Q496	L558	F496	P436	L376
T1160	T1160	V1100	K1040	R980	T920	R859	S799	T739	PRO	Q497	L559	Q497	V437	F377
Y1220	Y1161	V1101	W1041	R981	L921	L860	F800	N740	LEU	W620	P560	K498	Q438	V378
C1222	C1162	V1102	E1042	A982	G922	S863	W801	D741	LEU	W621	P561	K499	Q439	R379
I1223	S1103	S1103	V1043	Q983	G923	N864	R802	V742	THR	W622	P562	A500	N440	G380
I1224	F1164	S1104	V1044	Q984	G924	N865	N803	D743	ASP	W623	P563	K501	W441	T381
E1225	A1105	A1105	A1045	N985	G925	A865	H804	J744	ASN	W624	P564	K502	E442	K382
Y1166	V1106	V1106	E1046	N986	G926	R866	H805	P745	SER	W625	P565	K503	R443	K383
T1227	K1167	D1107	R1047	A987	A927	P867	K806	G746	ALA	W626	P566	K504	Y444	D384
L1168	L1168	Y1108	A1048	A988	G928	D868	R807	C747	ILE	R627	P567	K505	L445	L385
G1169	G1169	T1109	W1049	R989	T929	E869	I808	W748	TRP	R628	P568	E506	A446	R386
S1230	L1170	H1110	K1050	Q990	D930	V870	S809	F749	GLN	D629	H569	GLN	E447	E387
L1231	G1051	L1111	G1051	G991	L931	G871	S810	P750	THR	N630	P570	T509	A448	N388
D1172	G1052	M1112	G1052	L992	H932	S872	Q811	K751	VAL	L631	G571	A510	Q449	F389
L1173	T1053	L1113	T1053	R993	S933	E873	H812	L752	GLU	A632	H572	S511	Q450	Q390
P1174	V1114	V1114	E1054	R994	H934	K874	H813	P753	GLU	R633	H573	K512	T451	D391
S1235	A1115	A1115	S1055	Y995	T935	K875	W814	H754	LEU	L634	H574	L513	E453	L392
S1175	M1116	M1116	E1056	R996	A936	A876	W815	K755	ASP	P635	H575	P514	E454	K393
V1177	K1117	K1117	M1057	L997	T937	H877	L816	D756	TYR	T636	L576	I515	E454	D394
A1178	W1118	W1118	F1058	S998	T938	V878	P817	G757	LEU	G637	C577	E516	L455	K395
F1179	N1059	L1119	N1059	D999	V938	Q879	R818	H758	GLU	T638	P578	G517	Q456	C396
F1180	K1060	E1120	K1060	E1000	G940	A880	S819	S759	VAL	T639	H579	A518	R457	A397
S1181	L1061	E1121	L1061	G1001	I941	P881	A820	C760	GLU	L640	H580	G519	E458	K398
A1182	E1062	E1122	E1062	E1002	S942	P882	L821	N761	ALA	E641	D582	A520	M459	D399
V1183	S1063	F1123	S1063	W1003	R943	G883	P822	V762	GLU	S642	D583	P521	K460	V400
D1184	I1064	A1124	I1064	L1004	E944	V884	R823	G763	ALA	A643	P583	G522	K461	H401
L1185	I1065	T1125	I1065	V1005	H945	T885	A824	S764	LVS	G644	A584	D623	S462	H402
D1186	T1066	L1126	T1066	R1006	A946	P765	R825	P765	MET	V645	H585	P524	L463	T403
G1127	G1127	G1127	S1067	E1007	K947	V887	I826	F766	GLU	V646	H586	D525	M464	H404
R1188	D1068	L1128	D1068	L1008	I948	G888	R827	A767	ASN	C647	P587	D526	D465	E405
L1189	I1069	F1129	I1069	N1009	F949	A889	H828	K768	LEU	P648	G588	Q527	L466	V406
R1190	P1070	N1010	P1070	D890	N950	D890	P829	D769	R709	Y649	P589	E528	A467	F407
K1191	R1071	P1011	R1071	V881	V951	V881	D830	F770	A710	R650	S590	D529	N468	Q408
E1192	T1072	D1012	T1072	D892	G952	D892	Y831	L771	A711	A651	L591	G530	D469	Q409
V1193	D1013	D1013	S993	R953	R953	S993	D832	P772	V712	L652	L592	P532	A470	Q410
T1194	R1014	R1014	Q894	Q894	I954	Q894	E833	K773	F713	S653	S593	P532	C471	L411
M1195	T1015	V955	E835	E835	G956	L896	G835	N774	G714	S654	L594	G533	Q472	P412
D1196	E1016	E1016	L897	L897	A957	L897	L836	E775	Q715	L655	D595	G533	L473	L413
C1197	CYS	G1017	CYS	G1017	G956	H897	I837	D776	P716	Y656	H596	E536	L474	L414
K1198	CYS	G1018	CYS	G1018	G956	L898	Y837	G777	L717	R657	R597	E537	S475	L415
T1199	Y1139	Y1139	ILE	W1019	Q959	A899	G838	T778	A718	R658	V598	E538	G476	E416
P1200	L1140	L1140	SER	I1020	P960	A900	A839	L779	L719	H659	T599	F539	E477	R417
S1201	V1141	V1141	ARG	S1021	F961	V901	I840	Q780	T720	C660	P600	Q540	R478	C418
N1202	ALA	A1142	ALA	L1022	A962	L902	L841	A781	A721	L661	K601	Q541	Y479	P419
P1203	LEU	E1143	LEU	Q1023	E963	G903	P842	G782	R722	E662	L602	D542	K480	H420
T1204	GLU	D1024	GLU	D1024	R964	D904	Q843	P783	G723	Q663	H603	V543	E481	P421
G1205	P1025	L1025	PRO	L1025	L965	A905	Q843	G784	G724	G664	A604	V544	V422	V422
M1206	R1146	R1146	SER	R1026	L966	H906	W845	G785	P725	K665	L605	A545	P483	T423
E1207	Y1147	Y1147	ALA	K1027	N967	F907	T846	A786	K726	Q666	T606	R546	W484	L424
R1208	R1148	R1148	VAL	V1028	Q968	A908	A847	S787	D728	Q667	H607	A547	L485	A425
R1209	A1149	A1149	GLN	Q1029	P969	G909	G848	T728	T728	L668	D608	C548	W486	A426
Y1210	R1030	R1030	E1090	R1030	N970	N910	T849	P788	Q729	M689	G609	L549	D487	M427
G1211	L1151	L1151	E1091	E1031	H971	H911	I850	P789	P730	P670	F611	Q550	L488	L428
I1212	F1092	L1032	F1092	T1032	R972	G912	T851	A791	S731	Q671	P611	Q551	E489	E429
P1213	M1093	L1153	M1093	A1033	L973	C913	R852	L792	Y732	E672	L612	L552	W490	M430

- Molecule 2: DNA polymerase subunit gamma-2

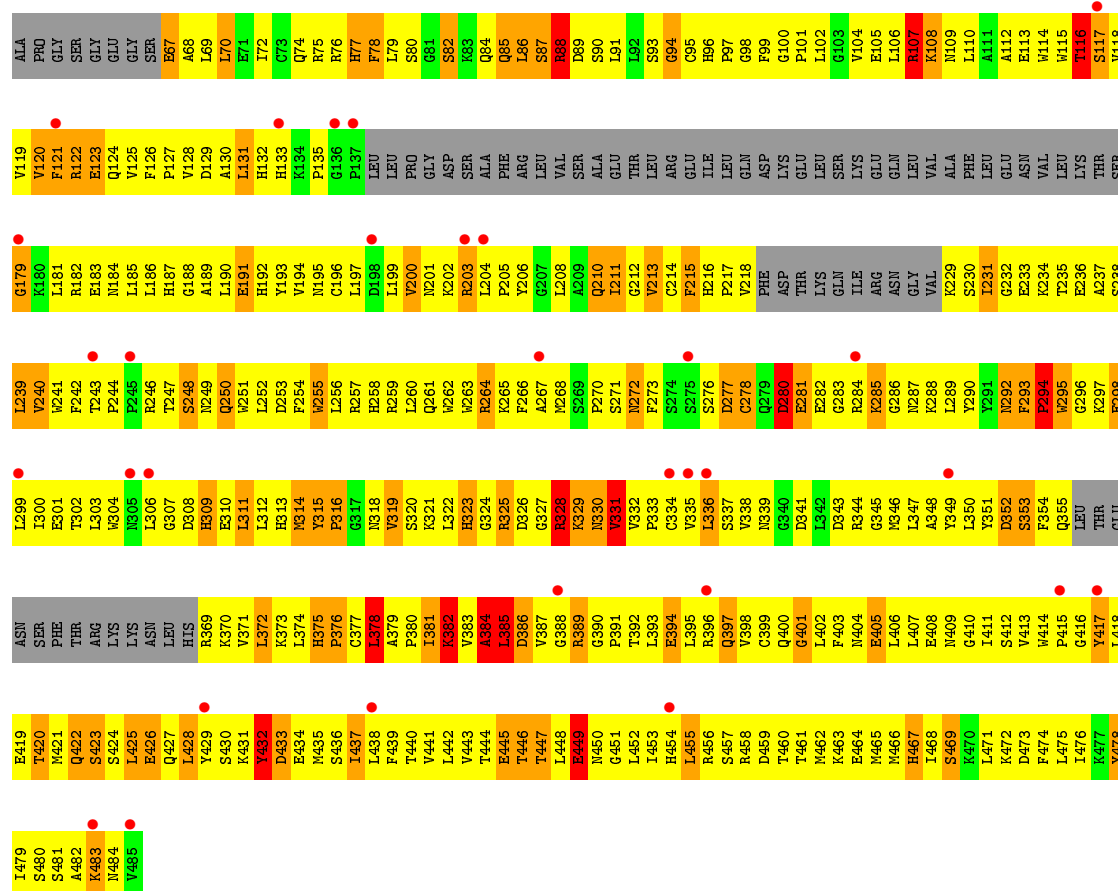


- Molecule 2: DNA polymerase subunit gamma-2

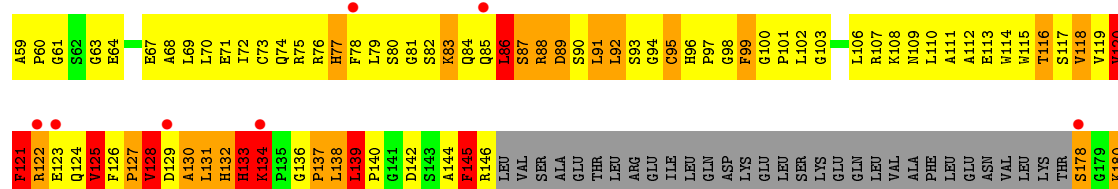




• Molecule 2: DNA polymerase subunit gamma-2



• Molecule 2: DNA polymerase subunit gamma-2



[illegible]

4 Data and refinement statistics

Property	Value	Source
Space group	P 32	Depositor
Cell constants a, b, c, α , β , γ	138.39Å 138.39Å 226.31Å 90.00° 90.00° 120.00°	Depositor
Resolution (Å)	46.92 – 3.24 46.92 – 3.25	Depositor EDS
% Data completeness (in resolution range)	89.5 (46.92-3.24) 89.5 (46.92-3.25)	Depositor EDS
R_{merge}	(Not available)	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	2.12 (at 3.25Å)	Xtriage
Refinement program	CNS 1.2	Depositor
R, R_{free}	0.284 , 0.303 0.362 , (Not available)	Depositor DCC
R_{free} test set	No test flags present.	DCC
Wilson B-factor (Å ²)	93.9	Xtriage
Anisotropy	0.065	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.28 , 64.7	EDS
Estimated twinning fraction	0.008 for -h,-k,l 0.459 for h,-h-k,-l 0.009 for -k,-h,-l	Xtriage
L-test for twinning ²	$\langle L \rangle = 0.52$, $\langle L^2 \rangle = 0.36$	Xtriage
Outliers	0 of 76508 reflections	Xtriage
F_o, F_c correlation	0.83	EDS
Total number of atoms	29480	wwPDB-VP
Average B, all atoms (Å ²)	85.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 2.12% 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, 0.375 respectively for untwinned datasets, and 0.333, 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	10/8913 (0.1%)	1.33	149/12100 (1.2%)
1	D	0.81	18/8927 (0.2%)	1.45	161/12118 (1.3%)
2	B	0.60	2/2944 (0.1%)	0.99	16/3981 (0.4%)
2	C	0.68	0/3262	1.31	47/4411 (1.1%)
2	E	0.67	6/2944 (0.2%)	1.11	23/3981 (0.6%)
2	F	0.64	0/3262	1.28	37/4411 (0.8%)
All	All	0.72	36/30252 (0.1%)	1.31	433/41002 (1.1%)

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	4
1	D	1	3
2	B	0	1
2	C	0	1
2	E	1	2
2	F	0	1
All	All	2	12

All (36) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	D	358	GLU	CG-CD	12.87	1.71	1.51
2	E	385	LEU	CA-CB	-11.49	1.27	1.53
1	D	352	SER	N-CA	9.45	1.65	1.46
1	A	232	ARG	N-CA	-8.85	1.28	1.46
2	E	382	LYS	N-CA	-8.69	1.28	1.46
2	B	280	ASP	C-O	-8.67	1.06	1.23
1	D	352	SER	CA-C	8.29	1.74	1.52
1	D	358	GLU	CB-CG	-8.14	1.36	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	D	249	VAL	CB-CG1	-7.88	1.36	1.52
1	A	232	ARG	CA-CB	-7.53	1.37	1.53
1	A	216	SER	CB-OG	-7.33	1.32	1.42
1	D	361	ARG	CZ-NH1	7.17	1.42	1.33
1	A	232	ARG	CB-CG	7.12	1.71	1.52
1	D	280	GLU	CG-CD	7.10	1.62	1.51
1	D	359	VAL	CB-CG2	-6.86	1.38	1.52
1	D	358	GLU	CD-OE1	6.86	1.33	1.25
1	A	870	VAL	CB-CG1	-6.53	1.39	1.52
1	A	231	GLU	C-O	-6.29	1.11	1.23
1	D	280	GLU	CB-CG	6.28	1.64	1.52
1	D	915	ALA	CA-CB	-6.13	1.39	1.52
1	A	1038	TRP	CB-CG	6.08	1.61	1.50
2	E	382	LYS	CB-CG	-5.93	1.36	1.52
2	E	331	VAL	CA-CB	5.91	1.67	1.54
2	E	382	LYS	CA-CB	5.84	1.66	1.53
1	A	239	LEU	CA-CB	-5.77	1.40	1.53
1	D	870	VAL	CA-CB	-5.77	1.42	1.54
2	B	281	GLU	CG-CD	5.65	1.60	1.51
1	D	248	GLU	N-CA	-5.54	1.35	1.46
2	E	381	ILE	C-O	-5.51	1.12	1.23
1	A	229	VAL	CA-CB	5.42	1.66	1.54
1	A	229	VAL	N-CA	5.40	1.57	1.46
1	D	503	VAL	CA-CB	5.36	1.66	1.54
1	D	870	VAL	CB-CG1	-5.33	1.41	1.52
1	D	361	ARG	CB-CG	5.31	1.66	1.52
1	D	869	ARG	CG-CD	-5.14	1.39	1.51
1	D	359	VAL	C-O	5.09	1.33	1.23

All (433) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	E	382	LYS	N-CA-C	16.56	155.72	111.00
2	E	385	LEU	N-CA-C	16.34	155.11	111.00
1	D	248	GLU	CA-C-N	-14.80	84.64	117.20
2	B	280	ASP	C-N-CA	14.32	157.51	121.70
1	A	1212	ILE	C-N-CD	-12.99	92.03	120.60
1	D	347	TRP	N-CA-C	12.67	145.20	111.00
1	D	171	TRP	N-CA-C	-12.65	76.84	111.00
1	D	721	ALA	N-CA-C	-12.53	77.17	111.00
1	D	810	SER	N-CA-C	-12.41	77.49	111.00
1	D	559	LEU	CA-CB-CG	12.39	143.80	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	1037	GLN	N-CA-C	11.88	143.06	111.00
1	A	379	LYS	N-CA-C	-11.75	79.27	111.00
2	E	382	LYS	N-CA-CB	-11.64	89.64	110.60
1	D	206	GLY	N-CA-C	-11.61	84.07	113.10
2	F	319	VAL	N-CA-C	-11.45	80.09	111.00
2	E	381	ILE	CA-C-N	-11.32	92.29	117.20
1	A	239	LEU	N-CA-C	11.07	140.90	111.00
1	D	627	ARG	N-CA-C	-11.03	81.21	111.00
1	D	358	GLU	OE1-CD-OE2	-11.03	110.07	123.30
1	A	1025	LEU	CA-CB-CG	-11.00	90.01	115.30
1	D	352	SER	O-C-N	-10.95	105.19	122.70
1	D	132	GLY	N-CA-C	10.77	140.03	113.10
2	E	384	ALA	N-CA-C	10.75	140.02	111.00
2	C	448	LEU	CA-CB-CG	10.65	139.80	115.30
1	A	238	GLN	CA-CB-CG	10.65	136.82	113.40
1	A	533	CYS	N-CA-C	-10.62	82.31	111.00
1	A	609	GLY	N-CA-C	-10.62	86.56	113.10
1	D	248	GLU	O-C-N	10.50	139.50	122.70
2	F	286	GLY	N-CA-C	-10.46	86.95	113.10
1	A	274	ASP	N-CA-C	-10.45	82.79	111.00
1	D	737	GLY	N-CA-C	-10.43	87.03	113.10
1	D	1069	ILE	C-N-CD	-10.22	98.12	120.60
2	F	208	LEU	CA-CB-CG	-10.11	92.05	115.30
1	D	274	ASP	N-CA-C	-10.05	83.85	111.00
2	C	460	THR	N-CA-C	-9.87	84.35	111.00
2	B	340	GLY	N-CA-C	-9.84	88.49	113.10
2	C	319	VAL	N-CA-C	-9.72	84.77	111.00
1	D	351	SER	N-CA-C	-9.71	84.80	111.00
1	D	736	ASN	N-CA-C	9.60	136.92	111.00
2	E	78	PHE	N-CA-C	-9.54	85.25	111.00
2	E	385	LEU	N-CA-CB	-9.49	91.41	110.40
1	A	1205	GLY	N-CA-C	-9.49	89.37	113.10
1	D	1196	ASP	N-CA-C	-9.49	85.38	111.00
2	E	381	ILE	C-N-CA	9.48	145.40	121.70
1	D	672	GLU	N-CA-C	-9.47	85.44	111.00
1	D	786	ALA	N-CA-C	-9.45	85.49	111.00
2	F	357	THR	N-CA-C	-9.38	85.69	111.00
2	C	208	LEU	CA-CB-CG	-9.27	93.97	115.30
1	D	167	PRO	N-CA-C	-9.25	88.06	112.10
2	F	121	PHE	N-CA-C	9.14	135.67	111.00
2	C	450	ASN	N-CA-C	-9.12	86.37	111.00
1	A	1010	LEU	CA-CB-CG	-9.08	94.41	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	C	303	LEU	N-CA-C	9.05	135.44	111.00
1	D	645	VAL	N-CA-C	8.99	135.27	111.00
1	D	115	GLN	C-N-CD	-8.98	100.84	120.60
1	D	178	TYR	N-CA-C	-8.97	86.78	111.00
2	C	342	LEU	N-CA-C	-8.96	86.81	111.00
1	D	726	LYS	N-CA-C	8.91	135.05	111.00
2	C	454	HIS	N-CA-C	-8.86	87.09	111.00
1	A	627	ARG	N-CA-C	-8.85	87.11	111.00
1	A	523	ASP	N-CA-C	-8.84	87.13	111.00
2	C	279	GLN	N-CA-C	8.80	134.76	111.00
1	A	238	GLN	N-CA-C	8.77	134.68	111.00
1	D	615	SER	N-CA-C	-8.74	87.41	111.00
1	D	346	ASP	C-N-CA	-8.71	99.93	121.70
2	C	399	CYS	N-CA-C	-8.66	87.62	111.00
2	F	460	THR	N-CA-C	-8.65	87.64	111.00
1	A	923	GLY	N-CA-C	-8.51	91.83	113.10
1	D	787	SER	N-CA-C	-8.48	88.11	111.00
1	D	879	GLN	N-CA-C	8.47	133.87	111.00
1	A	1044	VAL	N-CA-C	-8.43	88.23	111.00
1	A	178	TYR	N-CA-C	-8.39	88.35	111.00
2	C	452	LEU	CA-CB-CG	8.38	134.57	115.30
1	A	938	THR	N-CA-C	-8.31	88.55	111.00
1	A	1212	ILE	N-CA-C	8.29	133.37	111.00
2	F	471	LEU	CA-CB-CG	-8.28	96.27	115.30
2	C	390	GLY	N-CA-C	8.25	133.72	113.10
2	C	456	ARG	N-CA-C	8.24	133.26	111.00
1	D	360	HIS	N-CA-C	-8.21	88.84	111.00
1	D	523	ASP	N-CA-C	-8.20	88.85	111.00
1	D	279	ARG	N-CA-C	8.17	133.07	111.00
1	D	176	THR	N-CA-C	8.15	133.02	111.00
1	D	533	CYS	N-CA-C	-8.15	88.98	111.00
1	D	1010	LEU	CA-CB-CG	-8.15	96.55	115.30
2	C	357	THR	N-CA-C	-8.15	89.00	111.00
2	F	320	SER	N-CA-C	8.15	133.00	111.00
1	D	1027	LYS	N-CA-C	8.12	132.91	111.00
1	D	735	GLY	N-CA-C	8.10	133.35	113.10
2	F	342	LEU	N-CA-C	-8.09	89.16	111.00
1	D	1233	LYS	N-CA-C	-8.08	89.19	111.00
1	D	576	LEU	CA-CB-CG	8.06	133.85	115.30
2	F	390	GLY	N-CA-C	8.02	133.14	113.10
1	D	606	THR	N-CA-C	7.99	132.57	111.00
1	A	79	LEU	N-CA-C	-7.97	89.48	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	1069	ILE	N-CA-C	-7.95	89.53	111.00
2	F	279	GLN	N-CA-C	7.93	132.42	111.00
2	F	339	ASN	N-CA-C	-7.89	89.69	111.00
1	A	615	SER	N-CA-C	-7.88	89.72	111.00
1	D	243	ASP	N-CA-C	-7.86	89.77	111.00
1	D	912	GLY	N-CA-C	7.78	132.54	113.10
1	A	915	ALA	N-CA-C	-7.77	90.02	111.00
1	A	576	LEU	CA-CB-CG	7.76	133.16	115.30
1	A	188	ALA	N-CA-C	7.76	131.95	111.00
1	A	1127	GLY	N-CA-C	7.75	132.47	113.10
1	A	1210	TYR	N-CA-C	7.68	131.74	111.00
1	D	286	GLY	N-CA-C	-7.66	93.94	113.10
2	B	286	GLY	N-CA-C	-7.65	93.98	113.10
1	A	782	GLY	N-CA-C	-7.63	94.01	113.10
1	A	204	ALA	N-CA-C	-7.62	90.42	111.00
1	D	1038	TRP	N-CA-C	7.61	131.55	111.00
1	A	605	LEU	N-CA-C	7.61	131.53	111.00
1	D	232	ARG	N-CA-C	7.59	131.50	111.00
2	F	133	HIS	N-CA-C	7.58	131.47	111.00
1	A	537	GLU	N-CA-C	7.57	131.44	111.00
1	D	359	VAL	CB-CA-C	-7.56	97.03	111.40
1	A	182	GLY	N-CA-C	7.55	131.98	113.10
1	A	238	GLN	O-C-N	7.53	134.75	122.70
1	A	741	ASP	N-CA-C	-7.52	90.69	111.00
1	A	485	LEU	CA-CB-CG	7.49	132.53	115.30
1	D	249	VAL	N-CA-C	7.49	131.23	111.00
1	A	132	GLY	N-CA-C	7.48	131.81	113.10
1	D	922	GLN	N-CA-C	7.48	131.21	111.00
1	A	912	GLY	N-CA-C	7.46	131.76	113.10
1	D	177	ARG	N-CA-C	7.45	131.12	111.00
2	C	343	ASP	N-CA-C	-7.45	90.89	111.00
1	D	869	ARG	NE-CZ-NH1	-7.45	116.58	120.30
1	D	1236	GLN	N-CA-C	7.42	131.05	111.00
2	E	381	ILE	N-CA-C	-7.42	90.98	111.00
1	D	669	MET	N-CA-C	-7.41	90.98	111.00
1	D	728	THR	N-CA-C	-7.39	91.04	111.00
1	A	203	LEU	CA-CB-CG	-7.37	98.34	115.30
1	D	180	PRO	N-CA-C	-7.37	92.94	112.10
1	D	231	GLU	N-CA-C	7.37	130.90	111.00
1	D	1130	CYS	N-CA-C	7.33	130.79	111.00
1	A	872	SER	N-CA-C	-7.33	91.21	111.00
2	C	459	ASP	N-CA-C	7.33	130.79	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	602	LEU	N-CA-C	-7.32	91.24	111.00
1	D	566	LEU	N-CA-C	-7.30	91.28	111.00
2	E	381	ILE	O-C-N	7.29	134.36	122.70
1	D	568	GLY	N-CA-C	-7.28	94.89	113.10
2	E	433	ASP	N-CA-C	-7.27	91.37	111.00
1	A	195	LEU	CA-CB-CG	7.26	132.01	115.30
1	D	644	GLY	N-CA-C	-7.26	94.94	113.10
1	D	75	ASP	N-CA-C	-7.26	91.39	111.00
1	A	735	GLY	N-CA-C	7.23	131.18	113.10
1	A	996	ARG	N-CA-C	7.19	130.42	111.00
2	C	204	LEU	N-CA-C	7.18	130.39	111.00
2	F	196	CYS	CA-CB-SG	7.18	126.92	114.00
2	B	281	GLU	CA-C-N	-7.18	101.41	117.20
1	A	1207	GLU	N-CA-C	-7.11	91.80	111.00
1	A	1208	ARG	N-CA-C	-7.08	91.88	111.00
1	D	233	TYR	N-CA-C	7.07	130.10	111.00
2	F	127	PRO	N-CA-C	-7.07	93.72	112.10
2	B	280	ASP	CA-C-N	-7.05	101.69	117.20
1	D	1015	THR	N-CA-C	7.05	130.03	111.00
1	D	357	ALA	C-N-CA	-7.05	104.08	121.70
1	A	1236	GLN	N-CA-C	7.01	129.93	111.00
2	C	471	LEU	CA-CB-CG	-6.97	99.26	115.30
1	D	1237	PRO	N-CA-C	6.97	130.23	112.10
1	D	768	LYS	N-CA-C	-6.97	92.18	111.00
2	F	343	ASP	N-CA-C	-6.93	92.29	111.00
1	A	1211	GLY	N-CA-C	-6.93	95.78	113.10
1	D	991	GLY	N-CA-C	6.92	130.41	113.10
1	D	891	VAL	CA-CB-CG1	-6.91	100.53	110.90
2	C	320	SER	N-CA-C	6.90	129.62	111.00
1	A	759	SER	N-CA-C	6.88	129.57	111.00
2	F	457	SER	N-CA-C	6.87	129.55	111.00
1	D	633	LYS	N-CA-C	6.86	129.53	111.00
1	A	122	ASP	N-CA-C	6.83	129.43	111.00
2	C	318	ASN	N-CA-C	-6.82	92.60	111.00
1	A	1031	GLU	N-CA-C	-6.81	92.62	111.00
1	A	258	GLN	N-CA-C	-6.80	92.64	111.00
1	D	710	ALA	N-CA-C	6.79	129.35	111.00
1	D	846	THR	N-CA-C	-6.79	92.68	111.00
2	E	384	ALA	CA-C-N	-6.76	102.33	117.20
2	F	318	ASN	N-CA-C	-6.76	92.75	111.00
2	C	286	GLY	N-CA-C	-6.74	96.26	113.10
1	D	1180	PHE	N-CA-C	6.74	129.19	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	259	ARG	N-CA-C	6.73	129.18	111.00
1	A	1232	GLU	N-CA-C	-6.73	92.82	111.00
1	A	277	HIS	N-CA-C	-6.72	92.86	111.00
1	A	231	GLU	N-CA-C	6.72	129.13	111.00
2	F	318	ASN	C-N-CA	6.71	138.48	121.70
1	D	354	ASN	N-CA-C	6.69	129.07	111.00
2	C	133	HIS	N-CA-C	6.68	129.05	111.00
1	D	1070	PRO	CA-N-CD	-6.68	102.14	111.50
2	F	139	LEU	N-CA-C	6.68	129.03	111.00
1	A	92	GLY	N-CA-C	-6.68	96.41	113.10
2	C	457	SER	N-CA-C	6.66	128.99	111.00
1	D	516	GLU	N-CA-C	6.66	128.99	111.00
2	F	134	LYS	N-CA-C	6.65	128.94	111.00
2	F	247	THR	N-CA-C	-6.65	93.06	111.00
2	C	178	SER	N-CA-C	6.63	128.89	111.00
1	D	248	GLU	C-N-CA	6.63	138.27	121.70
2	E	280	ASP	N-CA-C	6.61	128.84	111.00
1	D	938	THR	N-CA-C	-6.60	93.17	111.00
1	D	759	SER	N-CA-C	6.56	128.70	111.00
1	A	1206	MET	N-CA-C	6.55	128.70	111.00
1	A	419	PRO	N-CA-C	6.54	129.11	112.10
2	F	120	VAL	N-CA-C	6.54	128.65	111.00
1	D	272	SER	N-CA-C	-6.48	93.49	111.00
2	B	281	GLU	O-C-N	6.47	133.06	122.70
2	C	127	PRO	N-CA-C	-6.46	95.29	112.10
2	F	178	SER	N-CA-C	6.46	128.44	111.00
2	C	247	THR	N-CA-C	-6.46	93.57	111.00
2	C	459	ASP	CA-C-N	-6.45	103.01	117.20
1	D	565	HIS	N-CA-C	-6.44	93.60	111.00
1	D	351	SER	O-C-N	6.43	132.99	122.70
2	C	335	VAL	CB-CA-C	-6.42	99.21	111.40
2	F	452	LEU	N-CA-C	-6.41	93.69	111.00
1	D	193	ARG	N-CA-C	-6.39	93.73	111.00
1	A	82	GLY	N-CA-C	-6.36	97.19	113.10
1	A	238	GLN	C-N-CA	6.34	137.55	121.70
1	A	346	ASP	N-CA-C	6.34	128.12	111.00
1	A	349	ASP	N-CA-C	-6.34	93.88	111.00
2	F	299	LEU	N-CA-C	6.33	128.10	111.00
1	A	1206	MET	CA-CB-CG	-6.33	102.54	113.30
1	A	1174	PRO	N-CA-C	6.32	128.54	112.10
1	D	275	ARG	N-CA-C	-6.32	93.92	111.00
2	C	428	LEU	CA-CB-CG	-6.32	100.76	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	291	PHE	N-CA-C	6.31	128.05	111.00
1	D	566	LEU	CA-CB-CG	6.31	129.81	115.30
1	A	658	LYS	N-CA-C	-6.31	93.97	111.00
1	A	233	TYR	N-CA-C	6.30	128.02	111.00
2	B	309	HIS	N-CA-C	6.30	128.00	111.00
1	D	116	PRO	N-CA-C	-6.27	95.79	112.10
1	A	1024	ASP	N-CA-C	6.27	127.92	111.00
1	A	72	ASN	N-CA-C	-6.26	94.11	111.00
1	A	956	GLY	N-CA-C	6.25	128.72	113.10
1	A	1198	LYS	N-CA-C	-6.23	94.18	111.00
1	A	1209	ARG	N-CA-C	-6.17	94.33	111.00
1	D	1008	LEU	CA-CB-CG	6.16	129.47	115.30
1	A	825	VAL	N-CA-C	-6.16	94.37	111.00
1	D	579	ARG	N-CA-C	-6.16	94.37	111.00
2	C	299	LEU	N-CA-C	6.14	127.59	111.00
1	A	1011	PRO	N-CA-C	-6.14	96.15	112.10
1	D	164	PRO	N-CA-C	-6.13	96.15	112.10
2	C	134	LYS	N-CA-C	6.13	127.56	111.00
1	D	1179	PHE	N-CA-C	6.13	127.55	111.00
2	E	381	ILE	CA-C-O	6.12	132.96	120.10
1	A	256	PRO	N-CA-C	-6.11	96.22	112.10
1	A	466	LEU	CA-CB-CG	6.09	129.30	115.30
1	A	392	LEU	N-CA-C	6.08	127.42	111.00
1	A	951	TYR	N-CA-C	-6.07	94.61	111.00
1	D	532	PRO	N-CA-C	6.06	127.86	112.10
1	D	744	ILE	N-CA-C	6.05	127.33	111.00
1	D	364	VAL	N-CA-C	6.03	127.28	111.00
1	A	632	ALA	N-CA-C	-6.02	94.75	111.00
1	A	846	THR	N-CA-C	-6.02	94.75	111.00
1	A	1026	ARG	N-CA-CB	-6.02	99.76	110.60
1	D	192	GLU	N-CA-C	6.01	127.24	111.00
1	A	130	LEU	N-CA-C	5.98	127.14	111.00
1	A	131	TYR	N-CA-C	5.97	127.13	111.00
2	E	281	GLU	N-CA-C	-5.97	94.88	111.00
1	D	648	PRO	N-CA-C	-5.96	96.61	112.10
2	F	450	ASN	N-CA-C	-5.93	94.98	111.00
1	D	348	LEU	CA-CB-CG	5.92	128.93	115.30
1	A	1030	ARG	N-CA-C	-5.91	95.05	111.00
1	A	518	ALA	N-CA-C	5.90	126.94	111.00
1	A	226	GLN	N-CA-C	-5.90	95.07	111.00
2	E	107	ARG	NE-CZ-NH1	5.89	123.25	120.30
1	D	346	ASP	CB-CA-C	5.89	122.17	110.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	220	TRP	N-CA-C	5.88	126.89	111.00
1	A	383	LYS	N-CA-C	-5.87	95.14	111.00
1	D	734	HIS	N-CA-C	-5.87	95.16	111.00
1	D	821	LEU	C-N-CD	-5.87	107.70	120.60
1	A	121	PRO	N-CA-C	5.85	127.31	112.10
1	A	966	LEU	CA-CB-CG	5.85	128.75	115.30
2	C	236	GLU	N-CA-C	5.85	126.79	111.00
2	E	385	LEU	C-N-CA	-5.84	107.09	121.70
1	A	275	ARG	N-CA-C	-5.84	95.24	111.00
1	D	537	GLU	N-CA-C	5.84	126.76	111.00
1	D	671	GLN	C-N-CA	5.84	136.29	121.70
1	A	1171	ASN	N-CA-C	5.83	126.75	111.00
2	C	400	GLN	N-CA-C	-5.82	95.29	111.00
1	A	1125	ILE	N-CA-C	5.82	126.70	111.00
1	D	418	CYS	C-N-CD	-5.81	107.82	120.60
1	A	209	PRO	N-CA-C	5.81	127.20	112.10
1	A	108	GLN	N-CA-C	-5.80	95.33	111.00
2	B	260	LEU	N-CA-C	-5.80	95.35	111.00
1	A	628	ARG	N-CA-C	-5.79	95.38	111.00
2	C	458	ARG	NE-CZ-NH2	-5.79	117.41	120.30
2	E	179	GLY	N-CA-C	-5.79	98.64	113.10
2	F	467	HIS	N-CA-C	-5.79	95.38	111.00
1	A	418	CYS	C-N-CD	-5.77	107.90	120.60
2	B	267	ALA	N-CA-C	5.76	126.56	111.00
1	D	1020	ILE	N-CA-C	5.75	126.52	111.00
1	A	238	GLN	CB-CA-C	-5.74	98.92	110.40
2	F	236	GLU	N-CA-C	5.72	126.45	111.00
2	C	139	LEU	N-CA-C	5.72	126.44	111.00
1	D	77	GLN	N-CA-C	5.71	126.42	111.00
1	A	348	LEU	N-CA-C	-5.70	95.61	111.00
1	A	1218	LEU	N-CA-C	5.70	126.39	111.00
1	A	498	LYS	N-CA-C	5.69	126.37	111.00
2	C	302	THR	N-CA-C	5.68	126.34	111.00
1	A	784	GLY	N-CA-C	5.67	127.29	113.10
2	E	386	ASP	N-CA-C	5.67	126.31	111.00
1	A	1204	THR	N-CA-C	5.67	126.31	111.00
1	D	874	LEU	N-CA-C	5.65	126.25	111.00
1	D	999	ASP	N-CA-CB	-5.64	100.44	110.60
1	A	1213	PRO	CA-N-CD	-5.63	103.61	111.50
1	D	1067	SER	N-CA-C	-5.63	95.81	111.00
1	A	660	CYS	N-CA-C	5.62	126.19	111.00
1	A	130	LEU	CA-CB-CG	-5.62	102.38	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	872	SER	C-N-CA	5.60	135.70	121.70
1	D	280	GLU	OE1-CD-OE2	-5.60	116.58	123.30
1	D	841	LEU	CA-CB-CG	5.60	128.17	115.30
1	D	735	GLY	C-N-CA	5.59	135.69	121.70
2	C	469	SER	N-CA-C	5.59	126.09	111.00
1	D	651	ALA	N-CA-C	-5.59	95.92	111.00
2	B	311	LEU	N-CA-C	-5.58	95.93	111.00
1	D	616	GLU	N-CA-C	5.58	126.07	111.00
1	A	1067	SER	N-CA-C	5.58	126.06	111.00
1	A	737	GLY	N-CA-C	5.58	127.04	113.10
1	D	189	ILE	N-CA-C	-5.57	95.96	111.00
1	A	560	PRO	N-CA-C	5.56	126.56	112.10
1	A	1158	LEU	CA-CB-CG	-5.54	102.56	115.30
1	A	996	ARG	NE-CZ-NH1	-5.52	117.54	120.30
1	D	1179	PHE	C-N-CA	5.52	135.51	121.70
1	D	352	SER	CA-C-O	5.52	131.70	120.10
1	D	510	ALA	N-CA-C	-5.52	96.09	111.00
1	D	581	ASP	N-CA-C	5.52	125.91	111.00
1	A	1214	GLN	N-CA-C	5.52	125.90	111.00
2	B	121	PHE	N-CA-C	5.51	125.89	111.00
1	A	180	PRO	N-CA-C	-5.51	97.77	112.10
1	D	351	SER	CA-C-O	-5.50	108.54	120.10
1	A	610	PHE	C-N-CA	-5.49	98.92	122.00
1	D	605	LEU	C-N-CA	-5.49	107.97	121.70
1	A	255	SER	N-CA-C	5.48	125.79	111.00
1	A	79	LEU	CB-CG-CD2	-5.47	101.70	111.00
2	C	341	ASP	N-CA-C	-5.47	96.23	111.00
2	F	280	ASP	N-CA-C	5.47	125.77	111.00
2	F	445	GLU	N-CA-C	5.47	125.77	111.00
1	A	286	GLY	N-CA-C	-5.46	99.46	113.10
1	A	907	PHE	N-CA-C	-5.45	96.28	111.00
1	A	218	SER	N-CA-CB	-5.44	102.33	110.50
1	D	212	ALA	N-CA-C	-5.44	96.31	111.00
1	A	172	ALA	N-CA-C	5.43	125.66	111.00
1	A	581	ASP	N-CA-C	5.42	125.64	111.00
2	C	279	GLN	CA-C-N	-5.42	105.26	117.20
2	F	86	LEU	CB-CG-CD2	-5.42	101.79	111.00
1	D	1142	ARG	N-CA-C	-5.40	96.41	111.00
1	A	1070	PRO	CA-N-CD	-5.40	103.94	111.50
1	D	221	TYR	N-CA-C	5.40	125.57	111.00
1	D	1171	ASN	N-CA-C	5.40	125.57	111.00
2	B	310	GLU	N-CA-C	-5.39	96.44	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	515	ILE	N-CA-C	5.38	125.53	111.00
2	C	129	ASP	N-CA-C	5.38	125.52	111.00
1	D	1177	VAL	N-CA-C	-5.38	96.48	111.00
1	D	775	GLU	N-CA-C	-5.37	96.50	111.00
2	F	367	LEU	N-CA-C	-5.37	96.51	111.00
1	A	238	GLN	CA-C-N	-5.36	105.40	117.20
1	A	74	LEU	C-N-CA	-5.36	108.30	121.70
1	D	247	LEU	CB-CG-CD1	-5.36	101.90	111.00
2	F	133	HIS	C-N-CA	5.34	135.05	121.70
2	F	355	GLN	N-CA-C	-5.34	96.58	111.00
1	D	71	HIS	N-CA-C	5.33	125.40	111.00
1	A	232	ARG	C-N-CA	5.33	135.03	121.70
1	D	1131	ILE	N-CA-C	5.33	125.38	111.00
1	A	919	MET	N-CA-C	-5.31	96.67	111.00
1	A	521	PRO	N-CA-C	5.30	125.88	112.10
1	D	1179	PHE	CA-C-N	-5.29	105.55	117.20
1	D	919	MET	N-CA-C	-5.29	96.71	111.00
1	D	257	THR	N-CA-C	5.29	125.29	111.00
1	A	873	GLU	N-CA-C	5.28	125.26	111.00
1	A	574	ARG	N-CA-C	5.28	125.25	111.00
1	D	528	GLU	N-CA-C	5.28	125.24	111.00
1	A	626	GLY	N-CA-C	5.27	126.28	113.10
2	F	459	ASP	N-CA-C	5.27	125.22	111.00
1	D	74	LEU	N-CA-C	5.26	125.21	111.00
1	A	1193	VAL	N-CA-C	5.26	125.20	111.00
1	D	530	LEU	CA-CB-CG	5.26	127.39	115.30
1	D	1072	THR	N-CA-C	5.25	125.17	111.00
1	D	353	VAL	N-CA-CB	-5.25	99.96	111.50
2	C	133	HIS	CA-C-N	-5.24	105.67	117.20
1	D	128	PRO	N-CA-C	5.23	125.71	112.10
1	D	239	LEU	N-CA-C	-5.23	96.87	111.00
1	D	198	ASP	N-CA-C	-5.23	96.87	111.00
1	A	563	PRO	N-CA-C	5.23	125.69	112.10
1	A	522	GLY	N-CA-C	5.22	126.16	113.10
1	A	1001	GLY	N-CA-C	-5.22	100.04	113.10
1	D	657	ARG	N-CA-C	5.21	125.08	111.00
2	E	325	ARG	N-CA-C	5.21	125.08	111.00
1	A	183	GLU	N-CA-C	5.21	125.06	111.00
1	A	74	LEU	CA-CB-CG	5.21	127.28	115.30
1	A	1210	TYR	CA-C-N	-5.20	105.80	116.20
1	D	881	PRO	C-N-CD	-5.20	109.17	120.60
2	C	92	LEU	CA-CB-CG	-5.20	103.35	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	498	LYS	N-CA-C	5.20	125.03	111.00
2	E	432	TYR	CA-CB-CG	5.20	123.27	113.40
2	E	378	LEU	CB-CG-CD2	-5.19	102.17	111.00
1	D	727	ASP	N-CA-C	-5.19	96.98	111.00
1	A	757	GLY	N-CA-C	-5.19	100.13	113.10
2	C	133	HIS	C-N-CA	5.18	134.66	121.70
2	F	449	GLU	N-CA-C	5.18	124.99	111.00
1	A	606	THR	CB-CA-C	-5.18	97.62	111.60
1	A	611	PRO	N-CA-C	5.18	125.56	112.10
1	D	1210	TYR	N-CA-C	-5.17	97.06	111.00
1	A	1142	ARG	N-CA-C	-5.16	97.08	111.00
2	E	381	ILE	CB-CA-C	5.15	121.90	111.60
1	A	1201	SER	N-CA-C	-5.15	97.09	111.00
1	D	130	LEU	CA-CB-CG	-5.15	103.46	115.30
2	B	215	PHE	N-CA-C	5.15	124.89	111.00
1	D	1132	SER	N-CA-C	-5.14	97.11	111.00
1	A	1231	LEU	N-CA-C	5.13	124.86	111.00
1	D	612	LEU	CB-CG-CD2	-5.13	102.28	111.00
1	D	1021	SER	N-CA-C	-5.12	97.18	111.00
1	A	612	LEU	N-CA-C	-5.12	97.18	111.00
1	D	292	LEU	CA-CB-CG	-5.11	103.54	115.30
2	C	458	ARG	NE-CZ-NH1	5.11	122.86	120.30
2	B	280	ASP	N-CA-C	5.11	124.79	111.00
2	C	467	HIS	N-CA-C	-5.09	97.26	111.00
1	D	1013	ASP	N-CA-C	5.08	124.73	111.00
1	D	411	LEU	N-CA-C	5.08	124.71	111.00
1	D	515	ILE	CB-CA-C	-5.07	101.46	111.60
1	A	187	VAL	N-CA-C	-5.06	97.33	111.00
2	C	197	LEU	CA-CB-CG	5.06	126.94	115.30
1	D	192	GLU	CA-C-N	-5.06	106.06	117.20
1	D	622	TYR	N-CA-C	5.06	124.67	111.00
1	D	992	LEU	CA-CB-CG	5.06	126.94	115.30
1	A	370	PRO	N-CA-C	-5.06	98.95	112.10
1	D	609	GLY	N-CA-C	-5.06	100.46	113.10
1	A	231	GLU	CA-C-N	-5.05	106.08	117.20
1	A	528	GLU	N-CA-C	5.03	124.58	111.00
1	A	189	ILE	CB-CA-C	-5.03	101.55	111.60
2	B	281	GLU	N-CA-C	5.03	124.57	111.00
1	D	1031	GLU	N-CA-C	-5.02	97.44	111.00
2	B	271	SER	N-CA-C	-5.02	97.44	111.00
1	A	224	CYS	CB-CA-C	-5.02	100.36	110.40
1	A	229	VAL	CA-CB-CG2	-5.01	103.38	110.90

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Mol	Chain	Res	Type	Atoms	Z	Observed($^{\circ}$)	Ideal($^{\circ}$)
1	D	1073	PRO	N-CA-C	5.01	125.13	112.10
2	C	138	LEU	N-CA-C	5.01	124.53	111.00
1	D	346	ASP	CB-CG-OD2	5.00	122.80	118.30

All (2) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
1	D	249	VAL	CA
2	E	385	LEU	CA

All (12) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	1179	PHE	Sidechain
1	A	221	TYR	Sidechain
1	A	239	LEU	Mainchain
1	A	395	TYR	Sidechain
2	B	417	TYR	Sidechain
2	C	351	TYR	Sidechain
1	D	368	PRO	Mainchain
1	D	656	TYR	Sidechain
1	D	995	TYR	Sidechain
2	E	432	TYR	Sidechain
2	E	478	TYR	Sidechain
2	F	206	TYR	Sidechain

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	8681	0	8521	3592	8
1	D	8695	0	8540	3900	7
2	B	2871	0	2862	809	0
2	C	3181	0	3168	1419	1
2	E	2871	0	2862	945	0
2	F	3181	0	3168	1467	4
All	All	29480	0	29121	11814	10

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 202.

All (11814) 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:D:352:SER:C	1:D:352:SER:CA	1.74	1.55
1:D:460:LYS:HE2	1:D:460:LYS:N	1.20	1.48
2:E:432:TYR:HB2	2:E:437:ILE:CD1	1.51	1.39
1:D:352:SER:CA	1:D:358:GLU:OE2	1.73	1.36
1:D:914:THR:O	1:D:918:TRP:HB3	1.24	1.35
1:D:352:SER:N	1:D:358:GLU:OE2	1.58	1.34
1:D:995:TYR:CE1	1:D:1004:LEU:HB3	1.66	1.31
2:C:356:LEU:HB3	2:C:359:ASN:ND2	1.44	1.30
1:D:460:LYS:H	1:D:460:LYS:CE	1.44	1.28
1:A:513:LEU:HB2	1:A:568:GLY:CA	1.64	1.28
1:A:1210:TYR:CE1	1:A:1213:PRO:HG2	1.68	1.28
2:F:241:TRP:HB2	2:F:336:LEU:CD1	1.64	1.27
1:A:784:GLY:HA2	2:C:362:THR:OG1	1.29	1.26
1:D:275:ARG:HB2	1:D:843:GLN:CB	1.64	1.25
1:D:275:ARG:NE	1:D:843:GLN:H	1.31	1.25
1:A:669:MET:O	1:A:673:ALA:HB2	1.29	1.25
1:D:1006:ARG:HA	1:D:1006:ARG:NH1	1.49	1.25
1:A:76:ILE:HB	1:A:911:HIS:NE2	1.49	1.24
1:A:210:THR:CB	1:A:224:CYS:HB2	1.66	1.24
1:D:352:SER:C	1:D:358:GLU:OE2	1.75	1.23
1:D:827:ARG:O	1:D:829:PRO:HD3	1.36	1.23
1:A:656:TYR:HB3	1:A:720:THR:CB	1.66	1.23
2:C:241:TRP:HB2	2:C:336:LEU:CD1	1.69	1.23
2:C:303:LEU:HA	2:C:337:SER:OG	1.31	1.23
1:D:891:VAL:HG11	1:D:1135:ASP:O	1.12	1.22
2:E:382:LYS:HE2	2:E:412:SER:O	1.37	1.22
1:D:259:ARG:HA	1:D:259:ARG:CZ	1.68	1.22
1:D:1131:ILE:HD11	1:D:1138:ARG:N	1.55	1.22
1:D:249:VAL:CG1	1:D:252:GLY:H	1.52	1.21
1:D:367:PRO:CD	1:D:368:PRO:HD3	1.70	1.21
1:D:420:HIS:ND1	1:D:421:PRO:HD2	1.54	1.21
1:D:801:TRP:CZ3	1:D:870:VAL:HG11	1.74	1.21
1:A:389:PHE:HA	1:A:393:MET:SD	1.79	1.21
1:D:548:CYS:O	1:D:552:LEU:HG	1.35	1.21
2:F:447:THR:O	2:F:452:LEU:HD12	1.40	1.21
2:E:78:PHE:O	2:E:79:LEU:HG	1.39	1.21
1:D:1029:GLN:CG	1:D:1030:ARG:HE	1.53	1.20

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1020:ILE:HD13	1:D:1021:SER:N	1.54	1.20
1:A:477:GLU:CG	1:A:478:ARG:H	1.54	1.20
1:D:278:ILE:HG22	1:D:279:ARG:HD3	1.23	1.20
1:A:213:VAL:HG12	1:A:401:TRP:NE1	1.55	1.20
1:D:504:LYS:CB	1:D:523:ASP:HB2	1.71	1.20
1:D:352:SER:O	1:D:358:GLU:CD	1.81	1.19
1:D:604:ALA:HA	1:D:613:HIS:CE1	1.77	1.19
1:A:996:ARG:NH1	1:A:1000:GLU:HB3	1.58	1.19
1:A:346:ASP:HB3	1:A:1040:LYS:HZ3	1.03	1.19
1:D:275:ARG:CB	1:D:843:GLN:HB3	1.71	1.18
1:A:666:GLN:O	1:A:670:PRO:HD3	1.36	1.18
1:A:927:ARG:HG2	1:A:928:GLY:H	1.04	1.18
1:D:610:PHE:H	1:D:611:PRO:HD3	1.07	1.18
1:D:294:THR:HG22	1:D:407:PHE:HE2	1.06	1.18
1:D:608:ASP:HB3	1:D:620:TRP:CD2	1.78	1.18
1:A:1026:ARG:CB	1:A:1029:GLN:HB2	1.74	1.18
2:E:432:TYR:CB	2:E:437:ILE:HD11	1.73	1.18
2:F:208:LEU:CD1	2:F:240:VAL:HB	1.72	1.18
1:D:992:LEU:HB3	1:D:996:ARG:HG2	1.18	1.18
1:A:498:LYS:HD2	1:A:518:ALA:HB3	1.24	1.17
1:D:1027:LYS:NZ	1:D:1091:GLU:HG2	1.60	1.17
2:B:383:VAL:HG13	2:B:413:VAL:CG1	1.75	1.17
2:C:424:SER:HB3	2:C:427:GLN:HE22	1.07	1.17
1:D:900:ALA:HB1	1:D:915:ALA:HB1	1.27	1.17
1:A:1005:VAL:HA	1:A:1008:LEU:CD1	1.73	1.17
1:A:1199:THR:HA	1:A:1200:PRO:C	1.64	1.17
2:F:446:THR:HB	2:F:450:ASN:HB2	1.18	1.16
1:D:642:SER:HA	1:D:645:VAL:HG11	1.26	1.16
1:D:891:VAL:CG1	1:D:1135:ASP:O	1.94	1.16
1:D:1165:ALA:HB2	1:D:1173:LEU:HD12	1.22	1.16
2:B:72:ILE:HA	2:B:75:ARG:HE	1.01	1.16
1:D:420:HIS:CD2	1:D:1020:ILE:HD12	1.79	1.16
1:A:477:GLU:HG3	1:A:478:ARG:N	1.55	1.16
1:D:1014:ARG:HD2	1:D:1014:ARG:O	1.45	1.16
1:A:213:VAL:CG1	1:A:401:TRP:HE1	1.59	1.16
1:A:612:LEU:HG	1:A:614:TYR:H	1.09	1.16
1:A:871:GLY:C	1:A:873:GLU:H	1.38	1.16
1:A:593:SER:HA	1:A:596:MET:HG3	1.28	1.16
1:D:1029:GLN:HG2	1:D:1030:ARG:HE	1.02	1.15
2:F:454:HIS:HB2	2:F:466:MET:O	1.46	1.15
2:F:78:PHE:CE1	2:F:103:GLY:HA2	1.80	1.15

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:454:HIS:HA	2:F:468:ILE:HD12	1.28	1.15
1:A:823:ARG:HB3	1:A:825:VAL:HG23	1.20	1.15
1:D:350:ILE:CG2	1:D:352:SER:HB2	1.74	1.15
2:E:376:PRO:O	2:E:380:PRO:HD3	1.42	1.15
2:B:383:VAL:CG1	2:B:413:VAL:HG13	1.76	1.15
2:C:78:PHE:CE1	2:C:103:GLY:HA2	1.81	1.15
1:D:616:GLU:O	1:D:725:PRO:HD3	1.47	1.15
2:E:372:LEU:HG	2:E:373:LYS:N	1.61	1.15
1:D:891:VAL:HG11	1:D:1135:ASP:C	1.67	1.15
1:A:612:LEU:HD22	1:A:621:GLY:N	1.61	1.15
1:D:548:CYS:C	1:D:551:LYS:HD3	1.67	1.14
1:D:891:VAL:HA	1:D:894:GLN:CD	1.66	1.14
2:C:127:PRO:HD2	2:C:210:GLN:HB2	1.26	1.14
1:D:551:LYS:H	1:D:551:LYS:CD	1.49	1.14
1:A:275:ARG:NH2	1:A:845:VAL:HA	1.62	1.14
2:C:386:ASP:HB2	2:C:442:LEU:HA	1.19	1.14
2:C:418:LEU:O	2:C:421:MET:HG2	1.46	1.14
1:A:472:GLN:HG3	2:C:460:THR:CA	1.77	1.14
1:D:509:THR:HB	1:D:512:LYS:HB3	1.27	1.14
2:B:309:HIS:HB3	2:B:312:LEU:HD22	1.29	1.14
1:D:1131:ILE:CD1	1:D:1138:ARG:H	1.58	1.14
1:D:356:LEU:HD13	1:D:369:LEU:CD2	1.75	1.14
1:A:1030:ARG:HG2	1:A:1040:LYS:HZ1	1.12	1.14
2:C:443:VAL:HA	2:C:447:THR:HG21	1.28	1.14
1:D:1204:THR:HA	1:D:1207:GLU:HB3	1.15	1.14
1:D:866:ARG:HG3	1:D:1069:ILE:HG21	1.15	1.14
1:A:253:ALA:HB2	1:A:280:GLU:HB2	1.16	1.14
1:A:996:ARG:HH12	1:A:1003:TRP:CB	1.58	1.14
1:D:300:ALA:HB2	1:D:847:ALA:HB1	1.18	1.13
1:D:81:ARG:NH1	1:D:81:ARG:HB2	1.63	1.13
1:D:504:LYS:HD2	1:D:523:ASP:HB3	1.23	1.13
1:D:801:TRP:CE3	1:D:870:VAL:HG11	1.82	1.13
1:D:242:ALA:HB1	1:D:245:ILE:HG12	1.21	1.13
1:D:267:VAL:HB	1:D:269:HIS:CD2	1.82	1.13
2:F:239:LEU:O	2:F:338:VAL:HG22	1.46	1.13
1:A:513:LEU:CB	1:A:568:GLY:HA2	1.78	1.13
2:C:82:SER:HA	2:C:86:LEU:HD21	1.18	1.13
1:D:1006:ARG:CA	1:D:1006:ARG:HH11	1.61	1.13
2:F:304:TRP:N	2:F:337:SER:HB2	1.60	1.13
2:F:387:VAL:CG2	2:F:417:TYR:HA	1.78	1.13
1:D:449:GLN:HG2	1:D:805:HIS:NE2	1.62	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:81:ARG:HH11	1:D:81:ARG:HB2	1.11	1.13
1:A:618:HIS:N	1:A:725:PRO:HG2	1.64	1.13
1:D:275:ARG:HE	1:D:842:PRO:HA	1.02	1.13
2:E:213:VAL:HA	2:E:235:THR:HA	1.28	1.13
1:D:1006:ARG:NH1	1:D:1009:ASN:HB2	1.61	1.13
1:D:891:VAL:HG13	1:D:1135:ASP:HB3	1.28	1.13
1:A:253:ALA:HB2	1:A:280:GLU:CB	1.77	1.13
1:A:502:LYS:HG3	1:A:503:VAL:H	1.08	1.13
1:A:444:TYR:HD2	1:A:874:LEU:HD23	1.07	1.13
2:C:125:VAL:HG22	2:C:208:LEU:HA	1.16	1.13
2:E:432:TYR:HA	2:E:435:MET:HB2	1.18	1.12
1:A:608:ASP:HB3	1:A:779:LEU:CD1	1.77	1.12
2:C:351:TYR:C	2:C:355:GLN:HB2	1.68	1.12
1:A:992:LEU:HD23	1:A:997:LEU:HD12	1.25	1.12
2:B:279:GLN:O	2:B:280:ASP:HB2	1.49	1.12
1:D:797:MET:HB2	1:D:869:ARG:HD2	1.30	1.12
2:C:236:GLU:HA	2:C:341:ASP:HB3	1.30	1.12
2:C:446:THR:HA	2:C:450:ASN:CB	1.78	1.12
1:A:959:GLN:HB2	1:A:960:PRO:CD	1.78	1.12
1:D:235:TRP:HD1	1:D:236:THR:HG22	1.06	1.12
1:D:1221:TYR:HA	1:D:1224:ILE:CG2	1.80	1.12
1:D:652:ILE:HA	1:D:655:LEU:HG	1.30	1.12
2:F:203:ARG:CA	2:F:325:ARG:HD3	1.79	1.12
1:A:756:ASP:HB3	1:A:1057:MET:HG2	1.20	1.12
1:A:245:ILE:HG13	1:A:246:PRO:HD3	1.31	1.12
1:D:1133:ILE:HB	1:D:1136:GLU:HB2	1.17	1.12
1:D:203:LEU:H	1:D:380:GLY:HA2	1.02	1.12
1:A:1043:VAL:HG13	1:A:1046:GLU:HB3	1.16	1.12
1:A:658:LYS:HG3	1:A:713:PRO:HB2	1.30	1.12
2:B:285:LYS:HG2	2:B:286:GLY:N	1.56	1.12
2:C:203:ARG:HA	2:C:325:ARG:HD3	1.20	1.12
1:A:346:ASP:HB3	1:A:1040:LYS:NZ	1.64	1.12
1:D:235:TRP:CD1	1:D:236:THR:HG22	1.84	1.12
1:D:668:LEU:O	1:D:671:GLN:HB2	1.46	1.12
1:D:1193:VAL:CG2	1:D:1194:THR:H	1.60	1.12
1:D:591:LEU:HD11	1:D:594:LEU:HG	1.31	1.12
1:D:775:GLU:HG2	1:D:780:GLN:HG3	1.31	1.12
1:A:503:VAL:HB	1:A:521:PRO:HA	1.30	1.12
1:D:649:TYR:HA	1:D:652:ILE:HB	1.19	1.11
2:F:387:VAL:HG21	2:F:417:TYR:HA	1.25	1.11
1:D:128:PRO:HA	1:D:130:LEU:CD2	1.80	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:79:LEU:HD21	1:A:1173:LEU:O	1.48	1.11
1:A:210:THR:HB	1:A:224:CYS:CB	1.79	1.11
1:A:70:ARG:HB2	1:A:81:ARG:HB3	1.32	1.11
1:A:135:LEU:HD23	1:A:135:LEU:H	1.05	1.11
1:D:367:PRO:HD2	1:D:368:PRO:CD	1.79	1.11
2:F:372:LEU:HD11	2:F:436:SER:HB2	1.12	1.11
1:A:775:GLU:HA	1:A:780:GLN:HG3	1.32	1.11
1:D:1200:PRO:O	1:D:1203:PRO:HD2	1.47	1.11
1:A:659:HIS:HB2	1:A:719:LEU:HD12	1.30	1.11
1:A:854:ALA:HB1	1:A:860:LEU:HD13	1.30	1.11
1:A:408:GLN:HA	1:A:411:LEU:HG	1.26	1.11
2:C:131:LEU:HG	2:C:133:HIS:CE1	1.85	1.11
1:A:502:LYS:HD2	1:A:506:GLU:OE1	1.49	1.11
1:D:375:GLU:HG3	1:D:379:LYS:HB3	1.29	1.11
2:F:247:THR:HB	2:F:251:TRP:CD1	1.85	1.11
2:F:354:PHE:CE1	2:F:370:LYS:HD3	1.84	1.11
1:A:436:PRO:HB3	1:A:881:PRO:HD3	1.13	1.11
1:A:656:TYR:HB3	1:A:720:THR:HB	1.24	1.11
1:D:420:HIS:HD2	1:D:1020:ILE:HD12	0.97	1.11
1:D:1131:ILE:HD11	1:D:1138:ARG:H	1.04	1.11
1:D:781:ALA:HB3	1:D:792:LEU:HD21	1.31	1.11
1:A:758:ASN:HB2	1:A:1058:PHE:HZ	1.09	1.11
1:D:1204:THR:CA	1:D:1207:GLU:HB3	1.81	1.10
1:D:179:GLY:HA3	1:D:182:GLY:HA3	1.13	1.10
1:D:592:LEU:HA	1:D:595:GLN:HG3	1.33	1.10
1:D:1209:ARG:HH21	2:F:253:ASP:HA	1.01	1.10
1:D:1206:MET:SD	1:D:1210:TYR:HA	1.90	1.10
1:A:170:ALA:HB1	1:A:175:TRP:CZ2	1.85	1.10
1:A:612:LEU:HG	1:A:613:HIS:N	1.59	1.10
2:C:372:LEU:HD11	2:C:436:SER:HB2	1.33	1.10
1:A:612:LEU:HG	1:A:613:HIS:H	1.06	1.10
1:A:249:VAL:HG11	1:A:817:PRO:HG3	1.30	1.10
2:B:407:LEU:HD21	2:C:120:VAL:HA	1.23	1.10
1:D:242:ALA:HB1	1:D:245:ILE:CG1	1.80	1.10
1:D:1231:LEU:HG	1:D:1232:GLU:H	0.98	1.10
1:D:212:ALA:O	1:D:213:VAL:HB	1.52	1.10
1:D:556:THR:HA	1:D:559:LEU:HD23	1.33	1.10
1:A:190:PRO:HB3	1:A:192:GLU:HG2	1.15	1.10
2:B:117:SER:HB2	2:B:262:TRP:NE1	1.64	1.10
1:A:363:ARG:HH22	1:A:409:GLN:HB2	1.16	1.10
1:D:798:ILE:HG22	1:D:869:ARG:HH22	1.06	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:241:TRP:CB	2:F:336:LEU:HD12	1.80	1.10
2:C:203:ARG:CA	2:C:325:ARG:HD3	1.81	1.10
1:A:176:THR:HA	1:A:218:SER:HB2	1.13	1.10
1:D:1193:VAL:HG22	1:D:1194:THR:N	1.66	1.10
1:A:490:TRP:HA	1:A:580:LEU:HD22	1.29	1.10
1:A:436:PRO:HB2	1:A:879:GLN:HA	1.33	1.10
1:D:504:LYS:HB2	1:D:523:ASP:HB2	1.10	1.10
1:D:270:ASN:OD1	1:D:294:THR:HG21	1.51	1.10
2:E:403:PHE:HD1	2:E:413:VAL:HB	1.01	1.10
2:C:382:LYS:HD3	2:C:411:ILE:HG13	1.34	1.10
1:D:1207:GLU:HA	1:D:1210:TYR:HB3	1.33	1.09
1:D:1210:TYR:OH	1:D:1212:ILE:HB	1.52	1.09
1:A:1148:ARG:HG2	1:A:1148:ARG:HH11	1.11	1.09
1:A:575:LYS:HG3	1:A:576:LEU:H	1.11	1.09
1:A:997:LEU:HD22	1:A:998:SER:H	1.00	1.09
2:C:446:THR:O	2:C:450:ASN:HB3	1.49	1.09
1:A:288:ARG:HA	1:A:288:ARG:HE	0.96	1.09
2:C:423:SER:OG	2:C:428:LEU:HD21	1.51	1.09
2:E:389:ARG:HE	2:E:390:GLY:N	1.48	1.09
2:F:466:MET:HB3	2:F:471:LEU:HD22	1.12	1.09
2:F:82:SER:HA	2:F:86:LEU:HD21	1.30	1.09
1:D:350:ILE:HG23	1:D:352:SER:HB2	1.30	1.09
2:F:127:PRO:CD	2:F:210:GLN:HB2	1.82	1.09
2:C:466:MET:HB3	2:C:471:LEU:HD22	1.24	1.09
1:D:551:LYS:H	1:D:551:LYS:HD2	0.95	1.09
1:D:607:TRP:HB2	1:D:779:LEU:HD23	1.33	1.09
1:A:515:ILE:HG21	1:A:565:HIS:HA	1.29	1.09
1:A:363:ARG:HD2	1:A:406:VAL:HB	1.29	1.09
1:A:606:THR:HG21	1:A:612:LEU:HB3	1.34	1.09
2:F:180:LYS:C	2:F:180:LYS:HE3	1.73	1.09
2:F:276:SER:O	2:F:287:ASN:HB2	1.51	1.09
1:D:458:GLU:HA	1:D:461:LYS:HD3	1.30	1.09
1:D:1206:MET:CG	2:F:257:ARG:HH11	1.66	1.09
1:A:1043:VAL:HA	1:A:1046:GLU:H	1.16	1.09
2:B:204:LEU:HG	2:B:325:ARG:HH21	1.18	1.09
2:F:236:GLU:HA	2:F:341:ASP:HB3	1.33	1.09
1:A:190:PRO:HB3	1:A:192:GLU:CG	1.81	1.09
2:B:262:TRP:O	2:B:265:LYS:HB3	1.52	1.08
1:D:245:ILE:HG13	1:D:246:PRO:N	1.63	1.08
1:D:1001:GLY:HA2	1:D:1004:LEU:HD22	1.09	1.08
2:F:301:GLU:HB3	2:F:339:ASN:HB3	1.31	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:72:ILE:HA	2:B:75:ARG:NE	1.67	1.08
2:C:456:ARG:HB2	2:C:463:LYS:CA	1.84	1.08
1:D:316:LYS:HG2	1:D:348:LEU:HG	1.36	1.08
1:D:546:ARG:HH21	2:F:404:ASN:HB3	1.11	1.08
2:E:322:LEU:O	2:E:324:GLY:N	1.86	1.08
1:D:797:MET:HB2	1:D:869:ARG:CD	1.82	1.08
1:A:371:LYS:HD3	1:A:396:CYS:HA	1.30	1.08
2:C:239:LEU:O	2:C:336:LEU:HD22	1.52	1.08
2:E:388:GLY:C	2:E:395:LEU:HD11	1.73	1.08
2:E:249:ASN:HD21	2:E:285:LYS:HE2	1.12	1.08
1:D:888:GLY:HA3	1:D:1138:ARG:HD3	1.24	1.08
2:E:382:LYS:HG2	2:E:412:SER:HB2	1.31	1.08
1:A:228:LEU:HD23	1:A:229:VAL:N	1.68	1.08
1:A:374:ARG:HE	1:A:392:LEU:HD13	1.17	1.08
2:C:421:MET:HG3	2:C:422:GLN:H	1.14	1.08
1:A:271:VAL:HG12	1:A:272:SER:H	1.14	1.08
1:A:549:LEU:HD21	2:C:398:VAL:HB	1.12	1.08
1:D:612:LEU:HD21	1:D:622:TYR:CD1	1.89	1.08
1:D:874:LEU:O	1:D:878:VAL:HG22	1.52	1.08
2:F:127:PRO:HD2	2:F:210:GLN:HB2	1.30	1.08
1:A:1068:ASP:H	1:A:1071:ARG:HG2	1.16	1.08
2:B:371:VAL:HA	2:B:433:ASP:HB3	1.09	1.08
2:C:435:MET:HE2	2:C:435:MET:HA	1.36	1.08
1:A:249:VAL:HG21	1:A:817:PRO:HD3	1.20	1.08
1:D:1141:VAL:HB	1:D:1146:ARG:NE	1.68	1.07
1:D:1172:ASP:O	1:D:1173:LEU:HG	1.54	1.07
2:C:375:HIS:H	2:C:378:LEU:HD22	1.16	1.07
1:A:297:MET:SD	1:A:411:LEU:HD22	1.94	1.07
2:C:385:LEU:HD11	2:C:402:LEU:HB2	1.36	1.07
2:E:371:VAL:HA	2:E:433:ASP:OD2	1.53	1.07
2:E:456:ARG:HB2	2:E:463:LYS:HG2	1.27	1.07
2:F:242:PHE:HB3	2:F:333:PRO:HB2	1.35	1.07
1:D:1067:SER:HA	1:D:1071:ARG:HA	1.12	1.07
1:A:213:VAL:HG13	1:A:221:TYR:OH	1.50	1.07
1:A:606:THR:HG23	1:A:613:HIS:C	1.72	1.07
1:A:439:GLN:HG2	1:A:835:GLY:CA	1.83	1.07
1:A:802:ARG:O	1:A:805:HIS:HB3	1.53	1.07
1:D:585:TRP:HB3	1:D:587:PRO:HD2	1.35	1.07
1:D:598:VAL:HA	1:D:601:LYS:HE2	1.36	1.07
1:D:869:ARG:H	1:D:1200:PRO:CB	1.66	1.07
1:D:356:LEU:HD13	1:D:369:LEU:HD21	1.37	1.07

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:389:PHE:HA	1:D:393:MET:CG	1.84	1.07
1:A:170:ALA:HB1	1:A:175:TRP:HZ2	1.10	1.07
2:C:208:LEU:CD1	2:C:240:VAL:HB	1.82	1.07
1:A:1131:ILE:HG21	1:A:1138:ARG:N	1.70	1.07
1:A:1145:ASP:HB3	1:A:1148:ARG:CZ	1.83	1.07
1:A:603:MET:O	1:A:605:LEU:HG	1.53	1.07
2:C:239:LEU:N	2:C:338:VAL:HG12	1.68	1.07
1:A:645:VAL:O	1:A:646:VAL:HG12	1.54	1.07
1:D:1069:ILE:HG23	1:D:1070:PRO:CD	1.85	1.07
1:D:389:PHE:HA	1:D:393:MET:HG3	1.34	1.07
1:A:249:VAL:HG21	1:A:817:PRO:CD	1.82	1.07
1:D:203:LEU:HB3	1:D:380:GLY:HA3	1.31	1.07
1:D:294:THR:HG22	1:D:407:PHE:CE2	1.89	1.07
2:B:323:HIS:HA	2:B:332:VAL:HA	1.36	1.07
2:C:301:GLU:HB3	2:C:339:ASN:HB3	1.36	1.07
1:A:80:SER:N	1:A:83:LEU:HD11	1.69	1.07
2:B:105:GLU:OE1	2:B:380:PRO:HA	1.55	1.07
2:B:309:HIS:HB3	2:B:312:LEU:CD2	1.83	1.07
1:A:503:VAL:CB	1:A:521:PRO:HA	1.85	1.07
1:D:670:PRO:HG2	1:D:673:ALA:HB2	1.29	1.07
2:E:203:ARG:HH12	2:F:418:LEU:HD22	1.14	1.06
2:F:372:LEU:HD11	2:F:436:SER:CB	1.83	1.06
1:D:181:GLU:HA	1:D:181:GLU:OE1	1.46	1.06
1:A:996:ARG:HH22	1:A:1003:TRP:HB3	1.13	1.06
2:B:213:VAL:HA	2:B:235:THR:O	1.55	1.06
1:A:232:ARG:HB3	1:A:239:LEU:HD22	1.13	1.06
1:A:618:HIS:HB2	1:A:725:PRO:HB2	1.36	1.06
2:C:443:VAL:CA	2:C:447:THR:HG21	1.84	1.06
2:E:418:LEU:HD22	2:F:122:ARG:HH21	1.14	1.06
1:D:107:LEU:HD21	1:D:913:CYS:HA	1.36	1.06
2:E:107:ARG:CG	2:E:107:ARG:HH11	1.67	1.06
1:D:210:THR:HG23	1:D:224:CYS:HB3	1.36	1.06
1:D:279:ARG:CD	1:D:841:LEU:HB3	1.85	1.06
2:F:234:LYS:HZ3	2:F:344:ARG:HD2	1.14	1.06
1:A:893:SER:HB2	1:A:896:LEU:HB2	1.31	1.06
1:A:1026:ARG:HB3	1:A:1029:GLN:CG	1.84	1.06
2:B:417:TYR:OH	2:C:122:ARG:HD3	1.52	1.06
1:A:503:VAL:HG11	1:A:520:ALA:C	1.76	1.06
1:D:856:GLU:OE1	1:D:859:TRP:HB2	1.54	1.06
1:D:989:THR:HA	1:D:998:SER:OG	1.56	1.06
1:A:711:ALA:C	1:A:713:PRO:HD3	1.75	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:903:GLY:CA	1:A:1002:GLU:HB3	1.85	1.06
1:A:869:ARG:HG2	1:A:870:VAL:H	1.16	1.06
1:A:784:GLY:H	2:C:363:ARG:CD	1.69	1.06
1:D:1055:SER:HA	1:D:1058:PHE:HB3	1.06	1.06
1:D:177:ARG:HB3	1:D:218:SER:HB3	1.37	1.06
2:F:247:THR:HB	2:F:251:TRP:HD1	1.15	1.06
2:F:454:HIS:CB	2:F:466:MET:HB2	1.86	1.06
1:D:1005:VAL:HA	1:D:1008:LEU:CD1	1.85	1.06
1:D:1209:ARG:HH21	2:F:253:ASP:CA	1.67	1.06
1:D:927:ARG:HG2	1:D:928:GLY:H	1.09	1.06
2:E:443:VAL:HG22	2:E:453:ILE:HG21	1.33	1.06
2:F:406:LEU:HA	2:F:409:ASN:ND2	1.70	1.06
1:A:1014:ARG:HB2	1:A:1026:ARG:CZ	1.85	1.06
1:A:875:LYS:O	1:A:878:VAL:HG23	1.54	1.06
2:B:202:LYS:O	2:B:325:ARG:HD2	1.53	1.06
1:A:363:ARG:NH1	1:A:406:VAL:HA	1.70	1.06
1:A:613:HIS:HB3	1:A:717:LEU:HG	1.35	1.06
1:A:845:VAL:HB	1:A:855:VAL:HB	1.36	1.06
1:D:203:LEU:H	1:D:380:GLY:CA	1.69	1.06
2:F:351:TYR:C	2:F:355:GLN:HB2	1.75	1.06
2:F:125:VAL:HG22	2:F:208:LEU:HA	1.07	1.06
2:C:242:PHE:HB3	2:C:333:PRO:HB2	1.36	1.06
1:D:956:GLY:HA2	1:D:1094:THR:HG23	1.08	1.05
2:F:446:THR:HG21	2:F:450:ASN:OD1	1.56	1.05
1:D:546:ARG:NH2	2:F:404:ASN:HB3	1.69	1.05
1:A:76:ILE:HG13	1:A:77:GLN:HG3	1.36	1.05
1:D:504:LYS:HB2	1:D:523:ASP:CB	1.85	1.05
2:E:390:GLY:N	2:E:395:LEU:HG	1.69	1.05
1:D:959:GLN:HB2	1:D:960:PRO:HD2	1.39	1.05
2:F:239:LEU:O	2:F:336:LEU:HD22	1.56	1.05
2:E:105:GLU:OE2	2:E:382:LYS:O	1.74	1.05
2:E:254:PHE:HA	2:E:257:ARG:HB3	1.37	1.05
2:B:285:LYS:CG	2:B:286:GLY:H	1.69	1.05
1:A:1005:VAL:HA	1:A:1008:LEU:HD13	1.34	1.05
1:A:76:ILE:HG23	1:A:77:GLN:H	0.95	1.05
2:C:180:LYS:HE3	2:C:180:LYS:C	1.77	1.05
1:D:885:THR:HB	1:D:1146:ARG:HD2	1.34	1.05
1:D:432:VAL:HG11	1:D:1128:ARG:HE	1.20	1.05
1:D:591:LEU:CD1	1:D:594:LEU:HG	1.86	1.05
1:D:612:LEU:HB2	1:D:620:TRP:HB2	1.34	1.05
2:F:208:LEU:HD11	2:F:240:VAL:HB	1.38	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:336:LEU:HD21	2:B:338:VAL:HG23	1.31	1.05
2:C:241:TRP:CB	2:C:336:LEU:HD12	1.85	1.05
1:A:607:TRP:HA	1:A:610:PHE:O	1.54	1.05
1:D:798:ILE:CG2	1:D:869:ARG:HH12	1.69	1.05
1:D:428:LEU:CD1	1:D:428:LEU:H	1.68	1.05
1:A:215:ILE:HG23	1:A:216:SER:H	1.20	1.05
2:B:101:PRO:HA	2:C:128:VAL:HG21	1.34	1.05
1:D:367:PRO:HD2	1:D:368:PRO:HD3	1.05	1.05
2:F:203:ARG:HA	2:F:325:ARG:HD3	1.07	1.04
1:D:620:TRP:O	1:D:748:TRP:HB2	1.56	1.04
2:F:387:VAL:HA	2:F:443:VAL:HG22	1.09	1.04
1:A:1026:ARG:HB3	1:A:1029:GLN:CB	1.86	1.04
1:A:1027:LYS:HB2	1:A:1092:PHE:HD2	1.22	1.04
2:E:382:LYS:HG3	2:E:383:VAL:N	1.72	1.04
1:D:822:PRO:HD2	1:D:825:VAL:CG2	1.87	1.04
1:A:167:PRO:HG2	1:A:177:ARG:HG3	1.38	1.04
1:A:667:GLN:O	1:A:670:PRO:HD2	1.58	1.04
2:C:431:LYS:HA	2:C:434:GLU:OE1	1.56	1.04
1:D:411:LEU:O	1:D:415:LEU:HB2	1.56	1.04
2:F:203:ARG:HA	2:F:325:ARG:CD	1.87	1.04
1:A:728:THR:HG22	1:A:729:GLN:H	1.20	1.04
2:C:342:LEU:CD2	2:C:345:GLY:H	1.71	1.04
2:C:85:GLN:HB3	2:C:95:CYS:O	1.56	1.04
1:A:239:LEU:O	1:A:241:PRO:HD2	1.58	1.04
1:A:959:GLN:CB	1:A:960:PRO:HD3	1.87	1.04
2:B:74:GLN:HG3	2:B:75:ARG:N	1.66	1.04
1:D:822:PRO:HD2	1:D:825:VAL:HG21	1.07	1.04
2:E:403:PHE:CD1	2:E:413:VAL:HB	1.92	1.04
1:D:428:LEU:HD12	1:D:428:LEU:H	0.93	1.04
2:F:342:LEU:CD2	2:F:345:GLY:H	1.71	1.04
1:A:1208:ARG:NH2	2:C:277:ASP:OD2	1.91	1.04
1:A:885:THR:CB	1:A:1146:ARG:HE	1.70	1.04
2:C:442:LEU:O	2:C:455:LEU:HD22	1.56	1.04
1:A:1056:GLU:HA	1:A:1059:ASN:ND2	1.73	1.04
2:F:239:LEU:H	2:F:338:VAL:CG1	1.70	1.03
1:D:1029:GLN:NE2	1:D:1030:ARG:HH21	1.55	1.03
1:D:798:ILE:HG22	1:D:869:ARG:NH2	1.71	1.03
1:A:1060:LYS:HE3	1:A:1065:ALA:H	1.18	1.03
2:C:406:LEU:HA	2:C:409:ASN:ND2	1.73	1.03
1:A:161:GLN:O	1:A:163:PRO:HD3	1.56	1.03
1:D:157:LEU:HD21	1:D:194:ALA:HB3	1.37	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:608:ASP:CB	1:A:779:LEU:HD13	1.87	1.03
1:A:916:PHE:HA	1:A:919:MET:HB2	1.35	1.03
1:A:1047:ARG:HA	1:A:1047:ARG:NH1	1.72	1.03
2:B:374:LEU:H	2:B:458:ARG:NH2	1.56	1.03
1:A:1236:GLN:HB3	1:A:1237:PRO:HD3	1.40	1.03
2:C:244:PRO:HD2	2:C:247:THR:HG21	1.39	1.03
1:D:348:LEU:HD21	1:D:350:ILE:HG13	1.35	1.03
2:E:107:ARG:HG3	2:E:107:ARG:HH11	0.89	1.03
1:A:1006:ARG:NH2	1:A:1010:LEU:HD11	1.74	1.03
1:A:1128:ARG:HG3	1:A:1129:PHE:N	1.69	1.03
1:A:175:TRP:HB3	1:A:186:PRO:HD3	1.41	1.03
1:A:669:MET:O	1:A:673:ALA:CB	2.06	1.03
2:F:382:LYS:HD3	2:F:411:ILE:HG13	1.34	1.03
1:D:371:LYS:HA	1:D:374:ARG:CD	1.88	1.03
1:D:798:ILE:HG23	1:D:869:ARG:HH12	1.17	1.03
2:F:473:ASP:HA	2:F:476:ILE:CG2	1.89	1.03
1:A:371:LYS:O	1:A:373:PRO:HD3	1.56	1.03
1:A:80:SER:H	1:A:83:LEU:HD11	1.19	1.03
1:D:866:ARG:HB3	1:D:867:PRO:HD3	1.41	1.02
2:E:262:TRP:HA	2:E:265:LYS:HE3	1.40	1.02
2:F:466:MET:CB	2:F:471:LEU:HD22	1.89	1.02
2:F:386:ASP:O	2:F:443:VAL:HG13	1.59	1.02
1:A:249:VAL:CG2	1:A:817:PRO:HD3	1.89	1.02
1:A:439:GLN:HG2	1:A:835:GLY:HA3	1.38	1.02
2:C:102:LEU:HD11	2:C:435:MET:HE1	1.41	1.02
1:A:997:LEU:CD2	1:A:998:SER:H	1.71	1.02
2:C:236:GLU:HA	2:C:341:ASP:CB	1.87	1.02
1:D:775:GLU:HA	1:D:780:GLN:HG2	1.40	1.02
1:D:815:TRP:HA	1:D:838:GLY:HA3	1.41	1.02
1:D:1209:ARG:NH1	2:F:252:LEU:HD13	1.74	1.02
1:D:213:VAL:HG22	1:D:401:TRP:CH2	1.93	1.02
1:D:854:ALA:HB1	1:D:860:LEU:CD1	1.88	1.02
1:A:125:LEU:HD23	1:A:125:LEU:H	1.19	1.02
2:E:428:LEU:HD13	2:E:428:LEU:H	1.25	1.02
2:E:447:THR:HG21	2:E:453:ILE:HG23	1.40	1.02
2:F:200:VAL:HG11	2:F:204:LEU:HB2	1.38	1.02
2:F:439:PHE:HD1	2:F:458:ARG:HB2	1.20	1.02
1:A:498:LYS:HE3	1:A:518:ALA:H	1.23	1.02
1:A:76:ILE:HG23	1:A:77:GLN:N	1.73	1.02
1:A:288:ARG:HA	1:A:288:ARG:NE	1.64	1.02
1:D:623:LEU:O	1:D:625:PRO:HD3	1.59	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:645:VAL:O	1:A:650:ARG:HD2	1.60	1.02
1:D:420:HIS:HD1	1:D:421:PRO:CD	1.71	1.02
1:D:605:LEU:HD22	1:D:781:ALA:CB	1.88	1.02
2:F:209:ALA:HB1	2:F:238:SER:O	1.60	1.02
2:C:300:ILE:HG23	2:C:342:LEU:HD11	1.40	1.02
2:C:430:SER:O	2:C:434:GLU:HG3	1.59	1.02
1:A:853:ARG:NH1	1:A:1105:ALA:HB3	1.73	1.02
1:A:479:TYR:O	1:A:480:LYS:HG2	1.57	1.02
1:D:135:LEU:HD23	1:D:135:LEU:H	1.22	1.02
2:F:386:ASP:HB2	2:F:442:LEU:HA	1.03	1.02
2:F:386:ASP:HB3	2:F:442:LEU:HD23	1.40	1.02
1:A:925:LYS:HG3	1:A:926:SER:N	1.73	1.02
1:A:1128:ARG:HG3	1:A:1129:PHE:H	0.89	1.02
1:A:192:GLU:HB3	1:A:217:PRO:HG2	1.39	1.02
1:A:444:TYR:CD2	1:A:874:LEU:HD23	1.94	1.02
1:A:502:LYS:HG3	1:A:503:VAL:N	1.74	1.02
2:E:186:LEU:O	2:E:190:LEU:HD13	1.58	1.02
1:D:1188:CYS:C	1:D:1189:LEU:HD22	1.80	1.01
1:D:351:SER:C	1:D:358:GLU:OE2	1.97	1.01
1:D:297:MET:HG2	1:D:411:LEU:HD22	1.40	1.01
1:D:717:LEU:O	1:D:721:ALA:HB2	1.59	1.01
1:D:607:TRP:CD1	1:D:780:GLN:N	2.27	1.01
1:D:880:ALA:HB2	1:D:1187:ARG:O	1.58	1.01
1:D:992:LEU:HB3	1:D:996:ARG:CG	1.90	1.01
1:D:1014:ARG:CG	1:D:1026:ARG:HE	1.72	1.01
1:D:1027:LYS:HZ2	1:D:1091:GLU:HG2	1.13	1.01
1:D:86:GLN:NE2	1:D:132:GLY:H	1.56	1.01
2:E:266:PHE:HB3	2:E:376:PRO:HD3	1.39	1.01
1:A:612:LEU:HD22	1:A:621:GLY:H	0.85	1.01
1:A:623:LEU:HB3	1:A:667:GLN:NE2	1.74	1.01
1:D:213:VAL:HG21	1:D:397:ALA:HB1	1.40	1.01
1:D:249:VAL:HG13	1:D:252:GLY:H	0.90	1.01
1:D:577:CYS:O	1:D:579:ARG:N	1.93	1.01
1:D:1131:ILE:CD1	1:D:1131:ILE:H	1.73	1.01
1:A:1190:ARG:CZ	1:A:1197:CYS:HB2	1.89	1.01
1:A:408:GLN:HA	1:A:411:LEU:CG	1.89	1.01
2:B:419:GLU:HB2	2:C:203:ARG:NH2	1.75	1.01
1:D:735:GLY:HA3	1:D:749:PHE:HB3	1.42	1.01
1:D:275:ARG:NE	1:D:843:GLN:N	2.07	1.01
2:F:126:PHE:HA	2:F:209:ALA:O	1.58	1.01
1:A:608:ASP:HB3	1:A:779:LEU:HD13	1.02	1.01

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1068:ASP:H	1:A:1071:ARG:CG	1.73	1.01
2:C:403:PHE:CE1	2:C:415:PRO:HG3	1.96	1.01
1:D:1141:VAL:HB	1:D:1146:ARG:HE	0.87	1.01
1:D:195:LEU:H	1:D:265:LEU:HD22	1.25	1.01
1:D:420:HIS:HD1	1:D:421:PRO:HD2	0.85	1.01
1:D:610:PHE:H	1:D:611:PRO:CD	1.73	1.01
1:D:249:VAL:HG13	1:D:252:GLY:N	1.75	1.01
1:D:483:PRO:O	1:D:486:TRP:HB2	1.61	1.01
1:A:369:PRO:HB2	1:A:372:GLU:HG3	1.03	1.01
2:C:200:VAL:HG21	2:C:204:LEU:HD22	1.42	1.01
2:C:386:ASP:HB3	2:C:442:LEU:HD23	1.38	1.01
1:D:503:VAL:O	1:D:505:LYS:N	1.92	1.01
1:D:179:GLY:HA3	1:D:182:GLY:CA	1.90	1.01
1:D:870:VAL:O	1:D:873:GLU:HG2	1.60	1.01
2:F:431:LYS:HA	2:F:434:GLU:OE1	1.60	1.01
1:D:439:GLN:HE21	1:D:835:GLY:HA3	1.23	1.01
1:D:612:LEU:HD21	1:D:622:TYR:CE1	1.95	1.01
1:D:736:ASN:HA	1:D:750:PHE:CE1	1.95	1.01
1:D:904:ASP:HB3	1:D:912:GLY:HA3	1.38	1.01
1:D:932:HIS:O	1:D:935:THR:HG22	1.59	1.01
2:F:446:THR:HB	2:F:450:ASN:CB	1.90	1.01
1:A:222:SER:HB2	1:A:223:TRP:CD1	1.96	1.01
1:A:408:GLN:O	1:A:412:PRO:HD3	1.61	1.01
1:A:615:SER:HB2	1:A:725:PRO:HB3	1.42	1.01
2:C:341:ASP:O	2:C:342:LEU:HD13	1.60	1.01
1:A:1060:LYS:HD3	1:A:1073:PRO:CB	1.90	1.01
2:C:277:ASP:OD2	2:C:287:ASN:HB3	1.59	1.01
2:C:454:HIS:HB3	2:C:466:MET:HB2	1.43	1.01
1:D:1019:TRP:CE3	1:D:1168:LEU:HG	1.94	1.00
1:D:1057:MET:HB3	1:D:1064:ILE:HD11	1.41	1.00
2:F:245:PRO:O	2:F:248:SER:HB2	1.61	1.00
2:F:387:VAL:HG12	2:F:443:VAL:HG21	1.43	1.00
2:E:82:SER:H	2:E:85:GLN:HG3	1.26	1.00
1:A:1179:PHE:HB3	1:A:1220:ILE:HG12	1.43	1.00
1:A:118:VAL:N	1:A:119:PRO:HD3	1.73	1.00
1:A:615:SER:CB	1:A:725:PRO:HB3	1.91	1.00
2:C:393:LEU:HD23	2:C:396:ARG:HG2	1.42	1.00
1:D:1131:ILE:HD11	1:D:1138:ARG:CA	1.91	1.00
2:F:190:LEU:O	2:F:193:TYR:HB3	1.62	1.00
2:F:241:TRP:CB	2:F:336:LEU:HB2	1.90	1.00
2:E:78:PHE:HA	2:E:102:LEU:HB3	1.41	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:304:TRP:H	2:F:337:SER:HB2	0.84	1.00
1:A:175:TRP:HB2	1:A:184:ALA:HA	1.44	1.00
1:A:420:HIS:CD2	1:A:1020:ILE:HG21	1.96	1.00
2:B:268:MET:H	2:B:270:PRO:HD2	1.19	1.00
1:A:639:THR:HB	1:A:640:LEU:HD23	1.39	1.00
1:D:275:ARG:NH1	1:D:843:GLN:HG2	1.76	1.00
2:E:331:VAL:HG22	2:E:333:PRO:HD3	1.42	1.00
1:D:1006:ARG:HH12	1:D:1009:ASN:HB2	0.83	1.00
1:D:269:HIS:O	1:D:294:THR:HG23	1.61	1.00
1:D:441:TRP:CE2	1:D:879:GLN:HA	1.97	1.00
2:E:432:TYR:HA	2:E:435:MET:CB	1.91	1.00
2:F:375:HIS:H	2:F:378:LEU:HD22	1.25	1.00
1:A:866:ARG:HG2	1:A:1069:ILE:HG21	1.42	1.00
2:E:432:TYR:HB2	2:E:437:ILE:HD11	1.00	1.00
2:F:125:VAL:HG22	2:F:208:LEU:CA	1.91	1.00
1:A:1047:ARG:HH11	1:A:1047:ARG:HA	1.20	1.00
2:C:234:LYS:HZ3	2:C:344:ARG:HD2	1.23	1.00
2:C:234:LYS:NZ	2:C:344:ARG:HD2	1.74	1.00
1:D:1197:CYS:H	1:D:1204:THR:HG23	1.26	1.00
1:D:374:ARG:HD3	1:D:392:LEU:HD21	1.41	1.00
2:F:301:GLU:CB	2:F:339:ASN:HB3	1.90	1.00
1:A:267:VAL:HG23	1:A:291:PHE:CD2	1.97	1.00
1:A:213:VAL:HG12	1:A:401:TRP:HE1	0.83	1.00
1:A:436:PRO:HG2	1:A:879:GLN:HB3	1.44	1.00
2:C:343:ASP:C	2:C:347:LEU:HD13	1.81	1.00
1:D:1165:ALA:CB	1:D:1173:LEU:HD12	1.90	1.00
1:D:946:ALA:HA	1:D:949:PHE:CE2	1.95	1.00
2:C:356:LEU:HB3	2:C:359:ASN:HD21	1.21	1.00
1:A:175:TRP:HB2	1:A:184:ALA:CA	1.92	1.00
2:C:301:GLU:CB	2:C:339:ASN:HB3	1.90	1.00
1:D:522:GLY:O	1:D:524:PRO:HD3	1.62	1.00
2:C:134:LYS:HB3	2:C:180:LYS:HD3	1.43	1.00
1:D:251:THR:OG1	1:D:254:SER:HA	1.62	0.99
2:E:128:VAL:HG13	2:E:210:GLN:HE21	1.24	0.99
2:E:283:GLY:HA3	2:E:306:LEU:HA	1.44	0.99
2:F:386:ASP:HB2	2:F:442:LEU:CA	1.90	0.99
2:F:454:HIS:HB3	2:F:466:MET:HB2	1.42	0.99
1:A:1141:VAL:HB	1:A:1146:ARG:HD2	1.44	0.99
1:A:871:GLY:C	1:A:873:GLU:N	2.10	0.99
1:D:504:LYS:CD	1:D:523:ASP:HB3	1.90	0.99
2:B:453:ILE:HD12	2:B:468:ILE:HB	1.43	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:242:ALA:CB	1:D:245:ILE:HG12	1.92	0.99
1:D:775:GLU:HB3	1:D:796:LYS:HE2	1.44	0.99
2:E:436:SER:C	2:E:437:ILE:HD12	1.82	0.99
2:F:468:ILE:O	2:F:472:LYS:HG2	1.59	0.99
1:A:792:LEU:HD12	1:A:793:GLU:H	1.22	0.99
2:C:325:ARG:HG3	2:C:325:ARG:HH11	1.23	0.99
2:B:285:LYS:HG2	2:B:286:GLY:H	0.85	0.99
2:C:387:VAL:HG11	2:C:399:CYS:HB3	1.44	0.99
1:D:1151:LEU:O	1:D:1155:ILE:HG12	1.63	0.99
1:D:1198:LYS:HA	1:D:1204:THR:OG1	1.59	0.99
2:F:473:ASP:HA	2:F:476:ILE:HG23	1.45	0.99
1:A:378:VAL:CB	1:A:381:THR:HG23	1.93	0.99
2:B:396:ARG:HD2	2:B:417:TYR:HB2	1.44	0.99
2:F:325:ARG:HG3	2:F:325:ARG:HH11	1.26	0.99
2:B:244:PRO:O	2:B:248:SER:HB2	1.61	0.99
2:B:127:PRO:HB2	2:C:104:VAL:HG11	1.41	0.99
1:A:634:LEU:HB3	1:A:635:PRO:HA	1.45	0.99
1:D:775:GLU:O	1:D:776:ASP:HB3	1.59	0.99
2:F:131:LEU:HG	2:F:133:HIS:CE1	1.97	0.99
1:A:623:LEU:HB3	1:A:667:GLN:HE22	1.24	0.99
2:B:374:LEU:HB2	2:B:458:ARG:HH12	1.26	0.99
1:A:152:GLU:HA	1:A:155:ASN:HD22	1.26	0.99
1:D:128:PRO:CA	1:D:130:LEU:HD21	1.91	0.99
1:D:641:GLU:C	1:D:645:VAL:HG21	1.81	0.99
2:F:244:PRO:HD2	2:F:247:THR:HG21	1.42	0.99
1:A:210:THR:O	1:A:211:LEU:HB2	1.62	0.99
1:D:490:TRP:CZ2	1:D:598:VAL:HB	1.97	0.99
1:D:74:LEU:CD1	1:D:77:GLN:HA	1.93	0.99
1:A:316:LYS:HD2	1:A:350:ILE:HG12	1.40	0.99
1:A:655:LEU:HB2	1:A:719:LEU:HB3	1.44	0.99
1:A:962:ALA:HB2	1:A:986:TYR:OH	1.60	0.99
1:D:1019:TRP:HE3	1:D:1168:LEU:HG	1.21	0.99
1:D:441:TRP:NE1	1:D:879:GLN:HA	1.74	0.99
2:F:102:LEU:HD11	2:F:435:MET:HE1	1.41	0.99
1:A:1057:MET:O	1:A:1064:ILE:HG21	1.62	0.99
1:A:624:VAL:O	1:A:626:GLY:N	1.95	0.99
1:A:793:GLU:HA	1:A:796:LYS:HD3	1.44	0.99
2:C:466:MET:SD	2:C:471:LEU:HD23	2.03	0.99
1:D:270:ASN:HA	1:D:294:THR:CG2	1.92	0.98
1:D:375:GLU:O	1:D:379:LYS:N	1.96	0.98
1:D:608:ASP:HB3	1:D:620:TRP:CE3	1.98	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:741:ASP:HB2	1:D:748:TRP:HH2	1.28	0.98
1:A:192:GLU:CB	1:A:217:PRO:HG2	1.93	0.98
2:C:396:ARG:HH11	2:C:397:GLN:HA	1.26	0.98
1:D:1005:VAL:HA	1:D:1008:LEU:HD13	1.41	0.98
1:D:114:GLY:C	1:D:116:PRO:HD3	1.83	0.98
1:D:994:TRP:O	1:D:995:TYR:HB3	1.62	0.98
2:F:236:GLU:CA	2:F:341:ASP:HB3	1.93	0.98
2:F:454:HIS:HD2	2:F:468:ILE:HA	1.24	0.98
1:A:879:GLN:HB2	1:A:886:LEU:HD11	1.45	0.98
1:D:1193:VAL:HG22	1:D:1194:THR:H	0.85	0.98
1:A:272:SER:HB3	1:A:273:PHE:HD2	1.27	0.98
2:C:468:ILE:O	2:C:472:LYS:HG2	1.60	0.98
2:F:317:GLY:CA	2:F:320:SER:HB2	1.92	0.98
1:D:245:ILE:HG13	1:D:246:PRO:CD	1.92	0.98
1:D:375:GLU:CG	1:D:379:LYS:HB3	1.93	0.98
1:D:551:LYS:N	1:D:551:LYS:HD2	1.63	0.98
1:D:73:PRO:HG3	1:D:90:GLN:HG2	1.43	0.98
1:A:192:GLU:OE2	1:A:194:ALA:N	1.97	0.98
1:A:869:ARG:HG2	1:A:870:VAL:N	1.71	0.98
1:A:382:MET:SD	1:A:383:LYS:N	2.36	0.98
1:A:621:GLY:HA2	1:A:748:TRP:HB3	1.42	0.98
2:B:308:ASP:HB3	2:B:309:HIS:ND1	1.79	0.98
2:C:203:ARG:HA	2:C:325:ARG:CD	1.92	0.98
2:F:372:LEU:HD21	2:F:435:MET:C	1.84	0.98
2:F:454:HIS:CD2	2:F:468:ILE:HA	1.97	0.98
1:A:222:SER:HB2	1:A:223:TRP:HD1	1.26	0.98
1:A:273:PHE:HA	1:A:276:ALA:CB	1.94	0.98
1:A:996:ARG:HB2	1:A:1004:LEU:HD22	1.44	0.98
2:B:324:GLY:N	2:B:331:VAL:O	1.95	0.98
2:B:70:LEU:HD22	2:B:88:ARG:HH11	1.24	0.98
2:C:247:THR:HB	2:C:251:TRP:CD1	1.99	0.98
1:D:886:LEU:C	1:D:1146:ARG:HH12	1.67	0.98
1:A:866:ARG:CG	1:A:1069:ILE:HG21	1.94	0.98
1:A:606:THR:HG23	1:A:614:TYR:N	1.79	0.98
1:A:771:LEU:HD22	1:A:771:LEU:H	1.28	0.98
1:A:815:TRP:HA	1:A:838:GLY:HA3	1.44	0.98
2:F:85:GLN:HB3	2:F:95:CYS:O	1.62	0.98
2:B:211:ILE:HA	2:B:237:ALA:HA	1.46	0.98
1:D:1055:SER:CA	1:D:1058:PHE:HB3	1.93	0.98
1:D:1069:ILE:HG23	1:D:1070:PRO:HD2	1.46	0.98
2:E:262:TRP:O	2:E:265:LYS:HB3	1.64	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:303:LEU:HA	2:F:337:SER:OG	1.63	0.98
2:F:236:GLU:HA	2:F:341:ASP:CB	1.93	0.98
1:A:1204:THR:O	1:A:1204:THR:HG22	1.61	0.98
1:A:188:ALA:O	1:A:189:ILE:HG23	1.62	0.98
1:A:196:VAL:HG13	1:A:269:HIS:CD2	1.99	0.98
2:C:372:LEU:HD11	2:C:436:SER:CB	1.94	0.98
2:C:473:ASP:HA	2:C:476:ILE:CG2	1.93	0.98
1:D:128:PRO:HA	1:D:130:LEU:HD21	0.99	0.98
1:D:275:ARG:HH11	1:D:843:GLN:NE2	1.60	0.98
1:D:459:MET:N	1:D:460:LYS:HZ1	1.62	0.98
1:D:926:SER:HB2	1:D:930:ASP:HB2	1.42	0.98
2:E:107:ARG:HG3	2:E:107:ARG:NH1	1.64	0.98
2:F:342:LEU:HD23	2:F:345:GLY:H	1.26	0.98
2:F:385:LEU:O	2:F:416:GLY:HA3	1.64	0.98
2:F:372:LEU:CD1	2:F:436:SER:HB2	1.94	0.98
1:A:297:MET:SD	1:A:411:LEU:HB2	2.04	0.98
1:A:914:THR:HA	1:A:918:TRP:HB3	1.42	0.98
2:C:363:ARG:HH11	2:C:363:ARG:HA	1.28	0.98
2:C:82:SER:CA	2:C:86:LEU:HD21	1.94	0.98
1:D:345:TRP:HA	1:D:1040:LYS:NZ	1.78	0.97
1:D:428:LEU:HD12	1:D:428:LEU:N	1.78	0.97
1:A:498:LYS:HE2	1:A:499:LYS:H	1.29	0.97
2:C:276:SER:O	2:C:287:ASN:HB2	1.62	0.97
2:C:317:GLY:HA3	2:C:320:SER:HB2	1.46	0.97
1:D:1124:ALA:HB1	1:D:1148:ARG:NH1	1.79	0.97
2:F:102:LEU:HD11	2:F:435:MET:CE	1.92	0.97
2:F:300:ILE:HG22	2:F:301:GLU:HG3	1.44	0.97
1:A:1207:GLU:HB3	1:A:1213:PRO:HG3	1.42	0.97
1:D:1200:PRO:HG2	1:D:1203:PRO:HG2	1.44	0.97
1:D:959:GLN:HB2	1:D:960:PRO:CD	1.93	0.97
2:E:407:LEU:CD2	2:E:412:SER:HA	1.94	0.97
1:A:959:GLN:HB2	1:A:960:PRO:HD3	0.99	0.97
1:A:996:ARG:HH12	1:A:1003:TRP:HB2	1.28	0.97
1:D:869:ARG:O	1:D:1200:PRO:HG3	1.63	0.97
1:D:1208:ARG:HH12	2:F:285:LYS:HD3	1.27	0.97
1:D:138:HIS:O	1:D:142:LEU:HG	1.64	0.97
2:F:386:ASP:CB	2:F:442:LEU:HA	1.93	0.97
2:B:106:LEU:HD12	2:B:109:ASN:HD22	1.27	0.97
2:C:241:TRP:CB	2:C:336:LEU:HB2	1.95	0.97
2:C:439:PHE:CD1	2:C:458:ARG:HG3	2.00	0.97
1:D:219:ALA:O	1:D:220:TRP:HB2	1.61	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:203:LEU:C	1:A:205:GLU:H	1.44	0.97
1:A:253:ALA:CB	1:A:280:GLU:HB2	1.94	0.97
2:B:101:PRO:CA	2:C:128:VAL:HG21	1.95	0.97
1:D:513:LEU:H	1:D:570:PRO:HG3	1.25	0.97
2:E:456:ARG:CZ	2:E:463:LYS:HE3	1.95	0.97
2:F:82:SER:CA	2:F:86:LEU:HD21	1.93	0.97
1:A:392:LEU:HG	1:A:395:TYR:CG	1.98	0.97
1:A:76:ILE:CB	1:A:911:HIS:NE2	2.26	0.97
1:A:526:ASP:O	1:A:527:GLN:HB2	1.64	0.97
1:D:1001:GLY:O	1:D:1004:LEU:HB2	1.64	0.97
1:D:1128:ARG:HG3	1:D:1129:PHE:H	1.28	0.97
1:D:1231:LEU:HD11	1:D:1236:GLN:NE2	1.80	0.97
1:D:656:TYR:HB3	1:D:744:ILE:HG21	1.47	0.97
1:D:876:ALA:HA	1:D:1189:LEU:HB2	1.47	0.97
1:A:308:GLN:NE2	1:A:1091:GLU:H	1.61	0.97
1:A:849:THR:HG22	1:A:851:THR:H	1.29	0.97
2:C:301:GLU:HB3	2:C:339:ASN:CB	1.93	0.97
2:C:419:GLU:OE2	2:C:428:LEU:HD13	1.64	0.97
1:D:115:GLN:N	1:D:116:PRO:HD3	1.76	0.97
1:D:1197:CYS:N	1:D:1204:THR:HG23	1.80	0.97
2:E:442:LEU:HD13	2:E:443:VAL:N	1.78	0.97
1:A:175:TRP:CB	1:A:184:ALA:HA	1.94	0.97
2:C:245:PRO:O	2:C:248:SER:HB2	1.63	0.97
1:D:275:ARG:HD2	1:D:843:GLN:CD	1.83	0.97
2:E:377:CYS:C	2:E:380:PRO:HD2	1.83	0.97
1:A:927:ARG:O	1:A:930:ASP:HB3	1.65	0.97
1:A:755:LYS:HD3	1:A:964:ARG:NH1	1.80	0.97
2:C:127:PRO:CD	2:C:210:GLN:HB2	1.94	0.97
1:D:350:ILE:H	1:D:350:ILE:HD12	1.30	0.97
1:D:601:LYS:HE3	1:D:721:ALA:HB1	1.45	0.97
1:D:618:HIS:HB2	1:D:725:PRO:HD2	1.47	0.97
1:D:996:ARG:NE	1:D:996:ARG:H	1.61	0.97
1:A:1002:GLU:O	1:A:1005:VAL:HG22	1.63	0.97
1:A:472:GLN:HG3	2:C:460:THR:HA	1.47	0.97
1:A:76:ILE:CG2	1:A:77:GLN:H	1.78	0.97
2:C:236:GLU:CA	2:C:341:ASP:HB3	1.95	0.97
2:C:441:VAL:HG11	2:C:454:HIS:CE1	1.99	0.97
1:D:495:PHE:O	1:D:496:LYS:HD2	1.64	0.96
1:D:801:TRP:CE3	1:D:870:VAL:CG1	2.46	0.96
2:E:264:ARG:HE	2:E:273:PHE:HB2	1.28	0.96
1:D:279:ARG:HD2	1:D:841:LEU:CB	1.95	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:302:SER:OG	1:D:351:SER:HB3	1.64	0.96
1:D:900:ALA:CB	1:D:915:ALA:HB1	1.94	0.96
1:A:1026:ARG:O	1:A:1030:ARG:N	1.95	0.96
1:A:225:SER:HA	1:A:228:LEU:HD13	1.46	0.96
1:D:1069:ILE:CG2	1:D:1070:PRO:HD2	1.95	0.96
1:D:375:GLU:HG3	1:D:379:LYS:CB	1.93	0.96
1:D:624:VAL:O	1:D:626:GLY:N	1.99	0.96
2:E:254:PHE:HB2	2:E:257:ARG:HH11	1.26	0.96
1:A:1198:LYS:HG3	1:A:1204:THR:CG2	1.94	0.96
1:A:166:PRO:CG	1:A:398:GLN:HE22	1.76	0.96
1:D:279:ARG:HD2	1:D:841:LEU:HB3	0.99	0.96
1:D:887:VAL:HG23	1:D:1146:ARG:CZ	1.95	0.96
1:A:811:GLN:HB2	1:A:840:ILE:HG21	1.45	0.96
2:C:344:ARG:HA	2:C:347:LEU:HB2	1.45	0.96
2:C:446:THR:HA	2:C:450:ASN:HB2	1.43	0.96
1:D:1198:LYS:CA	1:D:1204:THR:OG1	2.14	0.96
1:D:465:ASP:O	1:D:469:ASP:HB2	1.66	0.96
1:D:549:LEU:N	1:D:551:LYS:NZ	2.14	0.96
1:D:279:ARG:HB3	1:D:841:LEU:HB2	1.46	0.96
2:F:127:PRO:HD3	2:F:210:GLN:N	1.80	0.96
2:B:396:ARG:CG	2:B:417:TYR:HB2	1.93	0.96
1:D:101:ARG:HH11	1:D:101:ARG:HG3	1.30	0.96
1:A:589:PRO:HG3	1:A:805:HIS:CD2	1.99	0.96
1:D:1007:GLU:O	1:D:1011:PRO:HD3	1.65	0.96
1:D:798:ILE:HA	1:D:869:ARG:CZ	1.95	0.96
1:D:993:ARG:N	1:D:996:ARG:HD2	1.79	0.96
2:F:466:MET:HB3	2:F:471:LEU:CD2	1.96	0.96
1:A:1038:TRP:O	1:A:1042:GLU:HG2	1.65	0.96
1:A:211:LEU:CD1	1:A:221:TYR:HB3	1.96	0.96
1:A:605:LEU:HA	1:A:781:ALA:HB3	1.47	0.96
2:C:417:TYR:HD2	2:C:417:TYR:N	1.64	0.96
1:D:193:ARG:HA	1:D:193:ARG:CZ	1.96	0.96
1:D:866:ARG:HG3	1:D:1069:ILE:CG2	1.95	0.96
2:E:380:PRO:O	2:E:382:LYS:HB2	1.65	0.96
1:A:666:GLN:O	1:A:670:PRO:CD	2.14	0.96
2:C:239:LEU:H	2:C:338:VAL:HG12	1.18	0.96
1:D:1067:SER:CA	1:D:1071:ARG:HA	1.96	0.96
2:E:419:GLU:HA	2:F:203:ARG:NH2	1.81	0.96
1:A:493:GLN:O	1:A:494:GLU:HG2	1.65	0.96
2:C:317:GLY:HA3	2:C:320:SER:H	1.29	0.96
1:D:1069:ILE:CB	1:D:1070:PRO:HD2	1.96	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:285:GLN:NE2	1:D:1128:ARG:HG2	1.80	0.96
1:D:559:LEU:HG	1:D:560:PRO:HD2	1.45	0.96
1:D:604:ALA:HA	1:D:613:HIS:HE1	1.24	0.96
2:F:309:HIS:HA	2:F:312:LEU:HD12	1.43	0.96
2:F:455:LEU:HA	2:F:456:ARG:HH21	1.27	0.96
1:D:597:ARG:C	1:D:616:GLU:OE1	2.05	0.95
1:D:605:LEU:O	1:D:781:ALA:HA	1.65	0.95
1:A:660:CYS:SG	1:A:661:LEU:N	2.39	0.95
2:C:454:HIS:HA	2:C:468:ILE:HD12	1.47	0.95
2:C:317:GLY:CA	2:C:320:SER:HB2	1.95	0.95
2:F:114:TRP:HE1	2:F:259:ARG:CZ	1.78	0.95
2:F:241:TRP:N	2:F:336:LEU:CB	2.29	0.95
2:F:241:TRP:CD1	2:F:243:THR:HG23	2.01	0.95
1:D:1208:ARG:HH22	2:F:285:LYS:CD	1.78	0.95
2:F:423:SER:OG	2:F:428:LEU:HD21	1.66	0.95
2:F:453:ILE:HG22	2:F:455:LEU:H	1.30	0.95
2:B:261:GLN:NE2	2:B:264:ARG:HH21	1.62	0.95
2:B:311:LEU:CG	2:B:312:LEU:H	1.78	0.95
2:B:86:LEU:HD23	2:B:96:HIS:HB2	1.48	0.95
1:D:1138:ARG:HH11	1:D:1138:ARG:HG2	1.29	0.95
1:D:849:THR:HB	1:D:852:ARG:H	1.31	0.95
2:B:301:GLU:HA	2:B:340:GLY:HA3	1.47	0.95
2:C:431:LYS:NZ	2:C:435:MET:HG2	1.80	0.95
1:D:1023:GLN:OE1	1:D:1100:VAL:HA	1.64	0.95
1:D:241:PRO:O	1:D:244:LEU:HG	1.65	0.95
1:D:203:LEU:N	1:D:380:GLY:HA2	1.79	0.95
2:F:134:LYS:HB3	2:F:180:LYS:HD3	1.49	0.95
1:A:176:THR:CA	1:A:218:SER:HB2	1.96	0.95
1:A:758:ASN:HB2	1:A:1058:PHE:CZ	2.00	0.95
1:A:927:ARG:HG2	1:A:928:GLY:N	1.81	0.95
2:B:260:LEU:C	2:B:264:ARG:HD3	1.86	0.95
2:E:187:HIS:HA	2:E:190:LEU:HD22	1.47	0.95
2:E:181:LEU:HB3	2:E:215:PHE:HE1	1.28	0.95
2:F:127:PRO:HD3	2:F:209:ALA:C	1.87	0.95
2:F:350:LEU:HD12	2:F:350:LEU:N	1.80	0.95
2:F:435:MET:HE2	2:F:435:MET:HA	1.46	0.95
1:D:1006:ARG:HH12	1:D:1009:ASN:CB	1.79	0.95
1:D:576:LEU:CG	1:D:577:CYS:H	1.80	0.95
1:A:1218:LEU:O	1:A:1219:ASP:HB2	1.64	0.95
1:A:169:TRP:CZ3	1:A:391:ASP:HA	2.01	0.95
2:B:372:LEU:HD13	2:B:434:GLU:HA	1.45	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:206:TYR:HD2	2:C:206:TYR:H	0.96	0.95
2:C:466:MET:CB	2:C:471:LEU:HD22	1.97	0.95
1:D:486:TRP:HZ2	1:D:722:ARG:HA	1.31	0.95
1:A:1131:ILE:HD13	1:A:1138:ARG:N	1.82	0.95
1:A:490:TRP:HH2	1:A:598:VAL:HB	1.28	0.95
1:A:549:LEU:HD21	2:C:398:VAL:CB	1.97	0.95
2:E:255:TRP:O	2:E:259:ARG:HB2	1.66	0.95
2:E:382:LYS:HG2	2:E:412:SER:CB	1.96	0.95
2:E:449:GLU:HA	2:E:449:GLU:OE1	1.61	0.95
1:A:1131:ILE:HD11	1:A:1133:ILE:HG22	1.49	0.95
1:A:996:ARG:NH2	1:A:1003:TRP:HB3	1.81	0.95
1:A:203:LEU:C	1:A:205:GLU:N	2.19	0.95
1:A:194:ALA:HA	1:A:265:LEU:HB2	1.45	0.95
1:A:580:LEU:HD12	1:A:581:ASP:H	1.32	0.95
2:E:431:LYS:HA	2:E:434:GLU:CG	1.95	0.95
1:A:853:ARG:HH12	1:A:1105:ALA:HB3	1.29	0.95
1:A:176:THR:HG23	1:A:177:ARG:H	1.31	0.95
1:A:210:THR:HB	1:A:224:CYS:HB2	0.95	0.95
1:D:513:LEU:HB2	1:D:568:GLY:HA2	1.45	0.95
1:D:145:LYS:O	1:D:149:PRO:HD2	1.67	0.94
1:D:621:GLY:HA2	1:D:748:TRP:CB	1.96	0.94
1:D:441:TRP:HE1	1:D:880:ALA:H	1.14	0.94
2:E:117:SER:HA	2:E:121:PHE:CE1	2.01	0.94
1:A:1069:ILE:HG22	1:A:1070:PRO:HD3	1.47	0.94
1:A:348:LEU:HA	1:A:350:ILE:HD13	1.46	0.94
1:A:454:GLU:O	1:A:458:GLU:HG3	1.67	0.94
2:B:323:HIS:CA	2:B:332:VAL:HA	1.97	0.94
1:D:429:GLU:HG3	1:D:1128:ARG:NH2	1.82	0.94
1:D:607:TRP:HD1	1:D:780:GLN:H	1.00	0.94
1:D:914:THR:O	1:D:918:TRP:CB	2.15	0.94
1:A:756:ASP:CB	1:A:1057:MET:HG2	1.97	0.94
1:A:170:ALA:CB	1:A:175:TRP:HZ2	1.79	0.94
1:A:272:SER:HA	1:A:275:ARG:HB2	1.49	0.94
1:D:452:TYR:CB	1:D:801:TRP:HE1	1.79	0.94
1:D:490:TRP:HE3	1:D:580:LEU:HD22	1.29	0.94
1:D:813:VAL:HB	1:D:840:ILE:HG13	1.49	0.94
1:D:824:ALA:HA	1:D:827:ARG:CG	1.96	0.94
2:C:467:HIS:O	2:C:471:LEU:CD1	2.15	0.94
1:A:645:VAL:HG13	1:A:650:ARG:HB2	1.48	0.94
1:D:1118:TRP:O	1:D:1121:GLU:HG3	1.68	0.94
1:D:259:ARG:HA	1:D:259:ARG:NH1	1.80	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:447:THR:CG2	2:E:453:ILE:HG23	1.97	0.94
2:F:214:CYS:SG	2:F:236:GLU:HG3	2.07	0.94
1:A:1236:GLN:HB3	1:A:1237:PRO:CD	1.97	0.94
1:A:225:SER:OG	1:A:386:ARG:HB2	1.67	0.94
1:A:378:VAL:C	1:A:380:GLY:N	2.19	0.94
1:D:1069:ILE:O	1:D:1071:ARG:HG3	1.66	0.94
1:D:79:LEU:HD23	1:D:83:LEU:HD12	1.49	0.94
2:E:456:ARG:HD3	2:E:463:LYS:HG3	1.48	0.94
2:C:471:LEU:HD12	2:C:471:LEU:N	1.82	0.94
1:D:1029:GLN:HG2	1:D:1030:ARG:NE	1.82	0.94
1:D:933:SER:HB2	1:D:943:ARG:HD3	1.45	0.94
1:D:922:GLN:HG2	1:D:931:LEU:HD13	1.48	0.94
2:E:266:PHE:HB3	2:E:376:PRO:CD	1.96	0.94
2:E:425:LEU:HB3	2:E:426:GLU:OE2	1.68	0.94
2:F:466:MET:SD	2:F:471:LEU:HD23	2.08	0.94
1:A:515:ILE:HG22	1:A:516:GLU:H	1.26	0.94
2:C:190:LEU:O	2:C:193:TYR:HB3	1.67	0.94
2:C:303:LEU:CA	2:C:337:SER:OG	2.14	0.94
1:A:784:GLY:HA2	2:C:362:THR:HG1	1.20	0.94
2:C:439:PHE:HB3	2:C:457:SER:O	1.67	0.94
2:E:264:ARG:NE	2:E:273:PHE:HB2	1.82	0.94
1:D:1155:ILE:HA	1:D:1158:LEU:HD21	1.49	0.94
1:D:274:ASP:OD2	1:D:293:ASP:HA	1.66	0.94
1:D:577:CYS:C	1:D:579:ARG:H	1.70	0.94
2:F:125:VAL:HG23	2:F:126:PHE:N	1.81	0.94
2:B:343:ASP:HB3	2:B:344:ARG:HE	1.33	0.94
2:B:105:GLU:HB3	2:B:380:PRO:HB3	1.50	0.94
1:A:1001:GLY:O	1:A:1005:VAL:HG13	1.67	0.94
2:C:200:VAL:CG2	2:C:204:LEU:HD22	1.98	0.94
2:C:127:PRO:HD3	2:C:210:GLN:N	1.81	0.94
2:C:69:LEU:HD23	2:C:351:TYR:CE1	2.02	0.94
2:C:443:VAL:HA	2:C:447:THR:CG2	1.98	0.94
1:D:735:GLY:HA3	1:D:749:PHE:CB	1.97	0.94
2:E:430:SER:O	2:E:433:ASP:HB3	1.65	0.94
1:A:490:TRP:CH2	1:A:598:VAL:HB	2.02	0.94
2:C:357:THR:O	2:C:358:GLU:HG2	1.68	0.94
1:D:455:LEU:HD12	1:D:869:ARG:NH1	1.83	0.94
1:A:1023:GLN:NE2	1:A:1103:SER:OG	2.01	0.94
1:A:288:ARG:HE	1:A:288:ARG:CA	1.81	0.94
1:A:893:SER:HB2	1:A:896:LEU:CB	1.97	0.94
2:C:125:VAL:HG22	2:C:208:LEU:CA	1.97	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:623:LEU:HB3	1:D:667:GLN:NE2	1.83	0.94
1:A:640:LEU:H	1:A:640:LEU:HD23	1.33	0.94
1:D:887:VAL:N	1:D:1146:ARG:HH12	1.66	0.93
1:D:309:ARG:O	1:D:313:ILE:HG23	1.68	0.93
1:D:392:LEU:O	1:D:395:TYR:HB2	1.65	0.93
1:D:780:GLN:OE1	1:D:780:GLN:HA	1.67	0.93
2:F:353:SER:HB2	2:F:373:LYS:H	1.31	0.93
1:A:1128:ARG:CG	1:A:1129:PHE:H	1.77	0.93
1:A:201:VAL:HG22	1:A:208:CYS:O	1.69	0.93
1:A:166:PRO:HG3	1:A:398:GLN:NE2	1.83	0.93
1:A:897:TRP:O	1:A:900:ALA:HB3	1.68	0.93
2:B:72:ILE:O	2:B:75:ARG:HG3	1.67	0.93
1:D:1200:PRO:CG	1:D:1203:PRO:HG2	1.97	0.93
1:D:275:ARG:HH21	1:D:842:PRO:CB	1.81	0.93
1:D:356:LEU:HD13	1:D:369:LEU:HD22	1.50	0.93
1:D:407:PHE:O	1:D:411:LEU:HG	1.68	0.93
1:D:275:ARG:HH21	1:D:842:PRO:HB3	1.32	0.93
1:A:1006:ARG:NH1	1:A:1006:ARG:HA	1.83	0.93
1:A:298:HIS:CD2	1:A:358:LEU:HD22	2.02	0.93
2:B:114:TRP:HA	2:B:262:TRP:CZ2	2.03	0.93
2:F:238:SER:HA	2:F:338:VAL:HG11	1.48	0.93
2:F:289:LEU:HD12	2:F:301:GLU:HB2	1.48	0.93
2:F:391:PRO:HB2	2:F:394:GLU:OE2	1.68	0.93
1:A:1043:VAL:CG1	1:A:1046:GLU:HB3	1.97	0.93
1:A:1124:ALA:HB1	1:A:1148:ARG:NH1	1.82	0.93
1:A:196:VAL:HB	1:A:401:TRP:HZ3	1.29	0.93
1:A:485:LEU:HD12	1:A:489:GLU:OE2	1.69	0.93
1:A:893:SER:CB	1:A:896:LEU:HB2	1.98	0.93
2:C:322:LEU:HD13	2:C:322:LEU:O	1.68	0.93
1:A:566:LEU:HB3	1:A:567:PRO:HD3	1.51	0.93
2:E:277:ASP:HA	2:E:287:ASN:OD1	1.68	0.93
1:D:1204:THR:HA	1:D:1207:GLU:CB	1.99	0.93
1:D:606:THR:HG21	1:D:612:LEU:O	1.66	0.93
2:F:234:LYS:NZ	2:F:344:ARG:HD2	1.82	0.93
2:F:387:VAL:CA	2:F:443:VAL:HG22	1.97	0.93
1:A:582:ASP:N	1:A:583:PRO:HD2	1.83	0.93
2:C:376:PRO:HG3	2:C:482:ALA:HB1	1.49	0.93
1:D:1025:LEU:HD13	1:D:1027:LYS:HE2	1.47	0.93
1:D:1051:GLY:HA2	1:D:1054:GLU:HG3	1.49	0.93
1:D:1231:LEU:HG	1:D:1232:GLU:N	1.81	0.93
2:F:456:ARG:HG2	2:F:457:SER:H	1.33	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1026:ARG:HB2	1:A:1029:GLN:HB2	1.49	0.93
1:A:1131:ILE:HG23	1:A:1132:SER:H	1.31	0.93
1:A:925:LYS:HG3	1:A:926:SER:H	1.32	0.93
1:D:768:LYS:HA	1:D:772:PRO:HG3	1.51	0.93
1:A:309:ARG:O	1:A:313:ILE:HG23	1.68	0.93
2:B:396:ARG:CD	2:B:417:TYR:HB2	1.97	0.93
1:A:556:THR:HG23	2:C:452:LEU:HB3	1.50	0.93
1:D:1116:MET:HE3	1:D:1153:LEU:HA	1.46	0.93
2:F:203:ARG:HB3	2:F:204:LEU:CD1	1.97	0.93
1:A:744:ILE:HB	1:A:745:PRO:HD3	1.50	0.93
1:D:436:PRO:O	1:D:879:GLN:HB2	1.67	0.93
2:E:407:LEU:HD23	2:E:411:ILE:O	1.67	0.93
1:D:1206:MET:HG2	2:F:257:ARG:NH1	1.83	0.93
1:A:1057:MET:C	1:A:1064:ILE:HG21	1.89	0.93
1:A:606:THR:HG22	1:A:620:TRP:HE3	1.29	0.93
2:B:323:HIS:HB2	2:B:331:VAL:C	1.88	0.93
1:D:1231:LEU:HD11	1:D:1236:GLN:HE21	1.33	0.93
2:F:299:LEU:N	2:F:299:LEU:HD23	1.81	0.93
2:F:447:THR:O	2:F:452:LEU:HA	1.68	0.93
1:A:1188:CYS:C	1:A:1189:LEU:HD22	1.90	0.93
1:D:666:GLN:C	1:D:669:MET:O	2.07	0.93
1:D:1223:ILE:HG13	1:D:1224:ILE:N	1.83	0.93
1:D:580:LEU:HD11	1:D:593:SER:OG	1.67	0.93
1:D:608:ASP:OD1	1:D:611:PRO:HD2	1.69	0.93
1:D:549:LEU:HD21	2:F:398:VAL:HA	1.49	0.93
1:A:273:PHE:HA	1:A:276:ALA:HB2	1.46	0.93
1:D:509:THR:CB	1:D:512:LYS:HB3	1.97	0.93
1:D:897:TRP:CH2	1:D:1173:LEU:HD13	2.04	0.92
1:D:217:PRO:O	1:D:218:SER:OG	1.87	0.92
1:D:269:HIS:CD2	1:D:291:PHE:HD2	1.85	0.92
1:D:808:ILE:O	1:D:811:GLN:HB2	1.69	0.92
2:F:127:PRO:CD	2:F:128:VAL:H	1.78	0.92
2:F:301:GLU:CG	2:F:339:ASN:HB3	1.99	0.92
1:A:411:LEU:HD12	1:A:412:PRO:N	1.84	0.92
1:A:886:LEU:N	1:A:1146:ARG:NH2	2.17	0.92
2:C:454:HIS:HB2	2:C:468:ILE:HA	1.47	0.92
2:C:467:HIS:HB3	2:C:470:LYS:HE2	1.50	0.92
1:D:592:LEU:HA	1:D:595:GLN:CG	1.98	0.92
1:A:1131:ILE:HD13	1:A:1138:ARG:H	1.32	0.92
1:A:223:TRP:N	1:A:223:TRP:CD1	2.36	0.92
1:A:451:THR:OG1	1:A:1202:ASN:CG	2.07	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:350:LEU:N	2:C:350:LEU:HD12	1.82	0.92
1:D:270:ASN:HA	1:D:294:THR:HG23	1.49	0.92
1:D:371:LYS:HA	1:D:374:ARG:HD2	1.51	0.92
1:D:795:ASN:N	1:D:795:ASN:HD22	1.59	0.92
1:A:657:ARG:O	1:A:660:CYS:HB3	1.68	0.92
1:A:1030:ARG:CG	1:A:1040:LYS:HZ1	1.83	0.92
1:D:633:LYS:CB	1:D:638:THR:HG21	1.99	0.92
2:E:247:THR:HG22	2:E:250:GLN:HE21	1.32	0.92
1:A:434:TYR:O	1:A:435:LEU:HG	1.68	0.92
1:D:412:PRO:HG2	1:D:413:LEU:HG	1.51	0.92
2:E:431:LYS:C	2:E:433:ASP:H	1.67	0.92
2:F:453:ILE:HD13	2:F:465:MET:HB3	1.51	0.92
1:A:245:ILE:CG1	1:A:246:PRO:HD3	2.00	0.92
1:A:468:ASN:ND2	2:C:460:THR:HG21	1.84	0.92
2:C:303:LEU:C	2:C:337:SER:HB2	1.90	0.92
1:D:1131:ILE:HD12	1:D:1131:ILE:H	1.35	0.92
1:D:1208:ARG:NH1	2:F:285:LYS:HD3	1.83	0.92
1:D:127:LEU:HD22	1:D:129:PRO:HD3	1.51	0.92
1:D:438:ASN:O	1:D:441:TRP:HB2	1.69	0.92
1:D:275:ARG:NE	1:D:842:PRO:HA	1.84	0.92
1:A:112:LEU:HD23	1:A:920:THR:HB	1.50	0.92
1:A:125:LEU:H	1:A:125:LEU:CD2	1.82	0.92
1:A:169:TRP:HZ3	1:A:391:ASP:HA	1.32	0.92
2:C:341:ASP:CG	2:C:342:LEU:N	2.20	0.92
2:F:368:HIS:NE2	2:F:370:LYS:HD2	1.84	0.92
1:A:107:LEU:HB2	1:A:113:TRP:CD1	2.03	0.92
1:A:213:VAL:HA	1:A:221:TYR:HE2	1.35	0.92
2:C:209:ALA:HB1	2:C:238:SER:O	1.69	0.92
2:C:126:PHE:HA	2:C:209:ALA:O	1.70	0.92
2:F:317:GLY:HA2	2:F:320:SER:HB2	1.52	0.92
1:D:1027:LYS:HB3	1:D:1091:GLU:HB3	1.52	0.92
1:D:849:THR:HB	1:D:852:ARG:N	1.85	0.92
1:A:891:VAL:HA	1:A:1180:PHE:HE1	1.35	0.92
1:A:412:PRO:HG2	1:A:413:LEU:H	1.31	0.92
1:A:887:VAL:CG2	1:A:1146:ARG:HH11	1.83	0.92
2:F:222:LYS:O	2:F:223:GLN:HB2	1.70	0.92
1:D:1116:MET:CE	1:D:1153:LEU:HA	2.00	0.92
1:D:113:TRP:HE3	1:D:114:GLY:H	1.09	0.92
1:D:85:GLU:CD	1:D:130:LEU:HD22	1.90	0.92
1:D:283:LEU:HD23	1:D:839:ALA:HB3	1.52	0.92
1:D:1190:ARG:HD3	1:D:1195:MET:HE3	1.52	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:743:ASP:C	1:D:744:ILE:HD13	1.91	0.91
2:F:256:LEU:HD11	2:F:287:ASN:OD1	1.71	0.91
2:C:421:MET:HG3	2:C:422:GLN:N	1.85	0.91
1:D:873:GLU:HB3	1:D:1200:PRO:CD	2.00	0.91
1:D:470:ALA:O	1:D:474:LEU:HD13	1.71	0.91
1:A:195:LEU:CD2	1:A:217:PRO:HD2	2.01	0.91
1:D:513:LEU:H	1:D:570:PRO:CG	1.83	0.91
1:D:374:ARG:HB3	1:D:392:LEU:HD11	1.52	0.91
1:D:457:ARG:HA	1:D:460:LYS:HD2	1.52	0.91
1:D:272:SER:HB3	1:D:843:GLN:CD	1.91	0.91
1:A:1010:LEU:O	1:A:1096:ARG:NH2	2.02	0.91
2:C:342:LEU:HD23	2:C:345:GLY:H	1.33	0.91
1:D:350:ILE:O	1:D:361:ARG:HB2	1.70	0.91
1:D:424:LEU:O	1:D:428:LEU:HD11	1.69	0.91
2:E:383:VAL:O	2:E:384:ALA:CB	2.17	0.91
2:F:439:PHE:CD1	2:F:458:ARG:HB2	2.03	0.91
1:A:885:THR:C	1:A:886:LEU:HD23	1.91	0.91
2:C:303:LEU:HA	2:C:337:SER:HG	1.10	0.91
1:A:784:GLY:H	2:C:363:ARG:HD3	1.32	0.91
1:A:634:LEU:CB	1:A:635:PRO:HA	1.99	0.91
1:D:1218:LEU:HB3	1:D:1223:ILE:HG22	1.52	0.91
1:D:278:ILE:CG2	1:D:279:ARG:HD3	2.00	0.91
1:D:618:HIS:HB2	1:D:725:PRO:CD	2.00	0.91
1:D:73:PRO:CG	1:D:90:GLN:HG2	2.00	0.91
2:F:303:LEU:HA	2:F:337:SER:HG	1.32	0.91
1:A:228:LEU:CD1	1:A:386:ARG:HB3	2.01	0.91
1:A:618:HIS:HB2	1:A:725:PRO:CB	2.00	0.91
2:B:202:LYS:O	2:B:325:ARG:NH1	2.04	0.91
1:D:190:PRO:C	1:D:192:GLU:H	1.72	0.91
1:D:891:VAL:CG1	1:D:1135:ASP:HB3	2.00	0.91
2:E:128:VAL:HG13	2:E:210:GLN:NE2	1.84	0.91
2:F:125:VAL:CG2	2:F:208:LEU:HA	1.97	0.91
1:A:1060:LYS:CE	1:A:1065:ALA:H	1.82	0.91
1:A:240:SER:HA	1:A:243:ASP:OD2	1.70	0.91
1:A:316:LYS:CD	1:A:350:ILE:HG12	1.99	0.91
2:B:204:LEU:HD12	2:B:325:ARG:H	1.35	0.91
1:D:1002:GLU:O	1:D:1005:VAL:HG22	1.69	0.91
1:D:634:LEU:HD13	1:D:635:PRO:O	1.69	0.91
1:D:649:TYR:CA	1:D:652:ILE:HB	2.01	0.91
1:A:1046:GLU:O	1:A:1049:TRP:HD1	1.52	0.91
1:A:203:LEU:CD1	1:A:379:LYS:C	2.38	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:997:LEU:HD22	1:A:998:SER:N	1.85	0.91
2:C:454:HIS:N	2:C:466:MET:O	2.02	0.91
1:D:605:LEU:HD22	1:D:781:ALA:HB1	1.52	0.91
1:D:1209:ARG:HH22	2:F:252:LEU:HD22	1.33	0.91
1:A:996:ARG:NH1	1:A:1003:TRP:CB	2.33	0.91
1:A:374:ARG:HH11	1:A:374:ARG:HG2	1.35	0.91
1:A:498:LYS:CD	1:A:518:ALA:HB3	2.00	0.91
1:A:618:HIS:CB	1:A:725:PRO:HB2	2.00	0.91
2:C:241:TRP:HB2	2:C:336:LEU:HD12	0.94	0.91
2:C:306:LEU:N	2:C:335:VAL:HG11	1.85	0.91
2:C:302:THR:H	2:C:339:ASN:HA	1.36	0.91
1:A:784:GLY:CA	2:C:362:THR:OG1	2.18	0.91
2:C:471:LEU:HA	2:C:474:PHE:HB3	1.52	0.91
1:A:163:PRO:HB3	1:A:164:PRO:HD3	1.52	0.91
1:D:449:GLN:HA	1:D:452:TYR:HE2	1.32	0.91
2:E:428:LEU:CD1	2:E:428:LEU:H	1.83	0.91
2:F:342:LEU:HB2	2:F:344:ARG:CG	2.00	0.91
1:A:1200:PRO:HB2	1:A:1202:ASN:C	1.90	0.91
1:A:1236:GLN:H	1:A:1236:GLN:NE2	1.68	0.91
1:A:851:THR:HG22	1:A:853:ARG:H	1.36	0.91
2:C:435:MET:CE	2:C:435:MET:HA	2.01	0.91
2:C:81:GLY:HA3	2:C:97:PRO:HB2	1.53	0.91
2:B:211:ILE:HG12	2:B:237:ALA:HB1	1.51	0.91
1:D:630:ASN:HB2	1:D:744:ILE:C	1.91	0.91
1:D:436:PRO:O	1:D:879:GLN:CB	2.19	0.91
1:A:436:PRO:HD2	1:A:879:GLN:NE2	1.86	0.91
2:C:374:LEU:HB3	2:C:378:LEU:HB3	1.51	0.91
1:A:503:VAL:CG2	1:A:521:PRO:HA	2.01	0.91
1:A:774:MET:HA	1:A:777:GLY:HA3	1.52	0.91
1:D:1157:ASN:HB3	1:D:1161:ARG:NH1	1.85	0.90
1:D:297:MET:HG2	1:D:411:LEU:CD2	2.00	0.90
1:D:556:THR:O	1:D:559:LEU:HB3	1.71	0.90
2:E:422:GLN:HG2	2:E:424:SER:H	1.32	0.90
2:E:374:LEU:H	2:E:458:ARG:NH2	1.67	0.90
2:F:241:TRP:N	2:F:336:LEU:HB3	1.86	0.90
1:A:1191:LYS:O	1:A:1192:GLU:HB3	1.71	0.90
1:A:961:PHE:H	1:A:961:PHE:HD1	1.16	0.90
1:D:105:GLU:O	1:D:109:LYS:HG2	1.70	0.90
1:D:279:ARG:O	1:D:841:LEU:HG	1.69	0.90
2:F:241:TRP:HB2	2:F:336:LEU:HD12	0.91	0.90
1:A:1212:ILE:HG22	1:A:1213:PRO:N	1.81	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1131:ILE:N	1:D:1131:ILE:CD1	2.28	0.90
2:F:291:TYR:OH	2:F:355:GLN:HG3	1.71	0.90
1:A:1027:LYS:HB2	1:A:1092:PHE:CD2	2.05	0.90
1:A:477:GLU:HG2	1:A:482:ASP:CG	1.92	0.90
2:C:354:PHE:CE1	2:C:370:LYS:HD3	2.06	0.90
2:C:61:GLY:O	2:C:64:GLU:HG2	1.70	0.90
2:E:181:LEU:HB3	2:E:215:PHE:CE1	2.05	0.90
2:F:351:TYR:O	2:F:355:GLN:N	2.05	0.90
1:A:107:LEU:HB2	1:A:113:TRP:NE1	1.85	0.90
1:A:79:LEU:HB3	1:A:89:GLY:HA2	1.52	0.90
1:A:275:ARG:HH21	1:A:845:VAL:HA	1.23	0.90
2:C:454:HIS:HB3	2:C:466:MET:CB	2.01	0.90
1:D:503:VAL:C	1:D:505:LYS:H	1.74	0.90
1:D:74:LEU:HD12	1:D:77:GLN:HA	1.51	0.90
2:F:203:ARG:HB3	2:F:204:LEU:HD13	1.54	0.90
1:A:612:LEU:HB3	1:A:620:TRP:HB2	1.51	0.90
1:A:612:LEU:CD2	1:A:621:GLY:H	1.81	0.90
2:B:202:LYS:O	2:B:325:ARG:CD	2.18	0.90
1:D:371:LYS:HA	1:D:374:ARG:HD3	1.53	0.90
1:D:407:PHE:CD1	1:D:411:LEU:HD21	2.07	0.90
1:D:413:LEU:HA	1:D:416:GLU:OE1	1.70	0.90
1:A:87:ILE:HG22	1:A:88:PHE:N	1.86	0.90
2:B:347:LEU:O	2:B:350:LEU:HB3	1.71	0.90
2:E:325:ARG:HG2	2:E:330:ASN:HA	1.54	0.90
1:A:1231:LEU:HD11	1:A:1236:GLN:HE22	1.36	0.90
1:A:370:PRO:HB2	1:A:399:ASP:HB2	1.54	0.90
1:A:468:ASN:ND2	1:A:578:PRO:HG2	1.86	0.90
2:C:341:ASP:CG	2:C:342:LEU:H	1.72	0.90
2:C:388:GLY:O	2:C:444:THR:HA	1.70	0.90
2:C:446:THR:C	2:C:450:ASN:HB3	1.92	0.90
1:D:175:TRP:CE3	1:D:184:ALA:HA	2.06	0.90
1:D:214:ALA:N	1:D:401:TRP:CZ2	2.40	0.90
1:A:177:ARG:HA	1:A:182:GLY:O	1.69	0.90
2:B:374:LEU:H	2:B:458:ARG:HH22	1.12	0.90
1:D:1020:ILE:HD13	1:D:1021:SER:CA	2.02	0.90
1:D:1014:ARG:HD3	1:D:1026:ARG:HG3	1.54	0.90
1:D:823:ARG:HA	1:D:826:ILE:HD11	1.51	0.90
1:A:241:PRO:HA	1:A:244:LEU:HB2	1.53	0.90
1:A:197:PHE:N	1:A:269:HIS:NE2	2.19	0.90
2:C:336:LEU:HD23	2:C:337:SER:N	1.87	0.90
2:E:270:PRO:HA	2:E:273:PHE:CE2	2.05	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1208:ARG:HH22	2:F:285:LYS:HD3	1.36	0.90
1:D:265:LEU:HG	1:D:266:VAL:N	1.86	0.90
2:E:456:ARG:HB2	2:E:463:LYS:CG	2.02	0.90
2:F:208:LEU:HD12	2:F:208:LEU:O	1.72	0.90
2:F:447:THR:CA	2:F:452:LEU:HA	2.02	0.90
1:A:1224:ILE:O	1:A:1224:ILE:HD12	1.71	0.90
1:A:276:ALA:HA	1:A:279:ARG:HD3	1.53	0.90
2:C:395:LEU:HD21	2:C:443:VAL:HG12	1.52	0.90
1:D:966:LEU:HD23	1:D:970:ASN:HD21	1.34	0.90
1:D:486:TRP:CZ2	1:D:722:ARG:HA	2.07	0.89
1:D:795:ASN:H	1:D:795:ASN:ND2	1.68	0.89
1:A:653:GLU:HA	1:A:656:TYR:CZ	2.06	0.89
1:D:515:ILE:HD11	1:D:568:GLY:O	1.72	0.89
2:F:61:GLY:O	2:F:64:GLU:HG2	1.72	0.89
1:D:956:GLY:CA	1:D:1094:THR:HG23	2.00	0.89
1:D:460:LYS:N	1:D:460:LYS:CE	2.16	0.89
1:D:548:CYS:CA	1:D:551:LYS:HD3	2.01	0.89
2:F:305:ASN:C	2:F:335:VAL:HG11	1.92	0.89
2:F:341:ASP:CG	2:F:342:LEU:N	2.22	0.89
2:C:208:LEU:HD13	2:C:240:VAL:HB	1.53	0.89
2:C:350:LEU:HD12	2:C:350:LEU:H	1.37	0.89
1:D:548:CYS:N	1:D:551:LYS:HZ3	1.70	0.89
1:D:614:TYR:CE2	1:D:620:TRP:CE3	2.60	0.89
2:E:128:VAL:HB	2:E:208:LEU:HB3	1.53	0.89
2:C:256:LEU:HD11	2:C:287:ASN:OD1	1.73	0.89
2:C:241:TRP:N	2:C:336:LEU:CB	2.36	0.89
1:A:564:GLN:HA	1:A:567:PRO:HD2	1.53	0.89
1:D:1206:MET:O	1:D:1210:TYR:N	2.06	0.89
2:E:431:LYS:HA	2:E:434:GLU:HG2	1.52	0.89
2:F:403:PHE:CE1	2:F:415:PRO:HG3	2.06	0.89
1:A:1146:ARG:NH1	1:A:1185:ILE:HG23	1.86	0.89
1:A:214:ALA:HA	1:A:401:TRP:CZ2	2.07	0.89
2:B:324:GLY:HA2	2:B:325:ARG:CZ	2.01	0.89
2:C:102:LEU:HD11	2:C:435:MET:CE	2.01	0.89
2:C:396:ARG:NH1	2:C:397:GLN:HA	1.86	0.89
1:D:1066:THR:HG23	1:D:1068:ASP:OD1	1.71	0.89
1:D:179:GLY:CA	1:D:182:GLY:HA3	2.02	0.89
1:D:767:ALA:CB	1:D:771:LEU:HD22	2.03	0.89
1:D:804:ALA:O	1:D:808:ILE:HG13	1.72	0.89
1:A:1043:VAL:CA	1:A:1046:GLU:H	1.85	0.89
1:A:127:LEU:HD22	1:A:129:PRO:HD3	1.54	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:504:LYS:HD2	1:D:523:ASP:CB	2.01	0.89
1:D:457:ARG:N	1:D:460:LYS:HZ2	1.71	0.89
1:D:795:ASN:O	1:D:798:ILE:HG12	1.73	0.89
2:F:322:LEU:HD13	2:F:322:LEU:O	1.72	0.89
2:F:336:LEU:HD23	2:F:337:SER:N	1.87	0.89
2:B:396:ARG:NH1	2:B:417:TYR:CD2	2.40	0.89
1:D:366:GLY:N	1:D:367:PRO:HD3	1.88	0.89
1:D:371:LYS:O	1:D:373:PRO:HD3	1.72	0.89
1:D:213:VAL:HG22	1:D:401:TRP:CZ2	2.08	0.89
1:D:634:LEU:HD22	1:D:635:PRO:HD2	1.53	0.89
1:A:1066:THR:HG1	1:A:1071:ARG:HB3	1.37	0.89
1:A:256:PRO:HD2	1:A:257:THR:H	1.38	0.89
1:A:608:ASP:CA	1:A:779:LEU:HB2	2.03	0.89
1:A:621:GLY:CA	1:A:748:TRP:HB3	2.02	0.89
2:C:82:SER:HA	2:C:86:LEU:CD2	2.01	0.89
1:D:1166:TYR:HD1	1:D:1166:TYR:N	1.69	0.89
1:D:869:ARG:H	1:D:1200:PRO:HB3	1.37	0.89
1:A:1011:PRO:O	1:A:1028:VAL:HG21	1.71	0.89
1:A:604:ALA:HA	1:A:717:LEU:HD22	1.54	0.89
1:A:623:LEU:HG	1:A:623:LEU:O	1.73	0.89
1:A:912:GLY:C	1:A:917:GLY:HA3	1.92	0.89
1:D:95:PRO:HG2	1:D:99:ALA:HB3	1.54	0.89
1:D:213:VAL:HG13	1:D:401:TRP:CZ2	2.07	0.89
1:D:267:VAL:HB	1:D:269:HIS:HD2	1.28	0.89
1:D:374:ARG:CD	1:D:392:LEU:HD21	2.02	0.89
1:D:472:GLN:HB3	2:F:460:THR:O	1.72	0.89
1:D:772:PRO:HA	1:D:775:GLU:HB2	1.54	0.89
1:D:1209:ARG:NH2	2:F:253:ASP:HA	1.86	0.89
2:F:395:LEU:O	2:F:398:VAL:HB	1.73	0.89
1:A:297:MET:CE	1:A:411:LEU:HD13	2.03	0.89
1:A:425:ALA:HA	1:A:428:LEU:HD12	1.54	0.89
1:A:613:HIS:CB	1:A:717:LEU:HG	2.01	0.89
2:C:458:ARG:HH11	2:C:458:ARG:HG2	1.38	0.89
1:D:486:TRP:HA	1:D:489:GLU:OE2	1.73	0.89
1:D:608:ASP:O	1:D:609:GLY:C	2.12	0.89
1:D:879:GLN:OE1	1:D:1189:LEU:HD21	1.71	0.89
2:E:201:ASN:OD1	2:F:422:GLN:HB2	1.72	0.89
1:A:253:ALA:HB3	1:A:256:PRO:HB3	1.55	0.89
1:A:612:LEU:CG	1:A:613:HIS:N	2.33	0.89
1:A:656:TYR:HB3	1:A:720:THR:OG1	1.72	0.89
2:B:117:SER:HB2	2:B:262:TRP:CE2	2.07	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:254:PHE:HA	2:C:257:ARG:HH21	1.36	0.89
1:D:348:LEU:CD2	1:D:350:ILE:HG13	2.02	0.88
1:D:448:ALA:HB1	1:D:870:VAL:HB	1.51	0.88
1:D:914:THR:CB	1:D:999:ASP:HB3	2.03	0.88
2:E:185:LEU:HD11	2:E:236:GLU:O	1.73	0.88
2:F:474:PHE:HA	2:F:477:LYS:CD	2.01	0.88
1:A:1005:VAL:HA	1:A:1008:LEU:HD12	1.55	0.88
1:A:173:GLU:OE2	1:A:223:TRP:HZ2	1.55	0.88
1:A:352:SER:O	1:A:358:LEU:HD21	1.73	0.88
1:A:815:TRP:HA	1:A:838:GLY:CA	2.03	0.88
2:B:106:LEU:HD12	2:B:109:ASN:ND2	1.88	0.88
2:B:311:LEU:HG	2:B:312:LEU:H	1.35	0.88
1:D:1069:ILE:CG1	1:D:1070:PRO:HD2	2.02	0.88
1:D:349:ASP:O	1:D:361:ARG:HD3	1.73	0.88
2:F:454:HIS:N	2:F:466:MET:O	2.06	0.88
1:A:996:ARG:NH1	1:A:1003:TRP:HB2	1.89	0.88
2:B:115:TRP:O	2:B:119:VAL:HB	1.71	0.88
2:C:125:VAL:HG23	2:C:126:PHE:N	1.88	0.88
2:C:386:ASP:HB2	2:C:442:LEU:CA	2.02	0.88
1:A:638:THR:HG23	1:A:642:SER:OG	1.73	0.88
1:D:895:GLU:OE1	1:D:896:LEU:HD22	1.71	0.88
1:A:202:CYS:SG	1:A:210:THR:HG21	2.14	0.88
1:D:1131:ILE:HD12	1:D:1131:ILE:N	1.87	0.88
1:D:179:GLY:C	1:D:181:GLU:N	2.17	0.88
2:F:467:HIS:O	2:F:471:LEU:CD1	2.21	0.88
1:A:360:GLU:O	1:A:364:LEU:HD23	1.73	0.88
1:A:477:GLU:HG3	1:A:478:ARG:H	0.72	0.88
1:D:569:HIS:N	1:D:570:PRO:HD2	1.88	0.88
1:D:1208:ARG:NH2	2:F:285:LYS:HD3	1.89	0.88
1:D:352:SER:O	1:D:358:GLU:OE2	1.82	0.88
1:D:601:LYS:HD2	1:D:616:GLU:HG2	1.56	0.88
2:F:390:GLY:HA2	2:F:391:PRO:C	1.94	0.88
2:F:424:SER:HB3	2:F:427:GLN:HE22	1.39	0.88
2:F:471:LEU:HD12	2:F:471:LEU:N	1.88	0.88
1:A:215:ILE:CG2	1:A:216:SER:H	1.86	0.88
1:A:871:GLY:O	1:A:873:GLU:N	2.07	0.88
1:D:1010:LEU:CD2	1:D:1015:THR:HA	2.03	0.88
1:D:359:VAL:O	1:D:359:VAL:HG12	1.71	0.88
1:D:463:LEU:HA	1:D:466:LEU:CD2	2.04	0.88
1:D:656:TYR:HB3	1:D:744:ILE:CG2	2.04	0.88
1:D:83:LEU:HG	1:D:89:GLY:HA2	1.53	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:432:TYR:HB2	2:E:437:ILE:HD13	1.54	0.88
2:B:186:LEU:HD11	2:B:337:SER:HB2	1.55	0.88
1:D:1028:VAL:HG22	1:D:1096:ARG:NH1	1.87	0.88
1:D:1198:LYS:HA	1:D:1204:THR:CB	2.03	0.88
1:D:1206:MET:CG	2:F:257:ARG:NH1	2.36	0.88
1:D:214:ALA:HB3	1:D:218:SER:C	1.94	0.88
1:D:458:GLU:O	1:D:461:LYS:HG2	1.72	0.88
1:D:73:PRO:C	1:D:74:LEU:HD22	1.93	0.88
2:F:406:LEU:HA	2:F:409:ASN:HD21	1.39	0.88
1:A:1026:ARG:HB3	1:A:1029:GLN:CD	1.93	0.88
1:A:1124:ALA:HB1	1:A:1148:ARG:HH11	1.38	0.88
1:A:606:THR:CG2	1:A:620:TRP:HE3	1.86	0.88
1:A:925:LYS:CG	1:A:926:SER:H	1.85	0.88
2:B:67:GLU:HB3	2:B:70:LEU:HB2	1.53	0.88
1:D:278:ILE:O	1:D:282:TYR:CD1	2.27	0.88
1:D:645:VAL:O	1:D:645:VAL:HG13	1.74	0.88
1:D:737:GLY:H	1:D:750:PHE:HZ	1.16	0.88
1:A:515:ILE:HD11	1:A:568:GLY:O	1.72	0.88
1:A:792:LEU:HD12	1:A:793:GLU:N	1.88	0.88
2:B:400:GLN:N	2:B:400:GLN:OE1	2.07	0.88
2:C:342:LEU:HB2	2:C:344:ARG:CG	2.01	0.88
1:D:504:LYS:CB	1:D:523:ASP:CB	2.49	0.88
1:D:352:SER:O	1:D:358:GLU:CG	2.20	0.88
1:D:368:PRO:HA	1:D:369:LEU:HD23	1.56	0.88
1:D:611:PRO:HB2	1:D:612:LEU:HD13	1.56	0.88
1:D:854:ALA:HB1	1:D:860:LEU:HD12	1.55	0.88
1:A:243:ASP:O	1:A:246:PRO:HD2	1.73	0.88
1:A:297:MET:HE1	1:A:411:LEU:HD13	1.55	0.88
1:A:575:LYS:CG	1:A:576:LEU:H	1.86	0.88
2:B:447:THR:HG23	2:B:452:LEU:N	1.88	0.88
1:D:163:PRO:HB2	1:D:165:LYS:HG3	1.56	0.88
1:D:1220:ILE:O	1:D:1223:ILE:HG23	1.75	0.88
2:E:302:THR:C	2:E:303:LEU:HD12	1.94	0.88
1:A:1007:GLU:O	1:A:1011:PRO:HD3	1.73	0.88
1:A:436:PRO:HB3	1:A:881:PRO:CD	2.03	0.88
1:A:744:ILE:CB	1:A:745:PRO:HD3	2.04	0.88
2:C:320:SER:HB3	2:C:321:LYS:HG3	1.54	0.88
2:B:277:ASP:OD1	2:B:287:ASN:ND2	2.06	0.88
1:D:849:THR:HG22	1:D:851:THR:H	1.38	0.87
1:D:441:TRP:HE1	1:D:880:ALA:N	1.71	0.87
1:A:276:ALA:O	1:A:279:ARG:HB2	1.73	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:656:TYR:CB	1:A:720:THR:HB	2.05	0.87
1:A:845:VAL:HB	1:A:855:VAL:CB	2.04	0.87
2:F:217:PRO:HB2	2:F:219:PHE:CE1	2.10	0.87
1:D:1014:ARG:HG2	1:D:1026:ARG:HE	1.38	0.87
1:D:79:LEU:HD11	1:D:1173:LEU:O	1.74	0.87
1:D:897:TRP:CH2	1:D:1177:VAL:HG11	2.10	0.87
1:A:1041:TRP:NE1	1:A:1043:VAL:HB	1.90	0.87
1:A:296:SER:HB2	1:A:847:ALA:HB3	1.53	0.87
1:A:440:ASN:HB2	1:A:877:MET:SD	2.14	0.87
1:A:869:ARG:HA	1:A:1201:SER:H	1.36	0.87
2:C:241:TRP:N	2:C:336:LEU:HB2	1.89	0.87
1:D:1141:VAL:CB	1:D:1146:ARG:HE	1.82	0.87
1:D:599:THR:O	1:D:603:MET:HG3	1.74	0.87
1:D:989:THR:HG22	1:D:999:ASP:OD2	1.74	0.87
1:D:997:LEU:HD22	1:D:998:SER:H	1.38	0.87
2:F:341:ASP:O	2:F:342:LEU:HD13	1.74	0.87
2:C:301:GLU:CG	2:C:339:ASN:HB3	2.04	0.87
1:D:358:GLU:O	1:D:361:ARG:HB3	1.72	0.87
1:D:420:HIS:NE2	1:D:1020:ILE:HG21	1.88	0.87
1:D:610:PHE:N	1:D:611:PRO:HD3	1.85	0.87
2:F:417:TYR:HD2	2:F:417:TYR:N	1.71	0.87
2:F:424:SER:CB	2:F:427:GLN:HE22	1.87	0.87
1:A:444:TYR:HE1	1:A:1206:MET:SD	1.98	0.87
2:C:243:THR:OG1	2:C:334:CYS:HB2	1.73	0.87
2:C:304:TRP:N	2:C:337:SER:HB2	1.90	0.87
2:C:455:LEU:C	2:C:456:ARG:HE	1.78	0.87
2:B:453:ILE:HG21	2:B:468:ILE:HD12	1.57	0.87
1:D:196:VAL:HG21	1:D:215:ILE:HD12	1.53	0.87
1:D:198:ASP:CB	1:D:400:VAL:HG11	2.04	0.87
2:E:114:TRP:HE1	2:E:259:ARG:NH2	1.71	0.87
2:F:419:GLU:OE2	2:F:428:LEU:HD13	1.73	0.87
1:A:869:ARG:C	1:A:1200:PRO:HA	1.94	0.87
1:A:215:ILE:HG23	1:A:216:SER:N	1.85	0.87
2:B:261:GLN:HE21	2:B:264:ARG:NH2	1.71	0.87
2:C:113:GLU:O	2:C:116:THR:HG23	1.75	0.87
2:C:244:PRO:HD2	2:C:247:THR:CG2	2.05	0.87
2:C:471:LEU:HD12	2:C:471:LEU:H	1.33	0.87
2:B:83:LYS:HD2	2:B:83:LYS:H	1.36	0.87
1:D:1029:GLN:CG	1:D:1030:ARG:NE	2.38	0.87
1:D:449:GLN:HA	1:D:452:TYR:CE2	2.09	0.87
1:D:621:GLY:HA2	1:D:748:TRP:HB3	1.57	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:992:LEU:CB	1:D:996:ARG:HG2	2.04	0.87
1:A:1046:GLU:O	1:A:1049:TRP:CD1	2.28	0.87
1:A:612:LEU:CG	1:A:614:TYR:H	1.87	0.87
1:A:989:THR:HA	1:A:998:SER:OG	1.74	0.87
2:B:261:GLN:HE21	2:B:264:ARG:HH21	0.88	0.87
2:B:261:GLN:HA	2:B:264:ARG:HE	1.38	0.87
1:A:741:ASP:OD1	1:A:742:VAL:HG23	1.75	0.87
1:D:1067:SER:HA	1:D:1071:ARG:CA	2.03	0.87
1:D:612:LEU:CB	1:D:620:TRP:HB2	2.05	0.87
1:D:845:VAL:HB	1:D:855:VAL:HG21	1.57	0.87
1:D:798:ILE:CG2	1:D:869:ARG:NH1	2.36	0.87
1:A:1011:PRO:HD2	1:A:1012:VAL:HG22	1.54	0.87
1:A:249:VAL:HG23	1:A:815:TRP:O	1.75	0.87
1:A:618:HIS:H	1:A:725:PRO:HG2	1.30	0.87
1:A:502:LYS:CG	1:A:503:VAL:H	1.87	0.87
2:E:418:LEU:HD22	2:F:122:ARG:NH2	1.87	0.87
2:C:317:GLY:C	2:C:319:VAL:H	1.76	0.87
1:D:1019:TRP:HE3	1:D:1168:LEU:CG	1.88	0.87
1:D:193:ARG:HA	1:D:193:ARG:NH1	1.90	0.87
1:D:249:VAL:CG2	1:D:250:PRO:HD2	2.04	0.87
1:A:653:GLU:O	1:A:657:ARG:HG3	1.75	0.87
2:B:254:PHE:O	2:B:257:ARG:HB3	1.73	0.87
2:C:456:ARG:HB2	2:C:463:LYS:HA	1.55	0.87
1:D:95:PRO:HG2	1:D:99:ALA:CB	2.05	0.87
1:D:576:LEU:HG	1:D:577:CYS:H	1.37	0.87
1:D:660:CYS:CB	1:D:745:PRO:HG3	2.04	0.87
1:D:812:MET:O	1:D:840:ILE:HG23	1.73	0.87
1:D:815:TRP:HE3	1:D:838:GLY:H	1.19	0.87
1:D:927:ARG:HG2	1:D:928:GLY:N	1.88	0.87
2:E:205:PRO:HB3	2:E:243:THR:HA	1.57	0.87
1:A:515:ILE:HG12	1:A:565:HIS:O	1.75	0.87
2:B:261:GLN:HA	2:B:264:ARG:NE	1.89	0.87
2:C:77:HIS:O	2:C:100:GLY:HA3	1.75	0.87
1:D:456:GLN:C	1:D:460:LYS:NZ	2.29	0.86
1:D:768:LYS:HD3	1:D:769:ASP:N	1.87	0.86
1:D:795:ASN:H	1:D:795:ASN:HD22	0.88	0.86
1:D:996:ARG:CD	1:D:996:ARG:H	1.87	0.86
1:D:1209:ARG:HG3	2:F:249:ASN:HB3	1.55	0.86
1:A:869:ARG:HG2	1:A:870:VAL:HG23	1.55	0.86
2:C:208:LEU:HD22	2:C:242:PHE:CE2	2.10	0.86
1:A:230:GLU:CD	1:A:231:GLU:N	2.28	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:348:LEU:C	1:A:350:ILE:N	2.18	0.86
1:A:576:LEU:HD23	2:C:481:SER:OG	1.73	0.86
2:B:72:ILE:CA	2:B:75:ARG:HE	1.85	0.86
1:D:452:TYR:HB3	1:D:801:TRP:HE1	1.38	0.86
1:D:452:TYR:HB3	1:D:801:TRP:NE1	1.89	0.86
1:D:772:PRO:O	1:D:775:GLU:O	1.93	0.86
2:E:78:PHE:CD2	2:E:102:LEU:HD12	2.10	0.86
1:A:613:HIS:HB3	1:A:717:LEU:CG	2.04	0.86
2:B:396:ARG:HG3	2:B:417:TYR:HB2	1.56	0.86
1:D:1006:ARG:HA	1:D:1006:ARG:HH11	0.73	0.86
2:F:439:PHE:HD1	2:F:458:ARG:CB	1.88	0.86
1:A:903:GLY:HA2	1:A:1002:GLU:HB3	1.54	0.86
1:A:310:SER:O	1:A:313:ILE:HG13	1.75	0.86
2:C:252:LEU:HD12	2:C:305:ASN:HB2	1.56	0.86
1:D:163:PRO:HG2	1:D:165:LYS:HE3	1.57	0.86
1:D:995:TYR:O	1:D:1004:LEU:HD23	1.75	0.86
1:D:1066:THR:O	1:D:1073:PRO:HD3	1.76	0.86
1:D:1208:ARG:HH12	2:F:285:LYS:CD	1.88	0.86
2:E:455:LEU:O	2:E:463:LYS:HA	1.75	0.86
1:A:1146:ARG:CZ	1:A:1185:ILE:HG23	2.06	0.86
1:A:184:ALA:O	1:A:185:VAL:HG12	1.75	0.86
1:A:135:LEU:CD2	1:A:135:LEU:H	1.83	0.86
1:D:1019:TRP:CD1	1:D:1020:ILE:HG22	2.10	0.86
1:D:300:ALA:HB2	1:D:847:ALA:CB	2.03	0.86
1:D:439:GLN:NE2	1:D:835:GLY:HA3	1.90	0.86
1:D:891:VAL:HA	1:D:894:GLN:NE2	1.91	0.86
2:F:114:TRP:HB2	2:F:262:TRP:CZ2	2.10	0.86
2:F:238:SER:CA	2:F:338:VAL:HG11	2.06	0.86
1:D:549:LEU:HD11	2:F:401:GLY:HA3	1.57	0.86
1:A:190:PRO:CB	1:A:192:GLU:HG2	2.03	0.86
1:A:412:PRO:HG2	1:A:413:LEU:HG	1.54	0.86
1:D:485:LEU:HD12	1:D:489:GLU:OE2	1.76	0.86
1:D:490:TRP:CE3	1:D:580:LEU:HD22	2.10	0.86
2:F:203:ARG:HE	2:F:204:LEU:CD1	1.87	0.86
2:F:241:TRP:CZ3	2:F:255:TRP:NE1	2.43	0.86
2:F:418:LEU:O	2:F:421:MET:HG2	1.74	0.86
1:A:1158:LEU:O	1:A:1162:CYS:SG	2.33	0.86
1:A:293:ASP:HB3	1:A:296:SER:H	1.40	0.86
1:A:472:GLN:NE2	2:C:460:THR:O	2.07	0.86
2:C:326:ASP:OD2	2:C:329:LYS:N	2.08	0.86
2:C:390:GLY:HA2	2:C:391:PRO:C	1.95	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:391:PRO:HB2	2:C:394:GLU:OE2	1.76	0.86
1:D:1055:SER:HA	1:D:1058:PHE:CB	2.01	0.86
1:D:284:ILE:HD12	1:D:284:ILE:H	1.38	0.86
1:D:824:ALA:HA	1:D:827:ARG:HG3	1.55	0.86
2:E:374:LEU:O	2:E:375:HIS:HB2	1.74	0.86
2:E:387:VAL:HG12	2:E:416:GLY:HA3	1.57	0.86
2:C:289:LEU:HD12	2:C:301:GLU:HB2	1.56	0.86
2:C:387:VAL:HG12	2:C:395:LEU:CD2	2.06	0.86
2:E:270:PRO:HA	2:E:273:PHE:CD2	2.09	0.86
1:D:995:TYR:HE1	1:D:1004:LEU:HB3	1.41	0.86
1:D:242:ALA:HB1	1:D:245:ILE:CD1	2.05	0.86
1:D:383:LYS:O	1:D:386:ARG:HG3	1.76	0.86
1:D:457:ARG:HH22	2:F:485:VAL:HG23	1.40	0.86
1:D:869:ARG:N	1:D:1200:PRO:CB	2.38	0.86
2:E:387:VAL:HG12	2:E:416:GLY:CA	2.04	0.86
2:E:420:THR:HG23	2:E:421:MET:H	1.40	0.86
2:C:385:LEU:O	2:C:416:GLY:HA3	1.76	0.86
1:D:459:MET:H	1:D:460:LYS:HZ1	1.20	0.86
1:D:869:ARG:N	1:D:1200:PRO:HB3	1.89	0.86
2:F:343:ASP:C	2:F:347:LEU:HD13	1.94	0.86
1:A:1020:ILE:HD12	1:A:1021:SER:H	1.40	0.86
1:A:378:VAL:O	1:A:380:GLY:N	2.09	0.86
1:A:556:THR:HG23	2:C:452:LEU:CB	2.05	0.86
1:A:617:ARG:O	1:A:762:VAL:HG23	1.76	0.86
2:B:267:ALA:HB2	2:B:270:PRO:HB2	1.56	0.86
1:D:189:ILE:CG2	1:D:258:GLN:HB3	2.05	0.85
1:D:293:ASP:O	1:D:296:SER:HB3	1.76	0.85
1:D:598:VAL:HA	1:D:601:LYS:CE	2.06	0.85
2:F:190:LEU:HB2	2:F:314:MET:HE1	1.56	0.85
1:A:145:LYS:HG3	1:A:1118:TRP:CH2	2.11	0.85
1:A:272:SER:HB3	1:A:273:PHE:CD2	2.11	0.85
2:C:446:THR:OG1	2:C:455:LEU:HD21	1.76	0.85
1:A:472:GLN:HG3	2:C:460:THR:CB	2.05	0.85
1:D:564:GLN:HA	1:D:566:LEU:O	1.75	0.85
1:D:876:ALA:HB1	1:D:1188:CYS:SG	2.16	0.85
1:D:635:PRO:HA	1:D:639:THR:OG1	1.77	0.85
1:D:801:TRP:CH2	1:D:805:HIS:CG	2.65	0.85
2:F:306:LEU:O	2:F:306:LEU:HG	1.76	0.85
1:A:576:LEU:HD12	1:A:577:CYS:N	1.91	0.85
2:C:375:HIS:HB3	2:C:378:LEU:HD13	1.55	0.85
2:C:385:LEU:HD21	2:C:402:LEU:HD23	1.57	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:249:VAL:CG1	1:D:252:GLY:N	2.34	0.85
1:D:302:SER:HB2	1:D:351:SER:OG	1.76	0.85
1:D:612:LEU:HG	1:D:621:GLY:C	1.97	0.85
1:D:652:ILE:HA	1:D:655:LEU:CG	2.06	0.85
2:E:101:PRO:CD	2:F:128:VAL:HG21	2.06	0.85
2:F:215:PHE:CD2	2:F:233:GLU:HG2	2.11	0.85
1:D:1209:ARG:NH2	2:F:252:LEU:HD22	1.90	0.85
2:F:387:VAL:O	2:F:420:THR:OG1	1.94	0.85
1:A:369:PRO:CB	1:A:372:GLU:HG3	1.98	0.85
2:B:102:LEU:H	2:B:102:LEU:HD12	1.41	0.85
2:C:127:PRO:HD3	2:C:209:ALA:C	1.97	0.85
1:D:78:MET:SD	1:D:1175:GLN:NE2	2.49	0.85
1:D:1138:ARG:HH22	1:D:1191:LYS:CE	1.88	0.85
1:D:242:ALA:O	1:D:246:PRO:HD3	1.76	0.85
2:F:471:LEU:N	2:F:471:LEU:CD1	2.36	0.85
1:A:369:PRO:HB2	1:A:372:GLU:CG	1.98	0.85
1:A:359:ALA:HB2	1:A:407:PHE:HE2	1.39	0.85
1:A:604:ALA:HA	1:A:717:LEU:CD2	2.07	0.85
1:A:629:ASP:HB3	1:A:744:ILE:O	1.76	0.85
2:C:129:ASP:OD1	2:C:131:LEU:HD22	1.76	0.85
2:C:446:THR:HB	2:C:450:ASN:OD1	1.76	0.85
2:B:472:LYS:HG3	2:B:473:ASP:N	1.90	0.85
1:D:1191:LYS:O	1:D:1192:GLU:HB3	1.74	0.85
1:D:1212:ILE:HG23	1:D:1214:GLN:H	1.41	0.85
1:D:741:ASP:HB2	1:D:748:TRP:CH2	2.11	0.85
1:A:1141:VAL:HB	1:A:1146:ARG:CD	2.07	0.85
1:A:445:LEU:O	1:A:448:ALA:HB3	1.77	0.85
1:A:921:LEU:HD12	1:A:921:LEU:H	1.39	0.85
2:C:473:ASP:HA	2:C:476:ILE:HG23	1.56	0.85
1:D:175:TRP:CZ3	1:D:184:ALA:HA	2.10	0.85
1:D:309:ARG:HA	1:D:312:TRP:HB2	1.58	0.85
2:F:387:VAL:HG21	2:F:417:TYR:CA	2.07	0.85
2:F:394:GLU:HA	2:F:397:GLN:OE1	1.75	0.85
1:A:670:PRO:HG2	1:A:671:GLN:HE21	1.38	0.85
1:A:741:ASP:O	1:A:742:VAL:HB	1.77	0.85
2:F:326:ASP:OD1	2:F:329:LYS:HG2	1.76	0.85
1:D:444:TYR:OH	1:D:1206:MET:HB3	1.75	0.85
1:D:1221:TYR:HA	1:D:1224:ILE:HG23	1.59	0.85
1:D:995:TYR:H	1:D:996:ARG:NH1	1.74	0.85
1:A:1158:LEU:HD21	1:A:1178:ALA:O	1.77	0.85
1:A:784:GLY:N	2:C:363:ARG:CD	2.39	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:311:LEU:HG	2:B:312:LEU:N	1.89	0.85
2:C:244:PRO:CD	2:C:247:THR:HG21	2.05	0.85
1:D:524:PRO:O	1:D:525:MET:HB3	1.74	0.85
1:D:564:GLN:CA	1:D:566:LEU:O	2.25	0.85
1:D:569:HIS:H	1:D:570:PRO:HD2	1.40	0.85
1:A:832:ASP:O	1:A:833:GLU:HB2	1.77	0.85
1:D:1025:LEU:O	1:D:1027:LYS:HD3	1.75	0.85
2:F:342:LEU:HB2	2:F:344:ARG:HB2	1.59	0.85
1:A:127:LEU:HD23	1:A:128:PRO:HD2	1.57	0.85
1:A:606:THR:CG2	1:A:613:HIS:N	2.40	0.85
1:A:744:ILE:CG1	1:A:745:PRO:HD3	2.05	0.85
2:C:342:LEU:HG	2:C:344:ARG:HB2	1.57	0.85
2:C:180:LYS:HE3	2:C:181:LEU:N	1.91	0.85
2:F:317:GLY:HA3	2:F:320:SER:H	1.40	0.85
1:A:98:ALA:O	1:A:102:ARG:HG2	1.77	0.85
1:A:1145:ASP:HA	1:A:1148:ARG:HB2	1.59	0.85
1:A:1210:TYR:HE1	1:A:1213:PRO:HG2	1.35	0.85
1:A:612:LEU:HG	1:A:614:TYR:N	1.89	0.85
1:A:85:GLU:OE1	1:A:128:PRO:HA	1.76	0.85
2:F:326:ASP:OD2	2:F:329:LYS:N	2.08	0.85
1:D:1222:GLN:O	1:D:1226:LEU:HD22	1.74	0.85
1:D:549:LEU:H	1:D:551:LYS:NZ	1.71	0.85
1:D:559:LEU:HD12	1:D:560:PRO:HD3	1.57	0.85
2:E:109:ASN:HB2	2:E:378:LEU:CA	2.07	0.85
1:A:767:ALA:HB3	1:A:771:LEU:HD23	1.59	0.85
1:D:504:LYS:HB2	1:D:523:ASP:O	1.76	0.85
1:A:503:VAL:HG21	1:A:520:ALA:O	1.77	0.85
1:D:171:TRP:O	1:D:172:ALA:HB2	1.77	0.84
1:D:612:LEU:CD1	1:D:612:LEU:H	1.89	0.84
1:D:775:GLU:HA	1:D:780:GLN:CG	2.06	0.84
2:F:454:HIS:O	2:F:455:LEU:HB2	1.73	0.84
1:A:223:TRP:N	1:A:223:TRP:HD1	1.74	0.84
1:A:232:ARG:CB	1:A:239:LEU:HD22	2.04	0.84
1:A:452:TYR:HA	1:A:869:ARG:HH21	1.40	0.84
1:A:659:HIS:O	1:A:663:GLN:HG3	1.76	0.84
2:C:471:LEU:CD1	2:C:471:LEU:H	1.66	0.84
1:D:513:LEU:N	1:D:570:PRO:HG3	1.90	0.84
1:D:529:ASP:O	1:D:530:LEU:HB3	1.75	0.84
2:E:328:ARG:N	2:E:328:ARG:HD2	1.90	0.84
2:F:342:LEU:HB2	2:F:344:ARG:CB	2.06	0.84
1:A:1131:ILE:HG21	1:A:1137:VAL:C	1.96	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:213:VAL:HA	1:A:221:TYR:CE2	2.12	0.84
1:A:203:LEU:HD12	1:A:380:GLY:N	1.92	0.84
1:A:869:ARG:CG	1:A:870:VAL:H	1.89	0.84
1:A:992:LEU:CD1	1:A:993:ARG:H	1.90	0.84
2:C:354:PHE:HB2	2:C:372:LEU:HA	1.59	0.84
1:A:97:GLU:O	1:A:101:ARG:HB2	1.77	0.84
1:D:442:GLU:OE2	1:D:442:GLU:HA	1.77	0.84
1:D:797:MET:C	1:D:869:ARG:HD3	1.97	0.84
1:D:887:VAL:HG22	1:D:1185:ILE:HG22	1.57	0.84
2:E:432:TYR:CA	2:E:435:MET:HB2	2.05	0.84
2:F:130:ALA:HB1	2:F:182:ARG:HH12	1.42	0.84
2:F:214:CYS:SG	2:F:236:GLU:CG	2.65	0.84
1:A:569:HIS:CD2	1:A:570:PRO:HD2	2.13	0.84
1:A:612:LEU:CG	1:A:613:HIS:H	1.83	0.84
1:A:854:ALA:CB	1:A:860:LEU:HD13	2.07	0.84
1:D:269:HIS:C	1:D:294:THR:HG23	1.97	0.84
1:D:576:LEU:O	1:D:577:CYS:O	1.95	0.84
2:F:268:MET:HB2	2:F:352:ASP:OD1	1.76	0.84
1:A:1200:PRO:HB2	1:A:1203:PRO:N	1.90	0.84
1:A:308:GLN:HA	1:A:311:LEU:HD12	1.60	0.84
1:A:400:VAL:O	1:A:403:THR:HG23	1.77	0.84
2:C:291:TYR:OH	2:C:355:GLN:HG3	1.77	0.84
2:C:397:GLN:O	2:C:398:VAL:HG22	1.78	0.84
2:C:73:CYS:HB3	2:C:79:LEU:HD23	1.57	0.84
1:D:1197:CYS:O	1:D:1204:THR:OG1	1.95	0.84
1:D:1221:TYR:HA	1:D:1224:ILE:HG22	1.58	0.84
1:D:223:TRP:O	1:D:227:ARG:HD2	1.78	0.84
2:F:374:LEU:HB3	2:F:378:LEU:HB3	1.57	0.84
2:F:440:THR:O	2:F:457:SER:HB2	1.77	0.84
2:F:471:LEU:HA	2:F:474:PHE:HD2	1.41	0.84
1:A:879:GLN:HG2	1:A:1189:LEU:CD2	2.06	0.84
1:A:448:ALA:HB1	1:A:870:VAL:CG1	2.07	0.84
1:D:74:LEU:HA	1:D:76:ILE:O	1.77	0.84
1:D:779:LEU:HD22	1:D:780:GLN:N	1.92	0.84
1:D:987:ALA:O	1:D:991:GLY:N	2.10	0.84
1:A:286:GLY:N	1:A:1128:ARG:HH22	1.76	0.84
1:A:1145:ASP:HB3	1:A:1148:ARG:NH1	1.93	0.84
1:A:606:THR:HG22	1:A:620:TRP:CE3	2.12	0.84
1:A:655:LEU:O	1:A:659:HIS:HB2	1.77	0.84
1:A:842:PRO:HB2	1:A:856:GLU:OE1	1.77	0.84
1:D:210:THR:O	1:D:211:LEU:HB2	1.78	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:617:ARG:O	1:D:762:VAL:HG23	1.77	0.84
1:D:770:PHE:O	1:D:774:MET:HG2	1.77	0.84
2:F:180:LYS:HE3	2:F:181:LEU:N	1.91	0.84
2:F:347:LEU:C	2:F:350:LEU:HD11	1.98	0.84
2:F:346:MET:CE	2:F:347:LEU:HD12	2.08	0.84
1:A:118:VAL:H	1:A:119:PRO:HD3	1.42	0.84
1:A:854:ALA:HB1	1:A:860:LEU:CD1	2.07	0.84
1:A:904:ASP:HB2	1:A:910:MET:O	1.77	0.84
1:D:1027:LYS:HB3	1:D:1091:GLU:CB	2.08	0.84
1:D:80:SER:HB2	1:D:125:LEU:HB2	1.60	0.84
2:E:377:CYS:O	2:E:380:PRO:HD2	1.76	0.84
2:E:200:VAL:HG12	2:F:418:LEU:CD2	2.08	0.84
1:A:175:TRP:H	1:A:186:PRO:HB3	1.41	0.84
1:A:618:HIS:HA	1:A:762:VAL:CG2	2.08	0.84
1:A:825:VAL:O	1:A:829:PRO:CD	2.26	0.84
1:A:850:ILE:HD11	1:A:1099:TRP:CZ2	2.13	0.84
2:B:204:LEU:HG	2:B:325:ARG:NH2	1.91	0.84
2:C:129:ASP:OD1	2:C:131:LEU:HD13	1.77	0.84
1:D:157:LEU:CD2	1:D:194:ALA:HB3	2.08	0.84
1:D:257:THR:O	1:D:257:THR:HG22	1.77	0.84
1:D:282:TYR:HE2	1:D:432:VAL:HG22	1.41	0.84
1:D:302:SER:O	1:D:1024:ASP:OD2	1.95	0.84
1:D:854:ALA:HB1	1:D:860:LEU:HD13	1.58	0.84
2:F:336:LEU:HD23	2:F:336:LEU:C	1.98	0.84
1:A:256:PRO:CD	1:A:257:THR:H	1.87	0.84
1:A:397:ALA:O	1:A:400:VAL:HB	1.77	0.84
2:B:257:ARG:O	2:B:260:LEU:HB3	1.76	0.84
1:D:1236:GLN:HB2	1:D:1237:PRO:CD	2.07	0.84
2:F:197:LEU:HD11	2:F:322:LEU:HG	1.60	0.84
2:F:357:THR:HA	2:F:370:LYS:HZ1	1.43	0.84
1:A:308:GLN:HE21	1:A:1091:GLU:H	1.26	0.84
1:A:166:PRO:HG3	1:A:398:GLN:HE22	1.39	0.84
1:A:652:ILE:HA	1:A:655:LEU:HD21	1.60	0.84
1:A:801:TRP:CZ3	1:A:808:ILE:HD12	2.13	0.84
1:D:513:LEU:CB	1:D:568:GLY:HA2	2.07	0.84
2:F:317:GLY:HA3	2:F:320:SER:HB2	1.60	0.84
1:D:194:ALA:HB1	1:D:265:LEU:CD1	2.08	0.83
2:F:306:LEU:N	2:F:335:VAL:HG11	1.92	0.83
1:A:176:THR:HA	1:A:218:SER:CB	2.04	0.83
1:A:213:VAL:CG1	1:A:401:TRP:NE1	2.27	0.83
1:A:210:THR:CA	1:A:224:CYS:HB2	2.08	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:131:LEU:C	2:C:182:ARG:HD3	1.98	0.83
1:A:135:LEU:N	1:A:135:LEU:HD23	1.91	0.83
1:D:607:TRP:HD1	1:D:780:GLN:N	1.65	0.83
1:D:455:LEU:CD1	1:D:869:ARG:NH1	2.40	0.83
2:E:86:LEU:HA	2:E:90:SER:HB3	1.60	0.83
1:A:195:LEU:HG	1:A:265:LEU:HD22	1.60	0.83
1:A:192:GLU:CD	1:A:265:LEU:HD23	1.99	0.83
2:C:197:LEU:HD11	2:C:322:LEU:CG	2.08	0.83
1:A:784:GLY:N	2:C:363:ARG:HD2	1.94	0.83
1:D:195:LEU:H	1:D:265:LEU:CD2	1.90	0.83
1:D:547:ALA:HA	1:D:551:LYS:HE2	1.60	0.83
2:E:259:ARG:CZ	2:E:263:TRP:HE1	1.91	0.83
2:E:402:LEU:HD22	2:E:468:ILE:HD11	1.59	0.83
2:F:444:THR:O	2:F:447:THR:CG2	2.26	0.83
2:F:460:THR:HG23	2:F:462:MET:HB2	1.60	0.83
2:F:483:LYS:O	2:F:483:LYS:HE2	1.77	0.83
1:A:346:ASP:CB	1:A:1040:LYS:HZ3	1.88	0.83
1:A:1140:LEU:HD23	1:A:1141:VAL:H	1.43	0.83
1:A:192:GLU:CB	1:A:195:LEU:HD21	2.08	0.83
1:A:660:CYS:SG	1:A:745:PRO:HG3	2.18	0.83
1:A:608:ASP:HA	1:A:779:LEU:HB2	1.58	0.83
2:B:323:HIS:CB	2:B:332:VAL:HA	2.08	0.83
2:B:440:THR:HG22	2:B:441:VAL:H	1.42	0.83
1:D:1206:MET:SD	1:D:1210:TYR:CA	2.66	0.83
1:D:1208:ARG:HG3	1:D:1209:ARG:H	1.43	0.83
1:D:185:VAL:HG23	1:D:186:PRO:O	1.79	0.83
1:D:285:GLN:HE22	1:D:1128:ARG:HG2	1.38	0.83
1:D:436:PRO:HG3	1:D:881:PRO:HD3	1.60	0.83
1:D:959:GLN:O	1:D:963:GLU:HG2	1.78	0.83
2:F:417:TYR:HD2	2:F:417:TYR:H	0.89	0.83
2:F:451:GLY:O	2:F:452:LEU:HB2	1.77	0.83
2:B:383:VAL:HG13	2:B:413:VAL:HG13	0.87	0.83
2:C:217:PRO:HB2	2:C:219:PHE:CE1	2.13	0.83
1:D:1020:ILE:CD1	1:D:1021:SER:N	2.41	0.83
2:F:200:VAL:O	2:F:203:ARG:HB2	1.78	0.83
2:F:276:SER:O	2:F:287:ASN:CB	2.26	0.83
1:A:1026:ARG:HB3	1:A:1029:GLN:HB2	1.44	0.83
1:A:823:ARG:CB	1:A:825:VAL:HG23	2.06	0.83
1:A:83:LEU:HD22	1:A:89:GLY:HA2	1.60	0.83
2:B:323:HIS:HA	2:B:333:PRO:HD2	1.60	0.83
2:C:208:LEU:HD11	2:C:240:VAL:HB	1.59	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1027:LYS:HD2	1:D:1091:GLU:OE2	1.79	0.83
1:D:208:CYS:SG	1:D:244:LEU:HD12	2.19	0.83
1:D:887:VAL:HG23	1:D:1146:ARG:NH2	1.93	0.83
1:A:1041:TRP:C	1:A:1041:TRP:CD1	2.50	0.83
1:A:271:VAL:HG22	1:A:295:MET:SD	2.19	0.83
1:A:305:SER:HB2	1:A:308:GLN:CG	2.09	0.83
1:A:382:MET:SD	1:A:385:ILE:HG22	2.18	0.83
1:A:789:PRO:HA	1:A:792:LEU:HD11	1.60	0.83
2:B:269:SER:H	2:B:270:PRO:HD3	1.43	0.83
2:B:373:LYS:HE2	2:B:459:ASP:HA	1.61	0.83
2:C:364:LYS:H	2:C:364:LYS:HD3	1.42	0.83
2:C:395:LEU:HD21	2:C:443:VAL:CG1	2.08	0.83
2:C:424:SER:HB3	2:C:427:GLN:NE2	1.91	0.83
1:D:1206:MET:O	1:D:1206:MET:SD	2.37	0.83
1:D:436:PRO:CG	1:D:881:PRO:HD3	2.08	0.83
1:D:83:LEU:HD13	1:D:83:LEU:H	1.42	0.83
2:E:378:LEU:O	2:E:381:ILE:O	1.97	0.83
2:F:241:TRP:N	2:F:336:LEU:HB2	1.93	0.83
1:A:1043:VAL:HA	1:A:1046:GLU:N	1.92	0.83
1:A:393:MET:N	1:A:393:MET:SD	2.52	0.83
1:A:468:ASN:O	1:A:472:GLN:HG2	1.79	0.83
1:A:916:PHE:HA	1:A:919:MET:CB	2.09	0.83
2:C:308:ASP:OD1	2:C:332:VAL:O	1.95	0.83
2:C:483:LYS:HE2	2:C:483:LYS:O	1.78	0.83
1:D:990:LYS:HE2	1:D:1000:GLU:CD	1.99	0.83
2:E:106:LEU:HA	2:E:109:ASN:OD1	1.79	0.83
2:F:197:LEU:HD11	2:F:322:LEU:CG	2.07	0.83
1:A:1046:GLU:OE1	1:A:1049:TRP:HB3	1.79	0.83
1:A:1133:ILE:HG23	1:A:1136:GLU:HB2	1.61	0.83
2:E:249:ASN:ND2	2:E:285:LYS:HE2	1.93	0.83
1:D:670:PRO:HG2	1:D:673:ALA:CB	2.09	0.83
1:D:408:GLN:O	1:D:412:PRO:HD3	1.79	0.83
1:D:434:TYR:O	1:D:1130:CYS:HB3	1.79	0.83
2:E:347:LEU:O	2:E:350:LEU:HB3	1.79	0.83
2:F:197:LEU:HD23	2:F:198:ASP:N	1.93	0.83
1:A:1148:ARG:CG	1:A:1148:ARG:HH11	1.91	0.83
1:A:138:HIS:O	1:A:142:LEU:HG	1.77	0.83
1:A:573:TYR:O	1:A:574:ARG:HG3	1.78	0.83
1:A:664:GLY:O	1:A:667:GLN:HG2	1.79	0.83
1:A:853:ARG:CZ	1:A:1105:ALA:HB3	2.08	0.83
2:C:126:PHE:C	2:C:128:VAL:N	2.31	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:264:ARG:HA	2:C:270:PRO:HB3	1.60	0.83
1:D:380:GLY:O	1:D:381:THR:HG22	1.77	0.83
1:A:408:GLN:HA	1:A:411:LEU:CD2	2.09	0.83
1:A:786:ALA:HB1	1:A:790:ARG:NH1	1.94	0.83
1:D:270:ASN:O	1:D:272:SER:O	1.97	0.82
1:D:316:LYS:NZ	1:D:348:LEU:HD23	1.94	0.82
1:D:345:TRP:HA	1:D:1040:LYS:HZ3	1.40	0.82
1:D:475:SER:O	1:D:711:ALA:HB1	1.77	0.82
1:D:486:TRP:HZ2	1:D:722:ARG:CA	1.92	0.82
1:D:995:TYR:CD1	1:D:1004:LEU:HB3	2.14	0.82
1:D:914:THR:OG1	1:D:999:ASP:OD1	1.97	0.82
2:F:239:LEU:N	2:F:338:VAL:CG1	2.42	0.82
2:F:387:VAL:HG12	2:F:443:VAL:CG2	2.09	0.82
2:F:445:GLU:O	2:F:448:LEU:N	2.12	0.82
1:A:462:SER:O	1:A:466:LEU:HD23	1.78	0.82
1:A:484:TRP:O	1:A:488:LEU:HG	1.78	0.82
1:A:452:TYR:CA	1:A:869:ARG:HH21	1.91	0.82
1:A:927:ARG:CG	1:A:928:GLY:H	1.91	0.82
2:C:197:LEU:HD11	2:C:322:LEU:HG	1.58	0.82
2:C:431:LYS:HZ1	2:C:435:MET:HG2	1.40	0.82
1:D:167:PRO:C	1:D:169:TRP:H	1.79	0.82
1:D:271:VAL:HG12	1:D:272:SER:N	1.92	0.82
1:D:304:LEU:H	1:D:309:ARG:NH2	1.77	0.82
1:D:414:PHE:HA	1:D:417:ARG:HH11	1.42	0.82
1:A:1113:LEU:HB3	1:A:1117:LYS:HE3	1.61	0.82
1:A:1166:TYR:CD1	1:A:1166:TYR:N	2.46	0.82
1:A:379:LYS:HG2	1:A:379:LYS:O	1.79	0.82
1:A:490:TRP:O	1:A:580:LEU:HD13	1.78	0.82
1:A:743:ASP:C	1:A:744:ILE:HD13	1.99	0.82
2:B:261:GLN:HA	2:B:264:ARG:CD	2.09	0.82
2:B:313:HIS:O	2:B:316:PRO:HD3	1.79	0.82
2:C:247:THR:HB	2:C:251:TRP:HD1	1.40	0.82
2:C:356:LEU:HB3	2:C:359:ASN:CG	1.99	0.82
1:A:783:PRO:HG2	2:C:363:ARG:CZ	2.09	0.82
1:D:1065:ALA:C	1:D:1073:PRO:HG3	1.99	0.82
1:D:256:PRO:HG2	1:D:284:ILE:CD1	2.09	0.82
1:D:350:ILE:HG22	1:D:352:SER:HB2	1.57	0.82
1:D:496:LYS:HE3	1:D:571:GLY:CA	2.09	0.82
2:E:428:LEU:O	2:E:431:LYS:HB3	1.78	0.82
2:F:453:ILE:CG2	2:F:455:LEU:H	1.92	0.82
2:F:454:HIS:HA	2:F:468:ILE:CD1	2.08	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1047:ARG:NH2	1:A:1050:LYS:HD2	1.94	0.82
1:A:1060:LYS:HD3	1:A:1073:PRO:HB2	1.60	0.82
1:A:1109:LEU:HD12	1:A:1113:LEU:HD21	1.60	0.82
2:B:269:SER:N	2:B:270:PRO:CD	2.42	0.82
2:C:455:LEU:HB3	2:C:456:ARG:NE	1.94	0.82
1:D:213:VAL:C	1:D:401:TRP:HZ2	1.82	0.82
2:E:377:CYS:SG	2:E:378:LEU:HD12	2.18	0.82
2:E:451:GLY:O	2:E:452:LEU:HD13	1.79	0.82
2:F:447:THR:HA	2:F:452:LEU:HA	1.61	0.82
1:A:996:ARG:CZ	1:A:1000:GLU:HB3	2.09	0.82
1:A:213:VAL:C	1:A:401:TRP:HZ2	1.83	0.82
1:A:490:TRP:HA	1:A:580:LEU:CD2	2.07	0.82
1:A:649:TYR:O	1:A:653:GLU:N	2.11	0.82
1:A:827:ARG:HA	1:A:831:TYR:CZ	2.15	0.82
1:A:897:TRP:HE1	1:A:901:VAL:CG1	1.93	0.82
2:E:389:ARG:C	2:E:395:LEU:HG	1.98	0.82
2:F:320:SER:HB3	2:F:321:LYS:HG3	1.60	0.82
1:D:367:PRO:CG	1:D:368:PRO:HD3	2.09	0.82
1:A:223:TRP:O	1:A:227:ARG:HB2	1.79	0.82
1:A:601:LYS:NZ	1:A:721:ALA:HB1	1.95	0.82
2:C:208:LEU:HD11	2:C:240:VAL:C	1.99	0.82
2:C:398:VAL:O	2:C:402:LEU:HD13	1.79	0.82
2:C:446:THR:CA	2:C:450:ASN:HB3	2.09	0.82
1:D:312:TRP:C	1:D:316:LYS:HD3	2.00	0.82
1:D:960:PRO:HA	1:D:963:GLU:HG3	1.60	0.82
2:F:341:ASP:C	2:F:342:LEU:HD13	1.99	0.82
1:A:1047:ARG:HD3	1:A:1047:ARG:O	1.78	0.82
1:A:260:ASP:H	1:A:264:GLN:NE2	1.77	0.82
1:A:275:ARG:NH2	1:A:845:VAL:CA	2.43	0.82
1:A:471:CYS:HB3	1:A:472:GLN:OE1	1.78	0.82
1:D:1203:PRO:O	1:D:1207:GLU:N	2.12	0.82
1:D:1207:GLU:CA	1:D:1210:TYR:HD2	1.93	0.82
1:A:1025:LEU:C	1:A:1026:ARG:HG3	2.00	0.82
1:A:1198:LYS:HG3	1:A:1204:THR:HG21	1.60	0.82
1:D:1052:GLY:HA2	1:D:1055:SER:HB3	1.61	0.82
1:D:869:ARG:O	1:D:1200:PRO:CG	2.26	0.82
1:D:1207:GLU:O	1:D:1210:TYR:CD2	2.33	0.82
1:D:438:ASN:N	1:D:441:TRP:CD1	2.48	0.82
2:E:407:LEU:HD23	2:E:412:SER:HA	1.61	0.82
2:F:253:ASP:O	2:F:257:ARG:HD2	1.79	0.82
1:D:1206:MET:HG2	2:F:257:ARG:HH11	1.40	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:455:LEU:C	2:F:456:ARG:HE	1.83	0.82
1:A:1114:VAL:HA	1:A:1117:LYS:HD2	1.61	0.82
1:A:1131:ILE:CD1	1:A:1133:ILE:HG22	2.09	0.82
1:A:1220:ILE:HD12	1:A:1220:ILE:N	1.95	0.82
1:A:658:LYS:HG3	1:A:713:PRO:CB	2.06	0.82
2:C:372:LEU:CD1	2:C:436:SER:HB2	2.09	0.82
2:C:471:LEU:O	2:C:475:LEU:HG	1.80	0.82
1:D:890:ASP:OD1	1:D:1191:LYS:HD3	1.79	0.82
1:D:175:TRP:HB2	1:D:185:VAL:HG13	1.60	0.82
1:D:193:ARG:HA	1:D:193:ARG:NE	1.88	0.82
1:D:275:ARG:CD	1:D:843:GLN:H	1.92	0.82
1:D:375:GLU:O	1:D:379:LYS:HG3	1.79	0.82
1:D:549:LEU:H	1:D:551:LYS:HZ3	1.26	0.82
1:D:768:LYS:CE	1:D:769:ASP:H	1.93	0.82
1:D:808:ILE:O	1:D:811:GLN:CB	2.27	0.82
2:F:415:PRO:HB3	2:F:417:TYR:CE2	2.15	0.82
2:F:471:LEU:O	2:F:475:LEU:HG	1.80	0.82
1:A:885:THR:HB	1:A:1146:ARG:HE	1.44	0.82
1:A:165:LYS:HB2	1:A:178:TYR:HA	1.62	0.82
1:A:175:TRP:CD1	1:A:183:GLU:HB2	2.14	0.82
1:A:190:PRO:CB	1:A:192:GLU:CG	2.56	0.82
2:C:115:TRP:HH2	2:C:210:GLN:HG2	1.44	0.82
2:C:326:ASP:OD1	2:C:329:LYS:HG2	1.80	0.82
2:C:344:ARG:HA	2:C:347:LEU:HD22	1.62	0.82
1:D:498:LYS:HG2	1:D:499:LYS:H	1.45	0.82
1:D:1069:ILE:O	1:D:1071:ARG:CG	2.26	0.82
1:D:112:LEU:CD2	1:D:917:GLY:O	2.28	0.82
1:D:364:VAL:HG12	1:D:364:VAL:O	1.80	0.82
1:D:869:ARG:H	1:D:1200:PRO:HB2	1.45	0.82
1:A:228:LEU:HD12	1:A:386:ARG:HB3	1.62	0.82
1:A:626:GLY:HA3	1:A:628:ARG:O	1.80	0.82
1:A:784:GLY:H	2:C:363:ARG:HD2	1.41	0.82
1:D:422:VAL:HG11	1:D:1111:LEU:HD12	1.60	0.81
2:F:375:HIS:H	2:F:378:LEU:CD2	1.93	0.81
2:F:447:THR:O	2:F:452:LEU:CD1	2.26	0.81
1:A:1208:ARG:N	1:A:1210:TYR:CD1	2.48	0.81
1:D:851:THR:HG23	1:D:1102:GLN:HB3	1.63	0.81
1:D:931:LEU:HB3	1:D:934:LYS:HZ2	1.45	0.81
1:A:1040:LYS:HB3	1:A:1044:VAL:CG1	2.09	0.81
1:A:1160:THR:O	1:A:1164:PHE:HD1	1.63	0.81
1:A:115:GLN:N	1:A:116:PRO:HD2	1.93	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:490:TRP:CA	1:A:580:LEU:HD22	2.08	0.81
2:C:421:MET:HE2	2:C:422:GLN:HB2	1.62	0.81
1:D:1137:VAL:HG12	1:D:1139:TYR:CE1	2.15	0.81
1:D:196:VAL:CG2	1:D:215:ILE:HD12	2.11	0.81
1:D:847:ALA:HA	1:D:852:ARG:HA	1.62	0.81
2:F:125:VAL:HG23	2:F:126:PHE:H	1.42	0.81
2:F:455:LEU:CA	2:F:456:ARG:HE	1.92	0.81
1:D:1157:ASN:O	1:D:1161:ARG:HD2	1.80	0.81
1:D:1206:MET:HG3	2:F:257:ARG:HH11	1.46	0.81
1:D:347:TRP:CD1	1:D:349:ASP:OD2	2.32	0.81
1:D:457:ARG:HH11	1:D:457:ARG:HB3	1.44	0.81
1:D:618:HIS:HB2	1:D:725:PRO:CG	2.08	0.81
2:E:432:TYR:CB	2:E:437:ILE:CD1	2.44	0.81
1:A:213:VAL:CG1	1:A:397:ALA:HB1	2.10	0.81
1:A:512:LYS:O	1:A:513:LEU:HD23	1.80	0.81
2:C:402:LEU:HG	2:C:472:LYS:NZ	1.95	0.81
2:C:446:THR:CA	2:C:450:ASN:CB	2.58	0.81
2:C:467:HIS:O	2:C:471:LEU:HD11	1.80	0.81
1:A:608:ASP:O	1:A:620:TRP:CH2	2.33	0.81
1:A:794:ILE:HA	1:A:797:MET:SD	2.20	0.81
2:B:101:PRO:HA	2:C:128:VAL:CG2	2.09	0.81
2:B:192:HIS:CE1	2:C:80:SER:HB3	2.16	0.81
2:C:299:LEU:N	2:C:299:LEU:HD23	1.95	0.81
2:C:344:ARG:HA	2:C:347:LEU:CB	2.11	0.81
1:D:233:TYR:O	1:D:234:SER:HB2	1.79	0.81
1:D:718:ALA:O	1:D:722:ARG:HD2	1.80	0.81
1:D:798:ILE:HG23	1:D:869:ARG:NH1	1.95	0.81
2:F:208:LEU:HD22	2:F:242:PHE:CE2	2.15	0.81
2:F:341:ASP:CG	2:F:342:LEU:H	1.83	0.81
1:D:572:TRP:HZ2	2:F:470:LYS:HB3	1.45	0.81
1:A:282:TYR:O	1:A:283:LEU:HB2	1.79	0.81
1:A:575:LYS:HG3	1:A:576:LEU:N	1.94	0.81
1:A:628:ARG:O	1:A:629:ASP:HB2	1.79	0.81
1:A:653:GLU:HG3	1:A:656:TYR:CE1	2.15	0.81
1:A:866:ARG:HB3	1:A:867:PRO:HD3	1.59	0.81
1:A:900:ALA:HA	1:A:915:ALA:CB	2.11	0.81
2:B:344:ARG:H	2:B:344:ARG:HD2	1.43	0.81
2:C:384:ALA:HB2	2:C:437:ILE:HD12	1.60	0.81
2:B:133:HIS:CD2	2:C:231:ILE:HG21	2.14	0.81
1:D:265:LEU:HD21	1:D:267:VAL:HG13	1.61	0.81
1:D:375:GLU:C	1:D:379:LYS:HG3	2.01	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:493:GLN:O	1:D:494:GLU:HG2	1.81	0.81
1:A:1066:THR:OG1	1:A:1073:PRO:HD2	1.81	0.81
1:A:655:LEU:HD13	1:A:719:LEU:C	2.00	0.81
2:B:74:GLN:O	2:B:77:HIS:HA	1.80	0.81
1:D:1218:LEU:CB	1:D:1223:ILE:HG22	2.09	0.81
1:D:275:ARG:HE	1:D:842:PRO:CA	1.91	0.81
1:D:439:GLN:HG2	1:D:835:GLY:CA	2.11	0.81
1:D:903:GLY:HA3	1:D:1002:GLU:HB2	1.63	0.81
2:F:78:PHE:CD1	2:F:103:GLY:HA2	2.15	0.81
1:A:1060:LYS:HD3	1:A:1073:PRO:HB3	1.62	0.81
1:A:249:VAL:CG1	1:A:817:PRO:HG3	2.11	0.81
1:A:869:ARG:HG3	1:A:869:ARG:HH11	1.46	0.81
1:A:883:GLY:C	1:A:884:TYR:HD1	1.83	0.81
1:D:120:LEU:H	1:D:120:LEU:HD22	1.45	0.81
1:D:962:ALA:HB3	1:D:982:ALA:HB1	1.63	0.81
2:F:300:ILE:HG22	2:F:301:GLU:N	1.96	0.81
2:F:447:THR:HG23	2:F:448:LEU:HG	1.63	0.81
1:A:81:ARG:CZ	1:A:126:ARG:HA	2.09	0.81
2:B:323:HIS:HB2	2:B:331:VAL:O	1.80	0.81
2:C:374:LEU:HD12	2:C:379:ALA:HA	1.61	0.81
1:D:1128:ARG:HG3	1:D:1129:PHE:N	1.95	0.81
1:D:1208:ARG:CZ	2:F:285:LYS:HD3	2.11	0.81
1:D:135:LEU:CD2	1:D:135:LEU:H	1.94	0.81
2:E:79:LEU:HA	2:E:100:GLY:N	1.95	0.81
2:F:363:ARG:NH1	2:F:363:ARG:HB3	1.96	0.81
2:F:447:THR:C	2:F:452:LEU:HA	2.00	0.81
2:F:467:HIS:O	2:F:471:LEU:HD13	1.80	0.81
1:A:1158:LEU:CD2	1:A:1178:ALA:O	2.29	0.81
1:A:424:LEU:N	1:A:424:LEU:HD23	1.96	0.81
1:A:604:ALA:HB1	1:A:717:LEU:HD11	1.63	0.81
2:C:467:HIS:O	2:C:471:LEU:HD13	1.78	0.81
2:C:134:LYS:HB3	2:C:180:LYS:CD	2.09	0.81
1:D:170:ALA:HB1	1:D:171:TRP:O	1.81	0.81
1:D:177:ARG:HG3	1:D:178:TYR:N	1.96	0.81
1:D:549:LEU:N	1:D:551:LYS:HZ2	1.77	0.81
1:D:556:THR:HA	1:D:559:LEU:CD2	2.10	0.81
1:D:575:LYS:HE3	1:D:581:ASP:O	1.81	0.81
1:D:78:MET:O	1:D:79:LEU:HD12	1.79	0.81
1:A:370:PRO:O	1:A:371:LYS:HG2	1.79	0.81
1:A:665:LYS:HG2	1:A:666:GLN:N	1.96	0.81
2:B:267:ALA:CB	2:B:270:PRO:HB2	2.10	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:78:PHE:HE1	2:C:103:GLY:HA2	1.46	0.81
2:C:130:ALA:HB1	2:C:182:ARG:HH12	1.46	0.81
1:D:667:GLN:O	1:D:671:GLN:HG2	1.80	0.81
1:D:1029:GLN:HG3	1:D:1030:ARG:HE	1.46	0.81
1:D:194:ALA:HA	1:D:265:LEU:CB	2.10	0.81
1:D:350:ILE:C	1:D:352:SER:H	1.83	0.81
1:D:715:GLN:HG2	1:D:717:LEU:HG	1.62	0.81
1:D:116:PRO:HA	1:D:921:LEU:HD13	1.62	0.81
2:E:77:HIS:HB3	2:E:101:PRO:HD2	1.61	0.81
2:F:247:THR:CB	2:F:251:TRP:HD1	1.93	0.81
1:A:363:ARG:NH2	1:A:409:GLN:HB2	1.93	0.81
1:A:925:LYS:CG	1:A:926:SER:N	2.41	0.81
2:C:275:SER:HB2	2:C:287:ASN:ND2	1.95	0.81
1:A:163:PRO:CB	1:A:164:PRO:HD3	2.11	0.81
1:D:190:PRO:HB3	1:D:217:PRO:HB2	1.63	0.80
1:D:246:PRO:C	1:D:247:LEU:HD12	2.00	0.80
1:D:269:HIS:CE1	1:D:291:PHE:HA	2.16	0.80
1:D:435:LEU:O	1:D:839:ALA:HB1	1.81	0.80
1:D:719:LEU:HD12	1:D:720:THR:N	1.95	0.80
1:D:797:MET:CB	1:D:869:ARG:CD	2.59	0.80
2:F:299:LEU:HD23	2:F:299:LEU:H	1.45	0.80
1:A:606:THR:OG1	1:A:612:LEU:C	2.20	0.80
1:A:955:TYR:HB3	1:A:1098:ASN:CG	2.01	0.80
1:A:988:ALA:O	1:A:998:SER:OG	1.97	0.80
1:D:737:GLY:N	1:D:750:PHE:CZ	2.48	0.80
1:D:916:PHE:O	1:D:920:THR:N	2.14	0.80
1:A:312:TRP:HZ2	1:A:346:ASP:O	1.63	0.80
1:A:711:ALA:O	1:A:713:PRO:HD3	1.79	0.80
1:A:931:LEU:HD23	1:A:932:HIS:N	1.96	0.80
1:A:633:LYS:C	1:A:633:LYS:HD2	2.00	0.80
2:B:448:LEU:O	2:B:448:LEU:HD23	1.81	0.80
1:D:107:LEU:CD2	1:D:913:CYS:HA	2.12	0.80
1:D:134:ASN:HB3	1:D:137:GLN:HB2	1.62	0.80
1:D:256:PRO:HG2	1:D:284:ILE:HD11	1.62	0.80
1:D:288:ARG:HH11	1:D:288:ARG:HG3	1.46	0.80
1:D:389:PHE:HD1	1:D:390:GLN:N	1.77	0.80
1:D:717:LEU:HD23	1:D:717:LEU:N	1.95	0.80
1:D:767:ALA:HB2	1:D:771:LEU:HD22	1.60	0.80
2:F:430:SER:O	2:F:434:GLU:HG3	1.81	0.80
1:A:228:LEU:HD23	1:A:228:LEU:C	2.00	0.80
1:A:848:GLY:O	1:A:852:ARG:HA	1.81	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:147:SER:O	1:A:151:LEU:HG	1.80	0.80
1:D:1065:ALA:O	1:D:1073:PRO:HG3	1.81	0.80
1:D:173:GLU:O	1:D:220:TRP:HB3	1.81	0.80
1:D:597:ARG:NH1	1:D:724:GLY:O	2.14	0.80
1:D:612:LEU:CD1	1:D:612:LEU:N	2.44	0.80
2:E:382:LYS:CE	2:E:412:SER:O	2.25	0.80
1:A:552:LEU:HD22	2:C:448:LEU:HD23	1.62	0.80
2:C:310:GLU:O	2:C:313:HIS:HB3	1.81	0.80
1:D:1138:ARG:CG	1:D:1138:ARG:HH11	1.94	0.80
1:D:298:HIS:CD2	1:D:407:PHE:HB2	2.17	0.80
1:D:608:ASP:OD1	1:D:611:PRO:CD	2.30	0.80
1:D:608:ASP:HA	1:D:620:TRP:CH2	2.16	0.80
1:D:917:GLY:O	1:D:920:THR:HB	1.81	0.80
1:D:992:LEU:CD1	1:D:993:ARG:H	1.93	0.80
1:A:166:PRO:HG2	1:A:398:GLN:HE22	1.46	0.80
1:A:606:THR:CG2	1:A:613:HIS:H	1.95	0.80
1:A:607:TRP:CZ2	1:A:671:GLN:HA	2.16	0.80
1:A:793:GLU:CA	1:A:796:LYS:HD3	2.11	0.80
2:B:262:TRP:O	2:B:265:LYS:CB	2.29	0.80
2:B:385:LEU:HB3	2:B:441:VAL:HG23	1.63	0.80
1:D:1025:LEU:O	1:D:1026:ARG:C	2.17	0.80
1:D:1202:ASN:O	1:D:1206:MET:N	2.15	0.80
1:D:194:ALA:HB1	1:D:265:LEU:HD13	1.63	0.80
1:D:255:SER:HB3	1:D:256:PRO:C	2.02	0.80
1:D:252:GLY:O	1:D:256:PRO:HG3	1.82	0.80
1:D:282:TYR:CE2	1:D:432:VAL:HA	2.16	0.80
1:D:81:ARG:HG3	1:D:82:GLY:N	1.97	0.80
1:D:887:VAL:O	1:D:1139:TYR:N	2.15	0.80
2:E:247:THR:CG2	2:E:250:GLN:HE21	1.95	0.80
2:E:283:GLY:CA	2:E:306:LEU:HA	2.11	0.80
2:E:320:SER:O	2:E:323:HIS:CD2	2.35	0.80
1:A:1030:ARG:HG2	1:A:1040:LYS:NZ	1.97	0.80
1:A:577:CYS:HB3	1:A:578:PRO:CD	2.10	0.80
2:B:117:SER:CB	2:B:262:TRP:NE1	2.44	0.80
2:C:203:ARG:O	2:C:204:LEU:CB	2.29	0.80
2:C:397:GLN:HG2	2:C:398:VAL:HG13	1.62	0.80
2:C:454:HIS:CB	2:C:466:MET:O	2.30	0.80
2:E:388:GLY:C	2:E:395:LEU:CD1	2.49	0.80
1:D:424:LEU:O	1:D:428:LEU:CD1	2.29	0.80
1:D:621:GLY:HA2	1:D:748:TRP:HB2	1.62	0.80
2:E:384:ALA:O	2:E:414:TRP:CB	2.30	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:244:PRO:CD	2:F:247:THR:HG21	2.10	0.80
1:A:114:GLY:N	1:A:116:PRO:HD2	1.97	0.80
1:A:1196:ASP:HA	1:A:1207:GLU:OE1	1.81	0.80
1:A:176:THR:HG23	1:A:177:ARG:N	1.97	0.80
1:A:76:ILE:HB	1:A:911:HIS:CE1	2.16	0.80
2:F:317:GLY:HA3	2:F:320:SER:N	1.96	0.80
2:B:402:LEU:O	2:B:406:LEU:HD23	1.80	0.80
1:D:1010:LEU:HD13	1:D:1100:VAL:HG21	1.62	0.80
1:D:1124:ALA:HB1	1:D:1148:ARG:HH11	1.46	0.80
1:D:302:SER:CB	1:D:351:SER:HB3	2.11	0.80
1:D:823:ARG:HA	1:D:826:ILE:CD1	2.11	0.80
1:D:926:SER:CB	1:D:930:ASP:HB2	2.12	0.80
2:E:199:LEU:HG	2:E:200:VAL:HG23	1.64	0.80
1:A:897:TRP:HZ3	1:A:1108:TYR:HE1	1.29	0.80
2:C:357:THR:HA	2:C:370:LYS:NZ	1.96	0.80
1:D:1010:LEU:HD23	1:D:1015:THR:HA	1.64	0.80
1:D:267:VAL:CB	1:D:269:HIS:CD2	2.64	0.80
1:D:271:VAL:HG12	1:D:272:SER:H	1.46	0.80
2:E:380:PRO:N	2:E:438:LEU:HD11	1.97	0.80
2:F:357:THR:HA	2:F:370:LYS:NZ	1.96	0.80
1:A:1006:ARG:HA	1:A:1006:ARG:CZ	2.12	0.80
1:A:211:LEU:HG	1:A:221:TYR:CG	2.16	0.80
1:A:213:VAL:CA	1:A:221:TYR:HE2	1.93	0.80
1:A:606:THR:HG21	1:A:612:LEU:CB	2.10	0.80
1:A:863:SER:O	1:A:873:GLU:HG3	1.81	0.80
2:B:72:ILE:HG13	2:B:73:CYS:N	1.97	0.80
2:C:432:TYR:CG	2:C:437:ILE:HD11	2.17	0.80
1:D:1026:ARG:N	1:D:1026:ARG:HD3	1.97	0.80
1:D:1165:ALA:HB3	1:D:1166:TYR:CE1	2.16	0.80
1:D:259:ARG:HG3	1:D:261:TRP:HD1	1.46	0.80
1:D:256:PRO:HB3	1:D:281:GLN:HG3	1.64	0.80
1:D:636:THR:O	1:D:640:LEU:HG	1.82	0.80
2:F:241:TRP:HZ3	2:F:255:TRP:NE1	1.79	0.80
2:F:244:PRO:HD2	2:F:247:THR:CG2	2.12	0.80
1:A:365:TYR:O	1:A:366:VAL:HG22	1.82	0.80
1:A:725:PRO:O	1:A:726:LYS:HB2	1.81	0.80
2:C:474:PHE:HA	2:C:477:LYS:CD	2.12	0.80
1:D:201:VAL:HG22	1:D:208:CYS:O	1.81	0.79
1:D:606:THR:HG23	1:D:607:TRP:N	1.96	0.79
2:E:293:PHE:O	2:E:296:GLY:N	2.14	0.79
2:F:346:MET:HE3	2:F:347:LEU:HD12	1.63	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:821:LEU:O	1:A:826:ILE:HG23	1.80	0.79
1:A:893:SER:HA	1:A:896:LEU:HD23	1.64	0.79
1:A:898:ILE:HD11	1:A:1101:VAL:HG13	1.64	0.79
2:C:336:LEU:C	2:C:336:LEU:HD23	2.03	0.79
2:C:446:THR:HA	2:C:450:ASN:HB3	1.62	0.79
1:D:78:MET:HB3	1:D:1175:GLN:HB3	1.64	0.79
1:D:914:THR:HG21	1:D:999:ASP:HB3	1.63	0.79
1:A:1041:TRP:O	1:A:1042:GLU:HB2	1.81	0.79
1:A:486:TRP:CZ3	1:A:598:VAL:HG21	2.16	0.79
1:A:656:TYR:CD1	1:A:744:ILE:HD12	2.17	0.79
2:C:78:PHE:CD1	2:C:103:GLY:HA2	2.17	0.79
2:C:300:ILE:HG22	2:C:301:GLU:HG3	1.61	0.79
2:C:300:ILE:HG23	2:C:342:LEU:CD1	2.10	0.79
2:C:403:PHE:HE1	2:C:415:PRO:HG3	1.47	0.79
1:D:1027:LYS:HB3	1:D:1091:GLU:CA	2.11	0.79
1:D:1180:PHE:O	1:D:1182:ALA:N	2.15	0.79
1:D:1213:PRO:O	1:D:1214:GLN:HB2	1.82	0.79
1:D:271:VAL:CG1	1:D:272:SER:H	1.87	0.79
1:D:459:MET:HE2	1:D:594:LEU:HD11	1.64	0.79
1:D:649:TYR:CD2	1:D:652:ILE:HG21	2.16	0.79
1:D:741:ASP:CB	1:D:748:TRP:HH2	1.94	0.79
1:A:670:PRO:HG2	1:A:671:GLN:NE2	1.96	0.79
1:A:81:ARG:HD3	1:A:125:LEU:O	1.82	0.79
2:B:385:LEU:HD23	2:B:414:TRP:O	1.82	0.79
2:C:300:ILE:CG2	2:C:342:LEU:HD11	2.11	0.79
2:C:344:ARG:CA	2:C:347:LEU:HD22	2.11	0.79
2:B:450:ASN:HD22	2:B:452:LEU:HD23	1.47	0.79
1:D:206:GLY:HA2	1:D:245:ILE:CG2	2.12	0.79
1:D:797:MET:HB2	1:D:869:ARG:HD3	1.64	0.79
2:E:376:PRO:O	2:E:380:PRO:CD	2.26	0.79
2:F:417:TYR:CD2	2:F:417:TYR:N	2.43	0.79
1:A:196:VAL:CB	1:A:401:TRP:HZ3	1.95	0.79
1:A:513:LEU:HB2	1:A:568:GLY:HA2	0.85	0.79
1:A:78:MET:HG2	1:A:79:LEU:N	1.97	0.79
1:A:843:GLN:O	1:A:843:GLN:HG2	1.80	0.79
2:B:259:ARG:O	2:B:260:LEU:C	2.17	0.79
2:B:268:MET:N	2:B:270:PRO:HD2	1.97	0.79
2:C:303:LEU:HA	2:C:337:SER:CB	2.11	0.79
2:C:456:ARG:HB2	2:C:463:LYS:C	2.02	0.79
1:D:1133:ILE:O	1:D:1135:ASP:N	2.16	0.79
1:D:913:CYS:SG	1:D:914:THR:N	2.56	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1126:ASP:HB3	1:A:1141:VAL:HG13	1.64	0.79
1:A:1166:TYR:N	1:A:1166:TYR:HD1	1.79	0.79
1:A:374:ARG:HE	1:A:392:LEU:CD1	1.94	0.79
1:D:1141:VAL:N	1:D:1146:ARG:HH21	1.80	0.79
2:E:372:LEU:HG	2:E:373:LYS:H	1.46	0.79
2:F:442:LEU:N	2:F:455:LEU:HD22	1.97	0.79
2:F:454:HIS:CB	2:F:466:MET:O	2.29	0.79
1:A:1047:ARG:CA	1:A:1047:ARG:HH11	1.96	0.79
1:A:176:THR:CG2	1:A:177:ARG:H	1.87	0.79
1:A:273:PHE:O	1:A:276:ALA:HB3	1.83	0.79
1:A:150:TYR:OH	1:A:288:ARG:O	2.01	0.79
1:A:605:LEU:CA	1:A:781:ALA:HB3	2.13	0.79
1:A:944:GLU:HA	1:A:947:LYS:HE3	1.64	0.79
2:F:328:ARG:O	2:F:329:LYS:HD2	1.82	0.79
1:A:95:PRO:O	1:A:100:VAL:HG23	1.82	0.79
1:D:167:PRO:C	1:D:169:TRP:N	2.34	0.79
1:D:87:ILE:HG21	1:D:1166:TYR:OH	1.82	0.79
1:D:1209:ARG:HH22	2:F:256:LEU:HD22	1.48	0.79
1:A:468:ASN:ND2	1:A:578:PRO:CG	2.45	0.79
1:A:448:ALA:HB1	1:A:870:VAL:HG12	1.61	0.79
1:A:999:ASP:O	1:A:1000:GLU:HG3	1.83	0.79
2:B:412:SER:O	2:B:413:VAL:HG22	1.82	0.79
2:C:114:TRP:HE1	2:C:259:ARG:CZ	1.95	0.79
1:D:1207:GLU:HA	1:D:1210:TYR:CB	2.12	0.79
1:D:177:ARG:CG	1:D:178:TYR:N	2.45	0.79
1:D:284:ILE:HD12	1:D:284:ILE:N	1.98	0.79
1:D:645:VAL:HG13	1:D:650:ARG:HG2	1.64	0.79
1:D:649:TYR:HA	1:D:652:ILE:CB	2.09	0.79
1:D:768:LYS:CD	1:D:769:ASP:H	1.94	0.79
2:F:453:ILE:HG22	2:F:455:LEU:N	1.98	0.79
1:A:1041:TRP:CD1	1:A:1043:VAL:HB	2.18	0.79
1:A:1131:ILE:CG1	1:A:1133:ILE:HG22	2.13	0.79
1:A:118:VAL:N	1:A:119:PRO:CD	2.46	0.79
2:C:241:TRP:CZ3	2:C:255:TRP:NE1	2.51	0.79
2:C:251:TRP:HA	2:C:251:TRP:CE3	2.17	0.79
2:E:284:ARG:O	2:E:285:LYS:HB3	1.83	0.79
1:D:1027:LYS:HB2	1:D:1096:ARG:HE	1.48	0.79
1:D:135:LEU:HD23	1:D:135:LEU:N	1.98	0.79
1:D:845:VAL:HG12	1:D:846:THR:O	1.83	0.79
2:E:384:ALA:O	2:E:414:TRP:HB3	1.82	0.79
2:E:448:LEU:HD23	2:E:449:GLU:N	1.97	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1060:LYS:HE3	1:A:1065:ALA:N	1.96	0.79
1:A:1155:ILE:HA	1:A:1158:LEU:HD12	1.64	0.79
1:A:371:LYS:C	1:A:372:GLU:HG2	2.02	0.79
1:A:608:ASP:HA	1:A:779:LEU:CB	2.12	0.79
2:B:437:ILE:HG22	2:B:438:LEU:H	1.46	0.79
2:C:115:TRP:CH2	2:C:210:GLN:HG2	2.18	0.79
1:D:1026:ARG:NE	1:D:1029:GLN:OE1	2.16	0.79
1:D:195:LEU:N	1:D:265:LEU:HD22	1.98	0.79
1:D:546:ARG:C	1:D:551:LYS:HZ1	1.86	0.79
1:D:656:TYR:CD1	1:D:744:ILE:HD12	2.18	0.79
1:D:618:HIS:HA	1:D:762:VAL:CG2	2.13	0.79
2:E:123:GLU:HG2	2:F:415:PRO:HG2	1.64	0.79
2:F:241:TRP:HB3	2:F:336:LEU:HB2	1.65	0.79
2:F:303:LEU:CA	2:F:337:SER:OG	2.31	0.79
2:F:454:HIS:HB2	2:F:466:MET:C	2.02	0.79
1:A:1110:HIS:O	1:A:1114:VAL:HG23	1.82	0.79
1:A:253:ALA:HB3	1:A:256:PRO:CB	2.13	0.79
1:A:455:LEU:HD12	1:A:798:ILE:HG22	1.65	0.79
2:C:203:ARG:O	2:C:204:LEU:HB2	1.83	0.79
2:C:387:VAL:HG11	2:C:399:CYS:CB	2.11	0.79
1:D:1189:LEU:N	1:D:1189:LEU:HD22	1.97	0.78
1:D:612:LEU:N	1:D:612:LEU:HD13	1.98	0.78
1:D:605:LEU:HD22	1:D:781:ALA:HB2	1.63	0.78
1:D:996:ARG:N	1:D:996:ARG:CD	2.46	0.78
2:E:369:ARG:HA	2:E:430:SER:HB3	1.63	0.78
2:F:389:ARG:HG3	2:F:445:GLU:OE2	1.83	0.78
1:A:1016:GLU:HA	1:A:1021:SER:O	1.83	0.78
1:D:212:ALA:O	1:D:213:VAL:CB	2.31	0.78
1:D:412:PRO:O	1:D:416:GLU:HG3	1.84	0.78
1:D:995:TYR:CD2	1:D:995:TYR:O	2.35	0.78
2:F:245:PRO:HG2	2:F:246:ARG:HD3	1.64	0.78
1:A:411:LEU:O	1:A:415:LEU:HD22	1.83	0.78
1:A:477:GLU:HG2	1:A:482:ASP:OD2	1.83	0.78
1:A:653:GLU:HA	1:A:656:TYR:CE1	2.18	0.78
1:D:586:THR:N	1:D:587:PRO:CD	2.45	0.78
1:D:851:THR:CG2	1:D:1102:GLN:HB3	2.13	0.78
1:D:352:SER:CB	1:D:352:SER:C	2.51	0.78
1:D:559:LEU:CG	1:D:560:PRO:HD2	2.13	0.78
1:D:775:GLU:CA	1:D:780:GLN:HG2	2.13	0.78
2:E:109:ASN:HB2	2:E:378:LEU:HA	1.65	0.78
2:E:188:GLY:O	2:E:191:GLU:HG2	1.82	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1206:MET:N	1:A:1209:ARG:HB2	1.98	0.78
1:A:276:ALA:HA	1:A:279:ARG:CD	2.14	0.78
2:C:354:PHE:HE1	2:C:370:LYS:HB3	1.48	0.78
1:D:273:PHE:H	1:D:275:ARG:HB3	1.48	0.78
1:D:269:HIS:CD2	1:D:291:PHE:CD2	2.71	0.78
1:D:613:HIS:CE1	1:D:717:LEU:HD13	2.18	0.78
2:E:456:ARG:HA	2:E:462:MET:O	1.82	0.78
2:E:67:GLU:HG2	2:E:70:LEU:HD11	1.64	0.78
2:E:68:ALA:O	2:E:72:ILE:HD13	1.82	0.78
1:A:996:ARG:NH1	1:A:1004:LEU:N	2.31	0.78
1:A:1006:ARG:HH12	1:A:1009:ASN:HB2	1.48	0.78
1:A:1157:ASN:HB3	1:A:1161:ARG:CZ	2.12	0.78
1:A:316:LYS:HZ3	1:A:350:ILE:HG13	1.48	0.78
1:A:497:GLN:HA	1:A:516:GLU:HB3	1.63	0.78
2:B:110:LEU:CD2	2:B:342:LEU:HD21	2.14	0.78
2:C:439:PHE:HD1	2:C:458:ARG:HB2	1.47	0.78
2:E:389:ARG:NH2	2:E:391:PRO:O	2.16	0.78
2:C:317:GLY:HA3	2:C:320:SER:N	1.98	0.78
1:D:101:ARG:HH11	1:D:101:ARG:CG	1.97	0.78
1:A:563:PRO:C	1:A:564:GLN:HE21	1.86	0.78
1:D:1056:GLU:HA	1:D:1059:ASN:HD21	1.47	0.78
1:D:282:TYR:CD2	1:D:432:VAL:HG13	2.18	0.78
1:D:549:LEU:HA	1:D:552:LEU:HD12	1.66	0.78
1:D:616:GLU:HG3	1:D:724:GLY:O	1.82	0.78
2:E:193:TYR:CE2	2:E:322:LEU:HD22	2.19	0.78
2:E:199:LEU:HB2	2:F:77:HIS:NE2	1.98	0.78
1:A:1026:ARG:O	1:A:1030:ARG:HB2	1.83	0.78
1:A:879:GLN:HG2	1:A:1189:LEU:HG	1.63	0.78
2:B:99:PHE:HD2	2:C:129:ASP:CG	1.86	0.78
1:D:289:MET:C	1:D:289:MET:SD	2.62	0.78
1:D:879:GLN:OE1	1:D:1189:LEU:CD2	2.30	0.78
1:D:927:ARG:CG	1:D:928:GLY:H	1.92	0.78
2:F:78:PHE:HE1	2:F:103:GLY:HA2	1.49	0.78
2:F:354:PHE:HB2	2:F:372:LEU:HA	1.65	0.78
1:A:1226:LEU:HD22	1:A:1226:LEU:H	1.47	0.78
1:A:472:GLN:HG3	2:C:460:THR:HB	1.65	0.78
1:A:249:VAL:HG21	1:A:817:PRO:CG	2.13	0.78
2:C:375:HIS:O	2:C:378:LEU:HB2	1.84	0.78
1:D:659:HIS:ND1	1:D:719:LEU:HD22	1.99	0.78
2:E:115:TRP:O	2:E:116:THR:HG23	1.83	0.78
1:A:1231:LEU:HD11	1:A:1236:GLN:NE2	1.98	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:392:LEU:HG	1:A:395:TYR:CD1	2.18	0.78
1:A:601:LYS:HG2	1:A:721:ALA:CB	2.13	0.78
1:A:622:TYR:O	1:A:746:GLY:HA3	1.83	0.78
2:C:376:PRO:CG	2:C:482:ALA:HB1	2.13	0.78
2:C:424:SER:CB	2:C:427:GLN:HE22	1.93	0.78
2:C:472:LYS:HA	2:C:475:LEU:HD11	1.66	0.78
1:D:580:LEU:C	1:D:580:LEU:HD12	2.04	0.78
1:D:645:VAL:O	1:D:645:VAL:CG1	2.31	0.78
1:D:717:LEU:HD23	1:D:717:LEU:H	1.46	0.78
2:E:432:TYR:O	2:E:437:ILE:HD13	1.84	0.78
2:F:310:GLU:O	2:F:313:HIS:HB3	1.83	0.78
1:A:996:ARG:HH12	1:A:1003:TRP:CA	1.97	0.78
1:A:1041:TRP:C	1:A:1041:TRP:HD1	1.88	0.78
1:A:1131:ILE:CD1	1:A:1138:ARG:H	1.95	0.78
1:A:795:ASN:ND2	1:A:796:LYS:N	2.30	0.78
1:A:439:GLN:CG	1:A:836:LEU:H	1.96	0.78
1:A:933:SER:HB2	1:A:943:ARG:HD3	1.66	0.78
2:C:206:TYR:HD2	2:C:206:TYR:N	1.79	0.78
2:C:303:LEU:CD1	2:C:339:ASN:OD1	2.31	0.78
2:C:375:HIS:H	2:C:378:LEU:CD2	1.94	0.78
1:A:468:ASN:CG	2:C:460:THR:HG21	2.03	0.78
2:E:389:ARG:NE	2:E:390:GLY:N	2.29	0.78
1:D:194:ALA:HA	1:D:265:LEU:HB2	1.64	0.78
1:D:369:LEU:HB3	1:D:372:GLU:CG	2.14	0.78
1:D:613:HIS:CG	1:D:717:LEU:HD22	2.18	0.78
2:F:304:TRP:H	2:F:337:SER:CB	1.81	0.78
2:F:305:ASN:CA	2:F:335:VAL:HG11	2.14	0.78
2:F:375:HIS:O	2:F:378:LEU:HB2	1.84	0.78
2:F:385:LEU:HD11	2:F:402:LEU:HB2	1.66	0.78
2:F:435:MET:O	2:F:436:SER:CB	2.31	0.78
1:A:1131:ILE:CG2	1:A:1137:VAL:HA	2.14	0.78
1:A:271:VAL:HG12	1:A:272:SER:N	1.96	0.78
1:A:371:LYS:C	1:A:373:PRO:HD3	2.04	0.78
1:A:371:LYS:NZ	1:A:400:VAL:HG23	1.99	0.78
1:A:408:GLN:CA	1:A:411:LEU:HG	2.12	0.78
1:A:559:LEU:HB3	1:A:560:PRO:HD2	1.65	0.78
1:A:779:LEU:O	1:A:780:GLN:HG2	1.84	0.78
2:C:395:LEU:HD21	2:C:443:VAL:CB	2.12	0.78
1:D:1132:SER:HA	1:D:1136:GLU:O	1.83	0.78
1:D:1141:VAL:HG12	1:D:1142:ARG:O	1.84	0.78
1:D:660:CYS:HB2	1:D:745:PRO:HG3	1.65	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:275:ARG:HH11	1:D:843:GLN:CD	1.87	0.78
1:D:887:VAL:H	1:D:1146:ARG:HH22	1.31	0.78
1:D:76:ILE:HA	1:D:911:HIS:NE2	1.97	0.78
2:F:239:LEU:N	2:F:338:VAL:HG11	1.98	0.78
2:F:431:LYS:NZ	2:F:435:MET:HG2	1.98	0.78
2:F:82:SER:HA	2:F:86:LEU:CD2	2.10	0.78
1:A:482:ASP:HB2	1:A:483:PRO:CD	2.13	0.78
1:A:804:ALA:O	1:A:808:ILE:HG13	1.84	0.78
1:A:845:VAL:CB	1:A:855:VAL:HB	2.12	0.78
1:A:893:SER:HB2	1:A:896:LEU:CG	2.13	0.78
1:A:924:ARG:O	1:A:925:LYS:HG2	1.83	0.78
1:D:900:ALA:CA	1:D:915:ALA:HB1	2.13	0.77
1:D:922:GLN:HG2	1:D:931:LEU:CD1	2.13	0.77
2:F:182:ARG:NH2	2:F:213:VAL:O	2.15	0.77
1:A:1041:TRP:CD1	1:A:1041:TRP:O	2.37	0.77
1:A:241:PRO:HA	1:A:244:LEU:CB	2.14	0.77
1:D:206:GLY:HA2	1:D:245:ILE:HG22	1.65	0.77
1:D:371:LYS:NZ	1:D:396:CYS:HB2	1.99	0.77
1:D:414:PHE:HA	1:D:417:ARG:NH1	1.98	0.77
1:D:824:ALA:HA	1:D:827:ARG:CB	2.13	0.77
1:D:883:GLY:O	1:D:1143:GLU:HB2	1.84	0.77
2:E:94:GLY:HA2	2:E:96:HIS:CE1	2.20	0.77
1:A:758:ASN:CB	1:A:1058:PHE:HZ	1.95	0.77
1:A:1166:TYR:H	1:A:1166:TYR:HD1	1.31	0.77
1:A:238:GLN:HB2	1:A:239:LEU:HG	1.65	0.77
1:A:83:LEU:HD13	1:A:89:GLY:CA	2.15	0.77
2:E:418:LEU:CD1	2:F:122:ARG:HE	1.97	0.77
1:D:1001:GLY:HA2	1:D:1004:LEU:CD2	2.04	0.77
1:D:1134:HIS:O	1:D:1135:ASP:HB2	1.83	0.77
1:D:578:PRO:O	1:D:580:LEU:N	2.18	0.77
1:D:624:VAL:C	1:D:626:GLY:H	1.86	0.77
1:D:903:GLY:CA	1:D:1002:GLU:HB2	2.13	0.77
2:F:134:LYS:HB3	2:F:180:LYS:CD	2.13	0.77
1:A:196:VAL:HB	1:A:401:TRP:CZ3	2.18	0.77
1:A:213:VAL:HG13	1:A:397:ALA:HB1	1.67	0.77
1:A:813:VAL:CG2	1:A:814:VAL:N	2.47	0.77
2:B:231:ILE:CG1	2:C:135:PRO:HG3	2.14	0.77
1:D:1072:THR:N	1:D:1073:PRO:HD2	2.00	0.77
1:D:177:ARG:CG	1:D:178:TYR:H	1.97	0.77
1:D:247:LEU:O	1:D:248:GLU:HB2	1.85	0.77
1:D:348:LEU:O	1:D:348:LEU:HD13	1.84	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:775:GLU:HG2	1:D:780:GLN:CG	2.12	0.77
1:D:80:SER:O	1:D:82:GLY:N	2.17	0.77
2:E:254:PHE:O	2:E:258:HIS:N	2.17	0.77
2:F:114:TRP:HE1	2:F:259:ARG:NH1	1.82	0.77
2:F:233:GLU:O	2:F:234:LYS:HG2	1.85	0.77
2:F:243:THR:O	2:F:334:CYS:N	2.16	0.77
2:F:471:LEU:HA	2:F:474:PHE:CD2	2.19	0.77
2:B:323:HIS:HB2	2:B:332:VAL:CA	2.14	0.77
2:C:126:PHE:C	2:C:128:VAL:H	1.80	0.77
2:C:439:PHE:HD1	2:C:458:ARG:CB	1.98	0.77
2:C:367:LEU:O	2:C:369:ARG:HD2	1.83	0.77
1:D:866:ARG:CG	1:D:1069:ILE:HG21	2.08	0.77
1:D:214:ALA:N	1:D:401:TRP:HZ2	1.82	0.77
1:D:916:PHE:O	1:D:920:THR:CA	2.32	0.77
2:F:356:LEU:O	2:F:370:LYS:HE2	1.84	0.77
2:F:81:GLY:HA3	2:F:97:PRO:HD2	1.65	0.77
1:A:869:ARG:HA	1:A:1201:SER:N	1.98	0.77
1:A:175:TRP:CB	1:A:186:PRO:HD3	2.15	0.77
2:C:200:VAL:HG21	2:C:204:LEU:CD2	2.14	0.77
1:D:1166:TYR:CD1	1:D:1166:TYR:N	2.43	0.77
2:E:431:LYS:HA	2:E:434:GLU:HG3	1.64	0.77
1:A:308:GLN:NE2	1:A:1091:GLU:N	2.33	0.77
1:A:460:LYS:HD2	1:A:460:LYS:N	1.98	0.77
1:A:550:GLN:O	1:A:554:GLY:N	2.17	0.77
1:A:608:ASP:O	1:A:620:TRP:HH2	1.67	0.77
1:A:728:THR:HG22	1:A:729:GLN:N	1.99	0.77
1:A:805:HIS:ND1	1:A:806:LYS:N	2.32	0.77
1:D:504:LYS:HB3	1:D:523:ASP:HB2	1.66	0.77
2:C:81:GLY:CA	2:C:97:PRO:HB2	2.13	0.77
2:E:309:HIS:ND1	2:E:310:GLU:HG2	2.00	0.77
1:D:1221:TYR:HD1	1:D:1222:GLN:H	1.33	0.77
1:D:195:LEU:N	1:D:265:LEU:HD13	1.99	0.77
2:E:67:GLU:HG2	2:E:70:LEU:CD1	2.14	0.77
2:F:305:ASN:HA	2:F:335:VAL:HG11	1.65	0.77
1:A:79:LEU:CD2	1:A:1173:LEU:O	2.29	0.77
1:A:1221:TYR:O	1:A:1224:ILE:HG22	1.83	0.77
1:A:498:LYS:NZ	1:A:519:GLY:H	1.83	0.77
1:A:448:ALA:HA	1:A:870:VAL:HG13	1.65	0.77
1:A:962:ALA:HB2	1:A:986:TYR:CZ	2.19	0.77
2:B:324:GLY:CA	2:B:325:ARG:CZ	2.62	0.77
2:C:382:LYS:HD3	2:C:411:ILE:CG1	2.13	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:640:LEU:N	1:A:640:LEU:HD23	1.98	0.77
1:D:869:ARG:O	1:D:1200:PRO:CB	2.32	0.77
1:D:302:SER:HB2	1:D:351:SER:CB	2.15	0.77
2:E:78:PHE:O	2:E:79:LEU:CG	2.27	0.77
2:F:435:MET:CE	2:F:435:MET:HA	2.14	0.77
2:F:435:MET:O	2:F:436:SER:HB3	1.84	0.77
1:A:77:GLN:CB	1:A:1174:PRO:HB3	2.14	0.77
1:A:1175:GLN:HG3	1:A:1176:SER:N	2.00	0.77
1:A:358:LEU:O	1:A:361:VAL:HG13	1.85	0.77
1:A:460:LYS:O	1:A:464:MET:HG3	1.84	0.77
1:A:711:ALA:O	1:A:712:VAL:HG23	1.83	0.77
2:B:233:GLU:C	2:B:234:LYS:HE2	2.03	0.77
2:B:233:GLU:OE2	2:C:133:HIS:N	2.15	0.77
2:B:478:TYR:O	2:B:482:ALA:HB2	1.84	0.77
1:D:346:ASP:H	1:D:1040:LYS:HD3	1.48	0.77
1:D:294:THR:CG2	1:D:407:PHE:HE2	1.93	0.77
1:D:885:THR:C	1:D:886:LEU:HD23	2.05	0.77
2:E:69:LEU:H	2:E:69:LEU:HD12	1.48	0.77
1:A:1040:LYS:CG	1:A:1044:VAL:HG11	2.15	0.77
1:A:1210:TYR:CE1	1:A:1213:PRO:CG	2.61	0.77
2:B:311:LEU:CG	2:B:312:LEU:N	2.47	0.77
1:D:512:LYS:O	1:D:513:LEU:HD23	1.84	0.77
1:D:1008:LEU:C	1:D:1011:PRO:HD2	2.05	0.77
1:D:1143:GLU:O	1:D:1146:ARG:HG3	1.85	0.77
1:D:797:MET:CB	1:D:869:ARG:HD3	2.14	0.77
1:D:610:PHE:CE1	1:D:972:ARG:NH2	2.52	0.77
2:E:375:HIS:CD2	2:E:376:PRO:HB3	2.20	0.77
2:E:373:LYS:HB2	2:E:459:ASP:OD1	1.84	0.77
2:F:342:LEU:HG	2:F:344:ARG:HB2	1.67	0.77
1:A:885:THR:HB	1:A:1146:ARG:NE	1.99	0.77
1:A:608:ASP:CB	1:A:779:LEU:HB2	2.14	0.77
1:A:827:ARG:HH11	1:A:827:ARG:HG2	1.50	0.77
2:C:79:LEU:HA	2:C:99:PHE:HA	1.67	0.77
1:D:824:ALA:HA	1:D:827:ARG:HB3	1.67	0.76
2:F:383:VAL:HG23	2:F:439:PHE:O	1.83	0.76
1:A:562:ARG:HB3	1:A:565:HIS:CE1	2.21	0.76
1:A:584:ALA:C	1:A:585:TRP:HE3	1.87	0.76
1:A:593:SER:CA	1:A:596:MET:HG3	2.12	0.76
1:A:624:VAL:C	1:A:626:GLY:H	1.88	0.76
2:B:112:ALA:O	2:B:115:TRP:HB3	1.85	0.76
2:C:208:LEU:HD12	2:C:208:LEU:O	1.85	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:215:PHE:CD2	2:C:233:GLU:HG2	2.21	0.76
1:A:1037:GLN:C	1:A:1039:LYS:H	1.84	0.76
1:D:167:PRO:HD2	1:D:168:ALA:N	1.99	0.76
1:D:441:TRP:NE1	1:D:879:GLN:CA	2.48	0.76
2:F:302:THR:C	2:F:303:LEU:HD12	2.06	0.76
1:A:192:GLU:O	1:A:193:ARG:HG3	1.84	0.76
1:A:213:VAL:O	1:A:401:TRP:HZ2	1.68	0.76
1:A:462:SER:C	1:A:466:LEU:HD23	2.06	0.76
1:A:617:ARG:HG2	1:A:765:PRO:HD3	1.66	0.76
1:A:601:LYS:HA	1:A:717:LEU:HB3	1.65	0.76
1:A:732:TYR:HE1	1:A:755:LYS:CB	1.98	0.76
2:C:300:ILE:HG22	2:C:301:GLU:H	1.49	0.76
2:C:393:LEU:O	2:C:396:ARG:HG3	1.86	0.76
2:E:389:ARG:N	2:E:395:LEU:HD11	2.00	0.76
1:D:289:MET:SD	1:D:290:ARG:N	2.58	0.76
1:D:605:LEU:CD2	1:D:781:ALA:HB1	2.14	0.76
1:D:611:PRO:HB2	1:D:612:LEU:CD1	2.15	0.76
1:D:928:GLY:HA2	1:D:931:LEU:HD12	1.65	0.76
2:F:301:GLU:HB3	2:F:339:ASN:CB	2.12	0.76
1:A:1119:LEU:HA	1:A:1122:GLU:CG	2.15	0.76
1:A:441:TRP:NE1	1:A:877:MET:HB3	2.00	0.76
1:A:989:THR:HG22	1:A:999:ASP:HB2	1.65	0.76
2:B:69:LEU:O	2:B:72:ILE:HG12	1.83	0.76
2:C:201:ASN:C	2:C:203:ARG:H	1.88	0.76
2:C:301:GLU:HB3	2:C:339:ASN:CG	2.06	0.76
2:C:328:ARG:O	2:C:329:LYS:HD2	1.86	0.76
2:C:81:GLY:HA3	2:C:97:PRO:CB	2.15	0.76
1:A:643:ALA:O	1:A:645:VAL:HG23	1.85	0.76
1:D:1158:LEU:H	1:D:1158:LEU:HD23	1.50	0.76
1:D:842:PRO:HB2	1:D:856:GLU:CD	2.05	0.76
2:F:394:GLU:N	2:F:394:GLU:CD	2.39	0.76
1:A:1041:TRP:O	1:A:1041:TRP:HD1	1.68	0.76
2:B:323:HIS:HB2	2:B:332:VAL:N	2.00	0.76
2:C:241:TRP:N	2:C:336:LEU:HB3	2.00	0.76
2:C:389:ARG:CG	2:C:390:GLY:H	1.98	0.76
1:A:152:GLU:HA	1:A:155:ASN:ND2	2.00	0.76
2:C:258:HIS:HA	2:C:261:GLN:OE1	1.85	0.76
2:E:194:VAL:HG13	2:E:195:ASN:N	2.01	0.76
1:D:137:GLN:O	1:D:141:LEU:HD12	1.86	0.76
1:D:649:TYR:HD2	1:D:652:ILE:HG21	1.50	0.76
1:D:823:ARG:O	1:D:826:ILE:HG12	1.85	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:556:THR:HB	2:F:467:HIS:NE2	1.99	0.76
1:A:887:VAL:HG23	1:A:1146:ARG:HH11	1.50	0.76
1:A:885:THR:CB	1:A:1146:ARG:NE	2.48	0.76
1:A:1179:PHE:CD1	1:A:1220:ILE:HG21	2.20	0.76
1:A:244:LEU:O	1:A:247:LEU:HB3	1.86	0.76
1:A:591:LEU:HD12	1:A:594:LEU:HG	1.67	0.76
1:A:868:ASP:HB3	1:A:1199:THR:HB	1.68	0.76
2:B:74:GLN:O	2:B:77:HIS:CA	2.33	0.76
2:C:182:ARG:NH2	2:C:213:VAL:O	2.18	0.76
2:C:416:GLY:O	2:C:419:GLU:HB3	1.85	0.76
1:D:522:GLY:O	1:D:524:PRO:CD	2.33	0.76
1:D:513:LEU:O	1:D:570:PRO:HD2	1.86	0.76
1:D:437:VAL:HA	1:D:441:TRP:HD1	1.50	0.76
1:D:659:HIS:CD2	1:D:719:LEU:HB2	2.21	0.76
2:F:208:LEU:HD13	2:F:240:VAL:HB	1.64	0.76
1:A:582:ASP:HB3	1:A:592:LEU:HD22	1.68	0.76
1:A:600:PRO:O	1:A:602:LEU:O	2.04	0.76
1:A:79:LEU:HD12	1:A:89:GLY:O	1.85	0.76
2:C:241:TRP:HB3	2:C:336:LEU:HB2	1.67	0.76
1:D:1207:GLU:HA	1:D:1210:TYR:CD2	2.21	0.76
1:D:130:LEU:O	1:D:131:TYR:CD2	2.39	0.76
1:D:140:ARG:O	1:D:144:GLN:HB2	1.85	0.76
1:D:375:GLU:HG3	1:D:379:LYS:CG	2.16	0.76
1:D:456:GLN:C	1:D:460:LYS:HZ3	1.88	0.76
1:D:547:ALA:CA	1:D:551:LYS:HE2	2.16	0.76
1:D:606:THR:CG2	1:D:612:LEU:O	2.34	0.76
1:D:83:LEU:HG	1:D:89:GLY:CA	2.16	0.76
2:F:201:ASN:C	2:F:203:ARG:H	1.89	0.76
2:F:347:LEU:CA	2:F:350:LEU:HD11	2.15	0.76
2:F:350:LEU:H	2:F:350:LEU:HD12	1.51	0.76
2:F:449:GLU:O	2:F:450:ASN:C	2.24	0.76
1:A:610:PHE:N	1:A:610:PHE:HD2	1.83	0.76
2:C:239:LEU:N	2:C:338:VAL:CG1	2.46	0.76
2:C:352:ASP:N	2:C:355:GLN:HB2	2.00	0.76
2:C:466:MET:HB3	2:C:471:LEU:CD2	2.10	0.76
1:D:1102:GLN:O	1:D:1106:VAL:HG23	1.85	0.76
1:D:1165:ALA:C	1:D:1166:TYR:HD1	1.89	0.76
1:D:275:ARG:HD2	1:D:843:GLN:OE1	1.85	0.76
1:D:604:ALA:CA	1:D:613:HIS:CE1	2.67	0.76
1:D:630:ASN:HB2	1:D:744:ILE:O	1.86	0.76
1:D:645:VAL:CG1	1:D:650:ARG:HG2	2.15	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:717:LEU:CD2	1:D:717:LEU:H	1.92	0.76
1:D:748:TRP:CH2	1:D:750:PHE:HE2	2.02	0.76
1:D:914:THR:CG2	1:D:999:ASP:HB3	2.15	0.76
2:E:213:VAL:HG13	2:E:235:THR:HG22	1.67	0.76
2:F:197:LEU:O	2:F:202:LYS:HA	1.85	0.76
2:F:444:THR:O	2:F:447:THR:HG22	1.86	0.76
1:A:498:LYS:HE3	1:A:518:ALA:N	2.01	0.76
2:C:444:THR:O	2:C:447:THR:N	2.18	0.76
2:C:452:LEU:HD12	2:C:453:ILE:N	2.01	0.76
1:D:192:GLU:O	1:D:217:PRO:HD3	1.86	0.76
1:D:549:LEU:CA	1:D:551:LYS:HZ2	1.99	0.76
1:D:656:TYR:O	1:D:659:HIS:CE1	2.38	0.76
2:F:208:LEU:HD11	2:F:240:VAL:CB	2.16	0.76
2:F:241:TRP:HB2	2:F:336:LEU:CG	2.14	0.76
2:F:374:LEU:HD12	2:F:379:ALA:HA	1.65	0.76
1:A:1112:MET:HA	1:A:1156:THR:CG2	2.16	0.76
1:A:175:TRP:HB3	1:A:186:PRO:CD	2.15	0.76
1:A:274:ASP:N	1:A:276:ALA:H	1.83	0.76
1:A:618:HIS:HB2	1:A:725:PRO:CG	2.16	0.76
1:A:77:GLN:OE1	1:A:1174:PRO:CB	2.34	0.76
2:B:260:LEU:O	2:B:264:ARG:HD3	1.86	0.76
2:B:440:THR:HG22	2:B:441:VAL:N	2.01	0.76
2:C:244:PRO:CD	2:C:247:THR:CG2	2.64	0.76
2:C:300:ILE:HG22	2:C:301:GLU:N	2.00	0.76
2:C:303:LEU:HD11	2:C:339:ASN:OD1	1.86	0.76
2:C:385:LEU:HD11	2:C:402:LEU:CB	2.14	0.76
2:C:474:PHE:HE1	2:C:478:TYR:CD2	2.04	0.76
2:C:484:ASN:O	2:C:485:VAL:HG13	1.86	0.76
1:D:966:LEU:HD23	1:D:970:ASN:ND2	2.00	0.76
1:D:903:GLY:HA3	1:D:1002:GLU:CB	2.16	0.76
2:E:336:LEU:HD22	2:E:338:VAL:HG23	1.67	0.76
2:F:393:LEU:O	2:F:396:ARG:HG3	1.85	0.76
1:A:354:ASN:OD1	1:A:355:SER:N	2.18	0.76
1:A:471:CYS:CB	1:A:472:GLN:OE1	2.34	0.76
1:A:667:GLN:O	1:A:671:GLN:HG2	1.86	0.76
2:B:417:TYR:HH	2:C:122:ARG:HD3	1.48	0.76
1:A:120:LEU:H	1:A:120:LEU:HD22	1.50	0.76
1:D:907:PHE:CZ	1:D:1002:GLU:HB3	2.20	0.75
1:D:1027:LYS:HZ3	1:D:1091:GLU:HG2	1.48	0.75
1:D:1165:ALA:O	1:D:1169:GLY:HA3	1.87	0.75
1:D:203:LEU:C	1:D:203:LEU:HD23	2.07	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:463:LEU:HA	1:D:466:LEU:HD22	1.68	0.75
1:D:865:ALA:HA	1:D:872:SER:HA	1.68	0.75
1:D:993:ARG:C	1:D:996:ARG:HD2	2.05	0.75
2:E:102:LEU:HD13	2:E:102:LEU:O	1.87	0.75
2:E:383:VAL:O	2:E:384:ALA:HB2	1.85	0.75
2:F:200:VAL:CG1	2:F:204:LEU:HB2	2.15	0.75
1:A:1069:ILE:CG2	1:A:1070:PRO:HD3	2.15	0.75
1:A:1169:GLY:O	1:A:1170:LEU:HB3	1.86	0.75
1:A:913:CYS:C	1:A:918:TRP:H	1.89	0.75
2:C:254:PHE:CA	2:C:257:ARG:HH21	1.98	0.75
1:D:671:GLN:NE2	1:D:671:GLN:HA	2.00	0.75
1:D:948:ILE:HG23	1:D:1071:ARG:HB2	1.69	0.75
1:D:1145:ASP:HA	1:D:1148:ARG:CD	2.16	0.75
1:D:352:SER:O	1:D:358:GLU:HG2	1.85	0.75
1:D:389:PHE:CA	1:D:393:MET:HG3	2.16	0.75
2:E:336:LEU:HD23	2:E:337:SER:N	2.00	0.75
2:E:371:VAL:HA	2:E:433:ASP:CG	2.05	0.75
1:A:112:LEU:HD21	1:A:918:TRP:HA	1.69	0.75
2:C:415:PRO:HB3	2:C:417:TYR:CE2	2.21	0.75
1:A:560:PRO:HD2	2:C:452:LEU:HD23	1.66	0.75
2:C:382:LYS:O	2:C:438:LEU:HB2	1.86	0.75
1:D:432:VAL:HB	1:D:1128:ARG:HH21	1.51	0.75
1:D:744:ILE:HG12	1:D:745:PRO:HD2	1.68	0.75
1:D:823:ARG:NE	1:D:824:ALA:H	1.85	0.75
1:D:83:LEU:N	1:D:83:LEU:HD13	2.01	0.75
2:F:484:ASN:O	2:F:485:VAL:HG13	1.87	0.75
1:A:142:LEU:HD22	1:A:1118:TRP:CD1	2.21	0.75
1:A:374:ARG:NE	1:A:392:LEU:HD13	1.98	0.75
1:A:601:LYS:HG2	1:A:721:ALA:HB1	1.68	0.75
1:A:914:THR:O	1:A:915:ALA:CB	2.34	0.75
2:B:201:ASN:HB2	2:C:421:MET:HE1	1.68	0.75
2:B:269:SER:H	2:B:270:PRO:CD	1.98	0.75
1:D:1006:ARG:CZ	1:D:1010:LEU:HD11	2.15	0.75
1:D:1028:VAL:HG22	1:D:1096:ARG:CZ	2.16	0.75
1:D:113:TRP:HE3	1:D:114:GLY:N	1.81	0.75
1:D:239:LEU:CB	1:D:242:ALA:HB3	2.15	0.75
1:D:459:MET:N	1:D:460:LYS:NZ	2.34	0.75
1:D:956:GLY:HA2	1:D:1094:THR:CG2	2.04	0.75
2:E:351:TYR:C	2:E:353:SER:H	1.89	0.75
2:F:251:TRP:CE3	2:F:251:TRP:HA	2.20	0.75
1:A:1026:ARG:CB	1:A:1029:GLN:CB	2.49	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1151:LEU:HG	1:A:1223:ILE:HD11	1.68	0.75
1:A:879:GLN:CG	1:A:1189:LEU:HD21	2.15	0.75
1:A:1207:GLU:HA	1:A:1210:TYR:HD1	1.51	0.75
1:A:1231:LEU:HG	1:A:1232:GLU:H	1.51	0.75
2:B:90:SER:HA	2:B:93:SER:OG	1.85	0.75
2:B:99:PHE:CD2	2:C:129:ASP:CG	2.60	0.75
2:C:425:LEU:HD21	2:C:429:TYR:OH	1.86	0.75
2:C:372:LEU:HD21	2:C:435:MET:C	2.07	0.75
2:F:243:THR:O	2:F:334:CYS:HB2	1.87	0.75
2:F:454:HIS:HD2	2:F:468:ILE:CA	1.99	0.75
2:F:73:CYS:HB3	2:F:79:LEU:HD23	1.69	0.75
1:A:1041:TRP:CE2	1:A:1043:VAL:HB	2.21	0.75
1:A:594:LEU:HD23	1:A:594:LEU:N	2.01	0.75
1:A:961:PHE:N	1:A:961:PHE:CD1	2.51	0.75
2:C:128:VAL:HG12	2:C:192:HIS:CD2	2.21	0.75
2:C:239:LEU:O	2:C:338:VAL:CG1	2.34	0.75
1:A:940:GLY:O	1:A:941:ILE:HD12	1.86	0.75
1:D:1236:GLN:HB2	1:D:1237:PRO:HD3	1.68	0.75
1:D:134:ASN:HB3	1:D:137:GLN:H	1.51	0.75
1:D:259:ARG:HG3	1:D:261:TRP:CD1	2.22	0.75
1:D:434:TYR:O	1:D:435:LEU:HG	1.86	0.75
1:D:608:ASP:HB3	1:D:620:TRP:CE2	2.22	0.75
1:D:865:ALA:HB3	1:D:1197:CYS:SG	2.26	0.75
2:E:253:ASP:O	2:E:257:ARG:N	2.20	0.75
2:F:385:LEU:HD11	2:F:402:LEU:CB	2.17	0.75
2:B:306:LEU:HD22	2:B:310:GLU:HG2	1.67	0.75
1:D:1029:GLN:NE2	1:D:1030:ARG:NH2	2.32	0.75
1:D:1131:ILE:HD13	1:D:1131:ILE:H	1.51	0.75
1:D:1154:GLN:OE1	1:D:1220:ILE:HA	1.85	0.75
1:D:249:VAL:HG23	1:D:250:PRO:HD2	1.66	0.75
1:D:279:ARG:HG3	1:D:841:LEU:O	1.87	0.75
1:D:213:VAL:HG13	1:D:401:TRP:CE2	2.21	0.75
1:D:881:PRO:O	1:D:884:TYR:HB2	1.86	0.75
2:F:114:TRP:HE1	2:F:259:ARG:NH2	1.85	0.75
2:F:131:LEU:HB3	2:F:133:HIS:CG	2.21	0.75
2:F:239:LEU:H	2:F:338:VAL:CG2	1.99	0.75
1:A:1175:GLN:HG3	1:A:1176:SER:H	1.49	0.75
1:A:412:PRO:O	1:A:415:LEU:HB2	1.86	0.75
1:A:652:ILE:HA	1:A:655:LEU:CD2	2.16	0.75
1:A:823:ARG:O	1:A:823:ARG:HG3	1.87	0.75
1:D:356:LEU:CD1	1:D:369:LEU:HD22	2.16	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:417:ARG:O	1:D:418:CYS:SG	2.45	0.75
1:D:247:LEU:HA	1:D:815:TRP:CD1	2.22	0.75
2:F:277:ASP:OD2	2:F:287:ASN:HB3	1.87	0.75
2:F:446:THR:CA	2:F:450:ASN:H	1.99	0.75
1:A:879:GLN:HG2	1:A:1189:LEU:CG	2.17	0.75
1:A:172:ALA:HA	1:A:221:TYR:O	1.85	0.75
1:A:256:PRO:HB2	1:A:281:GLN:HA	1.67	0.75
1:A:473:LEU:HD23	1:A:473:LEU:H	1.50	0.75
1:A:922:GLN:HG3	1:A:923:GLY:N	2.01	0.75
1:A:926:SER:O	1:A:927:ARG:HB2	1.86	0.75
2:B:104:VAL:HG21	2:C:128:VAL:O	1.87	0.75
1:D:584:ALA:C	1:D:585:TRP:HE3	1.90	0.75
1:A:539:PHE:O	1:A:540:GLN:CG	2.35	0.75
1:D:249:VAL:HG22	1:D:250:PRO:CD	2.16	0.75
1:D:642:SER:CA	1:D:645:VAL:HG11	2.13	0.75
2:E:78:PHE:CA	2:E:102:LEU:HB3	2.17	0.75
2:E:353:SER:OG	2:E:372:LEU:HD11	1.86	0.75
2:E:422:GLN:C	2:E:424:SER:H	1.89	0.75
2:F:203:ARG:HE	2:F:204:LEU:HD13	1.48	0.75
2:F:254:PHE:CA	2:F:257:ARG:HH21	2.00	0.75
1:A:1011:PRO:HG2	1:A:1012:VAL:HG13	1.68	0.75
1:A:1119:LEU:HA	1:A:1122:GLU:HG2	1.69	0.75
1:A:870:VAL:HG22	1:A:1202:ASN:HD22	1.52	0.75
1:A:266:VAL:HG23	1:A:290:ARG:O	1.86	0.75
1:A:610:PHE:N	1:A:610:PHE:CD2	2.52	0.75
1:A:825:VAL:O	1:A:829:PRO:HD3	1.85	0.75
2:C:344:ARG:N	2:C:347:LEU:HD13	2.01	0.75
1:A:576:LEU:HD21	2:C:482:ALA:HB2	1.67	0.75
1:D:623:LEU:O	1:D:625:PRO:CD	2.34	0.75
1:D:665:LYS:O	1:D:669:MET:HG2	1.87	0.75
1:D:242:ALA:HA	1:D:245:ILE:HG23	1.68	0.74
1:D:352:SER:C	1:D:358:GLU:CD	2.37	0.74
1:D:542:ASP:O	1:D:546:ARG:N	2.18	0.74
1:D:275:ARG:NH1	1:D:843:GLN:CG	2.50	0.74
1:D:798:ILE:HG22	1:D:869:ARG:CZ	2.17	0.74
2:E:185:LEU:HD13	2:E:238:SER:OG	1.87	0.74
2:F:395:LEU:HB3	2:F:443:VAL:CG2	2.15	0.74
2:F:471:LEU:HA	2:F:474:PHE:HB3	1.69	0.74
2:F:402:LEU:HG	2:F:472:LYS:HZ3	1.52	0.74
1:A:481:GLU:HB3	1:A:482:ASP:OD1	1.86	0.74
1:A:608:ASP:C	1:A:620:TRP:HH2	1.90	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:821:LEU:HG	1:A:826:ILE:HG23	1.69	0.74
1:A:897:TRP:NE1	1:A:901:VAL:HG13	2.02	0.74
1:A:933:SER:HB2	1:A:943:ARG:CZ	2.17	0.74
1:A:933:SER:HB2	1:A:943:ARG:CD	2.15	0.74
1:A:958:GLY:O	1:A:961:PHE:CD1	2.39	0.74
1:A:958:GLY:O	1:A:961:PHE:HD1	1.69	0.74
2:C:386:ASP:CB	2:C:442:LEU:HA	2.10	0.74
1:D:1025:LEU:HD22	1:D:1025:LEU:O	1.87	0.74
1:D:1045:ALA:HB1	1:D:1090:GLU:HG2	1.69	0.74
1:D:131:TYR:CB	1:D:138:HIS:CE1	2.70	0.74
1:D:176:THR:N	1:D:185:VAL:HG11	2.01	0.74
1:D:865:ALA:HB1	1:D:872:SER:OG	1.86	0.74
1:D:904:ASP:OD2	1:D:915:ALA:HB3	1.86	0.74
2:E:215:PHE:HB3	2:E:233:GLU:HA	1.70	0.74
2:F:447:THR:HA	2:F:452:LEU:CA	2.17	0.74
2:F:445:GLU:HA	2:F:448:LEU:HD12	1.69	0.74
2:F:446:THR:HG22	2:F:449:GLU:OE1	1.87	0.74
1:A:1015:THR:HA	1:A:1016:GLU:OE2	1.86	0.74
1:A:1148:ARG:HG2	1:A:1148:ARG:NH1	1.90	0.74
1:A:76:ILE:HG13	1:A:77:GLN:CG	2.15	0.74
1:A:879:GLN:CB	1:A:886:LEU:HD11	2.16	0.74
2:B:306:LEU:C	2:B:308:ASP:N	2.37	0.74
2:B:344:ARG:CD	2:B:344:ARG:H	1.99	0.74
2:C:342:LEU:HB2	2:C:344:ARG:CB	2.17	0.74
1:D:885:THR:CB	1:D:1146:ARG:HD2	2.15	0.74
1:D:660:CYS:HB2	1:D:745:PRO:HD3	1.68	0.74
1:D:622:TYR:O	1:D:746:GLY:CA	2.36	0.74
2:E:380:PRO:O	2:E:382:LYS:CB	2.34	0.74
2:F:241:TRP:CA	2:F:336:LEU:HB2	2.18	0.74
2:F:351:TYR:O	2:F:352:ASP:C	2.24	0.74
1:A:261:TRP:HA	1:A:261:TRP:CE3	2.21	0.74
1:A:665:LYS:CG	1:A:666:GLN:N	2.50	0.74
1:A:717:LEU:O	1:A:721:ALA:HB3	1.88	0.74
1:A:813:VAL:HB	1:A:840:ILE:HG12	1.70	0.74
1:A:897:TRP:CZ3	1:A:1108:TYR:HE1	2.05	0.74
2:B:74:GLN:CG	2:B:75:ARG:N	2.50	0.74
2:C:273:PHE:CE1	2:C:291:TYR:HD1	2.04	0.74
1:D:872:SER:OG	1:D:1200:PRO:HD3	1.86	0.74
1:D:348:LEU:CD1	1:D:350:ILE:HD11	2.17	0.74
1:D:900:ALA:HB1	1:D:915:ALA:CB	2.13	0.74
2:F:389:ARG:HG3	2:F:390:GLY:H	1.52	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:622:TYR:HB2	1:A:747:CYS:O	1.87	0.74
2:C:276:SER:O	2:C:287:ASN:CB	2.36	0.74
2:C:343:ASP:O	2:C:347:LEU:HD13	1.87	0.74
1:D:1031:GLU:O	1:D:1031:GLU:HG3	1.87	0.74
1:D:1202:ASN:O	1:D:1203:PRO:C	2.25	0.74
1:D:220:TRP:CD1	1:D:221:TYR:HD2	2.05	0.74
1:D:269:HIS:HB3	1:D:277:HIS:CD2	2.22	0.74
1:D:622:TYR:O	1:D:746:GLY:HA3	1.87	0.74
2:F:245:PRO:O	2:F:248:SER:CB	2.35	0.74
1:A:282:TYR:CZ	1:A:432:VAL:HG22	2.21	0.74
1:A:451:THR:C	1:A:869:ARG:NH2	2.40	0.74
1:A:468:ASN:HD21	1:A:578:PRO:CG	2.00	0.74
1:A:490:TRP:CZ2	1:A:596:MET:HB3	2.22	0.74
1:A:601:LYS:HZ3	1:A:616:GLU:HG2	1.52	0.74
1:A:720:THR:CG2	1:A:721:ALA:N	2.51	0.74
1:A:932:HIS:HB3	1:A:947:LYS:HG2	1.68	0.74
1:A:958:GLY:HA2	1:A:986:TYR:CE2	2.22	0.74
2:B:284:ARG:O	2:B:285:LYS:HB3	1.87	0.74
2:C:194:VAL:O	2:C:197:LEU:HD22	1.86	0.74
2:C:207:GLY:HA3	2:C:239:LEU:HD11	1.69	0.74
1:D:513:LEU:H	1:D:570:PRO:CD	1.99	0.74
1:D:1014:ARG:CD	1:D:1026:ARG:NE	2.50	0.74
1:D:308:GLN:CD	1:D:1027:LYS:HZ1	1.91	0.74
1:D:610:PHE:HE1	1:D:972:ARG:NH2	1.86	0.74
1:D:744:ILE:HB	1:D:745:PRO:HD3	1.70	0.74
2:F:299:LEU:N	2:F:299:LEU:CD2	2.50	0.74
1:A:247:LEU:HD12	1:A:251:THR:CG2	2.17	0.74
1:A:448:ALA:O	1:A:870:VAL:HG11	1.87	0.74
2:B:350:LEU:O	2:B:350:LEU:HD13	1.88	0.74
2:C:387:VAL:O	2:C:420:THR:OG1	2.06	0.74
1:D:569:HIS:H	1:D:570:PRO:CD	2.01	0.74
1:D:1069:ILE:HG12	1:D:1070:PRO:HD2	1.68	0.74
1:D:80:SER:HB2	1:D:125:LEU:HG	1.69	0.74
1:D:220:TRP:CD1	1:D:221:TYR:CD2	2.76	0.74
1:D:312:TRP:CZ2	1:D:347:TRP:HA	2.21	0.74
1:D:551:LYS:N	1:D:551:LYS:CD	2.30	0.74
1:D:807:ARG:HB2	1:D:858:THR:HG23	1.69	0.74
1:D:931:LEU:O	1:D:934:LYS:HD2	1.88	0.74
1:D:972:ARG:O	1:D:974:THR:HG23	1.88	0.74
2:E:128:VAL:CG1	2:E:210:GLN:HE21	1.98	0.74
2:E:432:TYR:HD2	2:E:437:ILE:HG12	1.53	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:442:LEU:HD12	2:E:454:HIS:CE1	2.23	0.74
2:F:455:LEU:CB	2:F:456:ARG:HE	2.00	0.74
2:F:476:ILE:HD13	2:F:477:LYS:N	2.02	0.74
1:A:1026:ARG:N	1:A:1091:GLU:OE2	2.21	0.74
1:A:435:LEU:HD23	1:A:879:GLN:OE1	1.86	0.74
2:B:236:GLU:HG2	2:B:341:ASP:HB2	1.69	0.74
2:C:403:PHE:CD1	2:C:417:TYR:OH	2.41	0.74
1:D:879:GLN:O	1:D:1189:LEU:HD21	1.87	0.74
1:D:209:PRO:O	1:D:224:CYS:SG	2.45	0.74
1:D:189:ILE:HG23	1:D:258:GLN:HB3	1.67	0.74
2:E:373:LYS:HA	2:E:458:ARG:CZ	2.17	0.74
2:F:243:THR:OG1	2:F:334:CYS:HB2	1.87	0.74
2:F:344:ARG:HA	2:F:347:LEU:HD22	1.68	0.74
1:A:297:MET:HG3	1:A:298:HIS:N	2.03	0.74
1:A:452:TYR:N	1:A:869:ARG:HH21	1.86	0.74
1:A:497:GLN:HA	1:A:516:GLU:CB	2.18	0.74
1:A:606:THR:OG1	1:A:613:HIS:N	2.21	0.74
1:A:624:VAL:HG11	1:A:747:CYS:SG	2.26	0.74
2:B:311:LEU:CD1	2:B:312:LEU:H	2.00	0.74
1:D:1005:VAL:O	1:D:1009:ASN:N	2.21	0.74
1:D:897:TRP:CZ3	1:D:1177:VAL:HG11	2.22	0.74
1:D:1185:ILE:HD12	1:D:1185:ILE:O	1.87	0.74
1:D:1236:GLN:CB	1:D:1237:PRO:CD	2.66	0.74
1:D:189:ILE:HG21	1:D:258:GLN:HB3	1.70	0.74
1:D:270:ASN:CG	1:D:294:THR:HG21	2.08	0.74
1:D:614:TYR:HE2	1:D:620:TRP:CE3	2.05	0.74
1:D:842:PRO:HB2	1:D:856:GLU:OE2	1.88	0.74
2:F:193:TYR:HB2	2:F:242:PHE:CE2	2.22	0.74
2:F:389:ARG:CG	2:F:390:GLY:H	2.00	0.74
2:F:464:GLU:OE2	2:F:466:MET:HG2	1.87	0.74
1:A:903:GLY:HA3	1:A:1002:GLU:HB3	1.70	0.74
1:A:1014:ARG:HB2	1:A:1026:ARG:NH1	2.03	0.74
1:A:145:LYS:HG3	1:A:1118:TRP:HH2	1.51	0.74
1:A:252:GLY:O	1:A:280:GLU:HB3	1.88	0.74
1:A:261:TRP:HE3	1:A:261:TRP:HA	1.52	0.74
1:A:196:VAL:O	1:A:401:TRP:HH2	1.70	0.74
1:A:612:LEU:CB	1:A:620:TRP:HB2	2.17	0.74
1:A:765:PRO:HG2	1:A:766:PHE:CD1	2.23	0.74
1:A:873:GLU:OE1	1:A:875:LYS:CG	2.36	0.74
2:C:356:LEU:CB	2:C:359:ASN:HD21	2.00	0.74
1:D:101:ARG:HG3	1:D:101:ARG:NH1	1.91	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:195:LEU:O	1:D:267:VAL:HA	1.88	0.74
1:D:452:TYR:CG	1:D:801:TRP:NE1	2.55	0.74
1:D:478:ARG:NE	1:D:481:GLU:OE1	2.21	0.74
1:D:572:TRP:CZ2	2:F:470:LYS:HB3	2.23	0.74
1:D:913:CYS:O	1:D:917:GLY:HA3	1.88	0.74
2:F:387:VAL:HG23	2:F:417:TYR:HA	1.68	0.74
1:A:348:LEU:O	1:A:349:ASP:C	2.26	0.74
1:A:392:LEU:CD2	1:A:395:TYR:HB2	2.18	0.74
1:A:85:GLU:OE2	1:A:127:LEU:O	2.05	0.74
2:B:110:LEU:HD21	2:B:342:LEU:HD21	1.70	0.74
2:B:342:LEU:O	2:B:345:GLY:N	2.21	0.74
1:A:1209:ARG:NH2	2:C:256:LEU:HD23	2.03	0.74
2:B:279:GLN:O	2:B:280:ASP:CB	2.30	0.74
1:D:360:HIS:O	1:D:364:VAL:HG22	1.88	0.73
1:D:577:CYS:HB3	1:D:578:PRO:HD2	1.70	0.73
2:F:344:ARG:N	2:F:347:LEU:HD13	2.03	0.73
1:A:1236:GLN:CB	1:A:1237:PRO:CD	2.64	0.73
1:A:257:THR:HG23	1:A:258:GLN:O	1.87	0.73
1:A:581:ASP:C	1:A:583:PRO:HD2	2.08	0.73
2:C:197:LEU:HD23	2:C:198:ASP:N	2.03	0.73
2:C:394:GLU:CD	2:C:394:GLU:H	1.91	0.73
1:D:554:GLY:HA2	1:D:557:GLU:HG3	1.68	0.73
1:D:1072:THR:HG21	1:D:1134:HIS:CE1	2.23	0.73
1:D:1142:ARG:O	1:D:1144:GLU:O	2.06	0.73
1:D:556:THR:OG1	2:F:452:LEU:HD22	1.88	0.73
1:D:926:SER:HB2	1:D:930:ASP:CB	2.17	0.73
2:E:428:LEU:HD13	2:E:428:LEU:N	2.02	0.73
2:F:432:TYR:HB3	2:F:437:ILE:HD11	1.70	0.73
1:A:1205:GLY:O	1:A:1208:ARG:HB3	1.88	0.73
1:A:593:SER:HA	1:A:596:MET:CG	2.15	0.73
1:A:669:MET:C	1:A:673:ALA:HB2	2.09	0.73
1:A:815:TRP:CA	1:A:838:GLY:HA3	2.17	0.73
1:A:87:ILE:HG22	1:A:88:PHE:H	1.48	0.73
2:C:290:TYR:HA	2:C:298:GLU:O	1.88	0.73
2:C:323:HIS:O	2:C:333:PRO:HD3	1.88	0.73
1:D:1005:VAL:HA	1:D:1008:LEU:HD12	1.68	0.73
2:E:93:SER:C	2:E:95:CYS:H	1.90	0.73
2:F:115:TRP:CH2	2:F:210:GLN:HG2	2.22	0.73
2:F:431:LYS:HE2	2:F:435:MET:HB2	1.70	0.73
1:A:1024:ASP:HA	1:A:1026:ARG:NH1	2.03	0.73
1:A:887:VAL:HG22	1:A:1146:ARG:HH11	1.51	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:845:VAL:HB	1:A:855:VAL:CG2	2.19	0.73
2:B:306:LEU:C	2:B:308:ASP:H	1.91	0.73
2:C:192:HIS:O	2:C:196:CYS:SG	2.46	0.73
2:C:200:VAL:HG11	2:C:204:LEU:HD13	1.70	0.73
2:C:301:GLU:HG2	2:C:339:ASN:HB3	1.70	0.73
1:A:503:VAL:HB	1:A:521:PRO:CA	2.16	0.73
1:D:888:GLY:HA3	1:D:1138:ARG:HA	1.70	0.73
1:D:271:VAL:O	1:D:274:ASP:HB2	1.88	0.73
1:D:293:ASP:CG	1:D:296:SER:HB2	2.08	0.73
1:D:805:HIS:O	1:D:809:SER:N	2.20	0.73
1:D:992:LEU:HD12	1:D:993:ARG:H	1.52	0.73
2:F:247:THR:CB	2:F:251:TRP:CD1	2.67	0.73
2:F:353:SER:HB3	2:F:373:LYS:O	1.88	0.73
1:A:1200:PRO:O	1:A:1204:THR:OG1	2.06	0.73
1:A:205:GLU:HA	1:A:205:GLU:OE1	1.87	0.73
1:A:904:ASP:C	1:A:910:MET:HA	2.08	0.73
1:A:933:SER:HB2	1:A:943:ARG:NE	2.03	0.73
2:B:123:GLU:HB2	2:C:403:PHE:HZ	1.53	0.73
2:B:396:ARG:HD2	2:B:418:LEU:H	1.52	0.73
2:C:300:ILE:HA	2:C:342:LEU:HD12	1.69	0.73
1:A:739:TYR:HA	1:A:742:VAL:HB	1.71	0.73
1:D:1069:ILE:HG23	1:D:1070:PRO:N	2.02	0.73
1:D:86:GLN:NE2	1:D:132:GLY:N	2.34	0.73
1:D:179:GLY:CA	1:D:182:GLY:CA	2.65	0.73
1:D:369:LEU:O	1:D:371:LYS:O	2.05	0.73
1:D:598:VAL:CA	1:D:601:LYS:HE2	2.16	0.73
1:D:658:LYS:CE	1:D:713:PRO:HB2	2.19	0.73
1:D:891:VAL:O	1:D:894:GLN:HG2	1.88	0.73
1:D:996:ARG:N	1:D:996:ARG:NE	2.36	0.73
2:E:117:SER:HA	2:E:121:PHE:CD1	2.24	0.73
2:E:383:VAL:O	2:E:384:ALA:HB3	1.87	0.73
1:A:897:TRP:HZ3	1:A:1108:TYR:CE1	2.07	0.73
1:A:1160:THR:HG22	1:A:1164:PHE:HE1	1.51	0.73
1:A:297:MET:HG3	1:A:298:HIS:H	1.52	0.73
1:A:498:LYS:HZ2	1:A:519:GLY:H	1.35	0.73
1:A:500:ALA:HB1	1:A:520:ALA:HB3	1.70	0.73
1:D:429:GLU:HG3	1:D:1128:ARG:CZ	2.18	0.73
1:D:474:LEU:HD12	1:D:474:LEU:N	2.02	0.73
1:D:546:ARG:HH21	2:F:404:ASN:CB	1.95	0.73
1:D:546:ARG:O	1:D:551:LYS:NZ	2.19	0.73
1:D:618:HIS:NE2	1:D:761:ASN:HB3	2.02	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:761:ASN:O	1:D:763:GLY:N	2.21	0.73
1:D:989:THR:CA	1:D:998:SER:OG	2.34	0.73
1:D:993:ARG:CA	1:D:996:ARG:HD2	2.19	0.73
2:E:102:LEU:HD13	2:E:102:LEU:C	2.09	0.73
2:F:202:LYS:O	2:F:325:ARG:HG2	1.88	0.73
1:A:145:LYS:O	1:A:149:PRO:HD2	1.88	0.73
1:A:166:PRO:HG3	1:A:398:GLN:CD	2.08	0.73
1:A:278:ILE:HG12	1:A:291:PHE:CE1	2.22	0.73
1:A:386:ARG:HD3	1:A:387:GLU:H	1.53	0.73
1:A:712:VAL:O	1:A:719:LEU:HD22	1.88	0.73
1:A:868:ASP:O	1:A:869:ARG:HB2	1.88	0.73
1:A:875:LYS:C	1:A:878:VAL:HG23	2.09	0.73
1:A:87:ILE:CG2	1:A:88:PHE:H	2.01	0.73
2:B:128:VAL:HB	2:B:208:LEU:HB2	1.70	0.73
2:C:267:ALA:HB3	2:C:270:PRO:HB3	1.70	0.73
2:C:363:ARG:HH11	2:C:363:ARG:CA	1.99	0.73
1:D:410:GLN:O	1:D:412:PRO:HD2	1.88	0.73
1:A:1109:LEU:O	1:A:1112:MET:HG3	1.89	0.73
1:A:273:PHE:C	1:A:276:ALA:H	1.92	0.73
1:A:600:PRO:C	1:A:602:LEU:O	2.27	0.73
2:B:75:ARG:HD2	2:B:434:GLU:OE2	1.87	0.73
2:C:347:LEU:CA	2:C:350:LEU:HD11	2.18	0.73
1:A:634:LEU:HD12	1:A:638:THR:CB	2.19	0.73
1:D:495:PHE:HB2	1:D:573:TYR:CE1	2.23	0.73
1:D:645:VAL:O	1:D:650:ARG:HB3	1.88	0.73
2:F:452:LEU:O	2:F:453:ILE:HG13	1.88	0.73
1:A:1029:GLN:C	1:A:1031:GLU:O	2.26	0.73
1:A:1131:ILE:HG21	1:A:1137:VAL:CA	2.19	0.73
1:A:259:ARG:NE	1:A:259:ARG:HA	2.04	0.73
1:A:439:GLN:CG	1:A:835:GLY:HA3	2.17	0.73
2:C:347:LEU:HA	2:C:350:LEU:HD11	1.69	0.73
2:C:356:LEU:O	2:C:370:LYS:HE2	1.88	0.73
1:D:515:ILE:CD1	1:D:568:GLY:O	2.37	0.73
2:E:218:VAL:H	2:E:231:ILE:HA	1.53	0.73
2:E:272:ASN:HA	2:E:292:ASN:ND2	2.04	0.73
1:D:308:GLN:OE1	1:D:1027:LYS:NZ	2.21	0.73
1:D:1138:ARG:NH2	1:D:1191:LYS:CE	2.51	0.73
1:D:779:LEU:C	1:D:779:LEU:HD22	2.09	0.73
1:D:948:ILE:CG2	1:D:1071:ARG:HB2	2.19	0.73
1:A:1118:TRP:O	1:A:1122:GLU:HG2	1.89	0.73
1:A:219:ALA:O	1:A:220:TRP:CE3	2.41	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:79:LEU:CD1	1:A:89:GLY:O	2.37	0.73
2:B:99:PHE:CD2	2:C:129:ASP:OD2	2.42	0.73
1:A:1208:ARG:NH2	2:C:277:ASP:CG	2.42	0.73
2:C:398:VAL:HA	2:C:401:GLY:HA3	1.70	0.73
2:C:463:LYS:O	2:C:465:MET:HE2	1.88	0.73
1:D:1195:MET:HB3	1:D:1196:ASP:O	1.89	0.73
1:D:257:THR:OG1	1:D:281:GLN:NE2	2.21	0.73
1:D:278:ILE:O	1:D:282:TYR:HD1	1.72	0.73
1:D:432:VAL:CG1	1:D:1128:ARG:HE	1.99	0.73
1:D:81:ARG:HG2	1:D:125:LEU:O	1.88	0.73
2:E:377:CYS:SG	2:E:378:LEU:CD1	2.77	0.73
2:F:473:ASP:CA	2:F:476:ILE:HG23	2.17	0.73
1:A:241:PRO:HA	1:A:244:LEU:CG	2.18	0.73
1:A:436:PRO:CB	1:A:881:PRO:HD3	2.08	0.73
1:A:930:ASP:HA	1:A:933:SER:HB3	1.70	0.73
2:B:304:TRP:CE3	2:B:304:TRP:HA	2.24	0.73
2:B:400:GLN:NE2	2:B:417:TYR:CG	2.55	0.73
2:C:342:LEU:HB2	2:C:344:ARG:HD3	1.70	0.73
1:D:1151:LEU:HD23	1:D:1223:ILE:HD11	1.71	0.72
1:D:547:ALA:O	1:D:551:LYS:HG2	1.89	0.72
1:D:918:TRP:C	1:D:920:THR:N	2.41	0.72
2:E:431:LYS:C	2:E:433:ASP:N	2.38	0.72
2:E:76:ARG:HG3	2:E:435:MET:SD	2.29	0.72
2:F:342:LEU:HD22	2:F:342:LEU:O	1.88	0.72
1:A:1165:ALA:O	1:A:1170:LEU:N	2.22	0.72
1:A:223:TRP:H	1:A:223:TRP:HD1	1.32	0.72
1:A:821:LEU:HD12	1:A:825:VAL:HB	1.69	0.72
1:D:1113:LEU:N	1:D:1113:LEU:HD23	2.04	0.72
1:D:897:TRP:HH2	1:D:1173:LEU:HD13	1.50	0.72
2:F:403:PHE:HD1	2:F:417:TYR:HH	1.32	0.72
2:F:416:GLY:O	2:F:419:GLU:HB3	1.88	0.72
2:F:424:SER:HB3	2:F:427:GLN:NE2	2.03	0.72
1:A:449:GLN:HA	1:A:452:TYR:CE1	2.24	0.72
1:A:580:LEU:HD12	1:A:581:ASP:N	2.04	0.72
1:A:616:GLU:HG2	1:A:724:GLY:HA3	1.71	0.72
1:A:615:SER:HB3	1:A:725:PRO:HB3	1.69	0.72
2:B:324:GLY:N	2:B:333:PRO:HD3	2.03	0.72
2:B:99:PHE:HE2	2:C:131:LEU:HD11	1.54	0.72
1:A:1209:ARG:HH22	2:C:256:LEU:HD23	1.54	0.72
1:A:784:GLY:N	2:C:363:ARG:HD3	2.02	0.72
1:D:239:LEU:HB3	1:D:242:ALA:HB3	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:242:ALA:CA	1:D:245:ILE:HG12	2.17	0.72
1:D:453:GLU:O	1:D:457:ARG:HG2	1.90	0.72
1:D:735:GLY:CA	1:D:749:PHE:HB3	2.18	0.72
2:E:115:TRP:HZ3	2:E:210:GLN:HA	1.54	0.72
2:E:122:ARG:HD2	2:E:122:ARG:H	1.54	0.72
2:E:213:VAL:HG13	2:E:235:THR:CG2	2.18	0.72
2:F:120:VAL:HG23	2:F:121:PHE:N	2.03	0.72
1:D:553:LYS:HD3	2:F:468:ILE:HG22	1.70	0.72
1:A:1026:ARG:O	1:A:1030:ARG:CB	2.37	0.72
1:A:247:LEU:HD12	1:A:251:THR:HG21	1.71	0.72
1:A:913:CYS:N	1:A:917:GLY:HA3	2.03	0.72
2:B:114:TRP:CA	2:B:262:TRP:CZ2	2.71	0.72
1:D:670:PRO:HB2	1:D:672:GLU:O	1.89	0.72
1:A:1193:VAL:HG23	1:A:1194:THR:N	2.03	0.72
1:D:115:GLN:HE22	1:D:924:ARG:NH1	1.86	0.72
1:D:633:LYS:HB2	1:D:638:THR:HG21	1.70	0.72
2:F:77:HIS:O	2:F:100:GLY:HA3	1.89	0.72
2:F:342:LEU:O	2:F:342:LEU:CD2	2.37	0.72
2:F:402:LEU:HG	2:F:472:LYS:NZ	2.03	0.72
1:A:1131:ILE:HD11	1:A:1133:ILE:CG2	2.18	0.72
1:A:652:ILE:C	1:A:655:LEU:HG	2.09	0.72
1:A:73:PRO:O	1:A:74:LEU:HD13	1.90	0.72
1:A:87:ILE:CG2	1:A:88:PHE:N	2.51	0.72
2:B:304:TRP:HA	2:B:304:TRP:HE3	1.54	0.72
2:C:325:ARG:HG3	2:C:325:ARG:NH1	1.99	0.72
2:F:186:LEU:HD23	2:F:187:HIS:N	2.03	0.72
1:D:1000:GLU:O	1:D:1004:LEU:HD13	1.88	0.72
1:D:1005:VAL:CA	1:D:1008:LEU:HD13	2.15	0.72
1:D:429:GLU:HG2	1:D:1117:LYS:HE2	1.71	0.72
1:D:1126:ASP:OD2	1:D:1145:ASP:HB3	1.89	0.72
1:D:283:LEU:CD2	1:D:839:ALA:HB3	2.18	0.72
1:D:269:HIS:NE2	1:D:291:PHE:HD2	1.88	0.72
1:D:597:ARG:O	1:D:599:THR:N	2.22	0.72
1:D:600:PRO:HG2	1:D:616:GLU:HB3	1.72	0.72
1:D:657:ARG:O	1:D:660:CYS:HB3	1.89	0.72
1:D:997:LEU:CD2	1:D:998:SER:H	2.02	0.72
2:F:403:PHE:CD1	2:F:417:TYR:OH	2.43	0.72
2:F:485:VAL:OXT	2:F:485:VAL:CG2	2.38	0.72
1:A:1002:GLU:HG3	1:A:1003:TRP:CE3	2.24	0.72
1:A:1068:ASP:C	1:A:1070:PRO:HD3	2.10	0.72
1:A:87:ILE:HG21	1:A:1166:TYR:OH	1.89	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1179:PHE:CB	1:A:1220:ILE:HG12	2.19	0.72
1:A:451:THR:HG21	1:A:1202:ASN:HB3	1.72	0.72
1:A:80:SER:OG	1:A:125:LEU:HD11	1.89	0.72
1:A:175:TRP:HB2	1:A:184:ALA:C	2.10	0.72
1:A:175:TRP:N	1:A:186:PRO:HB3	2.03	0.72
1:A:615:SER:HB2	1:A:725:PRO:CB	2.19	0.72
1:A:753:PRO:HD2	1:A:770:PHE:CG	2.23	0.72
1:A:873:GLU:HB3	1:A:875:LYS:HG3	1.70	0.72
1:A:566:LEU:HB3	1:A:567:PRO:CD	2.19	0.72
2:E:307:GLY:HA2	2:E:334:CYS:HB3	1.71	0.72
1:D:443:ARG:HG2	1:D:443:ARG:HH11	1.55	0.72
1:D:907:PHE:HZ	1:D:1002:GLU:HB3	1.53	0.72
1:D:943:ARG:O	1:D:946:ALA:HB3	1.90	0.72
1:D:1209:ARG:NH2	2:F:256:LEU:HD22	2.05	0.72
1:A:1020:ILE:CD1	1:A:1021:SER:H	2.02	0.72
2:B:78:PHE:HE1	2:B:103:GLY:HA2	1.52	0.72
2:C:247:THR:CB	2:C:251:TRP:CD1	2.71	0.72
2:C:356:LEU:CB	2:C:359:ASN:ND2	2.39	0.72
2:B:199:LEU:HD23	2:C:77:HIS:HD2	1.53	0.72
1:D:513:LEU:O	1:D:568:GLY:CA	2.38	0.72
1:D:1113:LEU:HD23	1:D:1113:LEU:H	1.54	0.72
1:D:389:PHE:CD1	1:D:390:GLN:N	2.57	0.72
1:D:452:TYR:CB	1:D:801:TRP:NE1	2.48	0.72
1:D:599:THR:HB	1:D:600:PRO:HD3	1.72	0.72
1:D:656:TYR:HB3	1:D:744:ILE:HD12	1.70	0.72
2:E:114:TRP:CH2	2:E:238:SER:O	2.42	0.72
2:E:448:LEU:C	2:E:448:LEU:HD23	2.09	0.72
1:A:439:GLN:HG3	1:A:836:LEU:N	2.04	0.72
1:A:618:HIS:HA	1:A:762:VAL:HG21	1.72	0.72
1:A:621:GLY:N	1:A:748:TRP:HB3	2.03	0.72
1:A:786:ALA:O	1:A:789:PRO:HD2	1.90	0.72
2:C:303:LEU:N	2:C:303:LEU:HD12	2.05	0.72
2:C:355:GLN:O	2:C:357:THR:N	2.22	0.72
2:C:375:HIS:N	2:C:378:LEU:HD22	2.00	0.72
2:C:394:GLU:CD	2:C:394:GLU:N	2.43	0.72
2:C:485:VAL:CG2	2:C:485:VAL:OXT	2.38	0.72
2:B:469:SER:C	2:B:471:LEU:H	1.91	0.72
1:D:1116:MET:SD	1:D:1153:LEU:HD22	2.30	0.72
1:D:177:ARG:NE	1:D:216:SER:H	1.87	0.72
1:D:265:LEU:HD21	1:D:267:VAL:HA	1.72	0.72
1:D:421:PRO:O	1:D:425:ALA:HB3	1.90	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:620:TRP:NE1	1:D:751:LYS:HB2	2.04	0.72
1:D:795:ASN:HA	1:D:798:ILE:CD1	2.20	0.72
1:D:967:MET:HA	1:D:967:MET:HE3	1.72	0.72
2:F:114:TRP:NE1	2:F:259:ARG:NH1	2.36	0.72
2:F:209:ALA:CB	2:F:238:SER:O	2.38	0.72
2:F:305:ASN:HA	2:F:335:VAL:CG1	2.19	0.72
2:F:357:THR:O	2:F:358:GLU:HG2	1.90	0.72
2:F:423:SER:HB2	2:F:427:GLN:OE1	1.90	0.72
1:A:879:GLN:HG2	1:A:1189:LEU:HD21	1.72	0.72
1:A:228:LEU:HD11	1:A:386:ARG:HB3	1.70	0.72
1:A:286:GLY:N	1:A:1128:ARG:NH2	2.37	0.72
1:A:436:PRO:CG	1:A:879:GLN:HB3	2.19	0.72
1:A:658:LYS:CG	1:A:713:PRO:HB2	2.17	0.72
1:A:613:HIS:HB3	1:A:717:LEU:CD2	2.19	0.72
1:A:913:CYS:O	1:A:918:TRP:N	2.20	0.72
2:B:299:LEU:HD21	2:B:302:THR:HG23	1.72	0.72
2:B:67:GLU:HB2	2:B:88:ARG:NH1	2.03	0.72
2:C:243:THR:O	2:C:334:CYS:N	2.22	0.72
1:A:503:VAL:HG11	1:A:521:PRO:N	2.04	0.72
1:A:634:LEU:HB3	1:A:635:PRO:CA	2.19	0.72
1:A:739:TYR:C	1:A:741:ASP:H	1.92	0.72
1:A:539:PHE:O	1:A:540:GLN:HG2	1.88	0.72
2:C:272:ASN:HB3	2:C:292:ASN:HB2	1.72	0.72
1:D:1014:ARG:CD	1:D:1026:ARG:HE	2.02	0.72
1:D:368:PRO:CA	1:D:369:LEU:HD23	2.19	0.72
1:D:550:GLN:CB	1:D:551:LYS:HE3	2.19	0.72
1:D:726:LYS:O	1:D:727:ASP:CG	2.28	0.72
1:D:959:GLN:CB	1:D:960:PRO:CD	2.67	0.72
2:E:382:LYS:CG	2:E:383:VAL:N	2.47	0.72
2:F:360:SER:HB3	2:F:364:LYS:HD2	1.72	0.72
2:E:203:ARG:NH1	2:F:418:LEU:HD22	1.99	0.72
1:A:1113:LEU:HD23	1:A:1113:LEU:N	2.05	0.72
1:A:410:GLN:O	1:A:414:PHE:CD2	2.43	0.72
1:A:618:HIS:CE1	1:A:761:ASN:HD22	2.08	0.72
1:A:918:TRP:O	1:A:920:THR:HG22	1.90	0.72
2:C:454:HIS:HB3	2:C:466:MET:O	1.89	0.72
1:D:623:LEU:HB3	1:D:667:GLN:CD	2.10	0.72
2:B:468:ILE:O	2:B:468:ILE:HG23	1.89	0.72
1:D:1006:ARG:NH2	1:D:1010:LEU:HD11	2.05	0.72
1:D:371:LYS:HG3	1:D:396:CYS:CA	2.19	0.72
1:D:719:LEU:HG	1:D:720:THR:H	1.55	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:79:LEU:HA	2:E:100:GLY:H	1.55	0.72
2:E:255:TRP:CE3	2:E:255:TRP:HA	2.23	0.72
2:E:423:SER:O	2:E:424:SER:HB3	1.89	0.72
2:F:342:LEU:HB2	2:F:344:ARG:CD	2.20	0.72
1:A:955:TYR:CD2	1:A:1098:ASN:OD1	2.42	0.72
1:A:887:VAL:CG2	1:A:1146:ARG:NH1	2.51	0.72
1:A:1170:LEU:HD13	1:A:1170:LEU:O	1.87	0.72
1:A:402:ALA:C	1:A:406:VAL:HG13	2.10	0.72
1:A:866:ARG:HD3	1:A:866:ARG:O	1.89	0.72
1:A:893:SER:HB2	1:A:896:LEU:HD23	1.71	0.72
2:B:131:LEU:C	2:B:182:ARG:HH21	1.94	0.72
2:B:403:PHE:CE1	2:B:415:PRO:HD3	2.24	0.72
2:B:371:VAL:CA	2:B:433:ASP:HB3	2.04	0.72
2:C:186:LEU:HD23	2:C:187:HIS:N	2.05	0.72
2:C:374:LEU:CD1	2:C:379:ALA:HA	2.19	0.72
2:E:391:PRO:HG2	2:E:395:LEU:H	1.54	0.72
2:C:180:LYS:O	2:C:181:LEU:HD23	1.88	0.72
1:A:1056:GLU:HA	1:A:1059:ASN:HD21	1.52	0.72
1:D:1015:THR:HG23	1:D:1026:ARG:HB2	1.70	0.71
1:D:884:TYR:OH	1:D:1142:ARG:NH1	2.23	0.71
1:D:865:ALA:CB	1:D:1197:CYS:SG	2.78	0.71
1:D:744:ILE:HG12	1:D:745:PRO:CD	2.20	0.71
1:D:275:ARG:CZ	1:D:843:GLN:H	2.01	0.71
1:D:960:PRO:HA	1:D:963:GLU:CG	2.19	0.71
1:D:976:GLN:O	1:D:980:GLU:HG3	1.89	0.71
2:F:127:PRO:CD	2:F:128:VAL:N	2.48	0.71
2:F:185:LEU:HD12	2:F:236:GLU:HB2	1.72	0.71
2:F:402:LEU:O	2:F:406:LEU:HD13	1.89	0.71
1:A:1187:ARG:NH2	1:A:1212:ILE:O	2.23	0.71
1:A:452:TYR:N	1:A:869:ARG:NH2	2.37	0.71
1:A:955:TYR:CE2	1:A:1102:GLN:NE2	2.58	0.71
1:A:955:TYR:HD2	1:A:1098:ASN:OD1	1.72	0.71
2:B:234:LYS:O	2:B:236:GLU:HG3	1.89	0.71
2:C:344:ARG:HA	2:C:347:LEU:CG	2.20	0.71
2:C:372:LEU:HD21	2:C:436:SER:HB2	1.72	0.71
2:C:383:VAL:HG23	2:C:439:PHE:O	1.89	0.71
2:C:430:SER:O	2:C:434:GLU:CG	2.37	0.71
2:C:92:LEU:HD21	2:C:344:ARG:HE	1.55	0.71
1:D:145:LYS:O	1:D:149:PRO:CD	2.37	0.71
1:D:281:GLN:HE21	1:D:287:SER:CB	2.03	0.71
1:D:845:VAL:HB	1:D:855:VAL:CG2	2.20	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:865:ALA:HA	1:D:872:SER:CA	2.20	0.71
2:E:336:LEU:HD23	2:E:337:SER:H	1.54	0.71
2:E:350:LEU:HD23	2:E:350:LEU:O	1.90	0.71
2:E:431:LYS:O	2:E:433:ASP:N	2.23	0.71
2:F:458:ARG:NE	2:F:459:ASP:O	2.20	0.71
1:A:515:ILE:HG22	1:A:516:GLU:N	2.04	0.71
1:A:73:PRO:C	1:A:74:LEU:HD22	2.09	0.71
2:C:78:PHE:C	2:C:100:GLY:H	1.93	0.71
2:C:476:ILE:HD13	2:C:477:LYS:N	2.03	0.71
2:B:475:LEU:O	2:B:479:ILE:N	2.19	0.71
1:A:564:GLN:N	1:A:564:GLN:HE21	1.87	0.71
2:B:277:ASP:CG	2:B:287:ASN:HD21	1.93	0.71
1:D:1027:LYS:HB2	1:D:1096:ARG:HG3	1.70	0.71
1:D:626:GLY:O	1:D:627:ARG:HB3	1.89	0.71
1:D:279:ARG:HH21	1:D:844:VAL:CG2	2.02	0.71
2:F:354:PHE:CZ	2:F:370:LYS:HD3	2.25	0.71
1:A:1025:LEU:O	1:A:1026:ARG:HG3	1.89	0.71
1:A:129:PRO:O	1:A:131:TYR:CZ	2.44	0.71
1:A:157:LEU:HD11	1:A:195:LEU:HA	1.70	0.71
1:A:389:PHE:CD1	1:A:390:GLN:N	2.58	0.71
2:B:199:LEU:HD13	2:B:200:VAL:HG13	1.70	0.71
2:B:417:TYR:O	2:B:419:GLU:N	2.22	0.71
2:B:213:VAL:HG11	2:C:132:HIS:CE1	2.25	0.71
2:B:122:ARG:HB2	2:B:124:GLN:HE21	1.55	0.71
1:D:1108:TYR:HA	1:D:1164:PHE:CE1	2.25	0.71
1:D:260:ASP:OD2	1:D:261:TRP:CE2	2.44	0.71
1:D:367:PRO:CD	1:D:368:PRO:CD	2.54	0.71
1:D:457:ARG:CB	1:D:457:ARG:HH11	2.04	0.71
1:D:714:GLY:O	1:D:716:PRO:HD3	1.89	0.71
2:F:115:TRP:HH2	2:F:210:GLN:HG2	1.53	0.71
1:A:1040:LYS:HB3	1:A:1044:VAL:HG11	1.72	0.71
1:A:444:TYR:CE1	1:A:1206:MET:SD	2.83	0.71
1:A:612:LEU:CD2	1:A:620:TRP:HA	2.20	0.71
1:A:753:PRO:HB2	1:A:770:PHE:CZ	2.24	0.71
1:A:897:TRP:HE1	1:A:901:VAL:HG13	1.54	0.71
2:B:78:PHE:O	2:B:79:LEU:HB2	1.89	0.71
1:D:1008:LEU:O	1:D:1012:VAL:HG22	1.89	0.71
1:D:415:LEU:C	1:D:417:ARG:H	1.93	0.71
1:D:474:LEU:HD12	1:D:474:LEU:H	1.55	0.71
2:F:254:PHE:HE2	2:F:255:TRP:CE2	2.09	0.71
2:F:354:PHE:HE1	2:F:370:LYS:HB3	1.55	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1141:VAL:HG12	1:A:1142:ARG:O	1.91	0.71
1:A:348:LEU:HD13	1:A:350:ILE:HD13	1.72	0.71
1:A:476:GLY:CA	1:A:711:ALA:HB1	2.19	0.71
2:B:308:ASP:HB3	2:B:309:HIS:CE1	2.25	0.71
2:C:396:ARG:HH11	2:C:397:GLN:CA	2.02	0.71
2:C:317:GLY:HA3	2:C:320:SER:CB	2.19	0.71
1:D:1186:ASP:OD1	1:D:1187:ARG:N	2.23	0.71
1:D:739:TYR:CG	1:D:740:ASN:N	2.59	0.71
2:E:441:VAL:HG12	2:E:442:LEU:N	2.06	0.71
2:F:131:LEU:C	2:F:182:ARG:HD3	2.10	0.71
2:F:267:ALA:HB3	2:F:270:PRO:HB3	1.72	0.71
2:F:67:GLU:O	2:F:71:GLU:HB3	1.90	0.71
1:A:1019:TRP:HD1	1:A:1020:ILE:HG22	1.55	0.71
1:A:605:LEU:HA	1:A:781:ALA:CB	2.20	0.71
1:A:623:LEU:HD22	1:A:667:GLN:OE1	1.90	0.71
1:A:80:SER:OG	1:A:125:LEU:HD21	1.90	0.71
2:C:242:PHE:HA	2:C:334:CYS:O	1.91	0.71
2:C:468:ILE:HG13	2:C:472:LYS:HE3	1.72	0.71
2:C:317:GLY:HA2	2:C:320:SER:HB2	1.73	0.71
1:D:1206:MET:CE	1:D:1210:TYR:HA	2.20	0.71
1:D:275:ARG:CZ	1:D:843:GLN:N	2.53	0.71
1:D:553:LYS:HD3	2:F:468:ILE:CG2	2.20	0.71
1:D:827:ARG:HD3	1:D:828:HIS:N	2.06	0.71
1:D:885:THR:HG22	1:D:1186:ASP:O	1.91	0.71
2:F:190:LEU:CB	2:F:314:MET:HE1	2.20	0.71
1:A:297:MET:SD	1:A:411:LEU:CD2	2.77	0.71
1:A:496:LYS:O	1:A:516:GLU:HB2	1.90	0.71
1:A:576:LEU:HD12	1:A:577:CYS:HB2	1.71	0.71
1:A:606:THR:HG23	1:A:613:HIS:N	2.05	0.71
2:C:388:GLY:CA	2:C:444:THR:HG22	2.20	0.71
2:C:396:ARG:C	2:C:396:ARG:HD2	2.10	0.71
2:B:231:ILE:HD11	2:C:135:PRO:HG3	1.72	0.71
1:D:1197:CYS:C	1:D:1204:THR:OG1	2.29	0.71
1:D:305:SER:HB2	1:D:308:GLN:OE1	1.89	0.71
1:D:858:THR:HB	1:D:859:TRP:CD1	2.25	0.71
2:E:254:PHE:CE1	2:E:258:HIS:HB2	2.26	0.71
2:E:422:GLN:HG2	2:E:424:SER:N	2.03	0.71
2:E:86:LEU:HA	2:E:90:SER:CB	2.20	0.71
2:F:209:ALA:CB	2:F:239:LEU:HA	2.21	0.71
2:F:456:ARG:CG	2:F:457:SER:H	1.94	0.71
2:F:485:VAL:HG22	2:F:485:VAL:OXT	1.90	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1068:ASP:N	1:A:1071:ARG:CG	2.50	0.71
1:A:1153:LEU:HD13	1:A:1157:ASN:HD21	1.55	0.71
1:A:348:LEU:O	1:A:350:ILE:N	2.23	0.71
1:A:458:GLU:O	1:A:461:LYS:HG2	1.90	0.71
1:A:482:ASP:OD2	1:A:483:PRO:HD2	1.89	0.71
2:B:114:TRP:HE1	2:B:259:ARG:NE	1.88	0.71
2:C:342:LEU:HB2	2:C:344:ARG:CD	2.19	0.71
2:B:211:ILE:HG12	2:B:237:ALA:CB	2.18	0.71
1:D:1023:GLN:HE22	1:D:1099:TRP:C	1.93	0.71
1:D:1131:ILE:HD11	1:D:1138:ARG:CB	2.19	0.71
1:D:255:SER:OG	1:D:258:GLN:N	2.21	0.71
1:D:277:HIS:HB3	1:D:291:PHE:CE2	2.26	0.71
1:D:345:TRP:HA	1:D:1040:LYS:HZ2	1.53	0.71
1:D:633:LYS:HB3	1:D:638:THR:HG21	1.72	0.71
1:D:798:ILE:CG2	1:D:869:ARG:HH22	1.97	0.71
2:E:199:LEU:HG	2:E:200:VAL:N	2.04	0.71
2:E:387:VAL:O	2:E:442:LEU:HD22	1.91	0.71
2:E:456:ARG:HG3	2:E:462:MET:C	2.10	0.71
2:F:306:LEU:HD21	2:F:310:GLU:CD	2.10	0.71
1:A:853:ARG:HH22	1:A:1105:ALA:CB	2.04	0.71
1:A:1200:PRO:HB2	1:A:1202:ASN:O	1.89	0.71
1:A:174:GLY:HA2	1:A:220:TRP:N	2.04	0.71
1:A:577:CYS:CB	1:A:578:PRO:HD3	2.21	0.71
1:A:468:ASN:HD21	1:A:578:PRO:HG3	1.56	0.71
1:A:719:LEU:HD23	1:A:719:LEU:N	2.04	0.71
2:B:248:SER:O	2:B:252:LEU:HB2	1.91	0.71
2:C:125:VAL:CG2	2:C:208:LEU:HA	2.09	0.71
2:C:354:PHE:CZ	2:C:370:LYS:HD3	2.26	0.71
1:D:1165:ALA:HB3	1:D:1166:TYR:HE1	1.56	0.71
1:D:193:ARG:CA	1:D:193:ARG:NH1	2.54	0.71
1:D:420:HIS:NE2	1:D:1020:ILE:CG2	2.53	0.71
1:D:479:TYR:CD1	1:D:480:LYS:HG3	2.26	0.71
1:D:547:ALA:C	1:D:551:LYS:HE2	2.11	0.71
1:D:656:TYR:HD1	1:D:744:ILE:HD12	1.53	0.71
1:D:78:MET:CB	1:D:1175:GLN:HB3	2.19	0.71
1:D:815:TRP:HA	1:D:838:GLY:CA	2.20	0.71
2:E:353:SER:OG	2:E:374:LEU:HG	1.88	0.71
2:E:382:LYS:CG	2:E:412:SER:HB2	2.16	0.71
2:E:377:CYS:HB3	2:E:483:LYS:HE3	1.72	0.71
2:F:421:MET:HG3	2:F:422:GLN:H	1.56	0.71
1:A:1066:THR:OG1	1:A:1073:PRO:CD	2.39	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1141:VAL:HG11	1:A:1146:ARG:HA	1.73	0.71
1:A:1124:ALA:HB1	1:A:1148:ARG:HG2	1.71	0.71
1:A:188:ALA:O	1:A:189:ILE:CG2	2.37	0.71
1:A:298:HIS:ND1	1:A:299:MET:N	2.38	0.71
1:A:436:PRO:HB2	1:A:879:GLN:CA	2.15	0.71
1:A:444:TYR:OH	1:A:1200:PRO:HG3	1.90	0.71
2:C:120:VAL:HG23	2:C:121:PHE:N	2.06	0.71
2:C:240:VAL:HA	2:C:336:LEU:HB3	1.73	0.71
2:C:247:THR:CB	2:C:251:TRP:HD1	2.03	0.71
2:C:241:TRP:CA	2:C:336:LEU:HB2	2.21	0.71
1:D:510:ALA:O	1:D:512:LYS:N	2.24	0.71
1:D:765:PRO:HG2	1:D:766:PHE:CD1	2.26	0.71
1:D:550:GLN:N	1:D:551:LYS:CE	2.54	0.70
1:D:79:LEU:HB3	1:D:83:LEU:HD11	1.73	0.70
1:D:922:GLN:CG	1:D:931:LEU:HD13	2.18	0.70
2:E:79:LEU:O	2:E:100:GLY:HA3	1.91	0.70
2:E:239:LEU:HD23	2:E:338:VAL:HG21	1.72	0.70
2:E:294:PRO:HD2	2:E:351:TYR:CE2	2.25	0.70
1:D:458:GLU:OE2	2:F:264:ARG:HB3	1.90	0.70
1:A:1008:LEU:C	1:A:1008:LEU:HD22	2.11	0.70
1:A:1029:GLN:NE2	1:A:1096:ARG:CZ	2.54	0.70
1:A:1043:VAL:HG22	1:A:1046:GLU:HB2	1.72	0.70
1:A:214:ALA:CB	1:A:219:ALA:HB3	2.20	0.70
1:A:247:LEU:HD23	1:A:248:GLU:OE1	1.91	0.70
1:A:767:ALA:O	1:A:771:LEU:HB2	1.91	0.70
1:A:950:ASN:HA	1:A:953:ARG:HG3	1.73	0.70
2:B:259:ARG:HH11	2:B:259:ARG:HB3	1.56	0.70
2:C:200:VAL:O	2:C:203:ARG:HB2	1.90	0.70
2:C:241:TRP:HB2	2:C:336:LEU:CG	2.21	0.70
2:E:307:GLY:HA2	2:E:334:CYS:CB	2.21	0.70
1:D:754:HIS:O	1:D:758:ASN:N	2.21	0.70
1:D:1014:ARG:HD3	1:D:1026:ARG:NE	2.06	0.70
1:D:1232:GLU:HG3	1:D:1234:ARG:NE	2.06	0.70
1:D:265:LEU:HD21	1:D:267:VAL:CA	2.21	0.70
1:D:297:MET:SD	1:D:415:LEU:HD22	2.31	0.70
2:E:320:SER:O	2:E:323:HIS:HD2	1.73	0.70
1:A:630:ASN:HD22	1:A:631:LEU:N	1.90	0.70
2:B:202:LYS:O	2:B:325:ARG:CZ	2.39	0.70
2:B:383:VAL:HG22	2:B:384:ALA:H	1.56	0.70
2:B:387:VAL:HG11	2:B:396:ARG:HB2	1.73	0.70
1:D:576:LEU:HD12	1:D:577:CYS:N	2.06	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:732:TYR:CD2	1:D:732:TYR:N	2.59	0.70
1:D:801:TRP:HH2	1:D:805:HIS:CG	2.08	0.70
2:E:203:ARG:HH12	2:F:418:LEU:CD2	1.99	0.70
2:F:254:PHE:N	2:F:257:ARG:HH21	1.89	0.70
2:F:243:THR:OG1	2:F:334:CYS:CB	2.39	0.70
1:A:439:GLN:CG	1:A:836:LEU:N	2.54	0.70
1:A:549:LEU:HD23	1:A:552:LEU:HD12	1.73	0.70
1:A:601:LYS:HZ1	1:A:721:ALA:HB1	1.55	0.70
1:A:608:ASP:HB3	1:A:779:LEU:CG	2.21	0.70
1:A:813:VAL:HG22	1:A:814:VAL:N	2.06	0.70
1:A:933:SER:HA	1:A:943:ARG:HG2	1.71	0.70
2:B:373:LYS:HA	2:B:458:ARG:CZ	2.22	0.70
2:C:239:LEU:O	2:C:338:VAL:HG12	1.92	0.70
1:A:472:GLN:CG	2:C:460:THR:HA	2.20	0.70
1:A:634:LEU:HD12	1:A:638:THR:HB	1.72	0.70
1:D:1192:GLU:O	1:D:1192:GLU:HG2	1.89	0.70
1:D:225:SER:HA	1:D:228:LEU:HD21	1.72	0.70
1:D:203:LEU:CB	1:D:380:GLY:HA3	2.17	0.70
1:D:374:ARG:CB	1:D:392:LEU:HD11	2.21	0.70
1:D:598:VAL:HG12	1:D:602:LEU:HD11	1.73	0.70
1:D:885:THR:HB	1:D:1146:ARG:CD	2.17	0.70
2:E:109:ASN:ND2	2:E:378:LEU:HB3	2.06	0.70
2:F:239:LEU:H	2:F:338:VAL:HG13	1.55	0.70
2:F:454:HIS:CB	2:F:466:MET:CB	2.66	0.70
1:A:247:LEU:CD1	1:A:251:THR:HG21	2.21	0.70
1:A:355:SER:HA	1:A:407:PHE:CE1	2.26	0.70
1:A:515:ILE:CD1	1:A:568:GLY:O	2.40	0.70
1:A:627:ARG:O	1:A:628:ARG:HB3	1.90	0.70
1:A:655:LEU:O	1:A:719:LEU:HD12	1.91	0.70
1:A:771:LEU:H	1:A:771:LEU:CD2	2.02	0.70
2:C:299:LEU:H	2:C:299:LEU:HD23	1.54	0.70
2:C:439:PHE:CD1	2:C:458:ARG:HB2	2.25	0.70
2:C:470:LYS:HE3	2:C:471:LEU:HD21	1.74	0.70
2:E:249:ASN:HD21	2:E:285:LYS:CE	1.97	0.70
1:D:1158:LEU:HG	1:D:1159:LEU:H	1.56	0.70
1:D:190:PRO:C	1:D:192:GLU:N	2.44	0.70
1:D:266:VAL:O	1:D:266:VAL:HG13	1.91	0.70
1:D:278:ILE:HD13	1:D:291:PHE:HD1	1.55	0.70
1:D:401:TRP:HA	1:D:404:HIS:CD2	2.26	0.70
1:D:576:LEU:CG	1:D:579:ARG:HE	2.05	0.70
2:E:441:VAL:HG12	2:E:442:LEU:H	1.57	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:394:GLU:CD	2:F:394:GLU:H	1.92	0.70
1:A:1212:ILE:CG2	1:A:1213:PRO:N	2.55	0.70
2:C:214:CYS:SG	2:C:236:GLU:HG3	2.32	0.70
2:C:471:LEU:HA	2:C:474:PHE:HD2	1.54	0.70
1:D:623:LEU:HG	1:D:623:LEU:O	1.89	0.70
1:D:127:LEU:CD2	1:D:129:PRO:HD3	2.20	0.70
1:D:474:LEU:CD1	1:D:474:LEU:H	2.04	0.70
1:D:603:MET:HB3	1:D:605:LEU:CD2	2.22	0.70
1:D:601:LYS:HG2	1:D:721:ALA:CB	2.21	0.70
1:D:810:SER:OG	1:D:811:GLN:N	2.24	0.70
2:E:118:VAL:HG13	2:E:119:VAL:H	1.57	0.70
1:A:1066:THR:OG1	1:A:1071:ARG:HB3	1.91	0.70
1:A:1019:TRP:HZ3	1:A:1164:PHE:HA	1.56	0.70
1:A:125:LEU:N	1:A:125:LEU:CD2	2.50	0.70
1:A:239:LEU:O	1:A:241:PRO:CD	2.38	0.70
2:C:125:VAL:HG23	2:C:126:PHE:H	1.57	0.70
2:F:326:ASP:CG	2:F:327:GLY:N	2.42	0.70
1:D:597:ARG:HH11	1:D:724:GLY:C	1.94	0.70
1:D:612:LEU:HD13	1:D:612:LEU:H	1.53	0.70
2:E:114:TRP:HH2	2:E:238:SER:O	1.75	0.70
2:E:326:ASP:H	2:E:331:VAL:HG12	1.55	0.70
2:E:430:SER:O	2:E:434:GLU:HG2	1.91	0.70
1:A:169:TRP:CE3	1:A:390:GLN:O	2.45	0.70
1:A:606:THR:HG22	1:A:614:TYR:CD2	2.27	0.70
1:A:869:ARG:NH1	1:A:869:ARG:HG3	2.06	0.70
1:A:883:GLY:C	1:A:884:TYR:CD1	2.64	0.70
2:B:199:LEU:CD1	2:B:200:VAL:HG13	2.22	0.70
2:C:207:GLY:HA2	2:C:240:VAL:O	1.91	0.70
2:C:342:LEU:HB2	2:C:344:ARG:HB2	1.74	0.70
2:C:351:TYR:O	2:C:352:ASP:C	2.27	0.70
2:C:393:LEU:O	2:C:397:GLN:N	2.23	0.70
2:C:455:LEU:CA	2:C:456:ARG:HE	2.04	0.70
1:D:371:LYS:CA	1:D:374:ARG:HD2	2.22	0.70
1:D:420:HIS:CD2	1:D:1020:ILE:CD1	2.69	0.70
1:D:607:TRP:HD1	1:D:779:LEU:HA	1.56	0.70
1:D:892:ASP:OD2	1:D:892:ASP:C	2.30	0.70
2:F:446:THR:CG2	2:F:450:ASN:OD1	2.35	0.70
2:F:452:LEU:O	2:F:453:ILE:CG1	2.39	0.70
1:A:273:PHE:C	1:A:276:ALA:HB3	2.12	0.70
1:A:282:TYR:O	1:A:283:LEU:CB	2.39	0.70
1:A:412:PRO:CG	1:A:413:LEU:H	2.05	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:448:ALA:CA	1:A:870:VAL:HG13	2.21	0.70
1:A:783:PRO:HG2	2:C:363:ARG:NH1	2.07	0.70
2:B:204:LEU:HD12	2:B:325:ARG:N	2.05	0.70
1:D:107:LEU:HD23	1:D:107:LEU:N	2.06	0.70
1:D:1118:TRP:O	1:D:1122:GLU:HG2	1.92	0.70
1:D:889:ALA:HB3	1:D:1180:PHE:CE1	2.27	0.70
1:D:121:PRO:CB	1:D:1175:GLN:HE22	2.04	0.70
1:D:633:LYS:CD	1:D:638:THR:HG21	2.22	0.70
1:D:737:GLY:O	1:D:741:ASP:OD1	2.10	0.70
1:D:774:MET:O	1:D:779:LEU:O	2.10	0.70
1:D:993:ARG:HD3	1:D:993:ARG:O	1.92	0.70
2:F:241:TRP:HZ3	2:F:255:TRP:CE2	2.08	0.70
1:A:192:GLU:HB3	1:A:217:PRO:CG	2.18	0.70
1:A:211:LEU:HD12	1:A:221:TYR:HB3	1.73	0.70
1:A:402:ALA:O	1:A:406:VAL:HG13	1.92	0.70
1:A:417:ARG:HH11	1:A:417:ARG:HG2	1.57	0.70
1:A:451:THR:HG21	1:A:1202:ASN:CB	2.22	0.70
1:A:831:TYR:N	1:A:831:TYR:CD1	2.59	0.70
1:A:900:ALA:CA	1:A:915:ALA:HB1	2.22	0.70
2:B:215:PHE:O	2:B:216:HIS:HB2	1.91	0.70
2:B:117:SER:HB2	2:B:262:TRP:HE1	1.54	0.70
2:C:200:VAL:HG11	2:C:204:LEU:HB2	1.74	0.70
2:C:263:TRP:CZ2	2:C:301:GLU:OE2	2.45	0.70
2:C:344:ARG:HA	2:C:347:LEU:CD2	2.22	0.70
2:C:375:HIS:ND1	2:C:376:PRO:HD2	2.06	0.70
2:C:389:ARG:CZ	2:C:390:GLY:HA3	2.20	0.70
1:D:832:ASP:O	1:D:833:GLU:HB2	1.92	0.70
1:D:1014:ARG:HD3	1:D:1026:ARG:CG	2.21	0.70
1:D:1175:GLN:C	1:D:1177:VAL:H	1.94	0.70
1:D:1198:LYS:HD2	1:D:1204:THR:HB	1.73	0.70
1:D:267:VAL:HG21	1:D:291:PHE:HE2	1.56	0.70
1:D:374:ARG:HG2	1:D:374:ARG:HH11	1.57	0.70
1:D:294:THR:CG2	1:D:407:PHE:CE2	2.70	0.70
1:D:933:SER:HB2	1:D:943:ARG:CD	2.20	0.70
1:D:944:GLU:HB3	1:D:1071:ARG:HD3	1.74	0.70
2:E:437:ILE:HD12	2:E:437:ILE:N	2.04	0.70
2:F:213:VAL:HA	2:F:235:THR:HA	1.72	0.70
2:F:342:LEU:HB2	2:F:344:ARG:HD3	1.72	0.70
2:F:467:HIS:HB3	2:F:470:LYS:HE2	1.74	0.70
2:F:471:LEU:H	2:F:471:LEU:CD1	1.67	0.70
1:A:1031:GLU:O	1:A:1032:THR:HB	1.91	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:174:GLY:O	1:A:175:TRP:CG	2.44	0.70
1:A:603:MET:HE3	1:A:791:ALA:HB1	1.74	0.70
1:A:821:LEU:CD1	1:A:825:VAL:HB	2.22	0.70
1:A:827:ARG:HA	1:A:831:TYR:OH	1.91	0.70
2:C:127:PRO:CD	2:C:128:VAL:H	1.98	0.70
1:D:569:HIS:N	1:D:570:PRO:CD	2.55	0.70
2:F:382:LYS:HD3	2:F:411:ILE:CG1	2.18	0.70
1:D:500:ALA:HB2	1:D:521:PRO:HG3	1.72	0.70
1:D:527:GLN:HE21	1:D:528:GLU:HB3	1.56	0.70
1:D:370:GLU:O	1:D:371:LYS:HB2	1.92	0.69
1:D:600:PRO:CG	1:D:616:GLU:HB3	2.21	0.69
2:F:342:LEU:CB	2:F:344:ARG:HD3	2.22	0.69
1:A:107:LEU:HB2	1:A:113:TRP:HE1	1.54	0.69
1:A:77:GLN:HB2	1:A:1174:PRO:HB3	1.72	0.69
1:A:211:LEU:HG	1:A:221:TYR:HB3	1.74	0.69
1:A:886:LEU:HD23	1:A:886:LEU:N	2.07	0.69
1:A:927:ARG:CG	1:A:928:GLY:N	2.53	0.69
2:E:88:ARG:N	2:E:88:ARG:HD3	2.07	0.69
1:D:1008:LEU:HA	1:D:1011:PRO:HG3	1.73	0.69
1:D:205:GLU:HA	1:D:205:GLU:OE1	1.91	0.69
1:D:256:PRO:HB3	1:D:281:GLN:CG	2.22	0.69
1:D:381:THR:CG2	1:D:385:ILE:HB	2.21	0.69
1:D:547:ALA:C	1:D:551:LYS:CE	2.61	0.69
1:D:660:CYS:HB2	1:D:745:PRO:CD	2.21	0.69
1:D:658:LYS:HE3	1:D:713:PRO:CB	2.21	0.69
1:D:845:VAL:HG23	1:D:855:VAL:HB	1.74	0.69
1:D:436:PRO:O	1:D:879:GLN:HB3	1.92	0.69
2:F:81:GLY:HA3	2:F:97:PRO:HB2	1.75	0.69
1:A:891:VAL:CA	1:A:1180:PHE:HE1	2.03	0.69
1:A:1209:ARG:HH12	2:C:256:LEU:HB3	1.57	0.69
1:A:869:ARG:CA	1:A:1200:PRO:HA	2.22	0.69
2:C:241:TRP:HZ3	2:C:255:TRP:NE1	1.89	0.69
2:B:394:GLU:O	2:B:398:VAL:HG23	1.92	0.69
2:F:94:GLY:HA2	2:F:96:HIS:CE1	2.27	0.69
1:D:1111:LEU:HD22	1:D:1164:PHE:CE1	2.28	0.69
1:D:177:ARG:O	1:D:179:GLY:N	2.25	0.69
1:D:656:TYR:HE1	1:D:743:ASP:CG	1.96	0.69
1:D:798:ILE:CA	1:D:869:ARG:CZ	2.69	0.69
1:D:891:VAL:HG11	1:D:1135:ASP:CA	2.22	0.69
2:E:214:CYS:HB2	2:E:236:GLU:HB2	1.73	0.69
2:E:387:VAL:CG1	2:E:416:GLY:HA3	2.23	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:113:GLU:O	2:F:116:THR:HG23	1.93	0.69
2:F:389:ARG:O	2:F:392:THR:HA	1.92	0.69
2:F:371:VAL:HG23	2:F:433:ASP:C	2.12	0.69
1:A:483:PRO:HB3	1:A:712:VAL:HG11	1.73	0.69
1:A:869:ARG:O	1:A:1200:PRO:HA	1.91	0.69
1:A:886:LEU:N	1:A:1146:ARG:HH21	1.88	0.69
1:A:992:LEU:HD12	1:A:993:ARG:H	1.56	0.69
2:C:251:TRP:HA	2:C:251:TRP:HE3	1.57	0.69
2:C:363:ARG:NH1	2:C:363:ARG:HA	2.05	0.69
2:C:435:MET:O	2:C:436:SER:HB3	1.90	0.69
1:A:540:GLN:O	1:A:543:VAL:HG12	1.93	0.69
1:D:457:ARG:C	1:D:460:LYS:CE	2.60	0.69
1:D:556:THR:OG1	2:F:452:LEU:CD2	2.41	0.69
2:E:455:LEU:HD23	2:E:464:GLU:O	1.91	0.69
1:A:211:LEU:CG	1:A:221:TYR:HB3	2.22	0.69
1:A:195:LEU:HD23	1:A:217:PRO:HD2	1.73	0.69
1:A:658:LYS:O	1:A:662:GLU:HG2	1.93	0.69
1:A:827:ARG:HG3	1:A:828:HIS:N	2.08	0.69
2:B:269:SER:HB2	2:B:352:ASP:OD1	1.92	0.69
2:C:403:PHE:O	2:C:407:LEU:N	2.23	0.69
1:D:516:GLU:HA	1:D:565:HIS:CE1	2.26	0.69
2:C:134:LYS:HB3	2:C:180:LYS:CE	2.23	0.69
1:A:634:LEU:CG	1:A:635:PRO:HA	2.22	0.69
1:A:737:GLY:O	1:A:741:ASP:OD2	2.10	0.69
1:D:887:VAL:CG2	1:D:1146:ARG:CZ	2.69	0.69
1:D:348:LEU:HD11	1:D:350:ILE:HD11	1.75	0.69
1:D:788:GLY:O	1:D:791:ALA:N	2.24	0.69
2:E:116:THR:O	2:E:120:VAL:HG12	1.92	0.69
2:E:85:GLN:HB3	2:E:97:PRO:HG3	1.75	0.69
1:D:1209:ARG:NH2	2:F:253:ASP:N	2.40	0.69
2:F:254:PHE:HE2	2:F:255:TRP:CZ2	2.11	0.69
1:A:1199:THR:CA	1:A:1200:PRO:C	2.52	0.69
1:A:192:GLU:CA	1:A:217:PRO:HG2	2.22	0.69
1:A:667:GLN:C	1:A:670:PRO:HD2	2.12	0.69
1:A:732:TYR:HE1	1:A:755:LYS:CA	2.03	0.69
2:B:261:GLN:CA	2:B:264:ARG:HE	2.05	0.69
2:B:72:ILE:HD12	2:B:372:LEU:HD11	1.74	0.69
2:C:341:ASP:C	2:C:342:LEU:HD13	2.13	0.69
1:D:1192:GLU:OE2	1:D:1195:MET:SD	2.51	0.69
1:D:1197:CYS:N	1:D:1204:THR:CG2	2.54	0.69
1:D:127:LEU:HD22	1:D:129:PRO:CD	2.22	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:768:LYS:CD	1:D:769:ASP:N	2.55	0.69
1:D:768:LYS:C	1:D:772:PRO:HD2	2.13	0.69
1:D:85:GLU:OE1	1:D:130:LEU:HD22	1.92	0.69
1:D:992:LEU:HD23	1:D:997:LEU:HD12	1.73	0.69
2:E:254:PHE:HB2	2:E:257:ARG:NH1	2.03	0.69
2:E:384:ALA:O	2:E:414:TRP:N	2.26	0.69
1:A:115:GLN:N	1:A:116:PRO:CD	2.56	0.69
1:A:1199:THR:HA	1:A:1201:SER:N	2.07	0.69
1:A:624:VAL:C	1:A:626:GLY:N	2.43	0.69
1:A:653:GLU:HG3	1:A:656:TYR:HE1	1.58	0.69
1:A:769:ASP:O	1:A:773:LYS:HG3	1.91	0.69
1:A:792:LEU:HA	1:A:795:ASN:OD1	1.91	0.69
2:B:78:PHE:CG	2:B:102:LEU:HB2	2.27	0.69
2:B:324:GLY:CA	2:B:325:ARG:NH2	2.55	0.69
2:C:455:LEU:HD12	2:C:455:LEU:N	2.08	0.69
2:C:439:PHE:HD1	2:C:458:ARG:HG3	1.54	0.69
1:D:127:LEU:O	1:D:130:LEU:HD21	1.93	0.69
1:D:648:PRO:CD	1:D:649:TYR:H	2.01	0.69
2:E:289:LEU:O	2:E:299:LEU:HA	1.92	0.69
2:F:314:MET:HB3	2:F:315:TYR:CD2	2.28	0.69
2:F:374:LEU:CD1	2:F:379:ALA:HA	2.21	0.69
2:F:81:GLY:CA	2:F:97:PRO:HB2	2.22	0.69
1:A:285:GLN:C	1:A:1128:ARG:NH2	2.46	0.69
1:A:805:HIS:C	1:A:805:HIS:ND1	2.45	0.69
2:B:234:LYS:HE2	2:B:234:LYS:N	2.07	0.69
2:C:435:MET:O	2:C:436:SER:CB	2.40	0.69
2:E:391:PRO:HD2	2:E:395:LEU:HB2	1.73	0.69
1:A:741:ASP:OD1	1:A:742:VAL:N	2.25	0.69
1:D:1047:ARG:O	1:D:1050:LYS:N	2.26	0.69
1:D:167:PRO:O	1:D:169:TRP:N	2.25	0.69
1:D:279:ARG:CB	1:D:841:LEU:HB2	2.22	0.69
2:E:371:VAL:CG1	2:E:372:LEU:N	2.55	0.69
1:D:1107:ASP:O	1:D:1111:LEU:HD13	1.93	0.69
1:D:273:PHE:HA	1:D:276:ALA:H	1.58	0.69
1:D:606:THR:OG1	1:D:614:TYR:CZ	2.46	0.69
1:D:641:GLU:O	1:D:650:ARG:HD3	1.91	0.69
2:E:439:PHE:CZ	2:E:478:TYR:CE1	2.81	0.69
2:F:207:GLY:HA3	2:F:239:LEU:HD11	1.75	0.69
1:A:1006:ARG:HH22	1:A:1010:LEU:HD11	1.56	0.69
1:A:165:LYS:O	1:A:167:PRO:HD3	1.92	0.69
1:A:213:VAL:HG13	1:A:221:TYR:HH	1.53	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:352:SER:O	1:A:358:LEU:CD2	2.40	0.69
1:A:363:ARG:HD2	1:A:406:VAL:CB	2.16	0.69
1:A:559:LEU:HB3	1:A:560:PRO:CD	2.22	0.69
1:A:926:SER:O	1:A:927:ARG:CB	2.41	0.69
1:A:963:GLU:O	1:A:967:MET:HG3	1.93	0.69
2:B:99:PHE:CE1	2:B:343:ASP:OD1	2.46	0.69
2:C:241:TRP:CD1	2:C:243:THR:HG23	2.28	0.69
2:C:307:GLY:HA2	2:C:335:VAL:HG12	1.75	0.69
2:C:388:GLY:HA3	2:C:444:THR:HG22	1.73	0.69
1:A:451:THR:OG1	1:A:1202:ASN:ND2	2.26	0.69
1:A:183:GLU:OE1	1:A:184:ALA:HB2	1.93	0.69
1:A:992:LEU:HD13	1:A:993:ARG:H	1.57	0.69
2:B:383:VAL:HG22	2:B:384:ALA:N	2.07	0.69
2:C:357:THR:C	2:C:359:ASN:H	1.97	0.69
2:C:388:GLY:O	2:C:395:LEU:HD13	1.92	0.69
2:C:81:GLY:HA3	2:C:97:PRO:CG	2.23	0.69
1:A:1193:VAL:CG2	1:A:1194:THR:N	2.55	0.69
1:D:872:SER:OG	1:D:1200:PRO:CD	2.40	0.69
1:D:483:PRO:O	1:D:486:TRP:CB	2.41	0.69
1:D:725:PRO:O	1:D:726:LYS:HB2	1.92	0.69
1:D:742:VAL:HG13	1:D:743:ASP:N	2.07	0.69
1:D:656:TYR:CB	1:D:744:ILE:HG21	2.22	0.69
1:D:76:ILE:HB	1:D:911:HIS:CE1	2.27	0.69
1:D:801:TRP:HD1	1:D:869:ARG:HH21	1.38	0.69
1:D:826:ILE:HB	1:D:831:TYR:HE2	1.56	0.69
1:D:946:ALA:HA	1:D:949:PHE:CD2	2.28	0.69
1:D:996:ARG:CZ	1:D:996:ARG:H	2.05	0.69
2:E:183:GLU:HG3	2:E:216:HIS:CD2	2.28	0.69
2:E:193:TYR:CD2	2:E:322:LEU:HD22	2.27	0.69
2:F:254:PHE:HA	2:F:257:ARG:HH21	1.56	0.69
2:F:356:LEU:HD22	2:F:359:ASN:OD1	1.93	0.69
1:A:107:LEU:HD12	1:A:112:LEU:HD13	1.74	0.69
1:A:853:ARG:NH2	1:A:1105:ALA:CB	2.55	0.69
1:A:451:THR:C	1:A:869:ARG:HH22	1.95	0.69
1:A:462:SER:OG	1:A:463:LEU:N	2.21	0.69
1:A:719:LEU:O	1:A:722:ARG:HG3	1.93	0.69
1:A:827:ARG:HA	1:A:831:TYR:CE1	2.28	0.69
1:A:951:TYR:HB3	1:A:955:TYR:HE1	1.58	0.69
2:C:341:ASP:O	2:C:342:LEU:CD1	2.38	0.69
2:C:389:ARG:HG3	2:C:390:GLY:H	1.57	0.69
1:D:1197:CYS:C	1:D:1204:THR:HG1	1.95	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1231:LEU:CG	1:D:1232:GLU:H	1.83	0.69
1:D:167:PRO:HD2	1:D:168:ALA:H	1.55	0.69
1:D:307:PHE:HD1	1:D:307:PHE:H	1.39	0.69
2:E:371:VAL:HG12	2:E:372:LEU:N	2.07	0.69
2:E:450:ASN:HD22	2:E:452:LEU:HD21	1.57	0.69
2:F:114:TRP:NE1	2:F:259:ARG:NH2	2.40	0.69
2:E:419:GLU:HA	2:F:203:ARG:HH22	1.54	0.69
2:F:344:ARG:HA	2:F:347:LEU:HB2	1.73	0.69
2:F:344:ARG:HA	2:F:347:LEU:HD13	1.75	0.69
2:F:355:GLN:O	2:F:357:THR:N	2.25	0.69
2:F:389:ARG:CZ	2:F:390:GLY:HA3	2.23	0.69
1:A:1024:ASP:O	1:A:1026:ARG:HD2	1.92	0.69
1:A:1114:VAL:N	1:A:1117:LYS:HZ2	1.90	0.69
1:A:192:GLU:CD	1:A:193:ARG:N	2.46	0.69
1:A:214:ALA:CA	1:A:401:TRP:CZ2	2.75	0.69
1:A:373:PRO:HD2	1:A:374:ARG:HH11	1.58	0.69
1:A:363:ARG:CZ	1:A:406:VAL:HA	2.22	0.69
1:A:623:LEU:H	1:A:623:LEU:HD23	1.58	0.69
1:A:850:ILE:HG12	1:A:1099:TRP:NE1	2.07	0.69
1:A:86:GLN:O	1:A:87:ILE:HG13	1.92	0.69
2:C:303:LEU:C	2:C:337:SER:CB	2.60	0.69
2:B:199:LEU:HD23	2:C:77:HIS:CD2	2.27	0.69
2:F:272:ASN:HB3	2:F:292:ASN:HB2	1.73	0.69
1:D:1019:TRP:HD1	1:D:1020:ILE:HG22	1.57	0.68
1:D:1232:GLU:CG	1:D:1234:ARG:HB2	2.22	0.68
1:D:356:LEU:CD1	1:D:369:LEU:CD2	2.64	0.68
1:D:457:ARG:C	1:D:460:LYS:HE3	2.12	0.68
1:D:496:LYS:HE3	1:D:571:GLY:HA2	1.73	0.68
1:D:595:GLN:O	1:D:596:MET:C	2.31	0.68
1:D:283:LEU:HG	1:D:814:VAL:HG23	1.74	0.68
1:D:990:LYS:HG2	1:D:1000:GLU:HB2	1.73	0.68
2:E:456:ARG:NE	2:E:463:LYS:HE3	2.07	0.68
2:E:82:SER:N	2:E:85:GLN:HG3	2.04	0.68
1:A:853:ARG:NH2	1:A:1105:ALA:HB3	2.07	0.68
1:A:420:HIS:ND1	1:A:421:PRO:HD2	2.08	0.68
1:A:453:GLU:O	1:A:454:GLU:C	2.31	0.68
1:A:559:LEU:HD23	1:A:560:PRO:HD2	1.74	0.68
1:A:601:LYS:NZ	1:A:616:GLU:HG2	2.07	0.68
1:A:951:TYR:O	1:A:954:ILE:HB	1.93	0.68
2:B:115:TRP:CZ2	2:C:108:LYS:HD3	2.29	0.68
2:C:402:LEU:HG	2:C:472:LYS:HZ3	1.58	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:443:VAL:CG1	2:C:447:THR:HG21	2.22	0.68
1:A:230:GLU:CD	1:A:231:GLU:H	1.91	0.68
1:D:312:TRP:HH2	1:D:346:ASP:O	1.75	0.68
1:D:482:ASP:HB2	1:D:483:PRO:HD2	1.74	0.68
1:D:491:ASP:OD2	1:D:578:PRO:HB3	1.94	0.68
1:D:651:ALA:O	1:D:655:LEU:HD23	1.93	0.68
1:D:656:TYR:CD1	1:D:743:ASP:HB2	2.28	0.68
1:D:767:ALA:HB3	1:D:771:LEU:HD22	1.75	0.68
1:D:887:VAL:N	1:D:1146:ARG:NH1	2.39	0.68
2:F:244:PRO:CD	2:F:247:THR:CG2	2.71	0.68
1:A:1030:ARG:HD3	1:A:1091:GLU:OE1	1.91	0.68
1:A:1129:PHE:HD2	1:A:1138:ARG:O	1.76	0.68
1:A:1223:ILE:O	1:A:1227:THR:HG23	1.93	0.68
1:A:607:TRP:CH2	1:A:671:GLN:HA	2.27	0.68
1:A:659:HIS:CB	1:A:719:LEU:HD12	2.17	0.68
1:A:997:LEU:CD2	1:A:998:SER:N	2.51	0.68
2:C:353:SER:HB2	2:C:373:LYS:O	1.94	0.68
2:C:444:THR:H	2:C:447:THR:HG23	1.58	0.68
1:A:564:GLN:CA	1:A:567:PRO:HD2	2.23	0.68
1:D:1027:LYS:HG3	1:D:1091:GLU:HA	1.74	0.68
1:D:1202:ASN:O	1:D:1205:GLY:N	2.27	0.68
1:D:440:ASN:OD1	1:D:1210:TYR:O	2.11	0.68
1:D:369:LEU:HB3	1:D:372:GLU:HG3	1.74	0.68
1:D:414:PHE:O	1:D:417:ARG:HB2	1.94	0.68
1:D:424:LEU:N	1:D:424:LEU:HD23	2.07	0.68
1:D:814:VAL:O	1:D:838:GLY:HA3	1.94	0.68
1:D:856:GLU:HB2	1:D:860:LEU:H	1.58	0.68
1:D:952:GLY:O	1:D:954:ILE:O	2.10	0.68
2:E:460:THR:O	2:E:461:THR:HG22	1.93	0.68
2:F:203:ARG:HG2	2:F:325:ARG:NE	2.08	0.68
2:F:81:GLY:HA3	2:F:97:PRO:CD	2.23	0.68
1:A:190:PRO:CB	1:A:192:GLU:HG3	2.23	0.68
1:A:192:GLU:HB2	1:A:195:LEU:HD21	1.75	0.68
1:A:222:SER:CB	1:A:223:TRP:HD1	2.02	0.68
1:A:374:ARG:HH11	1:A:374:ARG:CG	2.06	0.68
1:A:203:LEU:HD12	1:A:379:LYS:C	2.13	0.68
1:A:476:GLY:HA3	1:A:711:ALA:HB1	1.76	0.68
1:A:734:HIS:HA	1:A:752:LEU:HD11	1.76	0.68
1:A:307:PHE:H	1:A:307:PHE:HD1	1.42	0.68
2:E:187:HIS:CA	2:E:190:LEU:HD22	2.22	0.68
2:B:217:PRO:O	2:B:230:SER:O	2.12	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1129:PHE:HB3	1:D:1139:TYR:HA	1.75	0.68
1:D:880:ALA:CB	1:D:1187:ARG:O	2.39	0.68
1:D:1138:ARG:HH22	1:D:1191:LYS:HE2	1.57	0.68
1:D:131:TYR:HB3	1:D:138:HIS:CE1	2.29	0.68
1:D:167:PRO:CD	1:D:168:ALA:N	2.56	0.68
1:D:203:LEU:O	1:D:203:LEU:HD23	1.93	0.68
1:D:659:HIS:NE2	1:D:719:LEU:HB2	2.08	0.68
2:E:318:ASN:O	2:E:321:LYS:N	2.18	0.68
2:E:443:VAL:CG2	2:E:453:ILE:HG21	2.20	0.68
2:F:78:PHE:CD1	2:F:102:LEU:HD22	2.29	0.68
1:A:1199:THR:HA	1:A:1200:PRO:O	1.93	0.68
1:A:299:MET:CE	1:A:353:VAL:HB	2.24	0.68
1:A:297:MET:SD	1:A:411:LEU:CB	2.80	0.68
1:A:582:ASP:N	1:A:583:PRO:CD	2.56	0.68
1:A:630:ASN:ND2	1:A:632:ALA:H	1.92	0.68
1:A:713:PRO:HA	1:A:719:LEU:HD13	1.75	0.68
1:A:767:ALA:HB3	1:A:771:LEU:CD2	2.24	0.68
1:A:974:THR:CG2	1:A:975:GLN:N	2.56	0.68
2:B:116:THR:HA	2:B:120:VAL:CG2	2.24	0.68
2:B:417:TYR:C	2:B:419:GLU:H	1.97	0.68
2:C:239:LEU:HG	2:C:336:LEU:HD13	1.75	0.68
2:C:357:THR:HA	2:C:370:LYS:HZ1	1.55	0.68
1:D:131:TYR:HB3	1:D:138:HIS:HE1	1.58	0.68
1:D:307:PHE:HA	1:D:310:SER:OG	1.94	0.68
1:D:645:VAL:HG13	1:D:650:ARG:CG	2.23	0.68
1:D:739:TYR:CD2	1:D:740:ASN:N	2.62	0.68
1:D:770:PHE:HA	1:D:773:LYS:HG3	1.76	0.68
1:D:275:ARG:NH1	1:D:843:GLN:NE2	2.40	0.68
2:E:377:CYS:HA	2:E:380:PRO:HG2	1.74	0.68
2:E:411:ILE:HD12	2:E:411:ILE:N	2.09	0.68
2:E:450:ASN:HB3	2:E:452:LEU:CD2	2.23	0.68
2:F:268:MET:N	2:F:352:ASP:OD2	2.27	0.68
1:A:167:PRO:HB2	1:A:177:ARG:CZ	2.24	0.68
1:A:286:GLY:O	1:A:287:SER:HB3	1.91	0.68
1:A:577:CYS:CB	1:A:578:PRO:CD	2.72	0.68
1:A:890:ASP:CG	1:A:1191:LYS:NZ	2.47	0.68
2:B:288:LYS:HG2	2:B:302:THR:HG22	1.74	0.68
1:D:623:LEU:HD22	1:D:667:GLN:OE1	1.94	0.68
2:B:211:ILE:HG23	2:B:237:ALA:HB2	1.73	0.68
1:D:347:TRP:HE1	1:D:349:ASP:CG	1.96	0.68
1:D:736:ASN:HA	1:D:750:PHE:CZ	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:922:GLN:NE2	1:D:931:LEU:HD13	2.08	0.68
2:F:203:ARG:HB3	2:F:204:LEU:HD12	1.76	0.68
2:F:455:LEU:HA	2:F:456:ARG:NH2	2.07	0.68
2:F:474:PHE:HA	2:F:477:LYS:HD2	1.75	0.68
1:A:245:ILE:O	1:A:248:GLU:CD	2.32	0.68
1:A:498:LYS:O	1:A:499:LYS:HB3	1.91	0.68
1:A:597:ARG:CB	1:A:616:GLU:OE1	2.40	0.68
1:A:865:ALA:O	1:A:868:ASP:OD2	2.12	0.68
1:A:884:TYR:N	1:A:884:TYR:HD1	1.91	0.68
2:C:132:HIS:HA	2:C:182:ARG:HG3	1.75	0.68
2:C:206:TYR:O	2:C:241:TRP:CD1	2.47	0.68
2:C:471:LEU:HA	2:C:474:PHE:CD2	2.28	0.68
2:B:479:ILE:HG22	2:B:480:SER:N	2.07	0.68
1:A:741:ASP:CG	1:A:742:VAL:HG23	2.13	0.68
1:D:316:LYS:HG2	1:D:348:LEU:CG	2.20	0.68
1:D:301:ILE:HD11	1:D:414:PHE:O	1.94	0.68
1:D:492:LEU:HD22	1:D:495:PHE:CZ	2.29	0.68
1:D:549:LEU:N	1:D:551:LYS:HZ3	1.84	0.68
1:D:622:TYR:N	1:D:747:CYS:O	2.26	0.68
1:D:607:TRP:CD1	1:D:779:LEU:HA	2.29	0.68
1:D:847:ALA:HA	1:D:852:ARG:HB3	1.75	0.68
2:E:107:ARG:CG	2:E:107:ARG:NH1	2.33	0.68
2:E:318:ASN:O	2:E:320:SER:N	2.27	0.68
2:E:400:GLN:HA	2:E:403:PHE:HB3	1.76	0.68
2:F:314:MET:HB3	2:F:315:TYR:CE2	2.28	0.68
2:F:322:LEU:HD22	2:F:322:LEU:C	2.13	0.68
1:A:869:ARG:CA	1:A:1201:SER:H	2.07	0.68
1:A:605:LEU:C	1:A:781:ALA:HB3	2.14	0.68
1:A:792:LEU:HD12	1:A:793:GLU:HG2	1.76	0.68
2:B:463:LYS:HD2	2:B:463:LYS:H	1.58	0.68
2:B:469:SER:C	2:B:471:LEU:N	2.47	0.68
1:A:973:LEU:HB3	1:A:977:GLU:CB	2.24	0.68
1:D:1206:MET:HA	2:F:253:ASP:OD2	1.94	0.68
1:D:79:LEU:HD23	1:D:83:LEU:CD1	2.22	0.68
1:D:914:THR:OG1	1:D:999:ASP:HB3	1.92	0.68
2:E:407:LEU:HA	2:E:411:ILE:O	1.93	0.68
2:F:259:ARG:NE	2:F:339:ASN:HD21	1.91	0.68
2:F:442:LEU:H	2:F:455:LEU:HD22	1.58	0.68
1:A:1220:ILE:CD1	1:A:1220:ILE:N	2.57	0.68
1:A:276:ALA:CA	1:A:279:ARG:HB2	2.24	0.68
1:A:313:ILE:HA	1:A:316:LYS:HE2	1.75	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:569:HIS:CG	1:A:570:PRO:HD2	2.29	0.68
1:A:916:PHE:CD1	1:A:919:MET:HB3	2.29	0.68
2:B:338:VAL:C	2:B:339:ASN:HD22	1.97	0.68
2:C:384:ALA:HB3	2:C:440:THR:HG22	1.76	0.68
1:A:739:TYR:C	1:A:741:ASP:N	2.45	0.68
1:A:532:PRO:HB3	1:A:534:SER:HB3	1.76	0.68
1:D:1209:ARG:HH21	2:F:253:ASP:N	1.92	0.68
1:D:147:SER:HB3	1:D:425:ALA:HB2	1.74	0.68
1:D:241:PRO:O	1:D:244:LEU:CG	2.40	0.68
1:D:407:PHE:HD1	1:D:411:LEU:HD21	1.57	0.68
1:D:603:MET:HB3	1:D:605:LEU:HD21	1.76	0.68
1:D:901:VAL:HG23	1:D:902:LEU:N	2.08	0.68
2:F:395:LEU:HB3	2:F:443:VAL:HG21	1.75	0.68
2:F:472:LYS:HA	2:F:475:LEU:HD11	1.76	0.68
1:A:1026:ARG:CB	1:A:1029:GLN:NE2	2.57	0.68
1:A:1165:ALA:HB2	1:A:1173:LEU:HG	1.75	0.68
1:A:194:ALA:HA	1:A:265:LEU:CB	2.21	0.68
1:A:393:MET:HE3	1:A:396:CYS:SG	2.33	0.68
2:C:485:VAL:HG22	2:C:485:VAL:OXT	1.94	0.68
1:D:236:THR:OG1	1:D:237:SER:N	2.27	0.68
2:C:222:LYS:O	2:C:223:GLN:HB2	1.93	0.68
1:D:1141:VAL:HG23	1:D:1146:ARG:HH21	1.58	0.68
1:D:371:LYS:HD3	1:D:392:LEU:CD2	2.23	0.68
1:D:610:PHE:N	1:D:611:PRO:CD	2.51	0.68
1:D:660:CYS:HB2	1:D:745:PRO:CG	2.23	0.68
1:D:824:ALA:CA	1:D:827:ARG:HG3	2.24	0.68
1:D:831:TYR:HB2	1:D:837:TYR:OH	1.93	0.68
1:D:427:MET:O	1:D:846:THR:CG2	2.41	0.68
2:E:353:SER:CB	2:E:372:LEU:HD11	2.23	0.68
2:E:431:LYS:HG3	2:E:432:TYR:HD1	1.59	0.68
2:F:474:PHE:C	2:F:474:PHE:CD1	2.68	0.68
1:A:583:PRO:O	1:A:585:TRP:N	2.25	0.68
1:A:744:ILE:CG1	1:A:745:PRO:CD	2.71	0.68
2:C:387:VAL:HA	2:C:443:VAL:O	1.93	0.68
2:C:456:ARG:HB2	2:C:463:LYS:CB	2.24	0.68
2:C:72:ILE:HD12	2:C:76:ARG:HE	1.57	0.68
1:D:499:LYS:HZ2	1:D:499:LYS:HB2	1.59	0.68
1:D:127:LEU:O	1:D:130:LEU:CD2	2.42	0.67
1:D:179:GLY:CA	1:D:182:GLY:N	2.57	0.67
1:D:312:TRP:CH2	1:D:346:ASP:O	2.47	0.67
1:D:600:PRO:HA	1:D:603:MET:HB2	1.77	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:726:LYS:O	1:D:727:ASP:CB	2.41	0.67
2:E:374:LEU:N	2:E:458:ARG:NH2	2.42	0.67
2:F:393:LEU:HD23	2:F:396:ARG:HG2	1.76	0.67
1:A:1005:VAL:CA	1:A:1008:LEU:HD13	2.18	0.67
1:A:359:ALA:HB2	1:A:407:PHE:CE2	2.26	0.67
1:A:382:MET:SD	1:A:385:ILE:CG2	2.82	0.67
1:A:76:ILE:CG1	1:A:911:HIS:NE2	2.57	0.67
2:C:306:LEU:O	2:C:306:LEU:HD12	1.94	0.67
2:C:439:PHE:CD1	2:C:458:ARG:CG	2.77	0.67
1:D:104:VAL:O	1:D:108:GLN:HG3	1.93	0.67
2:B:133:HIS:NE2	2:C:231:ILE:CG2	2.57	0.67
2:B:395:LEU:HD11	2:B:443:VAL:HG12	1.76	0.67
1:D:177:ARG:HG2	1:D:178:TYR:H	1.58	0.67
1:D:316:LYS:CD	1:D:348:LEU:HD23	2.24	0.67
1:D:642:SER:N	1:D:645:VAL:HG21	2.08	0.67
1:D:791:ALA:C	1:D:795:ASN:HD21	1.96	0.67
1:D:847:ALA:HA	1:D:852:ARG:CA	2.24	0.67
2:E:254:PHE:HA	2:E:257:ARG:HE	1.59	0.67
2:F:387:VAL:CG2	2:F:417:TYR:CA	2.65	0.67
2:E:129:ASP:CG	2:F:98:GLY:HA3	2.14	0.67
1:A:1117:LYS:O	1:A:1121:GLU:HG3	1.94	0.67
1:A:255:SER:HB2	1:A:257:THR:O	1.94	0.67
1:A:363:ARG:HH11	1:A:406:VAL:HA	1.58	0.67
1:A:472:GLN:HB3	2:C:461:THR:HG23	1.76	0.67
1:A:869:ARG:O	1:A:872:SER:HB2	1.94	0.67
2:B:213:VAL:CA	2:B:235:THR:O	2.38	0.67
2:E:218:VAL:C	2:E:230:SER:HG	1.96	0.67
1:D:119:PRO:C	1:D:121:PRO:HD2	2.15	0.67
1:D:177:ARG:H	1:D:218:SER:CB	2.06	0.67
1:D:273:PHE:HA	1:D:276:ALA:HB2	1.76	0.67
1:D:636:THR:HG22	1:D:637:GLY:N	2.10	0.67
1:D:770:PHE:CE1	1:D:968:GLN:HB3	2.29	0.67
2:F:134:LYS:HB3	2:F:180:LYS:CE	2.25	0.67
2:F:351:TYR:CD1	2:F:355:GLN:HG2	2.29	0.67
2:F:360:SER:CB	2:F:364:LYS:HD2	2.24	0.67
2:F:383:VAL:HG22	2:F:384:ALA:N	2.09	0.67
1:A:1043:VAL:HG22	1:A:1046:GLU:CB	2.23	0.67
1:A:1047:ARG:CZ	1:A:1050:LYS:HD2	2.24	0.67
1:A:955:TYR:CD2	1:A:1102:GLN:NE2	2.62	0.67
1:A:1236:GLN:H	1:A:1236:GLN:HE21	1.40	0.67
1:A:202:CYS:HB2	1:A:204:ALA:H	1.59	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:265:LEU:HD12	1:A:265:LEU:O	1.94	0.67
1:A:276:ALA:O	1:A:280:GLU:OE1	2.11	0.67
2:B:69:LEU:HD21	2:B:351:TYR:CZ	2.30	0.67
2:E:270:PRO:O	2:E:273:PHE:HD2	1.77	0.67
1:D:498:LYS:NZ	1:D:521:PRO:HB3	2.10	0.67
1:D:273:PHE:HA	1:D:276:ALA:CB	2.24	0.67
1:D:483:PRO:O	1:D:486:TRP:N	2.26	0.67
1:D:548:CYS:C	1:D:551:LYS:CD	2.54	0.67
1:D:607:TRP:CB	1:D:779:LEU:HD23	2.19	0.67
1:D:847:ALA:HA	1:D:852:ARG:CB	2.25	0.67
1:D:112:LEU:HD23	1:D:917:GLY:O	1.94	0.67
2:E:129:ASP:OD2	2:F:98:GLY:HA3	1.94	0.67
1:A:80:SER:CB	1:A:125:LEU:HD21	2.24	0.67
1:A:873:GLU:OE1	1:A:875:LYS:HG3	1.94	0.67
2:C:389:ARG:CD	2:C:390:GLY:H	2.06	0.67
2:C:78:PHE:CD1	2:C:102:LEU:HD22	2.30	0.67
1:D:515:ILE:CG1	1:D:568:GLY:O	2.42	0.67
1:A:500:ALA:HB1	1:A:520:ALA:CB	2.24	0.67
2:B:475:LEU:C	2:B:479:ILE:HD13	2.15	0.67
2:B:217:PRO:C	2:B:231:ILE:HA	2.14	0.67
2:E:272:ASN:HA	2:E:292:ASN:HD22	1.55	0.67
1:D:1224:ILE:HD13	1:D:1224:ILE:C	2.15	0.67
1:D:272:SER:HB3	1:D:843:GLN:NE2	2.10	0.67
1:D:792:LEU:HA	1:D:795:ASN:ND2	2.09	0.67
1:D:801:TRP:HH2	1:D:805:HIS:ND1	1.93	0.67
1:D:995:TYR:CG	1:D:995:TYR:O	2.47	0.67
2:E:335:VAL:HG22	2:E:336:LEU:N	2.10	0.67
2:F:208:LEU:CD1	2:F:240:VAL:CB	2.62	0.67
2:F:213:VAL:HG23	2:F:235:THR:OG1	1.95	0.67
2:F:314:MET:CE	2:F:315:TYR:HE2	2.07	0.67
2:F:387:VAL:HA	2:F:443:VAL:CG2	2.05	0.67
2:F:467:HIS:O	2:F:471:LEU:HD11	1.93	0.67
1:A:278:ILE:HD13	1:A:291:PHE:CD1	2.28	0.67
1:A:711:ALA:O	1:A:712:VAL:CG2	2.42	0.67
2:B:116:THR:O	2:B:121:PHE:HD2	1.77	0.67
2:B:125:VAL:HG22	2:B:207:GLY:H	1.57	0.67
2:B:336:LEU:HD21	2:B:338:VAL:CG2	2.19	0.67
2:B:426:GLU:N	2:B:426:GLU:OE1	2.27	0.67
2:C:309:HIS:HB3	2:C:310:GLU:OE1	1.94	0.67
2:C:326:ASP:CG	2:C:327:GLY:N	2.46	0.67
1:D:512:LYS:HG2	1:D:570:PRO:HG3	1.77	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:503:VAL:HG21	1:A:521:PRO:HA	1.74	0.67
1:A:564:GLN:CA	1:A:564:GLN:HE21	2.08	0.67
2:E:217:PRO:HA	2:E:231:ILE:HG23	1.77	0.67
1:D:1010:LEU:N	1:D:1011:PRO:HD2	2.08	0.67
1:D:297:MET:SD	1:D:411:LEU:HD13	2.35	0.67
1:D:439:GLN:HG2	1:D:835:GLY:HA3	1.76	0.67
1:D:461:LYS:HA	1:D:464:MET:CE	2.24	0.67
1:D:591:LEU:HD12	1:D:591:LEU:O	1.94	0.67
1:D:613:HIS:ND1	1:D:717:LEU:HD22	2.09	0.67
2:E:266:PHE:O	2:E:349:TYR:HE1	1.77	0.67
2:F:127:PRO:HD2	2:F:128:VAL:H	1.56	0.67
2:F:456:ARG:HG2	2:F:457:SER:N	2.10	0.67
1:A:1040:LYS:O	1:A:1044:VAL:HB	1.94	0.67
1:A:769:ASP:OD2	1:A:770:PHE:CD2	2.47	0.67
1:A:798:ILE:HD11	1:A:802:ARG:HE	1.59	0.67
1:A:296:SER:CB	1:A:846:THR:O	2.43	0.67
2:B:343:ASP:HB3	2:B:344:ARG:NE	2.08	0.67
2:B:69:LEU:HD12	2:B:70:LEU:N	2.09	0.67
2:C:306:LEU:H	2:C:335:VAL:HG11	1.56	0.67
1:D:1006:ARG:NH2	1:D:1017:GLY:HA3	2.09	0.67
1:D:904:ASP:CG	1:D:915:ALA:HB3	2.15	0.67
2:E:105:GLU:O	2:E:108:LYS:HG2	1.94	0.67
2:E:244:PRO:HG3	2:E:326:ASP:OD2	1.95	0.67
2:F:372:LEU:HD21	2:F:435:MET:O	1.95	0.67
1:A:1030:ARG:CD	1:A:1040:LYS:NZ	2.58	0.67
1:A:169:TRP:O	1:A:170:ALA:CB	2.43	0.67
1:A:414:PHE:HA	1:A:417:ARG:HB2	1.75	0.67
1:A:720:THR:HG23	1:A:721:ALA:N	2.08	0.67
2:B:343:ASP:CB	2:B:344:ARG:HE	2.07	0.67
1:D:516:GLU:HA	1:D:565:HIS:HE1	1.57	0.67
1:D:206:GLY:CA	1:D:245:ILE:HG22	2.24	0.67
1:D:550:GLN:N	1:D:551:LYS:HE3	2.10	0.67
1:D:652:ILE:O	1:D:656:TYR:HD2	1.76	0.67
1:D:660:CYS:SG	1:D:661:LEU:N	2.68	0.67
2:E:448:LEU:O	2:E:450:ASN:N	2.27	0.67
2:F:240:VAL:O	2:F:336:LEU:HD13	1.95	0.67
2:F:344:ARG:CA	2:F:347:LEU:HD13	2.25	0.67
2:F:454:HIS:NE2	2:F:472:LYS:HD3	2.09	0.67
2:F:79:LEU:HA	2:F:99:PHE:HA	1.77	0.67
1:A:245:ILE:CD1	1:A:246:PRO:HD3	2.24	0.67
1:A:308:GLN:HB3	1:A:1090:GLU:HA	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:386:ARG:HA	1:A:389:PHE:HB3	1.76	0.67
1:A:473:LEU:HD23	1:A:473:LEU:N	2.09	0.67
1:A:904:ASP:CB	1:A:910:MET:O	2.42	0.67
2:C:193:TYR:HB2	2:C:242:PHE:CE2	2.29	0.67
1:D:1029:GLN:HE21	1:D:1030:ARG:HH21	1.43	0.67
1:D:285:GLN:HE22	1:D:1128:ARG:CG	2.07	0.67
1:D:1154:GLN:O	1:D:1157:ASN:HB2	1.94	0.67
1:D:245:ILE:HG13	1:D:246:PRO:HD3	1.75	0.67
1:D:401:TRP:HD1	1:D:404:HIS:CD2	2.13	0.67
1:D:647:CYS:SG	1:D:648:PRO:HD2	2.35	0.67
1:D:853:ARG:NH1	1:D:1105:ALA:HB3	2.09	0.67
2:E:257:ARG:O	2:E:261:GLN:HG3	1.95	0.67
2:E:370:LYS:HG3	2:E:434:GLU:OE2	1.95	0.67
2:F:211:ILE:HG23	2:F:236:GLU:O	1.95	0.67
2:F:471:LEU:CA	2:F:474:PHE:HD2	2.08	0.67
1:A:1046:GLU:OE1	1:A:1049:TRP:CB	2.43	0.67
1:A:276:ALA:HA	1:A:279:ARG:CB	2.24	0.67
1:A:439:GLN:NE2	1:A:837:TYR:HE2	1.92	0.67
2:B:323:HIS:HB2	2:B:332:VAL:HA	1.73	0.67
2:B:93:SER:C	2:B:95:CYS:H	1.98	0.67
2:C:239:LEU:H	2:C:338:VAL:CG1	2.03	0.67
2:C:455:LEU:HB3	2:C:456:ARG:CZ	2.25	0.67
1:A:941:ILE:HG23	1:A:942:SER:N	2.10	0.67
1:A:973:LEU:HB3	1:A:977:GLU:HB2	1.76	0.67
1:D:364:VAL:O	1:D:364:VAL:CG1	2.43	0.67
1:D:466:LEU:HD13	1:D:466:LEU:N	2.10	0.67
1:D:847:ALA:CA	1:D:852:ARG:HA	2.24	0.67
1:D:902:LEU:HD11	1:D:1168:LEU:HD22	1.75	0.67
2:E:445:GLU:OE2	2:E:446:THR:N	2.26	0.67
2:F:277:ASP:CG	2:F:287:ASN:HB3	2.15	0.67
2:F:300:ILE:HG23	2:F:342:LEU:HD11	1.75	0.67
2:F:368:HIS:O	2:F:368:HIS:ND1	2.28	0.67
2:F:391:PRO:HD2	2:F:394:GLU:HG2	1.77	0.67
1:A:996:ARG:HH11	1:A:1000:GLU:HB3	1.57	0.67
1:A:426:GLY:HA2	1:A:1117:LYS:NZ	2.10	0.67
1:A:297:MET:O	1:A:300:ALA:HB3	1.95	0.67
1:A:953:ARG:HD3	1:A:986:TYR:CE2	2.30	0.67
1:D:194:ALA:CB	1:D:265:LEU:HB2	2.25	0.66
1:D:249:VAL:CG2	1:D:250:PRO:CD	2.73	0.66
1:D:259:ARG:HA	1:D:259:ARG:NE	2.05	0.66
2:E:410:GLY:C	2:E:411:ILE:HD12	2.15	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:241:TRP:CD1	2:F:243:THR:CG2	2.78	0.66
1:A:1000:GLU:O	1:A:1004:LEU:HB2	1.95	0.66
1:A:1131:ILE:HG23	1:A:1132:SER:N	2.07	0.66
1:A:1165:ALA:CB	1:A:1173:LEU:HG	2.25	0.66
1:A:270:ASN:C	1:A:271:VAL:HG23	2.16	0.66
1:A:660:CYS:HB2	1:A:745:PRO:CG	2.25	0.66
1:A:859:TRP:HZ3	1:A:878:VAL:HG13	1.60	0.66
1:A:875:LYS:O	1:A:878:VAL:CG2	2.39	0.66
1:A:893:SER:HB2	1:A:896:LEU:CD2	2.25	0.66
1:A:996:ARG:HD2	1:A:1000:GLU:O	1.94	0.66
2:B:101:PRO:N	2:C:128:VAL:HG21	2.10	0.66
2:B:183:GLU:O	2:B:214:CYS:SG	2.53	0.66
2:C:129:ASP:OD1	2:C:131:LEU:CD2	2.43	0.66
2:C:204:LEU:HD21	2:C:206:TYR:CG	2.31	0.66
2:C:305:ASN:HA	2:C:335:VAL:HG11	1.76	0.66
1:D:504:LYS:O	1:D:504:LYS:HG3	1.95	0.66
1:D:564:GLN:C	1:D:566:LEU:O	2.32	0.66
1:D:1009:ASN:C	1:D:1010:LEU:HG	2.13	0.66
1:D:1015:THR:OG1	1:D:1022:LEU:HD23	1.94	0.66
1:D:270:ASN:HA	1:D:294:THR:OG1	1.95	0.66
1:D:270:ASN:CA	1:D:294:THR:CG2	2.71	0.66
1:D:298:HIS:CD2	1:D:411:LEU:HD23	2.29	0.66
1:D:417:ARG:C	1:D:418:CYS:SG	2.74	0.66
1:D:822:PRO:O	1:D:826:ILE:HD13	1.94	0.66
1:D:923:GLY:C	1:D:931:LEU:HD11	2.15	0.66
2:E:72:ILE:HD11	2:E:370:LYS:CE	2.25	0.66
2:F:298:GLU:OE1	2:F:344:ARG:NH2	2.28	0.66
1:A:110:HIS:O	1:A:112:LEU:HD12	1.94	0.66
1:A:253:ALA:HB2	1:A:280:GLU:C	2.15	0.66
1:A:612:LEU:HD21	1:A:614:TYR:O	1.94	0.66
2:C:209:ALA:CB	2:C:238:SER:O	2.42	0.66
2:C:236:GLU:HA	2:C:341:ASP:CG	2.16	0.66
2:C:353:SER:HB2	2:C:373:LYS:H	1.59	0.66
1:A:633:LYS:HD2	1:A:634:LEU:N	2.11	0.66
2:E:394:GLU:O	2:E:397:GLN:HB2	1.94	0.66
1:D:224:CYS:C	1:D:228:LEU:HD23	2.15	0.66
1:D:269:HIS:CE1	1:D:291:PHE:CA	2.78	0.66
1:D:307:PHE:O	1:D:311:LEU:HD12	1.96	0.66
2:E:76:ARG:HE	2:E:431:LYS:HE3	1.60	0.66
2:F:125:VAL:CG2	2:F:126:PHE:N	2.55	0.66
2:F:303:LEU:CB	2:F:337:SER:OG	2.43	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:314:MET:HE3	2:F:315:TYR:HE2	1.59	0.66
1:A:1016:GLU:OE2	1:A:1022:LEU:CD2	2.42	0.66
1:A:1113:LEU:HB2	1:A:1117:LYS:HZ2	1.61	0.66
1:A:1129:PHE:CD2	1:A:1138:ARG:O	2.48	0.66
1:A:241:PRO:CA	1:A:244:LEU:HB2	2.23	0.66
1:A:355:SER:HA	1:A:407:PHE:HE1	1.61	0.66
1:A:609:GLY:H	1:A:610:PHE:HD2	1.44	0.66
1:A:74:LEU:C	1:A:76:ILE:N	2.42	0.66
1:A:756:ASP:HB3	1:A:1057:MET:CG	2.12	0.66
2:B:113:GLU:O	2:B:262:TRP:HZ2	1.78	0.66
2:C:245:PRO:HG2	2:C:246:ARG:HD3	1.77	0.66
2:C:342:LEU:CB	2:C:344:ARG:HD3	2.25	0.66
2:C:358:GLU:HA	2:C:361:PHE:HB2	1.78	0.66
2:C:403:PHE:HD1	2:C:417:TYR:OH	1.78	0.66
1:A:1037:GLN:C	1:A:1039:LYS:N	2.46	0.66
2:F:367:LEU:O	2:F:369:ARG:HD2	1.95	0.66
1:D:1049:TRP:C	1:D:1052:GLY:H	1.99	0.66
1:D:140:ARG:CG	1:D:140:ARG:HH11	2.08	0.66
1:D:768:LYS:O	1:D:772:PRO:HD2	1.95	0.66
1:D:949:PHE:O	1:D:953:ARG:N	2.28	0.66
1:D:1209:ARG:HG3	2:F:249:ASN:CB	2.26	0.66
1:D:1208:ARG:NH2	2:F:285:LYS:CD	2.50	0.66
2:F:325:ARG:HG3	2:F:325:ARG:NH1	1.99	0.66
1:A:402:ALA:O	1:A:406:VAL:N	2.22	0.66
2:B:260:LEU:O	2:B:264:ARG:CG	2.43	0.66
2:B:67:GLU:HB3	2:B:70:LEU:HD13	1.75	0.66
2:C:372:LEU:HD12	2:C:373:LYS:N	2.11	0.66
1:A:552:LEU:HD22	2:C:448:LEU:CD2	2.25	0.66
1:D:857:PRO:O	1:D:1062:GLU:OE2	2.13	0.66
1:D:271:VAL:HA	1:D:293:ASP:OD2	1.96	0.66
1:D:307:PHE:CD1	1:D:307:PHE:N	2.63	0.66
1:D:408:GLN:O	1:D:411:LEU:HD12	1.95	0.66
1:D:550:GLN:H	1:D:551:LYS:NZ	1.92	0.66
1:D:914:THR:OG1	1:D:999:ASP:CG	2.34	0.66
1:D:923:GLY:CA	1:D:931:LEU:HD11	2.26	0.66
2:E:109:ASN:HD22	2:E:378:LEU:C	1.99	0.66
2:F:209:ALA:HB2	2:F:239:LEU:HA	1.76	0.66
2:F:445:GLU:C	2:F:447:THR:N	2.43	0.66
1:A:1019:TRP:O	1:A:1020:ILE:HG22	1.94	0.66
1:A:1131:ILE:HG21	1:A:1138:ARG:H	1.55	0.66
1:A:213:VAL:O	1:A:401:TRP:CZ2	2.48	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:515:ILE:CG2	1:A:565:HIS:HA	2.15	0.66
1:A:744:ILE:HG12	1:A:745:PRO:CD	2.24	0.66
1:A:962:ALA:CB	1:A:986:TYR:OH	2.39	0.66
2:C:129:ASP:O	2:C:130:ALA:HB3	1.94	0.66
2:C:208:LEU:CD1	2:C:240:VAL:CB	2.68	0.66
2:C:335:VAL:CG2	2:C:336:LEU:N	2.57	0.66
2:C:435:MET:SD	2:C:435:MET:O	2.53	0.66
2:C:454:HIS:CA	2:C:468:ILE:HD12	2.23	0.66
1:D:240:SER:HB2	1:D:241:PRO:HD3	1.77	0.66
1:D:374:ARG:HD3	1:D:392:LEU:CD2	2.23	0.66
1:D:552:LEU:O	2:F:452:LEU:HD21	1.96	0.66
2:E:402:LEU:O	2:E:406:LEU:HD23	1.94	0.66
2:E:132:HIS:HB2	2:F:233:GLU:CD	2.16	0.66
2:F:353:SER:HB2	2:F:373:LYS:N	2.08	0.66
2:F:368:HIS:NE2	2:F:370:LYS:CD	2.58	0.66
2:F:81:GLY:HA3	2:F:97:PRO:CG	2.25	0.66
1:A:308:GLN:HB3	1:A:1090:GLU:CB	2.25	0.66
1:A:559:LEU:HD23	1:A:560:PRO:CD	2.26	0.66
2:B:323:HIS:HA	2:B:333:PRO:CD	2.25	0.66
2:C:288:LYS:HB2	2:C:290:TYR:OH	1.96	0.66
1:D:1175:GLN:O	1:D:1177:VAL:N	2.28	0.66
1:D:418:CYS:N	1:D:419:PRO:HD3	2.10	0.66
1:D:577:CYS:C	1:D:579:ARG:N	2.36	0.66
1:D:870:VAL:HG13	1:D:871:GLY:N	2.10	0.66
2:E:108:LYS:NZ	2:E:108:LYS:HB2	2.11	0.66
2:F:242:PHE:HB3	2:F:333:PRO:CB	2.18	0.66
1:D:1209:ARG:NH2	2:F:253:ASP:CA	2.50	0.66
2:F:344:ARG:CA	2:F:347:LEU:HD22	2.25	0.66
2:F:413:VAL:HG12	2:F:414:TRP:N	2.10	0.66
2:F:469:SER:O	2:F:472:LYS:HB2	1.95	0.66
1:A:515:ILE:N	1:A:515:ILE:HD12	2.11	0.66
1:A:597:ARG:O	1:A:598:VAL:HB	1.96	0.66
1:A:652:ILE:O	1:A:655:LEU:HG	1.95	0.66
1:A:436:PRO:HD2	1:A:879:GLN:CD	2.15	0.66
2:B:235:THR:H	2:B:343:ASP:HB2	1.60	0.66
2:B:425:LEU:HD23	2:B:425:LEU:O	1.96	0.66
1:D:1165:ALA:CB	1:D:1173:LEU:CD1	2.72	0.66
1:D:176:THR:N	1:D:185:VAL:CG1	2.59	0.66
1:D:386:ARG:HA	1:D:389:PHE:CD2	2.31	0.66
1:D:657:ARG:HA	1:D:660:CYS:HB3	1.77	0.66
1:D:771:LEU:HD12	1:D:771:LEU:N	2.11	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:802:ARG:CG	1:D:803:ASN:OD1	2.44	0.66
1:D:826:ILE:HG12	1:D:827:ARG:N	2.10	0.66
1:D:87:ILE:HG22	1:D:88:PHE:H	1.61	0.66
2:E:424:SER:C	2:E:428:LEU:HD11	2.16	0.66
2:E:380:PRO:HG3	2:E:438:LEU:HD13	1.77	0.66
2:E:475:LEU:HB3	2:E:479:ILE:HD11	1.78	0.66
2:F:415:PRO:HB3	2:F:417:TYR:HE2	1.57	0.66
1:A:1002:GLU:O	1:A:1006:ARG:N	2.26	0.66
1:A:717:LEU:O	1:A:721:ALA:CB	2.44	0.66
1:A:769:ASP:O	1:A:772:PRO:HD2	1.96	0.66
1:A:439:GLN:HG3	1:A:836:LEU:H	1.59	0.66
1:A:858:THR:HB	1:A:859:TRP:CD1	2.31	0.66
2:B:204:LEU:HD22	2:B:205:PRO:HB3	1.77	0.66
2:B:323:HIS:CB	2:B:332:VAL:CA	2.71	0.66
2:C:302:THR:C	2:C:303:LEU:HD12	2.16	0.66
2:C:395:LEU:CD1	2:C:444:THR:HA	2.25	0.66
1:D:652:ILE:CA	1:D:655:LEU:HG	2.18	0.66
2:F:306:LEU:HD11	2:F:310:GLU:OE1	1.96	0.66
1:A:1006:ARG:O	1:A:1010:LEU:HD12	1.95	0.66
1:A:1007:GLU:O	1:A:1011:PRO:CD	2.44	0.66
1:A:355:SER:O	1:A:358:LEU:HG	1.96	0.66
2:C:439:PHE:HD1	2:C:458:ARG:CG	2.07	0.66
1:A:522:GLY:O	1:A:524:PRO:HD3	1.95	0.66
1:D:194:ALA:CA	1:D:265:LEU:HB2	2.25	0.66
1:D:457:ARG:HA	1:D:460:LYS:CD	2.26	0.66
1:D:580:LEU:HD21	1:D:593:SER:OG	1.96	0.66
1:D:601:LYS:HD2	1:D:616:GLU:CG	2.26	0.66
1:D:658:LYS:HE2	1:D:713:PRO:HB2	1.76	0.66
1:D:719:LEU:HD12	1:D:720:THR:OG1	1.95	0.66
1:D:736:ASN:HB3	1:D:748:TRP:O	1.96	0.66
1:D:737:GLY:N	1:D:750:PHE:CE1	2.64	0.66
1:D:798:ILE:CG2	1:D:869:ARG:NH2	2.56	0.66
1:D:873:GLU:HB3	1:D:1200:PRO:HD2	1.78	0.66
1:A:1047:ARG:NH1	1:A:1050:LYS:HB2	2.11	0.66
1:A:885:THR:C	1:A:1146:ARG:HE	1.99	0.66
1:A:166:PRO:HG3	1:A:398:GLN:OE1	1.96	0.66
1:A:192:GLU:OE1	1:A:265:LEU:HD23	1.96	0.66
1:A:271:VAL:HG21	1:A:295:MET:HE3	1.77	0.66
1:A:298:HIS:HE2	1:A:351:SER:C	2.00	0.66
1:A:346:ASP:OD2	1:A:1040:LYS:HD2	1.97	0.66
1:A:732:TYR:CD1	1:A:755:LYS:NZ	2.62	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:830:ASP:C	1:A:831:TYR:HD1	2.00	0.66
1:A:989:THR:CA	1:A:998:SER:OG	2.44	0.66
2:B:70:LEU:HD22	2:B:88:ARG:NH1	2.05	0.66
2:B:74:GLN:O	2:B:77:HIS:N	2.29	0.66
2:B:453:ILE:H	2:B:453:ILE:HD13	1.60	0.66
2:F:317:GLY:CA	2:F:320:SER:CB	2.72	0.66
2:B:231:ILE:CD1	2:C:135:PRO:HG3	2.25	0.66
1:A:522:GLY:C	1:A:524:PRO:HD3	2.16	0.66
1:D:1015:THR:HG23	1:D:1026:ARG:CB	2.25	0.65
1:D:1131:ILE:CD1	1:D:1138:ARG:HB2	2.27	0.65
1:D:1165:ALA:HB2	1:D:1173:LEU:CD1	2.14	0.65
1:D:175:TRP:HB2	1:D:185:VAL:HG22	1.78	0.65
1:D:208:CYS:SG	1:D:244:LEU:CD1	2.84	0.65
1:D:292:LEU:HD23	1:D:297:MET:CE	2.26	0.65
1:D:307:PHE:HD1	1:D:307:PHE:N	1.93	0.65
1:D:481:GLU:O	1:D:482:ASP:CG	2.34	0.65
1:D:485:LEU:O	1:D:489:GLU:HG3	1.96	0.65
1:D:608:ASP:OD1	1:D:611:PRO:N	2.29	0.65
1:D:616:GLU:O	1:D:725:PRO:CD	2.37	0.65
1:D:618:HIS:CB	1:D:725:PRO:HG2	2.25	0.65
2:E:263:TRP:CZ3	2:E:300:ILE:HD12	2.31	0.65
2:E:79:LEU:C	2:E:100:GLY:HA3	2.17	0.65
2:F:347:LEU:HA	2:F:350:LEU:HD11	1.76	0.65
1:A:1207:GLU:CA	1:A:1210:TYR:HD1	2.09	0.65
1:A:432:VAL:HG12	1:A:432:VAL:O	1.95	0.65
1:A:606:THR:H	1:A:613:HIS:HA	1.60	0.65
1:A:659:HIS:ND1	1:A:716:PRO:HA	2.11	0.65
1:A:768:LYS:HG2	1:A:769:ASP:H	1.59	0.65
1:A:861:THR:HG22	1:A:861:THR:O	1.96	0.65
1:A:884:TYR:N	1:A:884:TYR:CD1	2.65	0.65
1:A:905:ALA:N	1:A:910:MET:HA	2.11	0.65
1:D:504:LYS:O	1:D:504:LYS:CG	2.44	0.65
1:D:664:GLY:HA2	1:D:667:GLN:HG3	1.77	0.65
1:D:1131:ILE:HD11	1:D:1138:ARG:HB2	1.78	0.65
1:D:1138:ARG:HG2	1:D:1138:ARG:NH1	2.07	0.65
1:D:78:MET:HE3	1:D:121:PRO:HB2	1.77	0.65
1:D:641:GLU:O	1:D:645:VAL:HG21	1.96	0.65
1:D:711:ALA:C	1:D:713:PRO:HD3	2.16	0.65
1:D:275:ARG:CZ	1:D:843:GLN:HG2	2.26	0.65
1:D:900:ALA:C	1:D:915:ALA:HB1	2.16	0.65
1:D:923:GLY:O	1:D:927:ARG:O	2.14	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:948:ILE:HG21	1:D:1071:ARG:CB	2.25	0.65
2:E:101:PRO:HD3	2:F:128:VAL:HG21	1.78	0.65
2:E:437:ILE:O	2:E:458:ARG:HD3	1.95	0.65
2:F:336:LEU:HD23	2:F:337:SER:C	2.16	0.65
2:F:69:LEU:HD23	2:F:351:TYR:CE1	2.29	0.65
2:F:395:LEU:HD23	2:F:448:LEU:HD11	1.77	0.65
1:A:271:VAL:CG2	1:A:295:MET:HE3	2.27	0.65
1:A:309:ARG:HH12	1:A:352:SER:HA	1.61	0.65
1:A:601:LYS:NZ	1:A:721:ALA:O	2.28	0.65
1:A:616:GLU:CD	1:A:724:GLY:H	2.00	0.65
1:A:588:GLY:CA	1:A:806:LYS:NZ	2.59	0.65
1:A:879:GLN:CA	1:A:879:GLN:HE21	2.08	0.65
2:B:215:PHE:HE2	2:B:233:GLU:HG2	1.60	0.65
2:B:407:LEU:HD21	2:C:120:VAL:CA	2.14	0.65
2:B:74:GLN:HB2	2:B:77:HIS:HD2	1.62	0.65
2:C:406:LEU:HA	2:C:409:ASN:HD22	1.57	0.65
2:C:447:THR:HG22	2:C:455:LEU:HD11	1.77	0.65
2:C:80:SER:OG	2:C:99:PHE:O	2.12	0.65
2:B:122:ARG:HD3	2:B:124:GLN:NE2	2.11	0.65
1:D:1008:LEU:O	1:D:1011:PRO:HD2	1.96	0.65
1:D:1141:VAL:HG23	1:D:1146:ARG:NH2	2.10	0.65
1:D:297:MET:CG	1:D:411:LEU:HD22	2.21	0.65
1:D:798:ILE:HG22	1:D:869:ARG:NH1	2.10	0.65
1:D:80:SER:HB2	1:D:125:LEU:CB	2.27	0.65
1:D:999:ASP:O	1:D:1001:GLY:N	2.30	0.65
1:A:303:GLY:O	1:A:1025:LEU:CD1	2.44	0.65
1:A:190:PRO:HB3	1:A:192:GLU:HG3	1.78	0.65
1:A:253:ALA:O	1:A:254:SER:HB3	1.95	0.65
1:A:298:HIS:NE2	1:A:358:LEU:HD22	2.11	0.65
1:A:474:LEU:CD1	1:A:474:LEU:N	2.59	0.65
1:A:601:LYS:HD2	1:A:616:GLU:HB3	1.78	0.65
1:A:612:LEU:HD23	1:A:620:TRP:CB	2.26	0.65
1:A:664:GLY:HA2	1:A:667:GLN:HE21	1.61	0.65
1:A:898:ILE:CG1	1:A:899:ALA:N	2.59	0.65
2:B:289:LEU:HB3	2:B:300:ILE:CG1	2.27	0.65
2:B:303:LEU:HD12	2:B:303:LEU:N	2.11	0.65
2:C:204:LEU:HG	2:C:206:TYR:HB3	1.77	0.65
2:C:270:PRO:HA	2:C:273:PHE:CD2	2.32	0.65
2:C:423:SER:HB2	2:C:427:GLN:OE1	1.96	0.65
2:C:439:PHE:HB3	2:C:458:ARG:HB2	1.78	0.65
1:D:1048:ALA:O	1:D:1052:GLY:N	2.29	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:267:VAL:O	1:D:269:HIS:CE1	2.49	0.65
1:D:458:GLU:N	1:D:460:LYS:HZ2	1.94	0.65
1:D:461:LYS:HA	1:D:464:MET:HE3	1.78	0.65
1:D:798:ILE:HA	1:D:869:ARG:NH2	2.12	0.65
2:E:130:ALA:N	2:E:210:GLN:HE22	1.94	0.65
2:E:400:GLN:O	2:E:403:PHE:N	2.30	0.65
2:F:197:LEU:HD11	2:F:322:LEU:CD2	2.27	0.65
2:F:294:PRO:HG3	2:F:355:GLN:NE2	2.11	0.65
2:F:342:LEU:CB	2:F:344:ARG:HB2	2.27	0.65
1:A:1026:ARG:HB3	1:A:1029:GLN:NE2	2.10	0.65
1:A:1200:PRO:C	1:A:1202:ASN:N	2.42	0.65
1:A:213:VAL:HG21	1:A:397:ALA:HA	1.78	0.65
1:A:498:LYS:HE2	1:A:499:LYS:N	2.08	0.65
1:A:612:LEU:CD2	1:A:621:GLY:N	2.50	0.65
2:B:269:SER:O	2:B:271:SER:N	2.27	0.65
2:C:209:ALA:HB2	2:C:239:LEU:HA	1.77	0.65
2:C:353:SER:CB	2:C:373:LYS:O	2.44	0.65
1:A:553:LYS:HD2	2:C:468:ILE:HG22	1.79	0.65
2:B:229:LYS:HD2	2:B:229:LYS:N	2.10	0.65
1:D:955:TYR:HB3	1:D:1098:ASN:CG	2.16	0.65
1:D:275:ARG:HH21	1:D:842:PRO:CA	2.09	0.65
1:D:279:ARG:HB3	1:D:841:LEU:CB	2.24	0.65
1:D:308:GLN:NE2	1:D:1027:LYS:NZ	2.44	0.65
1:D:407:PHE:CD1	1:D:407:PHE:C	2.70	0.65
1:D:721:ALA:O	1:D:722:ARG:HB2	1.96	0.65
1:D:732:TYR:HD2	1:D:732:TYR:N	1.92	0.65
1:D:783:PRO:HG2	1:D:784:GLY:H	1.61	0.65
2:E:283:GLY:HA3	2:E:306:LEU:CA	2.24	0.65
2:E:323:HIS:O	2:E:324:GLY:C	2.34	0.65
2:F:431:LYS:HZ1	2:F:435:MET:HG2	1.60	0.65
1:A:1060:LYS:CD	1:A:1073:PRO:HB2	2.25	0.65
1:A:1133:ILE:HD13	1:A:1134:HIS:HB2	1.78	0.65
1:A:1164:PHE:HD2	1:A:1168:LEU:HD11	1.60	0.65
1:A:556:THR:CG2	2:C:452:LEU:CB	2.75	0.65
1:A:601:LYS:HE2	1:A:616:GLU:OE2	1.96	0.65
1:A:624:VAL:O	1:A:745:PRO:O	2.15	0.65
1:A:87:ILE:CG2	1:A:1166:TYR:OH	2.44	0.65
2:C:200:VAL:HG12	2:C:200:VAL:O	1.95	0.65
2:C:387:VAL:HG12	2:C:443:VAL:HB	1.79	0.65
2:E:417:TYR:CD1	2:E:418:LEU:HG	2.31	0.65
1:D:177:ARG:HE	1:D:216:SER:H	1.43	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:203:LEU:O	1:D:204:ALA:C	2.35	0.65
1:D:313:ILE:HG13	1:D:314:ALA:H	1.62	0.65
1:D:371:LYS:HE2	1:D:396:CYS:SG	2.37	0.65
1:D:546:ARG:C	1:D:551:LYS:NZ	2.49	0.65
1:D:741:ASP:O	1:D:742:VAL:HB	1.95	0.65
1:D:768:LYS:CA	1:D:772:PRO:HG3	2.26	0.65
1:D:780:GLN:OE1	1:D:780:GLN:CA	2.45	0.65
1:D:776:ASP:CB	1:D:796:LYS:HZ3	2.10	0.65
1:D:798:ILE:N	1:D:869:ARG:NH1	2.45	0.65
1:D:929:THR:O	1:D:933:SER:HB3	1.96	0.65
1:D:983:GLN:HE21	1:D:983:GLN:HA	1.62	0.65
2:E:372:LEU:O	2:E:458:ARG:HD2	1.96	0.65
1:A:136:ASP:O	1:A:140:ARG:HG2	1.97	0.65
1:A:211:LEU:HG	1:A:221:TYR:CB	2.26	0.65
1:A:305:SER:HB2	1:A:308:GLN:HG3	1.77	0.65
1:A:660:CYS:SG	1:A:745:PRO:CG	2.83	0.65
1:A:862:ALA:HB1	1:A:875:LYS:HE3	1.79	0.65
2:B:306:LEU:O	2:B:308:ASP:N	2.29	0.65
2:C:204:LEU:CD2	2:C:206:TYR:CG	2.80	0.65
1:D:668:LEU:O	1:D:671:GLN:CB	2.35	0.65
1:D:876:ALA:CA	1:D:1189:LEU:HB2	2.23	0.65
1:D:177:ARG:HB3	1:D:218:SER:CB	2.22	0.65
1:D:175:TRP:CZ2	1:D:183:GLU:HG3	2.31	0.65
1:D:301:ILE:O	1:D:1024:ASP:HB2	1.97	0.65
1:D:406:VAL:HG22	1:D:407:PHE:N	2.10	0.65
1:D:913:CYS:O	1:D:917:GLY:CA	2.45	0.65
1:D:996:ARG:N	1:D:996:ARG:HD3	2.10	0.65
2:E:200:VAL:O	2:F:418:LEU:HD21	1.96	0.65
2:F:270:PRO:HA	2:F:273:PHE:CD2	2.32	0.65
2:F:375:HIS:HB3	2:F:378:LEU:HD22	1.77	0.65
1:A:392:LEU:O	1:A:395:TYR:N	2.24	0.65
1:A:479:TYR:O	1:A:480:LYS:CG	2.40	0.65
1:A:966:LEU:HD12	1:A:967:MET:N	2.12	0.65
1:A:784:GLY:HA3	2:C:363:ARG:HG2	1.78	0.65
1:D:550:GLN:N	1:D:551:LYS:HZ2	1.95	0.65
1:D:576:LEU:CG	1:D:577:CYS:N	2.58	0.65
1:D:804:ALA:HB1	1:D:808:ILE:HD11	1.78	0.65
1:D:957:ALA:HB3	1:D:961:PHE:CE1	2.32	0.65
2:E:437:ILE:HG22	2:E:439:PHE:N	2.12	0.65
2:F:311:LEU:HA	2:F:314:MET:HB2	1.78	0.65
2:F:385:LEU:O	2:F:416:GLY:CA	2.43	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:907:PHE:CZ	1:A:1001:GLY:N	2.65	0.65
1:A:1133:ILE:CG2	1:A:1136:GLU:HB2	2.26	0.65
1:A:1236:GLN:N	1:A:1236:GLN:NE2	2.41	0.65
1:A:301:ILE:HD13	1:A:414:PHE:CB	2.27	0.65
1:A:417:ARG:NH1	1:A:417:ARG:HG2	2.10	0.65
1:A:577:CYS:SG	1:A:578:PRO:HD3	2.37	0.65
1:A:851:THR:CG2	1:A:853:ARG:HB2	2.27	0.65
2:B:215:PHE:CE2	2:B:233:GLU:HG2	2.32	0.65
2:B:67:GLU:CB	2:B:70:LEU:HB2	2.24	0.65
1:D:944:GLU:CD	1:D:1071:ARG:NH1	2.51	0.65
1:D:113:TRP:HA	1:D:116:PRO:HG2	1.77	0.65
1:D:256:PRO:HB3	1:D:281:GLN:CB	2.26	0.65
1:D:460:LYS:CA	1:D:460:LYS:HE2	2.19	0.65
1:D:616:GLU:C	1:D:617:ARG:HG3	2.16	0.65
1:D:618:HIS:HA	1:D:762:VAL:HG21	1.78	0.65
1:D:612:LEU:HG	1:D:622:TYR:N	2.10	0.65
2:E:240:VAL:HG12	2:E:242:PHE:CE2	2.31	0.65
2:F:290:TYR:HA	2:F:298:GLU:O	1.97	0.65
2:F:445:GLU:C	2:F:447:THR:H	1.98	0.65
1:A:1108:TYR:HA	1:A:1164:PHE:HZ	1.61	0.65
1:A:1190:ARG:CZ	1:A:1197:CYS:CB	2.70	0.65
1:A:196:VAL:CG1	1:A:269:HIS:CD2	2.78	0.65
1:A:271:VAL:C	1:A:275:ARG:HD3	2.18	0.65
1:A:813:VAL:HG11	1:A:815:TRP:CZ2	2.32	0.65
1:A:823:ARG:C	1:A:825:VAL:N	2.49	0.65
1:A:848:GLY:O	1:A:849:THR:HB	1.96	0.65
2:B:309:HIS:N	2:B:309:HIS:ND1	2.43	0.65
2:C:309:HIS:HA	2:C:312:LEU:HD12	1.78	0.65
2:C:403:PHE:HD2	2:C:403:PHE:C	2.00	0.65
2:C:439:PHE:CB	2:C:457:SER:O	2.44	0.65
2:C:474:PHE:C	2:C:474:PHE:CD1	2.70	0.65
1:D:502:LYS:HG2	1:D:506:GLU:OE1	1.97	0.65
2:B:217:PRO:HB2	2:B:231:ILE:HG22	1.79	0.65
1:D:1136:GLU:OE2	1:D:1191:LYS:HE3	1.96	0.65
1:D:719:LEU:CG	1:D:720:THR:H	2.09	0.65
1:D:897:TRP:CZ2	1:D:1173:LEU:HD22	2.32	0.65
1:D:112:LEU:HD23	1:D:920:THR:HB	1.78	0.65
2:F:242:PHE:HA	2:F:334:CYS:O	1.96	0.65
2:F:309:HIS:CA	2:F:312:LEU:HD12	2.25	0.65
1:A:1209:ARG:HH22	2:C:256:LEU:CD2	2.10	0.65
1:A:608:ASP:C	1:A:620:TRP:CH2	2.71	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:114:TRP:O	2:B:117:SER:HB3	1.97	0.65
2:B:324:GLY:HA3	2:B:325:ARG:NH2	2.12	0.65
2:C:305:ASN:CA	2:C:335:VAL:HG11	2.27	0.65
2:C:464:GLU:OE2	2:C:466:MET:HG2	1.97	0.65
2:E:194:VAL:HG13	2:E:195:ASN:H	1.59	0.65
2:B:293:PHE:HB2	2:B:297:LYS:HA	1.79	0.65
1:D:404:HIS:O	1:D:408:GLN:N	2.29	0.64
1:D:462:SER:O	1:D:466:LEU:HD22	1.97	0.64
1:D:481:GLU:O	1:D:482:ASP:OD1	2.15	0.64
1:D:608:ASP:O	1:D:610:PHE:N	2.30	0.64
1:D:658:LYS:HE3	1:D:713:PRO:CG	2.27	0.64
1:D:770:PHE:O	1:D:774:MET:CG	2.45	0.64
1:D:854:ALA:O	1:D:855:VAL:HG22	1.96	0.64
2:E:205:PRO:CG	2:E:241:TRP:HE1	2.10	0.64
2:F:238:SER:HA	2:F:338:VAL:CG1	2.24	0.64
1:A:1122:GLU:C	1:A:1123:PHE:CD1	2.70	0.64
1:A:1232:GLU:OE2	1:A:1234:ARG:NH2	2.30	0.64
1:A:228:LEU:HD23	1:A:229:VAL:H	1.62	0.64
1:A:417:ARG:CG	1:A:417:ARG:HH11	2.10	0.64
1:A:418:CYS:N	1:A:419:PRO:CD	2.60	0.64
1:A:493:GLN:O	1:A:494:GLU:CG	2.42	0.64
1:A:553:LYS:O	1:A:556:THR:HB	1.97	0.64
1:A:612:LEU:HD23	1:A:620:TRP:CA	2.28	0.64
1:A:612:LEU:HD23	1:A:620:TRP:HA	1.79	0.64
2:B:396:ARG:HG3	2:B:417:TYR:CB	2.25	0.64
2:C:129:ASP:OD1	2:C:131:LEU:CD1	2.44	0.64
2:C:431:LYS:CA	2:C:434:GLU:OE1	2.38	0.64
2:C:473:ASP:CA	2:C:476:ILE:HG23	2.24	0.64
2:E:418:LEU:HD13	2:F:122:ARG:HE	1.62	0.64
1:D:1014:ARG:CD	1:D:1026:ARG:HG3	2.27	0.64
1:D:142:LEU:HD22	1:D:1118:TRP:CD1	2.33	0.64
1:D:1141:VAL:HG21	1:D:1149:ALA:CB	2.27	0.64
1:D:278:ILE:CD1	1:D:291:PHE:HD1	2.10	0.64
1:D:301:ILE:O	1:D:1024:ASP:OD1	2.15	0.64
1:D:408:GLN:HA	1:D:411:LEU:HD11	1.78	0.64
1:D:591:LEU:CD1	1:D:594:LEU:CG	2.70	0.64
1:D:732:TYR:HB3	1:D:755:LYS:NZ	2.12	0.64
2:E:239:LEU:HD23	2:E:338:VAL:CG2	2.26	0.64
2:E:261:GLN:O	2:E:265:LYS:HB2	1.97	0.64
2:E:451:GLY:C	2:E:452:LEU:HD22	2.17	0.64
1:A:907:PHE:CE1	1:A:1001:GLY:HA3	2.31	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1145:ASP:CA	1:A:1148:ARG:HB2	2.28	0.64
1:A:1153:LEU:HD13	1:A:1157:ASN:ND2	2.13	0.64
1:A:444:TYR:CE2	1:A:874:LEU:HA	2.32	0.64
1:A:744:ILE:HB	1:A:745:PRO:CD	2.24	0.64
1:A:849:THR:HG22	1:A:851:THR:N	2.08	0.64
1:A:907:PHE:HZ	1:A:1001:GLY:H	1.45	0.64
2:B:341:ASP:O	2:B:344:ARG:HD2	1.97	0.64
2:B:99:PHE:HE1	2:B:343:ASP:OD1	1.81	0.64
2:C:242:PHE:HB3	2:C:333:PRO:CB	2.21	0.64
2:C:371:VAL:HG23	2:C:433:ASP:HB3	1.78	0.64
1:A:645:VAL:O	1:A:646:VAL:CG1	2.38	0.64
1:D:1026:ARG:CD	1:D:1026:ARG:N	2.60	0.64
1:D:608:ASP:HA	1:D:620:TRP:CZ3	2.33	0.64
2:E:253:ASP:O	2:E:256:LEU:HB3	1.96	0.64
2:F:240:VAL:C	2:F:336:LEU:HB3	2.18	0.64
2:F:237:ALA:O	2:F:339:ASN:O	2.16	0.64
1:D:472:GLN:HG3	2:F:460:THR:OG1	1.97	0.64
1:A:1033:ALA:O	1:A:1034:ARG:HB2	1.97	0.64
1:A:1131:ILE:HG21	1:A:1137:VAL:HA	1.75	0.64
1:A:1190:ARG:HD3	1:A:1197:CYS:SG	2.38	0.64
1:A:301:ILE:HD13	1:A:414:PHE:HB2	1.79	0.64
1:A:813:VAL:CG2	1:A:814:VAL:H	2.10	0.64
1:A:898:ILE:HD11	1:A:1101:VAL:CG1	2.27	0.64
2:B:311:LEU:CD1	2:B:322:LEU:HD13	2.27	0.64
2:C:205:PRO:O	2:C:241:TRP:NE1	2.27	0.64
2:C:357:THR:O	2:C:358:GLU:CG	2.44	0.64
1:A:472:GLN:CG	2:C:460:THR:CA	2.67	0.64
1:D:309:ARG:O	1:D:313:ILE:N	2.25	0.64
1:D:302:SER:CB	1:D:351:SER:CB	2.73	0.64
1:D:653:GLU:O	1:D:656:TYR:HB2	1.97	0.64
1:D:618:HIS:HB2	1:D:725:PRO:HG2	1.78	0.64
2:E:101:PRO:N	2:F:128:VAL:HG21	2.12	0.64
2:E:288:LYS:HB3	2:E:299:LEU:HD12	1.79	0.64
2:F:446:THR:O	2:F:450:ASN:N	2.30	0.64
1:A:213:VAL:HG11	1:A:397:ALA:HB1	1.80	0.64
1:A:282:TYR:OH	1:A:432:VAL:HG22	1.97	0.64
1:A:484:TRP:CE3	1:A:484:TRP:HA	2.32	0.64
1:A:729:GLN:HB2	1:A:731:SER:OG	1.98	0.64
1:A:886:LEU:HD22	1:A:1140:LEU:HG	1.78	0.64
1:A:755:LYS:HD3	1:A:964:ARG:HH11	1.58	0.64
2:B:396:ARG:HH11	2:B:417:TYR:HD2	1.39	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:374:LEU:N	2:B:458:ARG:NH2	2.38	0.64
2:C:386:ASP:O	2:C:387:VAL:HG13	1.96	0.64
2:C:228:VAL:HG23	2:C:228:VAL:O	1.98	0.64
1:D:1067:SER:HB2	1:D:1071:ARG:HB3	1.79	0.64
1:D:270:ASN:N	1:D:274:ASP:OD1	2.31	0.64
1:D:298:HIS:CE1	1:D:362:LEU:HD11	2.33	0.64
1:D:282:TYR:CE2	1:D:432:VAL:HG13	2.32	0.64
2:E:301:GLU:HG2	2:E:303:LEU:HD11	1.78	0.64
2:E:324:GLY:N	2:E:331:VAL:O	2.30	0.64
2:F:194:VAL:O	2:F:197:LEU:HD22	1.98	0.64
2:F:273:PHE:CE1	2:F:291:TYR:HD1	2.15	0.64
2:F:301:GLU:HG2	2:F:339:ASN:HB3	1.78	0.64
2:F:74:GLN:HG3	2:F:75:ARG:N	2.10	0.64
1:A:1043:VAL:HG13	1:A:1046:GLU:CB	2.10	0.64
1:A:1129:PHE:HB3	1:A:1139:TYR:HA	1.80	0.64
1:A:113:TRP:HA	1:A:116:PRO:HD3	1.79	0.64
1:A:1188:CYS:O	1:A:1189:LEU:HD22	1.97	0.64
1:A:276:ALA:C	1:A:279:ARG:H	2.01	0.64
1:A:274:ASP:C	1:A:276:ALA:N	2.48	0.64
1:A:608:ASP:HB3	1:A:779:LEU:HB2	1.78	0.64
1:A:812:MET:C	1:A:813:VAL:HG12	2.18	0.64
1:A:880:ALA:H	1:A:881:PRO:HD3	1.61	0.64
2:B:114:TRP:NE1	2:B:259:ARG:NE	2.45	0.64
2:C:194:VAL:O	2:C:197:LEU:HB3	1.97	0.64
2:C:241:TRP:HZ3	2:C:255:TRP:CE2	2.16	0.64
2:C:263:TRP:HZ2	2:C:301:GLU:OE2	1.81	0.64
1:A:553:LYS:HD3	2:C:468:ILE:HG21	1.80	0.64
2:C:475:LEU:HA	2:C:478:TYR:HB2	1.78	0.64
1:A:163:PRO:CB	1:A:164:PRO:CD	2.75	0.64
2:B:389:ARG:HB2	2:B:445:GLU:H	1.62	0.64
2:C:317:GLY:CA	2:C:320:SER:CB	2.74	0.64
2:E:133:HIS:CE1	2:F:231:ILE:HG22	2.32	0.64
1:D:1204:THR:O	1:D:1205:GLY:C	2.36	0.64
1:D:142:LEU:O	1:D:146:GLN:HG2	1.97	0.64
1:D:196:VAL:HG12	1:D:268:GLY:H	1.62	0.64
1:D:265:LEU:CD2	1:D:267:VAL:HG13	2.27	0.64
1:D:455:LEU:HD12	1:D:869:ARG:CZ	2.27	0.64
2:E:200:VAL:CG1	2:F:418:LEU:CD2	2.76	0.64
2:E:386:ASP:OD2	2:E:442:LEU:HA	1.97	0.64
2:E:437:ILE:HG21	2:E:440:THR:OG1	1.98	0.64
2:E:475:LEU:O	2:E:479:ILE:N	2.31	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:375:HIS:N	2:F:378:LEU:HD22	2.06	0.64
2:F:416:GLY:CA	2:F:419:GLU:HB3	2.28	0.64
2:F:442:LEU:CA	2:F:455:LEU:HD22	2.27	0.64
2:F:82:SER:O	2:F:83:LYS:HG3	1.97	0.64
1:A:597:ARG:HB3	1:A:616:GLU:OE1	1.98	0.64
2:B:423:SER:OG	2:B:428:LEU:HD22	1.97	0.64
2:C:190:LEU:HB2	2:C:314:MET:HE1	1.79	0.64
2:C:204:LEU:CG	2:C:206:TYR:HB3	2.27	0.64
2:C:243:THR:O	2:C:334:CYS:SG	2.56	0.64
1:A:937:THR:O	1:A:940:GLY:N	2.30	0.64
1:D:1027:LYS:CB	1:D:1091:GLU:HA	2.27	0.64
1:D:1227:THR:O	1:D:1227:THR:OG1	2.10	0.64
1:D:223:TRP:N	1:D:223:TRP:CD1	2.65	0.64
1:D:288:ARG:NH1	1:D:288:ARG:HG3	2.13	0.64
1:D:597:ARG:HB3	1:D:616:GLU:OE2	1.97	0.64
1:D:80:SER:O	1:D:83:LEU:HD13	1.97	0.64
1:D:826:ILE:HG12	1:D:827:ARG:H	1.62	0.64
1:D:948:ILE:HD12	1:D:1071:ARG:HB3	1.79	0.64
2:F:384:ALA:HB2	2:F:437:ILE:HD12	1.77	0.64
2:F:474:PHE:HA	2:F:477:LYS:HD3	1.77	0.64
2:F:376:PRO:HG3	2:F:482:ALA:HB1	1.79	0.64
1:A:604:ALA:CB	1:A:717:LEU:HD11	2.27	0.64
1:A:624:VAL:C	1:A:745:PRO:O	2.36	0.64
1:A:829:PRO:HB3	1:A:882:PRO:C	2.17	0.64
1:A:913:CYS:O	1:A:914:THR:C	2.35	0.64
2:C:262:TRP:O	2:C:265:LYS:HB3	1.97	0.64
2:C:425:LEU:HD13	2:C:442:LEU:HD11	1.79	0.64
2:C:455:LEU:C	2:C:456:ARG:NE	2.51	0.64
2:F:317:GLY:HA2	2:F:320:SER:CB	2.24	0.64
1:D:1019:TRP:HD1	1:D:1020:ILE:N	1.94	0.64
1:D:1131:ILE:O	1:D:1137:VAL:HA	1.98	0.64
1:D:885:THR:O	1:D:1140:LEU:HD23	1.98	0.64
1:D:1207:GLU:OE2	1:D:1210:TYR:HE2	1.80	0.64
1:D:204:ALA:O	1:D:205:GLU:HG2	1.98	0.64
1:D:271:VAL:HA	1:D:293:ASP:CG	2.18	0.64
1:D:592:LEU:CA	1:D:595:GLN:HG3	2.19	0.64
1:D:608:ASP:OD2	1:D:612:LEU:HB2	1.97	0.64
1:D:73:PRO:HG3	1:D:90:GLN:CG	2.24	0.64
1:D:76:ILE:CB	1:D:911:HIS:NE2	2.61	0.64
1:D:873:GLU:HB3	1:D:1200:PRO:HD3	1.78	0.64
1:D:914:THR:OG1	1:D:999:ASP:CB	2.46	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:917:GLY:O	1:D:920:THR:CA	2.46	0.64
2:F:120:VAL:CG2	2:F:121:PHE:N	2.61	0.64
2:F:456:ARG:HB2	2:F:463:LYS:HA	1.79	0.64
1:A:1140:LEU:HD23	1:A:1141:VAL:N	2.12	0.64
1:A:83:LEU:O	1:A:85:GLU:N	2.30	0.64
2:C:200:VAL:CG1	2:C:204:LEU:HB2	2.28	0.64
2:C:351:TYR:O	2:C:355:GLN:N	2.24	0.64
2:C:354:PHE:CE1	2:C:370:LYS:HB3	2.32	0.64
2:C:396:ARG:NH1	2:C:397:GLN:CA	2.60	0.64
2:C:395:LEU:HD21	2:C:443:VAL:HB	1.80	0.64
1:D:586:THR:H	1:D:587:PRO:HD3	1.63	0.64
1:D:999:ASP:C	1:D:1001:GLY:H	2.01	0.64
1:D:116:PRO:HB3	1:D:921:LEU:HD11	1.78	0.64
1:D:179:GLY:C	1:D:181:GLU:H	1.86	0.64
1:D:247:LEU:O	1:D:248:GLU:CB	2.44	0.64
1:D:255:SER:HG	1:D:258:GLN:H	1.41	0.64
1:D:438:ASN:O	1:D:441:TRP:CB	2.45	0.64
1:D:550:GLN:H	1:D:551:LYS:HZ2	1.46	0.64
1:D:633:LYS:CB	1:D:638:THR:CG2	2.76	0.64
2:E:369:ARG:HA	2:E:430:SER:CB	2.28	0.64
2:F:241:TRP:CB	2:F:336:LEU:CB	2.71	0.64
2:F:403:PHE:O	2:F:407:LEU:N	2.28	0.64
2:F:455:LEU:C	2:F:456:ARG:NE	2.51	0.64
2:F:468:ILE:HG13	2:F:472:LYS:HE3	1.79	0.64
1:A:996:ARG:CZ	1:A:1003:TRP:HB3	2.28	0.64
1:A:192:GLU:C	1:A:193:ARG:CG	2.66	0.64
2:C:238:SER:HA	2:C:338:VAL:HB	1.80	0.64
2:C:356:LEU:CD1	2:C:359:ASN:HD21	2.10	0.64
2:C:456:ARG:HG2	2:C:457:SER:N	2.11	0.64
2:C:454:HIS:CB	2:C:466:MET:HB2	2.26	0.64
1:A:807:ARG:O	1:A:810:SER:OG	2.15	0.64
1:D:259:ARG:CZ	1:D:259:ARG:CA	2.62	0.64
1:D:556:THR:CA	1:D:559:LEU:HD23	2.21	0.64
1:D:767:ALA:HB3	1:D:771:LEU:HB2	1.80	0.64
1:D:865:ALA:HB2	1:D:872:SER:O	1.98	0.64
2:E:254:PHE:CB	2:E:257:ARG:HH11	2.08	0.64
2:E:259:ARG:NH1	2:E:263:TRP:HE1	1.95	0.64
2:F:246:ARG:O	2:F:247:THR:C	2.36	0.64
1:A:890:ASP:OD1	1:A:1136:GLU:HG3	1.98	0.64
1:A:606:THR:HG23	1:A:613:HIS:H	1.59	0.64
1:A:648:PRO:O	1:A:652:ILE:N	2.28	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:813:VAL:HG23	1:A:814:VAL:H	1.63	0.64
1:A:823:ARG:C	1:A:825:VAL:H	2.00	0.64
2:C:253:ASP:O	2:C:257:ARG:HD2	1.98	0.64
2:C:343:ASP:C	2:C:347:LEU:CD1	2.64	0.64
2:C:417:TYR:H	2:C:417:TYR:HD2	0.82	0.64
2:C:474:PHE:HA	2:C:477:LYS:HD2	1.79	0.64
1:A:528:GLU:OE2	1:A:528:GLU:HA	1.97	0.64
1:A:542:ASP:O	1:A:546:ARG:N	2.24	0.64
1:D:1116:MET:HE2	1:D:1116:MET:HA	1.80	0.63
1:D:1126:ASP:N	1:D:1126:ASP:OD1	2.30	0.63
1:D:1131:ILE:HD13	1:D:1131:ILE:N	2.11	0.63
1:D:1155:ILE:O	1:D:1158:LEU:HG	1.97	0.63
1:D:307:PHE:CD2	1:D:1048:ALA:O	2.51	0.63
1:D:349:ASP:O	1:D:351:SER:N	2.31	0.63
1:D:895:GLU:O	1:D:898:ILE:HG12	1.98	0.63
2:F:126:PHE:HA	2:F:209:ALA:C	2.18	0.63
2:F:308:ASP:OD1	2:F:332:VAL:O	2.16	0.63
2:F:323:HIS:O	2:F:333:PRO:HD3	1.98	0.63
1:A:1040:LYS:CB	1:A:1044:VAL:HG11	2.28	0.63
1:A:192:GLU:HB2	1:A:195:LEU:CD2	2.29	0.63
1:A:275:ARG:HH21	1:A:845:VAL:CA	2.04	0.63
1:A:624:VAL:CG1	1:A:747:CYS:SG	2.86	0.63
1:A:652:ILE:HA	1:A:655:LEU:CG	2.27	0.63
1:A:878:VAL:HG12	1:A:879:GLN:NE2	2.13	0.63
2:C:203:ARG:N	2:C:325:ARG:HD3	2.13	0.63
2:C:240:VAL:C	2:C:336:LEU:HB3	2.18	0.63
2:E:270:PRO:HA	2:E:273:PHE:HE2	1.59	0.63
1:D:1026:ARG:CZ	1:D:1029:GLN:OE1	2.46	0.63
1:D:869:ARG:N	1:D:1200:PRO:HB2	2.08	0.63
1:D:131:TYR:HB2	1:D:138:HIS:ND1	2.13	0.63
1:D:551:LYS:CE	1:D:551:LYS:H	2.10	0.63
1:D:576:LEU:CD1	1:D:577:CYS:H	2.12	0.63
1:D:632:ALA:CB	1:D:657:ARG:HD2	2.28	0.63
1:D:282:TYR:HB2	1:D:841:LEU:HD12	1.80	0.63
1:D:918:TRP:C	1:D:920:THR:H	1.98	0.63
2:E:268:MET:CE	2:E:373:LYS:HD3	2.28	0.63
2:F:300:ILE:HG22	2:F:301:GLU:H	1.62	0.63
2:F:351:TYR:HD1	2:F:355:GLN:HG2	1.62	0.63
2:F:432:TYR:CG	2:F:437:ILE:HD11	2.32	0.63
2:F:88:ARG:NH1	2:F:88:ARG:HB2	2.13	0.63
1:A:1210:TYR:CD1	1:A:1213:PRO:HG2	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:271:VAL:CG1	1:A:272:SER:H	1.83	0.63
1:A:509:THR:HB	1:A:513:LEU:HD21	1.80	0.63
1:A:556:THR:CG2	2:C:452:LEU:HB2	2.28	0.63
1:A:591:LEU:HD12	1:A:594:LEU:CG	2.28	0.63
1:A:769:ASP:CG	1:A:770:PHE:H	2.01	0.63
1:A:275:ARG:HH22	1:A:845:VAL:HA	1.60	0.63
1:A:944:GLU:CA	1:A:947:LYS:HE3	2.28	0.63
2:B:116:THR:HG22	2:B:121:PHE:CE2	2.34	0.63
2:B:313:HIS:CD2	2:B:314:MET:N	2.66	0.63
2:C:268:MET:N	2:C:352:ASP:OD2	2.27	0.63
1:D:101:ARG:O	1:D:105:GLU:HG3	1.97	0.63
1:D:308:GLN:CD	1:D:1027:LYS:NZ	2.51	0.63
1:D:1064:ILE:HG23	1:D:1064:ILE:O	1.97	0.63
1:D:113:TRP:CZ3	1:D:116:PRO:HD2	2.34	0.63
1:D:1197:CYS:O	1:D:1199:THR:O	2.17	0.63
1:D:131:TYR:HB2	1:D:138:HIS:CE1	2.32	0.63
1:D:371:LYS:HG3	1:D:396:CYS:HB2	1.80	0.63
1:D:425:ALA:O	1:D:428:LEU:HD13	1.97	0.63
1:D:275:ARG:NH2	1:D:842:PRO:HB3	2.09	0.63
1:D:967:MET:HA	1:D:967:MET:CE	2.27	0.63
2:E:132:HIS:HB2	2:F:233:GLU:OE2	1.98	0.63
2:E:121:PHE:HA	2:F:407:LEU:HD11	1.80	0.63
2:F:454:HIS:HB2	2:F:466:MET:CB	2.28	0.63
1:A:363:ARG:O	1:A:364:LEU:HD12	1.98	0.63
1:A:433:SER:HB2	1:A:1129:PHE:CE1	2.33	0.63
1:A:586:THR:N	1:A:587:PRO:HD2	2.14	0.63
1:A:652:ILE:O	1:A:655:LEU:CD1	2.46	0.63
1:A:657:ARG:HA	1:A:660:CYS:HB3	1.79	0.63
1:A:868:ASP:O	1:A:869:ARG:CB	2.46	0.63
2:B:261:GLN:O	2:B:265:LYS:N	2.30	0.63
2:C:347:LEU:C	2:C:350:LEU:HD11	2.18	0.63
2:C:387:VAL:HG12	2:C:395:LEU:HD22	1.80	0.63
2:C:88:ARG:NH1	2:C:88:ARG:HB2	2.14	0.63
2:B:231:ILE:HD11	2:C:135:PRO:CG	2.29	0.63
1:A:539:PHE:O	1:A:540:GLN:CB	2.45	0.63
1:D:293:ASP:CB	1:D:296:SER:HB2	2.28	0.63
1:D:283:LEU:HA	1:D:434:TYR:HE1	1.61	0.63
1:D:576:LEU:C	1:D:577:CYS:O	2.34	0.63
1:D:612:LEU:HD11	1:D:622:TYR:CD2	2.33	0.63
1:D:788:GLY:O	1:D:791:ALA:HB3	1.98	0.63
1:D:856:GLU:HG3	1:D:860:LEU:HG	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:876:ALA:HA	1:D:1189:LEU:CB	2.25	0.63
2:E:200:VAL:CG1	2:F:418:LEU:HD23	2.29	0.63
1:A:238:GLN:O	1:A:242:ALA:HB2	1.97	0.63
1:A:436:PRO:CD	1:A:879:GLN:NE2	2.60	0.63
1:A:893:SER:C	1:A:895:GLU:N	2.48	0.63
1:A:919:MET:O	1:A:920:THR:O	2.15	0.63
1:A:996:ARG:HB2	1:A:1004:LEU:CD2	2.26	0.63
2:B:269:SER:C	2:B:271:SER:H	2.02	0.63
2:B:374:LEU:N	2:B:458:ARG:HH22	1.90	0.63
2:C:241:TRP:CB	2:C:336:LEU:CB	2.75	0.63
2:C:240:VAL:CA	2:C:336:LEU:HB3	2.28	0.63
1:D:516:GLU:O	1:D:517:GLY:O	2.16	0.63
1:D:499:LYS:NZ	1:D:499:LYS:HB2	2.13	0.63
1:A:539:PHE:CE2	1:A:540:GLN:HG2	2.34	0.63
2:B:481:SER:HA	2:B:484:ASN:OD1	1.98	0.63
1:D:1060:LYS:HA	1:D:1060:LYS:HE3	1.81	0.63
1:D:1133:ILE:O	1:D:1136:GLU:N	2.27	0.63
1:D:495:PHE:HB2	1:D:573:TYR:HE1	1.64	0.63
1:D:658:LYS:O	1:D:662:GLU:HG2	1.98	0.63
1:D:659:HIS:CD2	1:D:719:LEU:CB	2.82	0.63
2:E:375:HIS:ND1	2:E:376:PRO:HA	2.13	0.63
2:F:263:TRP:CZ2	2:F:301:GLU:OE2	2.51	0.63
2:F:363:ARG:CZ	2:F:363:ARG:HB3	2.29	0.63
2:F:360:SER:HB3	2:F:364:LYS:CD	2.28	0.63
2:F:442:LEU:HB2	2:F:455:LEU:CD2	2.28	0.63
2:F:449:GLU:C	2:F:451:GLY:N	2.43	0.63
2:F:470:LYS:HE3	2:F:471:LEU:HD21	1.79	0.63
1:A:1006:ARG:CA	1:A:1006:ARG:HH11	2.12	0.63
1:A:1030:ARG:CZ	1:A:1040:LYS:HG2	2.29	0.63
1:A:1112:MET:HB2	1:A:1113:LEU:HD23	1.81	0.63
1:A:114:GLY:C	1:A:116:PRO:HD2	2.18	0.63
1:A:890:ASP:OD2	1:A:1191:LYS:NZ	2.30	0.63
1:A:260:ASP:N	1:A:264:GLN:NE2	2.46	0.63
1:A:570:PRO:CB	1:A:572:TRP:CZ3	2.81	0.63
1:A:81:ARG:NH1	1:A:126:ARG:HA	2.12	0.63
1:A:879:GLN:HA	1:A:879:GLN:HE21	1.63	0.63
1:A:885:THR:O	1:A:1140:LEU:HD23	1.98	0.63
2:B:270:PRO:HB3	2:B:273:PHE:CE2	2.33	0.63
2:C:389:ARG:NE	2:C:390:GLY:H	1.94	0.63
1:D:136:ASP:OD2	1:D:1167:LYS:HE3	1.99	0.63
1:D:196:VAL:HG21	1:D:215:ILE:CD1	2.26	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:221:TYR:O	1:D:222:SER:OG	2.13	0.63
1:D:350:ILE:C	1:D:352:SER:N	2.48	0.63
1:D:283:LEU:HA	1:D:434:TYR:CE1	2.33	0.63
1:D:601:LYS:HG2	1:D:721:ALA:HA	1.80	0.63
2:F:375:HIS:HB3	2:F:378:LEU:HD13	1.80	0.63
2:F:474:PHE:HE1	2:F:478:TYR:CD2	2.15	0.63
1:A:301:ILE:HG23	1:A:302:SER:N	2.13	0.63
1:A:412:PRO:HG2	1:A:413:LEU:N	2.11	0.63
1:A:658:LYS:HB2	1:A:713:PRO:O	1.99	0.63
1:A:898:ILE:HG12	1:A:899:ALA:N	2.14	0.63
1:A:914:THR:CA	1:A:918:TRP:HB3	2.23	0.63
2:C:338:VAL:O	2:C:339:ASN:C	2.34	0.63
2:C:425:LEU:HD11	2:C:429:TYR:CE2	2.34	0.63
1:A:503:VAL:HG21	1:A:521:PRO:CA	2.28	0.63
1:D:1000:GLU:O	1:D:1003:TRP:N	2.32	0.63
1:D:175:TRP:HB2	1:D:185:VAL:CG1	2.27	0.63
1:D:439:GLN:HG2	1:D:835:GLY:HA2	1.79	0.63
1:D:457:ARG:CA	1:D:460:LYS:HD2	2.27	0.63
1:D:576:LEU:CD2	1:D:579:ARG:HE	2.11	0.63
2:E:380:PRO:O	2:E:382:LYS:CD	2.47	0.63
2:E:75:ARG:NH1	2:E:434:GLU:OE2	2.32	0.63
2:F:203:ARG:NE	2:F:204:LEU:CD1	2.62	0.63
2:F:303:LEU:HA	2:F:337:SER:CB	2.29	0.63
2:F:347:LEU:C	2:F:350:LEU:CD1	2.67	0.63
2:F:387:VAL:CG1	2:F:399:CYS:HB2	2.29	0.63
1:A:1006:ARG:NH1	1:A:1006:ARG:CA	2.58	0.63
1:A:1131:ILE:HG23	1:A:1137:VAL:HG13	1.80	0.63
1:A:1190:ARG:NE	1:A:1197:CYS:HB2	2.12	0.63
1:A:545:ALA:HA	1:A:548:CYS:SG	2.38	0.63
1:A:559:LEU:CB	1:A:560:PRO:CD	2.77	0.63
1:A:831:TYR:HD1	1:A:831:TYR:N	1.97	0.63
1:A:83:LEU:CD2	1:A:89:GLY:HA2	2.28	0.63
1:A:974:THR:HG22	1:A:975:GLN:N	2.14	0.63
2:B:381:ILE:HD13	2:B:437:ILE:HG23	1.79	0.63
2:B:78:PHE:O	2:B:78:PHE:HD1	1.82	0.63
2:C:127:PRO:CD	2:C:128:VAL:N	2.60	0.63
2:C:472:LYS:HA	2:C:475:LEU:CD1	2.28	0.63
2:E:390:GLY:N	2:E:395:LEU:CG	2.54	0.63
1:D:1007:GLU:O	1:D:1011:PRO:CD	2.45	0.63
1:D:1190:ARG:HH11	1:D:1195:MET:HE3	1.62	0.63
1:D:294:THR:O	1:D:298:HIS:HB2	1.98	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:386:ARG:O	1:D:389:PHE:CG	2.51	0.63
1:D:420:HIS:HD2	1:D:1020:ILE:CD1	1.92	0.63
1:D:544:MET:O	1:D:548:CYS:HB3	1.99	0.63
1:D:598:VAL:O	1:D:600:PRO:N	2.32	0.63
1:D:760:CYS:O	1:D:761:ASN:C	2.36	0.63
1:D:770:PHE:CZ	1:D:968:GLN:O	2.52	0.63
1:D:948:ILE:HG21	1:D:1071:ARG:HB3	1.81	0.63
2:F:215:PHE:N	2:F:215:PHE:CD1	2.65	0.63
2:F:239:LEU:N	2:F:338:VAL:CG2	2.61	0.63
2:F:193:TYR:OH	2:F:333:PRO:HG3	1.99	0.63
2:F:353:SER:CB	2:F:373:LYS:O	2.45	0.63
1:A:1157:ASN:O	1:A:1161:ARG:HD2	1.99	0.63
1:A:224:CYS:O	1:A:227:ARG:HB3	1.98	0.63
1:A:612:LEU:HB2	1:A:621:GLY:O	1.99	0.63
2:B:127:PRO:HB2	2:C:104:VAL:CG1	2.24	0.63
2:C:406:LEU:HA	2:C:409:ASN:HD21	1.63	0.63
1:D:1044:VAL:O	1:D:1047:ARG:CG	2.47	0.63
1:D:1028:VAL:CG2	1:D:1096:ARG:NH1	2.61	0.63
1:D:1171:ASN:O	1:D:1172:ASP:CB	2.47	0.63
1:D:316:LYS:CE	1:D:348:LEU:HD23	2.28	0.63
1:D:375:GLU:CD	1:D:379:LYS:HB3	2.18	0.63
1:D:454:GLU:HA	1:D:457:ARG:HG3	1.81	0.63
1:D:715:GLN:HG2	1:D:717:LEU:CG	2.29	0.63
1:D:728:THR:HG22	1:D:729:GLN:H	1.64	0.63
1:D:957:ALA:HB2	1:D:1094:THR:OG1	1.99	0.63
1:A:490:TRP:HA	1:A:580:LEU:HB2	1.80	0.63
1:A:630:ASN:ND2	1:A:632:ALA:N	2.47	0.63
1:A:955:TYR:HB3	1:A:1098:ASN:OD1	1.98	0.63
2:B:234:LYS:HG2	2:B:341:ASP:OD2	1.99	0.63
2:B:461:THR:CG2	2:B:463:LYS:HZ2	2.12	0.63
2:C:235:THR:O	2:C:341:ASP:OD1	2.16	0.63
2:C:342:LEU:CD2	2:C:345:GLY:N	2.54	0.63
2:C:402:LEU:C	2:C:406:LEU:HD22	2.19	0.63
1:A:634:LEU:HG	1:A:635:PRO:HA	1.80	0.63
1:D:164:PRO:CD	1:D:165:LYS:H	2.10	0.63
1:D:561:LYS:O	1:D:562:ARG:HB2	1.97	0.63
1:D:1058:PHE:CE2	1:D:1059:ASN:HB3	2.33	0.62
1:D:359:VAL:CG2	1:D:406:VAL:HG21	2.29	0.62
1:D:612:LEU:CD2	1:D:622:TYR:CE1	2.79	0.62
1:D:549:LEU:CD1	2:F:401:GLY:HA3	2.26	0.62
1:A:754:HIS:O	1:A:758:ASN:N	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:812:MET:HG3	1:A:813:VAL:H	1.64	0.62
1:A:913:CYS:HB2	1:A:999:ASP:OD1	1.99	0.62
2:B:342:LEU:C	2:B:342:LEU:HD13	2.20	0.62
2:B:77:HIS:HB2	2:C:195:ASN:OD1	1.99	0.62
2:C:322:LEU:C	2:C:322:LEU:HD22	2.19	0.62
2:C:342:LEU:CG	2:C:344:ARG:HB2	2.29	0.62
1:D:584:ALA:C	1:D:585:TRP:CE3	2.72	0.62
1:A:539:PHE:CD2	1:A:540:GLN:HG2	2.34	0.62
1:D:1208:ARG:HG3	1:D:1209:ARG:N	2.12	0.62
1:D:134:ASN:CB	1:D:137:GLN:HB2	2.29	0.62
1:D:177:ARG:O	1:D:178:TYR:C	2.37	0.62
1:D:213:VAL:CG1	1:D:401:TRP:CZ2	2.81	0.62
1:D:251:THR:OG1	1:D:254:SER:CA	2.43	0.62
1:D:917:GLY:O	1:D:920:THR:CB	2.46	0.62
2:E:199:LEU:HG	2:E:200:VAL:CG2	2.29	0.62
2:F:403:PHE:CD2	2:F:403:PHE:C	2.72	0.62
1:A:1207:GLU:C	1:A:1210:TYR:CD1	2.72	0.62
1:A:157:LEU:HD11	1:A:195:LEU:CA	2.29	0.62
1:A:385:ILE:HA	1:A:388:ASN:ND2	2.14	0.62
1:A:652:ILE:CA	1:A:655:LEU:HG	2.29	0.62
1:A:734:HIS:CA	1:A:752:LEU:HD11	2.29	0.62
1:A:786:ALA:HB1	1:A:790:ARG:CZ	2.28	0.62
1:A:792:LEU:CD1	1:A:793:GLU:HG2	2.29	0.62
1:A:448:ALA:CB	1:A:870:VAL:CG1	2.77	0.62
2:B:125:VAL:HG13	2:B:207:GLY:C	2.19	0.62
2:B:261:GLN:HA	2:B:264:ARG:CG	2.29	0.62
2:B:231:ILE:HG13	2:C:135:PRO:HG3	1.79	0.62
1:D:1117:LYS:O	1:D:1121:GLU:HG2	1.99	0.62
1:D:1236:GLN:CB	1:D:1237:PRO:HD3	2.29	0.62
1:D:197:PHE:HB2	1:D:277:HIS:HE1	1.64	0.62
1:D:210:THR:HG23	1:D:224:CYS:CB	2.23	0.62
1:D:389:PHE:HA	1:D:393:MET:HG2	1.77	0.62
1:D:459:MET:CB	1:D:460:LYS:HZ1	2.12	0.62
1:D:456:GLN:C	1:D:460:LYS:HZ2	1.97	0.62
1:D:608:ASP:CA	1:D:620:TRP:CZ3	2.82	0.62
1:D:627:ARG:O	1:D:628:ARG:HB3	1.98	0.62
1:D:648:PRO:O	1:D:652:ILE:HG13	1.99	0.62
1:D:743:ASP:HB2	1:D:744:ILE:HD13	1.80	0.62
1:D:452:TYR:HB3	1:D:801:TRP:CD1	2.33	0.62
1:D:282:TYR:HB2	1:D:841:LEU:CD1	2.29	0.62
2:F:124:GLN:O	2:F:125:VAL:HG12	2.00	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:233:GLU:OE1	2:F:131:LEU:HD12	1.98	0.62
2:F:277:ASP:OD2	2:F:285:LYS:HD2	1.99	0.62
2:F:307:GLY:HA2	2:F:335:VAL:H	1.64	0.62
2:F:343:ASP:O	2:F:347:LEU:HD13	2.00	0.62
2:E:201:ASN:OD1	2:F:422:GLN:CB	2.47	0.62
1:A:885:THR:C	1:A:1146:ARG:NE	2.53	0.62
1:A:1124:ALA:CB	1:A:1148:ARG:NH1	2.59	0.62
1:A:223:TRP:O	1:A:227:ARG:CB	2.47	0.62
1:A:226:GLN:CD	1:A:226:GLN:O	2.37	0.62
1:A:228:LEU:N	1:A:228:LEU:HD22	2.15	0.62
1:A:203:LEU:HD13	1:A:379:LYS:O	1.99	0.62
1:A:392:LEU:HD22	1:A:393:MET:HE3	1.80	0.62
1:A:655:LEU:HD13	1:A:719:LEU:O	1.98	0.62
1:A:656:TYR:CE1	1:A:744:ILE:HD12	2.34	0.62
2:B:396:ARG:HD2	2:B:417:TYR:CB	2.25	0.62
2:C:128:VAL:CG1	2:C:192:HIS:CD2	2.82	0.62
2:C:215:PHE:N	2:C:215:PHE:CD1	2.67	0.62
2:C:399:CYS:SG	2:C:400:GLN:N	2.72	0.62
1:D:1133:ILE:HG22	1:D:1134:HIS:CD2	2.35	0.62
1:D:1133:ILE:O	1:D:1134:HIS:C	2.37	0.62
1:D:1200:PRO:HG2	1:D:1203:PRO:CG	2.25	0.62
1:D:434:TYR:CG	1:D:435:LEU:N	2.67	0.62
1:D:614:TYR:HA	1:D:620:TRP:HA	1.81	0.62
1:D:748:TRP:CZ3	1:D:750:PHE:HE2	2.17	0.62
2:E:233:GLU:O	2:E:234:LYS:HG3	2.00	0.62
2:E:475:LEU:O	2:E:478:TYR:N	2.31	0.62
2:F:372:LEU:HD23	2:F:434:GLU:O	1.99	0.62
2:F:403:PHE:HD2	2:F:403:PHE:C	2.03	0.62
1:A:1149:ALA:O	1:A:1152:ALA:HB3	2.00	0.62
1:A:365:TYR:C	1:A:367:GLY:N	2.52	0.62
1:A:756:ASP:OD1	1:A:770:PHE:CZ	2.53	0.62
1:A:795:ASN:HD22	1:A:796:LYS:N	1.97	0.62
1:A:80:SER:N	1:A:83:LEU:CD1	2.55	0.62
1:A:869:ARG:CG	1:A:870:VAL:HG23	2.28	0.62
2:B:131:LEU:HD22	2:C:233:GLU:OE2	1.99	0.62
2:B:322:LEU:O	2:B:323:HIS:O	2.17	0.62
2:C:209:ALA:CB	2:C:239:LEU:HA	2.29	0.62
2:C:395:LEU:CD2	2:C:443:VAL:O	2.47	0.62
1:D:80:SER:HB2	1:D:125:LEU:CG	2.29	0.62
1:D:257:THR:OG1	1:D:287:SER:HB2	1.98	0.62
1:D:282:TYR:CE2	1:D:432:VAL:HG22	2.29	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:633:LYS:HB2	1:D:638:THR:CG2	2.29	0.62
1:D:771:LEU:O	1:D:774:MET:HB2	1.99	0.62
1:D:300:ALA:CB	1:D:847:ALA:HB1	2.13	0.62
2:E:91:LEU:HB3	2:E:347:LEU:HD11	1.81	0.62
2:E:380:PRO:N	2:E:438:LEU:CD1	2.62	0.62
2:F:114:TRP:NE1	2:F:259:ARG:CZ	2.56	0.62
2:F:346:MET:HE1	2:F:347:LEU:HD12	1.81	0.62
2:F:68:ALA:O	2:F:72:ILE:HG12	2.00	0.62
1:A:1145:ASP:HB3	1:A:1148:ARG:NH2	2.14	0.62
1:A:145:LYS:O	1:A:149:PRO:CD	2.47	0.62
1:A:607:TRP:CA	1:A:610:PHE:O	2.40	0.62
1:A:767:ALA:O	1:A:768:LYS:O	2.18	0.62
1:A:851:THR:HG22	1:A:853:ARG:HB2	1.80	0.62
2:B:236:GLU:HA	2:B:342:LEU:H	1.63	0.62
2:C:114:TRP:HE1	2:C:259:ARG:NH1	1.97	0.62
2:C:114:TRP:HE1	2:C:259:ARG:NH2	1.95	0.62
2:C:126:PHE:HA	2:C:209:ALA:C	2.19	0.62
2:C:264:ARG:HA	2:C:270:PRO:CB	2.28	0.62
1:D:509:THR:HB	1:D:512:LYS:CB	2.18	0.62
2:B:474:PHE:O	2:B:475:LEU:HB2	1.97	0.62
1:D:1014:ARG:HB3	1:D:1014:ARG:HH11	1.65	0.62
1:D:1145:ASP:HA	1:D:1148:ARG:NE	2.14	0.62
1:D:1157:ASN:OD1	1:D:1161:ARG:CZ	2.47	0.62
1:D:1193:VAL:O	1:D:1195:MET:HE2	2.00	0.62
1:D:1200:PRO:CD	1:D:1203:PRO:HG2	2.28	0.62
1:D:282:TYR:O	1:D:434:TYR:CD1	2.52	0.62
1:D:350:ILE:N	1:D:350:ILE:HD12	2.09	0.62
1:D:631:LEU:HD23	1:D:631:LEU:N	2.15	0.62
1:D:649:TYR:HD2	1:D:652:ILE:CG2	2.12	0.62
1:D:656:TYR:CB	1:D:744:ILE:HD12	2.28	0.62
1:D:922:GLN:HE21	1:D:931:LEU:HD13	1.64	0.62
1:D:944:GLU:HG2	1:D:1071:ARG:NH1	2.14	0.62
1:D:546:ARG:CZ	2:F:404:ASN:HB3	2.30	0.62
2:F:428:LEU:O	2:F:429:TYR:C	2.37	0.62
1:A:1155:ILE:HA	1:A:1158:LEU:CD1	2.29	0.62
1:A:215:ILE:CG2	1:A:216:SER:N	2.51	0.62
1:A:157:LEU:HD21	1:A:216:SER:OG	1.99	0.62
1:A:241:PRO:O	1:A:244:LEU:HB2	1.99	0.62
1:A:376:LEU:C	1:A:378:VAL:H	2.02	0.62
1:A:382:MET:O	1:A:383:LYS:HB2	2.00	0.62
1:A:559:LEU:CD2	2:C:451:GLY:O	2.47	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:801:TRP:HZ3	1:A:808:ILE:HD12	1.62	0.62
1:A:931:LEU:HD23	1:A:931:LEU:C	2.19	0.62
1:A:990:LYS:HD3	1:A:1000:GLU:OE2	1.99	0.62
1:A:992:LEU:CD2	1:A:997:LEU:HD12	2.16	0.62
2:B:103:GLY:O	2:B:106:LEU:HB3	1.99	0.62
2:C:431:LYS:NZ	2:C:435:MET:CG	2.61	0.62
1:D:1163:MET:HG3	1:D:1164:PHE:N	2.13	0.62
1:D:265:LEU:HD21	1:D:267:VAL:CG1	2.29	0.62
1:D:290:ARG:HB3	1:D:428:LEU:HB3	1.81	0.62
1:D:359:VAL:HG22	1:D:406:VAL:HG21	1.82	0.62
1:D:496:LYS:HE3	1:D:571:GLY:HA3	1.81	0.62
1:D:575:LYS:CE	1:D:581:ASP:O	2.47	0.62
1:D:744:ILE:CG1	1:D:745:PRO:CD	2.77	0.62
2:E:106:LEU:CA	2:E:109:ASN:OD1	2.47	0.62
2:E:112:ALA:O	2:E:115:TRP:N	2.33	0.62
2:E:297:LYS:O	2:E:298:GLU:HB2	1.99	0.62
2:F:193:TYR:CD1	2:F:193:TYR:C	2.72	0.62
2:F:425:LEU:HD11	2:F:429:TYR:CE2	2.35	0.62
1:A:197:PHE:CD1	1:A:213:VAL:O	2.53	0.62
1:A:241:PRO:HA	1:A:244:LEU:HG	1.79	0.62
1:A:297:MET:HE1	1:A:415:LEU:HD21	1.80	0.62
1:A:452:TYR:CD2	1:A:453:GLU:HG2	2.35	0.62
1:A:482:ASP:HB2	1:A:483:PRO:HD3	1.80	0.62
1:A:732:TYR:HD1	1:A:755:LYS:HZ3	1.44	0.62
1:A:898:ILE:HG12	1:A:899:ALA:H	1.64	0.62
2:B:259:ARG:O	2:B:262:TRP:N	2.33	0.62
2:B:324:GLY:HA2	2:B:325:ARG:NH2	2.13	0.62
2:B:407:LEU:CD2	2:C:120:VAL:HA	2.14	0.62
2:C:208:LEU:HD11	2:C:240:VAL:CB	2.28	0.62
1:D:515:ILE:HG23	1:D:565:HIS:O	1.98	0.62
1:D:1190:ARG:NH1	1:D:1195:MET:CE	2.62	0.62
1:D:200:GLU:CB	1:D:211:LEU:HD13	2.30	0.62
1:D:214:ALA:HB2	1:D:219:ALA:HB3	1.81	0.62
1:D:229:VAL:HG12	1:D:382:MET:O	2.00	0.62
1:D:366:GLY:N	1:D:367:PRO:CD	2.61	0.62
1:D:429:GLU:HA	1:D:429:GLU:OE1	1.99	0.62
1:D:608:ASP:CB	1:D:620:TRP:CE3	2.78	0.62
1:D:823:ARG:O	1:D:827:ARG:N	2.18	0.62
1:D:964:ARG:O	1:D:968:GLN:HG2	1.99	0.62
2:E:131:LEU:C	2:E:182:ARG:HH21	2.03	0.62
2:F:336:LEU:CD2	2:F:336:LEU:C	2.68	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1153:LEU:CD1	1:A:1157:ASN:HD21	2.13	0.62
1:A:1200:PRO:CB	1:A:1202:ASN:O	2.46	0.62
1:A:256:PRO:HB3	1:A:280:GLU:O	1.99	0.62
1:A:624:VAL:CG2	1:A:624:VAL:O	2.48	0.62
2:B:396:ARG:NH1	2:B:417:TYR:HD2	1.93	0.62
2:E:249:ASN:HA	2:E:252:LEU:HB2	1.80	0.62
2:B:247:THR:O	2:B:251:TRP:HB2	2.00	0.62
1:D:175:TRP:HA	1:D:185:VAL:HG21	1.81	0.62
1:D:345:TRP:CA	1:D:1040:LYS:HZ2	2.13	0.62
1:D:295:MET:HE2	1:D:354:ASN:HA	1.80	0.62
1:D:74:LEU:HD13	1:D:76:ILE:O	2.00	0.62
1:D:860:LEU:HD22	1:D:1132:SER:O	2.00	0.62
1:D:877:MET:CE	1:D:1207:GLU:HB2	2.30	0.62
1:D:973:LEU:N	1:D:973:LEU:HD23	2.14	0.62
2:E:205:PRO:HB3	2:E:242:PHE:O	1.99	0.62
2:F:241:TRP:CG	2:F:243:THR:HG23	2.34	0.62
1:A:1068:ASP:O	1:A:1069:ILE:HG22	1.99	0.62
1:A:669:MET:O	1:A:673:ALA:CA	2.48	0.62
2:C:384:ALA:CB	2:C:437:ILE:HD12	2.29	0.62
1:D:1027:LYS:CB	1:D:1096:ARG:HE	2.12	0.62
1:D:1044:VAL:O	1:D:1047:ARG:HG3	1.99	0.62
1:D:1060:LYS:HD2	1:D:1060:LYS:N	2.15	0.62
1:D:193:ARG:HD2	1:D:263:GLU:CD	2.20	0.62
1:D:215:ILE:O	1:D:215:ILE:HG22	1.99	0.62
1:D:242:ALA:HB1	1:D:245:ILE:HD13	1.82	0.62
1:D:719:LEU:CD1	1:D:720:THR:N	2.62	0.62
1:D:597:ARG:HD3	1:D:724:GLY:O	1.99	0.62
1:D:735:GLY:CA	1:D:749:PHE:HA	2.30	0.62
1:D:739:TYR:O	1:D:741:ASP:N	2.33	0.62
1:D:767:ALA:O	1:D:768:LYS:HB3	2.00	0.62
1:D:868:ASP:C	1:D:869:ARG:HG3	2.20	0.62
2:F:240:VAL:HA	2:F:336:LEU:HB3	1.82	0.62
2:F:371:VAL:HG23	2:F:433:ASP:HB3	1.82	0.62
1:A:175:TRP:HB2	1:A:184:ALA:O	2.00	0.62
1:A:394:GLN:O	1:A:397:ALA:HB3	2.00	0.62
1:A:400:VAL:CG1	1:A:404:HIS:NE2	2.63	0.62
1:A:623:LEU:O	1:A:625:PRO:HD2	2.00	0.62
1:A:960:PRO:HD2	1:A:961:PHE:CE1	2.35	0.62
2:C:335:VAL:HG22	2:C:336:LEU:N	2.15	0.62
2:C:403:PHE:CD2	2:C:403:PHE:C	2.73	0.62
1:D:1010:LEU:O	1:D:1028:VAL:HG21	1.99	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:145:LYS:HB2	1:D:1118:TRP:CH2	2.34	0.61
1:D:213:VAL:CG2	1:D:401:TRP:CZ2	2.82	0.61
1:D:279:ARG:NH1	1:D:433:SER:O	2.32	0.61
1:D:470:ALA:O	1:D:474:LEU:CD1	2.47	0.61
1:D:738:PRO:O	1:D:740:ASN:ND2	2.33	0.61
1:D:656:TYR:CE1	1:D:743:ASP:CB	2.83	0.61
1:D:891:VAL:HG13	1:D:1135:ASP:CB	2.16	0.61
1:D:938:THR:O	1:D:939:VAL:CB	2.47	0.61
1:D:948:ILE:O	1:D:951:TYR:HB2	2.00	0.61
2:E:203:ARG:HH21	2:E:203:ARG:HG3	1.64	0.61
2:E:110:LEU:HA	2:E:378:LEU:CD2	2.30	0.61
2:E:406:LEU:HG	2:E:413:VAL:CG2	2.29	0.61
2:E:69:LEU:N	2:E:69:LEU:HD12	2.14	0.61
2:F:290:TYR:CD1	2:F:299:LEU:HA	2.35	0.61
2:F:314:MET:HE3	2:F:315:TYR:CE2	2.35	0.61
2:F:400:GLN:HE21	2:F:404:ASN:CG	2.03	0.61
2:F:385:LEU:HD21	2:F:402:LEU:HD23	1.81	0.61
1:A:1068:ASP:HA	1:A:1071:ARG:HD3	1.80	0.61
1:A:1155:ILE:O	1:A:1159:LEU:HG	1.99	0.61
1:A:273:PHE:CA	1:A:276:ALA:CB	2.74	0.61
1:A:299:MET:HE1	1:A:353:VAL:HB	1.82	0.61
1:A:385:ILE:HA	1:A:388:ASN:HD22	1.65	0.61
1:A:624:VAL:H	1:A:746:GLY:HA2	1.64	0.61
1:A:771:LEU:HD22	1:A:771:LEU:N	2.09	0.61
1:A:893:SER:CB	1:A:896:LEU:HD23	2.29	0.61
2:B:419:GLU:CB	2:C:203:ARG:NH2	2.59	0.61
1:D:1131:ILE:CD1	1:D:1138:ARG:N	2.33	0.61
1:D:1187:ARG:HD2	1:D:1212:ILE:HD11	1.81	0.61
1:D:869:ARG:CA	1:D:1200:PRO:HB2	2.30	0.61
1:D:1200:PRO:HG2	1:D:1200:PRO:O	2.00	0.61
1:D:1232:GLU:HG3	1:D:1234:ARG:HB2	1.80	0.61
1:D:252:GLY:O	1:D:253:ALA:HB3	2.00	0.61
1:D:453:GLU:O	1:D:456:GLN:N	2.32	0.61
1:D:576:LEU:HD23	1:D:579:ARG:HE	1.64	0.61
1:D:81:ARG:CZ	1:D:81:ARG:HB2	2.30	0.61
1:D:827:ARG:O	1:D:829:PRO:CD	2.30	0.61
1:D:831:TYR:O	1:D:831:TYR:HD1	1.83	0.61
1:D:902:LEU:O	1:D:905:ALA:HB3	1.99	0.61
2:E:407:LEU:C	2:E:409:ASN:H	2.04	0.61
2:F:466:MET:CA	2:F:471:LEU:HD22	2.31	0.61
1:A:169:TRP:O	1:A:170:ALA:HB3	1.99	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:434:TYR:C	1:A:435:LEU:HG	2.20	0.61
1:A:630:ASN:C	1:A:631:LEU:HD23	2.20	0.61
1:A:945:HIS:HA	1:A:948:ILE:HD11	1.82	0.61
2:B:113:GLU:C	2:B:262:TRP:HZ2	2.04	0.61
2:B:400:GLN:HE22	2:B:417:TYR:HB3	1.65	0.61
2:C:416:GLY:CA	2:C:419:GLU:HB3	2.30	0.61
2:C:395:LEU:HD13	2:C:444:THR:HA	1.80	0.61
1:D:499:LYS:HE3	1:D:499:LYS:O	2.00	0.61
1:D:1027:LYS:HB3	1:D:1091:GLU:HA	1.83	0.61
1:D:543:VAL:O	1:D:547:ALA:HB3	2.00	0.61
1:D:598:VAL:HA	1:D:601:LYS:CD	2.30	0.61
1:D:733:HIS:ND1	1:D:750:PHE:HD1	1.98	0.61
2:E:123:GLU:HB3	2:F:403:PHE:CE1	2.36	0.61
2:F:243:THR:O	2:F:334:CYS:CB	2.48	0.61
2:F:306:LEU:HD21	2:F:310:GLU:HG2	1.81	0.61
2:F:310:GLU:O	2:F:314:MET:N	2.30	0.61
2:F:342:LEU:CD2	2:F:345:GLY:N	2.55	0.61
2:F:455:LEU:O	2:F:465:MET:HA	1.99	0.61
1:A:1030:ARG:HE	1:A:1040:LYS:HE3	1.64	0.61
1:A:158:LEU:O	1:A:409:GLN:NE2	2.29	0.61
1:A:308:GLN:HB3	1:A:1090:GLU:HG3	1.83	0.61
1:A:719:LEU:HD23	1:A:719:LEU:H	1.64	0.61
1:A:743:ASP:CB	1:A:744:ILE:HD13	2.30	0.61
2:B:116:THR:HA	2:B:120:VAL:HG23	1.82	0.61
2:B:378:LEU:HD12	2:B:378:LEU:N	2.16	0.61
2:B:106:LEU:HB2	2:B:380:PRO:HG3	1.81	0.61
2:C:120:VAL:CG2	2:C:121:PHE:N	2.63	0.61
2:C:342:LEU:HB2	2:C:344:ARG:HG3	1.82	0.61
2:C:414:TRP:O	2:C:415:PRO:O	2.19	0.61
2:C:443:VAL:C	2:C:447:THR:HG21	2.21	0.61
1:D:1207:GLU:O	1:D:1208:ARG:C	2.39	0.61
1:D:270:ASN:HA	1:D:294:THR:CB	2.31	0.61
1:D:383:LYS:O	1:D:386:ARG:CG	2.46	0.61
1:D:486:TRP:HE1	1:D:722:ARG:HG3	1.65	0.61
1:D:742:VAL:HB	1:D:748:TRP:CZ3	2.35	0.61
1:D:795:ASN:HA	1:D:798:ILE:HD11	1.80	0.61
1:D:283:LEU:HG	1:D:814:VAL:CG2	2.30	0.61
1:D:977:GLU:HA	1:D:980:GLU:HG3	1.81	0.61
2:E:189:ALA:HB1	2:E:240:VAL:HG11	1.82	0.61
2:F:241:TRP:H	2:F:336:LEU:CB	2.11	0.61
1:A:1002:GLU:O	1:A:1005:VAL:CG2	2.44	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1048:ALA:HB3	1:A:1092:PHE:HE1	1.65	0.61
1:A:372:GLU:OE2	1:A:399:ASP:OD2	2.17	0.61
1:A:660:CYS:HB2	1:A:745:PRO:HG3	1.83	0.61
1:A:769:ASP:CG	1:A:770:PHE:N	2.52	0.61
2:B:93:SER:OG	2:B:95:CYS:HB2	2.01	0.61
1:D:666:GLN:O	1:D:669:MET:O	2.17	0.61
1:D:161:GLN:O	1:D:163:PRO:HD3	2.00	0.61
2:C:367:LEU:O	2:C:369:ARG:CD	2.47	0.61
2:B:294:PRO:HG2	2:B:295:TRP:H	1.66	0.61
1:D:1210:TYR:C	1:D:1210:TYR:CD1	2.74	0.61
1:D:176:THR:OG1	1:D:218:SER:HB2	2.00	0.61
1:D:307:PHE:HD2	1:D:1048:ALA:O	1.82	0.61
1:D:359:VAL:O	1:D:359:VAL:CG1	2.46	0.61
1:D:792:LEU:CA	1:D:795:ASN:ND2	2.64	0.61
1:D:996:ARG:HD3	1:D:996:ARG:H	1.60	0.61
2:E:189:ALA:CB	2:E:240:VAL:HG21	2.30	0.61
2:E:303:LEU:HA	2:E:337:SER:O	2.00	0.61
2:F:243:THR:HB	2:F:247:THR:HG1	1.66	0.61
2:F:439:PHE:HD1	2:F:458:ARG:CG	2.13	0.61
1:A:1030:ARG:CD	1:A:1040:LYS:HZ1	2.14	0.61
1:A:228:LEU:CD2	1:A:229:VAL:N	2.55	0.61
1:A:477:GLU:HG2	1:A:482:ASP:OD1	2.00	0.61
1:A:656:TYR:C	1:A:659:HIS:HB3	2.21	0.61
1:A:815:TRP:HE3	1:A:838:GLY:H	1.47	0.61
1:A:886:LEU:H	1:A:1146:ARG:NH2	1.96	0.61
2:B:106:LEU:O	2:B:109:ASN:HB3	2.00	0.61
2:B:238:SER:HA	2:B:339:ASN:ND2	2.14	0.61
2:B:311:LEU:O	2:B:313:HIS:N	2.33	0.61
2:C:402:LEU:HG	2:C:472:LYS:HZ1	1.63	0.61
2:C:431:LYS:HE2	2:C:435:MET:HB2	1.83	0.61
1:A:503:VAL:CG2	1:A:520:ALA:O	2.45	0.61
1:D:1025:LEU:CD2	1:D:1025:LEU:O	2.49	0.61
1:D:1031:GLU:CD	1:D:1096:ARG:HH22	2.03	0.61
1:D:380:GLY:O	1:D:385:ILE:HD12	2.00	0.61
1:D:595:GLN:O	1:D:597:ARG:N	2.33	0.61
1:D:626:GLY:HA3	1:D:628:ARG:O	2.00	0.61
1:D:938:THR:O	1:D:939:VAL:HB	2.00	0.61
2:E:478:TYR:CD1	2:E:478:TYR:O	2.54	0.61
2:E:90:SER:HA	2:E:93:SER:HB2	1.81	0.61
2:F:132:HIS:ND1	2:F:132:HIS:N	2.42	0.61
2:F:255:TRP:O	2:F:256:LEU:C	2.39	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1002:GLU:HG3	1:A:1003:TRP:HE3	1.65	0.61
1:A:169:TRP:HZ3	1:A:391:ASP:CA	2.11	0.61
1:A:225:SER:HG	1:A:386:ARG:HB2	1.64	0.61
1:A:572:TRP:O	1:A:573:TYR:C	2.36	0.61
1:A:894:GLN:O	1:A:898:ILE:HG23	2.00	0.61
2:B:342:LEU:HD13	2:B:342:LEU:O	2.00	0.61
2:C:307:GLY:C	2:C:308:ASP:OD2	2.39	0.61
2:C:78:PHE:HZ	2:C:346:MET:CE	2.14	0.61
2:C:447:THR:O	2:C:448:LEU:C	2.38	0.61
2:B:281:GLU:HG3	2:B:282:GLU:HG2	1.82	0.61
1:A:1055:SER:O	1:A:1059:ASN:OD1	2.19	0.61
1:D:1210:TYR:CG	1:D:1211:GLY:N	2.68	0.61
1:D:1187:ARG:CD	1:D:1212:ILE:HD11	2.29	0.61
1:D:1183:VAL:O	1:D:1217:ALA:HA	2.00	0.61
1:D:275:ARG:HB2	1:D:843:GLN:CA	2.30	0.61
1:D:353:VAL:HG23	1:D:354:ASN:N	2.15	0.61
1:D:369:LEU:HD13	1:D:372:GLU:HB3	1.82	0.61
1:D:434:TYR:O	1:D:435:LEU:CG	2.47	0.61
1:D:866:ARG:HB3	1:D:867:PRO:CD	2.24	0.61
2:F:301:GLU:OE1	2:F:339:ASN:ND2	2.33	0.61
2:F:309:HIS:HA	2:F:312:LEU:CD1	2.23	0.61
2:F:306:LEU:HD21	2:F:310:GLU:OE1	2.01	0.61
2:F:81:GLY:HA3	2:F:97:PRO:CB	2.31	0.61
1:A:1113:LEU:HB3	1:A:1117:LYS:CE	2.29	0.61
1:A:169:TRP:CZ3	1:A:391:ASP:CA	2.80	0.61
1:A:241:PRO:C	1:A:244:LEU:H	2.04	0.61
1:A:259:ARG:HG3	1:A:261:TRP:CD1	2.35	0.61
1:A:196:VAL:C	1:A:401:TRP:HH2	2.04	0.61
1:A:473:LEU:N	1:A:473:LEU:CD2	2.62	0.61
1:A:70:ARG:CB	1:A:81:ARG:HB3	2.20	0.61
1:A:893:SER:CA	1:A:896:LEU:HD23	2.30	0.61
2:C:131:LEU:O	2:C:133:HIS:HB2	2.01	0.61
2:C:244:PRO:N	2:C:247:THR:CG2	2.63	0.61
2:C:387:VAL:CG1	2:C:399:CYS:CB	2.78	0.61
2:C:394:GLU:O	2:C:397:GLN:HB3	2.01	0.61
1:D:1056:GLU:HB3	1:D:1057:MET:CE	2.31	0.61
1:D:1055:SER:O	1:D:1059:ASN:ND2	2.34	0.61
1:D:1210:TYR:HH	1:D:1212:ILE:HB	1.64	0.61
1:D:185:VAL:HB	1:D:186:PRO:HD2	1.83	0.61
1:D:190:PRO:HG3	1:D:219:ALA:HB2	1.81	0.61
1:D:313:ILE:HG13	1:D:314:ALA:N	2.16	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:367:PRO:HG2	1:D:368:PRO:HD3	1.82	0.61
1:D:632:ALA:HB1	1:D:657:ARG:HD2	1.83	0.61
2:E:456:ARG:CG	2:E:457:SER:N	2.64	0.61
2:E:456:ARG:CB	2:E:463:LYS:HG2	2.17	0.61
2:F:442:LEU:O	2:F:455:LEU:CD2	2.48	0.61
1:A:1026:ARG:CB	1:A:1029:GLN:CD	2.66	0.61
1:A:1189:LEU:HD22	1:A:1189:LEU:N	2.16	0.61
1:A:199:VAL:CG2	1:A:277:HIS:HB2	2.31	0.61
1:A:275:ARG:HH22	1:A:844:VAL:C	2.03	0.61
1:A:928:GLY:O	1:A:931:LEU:HB3	2.00	0.61
2:B:258:HIS:CG	2:B:259:ARG:N	2.66	0.61
2:C:124:GLN:O	2:C:125:VAL:HG12	2.01	0.61
2:C:208:LEU:CD2	2:C:242:PHE:CD2	2.84	0.61
2:C:298:GLU:OE1	2:C:344:ARG:NH2	2.33	0.61
2:E:217:PRO:HA	2:E:231:ILE:HA	1.82	0.61
2:C:282:GLU:C	2:C:284:ARG:H	2.04	0.61
1:D:1221:TYR:O	1:D:1222:GLN:C	2.36	0.61
1:D:289:MET:SD	1:D:291:PHE:CZ	2.94	0.61
1:D:412:PRO:HG2	1:D:413:LEU:H	1.66	0.61
1:D:420:HIS:O	1:D:424:LEU:HG	2.01	0.61
1:D:428:LEU:CD1	1:D:428:LEU:N	2.46	0.61
1:D:548:CYS:H	1:D:551:LYS:HZ3	1.45	0.61
1:D:494:GLU:HA	1:D:573:TYR:H	1.66	0.61
1:D:612:LEU:H	1:D:612:LEU:HD12	1.62	0.61
1:D:801:TRP:CG	1:D:802:ARG:N	2.69	0.61
2:F:241:TRP:H	2:F:336:LEU:HB3	1.65	0.61
2:F:456:ARG:CG	2:F:457:SER:N	2.63	0.61
1:A:1072:THR:N	1:A:1073:PRO:HD2	2.15	0.61
1:A:1236:GLN:N	1:A:1236:GLN:HE21	1.97	0.61
1:A:392:LEU:HD23	1:A:392:LEU:C	2.22	0.61
1:A:410:GLN:O	1:A:414:PHE:HD2	1.83	0.61
1:A:420:HIS:O	1:A:424:LEU:HG	2.01	0.61
1:A:653:GLU:HA	1:A:656:TYR:CE2	2.35	0.61
1:A:73:PRO:HB2	1:A:90:GLN:HB3	1.83	0.61
2:B:125:VAL:HG12	2:B:126:PHE:N	2.15	0.61
2:B:127:PRO:O	2:C:104:VAL:HG21	2.01	0.61
2:B:236:GLU:CD	2:B:341:ASP:HB3	2.21	0.61
2:C:402:LEU:O	2:C:406:LEU:HD13	2.00	0.61
1:D:500:ALA:HB2	1:D:521:PRO:CG	2.31	0.61
1:D:1030:ARG:C	1:D:1032:THR:H	2.04	0.61
1:D:275:ARG:HB2	1:D:843:GLN:HB3	0.75	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:293:ASP:HB3	1:D:296:SER:HB2	1.83	0.61
1:D:371:LYS:HG3	1:D:396:CYS:CB	2.31	0.61
1:D:402:ALA:O	1:D:406:VAL:HG13	2.00	0.61
1:D:618:HIS:CB	1:D:725:PRO:CG	2.77	0.61
1:D:914:THR:HG21	1:D:999:ASP:CB	2.30	0.61
2:E:456:ARG:HD3	2:E:463:LYS:CG	2.29	0.61
2:F:425:LEU:HD21	2:F:429:TYR:OH	2.00	0.61
2:F:449:GLU:C	2:F:449:GLU:CD	2.59	0.61
1:A:1204:THR:CG2	1:A:1204:THR:O	2.37	0.61
1:A:259:ARG:NE	1:A:259:ARG:CA	2.60	0.61
1:A:261:TRP:O	1:A:262:GLN:HB2	2.00	0.61
1:A:195:LEU:HG	1:A:265:LEU:CD2	2.31	0.61
1:A:411:LEU:N	1:A:412:PRO:HD2	2.16	0.61
1:A:441:TRP:CZ3	1:A:445:LEU:HD12	2.36	0.61
1:A:553:LYS:HD2	2:C:468:ILE:CG2	2.31	0.61
1:A:813:VAL:CG1	1:A:815:TRP:CZ2	2.84	0.61
1:A:81:ARG:HG3	1:A:81:ARG:HH11	1.66	0.61
2:C:200:VAL:CB	2:C:204:LEU:HD22	2.31	0.61
2:C:407:LEU:HD13	2:C:408:GLU:OE2	2.01	0.61
2:C:395:LEU:CD2	2:C:443:VAL:HG12	2.28	0.61
2:E:417:TYR:HD1	2:E:418:LEU:HG	1.66	0.61
2:B:466:MET:HG2	2:B:467:HIS:N	2.15	0.61
1:D:966:LEU:CD2	1:D:970:ASN:ND2	2.64	0.61
1:D:1031:GLU:CD	1:D:1096:ARG:NH2	2.54	0.60
1:D:1107:ASP:O	1:D:1110:HIS:HB2	2.01	0.60
1:D:1113:LEU:O	1:D:1116:MET:HB2	2.01	0.60
1:D:1190:ARG:CD	1:D:1195:MET:HE3	2.26	0.60
1:D:1221:TYR:O	1:D:1224:ILE:N	2.34	0.60
1:D:140:ARG:HG3	1:D:140:ARG:HH11	1.65	0.60
1:D:239:LEU:N	1:D:239:LEU:HD23	2.16	0.60
1:D:603:MET:O	1:D:605:LEU:N	2.34	0.60
1:D:909:GLY:O	1:D:912:GLY:HA2	2.00	0.60
2:E:375:HIS:NE2	2:E:376:PRO:HB3	2.16	0.60
2:F:303:LEU:HG	2:F:337:SER:OG	2.01	0.60
2:F:197:LEU:CD1	2:F:322:LEU:HD21	2.31	0.60
2:F:403:PHE:HE1	2:F:415:PRO:HG3	1.60	0.60
2:F:395:LEU:CB	2:F:443:VAL:HG21	2.30	0.60
1:A:1060:LYS:NZ	1:A:1073:PRO:HB2	2.16	0.60
1:A:210:THR:O	1:A:211:LEU:CB	2.38	0.60
1:A:257:THR:HA	1:A:258:GLN:OE1	2.01	0.60
1:A:411:LEU:N	1:A:412:PRO:CD	2.64	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:851:THR:HG22	1:A:853:ARG:N	2.13	0.60
1:A:887:VAL:N	1:A:1146:ARG:NH1	2.48	0.60
1:A:950:ASN:O	1:A:951:TYR:C	2.39	0.60
2:C:307:GLY:HA2	2:C:335:VAL:H	1.66	0.60
2:C:368:HIS:NE2	2:C:370:LYS:HD2	2.16	0.60
2:C:387:VAL:CG2	2:C:417:TYR:HA	2.31	0.60
2:C:317:GLY:C	2:C:319:VAL:N	2.47	0.60
2:F:326:ASP:CG	2:F:329:LYS:HB2	2.22	0.60
1:A:96:GLY:O	1:A:100:VAL:HG23	2.01	0.60
1:D:1006:ARG:HH21	1:D:1017:GLY:HA3	1.64	0.60
1:D:1014:ARG:C	1:D:1014:ARG:HD2	2.21	0.60
1:D:1198:LYS:HA	1:D:1204:THR:HB	1.83	0.60
1:D:129:PRO:O	1:D:131:TYR:CZ	2.54	0.60
1:D:273:PHE:O	1:D:277:HIS:HD2	1.84	0.60
1:D:347:TRP:NE1	1:D:349:ASP:OD2	2.33	0.60
1:D:790:ARG:O	1:D:794:ILE:HG12	2.01	0.60
2:E:385:LEU:CD1	2:E:441:VAL:HB	2.31	0.60
2:E:447:THR:HG23	2:E:452:LEU:N	2.16	0.60
1:A:1019:TRP:CZ2	1:A:1111:LEU:HD21	2.36	0.60
1:A:1128:ARG:CG	1:A:1129:PHE:N	2.43	0.60
1:A:1231:LEU:HD12	1:A:1235:SER:HB2	1.83	0.60
1:A:174:GLY:O	1:A:175:TRP:CD2	2.54	0.60
1:A:254:SER:O	1:A:255:SER:CB	2.49	0.60
1:A:313:ILE:HA	1:A:316:LYS:HG2	1.84	0.60
1:A:434:TYR:O	1:A:1129:PHE:CE1	2.54	0.60
1:A:513:LEU:CB	1:A:568:GLY:CA	2.56	0.60
1:A:582:ASP:O	1:A:583:PRO:C	2.39	0.60
2:B:182:ARG:HB3	2:B:215:PHE:CD1	2.36	0.60
2:C:114:TRP:NE1	2:C:259:ARG:NH1	2.49	0.60
2:C:387:VAL:HG12	2:C:395:LEU:HD23	1.81	0.60
1:A:739:TYR:C	1:A:741:ASP:O	2.39	0.60
2:C:272:ASN:HA	2:C:292:ASN:ND2	2.15	0.60
2:B:122:ARG:HB3	2:B:122:ARG:NH1	2.16	0.60
1:D:891:VAL:CG1	1:D:1135:ASP:CB	2.76	0.60
1:D:368:PRO:CB	1:D:369:LEU:HD23	2.31	0.60
2:E:214:CYS:SG	2:E:236:GLU:HB3	2.41	0.60
2:F:254:PHE:N	2:F:257:ARG:NH2	2.48	0.60
1:A:866:ARG:CG	1:A:1069:ILE:CG2	2.75	0.60
1:A:1093:MET:O	1:A:1097:VAL:HG23	2.01	0.60
1:A:352:SER:C	1:A:358:LEU:HD21	2.22	0.60
1:A:816:LEU:HD12	1:A:837:TYR:O	2.00	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:918:TRP:C	1:A:920:THR:N	2.52	0.60
1:A:932:HIS:O	1:A:935:THR:HB	2.01	0.60
1:A:964:ARG:HD2	1:A:1053:THR:HG21	1.82	0.60
2:B:437:ILE:HG22	2:B:438:LEU:N	2.15	0.60
2:C:239:LEU:O	2:C:338:VAL:HG13	2.01	0.60
2:C:338:VAL:O	2:C:338:VAL:HG23	2.01	0.60
1:D:1155:ILE:CA	1:D:1158:LEU:HD21	2.27	0.60
1:D:224:CYS:O	1:D:227:ARG:N	2.35	0.60
2:E:319:VAL:HG13	2:E:320:SER:N	2.16	0.60
2:F:200:VAL:O	2:F:201:ASN:HB2	2.01	0.60
1:D:1209:ARG:HH12	2:F:252:LEU:HD13	1.64	0.60
2:F:254:PHE:O	2:F:257:ARG:HB2	2.01	0.60
1:A:1133:ILE:HD13	1:A:1133:ILE:C	2.21	0.60
1:A:213:VAL:HG22	1:A:221:TYR:CE2	2.36	0.60
1:A:301:ILE:CG2	1:A:302:SER:N	2.64	0.60
1:A:878:VAL:C	1:A:879:GLN:NE2	2.54	0.60
2:B:244:PRO:O	2:B:248:SER:CB	2.45	0.60
2:B:260:LEU:O	2:B:264:ARG:CD	2.49	0.60
2:B:440:THR:O	2:B:441:VAL:HG13	2.01	0.60
2:C:110:LEU:O	2:C:111:ALA:C	2.40	0.60
2:C:204:LEU:HD11	2:C:206:TYR:HB3	1.81	0.60
2:C:336:LEU:CD2	2:C:337:SER:O	2.50	0.60
2:C:342:LEU:O	2:C:342:LEU:CD2	2.48	0.60
2:C:421:MET:HE2	2:C:422:GLN:CB	2.30	0.60
2:C:144:ALA:O	2:C:145:PHE:HB2	2.01	0.60
1:D:219:ALA:O	1:D:220:TRP:CB	2.44	0.60
1:D:242:ALA:O	1:D:245:ILE:HG12	2.02	0.60
1:D:267:VAL:O	1:D:269:HIS:NE2	2.34	0.60
1:D:436:PRO:HA	1:D:839:ALA:HB2	1.84	0.60
1:D:842:PRO:O	1:D:844:VAL:HG22	2.01	0.60
2:E:214:CYS:SG	2:E:236:GLU:CD	2.79	0.60
2:F:206:TYR:N	2:F:206:TYR:CD2	2.69	0.60
1:A:1019:TRP:CD1	1:A:1020:ILE:HG22	2.36	0.60
1:A:1028:VAL:O	1:A:1031:GLU:O	2.18	0.60
1:A:448:ALA:CA	1:A:870:VAL:CG1	2.80	0.60
1:A:572:TRP:HE3	1:A:572:TRP:HA	1.66	0.60
1:A:606:THR:HG23	1:A:613:HIS:CA	2.29	0.60
1:A:869:ARG:O	1:A:872:SER:CB	2.48	0.60
2:B:304:TRP:HB2	2:B:337:SER:HB3	1.83	0.60
2:B:400:GLN:NE2	2:B:417:TYR:HB3	2.16	0.60
2:C:305:ASN:C	2:C:335:VAL:HG11	2.21	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:344:ARG:CA	2:C:347:LEU:HD13	2.31	0.60
2:C:372:LEU:CD1	2:C:374:LEU:HG	2.32	0.60
2:C:456:ARG:HE	2:C:456:ARG:N	1.99	0.60
2:C:471:LEU:CA	2:C:474:PHE:HD2	2.15	0.60
1:D:301:ILE:O	1:D:1024:ASP:CG	2.39	0.60
1:D:944:GLU:HG2	1:D:1071:ARG:CZ	2.31	0.60
1:D:1144:GLU:O	1:D:1145:ASP:HB2	2.01	0.60
1:D:1232:GLU:HB2	1:D:1234:ARG:HB2	1.84	0.60
1:D:194:ALA:O	1:D:195:LEU:HG	2.02	0.60
1:D:316:LYS:HD2	1:D:348:LEU:HD23	1.83	0.60
1:D:359:VAL:O	1:D:360:HIS:C	2.37	0.60
1:D:371:LYS:HD3	1:D:392:LEU:HD21	1.81	0.60
1:D:379:LYS:O	1:D:379:LYS:HD3	2.00	0.60
1:D:414:PHE:C	1:D:417:ARG:HB2	2.21	0.60
1:D:491:ASP:O	1:D:492:LEU:O	2.20	0.60
1:D:718:ALA:O	1:D:722:ARG:CD	2.48	0.60
1:D:86:GLN:C	1:D:87:ILE:HG12	2.21	0.60
2:E:325:ARG:HG2	2:E:330:ASN:CA	2.27	0.60
2:E:437:ILE:CD1	2:E:437:ILE:N	2.64	0.60
2:E:443:VAL:HA	2:E:453:ILE:HG22	1.84	0.60
2:E:93:SER:O	2:E:95:CYS:N	2.34	0.60
2:F:311:LEU:O	2:F:314:MET:HB2	2.01	0.60
2:F:340:GLY:O	2:F:341:ASP:O	2.18	0.60
2:F:386:ASP:HB3	2:F:442:LEU:CD2	2.26	0.60
2:F:430:SER:OG	2:F:431:LYS:N	2.34	0.60
2:F:461:THR:OG1	2:F:461:THR:O	2.20	0.60
1:A:1118:TRP:CE3	1:A:1118:TRP:HA	2.36	0.60
2:B:431:LYS:O	2:B:435:MET:N	2.32	0.60
2:C:121:PHE:O	2:C:123:GLU:N	2.35	0.60
2:C:187:HIS:HA	2:C:190:LEU:HD12	1.84	0.60
2:C:389:ARG:CG	2:C:390:GLY:N	2.65	0.60
1:D:666:GLN:HG2	1:D:669:MET:CE	2.31	0.60
2:B:217:PRO:O	2:B:231:ILE:HA	2.02	0.60
2:B:231:ILE:N	2:B:231:ILE:HD13	2.15	0.60
2:F:94:GLY:HA2	2:F:96:HIS:ND1	2.16	0.60
1:D:1029:GLN:HE21	1:D:1030:ARG:NH2	1.97	0.60
1:D:214:ALA:N	1:D:401:TRP:CH2	2.69	0.60
1:D:251:THR:O	1:D:254:SER:C	2.39	0.60
1:D:350:ILE:HG23	1:D:352:SER:CB	2.20	0.60
1:D:422:VAL:HG11	1:D:1111:LEU:CD1	2.31	0.60
1:D:597:ARG:HB3	1:D:616:GLU:CD	2.22	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:618:HIS:HB3	1:D:725:PRO:HB2	1.83	0.60
1:D:427:MET:O	1:D:846:THR:HG21	2.01	0.60
1:D:894:GLN:O	1:D:898:ILE:HG23	2.02	0.60
1:D:961:PHE:HD1	1:D:961:PHE:H	1.48	0.60
2:E:199:LEU:HD22	2:F:77:HIS:CD2	2.35	0.60
2:E:94:GLY:N	2:E:234:LYS:HE2	2.16	0.60
2:E:371:VAL:CG1	2:E:372:LEU:H	2.14	0.60
2:E:385:LEU:O	2:E:440:THR:HG23	2.01	0.60
2:F:260:LEU:O	2:F:264:ARG:HG3	2.02	0.60
1:A:1069:ILE:HG22	1:A:1070:PRO:CD	2.28	0.60
1:A:136:ASP:OD1	1:A:137:GLN:N	2.34	0.60
1:A:216:SER:O	1:A:217:PRO:O	2.19	0.60
1:A:271:VAL:O	1:A:275:ARG:HD3	2.01	0.60
1:A:386:ARG:HB2	1:A:390:GLN:HE22	1.67	0.60
1:A:400:VAL:HG12	1:A:404:HIS:CD2	2.35	0.60
1:A:570:PRO:HB2	1:A:572:TRP:CE3	2.37	0.60
1:A:844:VAL:HG13	1:A:844:VAL:O	2.02	0.60
1:A:811:GLN:NE2	1:A:859:TRP:HZ2	1.99	0.60
1:A:451:THR:HB	1:A:869:ARG:NH2	2.17	0.60
1:A:894:GLN:O	1:A:898:ILE:CG2	2.50	0.60
2:B:116:THR:HA	2:B:120:VAL:HB	1.84	0.60
2:C:402:LEU:HB3	2:C:406:LEU:HD21	1.84	0.60
2:B:447:THR:HA	2:B:450:ASN:OD1	2.02	0.60
1:D:1065:ALA:O	1:D:1067:SER:N	2.35	0.60
1:D:1027:LYS:HB2	1:D:1096:ARG:NE	2.17	0.60
1:D:136:ASP:O	1:D:140:ARG:HD3	2.02	0.60
1:D:656:TYR:O	1:D:744:ILE:HG13	2.02	0.60
1:D:954:ILE:HG22	1:D:955:TYR:CD2	2.37	0.60
2:F:134:LYS:CB	2:F:180:LYS:HE2	2.32	0.60
2:F:447:THR:O	2:F:452:LEU:CA	2.46	0.60
1:A:1120:PHE:O	1:A:1125:ILE:HD13	2.02	0.60
1:A:177:ARG:HB2	1:A:183:GLU:HB3	1.83	0.60
1:A:177:ARG:CB	1:A:183:GLU:HB3	2.32	0.60
1:A:188:ALA:O	1:A:189:ILE:HG12	2.01	0.60
1:A:232:ARG:HB3	1:A:239:LEU:CD2	2.08	0.60
1:A:297:MET:CE	1:A:411:LEU:HD22	2.32	0.60
1:A:572:TRP:CE3	1:A:572:TRP:HA	2.37	0.60
1:A:655:LEU:HB2	1:A:719:LEU:CB	2.26	0.60
1:A:655:LEU:HD22	1:A:722:ARG:CZ	2.32	0.60
1:A:823:ARG:O	1:A:824:ALA:HB3	2.02	0.60
2:C:395:LEU:HD22	2:C:443:VAL:O	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:187:HIS:O	2:E:190:LEU:HB2	2.01	0.60
1:D:1006:ARG:CA	1:D:1006:ARG:NH1	2.39	0.60
1:D:173:GLU:O	1:D:175:TRP:N	2.34	0.60
1:D:273:PHE:O	1:D:277:HIS:CD2	2.54	0.60
1:D:457:ARG:CA	1:D:460:LYS:HZ2	2.15	0.60
1:D:620:TRP:CE2	1:D:751:LYS:HB2	2.37	0.60
1:D:656:TYR:HD1	1:D:743:ASP:HB2	1.65	0.60
1:D:815:TRP:CE3	1:D:836:LEU:HB3	2.36	0.60
1:D:883:GLY:HA2	1:D:1143:GLU:OE1	2.02	0.60
2:E:119:VAL:O	2:E:123:GLU:HA	2.00	0.60
2:E:189:ALA:CB	2:E:240:VAL:HG11	2.32	0.60
2:F:267:ALA:O	2:F:270:PRO:HD3	2.01	0.60
1:A:1016:GLU:HB2	1:A:1020:ILE:HA	1.84	0.60
1:A:256:PRO:CD	1:A:257:THR:N	2.61	0.60
1:A:353:VAL:O	1:A:354:ASN:HB3	2.02	0.60
1:A:228:LEU:HD11	1:A:386:ARG:CB	2.31	0.60
1:A:410:GLN:C	1:A:412:PRO:HD2	2.22	0.60
1:A:481:GLU:O	1:A:484:TRP:HB2	2.02	0.60
1:A:573:TYR:C	1:A:574:ARG:HG3	2.22	0.60
1:A:612:LEU:CD2	1:A:620:TRP:CA	2.78	0.60
1:A:766:PHE:O	1:A:767:ALA:C	2.40	0.60
1:A:891:VAL:HA	1:A:1180:PHE:CE1	2.27	0.60
2:B:208:LEU:HD21	2:B:240:VAL:HB	1.84	0.60
2:B:259:ARG:C	2:B:261:GLN:N	2.51	0.60
2:C:290:TYR:CD1	2:C:299:LEU:HA	2.36	0.60
2:C:309:HIS:HA	2:C:312:LEU:CD1	2.32	0.60
2:E:388:GLY:CA	2:E:395:LEU:HD11	2.32	0.60
2:E:393:LEU:O	2:E:396:ARG:HG3	2.02	0.60
1:D:85:GLU:OE1	1:D:130:LEU:HD13	2.01	0.60
2:E:193:TYR:O	2:E:197:LEU:CB	2.50	0.60
1:A:225:SER:OG	1:A:386:ARG:CB	2.47	0.60
1:A:348:LEU:C	1:A:350:ILE:H	1.99	0.60
1:A:389:PHE:HD1	1:A:390:GLN:NE2	2.00	0.60
1:A:592:LEU:O	1:A:596:MET:HG2	2.02	0.60
1:A:612:LEU:CD2	1:A:614:TYR:H	2.14	0.60
1:A:616:GLU:O	1:A:617:ARG:HB2	2.01	0.60
1:A:659:HIS:NE2	1:A:720:THR:HG22	2.17	0.60
1:A:812:MET:HE3	1:A:813:VAL:HG13	1.83	0.60
1:A:844:VAL:HA	1:A:856:GLU:HG3	1.82	0.60
2:B:265:LYS:CD	2:B:349:TYR:OH	2.50	0.60
2:B:412:SER:O	2:B:413:VAL:CG2	2.48	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:463:LYS:H	2:B:463:LYS:CD	2.15	0.60
2:C:204:LEU:CD1	2:C:206:TYR:HB3	2.31	0.60
1:A:156:LEU:N	1:A:156:LEU:HD23	2.16	0.60
1:D:1002:GLU:CG	1:D:1003:TRP:N	2.64	0.59
1:D:853:ARG:HH12	1:D:1102:GLN:HA	1.67	0.59
1:D:1116:MET:HG2	1:D:1156:THR:HG21	1.84	0.59
1:D:407:PHE:CD1	1:D:408:GLN:N	2.70	0.59
1:D:444:TYR:OH	1:D:1206:MET:CB	2.49	0.59
1:D:549:LEU:HD23	1:D:552:LEU:HD12	1.84	0.59
1:D:815:TRP:CZ3	1:D:836:LEU:HB3	2.36	0.59
1:D:807:ARG:CB	1:D:858:THR:HG23	2.31	0.59
2:E:262:TRP:O	2:E:265:LYS:CB	2.46	0.59
2:E:380:PRO:HA	2:E:438:LEU:HD12	1.84	0.59
2:E:444:THR:OG1	2:E:446:THR:CG2	2.49	0.59
2:E:455:LEU:HD23	2:E:455:LEU:N	2.16	0.59
2:E:86:LEU:HB2	2:E:91:LEU:CD2	2.31	0.59
2:F:306:LEU:C	2:F:335:VAL:HB	2.22	0.59
2:F:342:LEU:CG	2:F:344:ARG:HB2	2.31	0.59
2:F:352:ASP:N	2:F:355:GLN:HB2	2.17	0.59
2:F:266:PHE:HB3	2:F:378:LEU:HD11	1.83	0.59
2:F:395:LEU:CD2	2:F:448:LEU:HD11	2.32	0.59
1:A:1058:PHE:O	1:A:1060:LYS:N	2.35	0.59
1:A:1165:ALA:HB2	1:A:1173:LEU:CG	2.32	0.59
1:A:1210:TYR:HE1	1:A:1213:PRO:CG	2.09	0.59
1:A:184:ALA:C	1:A:186:PRO:HD3	2.23	0.59
1:A:204:ALA:C	1:A:206:GLY:H	2.04	0.59
1:A:215:ILE:HB	1:A:401:TRP:CZ3	2.37	0.59
1:A:476:GLY:HA2	1:A:711:ALA:HB1	1.84	0.59
1:A:515:ILE:H	1:A:515:ILE:HD12	1.66	0.59
1:A:769:ASP:C	1:A:772:PRO:HD2	2.22	0.59
2:B:213:VAL:HA	2:B:235:THR:C	2.22	0.59
2:C:336:LEU:HD23	2:C:337:SER:C	2.21	0.59
2:C:235:THR:O	2:C:341:ASP:CG	2.41	0.59
2:C:342:LEU:HD21	2:C:345:GLY:H	1.65	0.59
2:C:354:PHE:O	2:C:355:GLN:C	2.40	0.59
2:C:372:LEU:HD11	2:C:436:SER:OG	2.00	0.59
2:C:388:GLY:N	2:C:443:VAL:O	2.32	0.59
2:C:456:ARG:HB2	2:C:463:LYS:O	2.02	0.59
1:D:665:LYS:C	1:D:669:MET:HG2	2.22	0.59
1:D:1010:LEU:N	1:D:1011:PRO:CD	2.65	0.59
1:D:1204:THR:O	1:D:1207:GLU:N	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:184:ALA:O	1:D:185:VAL:HG12	2.02	0.59
1:D:309:ARG:C	1:D:313:ILE:HG23	2.22	0.59
1:D:389:PHE:CA	1:D:393:MET:CG	2.72	0.59
1:D:752:LEU:HB3	1:D:753:PRO:HB3	1.83	0.59
1:D:992:LEU:HB3	1:D:996:ARG:CD	2.32	0.59
2:E:104:VAL:HG21	2:F:128:VAL:CA	2.32	0.59
2:F:346:MET:SD	2:F:346:MET:C	2.81	0.59
2:F:387:VAL:HG13	2:F:399:CYS:HB2	1.83	0.59
2:F:446:THR:CB	2:F:450:ASN:H	2.15	0.59
2:F:472:LYS:O	2:F:475:LEU:HG	2.02	0.59
1:A:1119:LEU:HD23	1:A:1122:GLU:HG3	1.85	0.59
1:A:1112:MET:HA	1:A:1156:THR:HG23	1.83	0.59
1:A:309:ARG:O	1:A:313:ILE:N	2.34	0.59
1:A:198:ASP:CB	1:A:400:VAL:HG11	2.32	0.59
1:A:371:LYS:HZ3	1:A:400:VAL:HG23	1.66	0.59
1:A:435:LEU:HD12	1:A:842:PRO:HD3	1.83	0.59
1:A:488:LEU:N	1:A:488:LEU:HD23	2.17	0.59
1:A:556:THR:HG22	2:C:467:HIS:NE2	2.17	0.59
1:A:570:PRO:HB2	1:A:572:TRP:CZ3	2.37	0.59
1:A:845:VAL:HG11	1:A:848:GLY:HA3	1.84	0.59
1:A:887:VAL:HG22	1:A:1146:ARG:NH1	2.14	0.59
1:A:948:ILE:HG13	1:A:949:PHE:HD2	1.66	0.59
1:D:504:LYS:CB	1:D:523:ASP:O	2.47	0.59
1:D:1190:ARG:NE	1:D:1195:MET:HG2	2.17	0.59
1:D:1207:GLU:OE2	1:D:1210:TYR:CE2	2.54	0.59
1:D:285:GLN:NE2	1:D:1128:ARG:CG	2.61	0.59
1:D:257:THR:OG1	1:D:287:SER:CB	2.50	0.59
1:D:798:ILE:HG22	1:D:869:ARG:HH12	1.61	0.59
2:E:380:PRO:O	2:E:382:LYS:HD2	2.02	0.59
2:E:442:LEU:C	2:E:442:LEU:HD13	2.23	0.59
2:E:447:THR:HG21	2:E:453:ILE:CG2	2.24	0.59
2:F:264:ARG:HA	2:F:270:PRO:HB3	1.84	0.59
1:A:1153:LEU:CD1	1:A:1157:ASN:ND2	2.65	0.59
1:A:1190:ARG:NH2	1:A:1197:CYS:HB2	2.18	0.59
1:A:245:ILE:O	1:A:248:GLU:HG2	2.02	0.59
1:A:665:LYS:HG2	1:A:666:GLN:H	1.66	0.59
1:A:275:ARG:HG2	1:A:843:GLN:CB	2.32	0.59
2:B:440:THR:O	2:B:455:LEU:HB2	2.02	0.59
2:B:79:LEU:CA	2:B:100:GLY:HA3	2.33	0.59
2:C:208:LEU:CD2	2:C:242:PHE:CE2	2.84	0.59
2:C:456:ARG:CB	2:C:463:LYS:O	2.50	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:186:LEU:O	2:F:189:ALA:HB3	2.01	0.59
1:D:113:TRP:CE3	1:D:116:PRO:HG2	2.36	0.59
1:D:1193:VAL:CG2	1:D:1194:THR:N	2.36	0.59
1:D:192:GLU:OE1	1:D:193:ARG:NH1	2.35	0.59
1:D:247:LEU:CA	1:D:815:TRP:CD1	2.84	0.59
1:D:808:ILE:HG21	1:D:874:LEU:HD23	1.83	0.59
1:D:887:VAL:H	1:D:1146:ARG:NH2	1.98	0.59
1:D:930:ASP:OD2	1:D:930:ASP:C	2.40	0.59
2:F:78:PHE:C	2:F:100:GLY:H	2.06	0.59
2:E:101:PRO:CG	2:F:128:VAL:HG21	2.32	0.59
2:F:193:TYR:HB2	2:F:242:PHE:CZ	2.38	0.59
2:F:343:ASP:C	2:F:347:LEU:CD1	2.70	0.59
1:A:115:GLN:O	1:A:118:VAL:HG13	2.02	0.59
1:A:1232:GLU:HG3	1:A:1234:ARG:NE	2.17	0.59
1:A:275:ARG:NH2	1:A:844:VAL:C	2.55	0.59
1:A:869:ARG:O	1:A:872:SER:N	2.30	0.59
2:B:246:ARG:C	2:B:248:SER:H	2.06	0.59
2:B:344:ARG:N	2:B:344:ARG:CD	2.65	0.59
2:B:372:LEU:O	2:B:458:ARG:HD2	2.01	0.59
2:C:310:GLU:OE1	2:C:310:GLU:N	2.36	0.59
2:C:239:LEU:HG	2:C:336:LEU:CD1	2.33	0.59
2:C:353:SER:OG	2:C:354:PHE:N	2.33	0.59
1:D:97:GLU:O	1:D:101:ARG:HG3	2.02	0.59
2:B:83:LYS:N	2:B:83:LYS:HD2	2.15	0.59
1:D:1015:THR:CG2	1:D:1026:ARG:CB	2.80	0.59
1:D:866:ARG:NE	1:D:1069:ILE:HB	2.17	0.59
1:D:1198:LYS:N	1:D:1204:THR:HG21	2.18	0.59
1:D:127:LEU:HD23	1:D:128:PRO:CD	2.32	0.59
1:D:182:GLY:O	1:D:183:GLU:HB2	2.03	0.59
1:D:289:MET:HE2	1:D:290:ARG:H	1.68	0.59
1:D:416:GLU:C	1:D:418:CYS:H	2.03	0.59
1:D:458:GLU:N	1:D:460:LYS:NZ	2.50	0.59
1:D:770:PHE:HZ	1:D:968:GLN:O	1.85	0.59
1:D:793:GLU:O	1:D:797:MET:HG3	2.02	0.59
1:D:822:PRO:O	1:D:825:VAL:HB	2.03	0.59
1:D:938:THR:O	1:D:939:VAL:HG23	2.03	0.59
2:E:127:PRO:HG2	2:F:101:PRO:CA	2.33	0.59
2:E:211:ILE:HD13	2:E:211:ILE:N	2.18	0.59
2:E:456:ARG:HD2	2:E:461:THR:C	2.23	0.59
2:F:446:THR:C	2:F:449:GLU:H	2.06	0.59
1:A:1006:ARG:NH2	1:A:1010:LEU:CD1	2.57	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1200:PRO:HB2	1:A:1203:PRO:CD	2.32	0.59
1:A:608:ASP:CA	1:A:779:LEU:HD22	2.32	0.59
2:B:132:HIS:HA	2:B:182:ARG:NE	2.17	0.59
2:B:413:VAL:HG12	2:B:414:TRP:H	1.66	0.59
2:C:454:HIS:HD2	2:C:472:LYS:HE3	1.68	0.59
2:C:69:LEU:HD23	2:C:351:TYR:CD1	2.37	0.59
1:D:586:THR:N	1:D:587:PRO:HD3	2.16	0.59
1:D:1069:ILE:HG12	1:D:1070:PRO:CD	2.32	0.59
1:D:1175:GLN:HG3	1:D:1176:SER:N	2.16	0.59
1:D:872:SER:OG	1:D:1200:PRO:N	2.35	0.59
1:D:873:GLU:OE1	1:D:1203:PRO:CG	2.50	0.59
1:D:359:VAL:C	1:D:361:ARG:N	2.49	0.59
1:D:403:THR:HA	1:D:406:VAL:HG13	1.85	0.59
1:D:76:ILE:CA	1:D:911:HIS:NE2	2.66	0.59
2:E:193:TYR:HE2	2:E:322:LEU:HD22	1.66	0.59
2:E:447:THR:HG23	2:E:452:LEU:H	1.68	0.59
2:E:78:PHE:CD2	2:E:102:LEU:CD1	2.85	0.59
1:D:472:GLN:HB3	2:F:460:THR:C	2.23	0.59
1:A:1131:ILE:CG2	1:A:1138:ARG:N	2.57	0.59
1:A:1219:ASP:O	1:A:1223:ILE:HG22	2.02	0.59
1:A:974:THR:CG2	1:A:975:GLN:H	2.16	0.59
2:B:69:LEU:HD13	2:B:73:CYS:SG	2.43	0.59
2:C:206:TYR:O	2:C:241:TRP:HA	2.02	0.59
2:C:307:GLY:CA	2:C:335:VAL:HG12	2.32	0.59
2:C:197:LEU:HD11	2:C:322:LEU:CD2	2.32	0.59
2:C:341:ASP:OD1	2:C:342:LEU:N	2.30	0.59
1:D:1015:THR:CG2	1:D:1026:ARG:HB2	2.32	0.59
1:D:306:SER:CB	1:D:1058:PHE:CE2	2.86	0.59
1:D:875:LYS:HG3	1:D:1189:LEU:O	2.01	0.59
1:D:1148:ARG:CZ	1:D:1231:LEU:HD22	2.33	0.59
1:D:167:PRO:CD	1:D:168:ALA:H	2.16	0.59
1:D:245:ILE:CG1	1:D:246:PRO:HD3	2.33	0.59
1:D:312:TRP:CA	1:D:316:LYS:HD3	2.32	0.59
1:D:479:TYR:HD1	1:D:480:LYS:HG3	1.65	0.59
1:D:769:ASP:O	1:D:773:LYS:HG2	2.02	0.59
1:D:775:GLU:CG	1:D:780:GLN:HG3	2.20	0.59
1:D:795:ASN:N	1:D:795:ASN:ND2	2.34	0.59
2:E:106:LEU:O	2:E:109:ASN:OD1	2.20	0.59
2:E:205:PRO:HG2	2:E:241:TRP:HE1	1.67	0.59
2:E:475:LEU:HB3	2:E:479:ILE:CD1	2.33	0.59
2:F:251:TRP:HE3	2:F:251:TRP:HA	1.66	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:306:LEU:HD21	2:F:310:GLU:CG	2.33	0.59
2:F:343:ASP:HB2	2:F:347:LEU:HD11	1.83	0.59
2:F:428:LEU:HB3	2:F:432:TYR:CE1	2.38	0.59
2:F:432:TYR:CB	2:F:437:ILE:HD11	2.31	0.59
2:F:67:GLU:O	2:F:71:GLU:N	2.33	0.59
1:A:851:THR:OG1	1:A:1102:GLN:HB3	2.02	0.59
1:A:868:ASP:CG	1:A:1199:THR:HG21	2.23	0.59
1:A:175:TRP:CD1	1:A:176:THR:N	2.71	0.59
1:A:365:TYR:CD2	1:A:365:TYR:N	2.70	0.59
1:A:83:LEU:O	1:A:84:HIS:C	2.41	0.59
2:B:204:LEU:CG	2:B:325:ARG:HH21	2.05	0.59
1:D:509:THR:HG21	1:D:512:LYS:HD3	1.85	0.59
1:A:634:LEU:HD12	1:A:638:THR:CG2	2.33	0.59
1:A:776:ASP:C	1:A:776:ASP:OD1	2.41	0.59
2:E:282:GLU:HG3	2:E:309:HIS:CE1	2.37	0.59
2:B:82:SER:HB3	2:B:85:GLN:HG2	1.85	0.59
2:E:466:MET:HG2	2:E:467:HIS:H	1.67	0.59
1:D:176:THR:HG23	1:D:177:ARG:N	2.16	0.59
1:D:279:ARG:CZ	1:D:433:SER:O	2.51	0.59
1:D:460:LYS:O	1:D:461:LYS:C	2.38	0.59
1:D:646:VAL:HG12	1:D:647:CYS:N	2.18	0.59
1:D:742:VAL:CG2	1:D:748:TRP:CE3	2.86	0.59
1:D:868:ASP:OD1	1:D:1199:THR:HG21	2.03	0.59
1:D:898:ILE:CG1	1:D:899:ALA:N	2.66	0.59
2:F:208:LEU:O	2:F:208:LEU:CD1	2.48	0.59
2:F:208:LEU:HD11	2:F:240:VAL:C	2.23	0.59
1:D:472:GLN:CG	2:F:460:THR:OG1	2.51	0.59
1:A:996:ARG:CD	1:A:1000:GLU:O	2.51	0.59
1:A:1003:TRP:HA	1:A:1006:ARG:HB2	1.85	0.59
1:A:1231:LEU:HD12	1:A:1235:SER:CB	2.33	0.59
1:A:154:ALA:O	1:A:158:LEU:HD13	2.02	0.59
1:A:213:VAL:HG12	1:A:401:TRP:CE2	2.36	0.59
1:A:282:TYR:CZ	1:A:291:PHE:CE1	2.91	0.59
1:A:460:LYS:O	1:A:464:MET:N	2.36	0.59
1:A:605:LEU:HB2	1:A:614:TYR:CD1	2.38	0.59
1:A:601:LYS:HZ1	1:A:721:ALA:C	2.05	0.59
1:A:728:THR:CG2	1:A:729:GLN:H	2.03	0.59
1:A:875:LYS:O	1:A:1189:LEU:HB2	2.02	0.59
2:B:106:LEU:O	2:B:110:LEU:N	2.32	0.59
2:C:344:ARG:CA	2:C:347:LEU:HB2	2.25	0.59
2:C:393:LEU:HA	2:C:396:ARG:HG3	1.85	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1008:LEU:HD22	1:D:1008:LEU:C	2.22	0.59
1:D:1057:MET:HB3	1:D:1064:ILE:CD1	2.26	0.59
1:D:1069:ILE:CB	1:D:1070:PRO:CD	2.77	0.59
1:D:1109:LEU:HD21	1:D:1137:VAL:HG21	1.85	0.59
1:D:86:GLN:HE21	1:D:132:GLY:HA3	1.68	0.59
1:D:312:TRP:CE3	1:D:316:LYS:HD2	2.38	0.59
1:D:348:LEU:HD21	1:D:350:ILE:CG1	2.23	0.59
1:D:597:ARG:NH1	1:D:725:PRO:N	2.51	0.59
1:D:907:PHE:HE2	1:D:913:CYS:HG	1.50	0.59
1:D:973:LEU:HB3	1:D:977:GLU:OE1	2.01	0.59
1:D:993:ARG:C	1:D:996:ARG:CD	2.70	0.59
2:E:199:LEU:CG	2:E:200:VAL:N	2.65	0.59
2:E:247:THR:HB	2:E:251:TRP:NE1	2.18	0.59
2:E:379:ALA:HB3	2:E:438:LEU:HD21	1.85	0.59
2:E:76:ARG:CG	2:E:435:MET:SD	2.90	0.59
2:F:442:LEU:O	2:F:455:LEU:CD1	2.50	0.59
1:A:1060:LYS:HE2	1:A:1064:ILE:HG23	1.85	0.59
1:A:1113:LEU:HB2	1:A:1117:LYS:NZ	2.18	0.59
1:A:157:LEU:HD21	1:A:194:ALA:O	2.03	0.59
1:A:245:ILE:HG13	1:A:246:PRO:CD	2.22	0.59
1:A:560:PRO:HB2	2:C:452:LEU:HD21	1.85	0.59
1:A:464:MET:SD	1:A:579:ARG:HB2	2.43	0.59
1:A:617:ARG:HB3	1:A:763:GLY:O	2.03	0.59
1:A:604:ALA:CB	1:A:717:LEU:CD1	2.80	0.59
2:B:87:SER:C	2:B:89:ASP:H	2.06	0.59
2:C:466:MET:CA	2:C:471:LEU:HD22	2.33	0.59
2:E:218:VAL:HB	2:E:230:SER:OG	2.02	0.59
2:B:411:ILE:HD12	2:B:411:ILE:N	2.18	0.59
1:D:1165:ALA:HB3	1:D:1166:TYR:CD1	2.37	0.59
1:D:877:MET:HE1	1:D:1203:PRO:C	2.22	0.59
1:D:297:MET:HG3	1:D:415:LEU:CD1	2.32	0.59
1:D:482:ASP:CB	1:D:483:PRO:HD2	2.32	0.59
1:D:742:VAL:CG1	1:D:743:ASP:N	2.65	0.59
1:D:795:ASN:C	1:D:798:ILE:HG12	2.22	0.59
2:F:304:TRP:N	2:F:337:SER:CB	2.52	0.59
2:F:371:VAL:HG23	2:F:433:ASP:CA	2.32	0.59
1:A:1147:TYR:O	1:A:1150:ALA:HB3	2.03	0.59
1:A:868:ASP:CB	1:A:1199:THR:HB	2.33	0.59
1:A:1179:PHE:O	1:A:1220:ILE:HD11	2.03	0.59
1:A:173:GLU:OE2	1:A:223:TRP:CZ2	2.47	0.59
1:A:509:THR:CB	1:A:513:LEU:HD21	2.33	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:853:ARG:NH1	1:A:1102:GLN:O	2.36	0.59
1:A:896:LEU:HD21	1:A:951:TYR:CE2	2.38	0.59
2:B:182:ARG:HB3	2:B:215:PHE:HD1	1.67	0.59
2:C:114:TRP:NE1	2:C:259:ARG:NH2	2.51	0.59
2:C:361:PHE:HD2	2:C:362:THR:HG23	1.66	0.59
1:D:516:GLU:HG2	1:D:565:HIS:NE2	2.17	0.59
1:D:514:PRO:O	1:D:568:GLY:HA3	2.03	0.59
2:C:81:GLY:HA3	2:C:97:PRO:CD	2.33	0.59
2:F:382:LYS:O	2:F:438:LEU:HB2	2.02	0.59
1:D:502:LYS:CG	1:D:506:GLU:HB3	2.33	0.59
2:C:67:GLU:O	2:C:71:GLU:HB3	2.02	0.59
2:C:229:LYS:O	2:C:230:SER:HB2	2.02	0.59
1:D:1056:GLU:HB3	1:D:1057:MET:HE3	1.83	0.58
1:D:1188:CYS:SG	1:D:1189:LEU:N	2.75	0.58
1:D:291:PHE:O	1:D:293:ASP:N	2.36	0.58
1:D:213:VAL:HG13	1:D:401:TRP:NE1	2.17	0.58
1:D:439:GLN:NE2	1:D:837:TYR:HE2	2.01	0.58
1:D:474:LEU:CD1	1:D:474:LEU:N	2.65	0.58
1:D:620:TRP:N	1:D:620:TRP:CD1	2.69	0.58
1:D:967:MET:CE	1:D:978:ALA:HB2	2.33	0.58
1:D:981:LYS:O	1:D:985:MET:HG3	2.03	0.58
2:E:380:PRO:C	2:E:382:LYS:N	2.55	0.58
2:F:131:LEU:O	2:F:133:HIS:HB2	2.03	0.58
2:F:239:LEU:H	2:F:338:VAL:CB	2.14	0.58
2:F:294:PRO:HD2	2:F:295:TRP:CZ3	2.38	0.58
2:F:475:LEU:HA	2:F:478:TYR:HB2	1.85	0.58
1:A:1015:THR:C	1:A:1016:GLU:OE2	2.41	0.58
1:A:426:GLY:HA2	1:A:1117:LYS:HZ1	1.66	0.58
1:A:201:VAL:HG13	1:A:206:GLY:O	2.03	0.58
1:A:245:ILE:O	1:A:248:GLU:CG	2.51	0.58
1:A:371:LYS:HD3	1:A:396:CYS:CA	2.21	0.58
1:A:996:ARG:NH1	1:A:1003:TRP:HB3	2.17	0.58
2:B:203:ARG:HH11	2:B:206:TYR:HB3	1.68	0.58
2:B:385:LEU:CB	2:B:441:VAL:HG23	2.32	0.58
2:C:128:VAL:CG1	2:C:192:HIS:HD2	2.16	0.58
2:C:344:ARG:CG	2:C:347:LEU:HD22	2.33	0.58
2:C:351:TYR:HD1	2:C:355:GLN:HA	1.67	0.58
2:C:394:GLU:C	2:C:397:GLN:HB3	2.23	0.58
2:C:484:ASN:C	2:C:485:VAL:HG13	2.23	0.58
2:C:74:GLN:HG3	2:C:75:ARG:N	2.17	0.58
1:D:567:PRO:O	1:D:569:HIS:N	2.36	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:81:GLY:HA3	2:C:97:PRO:HD2	1.84	0.58
1:A:750:PHE:N	1:A:750:PHE:CD1	2.71	0.58
1:D:498:LYS:HZ3	1:D:521:PRO:HB3	1.68	0.58
1:A:937:THR:HA	1:A:940:GLY:HA2	1.84	0.58
2:B:122:ARG:O	2:B:124:GLN:N	2.30	0.58
1:D:1067:SER:HB3	1:D:1073:PRO:HG2	1.85	0.58
1:D:1196:ASP:OD2	1:D:1204:THR:HG21	2.03	0.58
1:D:177:ARG:HG2	1:D:218:SER:OG	2.03	0.58
1:D:371:LYS:HZ1	1:D:396:CYS:HB2	1.66	0.58
1:D:456:GLN:O	1:D:460:LYS:HD3	2.03	0.58
1:D:607:TRP:O	1:D:611:PRO:O	2.21	0.58
1:D:737:GLY:N	1:D:750:PHE:HZ	1.94	0.58
1:D:802:ARG:HG3	1:D:803:ASN:OD1	2.02	0.58
1:D:247:LEU:CA	1:D:815:TRP:HD1	2.16	0.58
1:D:829:PRO:HG3	1:D:831:TYR:CZ	2.38	0.58
2:E:79:LEU:CA	2:E:100:GLY:N	2.65	0.58
2:E:369:ARG:CA	2:E:430:SER:HB3	2.33	0.58
2:F:254:PHE:CE2	2:F:255:TRP:CE2	2.90	0.58
2:F:396:ARG:C	2:F:396:ARG:HD2	2.23	0.58
2:F:455:LEU:HB3	2:F:456:ARG:HE	1.66	0.58
1:A:886:LEU:C	1:A:1146:ARG:CZ	2.72	0.58
1:A:445:LEU:O	1:A:449:GLN:HG3	2.03	0.58
1:A:627:ARG:O	1:A:627:ARG:HG3	2.03	0.58
1:A:655:LEU:O	1:A:659:HIS:CB	2.48	0.58
1:A:80:SER:OG	1:A:81:ARG:N	2.35	0.58
1:A:296:SER:CB	1:A:847:ALA:HB3	2.30	0.58
1:A:877:MET:HE2	1:A:877:MET:O	2.03	0.58
1:A:76:ILE:CB	1:A:911:HIS:CE1	2.82	0.58
2:B:269:SER:CB	2:B:352:ASP:OD1	2.51	0.58
2:C:287:ASN:O	2:C:302:THR:O	2.21	0.58
2:C:403:PHE:HD1	2:C:417:TYR:HH	1.48	0.58
2:C:452:LEU:HD12	2:C:453:ILE:CG1	2.33	0.58
1:A:526:ASP:OD2	1:A:527:GLN:N	2.36	0.58
1:D:1069:ILE:CG2	1:D:1070:PRO:CD	2.61	0.58
1:D:1141:VAL:HG11	1:D:1146:ARG:HA	1.86	0.58
1:D:1151:LEU:CD2	1:D:1223:ILE:HD11	2.33	0.58
1:D:131:TYR:CB	1:D:138:HIS:HE1	2.13	0.58
1:D:375:GLU:O	1:D:378:VAL:N	2.34	0.58
1:D:617:ARG:O	1:D:762:VAL:CG2	2.49	0.58
1:D:801:TRP:O	1:D:804:ALA:N	2.36	0.58
1:D:91:GLY:O	1:D:92:GLY:C	2.42	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:102:LEU:CD1	2:E:102:LEU:C	2.71	0.58
2:E:123:GLU:HB3	2:F:403:PHE:CZ	2.39	0.58
2:E:239:LEU:CD2	2:E:338:VAL:HG21	2.34	0.58
2:F:130:ALA:HB1	2:F:182:ARG:NH1	2.17	0.58
2:F:241:TRP:HZ3	2:F:255:TRP:CZ2	2.21	0.58
2:F:368:HIS:CE1	2:F:370:LYS:HD2	2.37	0.58
2:F:388:GLY:O	2:F:444:THR:HA	2.03	0.58
2:F:456:ARG:HE	2:F:456:ARG:N	2.00	0.58
2:F:466:MET:CE	2:F:471:LEU:HB3	2.33	0.58
1:A:1046:GLU:OE1	1:A:1046:GLU:HA	2.03	0.58
1:A:167:PRO:O	1:A:168:ALA:C	2.42	0.58
1:A:371:LYS:HZ1	1:A:400:VAL:HG23	1.65	0.58
1:A:297:MET:HE3	1:A:411:LEU:HD13	1.84	0.58
1:A:579:ARG:HH11	1:A:579:ARG:HG2	1.68	0.58
1:A:606:THR:N	1:A:613:HIS:O	2.36	0.58
1:A:752:LEU:HB3	1:A:753:PRO:CA	2.33	0.58
2:C:201:ASN:C	2:C:203:ARG:N	2.56	0.58
2:C:354:PHE:HE1	2:C:370:LYS:CB	2.16	0.58
2:C:398:VAL:CA	2:C:401:GLY:H	2.15	0.58
2:C:405:GLU:OE1	2:C:406:LEU:HD12	2.03	0.58
2:E:390:GLY:H	2:E:395:LEU:HG	1.64	0.58
2:B:445:GLU:C	2:B:447:THR:H	2.05	0.58
2:B:133:HIS:CE1	2:C:231:ILE:CG2	2.86	0.58
1:A:976:GLN:O	1:A:979:ALA:HB3	2.02	0.58
1:D:1138:ARG:NH2	1:D:1191:LYS:NZ	2.51	0.58
1:D:259:ARG:O	1:D:260:ASP:OD2	2.21	0.58
1:D:401:TRP:CD1	1:D:404:HIS:HD2	2.21	0.58
1:D:597:ARG:CA	1:D:616:GLU:OE1	2.51	0.58
1:D:728:THR:HG23	1:D:741:ASP:OD2	2.04	0.58
1:D:868:ASP:O	1:D:869:ARG:HB2	2.02	0.58
1:D:916:PHE:HD1	1:D:916:PHE:O	1.86	0.58
1:D:995:TYR:CE1	1:D:1004:LEU:CB	2.62	0.58
2:E:130:ALA:HB2	2:E:210:GLN:NE2	2.19	0.58
2:E:72:ILE:HD11	2:E:370:LYS:HE2	1.85	0.58
2:F:208:LEU:CD2	2:F:242:PHE:CD2	2.87	0.58
2:F:400:GLN:NE2	2:F:404:ASN:CG	2.56	0.58
1:A:1014:ARG:O	1:A:1015:THR:HG23	2.03	0.58
1:A:1020:ILE:CG1	1:A:1021:SER:N	2.66	0.58
1:A:1123:PHE:C	1:A:1125:ILE:N	2.56	0.58
1:A:1209:ARG:HH12	2:C:256:LEU:CB	2.16	0.58
1:A:291:PHE:O	1:A:292:LEU:HB2	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:312:TRP:O	1:A:316:LYS:HB3	2.03	0.58
1:A:374:ARG:NE	1:A:392:LEU:CD1	2.62	0.58
1:A:385:ILE:HG23	1:A:386:ARG:N	2.17	0.58
1:A:616:GLU:HG3	1:A:724:GLY:O	2.03	0.58
1:A:744:ILE:CB	1:A:745:PRO:CD	2.74	0.58
1:A:74:LEU:C	1:A:76:ILE:H	2.05	0.58
1:A:79:LEU:CB	1:A:89:GLY:HA2	2.28	0.58
1:A:929:THR:HG23	1:A:947:LYS:HD3	1.85	0.58
1:A:958:GLY:CA	1:A:986:TYR:CE2	2.86	0.58
2:B:248:SER:O	2:B:252:LEU:HD13	2.03	0.58
2:C:383:VAL:HG22	2:C:384:ALA:N	2.18	0.58
2:C:435:MET:HE2	2:C:435:MET:CA	2.23	0.58
2:C:448:LEU:HA	2:C:451:GLY:HA2	1.83	0.58
2:E:88:ARG:H	2:E:88:ARG:HD3	1.66	0.58
2:C:59:ALA:HA	2:C:62:SER:OG	2.02	0.58
1:A:443:ARG:O	1:A:447:GLU:HG3	2.02	0.58
1:D:1068:ASP:O	1:D:1069:ILE:O	2.21	0.58
1:D:249:VAL:CG2	1:D:251:THR:HG22	2.33	0.58
1:D:223:TRP:CZ3	1:D:250:PRO:HB2	2.38	0.58
1:D:449:GLN:O	1:D:452:TYR:CD2	2.56	0.58
2:E:429:TYR:O	2:E:431:LYS:O	2.20	0.58
2:F:241:TRP:HB2	2:F:336:LEU:HB2	1.82	0.58
2:F:414:TRP:CG	2:F:414:TRP:O	2.56	0.58
1:A:228:LEU:CD2	1:A:228:LEU:C	2.69	0.58
1:A:316:LYS:HZ2	1:A:350:ILE:CG2	2.17	0.58
1:A:376:LEU:C	1:A:378:VAL:N	2.56	0.58
1:A:572:TRP:HB2	2:C:477:LYS:HE2	1.85	0.58
1:A:732:TYR:CE1	1:A:755:LYS:HG2	2.38	0.58
1:A:660:CYS:CB	1:A:745:PRO:HG3	2.33	0.58
2:B:309:HIS:HB3	2:B:312:LEU:HD23	1.78	0.58
2:B:385:LEU:HB3	2:B:441:VAL:CG2	2.33	0.58
2:C:456:ARG:HB2	2:C:463:LYS:HB3	1.84	0.58
2:C:88:ARG:HG2	2:C:89:ASP:N	2.17	0.58
2:C:146:ARG:CB	2:C:228:VAL:HG11	2.33	0.58
2:F:274:SER:OG	2:F:275:SER:N	2.35	0.58
1:D:857:PRO:C	1:D:1062:GLU:OE2	2.42	0.58
1:D:265:LEU:HG	1:D:266:VAL:C	2.24	0.58
1:D:381:THR:HG23	1:D:382:MET:N	2.19	0.58
1:D:401:TRP:O	1:D:405:GLU:HG2	2.03	0.58
1:D:415:LEU:C	1:D:417:ARG:N	2.57	0.58
1:D:472:GLN:O	2:F:461:THR:HG21	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:652:ILE:HG22	1:D:656:TYR:HE2	1.67	0.58
1:D:272:SER:CB	1:D:843:GLN:CD	2.68	0.58
1:D:948:ILE:CG2	1:D:1071:ARG:CB	2.80	0.58
2:E:377:CYS:C	2:E:380:PRO:CD	2.67	0.58
2:E:379:ALA:C	2:E:438:LEU:HD11	2.24	0.58
2:E:425:LEU:N	2:E:428:LEU:HD11	2.17	0.58
1:A:1131:ILE:HG12	1:A:1133:ILE:H	1.68	0.58
1:A:886:LEU:N	1:A:1146:ARG:CZ	2.67	0.58
1:A:1154:GLN:NE2	1:A:1220:ILE:HG13	2.19	0.58
1:A:77:GLN:HB2	1:A:1174:PRO:CB	2.34	0.58
1:A:257:THR:OG1	1:A:258:GLN:N	2.35	0.58
1:A:197:PHE:HB3	1:A:277:HIS:CD2	2.39	0.58
1:A:296:SER:HB2	1:A:846:THR:O	2.02	0.58
1:A:214:ALA:HA	1:A:401:TRP:HZ2	1.62	0.58
1:A:588:GLY:HA3	1:A:806:LYS:NZ	2.18	0.58
1:A:930:ASP:CA	1:A:933:SER:HB3	2.31	0.58
1:A:948:ILE:HG13	1:A:949:PHE:CD2	2.38	0.58
2:C:452:LEU:CD1	2:C:453:ILE:H	2.16	0.58
2:B:451:GLY:C	2:B:452:LEU:HD22	2.24	0.58
2:B:133:HIS:NE2	2:C:231:ILE:HG22	2.19	0.58
2:E:278:CYS:O	2:E:286:GLY:O	2.20	0.58
1:D:872:SER:HG	1:D:1200:PRO:N	2.02	0.58
1:D:357:ALA:O	1:D:358:GLU:C	2.40	0.58
1:D:464:MET:O	1:D:467:ALA:HB3	2.03	0.58
1:D:601:LYS:HZ1	1:D:722:ARG:H	1.51	0.58
1:D:744:ILE:HB	1:D:745:PRO:CD	2.34	0.58
1:D:823:ARG:NH1	1:D:824:ALA:HB2	2.18	0.58
1:D:279:ARG:CB	1:D:841:LEU:CB	2.82	0.58
1:D:273:PHE:N	1:D:843:GLN:OE1	2.37	0.58
2:E:456:ARG:HG2	2:E:457:SER:N	2.16	0.58
2:E:104:VAL:HG21	2:F:128:VAL:HA	1.84	0.58
2:F:323:HIS:C	2:F:331:VAL:O	2.41	0.58
2:F:435:MET:HE2	2:F:435:MET:CA	2.26	0.58
1:A:1006:ARG:CZ	1:A:1010:LEU:CD1	2.81	0.58
1:A:190:PRO:C	1:A:192:GLU:HG3	2.24	0.58
1:A:220:TRP:NE1	1:A:221:TYR:CD1	2.72	0.58
1:A:232:ARG:HD2	1:A:239:LEU:HD23	1.86	0.58
1:A:605:LEU:H	1:A:717:LEU:HD21	1.69	0.58
1:A:659:HIS:HD2	1:A:720:THR:HB	1.68	0.58
1:A:622:TYR:N	1:A:747:CYS:O	2.37	0.58
1:A:884:TYR:HB3	1:A:1140:LEU:HD21	1.84	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:914:THR:O	1:A:915:ALA:HB3	2.04	0.58
2:B:396:ARG:HD2	2:B:418:LEU:N	2.19	0.58
2:B:417:TYR:C	2:B:419:GLU:N	2.54	0.58
2:C:423:SER:CB	2:C:428:LEU:HD21	2.33	0.58
2:C:484:ASN:C	2:C:485:VAL:CG1	2.72	0.58
2:C:68:ALA:O	2:C:72:ILE:HG12	2.04	0.58
1:A:637:GLY:O	1:A:641:GLU:HG2	2.03	0.58
1:A:739:TYR:O	1:A:740:ASN:HB3	2.02	0.58
2:E:229:LYS:HG2	2:E:231:ILE:HG13	1.84	0.58
1:D:252:GLY:O	1:D:253:ALA:CB	2.49	0.58
1:D:312:TRP:O	1:D:316:LYS:HD3	2.04	0.58
1:D:371:LYS:O	1:D:372:GLU:HB2	2.03	0.58
1:D:441:TRP:NE1	1:D:880:ALA:N	2.47	0.58
1:D:798:ILE:CG2	1:D:869:ARG:CZ	2.80	0.58
1:D:900:ALA:C	1:D:915:ALA:CB	2.71	0.58
1:D:949:PHE:CD1	1:D:950:ASN:N	2.72	0.58
1:D:972:ARG:C	1:D:974:THR:N	2.55	0.58
2:E:104:VAL:HG21	2:F:128:VAL:C	2.24	0.58
2:F:342:LEU:HD22	2:F:342:LEU:N	2.19	0.58
2:F:458:ARG:HH11	2:F:458:ARG:HG3	1.69	0.58
2:F:474:PHE:CE2	2:F:475:LEU:HD23	2.39	0.58
1:A:110:HIS:O	1:A:112:LEU:CD1	2.51	0.58
1:A:296:SER:HB3	1:A:846:THR:O	2.03	0.58
1:A:389:PHE:HD2	1:A:393:MET:HG2	1.68	0.58
1:A:392:LEU:CD2	1:A:393:MET:HE3	2.33	0.58
1:A:588:GLY:CA	1:A:806:LYS:HZ1	2.16	0.58
1:A:623:LEU:O	1:A:625:PRO:CD	2.52	0.58
1:A:76:ILE:CG1	1:A:77:GLN:HG3	2.22	0.58
1:A:788:GLY:HA2	1:A:791:ALA:HB3	1.86	0.58
2:B:323:HIS:CB	2:B:331:VAL:O	2.52	0.58
2:C:444:THR:N	2:C:447:THR:CG2	2.67	0.58
2:C:454:HIS:CB	2:C:468:ILE:HD12	2.33	0.58
1:D:670:PRO:CG	1:D:673:ALA:HB2	2.20	0.58
2:B:475:LEU:O	2:B:479:ILE:HD13	2.04	0.58
1:D:1207:GLU:CA	1:D:1210:TYR:CD2	2.79	0.58
1:D:169:TRP:O	1:D:169:TRP:CD1	2.57	0.58
1:D:195:LEU:H	1:D:265:LEU:HD13	1.68	0.58
1:D:203:LEU:N	1:D:380:GLY:CA	2.52	0.58
1:D:549:LEU:N	1:D:551:LYS:HD3	2.17	0.58
1:D:742:VAL:HG21	1:D:748:TRP:CE3	2.38	0.58
1:D:770:PHE:HA	1:D:773:LYS:CG	2.34	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:891:VAL:CG2	1:D:1135:ASP:HA	2.34	0.58
1:D:993:ARG:O	1:D:996:ARG:CD	2.52	0.58
2:E:215:PHE:HB2	2:E:232:GLY:O	2.04	0.58
2:E:326:ASP:H	2:E:331:VAL:CG1	2.16	0.58
2:F:214:CYS:SG	2:F:236:GLU:HG2	2.43	0.58
2:F:253:ASP:HB3	2:F:257:ARG:CZ	2.34	0.58
2:F:197:LEU:HD11	2:F:322:LEU:HD21	1.85	0.58
2:F:241:TRP:HB2	2:F:336:LEU:CB	2.33	0.58
2:F:442:LEU:O	2:F:455:LEU:HD21	2.04	0.58
1:A:129:PRO:C	1:A:130:LEU:HD12	2.24	0.58
1:A:386:ARG:CZ	1:A:387:GLU:HG2	2.34	0.58
1:A:389:PHE:CD2	1:A:393:MET:HG2	2.39	0.58
1:A:618:HIS:HB2	1:A:725:PRO:CD	2.34	0.58
1:A:630:ASN:HD21	1:A:632:ALA:HB2	1.68	0.58
1:A:726:LYS:HD3	1:A:727:ASP:OD2	2.04	0.58
2:C:344:ARG:HA	2:C:347:LEU:HD13	1.86	0.58
2:C:395:LEU:HD11	2:C:447:THR:OG1	2.03	0.58
2:C:134:LYS:CB	2:C:180:LYS:CE	2.81	0.58
2:E:270:PRO:HA	2:E:273:PHE:HD2	1.66	0.58
1:A:230:GLU:OE1	1:A:231:GLU:N	2.23	0.58
1:D:1116:MET:HE3	1:D:1153:LEU:CA	2.28	0.58
1:D:145:LYS:CB	1:D:1118:TRP:CH2	2.86	0.58
1:D:371:LYS:HG3	1:D:396:CYS:N	2.18	0.58
1:D:490:TRP:HZ2	1:D:598:VAL:HB	1.66	0.58
1:D:652:ILE:O	1:D:653:GLU:C	2.42	0.58
1:D:727:ASP:OD2	1:D:727:ASP:C	2.42	0.58
2:E:403:PHE:CZ	2:E:407:LEU:HD11	2.39	0.58
2:E:422:GLN:C	2:E:424:SER:N	2.57	0.58
2:F:240:VAL:C	2:F:336:LEU:HD13	2.23	0.58
2:F:421:MET:HG3	2:F:422:GLN:N	2.18	0.58
2:F:439:PHE:CD1	2:F:458:ARG:HG3	2.39	0.58
1:A:1027:LYS:CB	1:A:1092:PHE:HD2	2.09	0.58
1:A:1236:GLN:CB	1:A:1237:PRO:HD3	2.22	0.58
1:A:167:PRO:HB2	1:A:177:ARG:NE	2.19	0.58
1:A:253:ALA:HB2	1:A:280:GLU:CA	2.34	0.58
1:A:276:ALA:C	1:A:279:ARG:HB2	2.24	0.58
1:A:316:LYS:NZ	1:A:350:ILE:HG13	2.18	0.58
1:A:392:LEU:HD21	1:A:395:TYR:HB2	1.86	0.58
1:A:448:ALA:O	1:A:870:VAL:CG1	2.51	0.58
1:A:620:TRP:N	1:A:620:TRP:CD1	2.70	0.58
1:A:775:GLU:HB3	1:A:796:LYS:HD2	1.84	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:900:ALA:HA	1:A:915:ALA:HB1	1.83	0.58
1:A:900:ALA:HA	1:A:915:ALA:HB2	1.84	0.58
2:B:93:SER:C	2:B:95:CYS:N	2.56	0.58
2:E:389:ARG:NH2	2:E:390:GLY:O	2.37	0.58
2:E:396:ARG:HA	2:E:399:CYS:SG	2.44	0.58
2:B:217:PRO:HA	2:B:232:GLY:H	1.69	0.58
2:E:218:VAL:N	2:E:231:ILE:HA	2.19	0.58
1:A:534:SER:OG	1:A:535:GLU:N	2.37	0.58
1:D:1124:ALA:CB	1:D:1148:ARG:NH1	2.61	0.57
1:D:86:GLN:HG3	1:D:132:GLY:O	2.04	0.57
1:D:176:THR:H	1:D:185:VAL:CG1	2.17	0.57
1:D:347:TRP:NE1	1:D:349:ASP:CG	2.57	0.57
1:D:815:TRP:CE3	1:D:838:GLY:N	2.67	0.57
2:E:322:LEU:C	2:E:324:GLY:N	2.55	0.57
2:E:404:ASN:O	2:E:406:LEU:N	2.37	0.57
2:E:94:GLY:O	2:E:96:HIS:CD2	2.57	0.57
2:F:308:ASP:OD2	2:F:308:ASP:N	2.36	0.57
2:F:354:PHE:O	2:F:355:GLN:C	2.42	0.57
1:A:1131:ILE:HG12	1:A:1137:VAL:HA	1.86	0.57
1:A:1207:GLU:CA	1:A:1210:TYR:CD1	2.87	0.57
1:A:214:ALA:N	1:A:401:TRP:HZ2	2.02	0.57
1:A:576:LEU:HD23	2:C:481:SER:HG	1.68	0.57
1:A:808:ILE:O	1:A:811:GLN:CD	2.42	0.57
1:A:847:ALA:HA	1:A:852:ARG:HB3	1.86	0.57
2:B:260:LEU:O	2:B:264:ARG:HG3	2.03	0.57
2:B:323:HIS:CE1	2:B:330:ASN:HB2	2.38	0.57
2:B:104:VAL:CG2	2:C:128:VAL:O	2.52	0.57
2:C:193:TYR:HB2	2:C:242:PHE:CZ	2.38	0.57
2:C:185:LEU:HG	2:C:214:CYS:SG	2.44	0.57
2:C:294:PRO:HD2	2:C:295:TRP:CZ3	2.39	0.57
2:C:413:VAL:HG12	2:C:414:TRP:N	2.18	0.57
1:D:1131:ILE:CG1	1:D:1138:ARG:H	2.15	0.57
1:D:1166:TYR:C	1:D:1169:GLY:H	2.06	0.57
1:D:1151:LEU:CD1	1:D:1227:THR:OG1	2.51	0.57
1:D:1236:GLN:HB2	1:D:1237:PRO:HD2	1.86	0.57
1:D:371:LYS:CE	1:D:396:CYS:HB2	2.34	0.57
1:D:459:MET:N	1:D:460:LYS:CE	2.67	0.57
1:D:484:TRP:CD1	1:D:484:TRP:C	2.77	0.57
1:D:580:LEU:O	1:D:580:LEU:HD12	2.04	0.57
1:D:607:TRP:HD1	1:D:779:LEU:CA	2.17	0.57
1:D:726:LYS:O	1:D:726:LYS:HD3	2.03	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:900:ALA:CA	1:D:915:ALA:CB	2.81	0.57
2:E:255:TRP:HE3	2:E:255:TRP:HA	1.67	0.57
2:E:430:SER:C	2:E:433:ASP:HB3	2.25	0.57
2:F:92:LEU:HD21	2:F:344:ARG:HE	1.69	0.57
2:F:78:PHE:HZ	2:F:346:MET:HE2	1.68	0.57
1:A:996:ARG:HH12	1:A:1003:TRP:C	2.08	0.57
1:A:1019:TRP:HZ2	1:A:1111:LEU:HD21	1.67	0.57
1:A:243:ASP:C	1:A:246:PRO:HD2	2.24	0.57
1:A:712:VAL:N	1:A:713:PRO:HD3	2.20	0.57
2:B:208:LEU:CD2	2:B:240:VAL:HB	2.33	0.57
2:C:206:TYR:CZ	2:C:208:LEU:HD23	2.39	0.57
2:C:255:TRP:O	2:C:256:LEU:C	2.41	0.57
2:C:299:LEU:N	2:C:299:LEU:CD2	2.64	0.57
2:C:363:ARG:NH1	2:C:363:ARG:CA	2.64	0.57
2:C:389:ARG:CZ	2:C:390:GLY:N	2.67	0.57
2:C:452:LEU:CD1	2:C:453:ILE:N	2.65	0.57
2:C:454:HIS:HA	2:C:468:ILE:CD1	2.27	0.57
2:B:472:LYS:HG3	2:B:473:ASP:H	1.68	0.57
2:F:216:HIS:O	2:F:231:ILE:HD13	2.03	0.57
2:C:278:CYS:SG	2:C:279:GLN:N	2.77	0.57
1:D:1091:GLU:O	1:D:1096:ARG:NH2	2.35	0.57
1:D:1171:ASN:O	1:D:1172:ASP:HB2	2.02	0.57
1:D:157:LEU:HG	1:D:194:ALA:HB1	1.84	0.57
1:D:175:TRP:CB	1:D:185:VAL:HG13	2.33	0.57
1:D:357:ALA:C	1:D:359:VAL:N	2.54	0.57
1:D:613:HIS:ND1	1:D:717:LEU:CD2	2.67	0.57
1:D:959:GLN:O	1:D:982:ALA:HB1	2.04	0.57
2:F:132:HIS:HA	2:F:182:ARG:HG3	1.86	0.57
2:F:311:LEU:HD22	2:F:311:LEU:N	2.18	0.57
2:F:359:ASN:O	2:F:360:SER:HB2	2.04	0.57
2:F:472:LYS:HA	2:F:475:LEU:CD1	2.34	0.57
1:A:272:SER:HA	1:A:275:ARG:HD3	1.86	0.57
1:A:274:ASP:C	1:A:277:HIS:H	2.08	0.57
1:A:389:PHE:CD1	1:A:390:GLN:NE2	2.72	0.57
1:A:403:THR:OG1	1:A:404:HIS:N	2.37	0.57
1:A:421:PRO:HG2	1:A:422:VAL:N	2.19	0.57
1:A:608:ASP:HB3	1:A:779:LEU:CB	2.34	0.57
1:A:79:LEU:HB3	1:A:89:GLY:CA	2.30	0.57
1:A:943:ARG:O	1:A:947:LYS:HG3	2.04	0.57
2:B:67:GLU:CB	2:B:70:LEU:HD13	2.35	0.57
2:C:352:ASP:C	2:C:352:ASP:OD1	2.41	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:352:ASP:O	2:C:356:LEU:HG	2.04	0.57
2:C:395:LEU:O	2:C:395:LEU:HD23	2.04	0.57
2:C:78:PHE:O	2:C:99:PHE:HB3	2.03	0.57
2:E:184:ASN:O	2:E:184:ASN:OD1	2.21	0.57
1:A:634:LEU:HD12	1:A:638:THR:HG21	1.85	0.57
1:D:442:GLU:CA	1:D:442:GLU:OE2	2.50	0.57
1:A:1183:VAL:O	1:A:1217:ALA:HB1	2.03	0.57
1:D:138:HIS:HA	1:D:141:LEU:HD13	1.84	0.57
1:D:145:LYS:CB	1:D:1118:TRP:HH2	2.17	0.57
1:D:400:VAL:C	1:D:403:THR:HG23	2.24	0.57
1:D:792:LEU:O	1:D:795:ASN:N	2.37	0.57
1:D:801:TRP:HH2	1:D:805:HIS:CE1	2.22	0.57
2:E:122:ARG:N	2:E:122:ARG:HD2	2.20	0.57
2:E:247:THR:HG22	2:E:250:GLN:NE2	2.13	0.57
2:F:347:LEU:O	2:F:350:LEU:CD1	2.52	0.57
1:A:1030:ARG:NH2	1:A:1040:LYS:HG2	2.19	0.57
1:A:1054:GLU:O	1:A:1058:PHE:HD2	1.86	0.57
1:A:612:LEU:O	1:A:613:HIS:ND1	2.37	0.57
2:B:104:VAL:O	2:B:105:GLU:C	2.41	0.57
2:B:213:VAL:HG23	2:B:235:THR:HA	1.86	0.57
2:C:198:ASP:OD1	2:C:198:ASP:C	2.42	0.57
2:C:336:LEU:C	2:C:336:LEU:CD2	2.72	0.57
1:D:1175:GLN:C	1:D:1177:VAL:N	2.58	0.57
1:D:382:MET:O	1:D:385:ILE:HG22	2.04	0.57
1:D:417:ARG:HG2	1:D:417:ARG:NH1	2.19	0.57
1:D:458:GLU:C	1:D:460:LYS:HE3	2.24	0.57
1:D:652:ILE:O	1:D:655:LEU:N	2.37	0.57
1:D:831:TYR:CB	1:D:837:TYR:OH	2.52	0.57
2:E:288:LYS:HB2	2:E:290:TYR:OH	2.05	0.57
2:F:114:TRP:CB	2:F:262:TRP:CZ2	2.87	0.57
2:F:342:LEU:HD23	2:F:345:GLY:N	2.09	0.57
1:A:1043:VAL:C	1:A:1045:ALA:N	2.48	0.57
1:A:211:LEU:HD12	1:A:222:SER:H	1.67	0.57
1:A:376:LEU:HD12	1:A:379:LYS:HB3	1.85	0.57
1:A:498:LYS:CE	1:A:499:LYS:H	2.12	0.57
1:A:775:GLU:CA	1:A:780:GLN:HG3	2.20	0.57
1:A:207:THR:HG21	1:A:812:MET:HB3	1.85	0.57
1:A:928:GLY:O	1:A:931:LEU:N	2.35	0.57
2:B:67:GLU:HB3	2:B:70:LEU:CB	2.29	0.57
2:C:115:TRP:O	2:C:119:VAL:HG23	2.05	0.57
2:C:471:LEU:CA	2:C:474:PHE:HB3	2.30	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:668:LEU:H	1:D:668:LEU:HD23	1.70	0.57
2:B:450:ASN:HB2	2:B:452:LEU:HD23	1.85	0.57
1:D:966:LEU:C	1:D:970:ASN:HD22	2.07	0.57
1:A:619:GLY:HA3	1:A:750:PHE:HA	1.86	0.57
2:F:326:ASP:OD2	2:F:327:GLY:N	2.35	0.57
1:D:1032:THR:O	1:D:1032:THR:HG23	2.04	0.57
1:D:1196:ASP:C	1:D:1196:ASP:OD2	2.41	0.57
1:D:272:SER:HA	1:D:843:GLN:CB	2.35	0.57
1:D:369:LEU:O	1:D:371:LYS:N	2.37	0.57
1:D:851:THR:HG22	1:D:851:THR:O	2.03	0.57
2:E:300:ILE:HG13	2:E:301:GLU:N	2.20	0.57
2:E:234:LYS:HA	2:E:343:ASP:OD2	2.05	0.57
2:F:200:VAL:O	2:F:200:VAL:HG12	2.04	0.57
2:F:416:GLY:HA2	2:F:419:GLU:CB	2.34	0.57
2:F:425:LEU:HD13	2:F:442:LEU:HD11	1.86	0.57
2:F:441:VAL:HG23	2:F:441:VAL:O	2.05	0.57
2:F:74:GLN:NE2	2:F:75:ARG:HH21	2.01	0.57
1:A:1069:ILE:N	1:A:1070:PRO:HD3	2.20	0.57
1:A:1125:ILE:HD12	1:A:1125:ILE:O	2.04	0.57
1:A:1141:VAL:HG11	1:A:1145:ASP:O	2.05	0.57
1:A:886:LEU:C	1:A:1146:ARG:NH2	2.58	0.57
1:A:130:LEU:HG	1:A:131:TYR:CE1	2.39	0.57
1:A:262:GLN:OE1	1:A:262:GLN:HA	2.02	0.57
1:A:275:ARG:O	1:A:278:ILE:C	2.42	0.57
1:A:373:PRO:HD2	1:A:374:ARG:NH1	2.19	0.57
1:A:408:GLN:O	1:A:411:LEU:HG	2.03	0.57
2:B:113:GLU:C	2:B:262:TRP:CZ2	2.78	0.57
2:B:309:HIS:CB	2:B:312:LEU:HD22	2.18	0.57
2:C:254:PHE:HA	2:C:257:ARG:HD2	1.86	0.57
2:C:372:LEU:CD2	2:C:436:SER:HB2	2.35	0.57
2:C:82:SER:O	2:C:83:LYS:CB	2.53	0.57
2:B:398:VAL:O	2:B:402:LEU:HG	2.05	0.57
1:D:961:PHE:CZ	1:D:1049:TRP:CE3	2.92	0.57
1:D:1090:GLU:O	1:D:1090:GLU:HG3	2.04	0.57
1:D:392:LEU:HD23	1:D:392:LEU:O	2.04	0.57
1:D:580:LEU:HD11	1:D:593:SER:CB	2.35	0.57
1:D:628:ARG:O	1:D:630:ASN:N	2.33	0.57
1:D:719:LEU:CG	1:D:720:THR:N	2.65	0.57
1:D:770:PHE:HE1	1:D:968:GLN:CB	2.18	0.57
2:E:108:LYS:HZ2	2:E:108:LYS:HB2	1.70	0.57
2:E:122:ARG:O	2:E:125:VAL:HG23	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:123:GLU:HG2	2:F:415:PRO:CG	2.33	0.57
2:E:200:VAL:HG13	2:F:418:LEU:HD23	1.86	0.57
2:E:215:PHE:CB	2:E:233:GLU:HA	2.35	0.57
2:E:243:THR:OG1	2:E:244:PRO:HD2	2.05	0.57
2:E:441:VAL:CG1	2:E:442:LEU:H	2.17	0.57
2:E:460:THR:O	2:E:461:THR:CG2	2.53	0.57
1:A:1020:ILE:HG13	1:A:1021:SER:N	2.19	0.57
1:A:1205:GLY:HA3	1:A:1209:ARG:NE	2.20	0.57
1:A:1231:LEU:CD1	1:A:1236:GLN:HE22	2.15	0.57
1:A:203:LEU:CA	1:A:205:GLU:H	2.16	0.57
1:A:267:VAL:HG23	1:A:291:PHE:CE2	2.40	0.57
1:A:582:ASP:CB	1:A:592:LEU:HD22	2.35	0.57
1:A:613:HIS:O	1:A:614:TYR:HB2	2.03	0.57
1:A:620:TRP:C	1:A:748:TRP:CB	2.73	0.57
1:A:876:ALA:O	1:A:877:MET:C	2.43	0.57
2:B:206:TYR:CZ	2:B:242:PHE:HZ	2.22	0.57
2:B:325:ARG:N	2:B:325:ARG:NE	2.53	0.57
2:B:372:LEU:O	2:B:436:SER:HA	2.05	0.57
1:A:120:LEU:N	1:A:120:LEU:HD22	2.18	0.57
2:C:94:GLY:HA2	2:C:96:HIS:ND1	2.19	0.57
1:D:274:ASP:OD2	1:D:293:ASP:CA	2.47	0.57
1:D:406:VAL:CG2	1:D:407:PHE:N	2.67	0.57
1:D:548:CYS:N	1:D:551:LYS:NZ	2.50	0.57
1:D:645:VAL:C	1:D:646:VAL:HG23	2.25	0.57
1:D:711:ALA:O	1:D:713:PRO:HD3	2.05	0.57
2:E:436:SER:O	2:E:437:ILE:HD12	2.04	0.57
2:F:300:ILE:CG2	2:F:342:LEU:HD11	2.34	0.57
1:A:1017:GLY:O	1:A:1019:TRP:N	2.37	0.57
1:A:1191:LYS:O	1:A:1192:GLU:CB	2.51	0.57
1:A:192:GLU:CB	1:A:217:PRO:CG	2.77	0.57
1:A:197:PHE:HA	1:A:213:VAL:O	2.04	0.57
1:A:290:ARG:C	1:A:291:PHE:CD2	2.78	0.57
1:A:299:MET:HG3	1:A:848:GLY:HA2	1.86	0.57
1:A:560:PRO:O	1:A:561:LYS:HD3	2.05	0.57
1:A:770:PHE:HB2	1:A:771:LEU:HD22	1.86	0.57
2:C:204:LEU:HD21	2:C:206:TYR:CD1	2.39	0.57
2:C:342:LEU:N	2:C:342:LEU:HD22	2.20	0.57
2:F:317:GLY:HA3	2:F:320:SER:CB	2.33	0.57
2:E:311:LEU:HD22	2:E:312:LEU:HD12	1.86	0.57
2:C:145:PHE:O	2:C:146:ARG:HG3	2.05	0.57
1:D:1016:GLU:HG3	1:D:1022:LEU:HD21	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1072:THR:N	1:D:1073:PRO:CD	2.66	0.57
1:D:1134:HIS:O	1:D:1135:ASP:CB	2.48	0.57
1:D:224:CYS:SG	1:D:228:LEU:CD2	2.93	0.57
1:D:206:GLY:HA2	1:D:245:ILE:HG21	1.87	0.57
2:E:387:VAL:HG12	2:E:416:GLY:N	2.20	0.57
2:E:426:GLU:HA	2:E:429:TYR:HB3	1.86	0.57
2:E:432:TYR:CD2	2:E:437:ILE:HG12	2.35	0.57
2:E:448:LEU:CD2	2:E:449:GLU:N	2.67	0.57
2:E:442:LEU:O	2:E:453:ILE:HB	2.04	0.57
2:E:93:SER:C	2:E:95:CYS:N	2.54	0.57
1:A:192:GLU:O	1:A:193:ARG:CG	2.52	0.57
1:A:407:PHE:O	1:A:411:LEU:N	2.38	0.57
1:A:515:ILE:CG1	1:A:568:GLY:O	2.53	0.57
1:A:616:GLU:O	1:A:617:ARG:CB	2.52	0.57
1:A:618:HIS:HB2	1:A:725:PRO:HD2	1.87	0.57
1:A:949:PHE:N	1:A:949:PHE:CD2	2.73	0.57
2:B:417:TYR:O	2:C:203:ARG:NH1	2.38	0.57
2:C:243:THR:O	2:C:334:CYS:CB	2.52	0.57
2:C:398:VAL:C	2:C:401:GLY:H	2.07	0.57
1:A:235:TRP:O	1:A:235:TRP:HD1	1.88	0.57
1:D:539:PHE:O	1:D:540:GLN:CB	2.53	0.57
1:D:136:ASP:OD2	1:D:1167:LYS:CE	2.53	0.57
1:D:1231:LEU:O	1:D:1232:GLU:HG2	2.05	0.57
1:D:211:LEU:HA	1:D:221:TYR:CD1	2.40	0.57
1:D:597:ARG:NH1	1:D:724:GLY:C	2.57	0.57
1:D:730:PRO:HB3	1:D:761:ASN:HA	1.86	0.57
2:E:384:ALA:C	2:E:414:TRP:HB3	2.26	0.57
2:F:406:LEU:CA	2:F:409:ASN:ND2	2.56	0.57
2:F:82:SER:O	2:F:83:LYS:CG	2.52	0.57
2:F:88:ARG:HG2	2:F:89:ASP:N	2.20	0.57
1:A:956:GLY:O	1:A:1094:THR:HA	2.05	0.57
1:A:1208:ARG:H	1:A:1210:TYR:N	2.02	0.57
1:A:276:ALA:HA	1:A:279:ARG:CG	2.35	0.57
1:A:608:ASP:OD1	1:A:779:LEU:O	2.22	0.57
1:A:655:LEU:CD1	1:A:719:LEU:C	2.72	0.57
1:A:866:ARG:CB	1:A:1069:ILE:HG21	2.35	0.57
2:C:351:TYR:C	2:C:355:GLN:CB	2.60	0.57
2:C:417:TYR:N	2:C:417:TYR:CD2	2.40	0.57
2:E:481:SER:O	2:E:484:ASN:HB2	2.04	0.57
1:D:1002:GLU:O	1:D:1005:VAL:N	2.37	0.56
1:D:1219:ASP:OD2	1:D:1222:GLN:NE2	2.37	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:211:LEU:HG	1:D:221:TYR:HE1	1.70	0.56
1:D:275:ARG:O	1:D:279:ARG:N	2.36	0.56
1:D:363:TYR:O	1:D:364:VAL:HB	2.05	0.56
1:D:550:GLN:HB2	1:D:551:LYS:HE3	1.86	0.56
1:D:293:ASP:OD1	1:D:845:VAL:HG13	2.04	0.56
1:D:914:THR:HA	1:D:918:TRP:HB2	1.85	0.56
2:E:431:LYS:HG3	2:E:432:TYR:N	2.20	0.56
1:A:1113:LEU:CB	1:A:1117:LYS:HE3	2.34	0.56
1:A:1154:GLN:O	1:A:1158:LEU:HG	2.05	0.56
1:A:249:VAL:HG21	1:A:817:PRO:HG3	1.87	0.56
1:A:472:GLN:OE1	1:A:472:GLN:N	2.38	0.56
1:A:516:GLU:O	1:A:517:GLY:O	2.21	0.56
1:A:869:ARG:N	1:A:1201:SER:H	2.03	0.56
2:B:262:TRP:O	2:B:265:LYS:N	2.38	0.56
2:C:115:TRP:CZ3	2:C:210:GLN:HA	2.40	0.56
2:C:125:VAL:CG2	2:C:126:PHE:N	2.61	0.56
2:C:344:ARG:HG2	2:C:347:LEU:HD22	1.86	0.56
2:C:406:LEU:O	2:C:407:LEU:C	2.43	0.56
2:C:413:VAL:C	2:C:415:PRO:HD2	2.24	0.56
2:C:446:THR:OG1	2:C:455:LEU:CD2	2.51	0.56
1:A:504:LYS:HD3	1:A:523:ASP:CB	2.34	0.56
1:D:99:ALA:O	1:D:100:VAL:C	2.43	0.56
2:C:144:ALA:O	2:C:145:PHE:CB	2.53	0.56
1:D:948:ILE:HD13	1:D:1071:ARG:HD2	1.87	0.56
1:D:891:VAL:HG21	1:D:1135:ASP:HA	1.87	0.56
1:D:1198:LYS:N	1:D:1204:THR:OG1	2.38	0.56
1:D:132:GLY:N	1:D:138:HIS:CE1	2.73	0.56
1:D:282:TYR:HD2	1:D:432:VAL:HG13	1.67	0.56
1:D:407:PHE:HD1	1:D:411:LEU:CD2	2.19	0.56
1:D:652:ILE:HG22	1:D:656:TYR:CE2	2.40	0.56
1:D:767:ALA:C	1:D:768:LYS:O	2.36	0.56
2:E:241:TRP:CZ3	2:E:255:TRP:HZ2	2.23	0.56
2:E:432:TYR:HB3	2:E:437:ILE:HD11	1.82	0.56
2:E:445:GLU:OE2	2:E:445:GLU:N	2.38	0.56
2:F:127:PRO:CG	2:F:128:VAL:N	2.67	0.56
2:F:395:LEU:HB3	2:F:443:VAL:HG23	1.86	0.56
2:F:403:PHE:HD1	2:F:417:TYR:OH	1.82	0.56
2:F:265:LYS:HE2	2:F:485:VAL:HG22	1.86	0.56
1:A:80:SER:OG	1:A:125:LEU:CG	2.53	0.56
1:A:258:GLN:OE1	1:A:258:GLN:N	2.38	0.56
1:A:474:LEU:HD13	1:A:474:LEU:N	2.20	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:80:SER:HG	1:A:125:LEU:HD11	1.70	0.56
1:A:435:LEU:O	1:A:839:ALA:HB1	2.06	0.56
2:B:461:THR:HG23	2:B:463:LYS:HZ2	1.70	0.56
2:C:202:LYS:O	2:C:325:ARG:HG2	2.05	0.56
1:D:513:LEU:O	1:D:568:GLY:HA2	2.05	0.56
2:E:395:LEU:O	2:E:399:CYS:SG	2.64	0.56
2:C:365:LYS:HB3	2:C:367:LEU:HD21	1.88	0.56
1:D:1027:LYS:CG	1:D:1091:GLU:HA	2.35	0.56
1:D:556:THR:HG23	2:F:451:GLY:O	2.04	0.56
1:D:559:LEU:HD12	1:D:560:PRO:CD	2.33	0.56
1:D:496:LYS:CE	1:D:571:GLY:HA2	2.36	0.56
1:D:850:ILE:HG12	1:D:851:THR:N	2.20	0.56
1:D:992:LEU:C	1:D:996:ARG:HD2	2.24	0.56
2:E:214:CYS:N	2:E:234:LYS:O	2.32	0.56
1:D:1202:ASN:O	1:D:1203:PRO:O	2.24	0.56
1:D:297:MET:HG3	1:D:415:LEU:HD11	1.87	0.56
1:D:418:CYS:N	1:D:419:PRO:CD	2.68	0.56
1:D:656:TYR:CD1	1:D:743:ASP:CB	2.89	0.56
1:D:813:VAL:HB	1:D:840:ILE:CG1	2.31	0.56
1:D:873:GLU:OE1	1:D:1203:PRO:HG3	2.05	0.56
1:D:112:LEU:HD21	1:D:917:GLY:O	2.06	0.56
1:D:919:MET:C	1:D:920:THR:O	2.43	0.56
2:E:431:LYS:CD	2:E:432:TYR:H	2.17	0.56
2:F:311:LEU:HD22	2:F:311:LEU:H	1.71	0.56
2:F:298:GLU:HG3	2:F:344:ARG:HH21	1.70	0.56
2:F:350:LEU:HD12	2:F:351:TYR:H	1.71	0.56
2:F:354:PHE:CE1	2:F:370:LYS:HB3	2.38	0.56
2:F:466:MET:CB	2:F:471:LEU:CD2	2.70	0.56
1:A:1060:LYS:CD	1:A:1073:PRO:CB	2.76	0.56
1:A:196:VAL:CG1	1:A:401:TRP:HZ3	2.18	0.56
1:A:232:ARG:HD2	1:A:239:LEU:CD2	2.34	0.56
1:A:293:ASP:HB2	1:A:296:SER:HB3	1.85	0.56
1:A:371:LYS:HG3	1:A:372:GLU:N	2.19	0.56
1:A:608:ASP:O	1:A:620:TRP:CZ2	2.58	0.56
1:A:730:PRO:HB3	1:A:761:ASN:HA	1.88	0.56
1:A:620:TRP:C	1:A:748:TRP:HB3	2.24	0.56
1:A:867:PRO:CG	1:A:1067:SER:OG	2.53	0.56
1:A:913:CYS:C	1:A:918:TRP:N	2.58	0.56
1:A:938:THR:O	1:A:939:VAL:HG23	2.05	0.56
2:B:88:ARG:HA	2:B:91:LEU:HD12	1.86	0.56
2:C:204:LEU:HG	2:C:206:TYR:CB	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:393:LEU:HA	2:C:396:ARG:CG	2.34	0.56
2:C:414:TRP:CG	2:C:414:TRP:O	2.57	0.56
2:C:386:ASP:CB	2:C:442:LEU:HD23	2.25	0.56
2:C:444:THR:O	2:C:446:THR:N	2.38	0.56
1:A:1040:LYS:HB3	1:A:1044:VAL:HG12	1.86	0.56
1:A:1172:ASP:O	1:A:1173:LEU:HG	2.05	0.56
1:A:1212:ILE:HD12	1:A:1212:ILE:N	2.19	0.56
1:A:176:THR:OG1	1:A:177:ARG:N	2.35	0.56
1:A:411:LEU:HD12	1:A:412:PRO:CD	2.35	0.56
1:A:482:ASP:CB	1:A:483:PRO:CD	2.83	0.56
1:A:877:MET:HA	1:A:877:MET:CE	2.36	0.56
1:A:905:ALA:HA	1:A:910:MET:HB3	1.86	0.56
2:B:417:TYR:O	2:B:418:LEU:HB3	2.06	0.56
2:B:78:PHE:CD1	2:B:78:PHE:O	2.58	0.56
2:C:234:LYS:HZ1	2:C:344:ARG:HD2	1.65	0.56
2:F:221:THR:HG22	2:F:222:LYS:N	2.20	0.56
1:D:970:ASN:C	1:D:971:HIS:ND1	2.59	0.56
1:D:757:GLY:O	1:D:758:ASN:HB2	2.05	0.56
1:A:981:LYS:O	1:A:985:MET:HG3	2.05	0.56
1:D:1133:ILE:HD13	1:D:1133:ILE:N	2.19	0.56
1:D:1202:ASN:O	1:D:1205:GLY:CA	2.53	0.56
1:D:1200:PRO:HD2	1:D:1203:PRO:HG2	1.88	0.56
1:D:253:ALA:HB2	1:D:280:GLU:HB2	1.88	0.56
1:D:278:ILE:HD13	1:D:291:PHE:CD1	2.37	0.56
1:D:278:ILE:HG21	1:D:431:GLY:O	2.04	0.56
1:D:559:LEU:CG	1:D:560:PRO:CD	2.82	0.56
1:D:76:ILE:HB	1:D:911:HIS:NE2	2.19	0.56
1:D:901:VAL:HA	1:D:904:ASP:OD2	2.06	0.56
2:F:308:ASP:CG	2:F:334:CYS:HA	2.26	0.56
2:F:483:LYS:C	2:F:483:LYS:HE2	2.25	0.56
2:F:82:SER:O	2:F:83:LYS:CB	2.52	0.56
1:A:1047:ARG:HA	1:A:1047:ARG:CZ	2.34	0.56
1:A:196:VAL:HG13	1:A:269:HIS:NE2	2.21	0.56
1:A:197:PHE:H	1:A:269:HIS:CD2	2.20	0.56
1:A:602:LEU:O	1:A:603:MET:C	2.41	0.56
1:A:721:ALA:O	1:A:722:ARG:C	2.42	0.56
1:A:870:VAL:C	1:A:872:SER:H	2.09	0.56
1:A:886:LEU:H	1:A:1146:ARG:HH21	1.50	0.56
2:B:311:LEU:HD12	2:B:312:LEU:H	1.68	0.56
2:C:308:ASP:OD2	2:C:334:CYS:SG	2.62	0.56
2:C:350:LEU:O	2:C:353:SER:OG	2.22	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:431:LYS:O	2:C:434:GLU:HB2	2.05	0.56
1:D:1092:PHE:O	1:D:1096:ARG:CZ	2.54	0.56
1:D:1154:GLN:O	1:D:1158:LEU:HD23	2.05	0.56
1:D:171:TRP:O	1:D:172:ALA:CB	2.44	0.56
1:D:193:ARG:HH11	1:D:193:ARG:HA	1.68	0.56
1:D:245:ILE:CG1	1:D:246:PRO:CD	2.77	0.56
1:D:251:THR:O	1:D:255:SER:N	2.39	0.56
1:D:463:LEU:CA	1:D:466:LEU:HD22	2.35	0.56
1:D:486:TRP:HZ2	1:D:722:ARG:N	2.03	0.56
1:D:490:TRP:O	1:D:580:LEU:HA	2.06	0.56
1:D:641:GLU:O	1:D:645:VAL:CG2	2.54	0.56
1:D:648:PRO:CD	1:D:649:TYR:N	2.68	0.56
1:D:770:PHE:HE1	1:D:968:GLN:HB3	1.67	0.56
1:D:74:LEU:HD13	1:D:77:GLN:HA	1.86	0.56
1:D:788:GLY:O	1:D:792:LEU:HD13	2.04	0.56
1:D:892:ASP:HA	1:D:1135:ASP:OD2	2.05	0.56
2:F:115:TRP:NE1	2:F:119:VAL:HG21	2.21	0.56
2:F:265:LYS:HG3	2:F:485:VAL:HG21	1.88	0.56
2:F:387:VAL:CG1	2:F:443:VAL:CG2	2.83	0.56
1:A:1027:LYS:HD3	1:A:1092:PHE:HE2	1.71	0.56
1:A:1143:GLU:O	1:A:1146:ARG:HB2	2.05	0.56
1:A:1212:ILE:N	1:A:1212:ILE:CD1	2.68	0.56
1:A:196:VAL:HG11	1:A:404:HIS:CG	2.41	0.56
1:A:354:ASN:O	1:A:355:SER:CB	2.54	0.56
1:A:386:ARG:CB	1:A:390:GLN:HE22	2.19	0.56
1:A:437:VAL:HG12	1:A:438:ASN:N	2.20	0.56
1:A:585:TRP:N	1:A:585:TRP:HE3	2.02	0.56
1:A:821:LEU:O	1:A:821:LEU:HG	2.05	0.56
2:C:241:TRP:HB2	2:C:336:LEU:HB2	1.85	0.56
2:C:416:GLY:C	2:C:419:GLU:HB3	2.25	0.56
2:C:426:GLU:OE2	2:C:429:TYR:HD1	1.87	0.56
2:C:371:VAL:HG23	2:C:433:ASP:CA	2.35	0.56
1:D:97:GLU:O	1:D:101:ARG:CG	2.53	0.56
1:D:1025:LEU:HD13	1:D:1027:LYS:CE	2.30	0.56
1:D:1129:PHE:CB	1:D:1139:TYR:HA	2.35	0.56
1:D:242:ALA:C	1:D:245:ILE:HG12	2.26	0.56
1:D:257:THR:C	1:D:258:GLN:HG3	2.26	0.56
1:D:194:ALA:HB1	1:D:265:LEU:HD12	1.85	0.56
1:D:269:HIS:C	1:D:274:ASP:OD1	2.44	0.56
1:D:289:MET:SD	1:D:291:PHE:CE2	2.99	0.56
1:D:412:PRO:HG2	1:D:413:LEU:N	2.21	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:580:LEU:CD1	1:D:593:SER:OG	2.47	0.56
1:D:914:THR:C	1:D:918:TRP:HB3	2.19	0.56
2:E:128:VAL:O	2:E:210:GLN:NE2	2.39	0.56
2:E:283:GLY:HA3	2:E:306:LEU:CD2	2.35	0.56
2:F:204:LEU:N	2:F:204:LEU:HD12	2.21	0.56
2:F:403:PHE:CE2	2:F:407:LEU:HB2	2.41	0.56
1:A:278:ILE:HD13	1:A:291:PHE:HD1	1.71	0.56
1:A:597:ARG:O	1:A:598:VAL:CB	2.54	0.56
1:A:598:VAL:HA	1:A:601:LYS:HD2	1.87	0.56
1:A:616:GLU:HG3	1:A:724:GLY:C	2.25	0.56
1:A:732:TYR:CE1	1:A:755:LYS:CA	2.88	0.56
1:A:83:LEU:C	1:A:87:ILE:O	2.44	0.56
1:A:870:VAL:O	1:A:872:SER:N	2.38	0.56
2:B:269:SER:O	2:B:272:ASN:ND2	2.38	0.56
2:B:385:LEU:HD23	2:B:385:LEU:N	2.20	0.56
2:B:430:SER:O	2:B:433:ASP:HB2	2.05	0.56
2:B:67:GLU:HG2	2:B:69:LEU:HG	1.88	0.56
2:B:72:ILE:CG1	2:B:73:CYS:N	2.67	0.56
2:C:185:LEU:HD21	2:C:214:CYS:SG	2.45	0.56
2:C:286:GLY:HA3	2:C:304:TRP:HA	1.88	0.56
2:C:303:LEU:CA	2:C:337:SER:CB	2.81	0.56
2:C:432:TYR:CD1	2:C:440:THR:HG21	2.41	0.56
2:C:454:HIS:CD2	2:C:472:LYS:HE3	2.40	0.56
1:A:520:ALA:HB3	1:A:521:PRO:HD3	1.87	0.56
1:D:1069:ILE:O	1:D:1071:ARG:NH1	2.39	0.56
1:D:1108:TYR:HE1	1:D:1160:THR:HG22	1.71	0.56
1:D:196:VAL:HG12	1:D:268:GLY:CA	2.36	0.56
1:D:369:LEU:HB3	1:D:372:GLU:CB	2.35	0.56
1:D:369:LEU:HB3	1:D:372:GLU:HB2	1.86	0.56
1:D:576:LEU:HD11	1:D:577:CYS:SG	2.46	0.56
1:D:660:CYS:SG	1:D:745:PRO:HG3	2.45	0.56
1:D:795:ASN:CA	1:D:798:ILE:HG12	2.35	0.56
1:D:797:MET:O	1:D:869:ARG:HD3	2.05	0.56
1:D:797:MET:CA	1:D:869:ARG:HD3	2.35	0.56
1:D:916:PHE:CE2	1:D:1174:PRO:HG2	2.40	0.56
2:E:263:TRP:O	2:E:265:LYS:N	2.38	0.56
2:F:236:GLU:CB	2:F:341:ASP:HB3	2.36	0.56
2:F:375:HIS:ND1	2:F:376:PRO:HD2	2.20	0.56
2:F:390:GLY:HA2	2:F:392:THR:N	2.20	0.56
2:F:76:ARG:CZ	2:F:434:GLU:OE1	2.54	0.56
2:F:483:LYS:O	2:F:485:VAL:CG1	2.54	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:78:PHE:HB2	2:F:102:LEU:HD22	1.87	0.56
1:A:107:LEU:CB	1:A:113:TRP:CD1	2.84	0.56
1:A:1019:TRP:CZ3	1:A:1164:PHE:HA	2.39	0.56
1:A:278:ILE:CD1	1:A:291:PHE:CD1	2.88	0.56
1:A:392:LEU:HD23	1:A:392:LEU:O	2.05	0.56
1:A:656:TYR:CB	1:A:720:THR:OG1	2.51	0.56
1:A:754:HIS:ND1	1:A:760:CYS:O	2.39	0.56
1:A:798:ILE:CD1	1:A:802:ARG:HE	2.19	0.56
1:A:440:ASN:CB	1:A:877:MET:SD	2.92	0.56
1:A:576:LEU:CD2	2:C:481:SER:OG	2.51	0.56
1:D:666:GLN:CA	1:D:669:MET:O	2.54	0.56
2:E:133:HIS:NE2	2:F:231:ILE:HG22	2.20	0.56
2:C:285:LYS:HB2	2:C:285:LYS:HZ1	1.71	0.56
1:D:1116:MET:O	1:D:1120:PHE:HD1	1.88	0.56
1:D:223:TRP:CZ3	1:D:250:PRO:CB	2.88	0.56
1:D:255:SER:CB	1:D:256:PRO:C	2.73	0.56
1:D:270:ASN:OD1	1:D:404:HIS:CE1	2.59	0.56
1:D:213:VAL:CG2	1:D:397:ALA:HB1	2.26	0.56
1:D:410:GLN:HA	1:D:410:GLN:OE1	2.06	0.56
1:D:457:ARG:HH22	2:F:485:VAL:CG2	2.15	0.56
1:D:458:GLU:CA	1:D:461:LYS:HD3	2.19	0.56
1:D:951:TYR:O	1:D:954:ILE:HB	2.05	0.56
2:E:193:TYR:HE2	2:E:322:LEU:HB3	1.70	0.56
2:E:478:TYR:O	2:E:482:ALA:HB2	2.05	0.56
2:E:86:LEU:HB2	2:E:91:LEU:HD22	1.88	0.56
2:F:415:PRO:CB	2:F:417:TYR:HE2	2.18	0.56
2:F:454:HIS:CA	2:F:468:ILE:HD12	2.20	0.56
2:F:473:ASP:O	2:F:477:LYS:HG3	2.06	0.56
1:A:1091:GLU:O	1:A:1092:PHE:HB2	2.06	0.56
1:A:1093:MET:O	1:A:1094:THR:C	2.44	0.56
1:A:1108:TYR:HA	1:A:1164:PHE:CZ	2.41	0.56
1:A:253:ALA:O	1:A:254:SER:CB	2.54	0.56
1:A:277:HIS:O	1:A:281:GLN:HG3	2.06	0.56
1:A:301:ILE:CD1	1:A:414:PHE:HB3	2.36	0.56
1:A:734:HIS:N	1:A:752:LEU:HD11	2.21	0.56
1:A:823:ARG:CG	1:A:823:ARG:O	2.53	0.56
1:A:279:ARG:NH1	1:A:841:LEU:O	2.39	0.56
1:A:897:TRP:NE1	1:A:901:VAL:CG1	2.64	0.56
2:B:132:HIS:CG	2:B:182:ARG:NH2	2.73	0.56
2:C:432:TYR:CB	2:C:437:ILE:HD11	2.35	0.56
1:D:1163:MET:HE1	1:D:1167:LYS:HE3	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1231:LEU:C	1:D:1232:GLU:HG2	2.25	0.56
1:D:196:VAL:CG2	1:D:215:ILE:CD1	2.83	0.56
1:D:270:ASN:CA	1:D:294:THR:HG23	2.29	0.56
1:D:437:VAL:HA	1:D:441:TRP:CD1	2.38	0.56
1:D:546:ARG:O	1:D:550:GLN:NE2	2.39	0.56
1:D:652:ILE:CG1	1:D:655:LEU:HG	2.36	0.56
1:D:721:ALA:O	1:D:722:ARG:CB	2.53	0.56
2:F:270:PRO:HA	2:F:273:PHE:CE2	2.41	0.56
1:A:1232:GLU:HG3	1:A:1234:ARG:HG2	1.88	0.56
1:A:253:ALA:CB	1:A:280:GLU:C	2.73	0.56
1:A:656:TYR:HB3	1:A:720:THR:HG1	1.68	0.56
1:A:817:PRO:HG2	1:A:818:ARG:H	1.71	0.56
1:A:955:TYR:HB3	1:A:1098:ASN:ND2	2.20	0.56
2:B:213:VAL:HA	2:B:235:THR:HA	1.88	0.56
2:B:79:LEU:HA	2:B:100:GLY:H	1.71	0.56
2:C:194:VAL:O	2:C:197:LEU:CD2	2.54	0.56
2:C:241:TRP:HB2	2:C:336:LEU:CB	2.36	0.56
2:C:254:PHE:HA	2:C:257:ARG:CD	2.36	0.56
2:C:268:MET:HB2	2:C:352:ASP:OD1	2.06	0.56
2:C:270:PRO:HA	2:C:273:PHE:CE2	2.41	0.56
2:C:325:ARG:HH11	2:C:325:ARG:CG	2.07	0.56
2:C:347:LEU:C	2:C:350:LEU:CD1	2.74	0.56
2:C:356:LEU:HD13	2:C:359:ASN:HD21	1.70	0.56
2:C:371:VAL:HG23	2:C:433:ASP:C	2.27	0.56
1:D:539:PHE:O	1:D:540:GLN:CG	2.54	0.56
1:D:430:MET:CE	1:D:1132:SER:HB3	2.35	0.56
1:D:1154:GLN:HG2	1:D:1179:PHE:CE2	2.40	0.56
1:D:189:ILE:HG12	1:D:190:PRO:HD2	1.88	0.56
1:D:753:PRO:HG3	1:D:770:PHE:CG	2.41	0.56
2:E:247:THR:O	2:E:250:GLN:HB3	2.06	0.56
2:E:322:LEU:C	2:E:324:GLY:H	2.07	0.56
2:F:134:LYS:CB	2:F:180:LYS:CE	2.83	0.56
2:F:423:SER:CB	2:F:428:LEU:HD21	2.36	0.56
1:A:1095:SER:O	1:A:1099:TRP:HE3	1.89	0.56
1:A:1109:LEU:O	1:A:1113:LEU:HG	2.05	0.56
1:A:215:ILE:HB	1:A:401:TRP:CE3	2.41	0.56
1:A:434:TYR:CE2	1:A:1128:ARG:HD2	2.40	0.56
1:A:441:TRP:CZ3	1:A:445:LEU:CD1	2.89	0.56
1:A:612:LEU:HD22	1:A:620:TRP:HA	1.88	0.56
1:A:249:VAL:CG2	1:A:815:TRP:O	2.52	0.56
1:A:885:THR:OG1	1:A:1146:ARG:NE	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:347:LEU:O	2:C:350:LEU:CD1	2.54	0.56
1:D:515:ILE:HG23	1:D:565:HIS:C	2.25	0.56
1:D:443:ARG:HG2	1:D:443:ARG:NH1	2.19	0.56
1:D:196:VAL:HG12	1:D:268:GLY:N	2.21	0.55
1:D:278:ILE:HA	1:D:282:TYR:CE1	2.41	0.55
1:D:253:ALA:N	1:D:280:GLU:OE1	2.39	0.55
1:D:282:TYR:HE2	1:D:432:VAL:CG2	2.14	0.55
1:D:278:ILE:HA	1:D:291:PHE:CE1	2.41	0.55
1:D:400:VAL:O	1:D:403:THR:HG23	2.07	0.55
1:D:449:GLN:CA	1:D:452:TYR:CE2	2.87	0.55
1:D:466:LEU:H	1:D:466:LEU:HD13	1.70	0.55
1:D:479:TYR:CE1	1:D:480:LYS:HE2	2.41	0.55
1:D:550:GLN:N	1:D:551:LYS:NZ	2.53	0.55
1:D:247:LEU:N	1:D:815:TRP:CD1	2.74	0.55
1:D:88:PHE:HB2	1:D:91:GLY:HA3	1.88	0.55
2:E:259:ARG:NH2	2:E:263:TRP:HE1	2.04	0.55
2:E:69:LEU:H	2:E:69:LEU:CD1	2.18	0.55
2:F:234:LYS:NZ	2:F:344:ARG:CD	2.63	0.55
2:F:303:LEU:CG	2:F:337:SER:OG	2.54	0.55
2:F:305:ASN:C	2:F:307:GLY:H	2.09	0.55
2:F:389:ARG:CG	2:F:390:GLY:N	2.69	0.55
1:A:1013:ASP:OD1	1:A:1014:ARG:N	2.36	0.55
1:A:1038:TRP:CE3	1:A:1041:TRP:HA	2.40	0.55
1:A:392:LEU:HG	1:A:395:TYR:CB	2.35	0.55
1:A:475:SER:O	1:A:711:ALA:HA	2.06	0.55
2:C:397:GLN:C	2:C:399:CYS:H	2.10	0.55
2:C:403:PHE:CE2	2:C:407:LEU:HB2	2.41	0.55
2:C:473:ASP:C	2:C:476:ILE:HG23	2.26	0.55
2:C:285:LYS:HB2	2:C:285:LYS:NZ	2.21	0.55
1:D:1001:GLY:C	1:D:1004:LEU:HB2	2.27	0.55
1:D:1031:GLU:OE2	1:D:1092:PHE:CD2	2.59	0.55
1:D:1108:TYR:CE1	1:D:1160:THR:HG22	2.40	0.55
1:D:877:MET:HE1	1:D:1203:PRO:O	2.06	0.55
1:D:173:GLU:OE2	1:D:223:TRP:CD1	2.59	0.55
1:D:177:ARG:HH21	1:D:216:SER:HA	1.70	0.55
1:D:656:TYR:CE1	1:D:743:ASP:CG	2.78	0.55
1:D:659:HIS:CE1	1:D:719:LEU:HD22	2.39	0.55
1:D:615:SER:C	1:D:725:PRO:HG3	2.26	0.55
1:D:744:ILE:N	1:D:744:ILE:HD13	2.21	0.55
1:D:753:PRO:HD3	1:D:770:PHE:CD2	2.41	0.55
1:D:854:ALA:C	1:D:855:VAL:CG2	2.74	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:885:THR:O	1:D:886:LEU:HD23	2.06	0.55
1:D:987:ALA:O	1:D:988:ALA:C	2.43	0.55
2:E:199:LEU:HD12	2:E:200:VAL:H	1.71	0.55
2:E:456:ARG:HG3	2:E:462:MET:N	2.21	0.55
2:F:358:GLU:O	2:F:360:SER:N	2.39	0.55
1:A:1189:LEU:CD2	1:A:1189:LEU:N	2.69	0.55
1:A:370:PRO:O	1:A:371:LYS:CG	2.52	0.55
1:A:421:PRO:HG2	1:A:422:VAL:H	1.71	0.55
1:A:667:GLN:HG3	1:A:668:LEU:N	2.20	0.55
2:B:233:GLU:OE2	2:C:133:HIS:ND1	2.35	0.55
2:B:263:TRP:O	2:B:265:LYS:HG2	2.05	0.55
2:B:341:ASP:OD1	2:B:344:ARG:HD3	2.07	0.55
2:C:339:ASN:O	2:C:340:GLY:O	2.25	0.55
2:E:186:LEU:C	2:E:190:LEU:HD13	2.25	0.55
2:E:315:TYR:HD1	2:E:316:PRO:HD2	1.71	0.55
2:C:221:THR:CG2	2:C:222:LYS:N	2.69	0.55
1:A:487:ASP:O	1:A:491:ASP:CG	2.45	0.55
1:D:857:PRO:CA	1:D:1062:GLU:OE2	2.55	0.55
1:D:421:PRO:HG2	1:D:422:VAL:H	1.71	0.55
1:D:575:LYS:CD	1:D:581:ASP:O	2.55	0.55
1:D:727:ASP:O	1:D:728:THR:CB	2.54	0.55
1:D:735:GLY:HA2	1:D:749:PHE:HA	1.89	0.55
2:E:204:LEU:HD22	2:E:242:PHE:O	2.05	0.55
2:F:115:TRP:NE1	2:F:119:VAL:CG2	2.70	0.55
2:F:238:SER:C	2:F:338:VAL:HG11	2.26	0.55
2:F:406:LEU:N	2:F:406:LEU:CD1	2.69	0.55
1:A:1016:GLU:CB	1:A:1020:ILE:HA	2.36	0.55
1:A:1060:LYS:CE	1:A:1073:PRO:HB2	2.36	0.55
1:A:1171:ASN:N	1:A:1171:ASN:OD1	2.38	0.55
1:A:221:TYR:OH	1:A:397:ALA:CB	2.54	0.55
1:A:665:LYS:O	1:A:669:MET:HG2	2.06	0.55
1:A:77:GLN:OE1	1:A:1174:PRO:HB3	2.06	0.55
1:A:938:THR:O	1:A:939:VAL:CB	2.54	0.55
1:A:932:HIS:CB	1:A:947:LYS:HG2	2.34	0.55
2:C:116:THR:O	2:C:120:VAL:HG13	2.06	0.55
2:C:205:PRO:C	2:C:241:TRP:HE1	2.08	0.55
2:C:245:PRO:C	2:C:248:SER:HB2	2.24	0.55
2:C:301:GLU:OE1	2:C:339:ASN:ND2	2.38	0.55
2:C:308:ASP:N	2:C:308:ASP:OD2	2.39	0.55
2:C:351:TYR:CD1	2:C:355:GLN:HG2	2.42	0.55
2:E:415:PRO:HB2	2:E:417:TYR:CE2	2.42	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:453:ILE:CG2	2:B:468:ILE:HD12	2.33	0.55
2:F:144:ALA:O	2:F:145:PHE:HB2	2.06	0.55
1:D:233:TYR:O	1:D:234:SER:CB	2.51	0.55
2:C:221:THR:HG22	2:C:222:LYS:N	2.21	0.55
1:D:1069:ILE:HG23	1:D:1070:PRO:CG	2.37	0.55
1:D:1113:LEU:HB3	1:D:1117:LYS:HE3	1.88	0.55
1:D:886:LEU:HD22	1:D:1140:LEU:HG	1.88	0.55
1:D:1207:GLU:HA	1:D:1210:TYR:HD2	1.55	0.55
1:D:213:VAL:HG13	1:D:401:TRP:HZ2	1.68	0.55
1:D:251:THR:C	1:D:254:SER:H	2.08	0.55
1:D:455:LEU:HD13	1:D:869:ARG:NH1	2.22	0.55
2:E:378:LEU:CD1	2:E:378:LEU:N	2.69	0.55
2:F:344:ARG:HA	2:F:347:LEU:CG	2.36	0.55
2:F:346:MET:O	2:F:350:LEU:CD1	2.55	0.55
2:F:88:ARG:HH11	2:F:88:ARG:HB2	1.70	0.55
1:A:106:HIS:C	1:A:107:LEU:HD22	2.26	0.55
1:A:203:LEU:CD1	1:A:379:LYS:O	2.54	0.55
1:A:196:VAL:HG11	1:A:404:HIS:HB3	1.89	0.55
1:A:444:TYR:HE2	1:A:874:LEU:HA	1.71	0.55
1:A:586:THR:N	1:A:587:PRO:CD	2.70	0.55
1:A:936:ALA:HB2	1:A:946:ALA:CB	2.35	0.55
2:B:123:GLU:CB	2:C:403:PHE:HZ	2.20	0.55
2:C:192:HIS:C	2:C:196:CYS:HG	2.09	0.55
2:C:291:TYR:OH	2:C:352:ASP:N	2.38	0.55
2:C:402:LEU:HB3	2:C:406:LEU:CD2	2.34	0.55
1:D:509:THR:C	1:D:510:ALA:O	2.38	0.55
1:A:739:TYR:HA	1:A:741:ASP:O	2.06	0.55
1:D:1188:CYS:O	1:D:1189:LEU:HD13	2.06	0.55
1:D:1190:ARG:HH11	1:D:1195:MET:CE	2.17	0.55
1:D:242:ALA:C	1:D:246:PRO:HD3	2.27	0.55
1:D:733:HIS:CE1	1:D:750:PHE:CD1	2.95	0.55
1:D:795:ASN:O	1:D:798:ILE:CG1	2.53	0.55
1:D:802:ARG:HG2	1:D:803:ASN:OD1	2.07	0.55
1:D:279:ARG:HH11	1:D:841:LEU:CB	2.19	0.55
1:D:951:TYR:O	1:D:955:TYR:CD1	2.60	0.55
1:D:967:MET:SD	1:D:978:ALA:HB2	2.47	0.55
2:F:180:LYS:O	2:F:181:LEU:HD23	2.06	0.55
2:F:300:ILE:HG23	2:F:342:LEU:CD1	2.36	0.55
2:F:300:ILE:CG2	2:F:301:GLU:H	2.19	0.55
2:F:397:GLN:HG2	2:F:398:VAL:N	2.22	0.55
2:F:403:PHE:O	2:F:406:LEU:HB2	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:120:VAL:O	2:F:407:LEU:HG	2.07	0.55
1:A:1008:LEU:HD22	1:A:1009:ASN:N	2.22	0.55
1:A:1009:ASN:C	1:A:1010:LEU:HG	2.25	0.55
1:A:1026:ARG:HG2	1:A:1029:GLN:NE2	2.21	0.55
1:A:1069:ILE:N	1:A:1070:PRO:CD	2.69	0.55
1:A:430:MET:HA	1:A:1130:CYS:HB2	1.89	0.55
1:A:1165:ALA:HB2	1:A:1173:LEU:CD1	2.37	0.55
1:A:127:LEU:CD2	1:A:129:PRO:HD3	2.34	0.55
1:A:283:LEU:C	1:A:283:LEU:HD13	2.26	0.55
1:A:386:ARG:HG2	1:A:387:GLU:N	2.21	0.55
1:A:411:LEU:O	1:A:415:LEU:HD13	2.05	0.55
1:A:416:GLU:O	1:A:419:PRO:HD3	2.06	0.55
1:A:442:GLU:OE1	1:A:815:TRP:CZ2	2.60	0.55
1:A:469:ASP:O	1:A:473:LEU:HD23	2.07	0.55
2:B:102:LEU:CD1	2:B:102:LEU:H	2.17	0.55
2:B:439:PHE:O	2:B:440:THR:OG1	2.21	0.55
2:C:290:TYR:CE1	2:C:299:LEU:HD22	2.42	0.55
2:C:351:TYR:O	2:C:353:SER:N	2.38	0.55
2:C:354:PHE:C	2:C:356:LEU:N	2.53	0.55
2:B:452:LEU:HD22	2:B:452:LEU:N	2.21	0.55
1:D:1006:ARG:HA	1:D:1006:ARG:CZ	2.31	0.55
1:D:312:TRP:CZ3	1:D:348:LEU:HB2	2.40	0.55
1:D:607:TRP:NE1	1:D:780:GLN:O	2.40	0.55
1:D:77:GLN:O	1:D:78:MET:CB	2.53	0.55
1:D:903:GLY:HA2	1:D:1002:GLU:HB2	1.86	0.55
2:E:77:HIS:CB	2:E:101:PRO:HD2	2.33	0.55
2:E:109:ASN:O	2:E:378:LEU:HG	2.06	0.55
2:E:211:ILE:HA	2:E:237:ALA:HA	1.89	0.55
2:E:335:VAL:CG2	2:E:336:LEU:N	2.70	0.55
2:F:290:TYR:CE1	2:F:299:LEU:HD22	2.40	0.55
2:F:311:LEU:O	2:F:315:TYR:N	2.32	0.55
2:F:322:LEU:HD22	2:F:323:HIS:C	2.27	0.55
2:F:341:ASP:O	2:F:342:LEU:CD1	2.50	0.55
1:A:1005:VAL:CA	1:A:1008:LEU:CD1	2.67	0.55
1:A:1068:ASP:O	1:A:1070:PRO:HD3	2.07	0.55
1:A:1091:GLU:O	1:A:1092:PHE:CB	2.54	0.55
1:A:1224:ILE:HD11	1:A:1229:GLY:H	1.71	0.55
1:A:414:PHE:O	1:A:417:ARG:CB	2.54	0.55
1:A:439:GLN:HG2	1:A:836:LEU:N	2.22	0.55
1:A:658:LYS:O	1:A:662:GLU:CG	2.53	0.55
1:A:786:ALA:CB	1:A:790:ARG:NH1	2.67	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:916:PHE:O	1:A:920:THR:HA	2.05	0.55
2:C:233:GLU:O	2:C:234:LYS:HG2	2.06	0.55
2:C:243:THR:HB	2:C:247:THR:OG1	2.07	0.55
2:C:342:LEU:O	2:C:342:LEU:HD22	2.06	0.55
2:C:415:PRO:HA	2:C:417:TYR:HE2	1.72	0.55
2:E:284:ARG:O	2:E:285:LYS:CB	2.53	0.55
1:D:1060:LYS:CE	1:D:1060:LYS:HA	2.36	0.55
1:D:77:GLN:O	1:D:78:MET:HB2	2.07	0.55
1:D:892:ASP:OD2	1:D:893:SER:N	2.40	0.55
1:D:903:GLY:O	1:D:906:HIS:N	2.40	0.55
2:E:119:VAL:HA	2:E:125:VAL:HB	1.87	0.55
2:E:201:ASN:CG	2:F:422:GLN:HB2	2.26	0.55
2:E:114:TRP:NE1	2:E:259:ARG:NH2	2.49	0.55
2:E:266:PHE:CD1	2:E:376:PRO:HG3	2.42	0.55
2:E:338:VAL:O	2:E:338:VAL:HG12	2.06	0.55
1:D:1209:ARG:CG	2:F:249:ASN:HB3	2.32	0.55
2:F:312:LEU:O	2:F:313:HIS:C	2.44	0.55
2:F:407:LEU:HD23	2:F:407:LEU:C	2.26	0.55
2:F:426:GLU:OE2	2:F:429:TYR:HD1	1.89	0.55
2:F:469:SER:C	2:F:471:LEU:HD12	2.27	0.55
1:A:1008:LEU:CD2	1:A:1008:LEU:C	2.74	0.55
1:A:1013:ASP:CG	1:A:1014:ARG:H	2.09	0.55
1:A:1017:GLY:C	1:A:1019:TRP:H	2.09	0.55
1:A:1030:ARG:HD3	1:A:1091:GLU:CD	2.27	0.55
1:A:312:TRP:CZ2	1:A:346:ASP:C	2.80	0.55
1:A:221:TYR:OH	1:A:397:ALA:HB1	2.07	0.55
1:A:897:TRP:O	1:A:897:TRP:CD1	2.59	0.55
1:A:931:LEU:HA	1:A:934:LYS:HE3	1.87	0.55
2:C:390:GLY:HA2	2:C:392:THR:N	2.21	0.55
2:C:444:THR:N	2:C:447:THR:HG23	2.20	0.55
2:E:418:LEU:CD2	2:F:122:ARG:NH2	2.67	0.55
1:D:1028:VAL:HG12	1:D:1028:VAL:O	2.07	0.55
1:D:121:PRO:C	1:D:122:ASP:OD2	2.44	0.55
1:D:1221:TYR:CA	1:D:1224:ILE:HG22	2.33	0.55
1:D:192:GLU:HG3	1:D:195:LEU:HD12	1.89	0.55
1:D:211:LEU:HG	1:D:221:TYR:CE1	2.42	0.55
1:D:259:ARG:CG	1:D:261:TRP:HD1	2.17	0.55
1:D:271:VAL:N	1:D:294:THR:OG1	2.39	0.55
1:D:407:PHE:CG	1:D:408:GLN:N	2.75	0.55
1:D:591:LEU:HD11	1:D:594:LEU:CG	2.20	0.55
1:D:598:VAL:HA	1:D:601:LYS:HD3	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:776:ASP:OD1	1:D:777:GLY:N	2.39	0.55
1:D:916:PHE:O	1:D:920:THR:HA	2.04	0.55
1:D:954:ILE:HG22	1:D:955:TYR:CG	2.42	0.55
1:D:995:TYR:N	1:D:996:ARG:NH1	2.50	0.55
2:E:239:LEU:HB3	2:E:338:VAL:HB	1.89	0.55
2:E:244:PRO:O	2:E:248:SER:N	2.40	0.55
2:E:301:GLU:HG2	2:E:303:LEU:CD1	2.36	0.55
2:F:112:ALA:O	2:F:116:THR:HG22	2.06	0.55
2:F:244:PRO:N	2:F:247:THR:CG2	2.70	0.55
2:F:358:GLU:HG2	2:F:359:ASN:N	2.22	0.55
2:F:431:LYS:NZ	2:F:435:MET:CG	2.70	0.55
2:F:78:PHE:CD2	2:F:79:LEU:HD22	2.41	0.55
1:A:1129:PHE:CE2	1:A:1131:ILE:HB	2.42	0.55
1:A:1131:ILE:HG13	1:A:1133:ILE:HG22	1.88	0.55
1:A:125:LEU:HD23	1:A:125:LEU:N	2.00	0.55
1:A:267:VAL:CG1	1:A:269:HIS:CE1	2.90	0.55
1:A:379:LYS:HD3	1:A:379:LYS:H	1.70	0.55
1:A:823:ARG:HB3	1:A:825:VAL:CG2	2.13	0.55
1:A:851:THR:C	1:A:852:ARG:HG3	2.27	0.55
1:A:859:TRP:N	1:A:859:TRP:CD1	2.73	0.55
2:C:245:PRO:O	2:C:248:SER:CB	2.45	0.55
2:C:243:THR:HB	2:C:247:THR:CG2	2.37	0.55
2:C:236:GLU:CA	2:C:341:ASP:CB	2.69	0.55
1:A:784:GLY:CA	2:C:362:THR:O	2.55	0.55
2:C:432:TYR:HB3	2:C:437:ILE:HD11	1.88	0.55
1:D:1165:ALA:C	1:D:1166:TYR:CD1	2.77	0.55
1:D:1163:MET:CE	1:D:1167:LYS:HG2	2.37	0.55
1:D:1138:ARG:HH21	1:D:1191:LYS:HZ3	1.55	0.55
1:D:278:ILE:HD13	1:D:282:TYR:OH	2.07	0.55
1:D:550:GLN:OE1	1:D:551:LYS:CE	2.54	0.55
1:D:576:LEU:HG	1:D:579:ARG:HE	1.70	0.55
1:D:769:ASP:O	1:D:773:LYS:NZ	2.29	0.55
1:D:867:PRO:O	1:D:868:ASP:CB	2.55	0.55
1:D:874:LEU:O	1:D:878:VAL:CG2	2.41	0.55
1:D:951:TYR:O	1:D:955:TYR:CE1	2.59	0.55
2:E:431:LYS:CA	2:E:434:GLU:HG2	2.34	0.55
2:F:243:THR:C	2:F:334:CYS:HB2	2.26	0.55
2:F:351:TYR:C	2:F:355:GLN:CB	2.64	0.55
1:A:1209:ARG:HH22	2:C:256:LEU:CG	2.20	0.55
1:A:433:SER:HB2	1:A:1129:PHE:HE1	1.72	0.55
1:A:460:LYS:O	1:A:463:LEU:HB3	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:616:GLU:CG	1:A:724:GLY:CA	2.84	0.55
1:A:825:VAL:O	1:A:829:PRO:CG	2.55	0.55
2:B:378:LEU:O	2:B:379:ALA:C	2.46	0.55
2:B:432:TYR:HA	2:B:437:ILE:CD1	2.37	0.55
2:C:241:TRP:HE3	2:C:255:TRP:HE1	1.52	0.55
2:C:393:LEU:HD23	2:C:396:ARG:CG	2.26	0.55
2:C:398:VAL:C	2:C:401:GLY:N	2.60	0.55
2:C:441:VAL:O	2:C:441:VAL:HG23	2.07	0.55
2:C:72:ILE:O	2:C:75:ARG:O	2.25	0.55
1:A:735:GLY:C	1:A:736:ASN:CG	2.66	0.55
1:D:401:TRP:HD1	1:D:404:HIS:HD2	1.53	0.55
1:D:402:ALA:O	1:D:406:VAL:CG1	2.55	0.55
1:D:613:HIS:CE1	1:D:717:LEU:CD1	2.89	0.55
1:D:659:HIS:CG	1:D:719:LEU:HD22	2.42	0.55
1:D:732:TYR:CG	1:D:755:LYS:NZ	2.75	0.55
1:D:891:VAL:CG1	1:D:1135:ASP:CA	2.85	0.55
1:D:922:GLN:OE1	1:D:934:LYS:NZ	2.39	0.55
2:E:263:TRP:C	2:E:265:LYS:N	2.58	0.55
2:E:324:GLY:O	2:E:331:VAL:HG13	2.06	0.55
2:E:380:PRO:CA	2:E:438:LEU:HD12	2.37	0.55
2:F:115:TRP:O	2:F:118:VAL:HG12	2.06	0.55
2:F:116:THR:O	2:F:120:VAL:HG13	2.07	0.55
2:F:240:VAL:CA	2:F:336:LEU:HB3	2.36	0.55
2:F:342:LEU:CD2	2:F:342:LEU:C	2.74	0.55
2:F:344:ARG:HA	2:F:347:LEU:CD2	2.34	0.55
1:A:996:ARG:NH1	1:A:1003:TRP:C	2.60	0.55
1:A:955:TYR:HD2	1:A:1102:GLN:HE22	1.52	0.55
1:A:430:MET:O	1:A:1130:CYS:SG	2.64	0.55
1:A:267:VAL:CG1	1:A:269:HIS:HE1	2.20	0.55
1:A:358:LEU:HD12	1:A:407:PHE:CE1	2.42	0.55
1:A:572:TRP:O	1:A:574:ARG:N	2.40	0.55
1:A:613:HIS:CG	1:A:717:LEU:HG	2.41	0.55
1:A:732:TYR:CD1	1:A:755:LYS:HG2	2.42	0.55
1:A:610:PHE:CE1	1:A:972:ARG:NH2	2.75	0.55
2:B:78:PHE:CD2	2:B:102:LEU:HD22	2.42	0.55
2:B:182:ARG:O	2:B:182:ARG:HG2	2.06	0.55
2:B:348:ALA:C	2:B:350:LEU:N	2.61	0.55
2:B:69:LEU:O	2:B:73:CYS:SG	2.64	0.55
2:C:208:LEU:CD1	2:C:208:LEU:O	2.54	0.55
2:C:244:PRO:N	2:C:247:THR:HG21	2.22	0.55
2:C:344:ARG:O	2:C:347:LEU:HB2	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:387:VAL:HG11	2:C:395:LEU:O	2.07	0.55
2:C:387:VAL:CG1	2:C:399:CYS:HB2	2.37	0.55
2:C:406:LEU:CA	2:C:409:ASN:ND2	2.61	0.55
2:B:388:GLY:O	2:B:444:THR:HA	2.07	0.55
1:D:1111:LEU:HD22	1:D:1164:PHE:HE1	1.68	0.54
1:D:203:LEU:C	1:D:203:LEU:CD2	2.74	0.54
1:D:273:PHE:CA	1:D:276:ALA:H	2.19	0.54
1:D:439:GLN:NE2	1:D:837:TYR:CE2	2.75	0.54
1:D:460:LYS:HG2	1:D:461:LYS:N	2.22	0.54
1:D:768:LYS:C	1:D:772:PRO:CD	2.74	0.54
1:D:771:LEU:CD1	1:D:771:LEU:N	2.70	0.54
1:D:904:ASP:OD1	1:D:915:ALA:CB	2.55	0.54
2:E:109:ASN:HB2	2:E:378:LEU:CB	2.37	0.54
2:E:427:GLN:HA	2:E:430:SER:OG	2.07	0.54
2:F:118:VAL:CG1	2:F:119:VAL:N	2.70	0.54
2:F:123:GLU:OE1	2:F:206:TYR:HA	2.07	0.54
1:D:1209:ARG:CZ	2:F:252:LEU:HB3	2.37	0.54
2:F:375:HIS:CD2	2:F:377:CYS:H	2.24	0.54
2:F:460:THR:CG2	2:F:462:MET:HB2	2.35	0.54
1:A:1108:TYR:HE2	1:A:1112:MET:CE	2.20	0.54
1:A:885:THR:HB	1:A:1146:ARG:HH21	1.72	0.54
1:A:196:VAL:C	1:A:401:TRP:CH2	2.81	0.54
1:A:363:ARG:O	1:A:365:TYR:CE2	2.59	0.54
1:A:616:GLU:O	1:A:616:GLU:HG3	2.06	0.54
1:A:788:GLY:O	1:A:792:LEU:HG	2.07	0.54
1:A:815:TRP:HA	1:A:838:GLY:HA2	1.87	0.54
1:A:79:LEU:CB	1:A:89:GLY:CA	2.84	0.54
1:A:974:THR:HG23	1:A:975:GLN:H	1.72	0.54
2:B:283:GLY:O	2:B:305:ASN:O	2.25	0.54
2:B:456:ARG:HG2	2:B:463:LYS:HG3	1.89	0.54
2:B:75:ARG:HD2	2:B:434:GLU:CD	2.28	0.54
2:C:239:LEU:C	2:C:338:VAL:HG12	2.26	0.54
1:D:1151:LEU:HD12	1:D:1230:SER:HA	1.89	0.54
1:D:1190:ARG:NH1	1:D:1195:MET:HE3	2.21	0.54
1:D:284:ILE:H	1:D:284:ILE:CD1	2.02	0.54
1:D:612:LEU:HB3	1:D:621:GLY:N	2.22	0.54
1:D:739:TYR:O	1:D:741:ASP:O	2.25	0.54
1:D:808:ILE:O	1:D:811:GLN:OE1	2.25	0.54
1:D:879:GLN:O	1:D:1189:LEU:CD2	2.54	0.54
2:E:239:LEU:O	2:E:240:VAL:HB	2.06	0.54
2:E:382:LYS:CG	2:E:383:VAL:H	2.20	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:385:LEU:HD13	2:F:399:CYS:SG	2.47	0.54
2:F:402:LEU:O	2:F:403:PHE:C	2.45	0.54
1:A:1006:ARG:N	1:A:1006:ARG:HH11	2.06	0.54
1:A:189:ILE:HD12	1:A:258:GLN:HA	1.89	0.54
1:A:456:GLN:O	1:A:460:LYS:HD3	2.07	0.54
1:A:490:TRP:CH2	1:A:598:VAL:CB	2.85	0.54
1:A:496:LYS:NZ	1:A:570:PRO:O	2.37	0.54
1:A:730:PRO:HD3	1:A:761:ASN:OD1	2.08	0.54
2:B:376:PRO:O	2:B:378:LEU:N	2.39	0.54
2:B:75:ARG:C	2:B:76:ARG:HG2	2.28	0.54
2:C:200:VAL:O	2:C:201:ASN:HB2	2.06	0.54
2:C:266:PHE:HB3	2:C:378:LEU:HD11	1.88	0.54
2:C:342:LEU:CB	2:C:344:ARG:HB2	2.37	0.54
1:D:1157:ASN:OD1	1:D:1161:ARG:NH2	2.40	0.54
1:D:260:ASP:C	1:D:260:ASP:OD2	2.45	0.54
1:D:396:CYS:O	1:D:400:VAL:HG23	2.06	0.54
1:D:437:VAL:HG12	1:D:438:ASN:N	2.21	0.54
1:D:484:TRP:CD1	1:D:488:LEU:HG	2.42	0.54
1:D:618:HIS:CE1	1:D:761:ASN:HB3	2.42	0.54
1:D:933:SER:OG	1:D:943:ARG:CZ	2.55	0.54
2:E:318:ASN:O	2:E:319:VAL:C	2.46	0.54
2:E:385:LEU:HD13	2:E:441:VAL:HB	1.88	0.54
2:F:307:GLY:C	2:F:308:ASP:OD2	2.46	0.54
2:F:309:HIS:HB3	2:F:310:GLU:OE1	2.06	0.54
2:F:425:LEU:O	2:F:425:LEU:HD12	2.08	0.54
1:A:371:LYS:HA	1:A:374:ARG:HD2	1.88	0.54
1:A:420:HIS:CE1	1:A:422:VAL:HG23	2.42	0.54
1:A:652:ILE:HG23	1:A:655:LEU:HD11	1.89	0.54
2:B:78:PHE:CD2	2:B:102:LEU:HB2	2.42	0.54
2:B:67:GLU:N	2:B:70:LEU:HB2	2.22	0.54
2:C:357:THR:C	2:C:359:ASN:N	2.61	0.54
1:A:520:ALA:N	1:A:521:PRO:CD	2.71	0.54
1:D:999:ASP:C	1:D:1001:GLY:N	2.59	0.54
1:D:1063:SER:OG	1:D:1066:THR:HB	2.08	0.54
1:D:1153:LEU:N	1:D:1153:LEU:HD23	2.22	0.54
1:D:143:ALA:HB1	1:D:421:PRO:O	2.08	0.54
1:D:257:THR:CG2	1:D:257:THR:O	2.49	0.54
1:D:421:PRO:O	1:D:425:ALA:CB	2.55	0.54
1:D:542:ASP:O	1:D:546:ARG:HB2	2.06	0.54
1:D:601:LYS:HZ2	1:D:721:ALA:HA	1.72	0.54
1:D:744:ILE:CB	1:D:745:PRO:HD3	2.36	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:734:HIS:ND1	1:D:752:LEU:HD11	2.22	0.54
1:D:776:ASP:N	1:D:796:LYS:NZ	2.55	0.54
1:D:774:MET:O	1:D:776:ASP:O	2.25	0.54
1:D:807:ARG:HB2	1:D:858:THR:CG2	2.37	0.54
1:D:873:GLU:CD	1:D:1203:PRO:HG3	2.28	0.54
2:E:240:VAL:HG12	2:E:242:PHE:CZ	2.43	0.54
2:E:460:THR:O	2:E:462:MET:HE2	2.07	0.54
2:E:127:PRO:O	2:F:101:PRO:HA	2.07	0.54
2:F:381:ILE:CG2	2:F:437:ILE:HG22	2.38	0.54
2:F:413:VAL:C	2:F:415:PRO:HD2	2.27	0.54
1:A:1008:LEU:C	1:A:1011:PRO:HD3	2.28	0.54
1:A:1043:VAL:C	1:A:1046:GLU:H	2.10	0.54
1:A:885:THR:HB	1:A:1146:ARG:CZ	2.37	0.54
1:A:188:ALA:C	1:A:189:ILE:HG12	2.27	0.54
1:A:196:VAL:CG1	1:A:401:TRP:CZ3	2.91	0.54
1:A:273:PHE:CA	1:A:276:ALA:HB3	2.37	0.54
1:A:915:ALA:O	1:A:919:MET:N	2.26	0.54
2:B:307:GLY:HA2	2:B:334:CYS:HB3	1.90	0.54
2:C:395:LEU:CD2	2:C:443:VAL:HB	2.37	0.54
1:A:503:VAL:O	1:A:504:LYS:HB3	2.07	0.54
2:E:280:ASP:O	2:E:304:TRP:CH2	2.60	0.54
1:A:639:THR:HB	1:A:640:LEU:CD2	2.27	0.54
2:E:271:SER:C	2:E:273:PHE:H	2.10	0.54
1:D:1027:LYS:HB2	1:D:1096:ARG:CG	2.37	0.54
1:D:1122:GLU:O	1:D:1123:PHE:HB2	2.08	0.54
1:D:193:ARG:CA	1:D:193:ARG:HH11	2.19	0.54
1:D:279:ARG:HH11	1:D:841:LEU:HD13	1.71	0.54
1:D:308:GLN:NE2	1:D:1091:GLU:N	2.55	0.54
1:D:429:GLU:HG3	1:D:1128:ARG:HH22	1.68	0.54
1:D:434:TYR:HA	1:D:841:LEU:HD13	1.89	0.54
1:D:483:PRO:HG3	1:D:712:VAL:CG1	2.38	0.54
1:D:581:ASP:OD1	1:D:592:LEU:HD13	2.08	0.54
1:D:730:PRO:HB3	1:D:761:ASN:CA	2.38	0.54
1:D:945:HIS:HA	1:D:948:ILE:HD11	1.88	0.54
2:E:427:GLN:O	2:E:430:SER:HB2	2.08	0.54
2:F:129:ASP:HB3	2:F:131:LEU:HD22	1.88	0.54
2:F:383:VAL:CG2	2:F:384:ALA:N	2.70	0.54
2:F:385:LEU:HD11	2:F:402:LEU:HB3	1.90	0.54
2:F:407:LEU:C	2:F:407:LEU:CD2	2.76	0.54
2:F:473:ASP:C	2:F:476:ILE:HG23	2.27	0.54
1:A:112:LEU:HD12	1:A:112:LEU:N	2.21	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1140:LEU:CD2	1:A:1141:VAL:N	2.71	0.54
1:A:175:TRP:HD1	1:A:183:GLU:HB2	1.71	0.54
1:A:371:LYS:NZ	1:A:400:VAL:CG2	2.70	0.54
1:A:611:PRO:HG3	1:A:671:GLN:CG	2.37	0.54
2:B:285:LYS:C	2:B:305:ASN:HD22	2.11	0.54
2:B:69:LEU:HA	2:B:72:ILE:HG12	1.90	0.54
2:C:208:LEU:HD12	2:C:208:LEU:C	2.26	0.54
2:C:381:ILE:CG2	2:C:437:ILE:HG22	2.38	0.54
2:C:456:ARG:NH2	2:C:465:MET:SD	2.80	0.54
2:E:187:HIS:HA	2:E:190:LEU:CD2	2.31	0.54
1:D:503:VAL:C	1:D:505:LYS:N	2.46	0.54
2:B:469:SER:O	2:B:471:LEU:N	2.40	0.54
2:B:122:ARG:HB3	2:B:122:ARG:HH11	1.72	0.54
2:B:82:SER:HB3	2:B:85:GLN:CG	2.38	0.54
1:D:1014:ARG:HD3	1:D:1026:ARG:CD	2.37	0.54
1:D:1052:GLY:O	1:D:1056:GLU:N	2.40	0.54
1:D:888:GLY:CA	1:D:1138:ARG:HD3	2.18	0.54
1:D:298:HIS:CD2	1:D:411:LEU:CD2	2.90	0.54
1:D:348:LEU:HD13	1:D:350:ILE:HD11	1.88	0.54
1:D:437:VAL:CA	1:D:441:TRP:HD1	2.16	0.54
1:D:459:MET:H	1:D:460:LYS:NZ	2.00	0.54
1:D:641:GLU:C	1:D:645:VAL:CG2	2.68	0.54
1:D:735:GLY:HA2	1:D:750:PHE:N	2.23	0.54
1:D:779:LEU:HD13	1:D:779:LEU:O	2.07	0.54
1:D:854:ALA:O	1:D:855:VAL:CG2	2.55	0.54
1:D:901:VAL:O	1:D:902:LEU:C	2.46	0.54
1:D:949:PHE:CD1	1:D:949:PHE:C	2.80	0.54
2:E:101:PRO:N	2:F:128:VAL:CG2	2.71	0.54
2:E:199:LEU:CD1	2:E:200:VAL:H	2.21	0.54
2:E:402:LEU:O	2:E:403:PHE:C	2.46	0.54
2:F:131:LEU:HB3	2:F:133:HIS:ND1	2.22	0.54
2:F:256:LEU:O	2:F:257:ARG:C	2.46	0.54
1:A:1006:ARG:CZ	1:A:1010:LEU:HD11	2.35	0.54
1:A:1198:LYS:C	1:A:1199:THR:HG23	2.27	0.54
1:A:385:ILE:CG2	1:A:386:ARG:N	2.70	0.54
1:A:196:VAL:HG12	1:A:401:TRP:CZ3	2.43	0.54
1:A:613:HIS:O	1:A:614:TYR:CB	2.55	0.54
1:A:659:HIS:CD2	1:A:720:THR:HB	2.43	0.54
1:A:845:VAL:CG1	1:A:848:GLY:HA3	2.37	0.54
2:B:315:TYR:HB2	2:B:322:LEU:HD11	1.90	0.54
2:B:417:TYR:OH	2:C:122:ARG:CD	2.40	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:373:LYS:CE	2:B:459:ASP:HA	2.33	0.54
2:C:201:ASN:O	2:C:202:LYS:HB3	2.08	0.54
2:C:308:ASP:OD1	2:C:333:PRO:C	2.46	0.54
2:C:197:LEU:CD1	2:C:322:LEU:HD21	2.37	0.54
2:C:241:TRP:CE3	2:C:336:LEU:HD12	2.43	0.54
1:A:560:PRO:HD2	2:C:452:LEU:CD2	2.37	0.54
2:C:466:MET:O	2:C:467:HIS:C	2.45	0.54
1:D:667:GLN:C	1:D:671:GLN:HG2	2.27	0.54
2:B:217:PRO:HB2	2:B:231:ILE:CG2	2.38	0.54
2:E:352:ASP:CG	2:E:352:ASP:O	2.45	0.54
1:D:301:ILE:O	1:D:1024:ASP:CB	2.55	0.54
1:D:1145:ASP:HA	1:D:1148:ARG:HD3	1.90	0.54
1:D:148:LEU:CB	1:D:149:PRO:HD3	2.38	0.54
1:D:286:GLY:O	1:D:287:SER:CB	2.56	0.54
1:D:371:LYS:NZ	1:D:393:MET:HA	2.21	0.54
1:D:627:ARG:O	1:D:628:ARG:CB	2.56	0.54
1:D:735:GLY:CA	1:D:749:PHE:CA	2.85	0.54
1:D:901:VAL:CG2	1:D:902:LEU:N	2.70	0.54
2:E:214:CYS:HB2	2:E:236:GLU:CB	2.37	0.54
2:E:428:LEU:CD1	2:E:428:LEU:N	2.60	0.54
2:E:91:LEU:C	2:E:93:SER:N	2.61	0.54
1:D:1209:ARG:CZ	2:F:252:LEU:HD22	2.38	0.54
1:A:1015:THR:CA	1:A:1016:GLU:OE2	2.55	0.54
1:A:850:ILE:CG1	1:A:1099:TRP:NE1	2.71	0.54
1:A:175:TRP:CG	1:A:184:ALA:HA	2.43	0.54
1:A:297:MET:HE1	1:A:415:LEU:CD2	2.38	0.54
1:A:730:PRO:HG3	1:A:760:CYS:O	2.08	0.54
1:A:866:ARG:HD3	1:A:866:ARG:C	2.27	0.54
1:A:982:ALA:O	1:A:986:TYR:CD1	2.61	0.54
2:B:112:ALA:O	2:B:115:TRP:CB	2.56	0.54
2:B:400:GLN:HG2	2:B:417:TYR:CE2	2.43	0.54
2:C:326:ASP:OD2	2:C:327:GLY:N	2.39	0.54
1:A:784:GLY:CA	2:C:363:ARG:CD	2.85	0.54
2:C:483:LYS:HE2	2:C:483:LYS:C	2.27	0.54
2:C:78:PHE:CD2	2:C:79:LEU:HD22	2.42	0.54
2:B:447:THR:HG23	2:B:452:LEU:H	1.70	0.54
2:F:221:THR:CG2	2:F:222:LYS:N	2.71	0.54
2:F:328:ARG:HG3	2:F:329:LYS:HD2	1.90	0.54
1:D:422:VAL:CG1	1:D:1111:LEU:HD12	2.36	0.54
1:D:1147:TYR:CD2	1:D:1147:TYR:N	2.74	0.54
1:D:278:ILE:HA	1:D:291:PHE:CD1	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:354:ASN:O	1:D:355:SER:OG	2.19	0.54
1:D:548:CYS:N	1:D:551:LYS:HD3	2.22	0.54
1:D:972:ARG:C	1:D:974:THR:HG23	2.28	0.54
2:E:193:TYR:O	2:E:197:LEU:HB3	2.08	0.54
2:E:299:LEU:HD11	2:E:302:THR:HG22	1.89	0.54
2:F:400:GLN:C	2:F:400:GLN:CD	2.66	0.54
1:A:1043:VAL:O	1:A:1043:VAL:HG12	2.08	0.54
1:A:1111:LEU:O	1:A:1112:MET:C	2.46	0.54
1:A:82:GLY:O	1:A:85:GLU:HG3	2.07	0.54
1:A:879:GLN:O	1:A:1187:ARG:O	2.26	0.54
2:B:252:LEU:O	2:B:255:TRP:HB2	2.07	0.54
2:B:204:LEU:HD11	2:B:333:PRO:HB3	1.88	0.54
2:B:100:GLY:HA2	2:C:128:VAL:HG11	1.90	0.54
2:C:234:LYS:NZ	2:C:344:ARG:CD	2.61	0.54
2:C:358:GLU:HA	2:C:361:PHE:CB	2.38	0.54
2:C:395:LEU:HD11	2:C:444:THR:CA	2.36	0.54
2:C:81:GLY:O	2:C:86:LEU:HD21	2.07	0.54
1:D:585:TRP:CB	1:D:587:PRO:HD2	2.23	0.54
2:B:476:ILE:O	2:B:480:SER:N	2.36	0.54
2:B:277:ASP:HA	2:B:287:ASN:CG	2.28	0.54
2:F:146:ARG:CB	2:F:228:VAL:HG11	2.38	0.54
2:F:328:ARG:C	2:F:329:LYS:HD2	2.27	0.54
2:E:218:VAL:O	2:E:230:SER:OG	2.22	0.54
1:A:977:GLU:HA	1:A:980:GLU:OE2	2.08	0.54
1:D:1224:ILE:HG12	1:D:1227:THR:O	2.07	0.54
1:D:381:THR:HG23	1:D:385:ILE:HB	1.90	0.54
1:D:395:TYR:H	1:D:395:TYR:HD1	1.56	0.54
1:D:744:ILE:CB	1:D:745:PRO:CD	2.85	0.54
1:D:605:LEU:CD2	1:D:781:ALA:CB	2.73	0.54
1:D:823:ARG:CZ	1:D:824:ALA:HB3	2.38	0.54
1:D:859:TRP:N	1:D:859:TRP:CD1	2.76	0.54
1:D:865:ALA:HB1	1:D:872:SER:HG	1.71	0.54
2:E:99:PHE:CD1	2:E:107:ARG:NH2	2.76	0.54
2:E:189:ALA:HB3	2:E:240:VAL:HG21	1.89	0.54
2:E:115:TRP:CZ3	2:E:210:GLN:HA	2.40	0.54
2:E:96:HIS:C	2:E:98:GLY:H	2.11	0.54
2:E:419:GLU:CA	2:F:203:ARG:HH22	2.20	0.54
2:F:447:THR:CG2	2:F:448:LEU:HG	2.35	0.54
1:A:1128:ARG:O	1:A:1140:LEU:N	2.40	0.54
1:A:113:TRP:HA	1:A:116:PRO:CD	2.37	0.54
1:A:298:HIS:C	1:A:298:HIS:ND1	2.61	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:792:LEU:O	1:A:796:LYS:HG3	2.08	0.54
1:A:442:GLU:OE1	1:A:815:TRP:HZ2	1.91	0.54
1:A:964:ARG:O	1:A:968:GLN:CG	2.56	0.54
2:B:249:ASN:HA	2:B:252:LEU:HD22	1.89	0.54
2:B:265:LYS:HD3	2:B:349:TYR:OH	2.08	0.54
2:B:265:LYS:O	2:B:266:PHE:HB2	2.08	0.54
2:B:267:ALA:HB1	2:B:270:PRO:HB2	1.90	0.54
2:B:418:LEU:C	2:B:418:LEU:HD23	2.28	0.54
2:C:314:MET:HB3	2:C:315:TYR:CD2	2.43	0.54
2:C:336:LEU:HD23	2:C:337:SER:O	2.08	0.54
1:A:645:VAL:HA	1:A:650:ARG:CD	2.38	0.54
1:A:307:PHE:CD2	1:A:1051:GLY:HA3	2.43	0.54
1:D:1113:LEU:CD2	1:D:1113:LEU:N	2.71	0.54
1:D:1129:PHE:CE2	1:D:1139:TYR:CE2	2.96	0.54
1:D:1134:HIS:O	1:D:1134:HIS:CD2	2.60	0.54
1:D:385:ILE:O	1:D:388:ASN:HB2	2.08	0.54
1:D:543:VAL:O	1:D:547:ALA:CB	2.56	0.54
1:D:603:MET:O	1:D:605:LEU:HG	2.07	0.54
1:D:845:VAL:H	1:D:855:VAL:HB	1.73	0.54
1:D:853:ARG:NH1	1:D:1102:GLN:HA	2.23	0.54
2:E:353:SER:HG	2:E:372:LEU:HD11	1.73	0.54
2:E:75:ARG:O	2:E:76:ARG:HB2	2.08	0.54
2:F:311:LEU:CA	2:F:314:MET:HB2	2.38	0.54
2:F:241:TRP:CG	2:F:336:LEU:HD12	2.41	0.54
2:F:344:ARG:HA	2:F:347:LEU:CB	2.36	0.54
2:F:74:GLN:HE21	2:F:75:ARG:HH21	1.56	0.54
1:A:1046:GLU:OE1	1:A:1049:TRP:CG	2.60	0.54
1:A:136:ASP:HA	1:A:139:PHE:HB2	1.88	0.54
1:A:196:VAL:HG13	1:A:269:HIS:HD2	1.67	0.54
2:B:116:THR:HA	2:B:120:VAL:CB	2.38	0.54
2:B:200:VAL:O	2:B:202:LYS:N	2.41	0.54
2:B:267:ALA:HB1	2:B:270:PRO:HD2	1.89	0.54
2:C:305:ASN:HA	2:C:335:VAL:CG1	2.38	0.54
2:C:371:VAL:HG23	2:C:433:ASP:CB	2.37	0.54
2:C:444:THR:O	2:C:445:GLU:C	2.47	0.54
1:A:556:THR:HG22	2:C:467:HIS:CD2	2.43	0.54
2:C:466:MET:CE	2:C:471:LEU:HB3	2.38	0.54
1:A:563:PRO:C	1:A:564:GLN:NE2	2.60	0.54
1:A:733:HIS:C	1:A:735:GLY:N	2.60	0.54
1:D:885:THR:HG22	1:D:1186:ASP:C	2.28	0.53
1:D:1189:LEU:CD2	1:D:1189:LEU:N	2.68	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1204:THR:C	1:D:1207:GLU:HB3	2.29	0.53
1:D:441:TRP:CZ3	1:D:1210:TYR:CE1	2.96	0.53
1:D:211:LEU:O	1:D:212:ALA:HB2	2.08	0.53
1:D:189:ILE:HD13	1:D:258:GLN:HB3	1.91	0.53
1:D:269:HIS:HB2	1:D:274:ASP:OD1	2.08	0.53
1:D:279:ARG:HH21	1:D:844:VAL:HG22	1.72	0.53
1:D:472:GLN:O	1:D:473:LEU:C	2.47	0.53
1:D:495:PHE:O	1:D:496:LYS:CD	2.48	0.53
1:D:642:SER:C	1:D:645:VAL:HB	2.29	0.53
2:E:123:GLU:C	2:E:125:VAL:N	2.59	0.53
2:E:441:VAL:CG1	2:E:442:LEU:N	2.71	0.53
2:F:208:LEU:C	2:F:208:LEU:HD12	2.28	0.53
2:F:371:VAL:HG23	2:F:433:ASP:CB	2.38	0.53
2:F:454:HIS:NE2	2:F:472:LYS:CD	2.70	0.53
2:F:80:SER:OG	2:F:99:PHE:O	2.25	0.53
1:A:1169:GLY:O	1:A:1170:LEU:CB	2.53	0.53
1:A:244:LEU:O	1:A:247:LEU:CD2	2.56	0.53
1:A:316:LYS:HD3	1:A:350:ILE:HG12	1.89	0.53
1:A:301:ILE:CD1	1:A:414:PHE:CB	2.86	0.53
1:A:450:GLY:O	1:A:454:GLU:HG3	2.09	0.53
1:A:513:LEU:O	1:A:568:GLY:CA	2.56	0.53
1:A:498:LYS:CE	1:A:518:ALA:H	2.09	0.53
1:A:549:LEU:O	1:A:553:LYS:HG3	2.08	0.53
1:A:582:ASP:OD1	1:A:592:LEU:HD22	2.08	0.53
1:A:783:PRO:CG	2:C:363:ARG:CZ	2.84	0.53
1:A:827:ARG:NH1	1:A:827:ARG:HG2	2.19	0.53
2:B:182:ARG:HG2	2:B:214:CYS:HA	1.90	0.53
2:B:206:TYR:HA	2:B:241:TRP:CZ3	2.43	0.53
2:B:284:ARG:NE	2:B:284:ARG:HA	2.22	0.53
2:B:78:PHE:CE1	2:B:103:GLY:HA2	2.38	0.53
2:C:253:ASP:HA	2:C:256:LEU:HD23	1.90	0.53
2:C:375:HIS:HB3	2:C:378:LEU:CD1	2.35	0.53
2:C:416:GLY:HA2	2:C:419:GLU:CB	2.37	0.53
2:C:446:THR:CB	2:C:450:ASN:OD1	2.51	0.53
2:E:260:LEU:CD1	2:E:264:ARG:HD2	2.38	0.53
1:D:1116:MET:HE2	1:D:1156:THR:OG1	2.08	0.53
1:D:1131:ILE:HD11	1:D:1138:ARG:C	2.28	0.53
1:D:456:GLN:O	1:D:460:LYS:NZ	2.41	0.53
1:D:571:GLY:O	1:D:572:TRP:HB2	2.08	0.53
1:D:731:SER:O	1:D:732:TYR:HB2	2.09	0.53
2:E:127:PRO:HG2	2:F:101:PRO:HA	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:483:LYS:NZ	2:E:483:LYS:HB2	2.23	0.53
2:F:350:LEU:N	2:F:350:LEU:CD1	2.56	0.53
2:F:371:VAL:HG23	2:F:433:ASP:O	2.08	0.53
2:F:454:HIS:CG	2:F:466:MET:HB2	2.40	0.53
1:A:1115:ALA:O	1:A:1118:TRP:HB3	2.08	0.53
1:A:172:ALA:O	1:A:173:GLU:HB2	2.07	0.53
1:A:313:ILE:HD12	1:A:314:ALA:N	2.23	0.53
1:A:371:LYS:HZ1	1:A:400:VAL:CG2	2.21	0.53
1:A:899:ALA:O	1:A:1002:GLU:OE1	2.27	0.53
2:B:202:LYS:O	2:B:325:ARG:NE	2.41	0.53
2:C:291:TYR:HH	2:C:355:GLN:HG3	1.69	0.53
2:C:364:LYS:H	2:C:364:LYS:CD	2.08	0.53
2:C:426:GLU:O	2:C:429:TYR:HB2	2.08	0.53
2:C:451:GLY:O	2:C:452:LEU:HG	2.09	0.53
2:E:418:LEU:HD11	2:F:122:ARG:CG	2.38	0.53
1:D:210:THR:O	1:D:211:LEU:CB	2.54	0.53
1:D:312:TRP:CE3	1:D:316:LYS:CD	2.92	0.53
1:D:405:GLU:O	1:D:406:VAL:C	2.47	0.53
1:D:421:PRO:HG2	1:D:422:VAL:N	2.23	0.53
1:D:457:ARG:CB	1:D:457:ARG:NH1	2.71	0.53
1:D:491:ASP:O	1:D:492:LEU:C	2.45	0.53
1:D:603:MET:O	1:D:604:ALA:C	2.46	0.53
1:D:732:TYR:CB	1:D:755:LYS:NZ	2.71	0.53
1:D:776:ASP:CG	1:D:777:GLY:N	2.61	0.53
1:D:279:ARG:HA	1:D:841:LEU:HD12	1.91	0.53
1:D:932:HIS:ND1	1:D:933:SER:N	2.56	0.53
2:E:379:ALA:N	2:E:380:PRO:CD	2.72	0.53
2:F:199:LEU:HD12	2:F:200:VAL:H	1.73	0.53
2:F:127:PRO:HD3	2:F:210:GLN:HB2	1.82	0.53
2:F:286:GLY:HA3	2:F:303:LEU:O	2.08	0.53
2:F:306:LEU:CG	2:F:306:LEU:O	2.50	0.53
2:F:372:LEU:CD1	2:F:374:LEU:HG	2.38	0.53
2:F:416:GLY:HA2	2:F:419:GLU:HB3	1.91	0.53
1:A:1019:TRP:O	1:A:1020:ILE:CB	2.56	0.53
1:A:347:TRP:HD1	1:A:1030:ARG:HG2	1.73	0.53
1:A:885:THR:HB	1:A:1146:ARG:NH2	2.22	0.53
1:A:312:TRP:HZ2	1:A:346:ASP:C	2.12	0.53
1:A:400:VAL:HG13	1:A:404:HIS:NE2	2.22	0.53
1:A:420:HIS:O	1:A:424:LEU:CG	2.56	0.53
1:A:627:ARG:O	1:A:628:ARG:CB	2.56	0.53
1:A:814:VAL:O	1:A:838:GLY:HA3	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:435:LEU:HD12	1:A:842:PRO:CD	2.39	0.53
2:C:250:GLN:OE1	2:C:257:ARG:NH2	2.29	0.53
2:C:343:ASP:O	2:C:346:MET:HB3	2.09	0.53
1:A:644:GLY:O	1:A:645:VAL:HB	2.07	0.53
1:D:1016:GLU:HG3	1:D:1022:LEU:CD2	2.39	0.53
1:D:116:PRO:O	1:D:118:VAL:HG12	2.08	0.53
1:D:136:ASP:OD1	1:D:140:ARG:NE	2.41	0.53
1:D:278:ILE:HD13	1:D:282:TYR:CZ	2.43	0.53
1:D:298:HIS:NE2	1:D:407:PHE:HB2	2.24	0.53
1:D:434:TYR:C	1:D:1130:CYS:HB3	2.29	0.53
1:D:744:ILE:CG1	1:D:745:PRO:HD3	2.37	0.53
1:D:786:ALA:O	1:D:787:SER:CB	2.55	0.53
1:D:849:THR:HG21	1:D:851:THR:HB	1.89	0.53
1:D:932:HIS:O	1:D:935:THR:CG2	2.46	0.53
2:E:109:ASN:HD22	2:E:378:LEU:HB3	1.73	0.53
2:F:347:LEU:H	2:F:347:LEU:CD1	2.21	0.53
2:F:78:PHE:CD1	2:F:102:LEU:CD2	2.90	0.53
1:A:87:ILE:HG12	1:A:1166:TYR:CE2	2.43	0.53
1:A:316:LYS:HB2	1:A:348:LEU:CD1	2.39	0.53
1:A:408:GLN:HG3	1:A:411:LEU:HD21	1.90	0.53
1:A:441:TRP:HE1	1:A:877:MET:HB3	1.72	0.53
1:A:606:THR:CB	1:A:613:HIS:N	2.72	0.53
1:A:828:HIS:N	1:A:829:PRO:CD	2.72	0.53
1:A:857:PRO:O	1:A:861:THR:OG1	2.20	0.53
1:A:920:THR:OG1	1:A:921:LEU:HD12	2.08	0.53
2:B:348:ALA:O	2:B:351:TYR:N	2.36	0.53
2:B:412:SER:C	2:B:413:VAL:CG2	2.77	0.53
2:C:246:ARG:O	2:C:247:THR:C	2.45	0.53
2:C:274:SER:OG	2:C:275:SER:N	2.41	0.53
2:C:266:PHE:HD1	2:C:378:LEU:CD1	2.21	0.53
2:C:84:GLN:C	2:C:86:LEU:H	2.11	0.53
2:B:133:HIS:NE2	2:C:231:ILE:HG21	2.20	0.53
1:A:528:GLU:O	1:A:529:ASP:OD2	2.27	0.53
1:D:308:GLN:HE22	1:D:1027:LYS:NZ	2.06	0.53
1:D:576:LEU:HD22	2:F:481:SER:HB2	1.89	0.53
1:D:850:ILE:HG12	1:D:851:THR:H	1.73	0.53
1:D:441:TRP:CD1	1:D:878:VAL:O	2.61	0.53
1:D:938:THR:C	1:D:939:VAL:HG23	2.29	0.53
2:E:299:LEU:HD23	2:E:300:ILE:N	2.24	0.53
2:E:400:GLN:CA	2:E:403:PHE:HB3	2.38	0.53
1:D:472:GLN:CD	2:F:460:THR:O	2.47	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:86:LEU:O	2:F:87:SER:O	2.26	0.53
1:A:1019:TRP:O	1:A:1020:ILE:CG2	2.56	0.53
1:A:1124:ALA:CA	1:A:1148:ARG:HH12	2.21	0.53
1:A:247:LEU:HG	1:A:248:GLU:N	2.12	0.53
1:A:411:LEU:C	1:A:411:LEU:HD12	2.28	0.53
1:A:297:MET:CE	1:A:415:LEU:HD21	2.39	0.53
1:A:743:ASP:HB2	1:A:744:ILE:HD13	1.91	0.53
1:A:207:THR:HG21	1:A:812:MET:CB	2.38	0.53
1:A:831:TYR:HB2	1:A:837:TYR:OH	2.09	0.53
1:A:849:THR:HB	1:A:852:ARG:H	1.73	0.53
2:C:376:PRO:HG3	2:C:482:ALA:CB	2.31	0.53
2:C:403:PHE:CD1	2:C:415:PRO:HG3	2.43	0.53
2:E:389:ARG:HE	2:E:390:GLY:H	1.48	0.53
2:C:134:LYS:CB	2:C:180:LYS:HE2	2.38	0.53
1:D:104:VAL:HG12	1:D:105:GLU:N	2.24	0.53
2:C:272:ASN:CA	2:C:292:ASN:ND2	2.72	0.53
1:D:1029:GLN:CD	1:D:1030:ARG:HH21	2.09	0.53
1:D:1143:GLU:O	1:D:1146:ARG:CG	2.54	0.53
1:D:145:LYS:HB3	1:D:1118:TRP:HH2	1.73	0.53
1:D:430:MET:HE1	1:D:1132:SER:HB3	1.91	0.53
1:D:438:ASN:N	1:D:441:TRP:HD1	2.00	0.53
1:D:449:GLN:CA	1:D:452:TYR:HE2	2.13	0.53
1:D:588:GLY:HA3	1:D:806:LYS:HE3	1.90	0.53
2:E:132:HIS:HB2	2:F:233:GLU:OE1	2.08	0.53
2:E:332:VAL:N	2:E:333:PRO:HD3	2.23	0.53
2:E:101:PRO:HG3	2:F:128:VAL:HG21	1.90	0.53
2:F:399:CYS:SG	2:F:417:TYR:CD2	3.00	0.53
2:F:440:THR:O	2:F:457:SER:CB	2.54	0.53
1:A:1019:TRP:CH2	1:A:1163:MET:HG2	2.43	0.53
1:A:1208:ARG:O	2:C:253:ASP:OD2	2.26	0.53
1:A:286:GLY:O	1:A:287:SER:CB	2.56	0.53
1:A:355:SER:O	1:A:407:PHE:HZ	1.92	0.53
1:A:513:LEU:O	1:A:568:GLY:HA3	2.09	0.53
1:A:659:HIS:ND1	1:A:716:PRO:CA	2.71	0.53
1:A:843:GLN:O	1:A:856:GLU:HG2	2.08	0.53
2:B:235:THR:OG1	2:B:343:ASP:N	2.41	0.53
2:B:67:GLU:CG	2:B:69:LEU:HG	2.39	0.53
2:B:451:GLY:O	2:B:468:ILE:HG22	2.09	0.53
1:D:1014:ARG:HG2	1:D:1026:ARG:NE	2.18	0.53
1:D:1019:TRP:CD1	1:D:1020:ILE:N	2.76	0.53
1:D:249:VAL:HG22	1:D:250:PRO:N	2.24	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:267:VAL:C	1:D:269:HIS:NE2	2.62	0.53
1:D:287:SER:C	1:D:288:ARG:NE	2.62	0.53
1:D:357:ALA:C	1:D:359:VAL:H	2.12	0.53
1:D:549:LEU:O	1:D:553:LYS:HG3	2.08	0.53
1:D:656:TYR:C	1:D:659:HIS:CE1	2.81	0.53
1:D:808:ILE:O	1:D:811:GLN:HB3	2.08	0.53
1:D:973:LEU:O	1:D:977:GLU:HG3	2.09	0.53
2:E:351:TYR:C	2:E:353:SER:N	2.59	0.53
2:E:400:GLN:O	2:E:402:LEU:N	2.42	0.53
2:F:131:LEU:CG	2:F:133:HIS:CE1	2.82	0.53
1:A:1100:VAL:HG12	1:A:1101:VAL:N	2.24	0.53
1:A:1115:ALA:O	1:A:1119:LEU:HG	2.08	0.53
1:A:1198:LYS:CG	1:A:1204:THR:CG2	2.80	0.53
1:A:202:CYS:SG	1:A:210:THR:CG2	2.94	0.53
1:A:371:LYS:CG	1:A:372:GLU:HG2	2.39	0.53
1:A:594:LEU:CD2	1:A:594:LEU:N	2.72	0.53
1:A:606:THR:HG21	1:A:613:HIS:H	1.69	0.53
1:A:950:ASN:O	1:A:953:ARG:N	2.41	0.53
2:B:189:ALA:HB1	2:B:208:LEU:HD21	1.91	0.53
2:C:208:LEU:HD12	2:C:208:LEU:N	2.24	0.53
2:C:114:TRP:NE1	2:C:259:ARG:CZ	2.70	0.53
2:C:72:ILE:HG23	2:C:76:ARG:HD2	1.91	0.53
2:C:82:SER:O	2:C:83:LYS:HG3	2.09	0.53
1:A:646:VAL:O	1:A:646:VAL:HG13	2.09	0.53
2:B:217:PRO:HB3	2:B:232:GLY:H	1.74	0.53
1:D:1196:ASP:HB2	1:D:1204:THR:HG22	1.91	0.53
1:D:255:SER:OG	1:D:257:THR:N	2.41	0.53
1:D:369:LEU:CB	1:D:372:GLU:HB2	2.39	0.53
1:D:472:GLN:O	1:D:474:LEU:O	2.27	0.53
1:D:493:GLN:O	1:D:494:GLU:CG	2.53	0.53
1:D:776:ASP:CA	1:D:796:LYS:HZ3	2.22	0.53
1:D:910:MET:HE1	1:D:911:HIS:CE1	2.43	0.53
1:D:967:MET:SD	1:D:975:GLN:N	2.82	0.53
1:D:993:ARG:NE	1:D:994:TRP:HB2	2.24	0.53
2:E:116:THR:O	2:E:118:VAL:N	2.41	0.53
2:E:424:SER:OG	2:E:425:LEU:N	2.42	0.53
2:E:96:HIS:O	2:E:96:HIS:HD2	1.92	0.53
2:F:206:TYR:CE1	2:F:208:LEU:HD23	2.43	0.53
2:F:243:THR:HB	2:F:247:THR:OG1	2.08	0.53
2:F:241:TRP:CE3	2:F:336:LEU:HD12	2.44	0.53
2:F:348:ALA:O	2:F:351:TYR:CB	2.56	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:348:ALA:O	2:F:351:TYR:HB2	2.07	0.53
2:F:403:PHE:CD1	2:F:415:PRO:HG3	2.44	0.53
2:F:442:LEU:HB2	2:F:455:LEU:HD22	1.91	0.53
2:F:484:ASN:C	2:F:485:VAL:CG1	2.77	0.53
1:A:211:LEU:HD11	1:A:221:TYR:HB3	1.89	0.53
1:A:254:SER:O	1:A:255:SER:HB2	2.08	0.53
1:A:451:THR:OG1	1:A:1202:ASN:OD1	2.27	0.53
1:A:853:ARG:O	1:A:854:ALA:HB2	2.09	0.53
1:A:914:THR:O	1:A:915:ALA:HB2	2.09	0.53
2:B:190:LEU:HB3	2:B:315:TYR:OH	2.08	0.53
1:D:666:GLN:HG2	1:D:669:MET:HE2	1.91	0.53
2:F:61:GLY:C	2:F:64:GLU:HG2	2.29	0.53
2:E:311:LEU:O	2:E:315:TYR:N	2.42	0.53
2:C:285:LYS:CB	2:C:285:LYS:NZ	2.71	0.53
1:D:152:GLU:HA	1:D:155:ASN:HD22	1.74	0.53
1:D:1174:PRO:HB2	1:D:1177:VAL:CG2	2.39	0.53
1:D:1195:MET:O	1:D:1196:ASP:CG	2.48	0.53
1:D:1232:GLU:CB	1:D:1234:ARG:HB2	2.39	0.53
1:D:195:LEU:O	1:D:267:VAL:HG12	2.09	0.53
1:D:293:ASP:C	1:D:293:ASP:OD2	2.46	0.53
1:D:305:SER:OG	1:D:1099:TRP:NE1	2.42	0.53
1:D:312:TRP:HE3	1:D:316:LYS:CD	2.22	0.53
1:D:604:ALA:HA	1:D:613:HIS:NE2	2.20	0.53
1:D:618:HIS:HA	1:D:762:VAL:HG22	1.91	0.53
1:D:620:TRP:CZ2	1:D:751:LYS:HD3	2.44	0.53
2:E:106:LEU:C	2:E:109:ASN:OD1	2.47	0.53
2:E:247:THR:HB	2:E:251:TRP:CD1	2.42	0.53
2:E:91:LEU:C	2:E:93:SER:H	2.12	0.53
2:F:243:THR:CG2	2:F:247:THR:OG1	2.57	0.53
2:F:456:ARG:NE	2:F:456:ARG:N	2.56	0.53
1:A:1011:PRO:CD	1:A:1012:VAL:N	2.65	0.53
1:A:1143:GLU:O	1:A:1146:ARG:CB	2.57	0.53
1:A:213:VAL:CG1	1:A:397:ALA:CB	2.84	0.53
1:A:653:GLU:CA	1:A:656:TYR:CZ	2.88	0.53
1:A:658:LYS:CB	1:A:713:PRO:O	2.57	0.53
1:A:659:HIS:CE1	1:A:663:GLN:CD	2.82	0.53
1:A:812:MET:CG	1:A:813:VAL:H	2.21	0.53
2:B:301:GLU:HG2	2:B:303:LEU:CD1	2.39	0.53
2:B:76:ARG:HH21	2:B:431:LYS:HG3	1.73	0.53
2:B:69:LEU:CD1	2:B:73:CYS:SG	2.97	0.53
2:B:74:GLN:HB2	2:B:77:HIS:CD2	2.43	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:131:LEU:HB3	2:C:133:HIS:CG	2.44	0.53
2:C:387:VAL:HG13	2:C:399:CYS:HB2	1.90	0.53
2:C:414:TRP:O	2:C:414:TRP:CD1	2.62	0.53
1:D:567:PRO:O	1:D:568:GLY:C	2.45	0.53
1:D:1138:ARG:HH22	1:D:1191:LYS:HE3	1.73	0.53
1:D:172:ALA:O	1:D:173:GLU:HB2	2.09	0.53
1:D:243:ASP:O	1:D:246:PRO:HD2	2.08	0.53
1:D:271:VAL:N	1:D:274:ASP:OD2	2.42	0.53
1:D:349:ASP:O	1:D:350:ILE:C	2.47	0.53
1:D:472:GLN:HB3	2:F:461:THR:HG23	1.90	0.53
1:D:491:ASP:OD2	1:D:578:PRO:CB	2.56	0.53
1:D:608:ASP:N	1:D:620:TRP:CZ3	2.77	0.53
1:D:649:TYR:C	1:D:652:ILE:HB	2.28	0.53
1:D:711:ALA:C	1:D:713:PRO:CD	2.77	0.53
1:D:827:ARG:HD3	1:D:828:HIS:CA	2.38	0.53
1:D:890:ASP:C	1:D:891:VAL:HG12	2.23	0.53
1:D:920:THR:HG21	1:D:922:GLN:OE1	2.10	0.53
2:F:127:PRO:CD	2:F:209:ALA:C	2.71	0.53
2:F:207:GLY:HA2	2:F:240:VAL:O	2.08	0.53
2:F:360:SER:HA	2:F:363:ARG:HG3	1.91	0.53
2:F:88:ARG:NH1	2:F:88:ARG:CB	2.72	0.53
1:A:107:LEU:O	1:A:111:GLY:N	2.41	0.53
1:A:1131:ILE:HG12	1:A:1132:SER:N	2.24	0.53
1:A:1164:PHE:HB3	1:A:1168:LEU:HD12	1.91	0.53
1:A:129:PRO:O	1:A:130:LEU:HD12	2.09	0.53
1:A:213:VAL:C	1:A:401:TRP:CZ2	2.74	0.53
1:A:354:ASN:O	1:A:355:SER:HB2	2.08	0.53
1:A:389:PHE:O	1:A:390:GLN:C	2.47	0.53
1:A:545:ALA:HB1	2:C:400:GLN:OE1	2.09	0.53
1:A:821:LEU:HD13	1:A:825:VAL:CG1	2.39	0.53
1:A:815:TRP:HE3	1:A:838:GLY:N	2.07	0.53
1:A:887:VAL:HA	1:A:1146:ARG:HH12	1.74	0.53
1:A:949:PHE:O	1:A:953:ARG:N	2.42	0.53
1:A:992:LEU:CD1	1:A:993:ARG:N	2.66	0.53
2:B:105:GLU:HB3	2:B:380:PRO:CB	2.33	0.53
2:B:72:ILE:HD12	2:B:372:LEU:HD21	1.90	0.53
2:B:69:LEU:HA	2:B:72:ILE:CD1	2.39	0.53
2:C:273:PHE:CE1	2:C:291:TYR:CD1	2.92	0.53
2:C:359:ASN:O	2:C:360:SER:HB2	2.07	0.53
2:E:389:ARG:CA	2:E:395:LEU:HG	2.39	0.53
1:A:1183:VAL:C	1:A:1217:ALA:HB1	2.29	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:235:TRP:CD1	1:A:236:THR:OG1	2.62	0.53
1:D:944:GLU:CG	1:D:1071:ARG:NH1	2.71	0.52
1:D:224:CYS:SG	1:D:228:LEU:HD22	2.48	0.52
1:D:275:ARG:HH11	1:D:843:GLN:HE21	1.48	0.52
1:D:279:ARG:NH2	1:D:433:SER:HB3	2.25	0.52
1:D:404:HIS:O	1:D:408:GLN:HB2	2.09	0.52
1:D:795:ASN:HA	1:D:798:ILE:HG12	1.91	0.52
1:D:836:LEU:O	1:D:837:TYR:HD2	1.93	0.52
2:E:115:TRP:CH2	2:E:127:PRO:HB3	2.44	0.52
2:E:123:GLU:C	2:E:125:VAL:H	2.11	0.52
2:E:431:LYS:CG	2:E:432:TYR:N	2.72	0.52
2:E:445:GLU:CD	2:E:446:THR:N	2.62	0.52
2:F:127:PRO:O	2:F:192:HIS:CD2	2.62	0.52
2:F:196:CYS:HB3	2:F:206:TYR:OH	2.09	0.52
2:F:454:HIS:CD2	2:F:468:ILE:CA	2.83	0.52
1:A:1026:ARG:O	1:A:1030:ARG:CA	2.57	0.52
1:A:1133:ILE:O	1:A:1136:GLU:N	2.37	0.52
1:A:1157:ASN:HA	1:A:1160:THR:OG1	2.08	0.52
1:A:1206:MET:O	1:A:1207:GLU:HB2	2.09	0.52
1:A:1206:MET:H	1:A:1209:ARG:HB2	1.72	0.52
1:A:175:TRP:HB3	1:A:184:ALA:HA	1.90	0.52
1:A:308:GLN:HB3	1:A:1090:GLU:CG	2.38	0.52
1:A:353:VAL:HG22	1:A:354:ASN:N	2.23	0.52
1:A:424:LEU:H	1:A:424:LEU:HD23	1.69	0.52
1:A:515:ILE:CG2	1:A:516:GLU:H	2.08	0.52
1:A:659:HIS:CE1	1:A:716:PRO:C	2.82	0.52
1:A:657:ARG:C	1:A:660:CYS:HB3	2.27	0.52
1:A:662:GLU:O	1:A:666:GLN:HB2	2.08	0.52
1:A:770:PHE:O	1:A:773:LYS:HB2	2.09	0.52
1:A:80:SER:OG	1:A:125:LEU:CD1	2.56	0.52
1:A:829:PRO:O	1:A:830:ASP:CB	2.56	0.52
1:A:938:THR:C	1:A:939:VAL:HG23	2.28	0.52
2:B:123:GLU:OE2	2:B:126:PHE:CE2	2.62	0.52
2:C:415:PRO:CB	2:C:417:TYR:CE2	2.90	0.52
1:A:645:VAL:HG12	1:A:646:VAL:N	2.22	0.52
1:A:633:LYS:O	1:A:634:LEU:HD13	2.10	0.52
1:D:206:GLY:CA	1:D:245:ILE:CG2	2.84	0.52
1:D:598:VAL:O	1:D:601:LYS:HD3	2.09	0.52
1:D:601:LYS:NZ	1:D:616:GLU:OE2	2.39	0.52
1:D:630:ASN:C	1:D:630:ASN:HD22	2.13	0.52
1:D:632:ALA:CB	1:D:657:ARG:CD	2.87	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:732:TYR:CE1	1:D:755:LYS:HG3	2.44	0.52
1:D:823:ARG:O	1:D:827:ARG:HB3	2.10	0.52
1:D:851:THR:HG21	1:D:853:ARG:HD3	1.89	0.52
2:E:78:PHE:CE1	2:E:350:LEU:HD12	2.44	0.52
2:E:94:GLY:O	2:E:96:HIS:NE2	2.41	0.52
2:F:129:ASP:OD1	2:F:131:LEU:CD1	2.57	0.52
2:F:196:CYS:CB	2:F:206:TYR:OH	2.56	0.52
2:F:336:LEU:HD23	2:F:337:SER:CA	2.38	0.52
2:F:352:ASP:OD1	2:F:352:ASP:C	2.48	0.52
2:F:372:LEU:HD12	2:F:373:LYS:N	2.25	0.52
2:F:81:GLY:O	2:F:86:LEU:HD21	2.09	0.52
1:A:1072:THR:N	1:A:1073:PRO:CD	2.72	0.52
1:A:308:GLN:HB3	1:A:1090:GLU:CA	2.37	0.52
1:A:1109:LEU:HD11	1:A:1137:VAL:HG21	1.91	0.52
1:A:203:LEU:HD12	1:A:380:GLY:CA	2.39	0.52
1:A:813:VAL:HG11	1:A:815:TRP:CH2	2.44	0.52
1:A:893:SER:OG	1:A:896:LEU:HB2	2.09	0.52
2:B:288:LYS:C	2:B:289:LEU:HD22	2.29	0.52
2:B:379:ALA:O	2:B:380:PRO:C	2.46	0.52
2:C:190:LEU:O	2:C:193:TYR:CB	2.51	0.52
1:D:100:VAL:O	1:D:103:SER:OG	2.23	0.52
1:A:1193:VAL:CG2	1:A:1194:THR:H	2.21	0.52
2:B:180:LYS:HG3	2:B:181:LEU:H	1.73	0.52
1:D:1014:ARG:HG2	1:D:1026:ARG:CG	2.39	0.52
1:D:1135:ASP:O	1:D:1136:GLU:CD	2.48	0.52
1:D:1133:ILE:CB	1:D:1136:GLU:HB2	2.12	0.52
1:D:128:PRO:HA	1:D:130:LEU:CG	2.37	0.52
1:D:134:ASN:O	1:D:137:GLN:N	2.42	0.52
1:D:247:LEU:HA	1:D:815:TRP:HD1	1.71	0.52
1:D:195:LEU:H	1:D:265:LEU:CD1	2.22	0.52
1:D:271:VAL:HG22	1:D:295:MET:HG3	1.92	0.52
1:D:312:TRP:CB	1:D:316:LYS:HD3	2.40	0.52
1:D:370:GLU:HB2	1:D:399:ASP:OD2	2.09	0.52
1:D:735:GLY:HA3	1:D:749:PHE:CG	2.44	0.52
1:D:801:TRP:CE3	1:D:870:VAL:HG12	2.42	0.52
1:D:79:LEU:HB3	1:D:83:LEU:CD1	2.40	0.52
2:E:188:GLY:O	2:E:191:GLU:CG	2.54	0.52
2:E:248:SER:O	2:E:251:TRP:N	2.42	0.52
2:E:331:VAL:HG22	2:E:333:PRO:CD	2.28	0.52
2:E:349:TYR:O	2:E:374:LEU:HD21	2.09	0.52
2:E:373:LYS:HG2	2:E:374:LEU:N	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:114:TRP:HB2	2:F:262:TRP:CH2	2.43	0.52
2:E:104:VAL:HG21	2:F:128:VAL:O	2.09	0.52
2:F:433:ASP:C	2:F:435:MET:N	2.61	0.52
2:F:442:LEU:O	2:F:455:LEU:HD11	2.10	0.52
2:F:80:SER:H	2:F:99:PHE:CA	2.22	0.52
1:A:1004:LEU:O	1:A:1008:LEU:HB3	2.09	0.52
1:A:133:ASP:O	1:A:137:GLN:HB3	2.10	0.52
1:A:179:GLY:C	1:A:181:GLU:N	2.60	0.52
1:A:202:CYS:C	1:A:204:ALA:H	2.09	0.52
1:A:192:GLU:HB2	1:A:217:PRO:HD2	1.91	0.52
1:A:463:LEU:O	1:A:464:MET:C	2.47	0.52
1:A:652:ILE:HA	1:A:655:LEU:HG	1.88	0.52
1:A:787:SER:O	1:A:790:ARG:HB2	2.09	0.52
1:A:850:ILE:HG12	1:A:1099:TRP:HE1	1.72	0.52
2:B:303:LEU:HA	2:B:337:SER:O	2.10	0.52
2:C:82:SER:O	2:C:83:LYS:CG	2.57	0.52
1:A:1037:GLN:O	1:A:1039:LYS:N	2.42	0.52
2:C:258:HIS:CA	2:C:261:GLN:OE1	2.56	0.52
1:D:757:GLY:O	1:D:758:ASN:CB	2.57	0.52
1:D:1014:ARG:HG2	1:D:1026:ARG:HG2	1.90	0.52
1:D:897:TRP:HH2	1:D:1173:LEU:HB3	1.74	0.52
1:D:1221:TYR:O	1:D:1225:GLU:N	2.34	0.52
1:D:192:GLU:HA	1:D:195:LEU:HD11	1.92	0.52
1:D:239:LEU:CA	1:D:242:ALA:HB3	2.39	0.52
1:D:424:LEU:HA	1:D:427:MET:SD	2.49	0.52
1:D:457:ARG:O	1:D:461:LYS:CD	2.57	0.52
1:D:605:LEU:C	1:D:781:ALA:HA	2.28	0.52
1:D:794:ILE:HA	1:D:797:MET:SD	2.49	0.52
1:D:933:SER:C	1:D:935:THR:N	2.62	0.52
1:D:959:GLN:NE2	1:D:983:GLN:HE22	2.08	0.52
1:D:996:ARG:O	1:D:996:ARG:NE	2.34	0.52
2:E:119:VAL:HG23	2:E:120:VAL:N	2.24	0.52
2:E:254:PHE:CA	2:E:257:ARG:HB3	2.24	0.52
2:E:109:ASN:HB2	2:E:378:LEU:HB3	1.91	0.52
2:E:420:THR:HG23	2:E:421:MET:N	2.17	0.52
2:E:90:SER:O	2:E:94:GLY:N	2.42	0.52
1:A:1200:PRO:CB	1:A:1203:PRO:HD2	2.40	0.52
1:A:302:SER:O	1:A:351:SER:HB2	2.09	0.52
1:A:440:ASN:O	1:A:441:TRP:C	2.47	0.52
1:A:455:LEU:HD12	1:A:798:ILE:CG2	2.39	0.52
1:A:553:LYS:CD	2:C:468:ILE:CG2	2.87	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:624:VAL:O	1:A:624:VAL:HG23	2.09	0.52
1:A:616:GLU:CG	1:A:724:GLY:HA3	2.39	0.52
1:A:767:ALA:C	1:A:768:LYS:O	2.43	0.52
1:A:949:PHE:HD2	1:A:949:PHE:N	2.07	0.52
2:C:199:LEU:HD12	2:C:200:VAL:H	1.75	0.52
2:C:304:TRP:N	2:C:304:TRP:CD1	2.77	0.52
2:C:342:LEU:C	2:C:342:LEU:CD2	2.78	0.52
2:C:80:SER:H	2:C:99:PHE:CA	2.23	0.52
2:C:88:ARG:HB2	2:C:88:ARG:HH11	1.74	0.52
1:A:491:ASP:OD1	1:A:491:ASP:N	2.41	0.52
1:D:116:PRO:HA	1:D:921:LEU:CD1	2.35	0.52
1:D:1174:PRO:HB2	1:D:1177:VAL:HG21	1.91	0.52
1:D:120:LEU:H	1:D:120:LEU:CD2	2.21	0.52
1:D:217:PRO:O	1:D:218:SER:CB	2.57	0.52
1:D:223:TRP:HZ3	1:D:250:PRO:HB3	1.74	0.52
1:D:248:GLU:HB3	1:D:249:VAL:CB	2.38	0.52
1:D:427:MET:HB2	1:D:846:THR:HG21	1.91	0.52
1:D:490:TRP:CZ2	1:D:598:VAL:CB	2.84	0.52
1:D:614:TYR:CE2	1:D:751:LYS:NZ	2.70	0.52
1:D:630:ASN:HD22	1:D:631:LEU:N	2.08	0.52
1:D:638:THR:HG23	1:D:641:GLU:OE1	2.09	0.52
1:D:808:ILE:HG21	1:D:874:LEU:CD2	2.40	0.52
2:E:201:ASN:OD1	2:F:422:GLN:OE1	2.26	0.52
2:E:290:TYR:HA	2:E:298:GLU:O	2.10	0.52
2:E:371:VAL:HG13	2:E:372:LEU:H	1.74	0.52
1:D:1206:MET:CE	2:F:257:ARG:NH1	2.73	0.52
2:F:266:PHE:HD1	2:F:378:LEU:CD1	2.22	0.52
1:A:1131:ILE:CG2	1:A:1138:ARG:H	2.20	0.52
1:A:1164:PHE:CD2	1:A:1168:LEU:HD11	2.43	0.52
1:A:86:GLN:C	1:A:87:ILE:HG13	2.30	0.52
2:B:104:VAL:CG2	2:C:128:VAL:HG22	2.39	0.52
2:B:312:LEU:O	2:B:315:TYR:C	2.48	0.52
1:D:664:GLY:O	1:D:667:GLN:HG3	2.10	0.52
2:B:217:PRO:HB2	2:B:231:ILE:CB	2.39	0.52
1:D:151:LEU:O	1:D:154:ALA:N	2.41	0.52
1:D:87:ILE:HD11	1:D:135:LEU:HD13	1.91	0.52
1:D:462:SER:O	1:D:466:LEU:CD2	2.56	0.52
1:D:597:ARG:CG	1:D:617:ARG:HD2	2.40	0.52
1:D:618:HIS:CB	1:D:725:PRO:HB2	2.39	0.52
1:D:655:LEU:O	1:D:656:TYR:C	2.48	0.52
1:D:722:ARG:CG	1:D:722:ARG:HH11	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:452:TYR:CD2	1:D:801:TRP:NE1	2.76	0.52
1:D:823:ARG:CZ	1:D:824:ALA:CB	2.88	0.52
1:D:840:ILE:O	1:D:841:LEU:HD23	2.09	0.52
1:D:851:THR:O	1:D:851:THR:CG2	2.57	0.52
1:D:988:ALA:O	1:D:998:SER:OG	2.21	0.52
2:E:120:VAL:HG22	2:F:407:LEU:HG	1.91	0.52
2:F:376:PRO:CG	2:F:482:ALA:HB1	2.39	0.52
2:F:484:ASN:C	2:F:485:VAL:HG13	2.30	0.52
1:A:1225:GLU:O	1:A:1228:LYS:HG2	2.10	0.52
1:A:80:SER:OG	1:A:125:LEU:CD2	2.57	0.52
1:A:214:ALA:HB2	1:A:219:ALA:HB3	1.90	0.52
1:A:276:ALA:C	1:A:279:ARG:N	2.62	0.52
1:A:392:LEU:HG	1:A:395:TYR:HB2	1.90	0.52
1:A:414:PHE:O	1:A:417:ARG:HB3	2.09	0.52
1:A:437:VAL:N	1:A:878:VAL:O	2.42	0.52
1:A:783:PRO:HG2	2:C:363:ARG:NE	2.25	0.52
1:A:83:LEU:HD13	1:A:89:GLY:HA3	1.92	0.52
1:A:995:TYR:O	1:A:996:ARG:HG3	2.10	0.52
2:B:114:TRP:N	2:B:262:TRP:CZ2	2.77	0.52
2:B:311:LEU:HB2	2:B:315:TYR:HD2	1.75	0.52
2:B:342:LEU:O	2:B:344:ARG:N	2.43	0.52
2:B:76:ARG:NH2	2:B:431:LYS:HB2	2.24	0.52
2:C:193:TYR:CD1	2:C:242:PHE:CD2	2.97	0.52
2:C:125:VAL:HG13	2:C:206:TYR:HD1	1.74	0.52
2:C:307:GLY:N	2:C:335:VAL:CG1	2.73	0.52
2:C:398:VAL:O	2:C:402:LEU:N	2.42	0.52
1:D:584:ALA:O	1:D:585:TRP:HE3	1.92	0.52
1:D:1171:ASN:N	1:D:1171:ASN:OD1	2.41	0.52
1:D:1207:GLU:CA	1:D:1210:TYR:HB3	2.23	0.52
1:D:194:ALA:HA	1:D:265:LEU:HB3	1.90	0.52
1:D:281:GLN:HE21	1:D:287:SER:HB2	1.75	0.52
1:D:598:VAL:HG12	1:D:602:LEU:CD1	2.40	0.52
1:D:601:LYS:HG2	1:D:721:ALA:CA	2.40	0.52
1:D:605:LEU:O	1:D:781:ALA:CA	2.50	0.52
1:D:633:LYS:HB3	1:D:638:THR:CG2	2.40	0.52
1:D:775:GLU:CG	1:D:780:GLN:CG	2.86	0.52
1:D:279:ARG:HH11	1:D:841:LEU:HB3	1.75	0.52
1:D:85:GLU:OE2	1:D:130:LEU:HD22	2.08	0.52
2:E:354:PHE:CG	2:E:355:GLN:N	2.78	0.52
2:E:425:LEU:CA	2:E:428:LEU:HD11	2.39	0.52
2:E:456:ARG:CG	2:E:457:SER:H	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:256:LEU:O	2:F:257:ARG:O	2.27	0.52
1:D:1208:ARG:NH1	2:F:285:LYS:HE2	2.24	0.52
2:F:418:LEU:HD11	2:F:421:MET:SD	2.49	0.52
2:F:446:THR:HA	2:F:450:ASN:H	1.73	0.52
1:A:1131:ILE:CG1	1:A:1137:VAL:HA	2.40	0.52
1:A:157:LEU:HD21	1:A:216:SER:CB	2.40	0.52
1:A:188:ALA:O	1:A:189:ILE:CB	2.56	0.52
1:A:404:HIS:O	1:A:408:GLN:HB2	2.10	0.52
1:A:437:VAL:CG1	1:A:438:ASN:N	2.72	0.52
1:A:734:HIS:HA	1:A:752:LEU:CD1	2.39	0.52
1:A:749:PHE:N	1:A:749:PHE:CD1	2.77	0.52
1:A:77:GLN:OE1	1:A:1174:PRO:HB2	2.10	0.52
1:A:795:ASN:C	1:A:795:ASN:HD22	2.13	0.52
2:B:436:SER:OG	2:B:458:ARG:NH1	2.43	0.52
2:C:246:ARG:HD3	2:C:246:ARG:N	2.24	0.52
2:C:351:TYR:CD1	2:C:355:GLN:HA	2.43	0.52
2:C:294:PRO:HG3	2:C:355:GLN:HE21	1.74	0.52
2:C:425:LEU:HD11	2:C:429:TYR:CZ	2.45	0.52
2:F:369:ARG:HD2	2:F:369:ARG:H	1.75	0.52
2:C:107:ARG:HH11	2:C:107:ARG:HG2	1.75	0.52
1:D:1163:MET:HE3	1:D:1167:LYS:HG2	1.92	0.52
1:D:1206:MET:HE3	1:D:1210:TYR:HA	1.89	0.52
1:D:286:GLY:O	1:D:287:SER:OG	2.27	0.52
1:D:299:MET:O	1:D:303:GLY:N	2.36	0.52
1:D:454:GLU:O	1:D:458:GLU:HG3	2.09	0.52
1:D:491:ASP:N	1:D:491:ASP:OD1	2.43	0.52
1:D:597:ARG:CB	1:D:616:GLU:OE1	2.57	0.52
1:D:275:ARG:CG	1:D:843:GLN:H	2.23	0.52
2:F:416:GLY:C	2:F:419:GLU:HB3	2.29	0.52
1:A:1020:ILE:O	1:A:1021:SER:HB2	2.08	0.52
1:A:1066:THR:CB	1:A:1071:ARG:HA	2.40	0.52
1:A:308:GLN:CB	1:A:1090:GLU:HG3	2.39	0.52
1:A:253:ALA:O	1:A:256:PRO:O	2.27	0.52
1:A:732:TYR:CE1	1:A:755:LYS:CB	2.87	0.52
2:B:196:CYS:O	2:B:199:LEU:CB	2.57	0.52
2:B:261:GLN:HA	2:B:264:ARG:HG3	1.90	0.52
2:B:311:LEU:HG	2:B:312:LEU:HD13	1.91	0.52
2:B:313:HIS:C	2:B:313:HIS:CD2	2.82	0.52
2:B:383:VAL:O	2:B:413:VAL:HA	2.10	0.52
2:C:130:ALA:CB	2:C:182:ARG:HH12	2.21	0.52
2:C:196:CYS:HB2	2:C:206:TYR:OH	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:252:LEU:HD23	2:C:252:LEU:C	2.30	0.52
1:A:784:GLY:HA3	2:C:362:THR:O	2.10	0.52
2:C:466:MET:SD	2:C:471:LEU:CD2	2.89	0.52
2:C:82:SER:O	2:C:83:LYS:HB2	2.10	0.52
1:D:502:LYS:HD2	1:D:506:GLU:CB	2.39	0.52
1:A:973:LEU:N	1:A:973:LEU:HD23	2.25	0.52
2:B:318:ASN:O	2:B:321:LYS:HG2	2.10	0.52
1:D:1054:GLU:O	1:D:1058:PHE:N	2.42	0.52
1:D:1207:GLU:C	1:D:1210:TYR:HD2	2.13	0.52
1:D:220:TRP:NE1	1:D:221:TYR:CD2	2.78	0.52
1:D:312:TRP:HB3	1:D:316:LYS:HD3	1.92	0.52
1:D:374:ARG:O	1:D:378:VAL:N	2.43	0.52
1:D:656:TYR:HA	1:D:659:HIS:CE1	2.44	0.52
1:D:939:VAL:O	1:D:940:GLY:C	2.47	0.52
2:E:263:TRP:C	2:E:265:LYS:H	2.13	0.52
2:E:328:ARG:H	2:E:328:ARG:HD2	1.70	0.52
2:E:372:LEU:CG	2:E:373:LYS:N	2.52	0.52
2:E:450:ASN:HB3	2:E:452:LEU:HD21	1.90	0.52
2:E:465:MET:SD	2:E:465:MET:N	2.83	0.52
2:F:118:VAL:HG12	2:F:119:VAL:N	2.25	0.52
2:F:201:ASN:C	2:F:203:ARG:N	2.58	0.52
2:F:342:LEU:HB2	2:F:344:ARG:HG3	1.87	0.52
2:F:404:ASN:O	2:F:405:GLU:C	2.49	0.52
2:F:406:LEU:HA	2:F:409:ASN:HD22	1.68	0.52
1:D:546:ARG:CZ	2:F:408:GLU:OE1	2.58	0.52
2:F:414:TRP:O	2:F:414:TRP:CD1	2.63	0.52
1:A:1005:VAL:C	1:A:1006:ARG:HH11	2.13	0.52
1:A:1095:SER:HA	1:A:1099:TRP:CZ3	2.45	0.52
1:A:293:ASP:HB2	1:A:296:SER:CB	2.40	0.52
1:A:365:TYR:C	1:A:367:GLY:H	2.13	0.52
1:A:509:THR:HG21	1:A:513:LEU:CD2	2.39	0.52
1:A:553:LYS:CD	2:C:468:ILE:HG21	2.39	0.52
1:A:80:SER:HG	1:A:81:ARG:H	1.57	0.52
2:B:125:VAL:HG13	2:B:207:GLY:O	2.09	0.52
2:B:249:ASN:O	2:B:252:LEU:CB	2.58	0.52
2:C:322:LEU:HD22	2:C:323:HIS:C	2.30	0.52
2:C:323:HIS:C	2:C:331:VAL:O	2.48	0.52
2:C:327:GLY:O	2:C:328:ARG:CB	2.58	0.52
2:C:395:LEU:O	2:C:397:GLN:O	2.27	0.52
2:E:393:LEU:HA	2:E:396:ARG:CG	2.40	0.52
1:D:500:ALA:CB	1:D:521:PRO:HD3	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:146:ARG:HB2	2:C:228:VAL:HG11	1.91	0.52
1:D:1129:PHE:CD2	1:D:1139:TYR:CD2	2.98	0.52
1:D:1138:ARG:NH2	1:D:1191:LYS:HE2	2.20	0.52
1:D:157:LEU:HG	1:D:194:ALA:CB	2.40	0.52
1:D:655:LEU:O	1:D:659:HIS:NE2	2.44	0.52
1:D:83:LEU:O	1:D:87:ILE:N	2.34	0.52
1:D:854:ALA:CB	1:D:860:LEU:HD13	2.36	0.52
1:D:874:LEU:O	1:D:875:LYS:C	2.46	0.52
2:F:342:LEU:HD23	2:F:342:LEU:O	2.09	0.52
2:F:294:PRO:HG3	2:F:355:GLN:HE21	1.74	0.52
2:F:454:HIS:CD2	2:F:468:ILE:HD12	2.45	0.52
1:A:1175:GLN:CG	1:A:1176:SER:N	2.71	0.52
1:A:213:VAL:HG11	1:A:397:ALA:CB	2.40	0.52
1:A:192:GLU:HA	1:A:217:PRO:HG2	1.92	0.52
1:A:545:ALA:O	1:A:548:CYS:SG	2.65	0.52
1:A:576:LEU:HD12	1:A:577:CYS:CB	2.37	0.52
1:A:670:PRO:HA	1:A:673:ALA:HB3	1.91	0.52
1:A:618:HIS:CD2	1:A:726:LYS:O	2.63	0.52
1:A:727:ASP:O	1:A:728:THR:CB	2.56	0.52
1:A:964:ARG:O	1:A:968:GLN:HG2	2.10	0.52
2:B:257:ARG:O	2:B:258:HIS:C	2.48	0.52
2:C:129:ASP:O	2:C:130:ALA:CB	2.58	0.52
2:C:455:LEU:CB	2:C:456:ARG:NE	2.72	0.52
2:E:280:ASP:OD1	2:E:280:ASP:N	2.42	0.52
2:C:217:PRO:CB	2:C:219:PHE:CE1	2.90	0.52
1:D:866:ARG:CG	1:D:1069:ILE:CG2	2.78	0.51
1:D:1092:PHE:O	1:D:1096:ARG:NH1	2.43	0.51
1:D:1138:ARG:CG	1:D:1138:ARG:NH1	2.63	0.51
1:D:130:LEU:HD23	1:D:130:LEU:H	1.75	0.51
1:D:280:GLU:O	1:D:284:ILE:HD11	2.10	0.51
1:D:410:GLN:C	1:D:412:PRO:HD2	2.30	0.51
1:D:414:PHE:O	1:D:417:ARG:CB	2.56	0.51
1:D:601:LYS:HZ2	1:D:721:ALA:CA	2.24	0.51
1:D:787:SER:O	1:D:789:PRO:HD2	2.09	0.51
1:D:865:ALA:HB2	1:D:1197:CYS:SG	2.50	0.51
1:D:867:PRO:O	1:D:868:ASP:CG	2.49	0.51
1:D:994:TRP:O	1:D:994:TRP:HE3	1.93	0.51
2:E:118:VAL:HG13	2:E:119:VAL:N	2.25	0.51
2:E:205:PRO:HB3	2:E:243:THR:CA	2.36	0.51
2:E:130:ALA:HB3	2:E:210:GLN:OE1	2.10	0.51
2:E:443:VAL:HA	2:E:453:ILE:CG2	2.40	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:200:VAL:HG13	2:F:204:LEU:HD22	1.92	0.51
2:F:243:THR:OG1	2:F:334:CYS:HB3	2.08	0.51
2:F:310:GLU:OE1	2:F:310:GLU:N	2.43	0.51
2:F:306:LEU:H	2:F:335:VAL:HG21	1.75	0.51
2:F:415:PRO:CA	2:F:417:TYR:HE2	2.23	0.51
2:F:67:GLU:O	2:F:71:GLU:CB	2.57	0.51
2:F:76:ARG:CG	2:F:76:ARG:HH11	2.23	0.51
1:A:1068:ASP:N	1:A:1071:ARG:CD	2.73	0.51
1:A:192:GLU:C	1:A:193:ARG:HG2	2.30	0.51
1:A:292:LEU:HD11	1:A:428:LEU:HG	1.92	0.51
1:A:448:ALA:C	1:A:870:VAL:HG11	2.31	0.51
1:A:490:TRP:CZ2	1:A:596:MET:CB	2.91	0.51
1:A:726:LYS:O	1:A:727:ASP:O	2.27	0.51
2:B:114:TRP:N	2:B:262:TRP:CH2	2.78	0.51
2:B:309:HIS:HB2	2:B:312:LEU:HB2	1.93	0.51
2:B:373:LYS:HE2	2:B:459:ASP:CA	2.37	0.51
2:B:392:THR:HG22	2:B:393:LEU:HG	1.91	0.51
2:E:388:GLY:O	2:E:395:LEU:HD12	2.10	0.51
1:D:1029:GLN:HG3	1:D:1030:ARG:NE	2.14	0.51
1:D:214:ALA:O	1:D:401:TRP:CH2	2.63	0.51
1:D:229:VAL:HG23	1:D:230:GLU:N	2.25	0.51
1:D:246:PRO:O	1:D:247:LEU:HD12	2.10	0.51
1:D:251:THR:HG23	1:D:280:GLU:OE1	2.10	0.51
1:D:549:LEU:CD2	1:D:552:LEU:HD12	2.40	0.51
1:D:612:LEU:HD11	1:D:622:TYR:CG	2.45	0.51
2:E:419:GLU:OE2	2:F:201:ASN:ND2	2.43	0.51
1:A:1029:GLN:HA	1:A:1032:THR:HB	1.92	0.51
1:A:601:LYS:HZ1	1:A:721:ALA:CB	2.22	0.51
1:A:786:ALA:CB	1:A:790:ARG:CZ	2.87	0.51
1:A:865:ALA:HB2	1:A:872:SER:O	2.10	0.51
1:A:610:PHE:CZ	1:A:972:ARG:NH2	2.78	0.51
2:B:235:THR:N	2:B:343:ASP:HB2	2.25	0.51
2:B:323:HIS:HB3	2:B:332:VAL:HB	1.92	0.51
2:C:344:ARG:HA	2:C:347:LEU:CD1	2.40	0.51
2:C:389:ARG:CD	2:C:390:GLY:N	2.71	0.51
2:C:459:ASP:O	2:C:460:THR:HG23	2.11	0.51
2:C:454:HIS:HD2	2:C:468:ILE:HG13	1.73	0.51
2:C:471:LEU:C	2:C:474:PHE:H	2.13	0.51
2:C:472:LYS:O	2:C:475:LEU:HG	2.10	0.51
1:A:640:LEU:H	1:A:640:LEU:CD2	1.92	0.51
2:E:270:PRO:O	2:E:273:PHE:CD2	2.60	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:564:GLN:HA	1:A:564:GLN:HE21	1.73	0.51
2:E:282:GLU:HB3	2:E:309:HIS:ND1	2.25	0.51
1:D:1020:ILE:C	1:D:1020:ILE:HD13	2.27	0.51
1:D:1193:VAL:HG21	1:D:1215:GLY:H	1.74	0.51
1:D:148:LEU:HB2	1:D:149:PRO:HD3	1.92	0.51
1:D:275:ARG:CB	1:D:843:GLN:OE1	2.59	0.51
1:D:279:ARG:N	1:D:279:ARG:HD3	2.25	0.51
1:D:576:LEU:CD1	1:D:577:CYS:N	2.69	0.51
1:D:868:ASP:O	1:D:869:ARG:CB	2.58	0.51
1:D:888:GLY:HA3	1:D:1138:ARG:CD	2.18	0.51
1:D:931:LEU:HA	1:D:934:LYS:HE3	1.92	0.51
1:D:992:LEU:HD13	1:D:993:ARG:H	1.74	0.51
2:E:116:THR:C	2:E:118:VAL:H	2.12	0.51
2:E:429:TYR:O	2:E:433:ASP:HB2	2.10	0.51
2:F:115:TRP:CZ3	2:F:210:GLN:HA	2.46	0.51
2:F:445:GLU:O	2:F:447:THR:N	2.43	0.51
1:A:1175:GLN:CG	1:A:1176:SER:H	2.21	0.51
1:A:244:LEU:O	1:A:247:LEU:HD23	2.10	0.51
1:A:606:THR:CG2	1:A:614:TYR:N	2.66	0.51
1:A:870:VAL:C	1:A:872:SER:N	2.62	0.51
1:A:887:VAL:CA	1:A:1146:ARG:NH1	2.74	0.51
1:A:896:LEU:HD13	1:A:954:ILE:HD12	1.91	0.51
1:A:930:ASP:C	1:A:930:ASP:OD2	2.48	0.51
1:A:960:PRO:O	1:A:963:GLU:HG3	2.11	0.51
2:B:236:GLU:HG2	2:B:341:ASP:CB	2.37	0.51
2:B:259:ARG:CD	2:B:262:TRP:CG	2.93	0.51
2:C:266:PHE:HB2	2:C:349:TYR:OH	2.11	0.51
2:C:406:LEU:N	2:C:406:LEU:CD1	2.73	0.51
2:E:418:LEU:HD11	2:F:122:ARG:HG3	1.91	0.51
1:D:1121:GLU:O	1:D:1123:PHE:N	2.40	0.51
1:D:1175:GLN:O	1:D:1177:VAL:HG23	2.10	0.51
1:D:175:TRP:HB2	1:D:185:VAL:CG2	2.40	0.51
1:D:179:GLY:O	1:D:181:GLU:N	2.44	0.51
1:D:542:ASP:OD2	2:F:404:ASN:ND2	2.44	0.51
1:D:826:ILE:HB	1:D:831:TYR:CE2	2.42	0.51
1:D:937:THR:O	1:D:940:GLY:N	2.44	0.51
2:E:113:GLU:O	2:E:116:THR:OG1	2.28	0.51
2:E:113:GLU:HG3	2:E:378:LEU:HD11	1.92	0.51
2:E:407:LEU:O	2:E:409:ASN:N	2.43	0.51
2:E:72:ILE:HD12	2:E:72:ILE:N	2.25	0.51
2:F:115:TRP:O	2:F:119:VAL:HG23	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:180:LYS:CE	2:F:180:LYS:C	2.64	0.51
2:F:134:LYS:HB2	2:F:180:LYS:HE2	1.92	0.51
2:F:372:LEU:HD21	2:F:436:SER:HB2	1.92	0.51
2:F:381:ILE:O	2:F:437:ILE:HA	2.10	0.51
2:F:381:ILE:HB	2:F:437:ILE:HG22	1.93	0.51
1:A:1033:ALA:O	1:A:1034:ARG:CB	2.58	0.51
1:A:1041:TRP:NE1	1:A:1043:VAL:CB	2.68	0.51
1:A:1069:ILE:CG2	1:A:1070:PRO:CD	2.87	0.51
1:A:169:TRP:HD1	1:A:170:ALA:H	1.57	0.51
1:A:441:TRP:CE3	1:A:445:LEU:HD12	2.44	0.51
1:A:73:PRO:O	1:A:74:LEU:HD22	2.10	0.51
1:A:868:ASP:OD2	1:A:1199:THR:OG1	2.26	0.51
2:B:249:ASN:O	2:B:252:LEU:HB2	2.09	0.51
2:B:270:PRO:C	2:B:272:ASN:H	2.10	0.51
2:C:300:ILE:CG2	2:C:301:GLU:H	2.15	0.51
2:C:383:VAL:CG2	2:C:384:ALA:N	2.74	0.51
2:C:471:LEU:O	2:C:474:PHE:N	2.34	0.51
1:A:503:VAL:CG2	1:A:504:LYS:N	2.72	0.51
1:D:666:GLN:HG2	1:D:669:MET:HE3	1.93	0.51
1:D:163:PRO:O	1:D:165:LYS:HD2	2.11	0.51
1:D:1010:LEU:C	1:D:1012:VAL:H	2.10	0.51
1:D:1065:ALA:CA	1:D:1073:PRO:HG3	2.39	0.51
1:D:249:VAL:HG21	1:D:251:THR:HG22	1.91	0.51
1:D:275:ARG:HG3	1:D:843:GLN:N	2.26	0.51
1:D:275:ARG:NH1	1:D:843:GLN:HE21	2.06	0.51
1:D:316:LYS:HZ2	1:D:348:LEU:HD23	1.75	0.51
1:D:409:GLN:O	1:D:412:PRO:CD	2.59	0.51
1:D:80:SER:O	1:D:83:LEU:HD22	2.11	0.51
1:D:887:VAL:CG1	1:D:1183:VAL:CG1	2.89	0.51
1:D:888:GLY:CA	1:D:1138:ARG:HA	2.39	0.51
1:D:898:ILE:HG13	1:D:899:ALA:N	2.25	0.51
2:E:185:LEU:CD1	2:E:237:ALA:O	2.58	0.51
2:E:373:LYS:HG2	2:E:374:LEU:H	1.75	0.51
2:E:86:LEU:CA	2:E:90:SER:HB3	2.36	0.51
2:F:301:GLU:HG2	2:F:339:ASN:CB	2.40	0.51
1:A:1031:GLU:O	1:A:1032:THR:CB	2.56	0.51
1:A:1041:TRP:O	1:A:1042:GLU:CB	2.55	0.51
1:A:1113:LEU:N	1:A:1113:LEU:CD2	2.72	0.51
1:A:1207:GLU:HA	1:A:1210:TYR:CD1	2.39	0.51
1:A:1208:ARG:H	1:A:1210:TYR:H	1.58	0.51
1:A:305:SER:HB2	1:A:308:GLN:HG2	1.89	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:376:LEU:O	1:A:378:VAL:N	2.30	0.51
1:A:556:THR:HG21	2:C:452:LEU:HB2	1.93	0.51
1:A:592:LEU:HA	1:A:595:GLN:CG	2.40	0.51
1:A:752:LEU:CB	1:A:753:PRO:HA	2.39	0.51
1:A:81:ARG:HG3	1:A:81:ARG:NH1	2.25	0.51
1:A:920:THR:HG23	1:A:922:GLN:HB3	1.93	0.51
1:A:938:THR:O	1:A:939:VAL:HB	2.10	0.51
2:B:105:GLU:O	2:B:109:ASN:HB3	2.10	0.51
2:B:114:TRP:HZ2	2:B:259:ARG:HH21	1.59	0.51
2:B:123:GLU:HB2	2:C:403:PHE:CZ	2.40	0.51
2:B:125:VAL:HG12	2:B:126:PHE:H	1.75	0.51
2:B:320:SER:HA	2:B:323:HIS:HD2	1.76	0.51
2:B:69:LEU:O	2:B:72:ILE:CG1	2.56	0.51
2:C:235:THR:C	2:C:341:ASP:CG	2.69	0.51
1:A:93:GLU:O	1:A:95:PRO:HD3	2.11	0.51
1:D:941:ILE:HG12	1:D:942:SER:H	1.75	0.51
1:D:1027:LYS:NZ	1:D:1091:GLU:CG	2.53	0.51
1:D:1112:MET:HB3	1:D:1113:LEU:HD23	1.91	0.51
1:D:242:ALA:HA	1:D:245:ILE:H	1.75	0.51
1:D:278:ILE:HG22	1:D:279:ARG:N	2.26	0.51
1:D:287:SER:O	1:D:288:ARG:NH2	2.43	0.51
1:D:423:THR:O	1:D:427:MET:HG2	2.11	0.51
1:D:656:TYR:HE1	1:D:743:ASP:CB	2.23	0.51
1:D:761:ASN:O	1:D:762:VAL:C	2.48	0.51
1:D:792:LEU:O	1:D:795:ASN:ND2	2.43	0.51
1:D:83:LEU:N	1:D:83:LEU:CD1	2.72	0.51
1:D:279:ARG:HG3	1:D:841:LEU:C	2.30	0.51
1:D:993:ARG:O	1:D:996:ARG:HD2	2.11	0.51
2:E:205:PRO:CB	2:E:242:PHE:O	2.59	0.51
2:E:293:PHE:HD1	2:E:295:TRP:NE1	2.08	0.51
2:E:72:ILE:HD11	2:E:370:LYS:NZ	2.25	0.51
2:E:89:ASP:O	2:E:93:SER:N	2.43	0.51
2:F:201:ASN:O	2:F:202:LYS:HB3	2.11	0.51
2:F:364:LYS:HD3	2:F:364:LYS:H	1.76	0.51
1:A:1068:ASP:C	1:A:1070:PRO:CD	2.78	0.51
1:A:1224:ILE:O	1:A:1224:ILE:CD1	2.53	0.51
1:A:374:ARG:NH1	1:A:374:ARG:CG	2.69	0.51
1:A:821:LEU:O	1:A:821:LEU:CG	2.58	0.51
1:A:275:ARG:NH2	1:A:844:VAL:O	2.36	0.51
2:B:255:TRP:O	2:B:256:LEU:C	2.49	0.51
2:B:263:TRP:O	2:B:264:ARG:C	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:456:ARG:NH2	2:B:457:SER:O	2.44	0.51
2:C:193:TYR:CD1	2:C:193:TYR:C	2.83	0.51
2:B:77:HIS:ND1	2:C:195:ASN:OD1	2.43	0.51
2:C:239:LEU:HD21	2:C:255:TRP:CZ2	2.45	0.51
2:C:241:TRP:H	2:C:336:LEU:CB	2.22	0.51
2:C:351:TYR:HD1	2:C:355:GLN:HG2	1.75	0.51
2:C:399:CYS:O	2:C:400:GLN:C	2.47	0.51
2:C:451:GLY:O	2:C:452:LEU:CB	2.58	0.51
2:C:82:SER:CA	2:C:86:LEU:CD2	2.77	0.51
1:D:235:TRP:CD1	1:D:235:TRP:C	2.84	0.51
2:E:270:PRO:CA	2:E:273:PHE:HD2	2.21	0.51
2:B:410:GLY:C	2:B:411:ILE:HD12	2.31	0.51
1:D:1131:ILE:O	1:D:1136:GLU:O	2.29	0.51
1:D:886:LEU:CA	1:D:1146:ARG:HH12	2.23	0.51
1:D:1162:CYS:O	1:D:1165:ALA:HB3	2.10	0.51
1:D:902:LEU:HD13	1:D:1168:LEU:HD13	1.93	0.51
1:D:1147:TYR:HB3	1:D:1227:THR:CG2	2.41	0.51
1:D:610:PHE:CD1	1:D:972:ARG:NH2	2.77	0.51
1:D:735:GLY:HA3	1:D:749:PHE:CA	2.40	0.51
2:E:447:THR:HG23	2:E:452:LEU:C	2.31	0.51
2:F:193:TYR:CD1	2:F:242:PHE:CD2	2.98	0.51
2:F:290:TYR:HD1	2:F:299:LEU:HA	1.76	0.51
2:F:354:PHE:HE1	2:F:370:LYS:CB	2.23	0.51
2:F:464:GLU:OE2	2:F:466:MET:CG	2.58	0.51
2:F:78:PHE:CE2	2:F:350:LEU:HD23	2.46	0.51
1:A:389:PHE:HB2	1:A:393:MET:HG2	1.92	0.51
1:A:424:LEU:H	1:A:424:LEU:CD2	2.23	0.51
1:A:616:GLU:HG2	1:A:724:GLY:CA	2.39	0.51
1:A:878:VAL:O	1:A:879:GLN:NE2	2.43	0.51
1:A:829:PRO:HG3	1:A:883:GLY:HA2	1.92	0.51
2:B:126:PHE:HB3	2:B:127:PRO:HD2	1.92	0.51
2:B:324:GLY:H	2:B:333:PRO:HD3	1.76	0.51
2:B:86:LEU:HD23	2:B:97:PRO:HD2	1.92	0.51
2:C:118:VAL:CG1	2:C:119:VAL:N	2.74	0.51
2:C:331:VAL:HG13	2:C:332:VAL:N	2.24	0.51
2:C:404:ASN:O	2:C:405:GLU:C	2.49	0.51
2:C:371:VAL:HG23	2:C:433:ASP:O	2.11	0.51
2:C:76:ARG:HH11	2:C:76:ARG:CG	2.24	0.51
2:C:180:LYS:C	2:C:180:LYS:CE	2.65	0.51
1:A:564:GLN:C	1:A:567:PRO:HD2	2.30	0.51
1:A:973:LEU:O	1:A:977:GLU:HG3	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1008:LEU:HA	1:D:1011:PRO:CG	2.39	0.51
1:D:1020:ILE:HD13	1:D:1021:SER:H	1.61	0.51
1:D:1122:GLU:C	1:D:1123:PHE:CD1	2.84	0.51
1:D:251:THR:O	1:D:254:SER:CA	2.58	0.51
1:D:272:SER:HA	1:D:843:GLN:CG	2.41	0.51
1:D:374:ARG:HE	1:D:392:LEU:HG	1.76	0.51
1:D:392:LEU:O	1:D:395:TYR:CB	2.51	0.51
1:D:484:TRP:NE1	1:D:488:LEU:HG	2.26	0.51
1:D:715:GLN:HG2	1:D:717:LEU:CD1	2.40	0.51
1:D:620:TRP:C	1:D:748:TRP:HB2	2.29	0.51
1:D:775:GLU:O	1:D:776:ASP:CB	2.44	0.51
1:D:875:LYS:HG3	1:D:876:ALA:N	2.25	0.51
1:D:916:PHE:CE1	1:D:920:THR:O	2.63	0.51
1:D:964:ARG:O	1:D:968:GLN:CG	2.59	0.51
1:D:990:LYS:O	1:D:996:ARG:NH2	2.43	0.51
2:E:121:PHE:HB2	2:E:122:ARG:HD2	1.93	0.51
2:E:128:VAL:HG22	2:E:129:ASP:N	2.24	0.51
2:E:378:LEU:CD1	2:E:378:LEU:H	2.24	0.51
1:D:1209:ARG:NH2	2:F:256:LEU:CD2	2.74	0.51
2:F:239:LEU:N	2:F:338:VAL:HG21	2.25	0.51
2:F:235:THR:O	2:F:341:ASP:OD1	2.28	0.51
2:F:372:LEU:HD21	2:F:436:SER:N	2.25	0.51
1:D:546:ARG:HH21	2:F:404:ASN:HD22	1.58	0.51
2:F:454:HIS:NE2	2:F:472:LYS:HE3	2.25	0.51
2:F:78:PHE:CD1	2:F:103:GLY:CA	2.91	0.51
1:A:1010:LEU:O	1:A:1096:ARG:CZ	2.58	0.51
1:A:1040:LYS:HG2	1:A:1044:VAL:HG11	1.92	0.51
1:A:1109:LEU:HD11	1:A:1137:VAL:CG2	2.40	0.51
1:A:868:ASP:OD1	1:A:1199:THR:HG21	2.11	0.51
1:A:1187:ARG:HB2	1:A:1212:ILE:CG1	2.40	0.51
1:A:259:ARG:HB3	1:A:261:TRP:HB2	1.93	0.51
1:A:283:LEU:O	1:A:283:LEU:HD13	2.10	0.51
1:A:386:ARG:CG	1:A:387:GLU:N	2.74	0.51
1:A:460:LYS:C	1:A:464:MET:HG3	2.31	0.51
1:A:472:GLN:HG3	2:C:460:THR:C	2.29	0.51
1:A:891:VAL:CA	1:A:1180:PHE:CE1	2.91	0.51
2:B:132:HIS:ND1	2:B:182:ARG:NH2	2.59	0.51
2:B:283:GLY:O	2:B:284:ARG:HB2	2.08	0.51
2:C:443:VAL:HG13	2:C:447:THR:CG2	2.41	0.51
2:C:458:ARG:NH1	2:C:458:ARG:HG2	2.13	0.51
2:C:466:MET:CB	2:C:471:LEU:CD2	2.77	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:941:ILE:CG2	1:A:942:SER:N	2.74	0.51
2:E:218:VAL:H	2:E:231:ILE:CA	2.20	0.51
1:D:224:CYS:O	1:D:228:LEU:HD23	2.11	0.51
1:D:313:ILE:CG1	1:D:314:ALA:N	2.73	0.51
1:D:424:LEU:O	1:D:427:MET:HG2	2.11	0.51
1:D:432:VAL:O	1:D:433:SER:C	2.49	0.51
1:D:545:ALA:HA	1:D:548:CYS:SG	2.51	0.51
1:D:917:GLY:O	1:D:920:THR:HA	2.11	0.51
1:D:989:THR:HA	1:D:998:SER:HG	1.68	0.51
2:E:193:TYR:HA	2:E:206:TYR:OH	2.11	0.51
2:F:197:LEU:C	2:F:197:LEU:HD23	2.30	0.51
2:F:203:ARG:N	2:F:325:ARG:HD3	2.24	0.51
2:F:439:PHE:CD1	2:F:458:ARG:CG	2.93	0.51
2:F:454:HIS:CD2	2:F:468:ILE:O	2.64	0.51
1:A:1099:TRP:O	1:A:1103:SER:N	2.37	0.51
1:A:1111:LEU:CB	1:A:1160:THR:HG23	2.40	0.51
1:A:1111:LEU:HB2	1:A:1160:THR:HG23	1.93	0.51
1:A:157:LEU:HG	1:A:194:ALA:HB1	1.92	0.51
1:A:192:GLU:CD	1:A:193:ARG:H	2.13	0.51
1:A:313:ILE:O	1:A:316:LYS:HG2	2.11	0.51
2:B:132:HIS:HA	2:B:182:ARG:HE	1.75	0.51
2:B:259:ARG:C	2:B:262:TRP:H	2.14	0.51
2:C:112:ALA:O	2:C:116:THR:HG22	2.10	0.51
2:C:306:LEU:HD12	2:C:335:VAL:HB	1.92	0.51
2:C:450:ASN:O	2:C:451:GLY:C	2.48	0.51
2:C:456:ARG:HG2	2:C:457:SER:H	1.75	0.51
1:A:505:LYS:O	1:A:506:GLU:HB2	2.11	0.51
1:D:501:LYS:HG2	1:D:502:LYS:H	1.76	0.51
2:F:367:LEU:O	2:F:369:ARG:CD	2.59	0.51
2:B:84:GLN:NE2	2:B:84:GLN:O	2.44	0.51
1:D:198:ASP:H	1:D:213:VAL:HG12	1.76	0.51
1:D:294:THR:HA	1:D:407:PHE:CZ	2.46	0.51
1:D:429:GLU:CA	1:D:429:GLU:OE1	2.59	0.51
2:F:255:TRP:HA	2:F:255:TRP:CE3	2.46	0.51
2:F:357:THR:C	2:F:359:ASN:H	2.13	0.51
2:F:84:GLN:C	2:F:86:LEU:H	2.15	0.51
1:A:1160:THR:O	1:A:1164:PHE:CD1	2.54	0.51
1:A:205:GLU:OE1	1:A:205:GLU:CA	2.57	0.51
1:A:213:VAL:HG11	1:A:397:ALA:CA	2.40	0.51
1:A:439:GLN:HG2	1:A:835:GLY:HA2	1.83	0.51
1:A:605:LEU:HB2	1:A:614:TYR:HD1	1.74	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:607:TRP:CZ2	1:A:671:GLN:CA	2.93	0.51
1:A:612:LEU:HD23	1:A:614:TYR:CA	2.40	0.51
1:A:669:MET:O	1:A:673:ALA:N	2.44	0.51
1:A:788:GLY:O	1:A:791:ALA:N	2.44	0.51
2:B:205:PRO:HG2	2:B:241:TRP:HE1	1.76	0.51
2:B:86:LEU:HB3	2:B:91:LEU:HD21	1.92	0.51
2:C:415:PRO:CA	2:C:417:TYR:HE2	2.24	0.51
2:C:455:LEU:O	2:C:456:ARG:HD2	2.11	0.51
2:E:389:ARG:HE	2:E:389:ARG:C	2.10	0.51
1:A:646:VAL:CG1	1:A:646:VAL:O	2.59	0.51
1:D:500:ALA:HB1	1:D:520:ALA:C	2.31	0.51
1:D:500:ALA:CB	1:D:520:ALA:HB3	2.41	0.51
1:D:1029:GLN:O	1:D:1032:THR:HB	2.11	0.50
1:D:1056:GLU:CB	1:D:1057:MET:HE3	2.40	0.50
1:D:115:GLN:NE2	1:D:115:GLN:O	2.44	0.50
1:D:121:PRO:HB2	1:D:1175:GLN:HE22	1.74	0.50
1:D:1198:LYS:C	1:D:1204:THR:OG1	2.49	0.50
1:D:147:SER:O	1:D:150:TYR:HB3	2.11	0.50
1:D:298:HIS:NE2	1:D:407:PHE:HA	2.25	0.50
1:D:373:PRO:HD2	1:D:374:ARG:HG2	1.93	0.50
1:D:410:GLN:O	1:D:414:PHE:CD2	2.62	0.50
1:D:417:ARG:HG2	1:D:417:ARG:HH11	1.75	0.50
1:D:550:GLN:CA	1:D:551:LYS:HE3	2.41	0.50
1:D:624:VAL:C	1:D:626:GLY:N	2.49	0.50
1:D:82:GLY:HA3	1:D:125:LEU:HD12	1.93	0.50
1:D:886:LEU:C	1:D:1146:ARG:NH1	2.50	0.50
1:D:939:VAL:HG12	1:D:939:VAL:O	2.10	0.50
1:D:994:TRP:N	1:D:996:ARG:HH11	2.09	0.50
2:E:369:ARG:CZ	2:E:369:ARG:HB3	2.40	0.50
2:E:384:ALA:HB1	2:E:385:LEU:HB3	1.89	0.50
2:E:380:PRO:CD	2:E:438:LEU:HD11	2.40	0.50
2:E:93:SER:HB3	2:E:95:CYS:SG	2.51	0.50
2:F:123:GLU:OE1	2:F:205:PRO:O	2.28	0.50
2:F:253:ASP:HB3	2:F:257:ARG:NH2	2.26	0.50
2:F:262:TRP:HZ3	2:F:263:TRP:CZ2	2.28	0.50
2:F:483:LYS:HG3	2:F:484:ASN:N	2.26	0.50
1:A:1027:LYS:HG3	1:A:1028:VAL:N	2.26	0.50
1:A:891:VAL:HB	1:A:1180:PHE:CE1	2.46	0.50
1:A:890:ASP:CG	1:A:1191:LYS:HZ3	2.10	0.50
1:A:199:VAL:HG21	1:A:277:HIS:HB2	1.92	0.50
1:A:358:LEU:HD12	1:A:407:PHE:HE1	1.76	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:490:TRP:CE2	1:A:596:MET:HE1	2.46	0.50
1:A:495:PHE:O	1:A:496:LYS:HD2	2.11	0.50
1:A:604:ALA:HA	1:A:717:LEU:HD21	1.93	0.50
1:A:844:VAL:HB	1:A:856:GLU:OE2	2.10	0.50
1:A:83:LEU:HD13	1:A:89:GLY:N	2.25	0.50
2:B:99:PHE:CE2	2:C:131:LEU:HD21	2.46	0.50
2:C:197:LEU:HD11	2:C:322:LEU:HD21	1.92	0.50
2:C:309:HIS:CA	2:C:312:LEU:HD12	2.40	0.50
2:C:431:LYS:O	2:C:435:MET:N	2.44	0.50
1:D:237:SER:OG	1:D:238:GLN:N	2.44	0.50
1:A:1193:VAL:O	1:A:1214:GLN:HB3	2.10	0.50
1:D:1161:ARG:NH1	1:D:1178:ALA:O	2.44	0.50
1:D:127:LEU:HD22	1:D:129:PRO:CG	2.42	0.50
1:D:280:GLU:HB3	1:D:814:VAL:HG11	1.93	0.50
1:D:256:PRO:CG	1:D:284:ILE:HD11	2.39	0.50
1:D:550:GLN:OE1	1:D:551:LYS:HE2	2.10	0.50
1:D:798:ILE:CA	1:D:869:ARG:NH1	2.74	0.50
1:D:920:THR:CG2	1:D:922:GLN:OE1	2.59	0.50
2:E:324:GLY:O	2:E:331:VAL:CG1	2.59	0.50
2:E:445:GLU:OE2	2:E:446:THR:HG22	2.12	0.50
2:F:270:PRO:HA	2:F:273:PHE:HD2	1.76	0.50
2:F:466:MET:SD	2:F:471:LEU:CD2	2.93	0.50
2:F:83:LYS:N	2:F:86:LEU:HD22	2.27	0.50
1:A:1187:ARG:HB2	1:A:1212:ILE:HG13	1.93	0.50
1:A:270:ASN:OD1	1:A:270:ASN:N	2.43	0.50
1:A:348:LEU:CA	1:A:350:ILE:HD13	2.28	0.50
1:A:434:TYR:H	1:A:1129:PHE:HD1	1.57	0.50
1:A:490:TRP:HH2	1:A:598:VAL:CB	2.13	0.50
1:A:576:LEU:CD1	1:A:577:CYS:N	2.68	0.50
1:A:914:THR:HB	1:A:999:ASP:HB3	1.93	0.50
2:B:375:HIS:CE1	2:B:377:CYS:SG	3.04	0.50
2:C:314:MET:HE3	2:C:315:TYR:HE2	1.77	0.50
2:C:314:MET:HB3	2:C:315:TYR:CE2	2.45	0.50
2:C:78:PHE:HZ	2:C:346:MET:HE2	1.75	0.50
2:C:266:PHE:O	2:C:349:TYR:CE1	2.64	0.50
2:C:455:LEU:N	2:C:455:LEU:CD1	2.73	0.50
2:C:474:PHE:CE2	2:C:475:LEU:HD23	2.46	0.50
1:A:539:PHE:CG	1:A:539:PHE:O	2.64	0.50
1:D:1108:TYR:HA	1:D:1164:PHE:HE1	1.76	0.50
1:D:1209:ARG:HE	2:F:253:ASP:CB	2.24	0.50
1:D:437:VAL:CG1	1:D:438:ASN:N	2.73	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:887:VAL:HG22	1:D:1185:ILE:CG2	2.37	0.50
1:D:981:LYS:HA	1:D:984:GLN:OE1	2.11	0.50
2:E:122:ARG:HB2	2:E:124:GLN:HG2	1.93	0.50
2:E:375:HIS:CG	2:E:376:PRO:HA	2.46	0.50
2:E:403:PHE:O	2:E:406:LEU:HB2	2.11	0.50
2:E:426:GLU:CD	2:E:426:GLU:N	2.65	0.50
2:E:371:VAL:CA	2:E:433:ASP:OD2	2.43	0.50
2:F:123:GLU:HB3	2:F:206:TYR:HB2	1.93	0.50
2:F:245:PRO:HG2	2:F:246:ARG:N	2.26	0.50
2:F:431:LYS:C	2:F:431:LYS:HD3	2.31	0.50
1:A:1027:LYS:HD3	1:A:1092:PHE:CE2	2.46	0.50
1:A:1153:LEU:O	1:A:1154:GLN:C	2.50	0.50
1:A:276:ALA:C	1:A:278:ILE:N	2.55	0.50
1:A:304:LEU:HD11	1:A:350:ILE:O	2.11	0.50
1:A:606:THR:H	1:A:613:HIS:CA	2.25	0.50
1:A:620:TRP:HD1	1:A:620:TRP:H	1.58	0.50
1:A:652:ILE:O	1:A:655:LEU:CG	2.58	0.50
1:A:732:TYR:CE1	1:A:755:LYS:HA	2.47	0.50
2:B:115:TRP:O	2:B:120:VAL:HG23	2.11	0.50
2:B:284:ARG:O	2:B:285:LYS:CB	2.57	0.50
2:B:372:LEU:HB2	2:B:433:ASP:O	2.10	0.50
2:C:349:TYR:C	2:C:351:TYR:N	2.62	0.50
2:C:375:HIS:N	2:C:378:LEU:HB2	2.26	0.50
1:A:737:GLY:O	1:A:738:PRO:C	2.49	0.50
1:D:1134:HIS:O	1:D:1134:HIS:HD2	1.93	0.50
1:D:220:TRP:HD1	1:D:221:TYR:CD2	2.30	0.50
1:D:269:HIS:CE1	1:D:291:PHE:C	2.84	0.50
1:D:271:VAL:O	1:D:274:ASP:CB	2.58	0.50
1:D:287:SER:C	1:D:288:ARG:HE	2.15	0.50
1:D:403:THR:O	1:D:407:PHE:HB3	2.11	0.50
1:D:453:GLU:HB3	1:D:457:ARG:HD3	1.94	0.50
1:D:768:LYS:N	1:D:772:PRO:HD3	2.27	0.50
1:D:78:MET:CE	1:D:121:PRO:HB2	2.40	0.50
1:D:790:ARG:C	1:D:792:LEU:H	2.13	0.50
1:D:827:ARG:CD	1:D:828:HIS:N	2.73	0.50
1:D:891:VAL:HG22	1:D:891:VAL:O	2.12	0.50
2:E:115:TRP:CZ2	2:E:127:PRO:HB3	2.47	0.50
2:E:199:LEU:CG	2:E:200:VAL:H	2.25	0.50
2:E:378:LEU:HD12	2:E:378:LEU:N	2.26	0.50
2:F:123:GLU:CD	2:F:205:PRO:O	2.49	0.50
2:F:206:TYR:CZ	2:F:208:LEU:HD23	2.47	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:325:ARG:CG	2:F:325:ARG:NH1	2.72	0.50
1:A:1029:GLN:CA	1:A:1031:GLU:O	2.59	0.50
1:A:1066:THR:HB	1:A:1071:ARG:HA	1.92	0.50
1:A:167:PRO:HG2	1:A:177:ARG:CG	2.25	0.50
1:A:276:ALA:HA	1:A:279:ARG:HB2	1.88	0.50
1:A:296:SER:O	1:A:299:MET:HB2	2.12	0.50
1:A:353:VAL:CG2	1:A:354:ASN:N	2.75	0.50
1:A:648:PRO:O	1:A:652:ILE:HG13	2.10	0.50
1:A:848:GLY:HA3	1:A:855:VAL:HG21	1.92	0.50
1:A:870:VAL:HG22	1:A:1202:ASN:ND2	2.24	0.50
1:A:916:PHE:CA	1:A:919:MET:HB2	2.24	0.50
1:A:972:ARG:O	1:A:974:THR:N	2.45	0.50
1:A:999:ASP:O	1:A:1000:GLU:CG	2.57	0.50
2:B:206:TYR:HA	2:B:241:TRP:CH2	2.45	0.50
2:B:105:GLU:CB	2:B:380:PRO:HB3	2.32	0.50
2:B:374:LEU:N	2:B:458:ARG:CZ	2.75	0.50
1:D:504:LYS:NZ	1:D:525:MET:HG2	2.26	0.50
1:D:512:LYS:CE	1:D:514:PRO:HD3	2.41	0.50
2:B:447:THR:CA	2:B:450:ASN:OD1	2.59	0.50
2:E:194:VAL:CG1	2:E:195:ASN:N	2.70	0.50
1:D:266:VAL:HG23	1:D:290:ARG:O	2.12	0.50
1:D:270:ASN:CA	1:D:294:THR:OG1	2.59	0.50
1:D:436:PRO:HA	1:D:839:ALA:CB	2.42	0.50
1:D:660:CYS:CA	1:D:745:PRO:HG3	2.41	0.50
1:D:783:PRO:CG	1:D:784:GLY:H	2.23	0.50
1:D:795:ASN:HA	1:D:798:ILE:CG1	2.41	0.50
1:D:831:TYR:O	1:D:831:TYR:CD1	2.62	0.50
1:D:873:GLU:OE1	1:D:1203:PRO:HG2	2.12	0.50
1:D:915:ALA:O	1:D:919:MET:SD	2.69	0.50
1:D:927:ARG:CG	1:D:928:GLY:N	2.62	0.50
1:D:980:GLU:O	1:D:984:GLN:HG3	2.11	0.50
2:E:203:ARG:HH21	2:E:203:ARG:CG	2.22	0.50
2:E:302:THR:O	2:E:303:LEU:HD12	2.11	0.50
2:F:193:TYR:O	2:F:196:CYS:HB2	2.12	0.50
2:F:203:ARG:NE	2:F:204:LEU:HD11	2.27	0.50
2:F:123:GLU:HB3	2:F:206:TYR:CB	2.41	0.50
2:F:259:ARG:NE	2:F:339:ASN:ND2	2.59	0.50
2:F:239:LEU:C	2:F:338:VAL:HG22	2.27	0.50
2:F:357:THR:O	2:F:358:GLU:CG	2.60	0.50
2:E:120:VAL:HG13	2:F:407:LEU:HD21	1.93	0.50
2:F:418:LEU:O	2:F:421:MET:CG	2.54	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:466:MET:HE2	2:F:471:LEU:HB3	1.92	0.50
1:A:1131:ILE:CB	1:A:1138:ARG:H	2.24	0.50
1:A:266:VAL:HG22	1:A:267:VAL:N	2.26	0.50
1:A:607:TRP:CE2	1:A:671:GLN:NE2	2.79	0.50
1:A:771:LEU:O	1:A:775:GLU:HG3	2.11	0.50
1:A:786:ALA:O	1:A:790:ARG:HG2	2.12	0.50
1:A:86:GLN:OE1	1:A:130:LEU:HB3	2.11	0.50
1:A:886:LEU:CA	1:A:1146:ARG:NH2	2.74	0.50
1:A:905:ALA:HA	1:A:910:MET:CA	2.42	0.50
1:A:945:HIS:CE1	1:A:969:PHE:CB	2.94	0.50
1:A:966:LEU:O	1:A:970:ASN:HB2	2.12	0.50
1:A:996:ARG:NH1	1:A:1000:GLU:CB	2.52	0.50
1:A:998:SER:OG	1:A:999:ASP:N	2.42	0.50
2:B:259:ARG:HG2	2:B:262:TRP:CD1	2.47	0.50
2:B:348:ALA:O	2:B:350:LEU:N	2.45	0.50
2:B:67:GLU:CA	2:B:70:LEU:HB2	2.41	0.50
2:B:93:SER:O	2:B:95:CYS:N	2.44	0.50
2:C:126:PHE:HB2	2:C:128:VAL:HA	1.92	0.50
2:C:241:TRP:HE3	2:C:336:LEU:CD1	2.24	0.50
2:C:241:TRP:CG	2:C:336:LEU:HD12	2.45	0.50
2:E:388:GLY:O	2:E:395:LEU:CD1	2.59	0.50
2:B:447:THR:O	2:B:451:GLY:N	2.44	0.50
2:B:217:PRO:CA	2:B:232:GLY:H	2.24	0.50
1:A:543:VAL:HG13	1:A:544:MET:N	2.27	0.50
1:D:1131:ILE:HD13	1:D:1131:ILE:O	2.11	0.50
1:D:1147:TYR:HD2	1:D:1147:TYR:N	2.10	0.50
1:D:1187:ARG:HG3	1:D:1212:ILE:CG1	2.42	0.50
1:D:177:ARG:CG	1:D:218:SER:OG	2.60	0.50
1:D:194:ALA:CB	1:D:265:LEU:HD13	2.39	0.50
1:D:272:SER:O	1:D:273:PHE:HB2	2.10	0.50
1:D:307:PHE:O	1:D:308:GLN:C	2.50	0.50
1:D:375:GLU:OE1	1:D:379:LYS:HB3	2.11	0.50
1:D:472:GLN:C	2:F:461:THR:CG2	2.80	0.50
1:D:547:ALA:O	1:D:551:LYS:CG	2.58	0.50
1:D:800:PHE:CE2	1:D:867:PRO:HB3	2.47	0.50
1:D:85:GLU:OE2	1:D:127:LEU:O	2.30	0.50
2:E:425:LEU:HD23	2:E:426:GLU:OE1	2.12	0.50
2:E:449:GLU:CA	2:E:449:GLU:OE1	2.45	0.50
2:E:373:LYS:HA	2:E:458:ARG:NE	2.25	0.50
2:F:198:ASP:OD1	2:F:198:ASP:C	2.48	0.50
1:A:1121:GLU:O	1:A:1123:PHE:N	2.39	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1175:GLN:O	1:A:1178:ALA:HB3	2.11	0.50
1:A:486:TRP:HE1	1:A:722:ARG:HG2	1.77	0.50
1:A:657:ARG:CA	1:A:660:CYS:HB3	2.41	0.50
1:A:662:GLU:O	1:A:666:GLN:CB	2.60	0.50
1:A:824:ALA:HB1	1:A:827:ARG:HG2	1.93	0.50
1:A:931:LEU:HD23	1:A:932:HIS:CA	2.42	0.50
1:A:936:ALA:O	1:A:939:VAL:C	2.50	0.50
2:B:320:SER:HA	2:B:323:HIS:CD2	2.47	0.50
2:B:383:VAL:CG2	2:B:384:ALA:H	2.23	0.50
2:C:193:TYR:CG	2:C:194:VAL:N	2.80	0.50
2:C:326:ASP:CG	2:C:329:LYS:HB2	2.32	0.50
2:C:353:SER:HG	2:C:354:PHE:H	1.55	0.50
2:C:389:ARG:CZ	2:C:390:GLY:CA	2.90	0.50
2:C:78:PHE:HZ	2:C:346:MET:SD	2.35	0.50
2:E:88:ARG:CD	2:E:88:ARG:H	2.24	0.50
1:D:1124:ALA:HB1	1:D:1148:ARG:HH12	1.70	0.50
1:D:204:ALA:O	1:D:205:GLU:CG	2.60	0.50
1:D:245:ILE:CG1	1:D:246:PRO:N	2.51	0.50
1:D:439:GLN:CG	1:D:836:LEU:H	2.25	0.50
1:D:453:GLU:O	1:D:454:GLU:C	2.49	0.50
1:D:549:LEU:O	1:D:552:LEU:HB2	2.12	0.50
1:D:657:ARG:O	1:D:660:CYS:CB	2.58	0.50
1:D:657:ARG:C	1:D:660:CYS:HB3	2.31	0.50
1:D:732:TYR:HB3	1:D:755:LYS:HZ1	1.76	0.50
1:D:802:ARG:HG3	1:D:803:ASN:N	2.27	0.50
1:D:827:ARG:CD	1:D:827:ARG:C	2.80	0.50
1:D:938:THR:O	1:D:939:VAL:CG2	2.59	0.50
2:E:372:LEU:O	2:E:458:ARG:HG2	2.12	0.50
2:F:432:TYR:C	2:F:435:MET:H	2.15	0.50
2:F:447:THR:CG2	2:F:448:LEU:N	2.73	0.50
2:F:460:THR:O	2:F:461:THR:C	2.50	0.50
1:A:1016:GLU:OE2	1:A:1022:LEU:HD22	2.12	0.50
1:A:1158:LEU:HD13	1:A:1179:PHE:CZ	2.47	0.50
1:A:81:ARG:NH1	1:A:125:LEU:O	2.45	0.50
1:A:660:CYS:CB	1:A:745:PRO:CG	2.90	0.50
2:B:338:VAL:C	2:B:339:ASN:ND2	2.65	0.50
2:C:213:VAL:HA	2:C:235:THR:HA	1.94	0.50
2:C:389:ARG:NH2	2:C:390:GLY:HA3	2.26	0.50
2:C:471:LEU:CD1	2:C:471:LEU:N	2.38	0.50
1:A:1055:SER:O	1:A:1059:ASN:CG	2.50	0.50
2:C:224:ILE:N	2:C:224:ILE:HD13	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1068:ASP:OD1	1:D:1068:ASP:N	2.44	0.50
1:D:429:GLU:CG	1:D:1128:ARG:NH2	2.66	0.50
1:D:1126:ASP:CB	1:D:1141:VAL:HG13	2.42	0.50
1:D:239:LEU:C	1:D:242:ALA:H	2.14	0.50
1:D:295:MET:O	1:D:299:MET:N	2.41	0.50
1:D:308:GLN:O	1:D:312:TRP:HB2	2.12	0.50
1:D:428:LEU:O	1:D:432:VAL:HG23	2.11	0.50
1:D:457:ARG:C	1:D:460:LYS:HZ2	2.15	0.50
1:D:727:ASP:O	1:D:728:THR:OG1	2.25	0.50
1:D:907:PHE:HE2	1:D:913:CYS:SG	2.35	0.50
2:E:90:SER:OG	2:E:96:HIS:HA	2.12	0.50
2:F:196:CYS:O	2:F:199:LEU:HB3	2.12	0.50
2:F:204:LEU:CD1	2:F:204:LEU:N	2.75	0.50
2:F:307:GLY:HA2	2:F:335:VAL:HG12	1.94	0.50
2:F:354:PHE:C	2:F:356:LEU:N	2.60	0.50
2:F:447:THR:HG22	2:F:448:LEU:HD12	1.94	0.50
1:A:1207:GLU:O	1:A:1208:ARG:HB2	2.12	0.50
1:A:291:PHE:O	1:A:292:LEU:CB	2.59	0.50
1:A:371:LYS:C	1:A:373:PRO:CD	2.79	0.50
1:A:355:SER:C	1:A:407:PHE:HZ	2.15	0.50
1:A:658:LYS:O	1:A:659:HIS:C	2.50	0.50
1:A:789:PRO:O	1:A:792:LEU:HD12	2.12	0.50
1:A:863:SER:O	1:A:873:GLU:CG	2.57	0.50
1:A:953:ARG:HD3	1:A:986:TYR:CD2	2.47	0.50
2:B:194:VAL:HG13	2:B:195:ASN:N	2.27	0.50
2:B:105:GLU:HG3	2:B:380:PRO:C	2.31	0.50
2:B:437:ILE:N	2:B:458:ARG:HD3	2.26	0.50
2:C:118:VAL:HG12	2:C:119:VAL:N	2.27	0.50
2:C:307:GLY:N	2:C:335:VAL:HG12	2.27	0.50
2:C:469:SER:C	2:C:471:LEU:HD12	2.31	0.50
2:C:377:CYS:SG	2:C:483:LYS:HA	2.51	0.50
2:C:82:SER:C	2:C:83:LYS:HG3	2.32	0.50
2:E:389:ARG:HE	2:E:390:GLY:CA	2.23	0.50
2:B:389:ARG:HG2	2:B:444:THR:CG2	2.42	0.50
1:D:106:HIS:HA	1:D:109:LYS:CG	2.41	0.50
1:A:564:GLN:HA	1:A:567:PRO:CD	2.35	0.50
1:D:1090:GLU:HG3	1:D:1092:PHE:CZ	2.46	0.50
1:D:1106:VAL:O	1:D:1107:ASP:C	2.50	0.50
1:D:121:PRO:CB	1:D:1175:GLN:NE2	2.73	0.50
1:D:131:TYR:C	1:D:138:HIS:CE1	2.86	0.50
1:D:278:ILE:CG1	1:D:291:PHE:HD1	2.24	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:369:LEU:HD13	1:D:372:GLU:CB	2.41	0.50
1:D:452:TYR:CB	1:D:801:TRP:CD1	2.94	0.50
1:D:592:LEU:O	1:D:595:GLN:N	2.40	0.50
1:D:885:THR:HG22	1:D:886:LEU:H	1.75	0.50
2:E:192:HIS:O	2:E:193:TYR:C	2.49	0.50
2:F:241:TRP:CZ3	2:F:255:TRP:CZ2	2.99	0.50
2:F:348:ALA:O	2:F:352:ASP:N	2.41	0.50
2:F:375:HIS:CB	2:F:378:LEU:HD22	2.40	0.50
2:F:388:GLY:N	2:F:443:VAL:O	2.42	0.50
1:A:1068:ASP:CA	1:A:1071:ARG:HD3	2.41	0.50
1:A:1140:LEU:CD2	1:A:1141:VAL:H	2.20	0.50
1:A:183:GLU:C	1:A:183:GLU:CD	2.70	0.50
1:A:204:ALA:O	1:A:206:GLY:N	2.40	0.50
1:A:494:GLU:HB2	1:A:572:TRP:CD2	2.47	0.50
1:A:605:LEU:C	1:A:781:ALA:CB	2.81	0.50
1:A:754:HIS:HD2	1:A:757:GLY:HA3	1.77	0.50
1:A:892:ASP:HB3	1:A:927:ARG:HH22	1.77	0.50
2:B:440:THR:CG2	2:B:441:VAL:H	2.19	0.50
2:C:414:TRP:N	2:C:415:PRO:HD2	2.25	0.50
2:C:428:LEU:HD23	2:C:428:LEU:N	2.22	0.50
2:C:443:VAL:CG1	2:C:447:THR:CG2	2.89	0.50
1:A:645:VAL:HA	1:A:650:ARG:HD2	1.94	0.50
1:A:742:VAL:O	1:A:742:VAL:HG12	2.11	0.50
2:F:272:ASN:HA	2:F:292:ASN:ND2	2.26	0.50
1:D:262:GLN:OE1	1:D:262:GLN:HA	2.11	0.50
1:D:162:LEU:HG	1:D:162:LEU:O	2.12	0.50
1:D:1025:LEU:O	1:D:1027:LYS:N	2.43	0.49
1:D:1055:SER:O	1:D:1058:PHE:CD2	2.65	0.49
1:D:886:LEU:CA	1:D:1146:ARG:NH1	2.75	0.49
1:D:116:PRO:O	1:D:118:VAL:CG1	2.61	0.49
1:D:1207:GLU:HA	1:D:1210:TYR:CG	2.47	0.49
1:D:1213:PRO:O	1:D:1214:GLN:CB	2.57	0.49
1:D:298:HIS:CD2	1:D:407:PHE:CB	2.90	0.49
1:D:389:PHE:CD1	1:D:389:PHE:C	2.85	0.49
1:D:712:VAL:N	1:D:713:PRO:CD	2.75	0.49
1:D:427:MET:HB2	1:D:846:THR:CG2	2.42	0.49
1:D:847:ALA:C	1:D:852:ARG:HA	2.32	0.49
1:D:860:LEU:HD22	1:D:1133:ILE:HD12	1.94	0.49
1:D:891:VAL:HA	1:D:894:GLN:CG	2.41	0.49
2:E:109:ASN:HD22	2:E:378:LEU:CB	2.26	0.49
2:E:374:LEU:O	2:E:375:HIS:CB	2.52	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:96:HIS:CD2	2:E:96:HIS:O	2.65	0.49
2:F:244:PRO:N	2:F:247:THR:HG21	2.27	0.49
2:F:349:TYR:C	2:F:351:TYR:N	2.62	0.49
2:F:346:MET:O	2:F:350:LEU:HD11	2.12	0.49
2:F:357:THR:O	2:F:359:ASN:N	2.42	0.49
2:F:447:THR:CA	2:F:452:LEU:CA	2.81	0.49
2:F:457:SER:O	2:F:458:ARG:C	2.50	0.49
2:F:483:LYS:O	2:F:485:VAL:HG12	2.12	0.49
1:A:1092:PHE:O	1:A:1096:ARG:HG3	2.12	0.49
1:A:286:GLY:CA	1:A:1128:ARG:HH22	2.24	0.49
1:A:1163:MET:HE2	1:A:1167:LYS:HE3	1.94	0.49
1:A:174:GLY:HA2	1:A:220:TRP:HB2	1.94	0.49
1:A:192:GLU:CG	1:A:195:LEU:HD21	2.42	0.49
1:A:227:ARG:HD3	1:A:227:ARG:C	2.31	0.49
1:A:272:SER:C	1:A:274:ASP:H	2.16	0.49
1:A:392:LEU:O	1:A:394:GLN:N	2.45	0.49
1:A:441:TRP:O	1:A:445:LEU:HB2	2.12	0.49
1:A:452:TYR:CE2	1:A:453:GLU:HG2	2.47	0.49
1:A:730:PRO:HB3	1:A:760:CYS:O	2.12	0.49
1:A:764:SER:OG	1:A:767:ALA:HB2	2.11	0.49
2:B:114:TRP:CA	2:B:262:TRP:CH2	2.95	0.49
2:C:306:LEU:H	2:C:335:VAL:CG1	2.24	0.49
2:C:347:LEU:CD1	2:C:347:LEU:H	2.24	0.49
2:C:474:PHE:HA	2:C:477:LYS:HD3	1.92	0.49
1:A:503:VAL:HG11	1:A:520:ALA:CB	2.42	0.49
2:E:418:LEU:HD11	2:F:122:ARG:HE	1.74	0.49
2:B:472:LYS:HZ2	2:B:473:ASP:HA	1.77	0.49
2:C:311:LEU:HD22	2:C:311:LEU:N	2.26	0.49
2:E:392:THR:HG22	2:E:392:THR:O	2.11	0.49
1:D:1027:LYS:O	1:D:1031:GLU:HG2	2.12	0.49
1:D:1031:GLU:OE2	1:D:1092:PHE:HD2	1.95	0.49
1:D:1116:MET:HE3	1:D:1156:THR:HB	1.93	0.49
1:D:252:GLY:O	1:D:256:PRO:CG	2.58	0.49
1:D:270:ASN:C	1:D:274:ASP:OD1	2.50	0.49
1:D:298:HIS:CE1	1:D:362:LEU:CD1	2.95	0.49
1:D:359:VAL:HG22	1:D:406:VAL:CG2	2.42	0.49
1:D:371:LYS:CA	1:D:374:ARG:CD	2.77	0.49
1:D:407:PHE:CE1	1:D:411:LEU:HD21	2.46	0.49
1:D:607:TRP:HA	1:D:607:TRP:CE3	2.47	0.49
1:D:607:TRP:O	1:D:608:ASP:OD1	2.30	0.49
1:D:275:ARG:CG	1:D:843:GLN:N	2.75	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:441:TRP:CD1	1:D:879:GLN:HA	2.45	0.49
2:E:200:VAL:HG12	2:F:418:LEU:HD21	1.92	0.49
2:E:239:LEU:CB	2:E:338:VAL:HB	2.42	0.49
2:F:308:ASP:OD1	2:F:333:PRO:C	2.50	0.49
2:F:358:GLU:CG	2:F:359:ASN:N	2.75	0.49
2:F:413:VAL:CG1	2:F:414:TRP:N	2.76	0.49
2:F:384:ALA:CB	2:F:437:ILE:HD12	2.42	0.49
2:F:471:LEU:O	2:F:474:PHE:N	2.39	0.49
1:A:246:PRO:O	1:A:250:PRO:HD2	2.12	0.49
1:A:304:LEU:HB2	1:A:309:ARG:HB3	1.94	0.49
1:A:421:PRO:CG	1:A:422:VAL:N	2.75	0.49
1:A:789:PRO:O	1:A:793:GLU:HB2	2.12	0.49
1:A:960:PRO:HA	1:A:963:GLU:CG	2.43	0.49
1:A:755:LYS:CD	1:A:964:ARG:HH11	2.23	0.49
2:B:208:LEU:N	2:B:208:LEU:HD23	2.26	0.49
2:B:425:LEU:HB3	2:B:426:GLU:OE1	2.11	0.49
2:C:127:PRO:O	2:C:192:HIS:CD2	2.65	0.49
2:E:249:ASN:O	2:E:252:LEU:HB3	2.12	0.49
2:E:307:GLY:HA2	2:E:334:CYS:HB2	1.94	0.49
1:D:1092:PHE:HA	1:D:1096:ARG:NH2	2.27	0.49
1:D:260:ASP:OD2	1:D:261:TRP:CD2	2.65	0.49
1:D:298:HIS:HD2	1:D:407:PHE:CD1	2.30	0.49
1:D:643:ALA:N	1:D:645:VAL:HB	2.26	0.49
1:D:275:ARG:HB3	1:D:843:GLN:OE1	2.12	0.49
1:D:853:ARG:HH12	1:D:1105:ALA:HB3	1.76	0.49
1:D:931:LEU:HA	1:D:934:LYS:HB3	1.93	0.49
2:E:216:HIS:N	2:E:216:HIS:CD2	2.79	0.49
2:E:385:LEU:HB2	2:E:440:THR:HA	1.93	0.49
2:E:380:PRO:HG3	2:E:438:LEU:CD1	2.40	0.49
2:E:74:GLN:O	2:E:77:HIS:N	2.44	0.49
2:F:258:HIS:HA	2:F:261:GLN:OE1	2.12	0.49
2:F:306:LEU:O	2:F:335:VAL:HB	2.11	0.49
2:F:350:LEU:O	2:F:353:SER:OG	2.30	0.49
2:F:389:ARG:O	2:F:395:LEU:HD11	2.12	0.49
2:F:393:LEU:O	2:F:396:ARG:N	2.46	0.49
2:F:405:GLU:OE1	2:F:405:GLU:C	2.51	0.49
2:F:76:ARG:NH1	2:F:434:GLU:OE1	2.45	0.49
1:A:1065:ALA:O	1:A:1066:THR:OG1	2.22	0.49
1:A:1144:GLU:O	1:A:1146:ARG:N	2.45	0.49
1:A:1209:ARG:HE	1:A:1209:ARG:HA	1.78	0.49
1:A:228:LEU:N	1:A:228:LEU:CD2	2.73	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:421:PRO:HG2	1:A:422:VAL:HG23	1.93	0.49
1:A:583:PRO:O	1:A:584:ALA:HB3	2.12	0.49
1:A:599:THR:O	1:A:602:LEU:HB2	2.12	0.49
1:A:630:ASN:HD21	1:A:632:ALA:N	2.11	0.49
1:A:732:TYR:HE1	1:A:755:LYS:CG	2.24	0.49
1:A:762:VAL:O	1:A:764:SER:N	2.38	0.49
1:A:821:LEU:HD11	1:A:826:ILE:N	2.27	0.49
1:A:275:ARG:NH2	1:A:845:VAL:N	2.60	0.49
1:A:866:ARG:HB2	1:A:1069:ILE:HG21	1.94	0.49
1:A:878:VAL:HG12	1:A:879:GLN:HE22	1.77	0.49
1:A:919:MET:C	1:A:920:THR:O	2.51	0.49
2:B:205:PRO:HG2	2:B:241:TRP:NE1	2.27	0.49
2:B:259:ARG:HB3	2:B:259:ARG:NH1	2.27	0.49
2:B:336:LEU:HD23	2:B:336:LEU:C	2.32	0.49
2:B:463:LYS:HD2	2:B:463:LYS:N	2.27	0.49
2:C:270:PRO:HA	2:C:273:PHE:HD2	1.77	0.49
2:C:425:LEU:O	2:C:428:LEU:HB2	2.12	0.49
1:A:95:PRO:O	1:A:96:GLY:O	2.30	0.49
1:A:541:GLN:O	1:A:544:MET:HB3	2.12	0.49
1:D:561:LYS:O	1:D:562:ARG:CB	2.60	0.49
1:D:756:ASP:N	1:D:756:ASP:OD1	2.44	0.49
1:D:1051:GLY:O	1:D:1055:SER:N	2.45	0.49
1:D:1116:MET:HG3	1:D:1139:TYR:OH	2.13	0.49
1:D:1218:LEU:HB2	1:D:1223:ILE:HG22	1.90	0.49
1:D:371:LYS:O	1:D:372:GLU:CB	2.60	0.49
1:D:735:GLY:HA2	1:D:749:PHE:CA	2.43	0.49
1:D:779:LEU:CD2	1:D:779:LEU:C	2.79	0.49
1:D:840:ILE:O	1:D:841:LEU:CD2	2.60	0.49
2:E:432:TYR:HA	2:E:435:MET:CA	2.41	0.49
2:E:456:ARG:HD2	2:E:461:THR:CA	2.42	0.49
2:F:239:LEU:H	2:F:338:VAL:HG11	1.53	0.49
2:F:239:LEU:HG	2:F:336:LEU:HD13	1.94	0.49
2:F:424:SER:N	2:F:427:GLN:OE1	2.45	0.49
2:F:442:LEU:O	2:F:443:VAL:C	2.50	0.49
2:F:389:ARG:CG	2:F:445:GLU:OE2	2.58	0.49
2:F:74:GLN:CG	2:F:75:ARG:N	2.75	0.49
2:F:78:PHE:CD2	2:F:79:LEU:CD2	2.95	0.49
2:F:80:SER:H	2:F:99:PHE:HA	1.78	0.49
2:F:82:SER:C	2:F:83:LYS:HG3	2.33	0.49
1:A:1026:ARG:CG	1:A:1029:GLN:CD	2.81	0.49
1:A:1093:MET:C	1:A:1097:VAL:HG23	2.32	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1113:LEU:CB	1:A:1117:LYS:NZ	2.74	0.49
1:A:1232:GLU:CD	1:A:1234:ARG:NH2	2.65	0.49
1:A:240:SER:HB3	1:A:241:PRO:CD	2.42	0.49
1:A:298:HIS:NE2	1:A:351:SER:O	2.45	0.49
1:A:353:VAL:O	1:A:354:ASN:CB	2.60	0.49
1:A:434:TYR:HA	1:A:841:LEU:CD2	2.41	0.49
1:A:580:LEU:HD23	1:A:596:MET:CE	2.42	0.49
1:A:76:ILE:HG21	1:A:911:HIS:CE1	2.47	0.49
1:A:772:PRO:HG2	1:A:773:LYS:H	1.77	0.49
1:A:79:LEU:HB2	1:A:89:GLY:C	2.32	0.49
2:B:342:LEU:O	2:B:343:ASP:C	2.50	0.49
2:B:414:TRP:C	2:B:414:TRP:CD1	2.86	0.49
1:A:572:TRP:CD1	2:C:477:LYS:HE2	2.47	0.49
2:C:477:LYS:O	2:C:478:TYR:C	2.49	0.49
2:E:270:PRO:C	2:E:273:PHE:HD2	2.15	0.49
1:D:1010:LEU:C	1:D:1012:VAL:N	2.63	0.49
1:D:273:PHE:N	1:D:275:ARG:HB3	2.23	0.49
1:D:412:PRO:CG	1:D:413:LEU:H	2.23	0.49
1:D:748:TRP:CZ3	1:D:750:PHE:CE2	2.99	0.49
1:D:829:PRO:HG3	1:D:831:TYR:CE1	2.47	0.49
1:D:904:ASP:OD1	1:D:915:ALA:HB2	2.13	0.49
2:E:404:ASN:C	2:E:406:LEU:N	2.66	0.49
2:E:79:LEU:HD12	2:E:79:LEU:C	2.33	0.49
2:F:289:LEU:HD12	2:F:301:GLU:CB	2.32	0.49
2:F:291:TYR:HB2	2:F:300:ILE:HD11	1.95	0.49
2:F:368:HIS:NE2	2:F:370:LYS:CE	2.75	0.49
2:F:372:LEU:HD12	2:F:372:LEU:C	2.33	0.49
2:F:453:ILE:CG2	2:F:454:HIS:N	2.75	0.49
2:F:458:ARG:HD3	2:F:459:ASP:N	2.28	0.49
1:A:185:VAL:HG22	1:A:185:VAL:O	2.12	0.49
1:A:219:ALA:O	1:A:220:TRP:HE3	1.93	0.49
1:A:420:HIS:CG	1:A:421:PRO:HD2	2.47	0.49
1:A:467:ALA:CB	1:A:602:LEU:HD13	2.42	0.49
1:A:591:LEU:HD21	1:A:802:ARG:CZ	2.42	0.49
1:A:713:PRO:CA	1:A:719:LEU:HD13	2.42	0.49
1:A:773:LYS:NZ	1:A:1057:MET:CE	2.75	0.49
1:A:856:GLU:OE1	1:A:859:TRP:HB2	2.11	0.49
2:B:78:PHE:CD1	2:B:102:LEU:HB2	2.48	0.49
2:B:403:PHE:CE1	2:B:413:VAL:HB	2.47	0.49
2:B:79:LEU:HA	2:B:100:GLY:N	2.27	0.49
2:C:123:GLU:HG2	2:C:204:LEU:CD1	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:289:LEU:HD12	2:C:301:GLU:CB	2.35	0.49
2:C:347:LEU:O	2:C:350:LEU:HD13	2.12	0.49
2:C:431:LYS:HZ3	2:C:435:MET:CG	2.24	0.49
2:C:456:ARG:CB	2:C:463:LYS:C	2.79	0.49
1:A:504:LYS:HD3	1:A:523:ASP:HB3	1.92	0.49
2:E:391:PRO:CD	2:E:395:LEU:HB2	2.39	0.49
2:E:466:MET:HG2	2:E:467:HIS:N	2.27	0.49
2:E:481:SER:HA	2:E:484:ASN:CG	2.33	0.49
2:F:90:SER:HA	2:F:93:SER:OG	2.12	0.49
1:D:1047:ARG:O	1:D:1051:GLY:N	2.43	0.49
1:D:1114:VAL:HA	1:D:1117:LYS:HD2	1.95	0.49
1:D:1118:TRP:CE2	1:D:1122:GLU:OE1	2.65	0.49
1:D:444:TYR:HE2	1:D:1210:TYR:HB2	1.78	0.49
1:D:1210:TYR:CD1	1:D:1211:GLY:N	2.80	0.49
1:D:1224:ILE:CD1	1:D:1228:LYS:HE3	2.42	0.49
1:D:194:ALA:C	1:D:195:LEU:HG	2.33	0.49
1:D:361:ARG:NH2	1:D:362:LEU:HG	2.28	0.49
1:D:430:MET:O	1:D:430:MET:SD	2.70	0.49
1:D:435:LEU:O	1:D:839:ALA:CB	2.58	0.49
1:D:494:GLU:H	1:D:573:TYR:HB2	1.77	0.49
1:D:635:PRO:O	1:D:636:THR:HB	2.11	0.49
1:D:655:LEU:O	1:D:657:ARG:N	2.46	0.49
1:D:615:SER:O	1:D:725:PRO:HG3	2.12	0.49
1:D:768:LYS:HE2	1:D:769:ASP:H	1.76	0.49
1:D:868:ASP:OD1	1:D:1199:THR:CG2	2.60	0.49
1:D:891:VAL:HA	1:D:894:GLN:OE1	2.10	0.49
2:E:323:HIS:O	2:E:325:ARG:N	2.45	0.49
2:E:328:ARG:N	2:E:328:ARG:CD	2.68	0.49
2:E:443:VAL:HG12	2:E:444:THR:N	2.28	0.49
2:E:462:MET:CE	2:E:462:MET:HA	2.42	0.49
2:E:86:LEU:HD23	2:E:97:PRO:HD2	1.95	0.49
2:F:251:TRP:CZ3	2:F:254:PHE:CG	3.00	0.49
2:F:336:LEU:CD2	2:F:337:SER:O	2.61	0.49
2:F:371:VAL:CG2	2:F:433:ASP:O	2.60	0.49
2:F:449:GLU:OE1	2:F:450:ASN:N	2.46	0.49
1:A:1043:VAL:CB	1:A:1046:GLU:HB3	2.41	0.49
1:A:1148:ARG:CG	1:A:1148:ARG:NH1	2.62	0.49
1:A:1223:ILE:HD13	1:A:1224:ILE:N	2.27	0.49
1:A:260:ASP:H	1:A:264:GLN:HE22	1.57	0.49
1:A:420:HIS:CE1	1:A:421:PRO:HD2	2.47	0.49
1:A:485:LEU:O	1:A:489:GLU:OE2	2.31	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:719:LEU:CD2	1:A:719:LEU:H	2.23	0.49
1:A:657:ARG:HG2	1:A:744:ILE:HG21	1.94	0.49
1:A:795:ASN:ND2	1:A:795:ASN:C	2.65	0.49
1:A:853:ARG:HH22	1:A:1105:ALA:HB3	1.72	0.49
1:A:869:ARG:O	1:A:1200:PRO:CA	2.58	0.49
2:B:206:TYR:HD1	2:B:207:GLY:O	1.94	0.49
2:C:387:VAL:HG23	2:C:417:TYR:HA	1.95	0.49
2:C:414:TRP:O	2:C:415:PRO:C	2.50	0.49
2:C:78:PHE:CD1	2:C:103:GLY:CA	2.93	0.49
1:D:95:PRO:HG2	1:D:99:ALA:HB2	1.89	0.49
2:F:145:PHE:O	2:F:146:ARG:HG3	2.12	0.49
1:D:443:ARG:O	1:D:447:GLU:N	2.43	0.49
2:E:88:ARG:CD	2:E:88:ARG:N	2.75	0.49
1:D:1067:SER:CB	1:D:1073:PRO:HG2	2.42	0.49
1:D:1162:CYS:O	1:D:1165:ALA:N	2.46	0.49
1:D:175:TRP:C	1:D:185:VAL:HG11	2.32	0.49
1:D:213:VAL:CB	1:D:401:TRP:CZ2	2.96	0.49
1:D:288:ARG:NH1	1:D:290:ARG:NH2	2.61	0.49
1:D:203:LEU:HB2	1:D:379:LYS:O	2.13	0.49
1:D:383:LYS:HD3	1:D:383:LYS:N	2.27	0.49
1:D:453:GLU:O	1:D:457:ARG:CG	2.59	0.49
1:D:459:MET:HB3	1:D:460:LYS:NZ	2.27	0.49
1:D:578:PRO:O	1:D:579:ARG:C	2.50	0.49
1:D:719:LEU:HD12	1:D:720:THR:CB	2.43	0.49
1:D:735:GLY:HA2	1:D:750:PHE:H	1.77	0.49
1:D:854:ALA:C	1:D:855:VAL:HG23	2.33	0.49
2:E:129:ASP:HB2	2:F:99:PHE:O	2.12	0.49
2:E:183:GLU:CG	2:E:216:HIS:CD2	2.96	0.49
2:F:128:VAL:HG12	2:F:129:ASP:N	2.27	0.49
2:F:246:ARG:N	2:F:246:ARG:HD3	2.27	0.49
2:F:344:ARG:HA	2:F:347:LEU:CD1	2.39	0.49
2:F:466:MET:O	2:F:467:HIS:C	2.50	0.49
1:A:1029:GLN:CD	1:A:1096:ARG:NH2	2.65	0.49
1:A:296:SER:HB2	1:A:847:ALA:CB	2.36	0.49
1:A:515:ILE:O	1:A:516:GLU:C	2.50	0.49
1:A:659:HIS:CG	1:A:716:PRO:HA	2.47	0.49
1:A:613:HIS:NE2	1:A:716:PRO:HG3	2.28	0.49
1:A:455:LEU:CD1	1:A:798:ILE:HG22	2.40	0.49
1:A:869:ARG:N	1:A:872:SER:CB	2.75	0.49
1:A:877:MET:O	1:A:878:VAL:C	2.51	0.49
1:A:947:LYS:O	1:A:951:TYR:HD1	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:186:LEU:O	2:C:189:ALA:HB3	2.12	0.49
2:C:192:HIS:C	2:C:196:CYS:SG	2.91	0.49
2:C:123:GLU:HG2	2:C:204:LEU:HD12	1.95	0.49
2:C:397:GLN:O	2:C:398:VAL:CG2	2.55	0.49
2:C:447:THR:HG22	2:C:455:LEU:CD1	2.43	0.49
2:C:452:LEU:O	2:C:453:ILE:HB	2.13	0.49
1:A:500:ALA:O	1:A:501:LYS:HB2	2.12	0.49
1:A:307:PHE:HB3	1:A:1055:SER:OG	2.13	0.49
1:D:765:PRO:HG2	1:D:766:PHE:CE1	2.48	0.49
1:A:235:TRP:C	1:A:235:TRP:CD1	2.85	0.49
1:D:1065:ALA:O	1:D:1073:PRO:CG	2.59	0.49
1:D:873:GLU:OE1	1:D:1200:PRO:HG3	2.13	0.49
1:D:177:ARG:CB	1:D:218:SER:HB3	2.26	0.49
1:D:374:ARG:CG	1:D:374:ARG:HH11	2.25	0.49
1:D:374:ARG:NE	1:D:392:LEU:HG	2.27	0.49
1:D:439:GLN:CG	1:D:835:GLY:HA3	2.40	0.49
1:D:453:GLU:O	1:D:457:ARG:N	2.46	0.49
1:D:483:PRO:HG3	1:D:712:VAL:HG13	1.95	0.49
1:D:493:GLN:HG2	1:D:494:GLU:N	2.28	0.49
1:D:656:TYR:CA	1:D:659:HIS:CE1	2.95	0.49
1:D:791:ALA:C	1:D:792:LEU:HD12	2.32	0.49
1:D:801:TRP:CH2	1:D:805:HIS:ND1	2.75	0.49
1:D:950:ASN:OD1	1:D:953:ARG:HD2	2.13	0.49
1:D:959:GLN:HG2	1:D:983:GLN:NE2	2.27	0.49
2:E:325:ARG:HG2	2:E:330:ASN:N	2.27	0.49
2:E:113:GLU:HG3	2:E:378:LEU:CD1	2.42	0.49
2:F:354:PHE:CD2	2:F:354:PHE:O	2.66	0.49
2:F:385:LEU:CD2	2:F:402:LEU:HD23	2.42	0.49
2:F:432:TYR:HB3	2:F:437:ILE:CD1	2.41	0.49
1:A:196:VAL:CB	1:A:401:TRP:CZ3	2.86	0.49
1:A:212:ALA:H	1:A:221:TYR:HA	1.78	0.49
1:A:215:ILE:O	1:A:216:SER:C	2.51	0.49
1:A:355:SER:O	1:A:407:PHE:CZ	2.66	0.49
1:A:616:GLU:C	1:A:725:PRO:HG3	2.33	0.49
1:A:732:TYR:CE1	1:A:755:LYS:CG	2.95	0.49
1:A:900:ALA:C	1:A:915:ALA:HB1	2.32	0.49
1:A:992:LEU:HD13	1:A:993:ARG:N	2.26	0.49
2:B:128:VAL:HG21	2:B:208:LEU:HD12	1.94	0.49
2:B:432:TYR:HA	2:B:437:ILE:HD12	1.95	0.49
2:B:67:GLU:HB3	2:B:70:LEU:CD1	2.43	0.49
1:D:513:LEU:HB2	1:D:568:GLY:CA	2.32	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:515:ILE:HG12	1:D:568:GLY:O	2.13	0.49
1:D:564:GLN:O	1:D:565:HIS:C	2.51	0.49
2:B:280:ASP:HB3	2:B:281:GLU:HB2	1.93	0.49
2:B:450:ASN:HD22	2:B:452:LEU:CD2	2.22	0.49
1:A:776:ASP:OD1	1:A:777:GLY:N	2.46	0.49
1:D:259:ARG:NE	1:D:259:ARG:CA	2.72	0.49
1:D:267:VAL:CG2	1:D:291:PHE:HE2	2.23	0.49
1:D:412:PRO:CG	1:D:413:LEU:N	2.76	0.49
1:D:455:LEU:O	1:D:459:MET:HB2	2.12	0.49
1:D:630:ASN:HD21	1:D:632:ALA:HB2	1.76	0.49
2:E:268:MET:SD	2:E:373:LYS:HD3	2.52	0.49
2:E:67:GLU:N	2:E:67:GLU:CD	2.67	0.49
2:E:67:GLU:CG	2:E:70:LEU:HD11	2.37	0.49
2:F:130:ALA:O	2:F:131:LEU:HD13	2.13	0.49
2:F:208:LEU:CD2	2:F:242:PHE:CE2	2.93	0.49
2:F:293:PHE:C	2:F:295:TRP:H	2.17	0.49
2:F:363:ARG:CZ	2:F:363:ARG:CB	2.90	0.49
1:D:546:ARG:NH2	2:F:408:GLU:OE2	2.45	0.49
2:F:435:MET:O	2:F:435:MET:SD	2.71	0.49
1:A:274:ASP:O	1:A:277:HIS:N	2.45	0.49
1:A:369:PRO:O	1:A:371:LYS:O	2.31	0.49
1:A:214:ALA:CA	1:A:401:TRP:HZ2	2.20	0.49
1:A:421:PRO:CG	1:A:422:VAL:H	2.26	0.49
1:A:438:ASN:C	1:A:440:ASN:N	2.64	0.49
1:A:548:CYS:O	1:A:552:LEU:HG	2.12	0.49
1:A:616:GLU:O	1:A:725:PRO:HD3	2.13	0.49
1:A:618:HIS:HB3	1:A:725:PRO:HB2	1.90	0.49
1:A:730:PRO:HG3	1:A:760:CYS:C	2.33	0.49
2:B:403:PHE:CD1	2:B:415:PRO:HD3	2.48	0.49
2:C:203:ARG:O	2:C:204:LEU:HB3	2.09	0.49
2:C:243:THR:CG2	2:C:247:THR:OG1	2.61	0.49
2:C:306:LEU:HG	2:C:335:VAL:CG2	2.43	0.49
2:C:351:TYR:CA	2:C:355:GLN:HB2	2.42	0.49
2:C:357:THR:HA	2:C:370:LYS:HZ3	1.73	0.49
2:C:389:ARG:NE	2:C:390:GLY:N	2.60	0.49
2:C:416:GLY:HA2	2:C:419:GLU:HB2	1.93	0.49
1:A:233:TYR:O	1:A:234:SER:HB2	2.13	0.49
1:D:1026:ARG:O	1:D:1029:GLN:HB2	2.13	0.49
1:D:1141:VAL:N	1:D:1146:ARG:NH2	2.57	0.49
1:D:1208:ARG:HH22	2:F:285:LYS:HD2	1.72	0.49
1:D:169:TRP:O	1:D:169:TRP:CG	2.66	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:406:VAL:O	1:D:410:GLN:HB2	2.12	0.49
1:D:297:MET:HG2	1:D:411:LEU:CD1	2.43	0.49
1:D:649:TYR:CE2	1:D:652:ILE:HG21	2.47	0.49
1:D:749:PHE:N	1:D:749:PHE:CD1	2.80	0.49
1:D:807:ARG:HH22	1:D:1061:LEU:HD12	1.78	0.49
1:D:279:ARG:CG	1:D:841:LEU:HB3	2.43	0.49
1:D:907:PHE:CD1	1:D:907:PHE:N	2.80	0.49
2:F:115:TRP:CD1	2:F:119:VAL:HG21	2.48	0.49
2:F:341:ASP:OD1	2:F:342:LEU:N	2.41	0.49
2:F:342:LEU:HB3	2:F:344:ARG:HD3	1.92	0.49
1:A:1024:ASP:HA	1:A:1026:ARG:CZ	2.43	0.49
1:A:1064:ILE:HD12	1:A:1064:ILE:N	2.27	0.49
1:A:301:ILE:CG2	1:A:302:SER:H	2.26	0.49
1:A:371:LYS:O	1:A:372:GLU:HG2	2.13	0.49
1:A:384:ASP:O	1:A:387:GLU:HG3	2.12	0.49
1:A:410:GLN:O	1:A:414:PHE:CE2	2.66	0.49
1:A:614:TYR:CE2	1:A:620:TRP:CE3	3.01	0.49
1:A:789:PRO:O	1:A:793:GLU:CG	2.61	0.49
1:A:848:GLY:C	1:A:849:THR:O	2.49	0.49
1:A:879:GLN:CB	1:A:1189:LEU:HD21	2.43	0.49
1:A:436:PRO:HB2	1:A:879:GLN:HE21	1.78	0.49
1:A:914:THR:OG1	1:A:999:ASP:CB	2.60	0.49
2:C:336:LEU:HD23	2:C:337:SER:CA	2.43	0.49
2:C:375:HIS:CD2	2:C:377:CYS:HB2	2.47	0.49
2:C:407:LEU:C	2:C:407:LEU:CD2	2.81	0.49
2:C:70:LEU:C	2:C:70:LEU:HD13	2.33	0.49
1:D:505:LYS:O	1:D:506:GLU:HB3	2.13	0.49
2:B:217:PRO:CB	2:B:232:GLY:H	2.26	0.49
1:D:306:SER:HB2	1:D:1058:PHE:CZ	2.48	0.48
1:D:1069:ILE:CG2	1:D:1070:PRO:N	2.74	0.48
1:D:113:TRP:HZ3	1:D:116:PRO:HD2	1.76	0.48
1:D:192:GLU:OE2	1:D:258:GLN:NE2	2.46	0.48
1:D:281:GLN:HA	1:D:284:ILE:HD11	1.95	0.48
1:D:301:ILE:HG22	1:D:302:SER:N	2.28	0.48
1:D:768:LYS:H	1:D:772:PRO:HD3	1.77	0.48
1:D:920:THR:OG1	1:D:921:LEU:HD12	2.13	0.48
2:E:407:LEU:CA	2:E:411:ILE:O	2.61	0.48
2:F:127:PRO:CD	2:F:210:GLN:CB	2.72	0.48
2:F:88:ARG:CZ	2:F:88:ARG:HB3	2.42	0.48
1:A:1024:ASP:O	1:A:1026:ARG:CD	2.61	0.48
1:A:1164:PHE:O	1:A:1168:LEU:HG	2.12	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:127:LEU:HD23	1:A:128:PRO:CD	2.37	0.48
1:A:175:TRP:HB2	1:A:184:ALA:N	2.27	0.48
1:A:201:VAL:HG22	1:A:208:CYS:C	2.33	0.48
1:A:214:ALA:HA	1:A:401:TRP:CH2	2.47	0.48
1:A:461:LYS:O	1:A:462:SER:C	2.52	0.48
1:A:605:LEU:N	1:A:717:LEU:HD21	2.27	0.48
1:A:656:TYR:CD1	1:A:744:ILE:CD1	2.92	0.48
1:A:744:ILE:HG12	1:A:745:PRO:HD2	1.94	0.48
1:A:754:HIS:CE1	1:A:762:VAL:HA	2.48	0.48
2:B:182:ARG:O	2:B:215:PHE:N	2.46	0.48
2:B:205:PRO:CG	2:B:241:TRP:HE1	2.26	0.48
2:B:387:VAL:CG1	2:B:396:ARG:HB2	2.43	0.48
2:B:81:GLY:CA	2:B:86:LEU:HD11	2.44	0.48
2:C:127:PRO:CD	2:C:209:ALA:C	2.76	0.48
2:C:325:ARG:NH1	2:C:325:ARG:CG	2.70	0.48
2:C:243:THR:N	2:C:334:CYS:H	2.11	0.48
1:A:549:LEU:CD1	2:C:401:GLY:HA3	2.43	0.48
1:A:564:GLN:CA	1:A:564:GLN:NE2	2.75	0.48
2:B:217:PRO:CB	2:B:231:ILE:HB	2.43	0.48
2:C:145:PHE:O	2:C:146:ARG:CG	2.61	0.48
1:D:1212:ILE:HD12	1:D:1213:PRO:HD3	1.94	0.48
1:D:192:GLU:CD	1:D:258:GLN:NE2	2.66	0.48
1:D:274:ASP:O	1:D:278:ILE:HG12	2.13	0.48
1:D:549:LEU:N	1:D:551:LYS:CD	2.75	0.48
1:D:592:LEU:H	1:D:592:LEU:HD12	1.77	0.48
1:D:712:VAL:O	1:D:712:VAL:HG12	2.13	0.48
1:D:849:THR:CG2	1:D:851:THR:HB	2.43	0.48
1:D:887:VAL:CG2	1:D:1146:ARG:NH1	2.76	0.48
1:D:987:ALA:O	1:D:991:GLY:CA	2.61	0.48
2:E:306:LEU:HB2	2:E:335:VAL:CG1	2.43	0.48
2:F:125:VAL:CG2	2:F:208:LEU:CA	2.75	0.48
2:F:264:ARG:HA	2:F:270:PRO:CB	2.43	0.48
2:F:308:ASP:OD2	2:F:334:CYS:HA	2.12	0.48
2:F:395:LEU:CB	2:F:443:VAL:CG2	2.86	0.48
1:A:1016:GLU:OE2	1:A:1022:LEU:HD21	2.14	0.48
1:A:1156:THR:O	1:A:1160:THR:OG1	2.31	0.48
1:A:659:HIS:CD2	1:A:720:THR:HG22	2.48	0.48
1:A:789:PRO:HA	1:A:792:LEU:CD1	2.38	0.48
1:A:853:ARG:NH2	1:A:1105:ALA:HB1	2.28	0.48
1:A:916:PHE:CD1	1:A:919:MET:CB	2.96	0.48
1:A:967:MET:CG	1:A:978:ALA:HB2	2.43	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:234:LYS:CG	2:B:341:ASP:OD2	2.61	0.48
2:C:203:ARG:HG2	2:C:325:ARG:NE	2.28	0.48
2:C:251:TRP:HE3	2:C:254:PHE:HB3	1.78	0.48
2:C:357:THR:O	2:C:359:ASN:N	2.43	0.48
2:C:385:LEU:O	2:C:416:GLY:CA	2.54	0.48
1:D:1003:TRP:CH2	1:D:1101:VAL:HG22	2.48	0.48
1:D:1141:VAL:O	1:D:1142:ARG:C	2.52	0.48
1:D:1155:ILE:O	1:D:1156:THR:C	2.52	0.48
1:D:423:THR:HG22	1:D:424:LEU:HD23	1.95	0.48
1:D:432:VAL:O	1:D:433:SER:O	2.31	0.48
1:D:484:TRP:NE1	1:D:488:LEU:HD11	2.28	0.48
1:D:791:ALA:O	1:D:795:ASN:ND2	2.38	0.48
1:D:993:ARG:HD3	1:D:993:ARG:C	2.34	0.48
2:F:257:ARG:O	2:F:261:GLN:OE1	2.31	0.48
2:F:289:LEU:O	2:F:299:LEU:O	2.32	0.48
2:F:335:VAL:HG13	2:F:335:VAL:O	2.13	0.48
1:A:1027:LYS:O	1:A:1031:GLU:HB2	2.13	0.48
1:A:1198:LYS:HG3	1:A:1204:THR:HG23	1.89	0.48
1:A:1200:PRO:O	1:A:1201:SER:C	2.51	0.48
1:A:347:TRP:O	1:A:348:LEU:O	2.31	0.48
1:A:422:VAL:O	1:A:423:THR:C	2.49	0.48
1:A:449:GLN:CA	1:A:452:TYR:CE1	2.95	0.48
1:A:556:THR:HG23	2:C:452:LEU:CD2	2.43	0.48
1:A:610:PHE:HB3	1:A:671:GLN:OE1	2.13	0.48
1:A:788:GLY:HA2	1:A:791:ALA:CB	2.43	0.48
1:A:866:ARG:HB3	1:A:867:PRO:CD	2.35	0.48
1:A:907:PHE:O	1:A:908:ALA:HB3	2.14	0.48
2:B:259:ARG:HD2	2:B:262:TRP:CG	2.49	0.48
2:B:236:GLU:CG	2:B:341:ASP:CB	2.91	0.48
2:B:234:LYS:HB3	2:B:341:ASP:OD2	2.14	0.48
2:B:391:PRO:HD2	2:B:392:THR:H	1.78	0.48
2:B:400:GLN:NE2	2:B:417:TYR:CB	2.77	0.48
2:B:87:SER:O	2:B:89:ASP:N	2.46	0.48
2:C:342:LEU:O	2:C:342:LEU:HD23	2.13	0.48
2:C:371:VAL:CG2	2:C:433:ASP:O	2.62	0.48
2:C:395:LEU:HD21	2:C:443:VAL:C	2.34	0.48
1:A:560:PRO:CG	2:C:452:LEU:HD21	2.43	0.48
2:C:454:HIS:CA	2:C:466:MET:O	2.61	0.48
1:D:1008:LEU:C	1:D:1011:PRO:CD	2.79	0.48
1:D:110:HIS:O	1:D:112:LEU:HD12	2.13	0.48
1:D:1191:LYS:HE2	1:D:1191:LYS:HA	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:376:LEU:HD12	1:D:379:LYS:NZ	2.28	0.48
1:D:407:PHE:CE1	1:D:411:LEU:HD11	2.49	0.48
2:E:456:ARG:CA	2:E:462:MET:O	2.57	0.48
2:E:471:LEU:HA	2:E:474:PHE:HB3	1.95	0.48
2:E:69:LEU:O	2:E:70:LEU:C	2.50	0.48
2:F:196:CYS:O	2:F:199:LEU:CB	2.61	0.48
2:F:254:PHE:HA	2:F:257:ARG:CD	2.43	0.48
2:F:371:VAL:CG2	2:F:433:ASP:HB3	2.43	0.48
2:F:459:ASP:C	2:F:460:THR:HG22	2.34	0.48
1:D:473:LEU:HA	2:F:461:THR:CG2	2.44	0.48
2:F:471:LEU:CA	2:F:474:PHE:HB3	2.42	0.48
1:A:1029:GLN:H	1:A:1029:GLN:HG3	1.16	0.48
1:A:1196:ASP:OD2	1:A:1197:CYS:N	2.46	0.48
1:A:1219:ASP:C	1:A:1220:ILE:HD12	2.34	0.48
1:A:275:ARG:O	1:A:278:ILE:HB	2.14	0.48
1:A:301:ILE:O	1:A:1024:ASP:OD2	2.32	0.48
1:A:309:ARG:C	1:A:313:ILE:HG23	2.31	0.48
1:A:393:MET:CE	1:A:396:CYS:SG	3.02	0.48
1:A:490:TRP:CA	1:A:580:LEU:HB2	2.41	0.48
1:A:596:MET:O	1:A:597:ARG:C	2.51	0.48
1:A:793:GLU:O	1:A:797:MET:HG3	2.13	0.48
1:A:863:SER:O	1:A:875:LYS:HD2	2.14	0.48
1:A:930:ASP:OD2	1:A:930:ASP:O	2.32	0.48
1:A:960:PRO:HD2	1:A:961:PHE:CD1	2.48	0.48
1:A:992:LEU:HB2	1:A:997:LEU:CB	2.43	0.48
2:B:200:VAL:HB	2:B:203:ARG:HG2	1.95	0.48
2:B:309:HIS:CB	2:B:312:LEU:HB2	2.43	0.48
2:B:418:LEU:O	2:B:419:GLU:C	2.50	0.48
2:B:419:GLU:HA	2:C:203:ARG:HD2	1.95	0.48
2:C:99:PHE:HB2	2:C:103:GLY:HA3	1.95	0.48
1:D:512:LYS:HE2	1:D:514:PRO:HD3	1.95	0.48
1:D:623:LEU:H	1:D:623:LEU:HD23	1.78	0.48
1:A:638:THR:HG23	1:A:642:SER:HG	1.75	0.48
1:A:235:TRP:HD1	1:A:236:THR:HG1	1.61	0.48
1:D:1015:THR:HG21	1:D:1026:ARG:HB3	1.95	0.48
1:D:347:TRP:CZ2	1:D:1030:ARG:NH1	2.82	0.48
1:D:1060:LYS:O	1:D:1063:SER:O	2.31	0.48
1:D:1131:ILE:HD11	1:D:1138:ARG:O	2.13	0.48
1:D:1133:ILE:HB	1:D:1136:GLU:CB	2.12	0.48
1:D:248:GLU:OE1	1:D:250:PRO:HD2	2.12	0.48
1:D:277:HIS:HB3	1:D:291:PHE:CZ	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:547:ALA:CA	1:D:551:LYS:CE	2.88	0.48
1:D:634:LEU:HD22	1:D:635:PRO:CD	2.34	0.48
1:D:776:ASP:CA	1:D:796:LYS:NZ	2.76	0.48
1:D:920:THR:CG2	1:D:922:GLN:CD	2.82	0.48
1:D:946:ALA:CA	1:D:949:PHE:CE2	2.83	0.48
2:E:382:LYS:O	2:E:383:VAL:HG23	2.13	0.48
2:E:386:ASP:O	2:E:387:VAL:HG13	2.14	0.48
2:E:380:PRO:CA	2:E:438:LEU:CD1	2.92	0.48
2:E:448:LEU:O	2:E:449:GLU:C	2.52	0.48
2:F:127:PRO:HG2	2:F:128:VAL:N	2.28	0.48
2:E:101:PRO:HG3	2:F:128:VAL:CG2	2.43	0.48
2:F:241:TRP:CB	2:F:336:LEU:CD1	2.59	0.48
2:F:360:SER:HA	2:F:363:ARG:CG	2.44	0.48
2:F:415:PRO:CB	2:F:417:TYR:CE2	2.91	0.48
2:F:265:LYS:HG3	2:F:485:VAL:HG11	1.95	0.48
2:F:86:LEU:O	2:F:91:LEU:HB2	2.13	0.48
1:A:1026:ARG:HG2	1:A:1029:GLN:CD	2.33	0.48
1:A:278:ILE:O	1:A:278:ILE:HG22	2.14	0.48
1:A:389:PHE:CG	1:A:390:GLN:N	2.82	0.48
1:A:421:PRO:HG2	1:A:422:VAL:CG2	2.43	0.48
1:A:570:PRO:HG2	1:A:571:GLY:H	1.78	0.48
2:C:248:SER:OG	2:C:249:ASN:N	2.46	0.48
2:C:306:LEU:HG	2:C:335:VAL:HG21	1.94	0.48
2:C:346:MET:O	2:C:350:LEU:CD1	2.62	0.48
2:C:399:CYS:SG	2:C:417:TYR:CG	3.07	0.48
1:A:549:LEU:HD11	2:C:401:GLY:HA3	1.94	0.48
2:C:440:THR:O	2:C:457:SER:CB	2.61	0.48
2:E:267:ALA:O	2:E:270:PRO:HD3	2.13	0.48
2:B:277:ASP:HA	2:B:287:ASN:OD1	2.12	0.48
2:B:134:LYS:NZ	2:B:137:PRO:HD2	2.29	0.48
1:D:1193:VAL:CG2	1:D:1215:GLY:H	2.26	0.48
1:D:253:ALA:O	1:D:254:SER:CB	2.60	0.48
1:D:375:GLU:O	1:D:379:LYS:CG	2.55	0.48
1:D:410:GLN:O	1:D:414:PHE:HD2	1.97	0.48
1:D:439:GLN:C	1:D:441:TRP:N	2.66	0.48
1:D:604:ALA:CA	1:D:613:HIS:HE1	2.11	0.48
1:D:829:PRO:O	1:D:829:PRO:HD2	2.14	0.48
1:D:869:ARG:CA	1:D:1200:PRO:CB	2.91	0.48
2:E:191:GLU:HG3	2:E:192:HIS:CD2	2.49	0.48
2:E:346:MET:SD	2:E:346:MET:C	2.91	0.48
2:E:347:LEU:C	2:E:350:LEU:HB3	2.34	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:407:LEU:C	2:E:409:ASN:N	2.67	0.48
2:E:429:TYR:C	2:E:431:LYS:N	2.64	0.48
2:F:126:PHE:C	2:F:128:VAL:N	2.59	0.48
2:F:266:PHE:HB2	2:F:349:TYR:OH	2.14	0.48
1:D:1208:ARG:NH1	2:F:285:LYS:CD	2.57	0.48
2:F:414:TRP:N	2:F:415:PRO:HD2	2.28	0.48
1:A:996:ARG:CZ	1:A:1003:TRP:CB	2.87	0.48
1:A:1006:ARG:HA	1:A:1006:ARG:HH11	1.62	0.48
1:A:214:ALA:N	1:A:401:TRP:CZ2	2.82	0.48
1:A:176:THR:O	1:A:220:TRP:CZ3	2.66	0.48
1:A:267:VAL:HG12	1:A:269:HIS:CE1	2.49	0.48
1:A:269:HIS:O	1:A:294:THR:N	2.45	0.48
1:A:274:ASP:C	1:A:276:ALA:H	2.12	0.48
1:A:400:VAL:C	1:A:403:THR:HG23	2.33	0.48
1:A:556:THR:O	1:A:559:LEU:HB3	2.13	0.48
1:A:914:THR:N	1:A:918:TRP:HB2	2.29	0.48
2:B:116:THR:O	2:B:121:PHE:CD2	2.63	0.48
2:B:80:SER:N	2:B:100:GLY:HA3	2.29	0.48
2:B:90:SER:CA	2:B:93:SER:OG	2.59	0.48
2:C:127:PRO:CD	2:C:210:GLN:CB	2.78	0.48
2:C:131:LEU:O	2:C:182:ARG:HD3	2.13	0.48
1:A:783:PRO:HG2	2:C:363:ARG:HD2	1.96	0.48
2:C:375:HIS:HB3	2:C:378:LEU:HD22	1.94	0.48
2:C:90:SER:HA	2:C:93:SER:OG	2.13	0.48
1:A:640:LEU:N	1:A:640:LEU:CD2	2.64	0.48
2:F:146:ARG:HB2	2:F:228:VAL:HG11	1.95	0.48
1:D:860:LEU:CD2	1:D:1133:ILE:HD12	2.44	0.48
1:D:1133:ILE:HG22	1:D:1134:HIS:HD2	1.79	0.48
1:D:1159:LEU:O	1:D:1162:CYS:HB2	2.14	0.48
1:D:1222:GLN:O	1:D:1226:LEU:CD2	2.55	0.48
1:D:168:ALA:O	1:D:169:TRP:C	2.52	0.48
1:D:316:LYS:HZ1	1:D:348:LEU:HD23	1.76	0.48
1:D:779:LEU:HD22	1:D:780:GLN:HA	1.94	0.48
1:D:829:PRO:O	1:D:830:ASP:OD2	2.32	0.48
1:D:868:ASP:OD1	1:D:1199:THR:HB	2.14	0.48
1:D:923:GLY:O	1:D:931:LEU:HD11	2.12	0.48
2:E:120:VAL:HG22	2:F:407:LEU:CD2	2.43	0.48
2:E:243:THR:CG2	2:E:248:SER:HA	2.44	0.48
2:E:247:THR:HG23	2:E:250:GLN:HG3	1.96	0.48
2:E:294:PRO:HD2	2:E:351:TYR:CD2	2.49	0.48
2:F:190:LEU:O	2:F:193:TYR:CB	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:347:LEU:N	2:F:347:LEU:CD1	2.77	0.48
2:E:203:ARG:CZ	2:F:418:LEU:HD13	2.43	0.48
1:A:106:HIS:O	1:A:113:TRP:NE1	2.32	0.48
1:A:1157:ASN:HB3	1:A:1161:ARG:NH1	2.28	0.48
1:A:128:PRO:N	1:A:129:PRO:CD	2.76	0.48
1:A:165:LYS:CG	1:A:178:TYR:HB2	2.44	0.48
1:A:358:LEU:N	1:A:358:LEU:HD23	2.28	0.48
1:A:414:PHE:CA	1:A:417:ARG:HB2	2.44	0.48
1:A:954:ILE:HG22	1:A:955:TYR:N	2.28	0.48
2:B:110:LEU:O	2:B:111:ALA:C	2.52	0.48
2:B:245:PRO:O	2:B:248:SER:HB3	2.14	0.48
2:B:336:LEU:HG	2:B:337:SER:H	1.79	0.48
2:C:193:TYR:OH	2:C:333:PRO:HG3	2.13	0.48
2:C:239:LEU:CA	2:C:338:VAL:HG12	2.41	0.48
2:E:389:ARG:NH2	2:E:391:PRO:C	2.67	0.48
1:D:102:ARG:O	1:D:106:HIS:HD2	1.96	0.48
1:D:1005:VAL:HG23	1:D:1006:ARG:N	2.28	0.48
1:D:1058:PHE:CD2	1:D:1059:ASN:N	2.82	0.48
1:D:886:LEU:HA	1:D:1146:ARG:NH2	2.28	0.48
1:D:293:ASP:C	1:D:296:SER:HB3	2.34	0.48
1:D:297:MET:HG2	1:D:411:LEU:HD13	1.96	0.48
1:D:575:LYS:HG2	1:D:581:ASP:O	2.14	0.48
1:D:624:VAL:C	1:D:745:PRO:O	2.52	0.48
1:D:630:ASN:C	1:D:630:ASN:ND2	2.67	0.48
1:D:633:LYS:HD3	1:D:638:THR:HG21	1.92	0.48
2:E:105:GLU:O	2:E:109:ASN:OD1	2.31	0.48
2:E:120:VAL:HG12	2:E:121:PHE:HD1	1.79	0.48
2:E:332:VAL:N	2:E:333:PRO:CD	2.76	0.48
2:F:239:LEU:HB3	2:F:338:VAL:HG13	1.96	0.48
2:F:245:PRO:C	2:F:248:SER:HB2	2.32	0.48
2:F:353:SER:OG	2:F:354:PHE:N	2.47	0.48
2:F:351:TYR:O	2:F:355:GLN:HB2	2.10	0.48
2:F:375:HIS:CD2	2:F:377:CYS:HB2	2.49	0.48
2:E:203:ARG:NH2	2:F:418:LEU:HD11	2.29	0.48
2:F:416:GLY:HA2	2:F:419:GLU:HB2	1.95	0.48
2:F:88:ARG:CZ	2:F:88:ARG:CB	2.91	0.48
1:A:1144:GLU:HG3	1:A:1145:ASP:OD1	2.14	0.48
1:A:1224:ILE:HD12	1:A:1224:ILE:C	2.33	0.48
1:A:287:SER:O	1:A:288:ARG:NE	2.47	0.48
1:A:377:PHE:O	1:A:378:VAL:CB	2.61	0.48
1:A:423:THR:O	1:A:427:MET:HG2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:547:ALA:HA	1:A:550:GLN:OE1	2.14	0.48
1:A:626:GLY:O	1:A:627:ARG:HB3	2.13	0.48
1:A:727:ASP:O	1:A:728:THR:OG1	2.31	0.48
1:A:795:ASN:ND2	1:A:796:LYS:HG3	2.28	0.48
1:A:822:PRO:O	1:A:823:ARG:HB3	2.14	0.48
1:A:79:LEU:HB3	1:A:83:LEU:HD21	1.96	0.48
1:A:867:PRO:HG2	1:A:1067:SER:OG	2.12	0.48
2:B:285:LYS:O	2:B:305:ASN:HB2	2.13	0.48
2:C:246:ARG:C	2:C:248:SER:N	2.59	0.48
2:C:190:LEU:CB	2:C:314:MET:HE1	2.44	0.48
2:C:327:GLY:O	2:C:328:ARG:HB3	2.14	0.48
2:C:397:GLN:O	2:C:399:CYS:N	2.46	0.48
2:C:419:GLU:CG	2:C:432:TYR:OH	2.61	0.48
2:C:435:MET:CE	2:C:435:MET:CA	2.84	0.48
2:C:70:LEU:O	2:C:73:CYS:HB2	2.14	0.48
1:D:520:ALA:N	1:D:521:PRO:HD3	2.28	0.48
2:E:194:VAL:CG1	2:E:195:ASN:H	2.26	0.48
1:D:1207:GLU:C	1:D:1210:TYR:CD2	2.86	0.48
1:D:256:PRO:CG	1:D:281:GLN:HA	2.44	0.48
1:D:454:GLU:CA	1:D:457:ARG:HG3	2.43	0.48
1:D:895:GLU:CD	1:D:895:GLU:C	2.73	0.48
1:D:900:ALA:O	1:D:915:ALA:CB	2.62	0.48
2:E:441:VAL:HG22	2:E:455:LEU:HB3	1.96	0.48
2:F:243:THR:CA	2:F:334:CYS:HB2	2.44	0.48
2:F:290:TYR:CD1	2:F:299:LEU:HD22	2.49	0.48
2:F:372:LEU:HD11	2:F:436:SER:OG	2.11	0.48
2:F:387:VAL:HG11	2:F:396:ARG:HA	1.96	0.48
2:F:402:LEU:C	2:F:406:LEU:HD22	2.35	0.48
1:A:996:ARG:NH1	1:A:1000:GLU:O	2.25	0.48
1:A:1169:GLY:C	1:A:1170:LEU:HD12	2.34	0.48
1:A:304:LEU:HD23	1:A:1025:LEU:HD13	1.96	0.48
1:A:595:GLN:O	1:A:597:ARG:N	2.47	0.48
1:A:755:LYS:O	1:A:758:ASN:HA	2.14	0.48
1:A:879:GLN:HB2	1:A:1189:LEU:HD21	1.96	0.48
2:C:303:LEU:HD12	2:C:339:ASN:OD1	2.14	0.48
2:C:399:CYS:SG	2:C:417:TYR:CD2	3.07	0.48
2:C:456:ARG:CB	2:C:463:LYS:HB3	2.44	0.48
2:C:84:GLN:C	2:C:86:LEU:N	2.67	0.48
1:D:671:GLN:CA	1:D:671:GLN:NE2	2.75	0.48
2:B:453:ILE:H	2:B:453:ILE:CD1	2.24	0.48
2:F:217:PRO:CB	2:F:219:PHE:CE1	2.90	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:807:ARG:C	1:A:810:SER:OG	2.53	0.48
1:D:1002:GLU:HG2	1:D:1003:TRP:H	1.79	0.48
1:D:1198:LYS:CA	1:D:1204:THR:CB	2.85	0.48
1:D:195:LEU:H	1:D:265:LEU:CG	2.27	0.48
1:D:195:LEU:HD21	1:D:217:PRO:HD2	1.96	0.48
1:D:242:ALA:CB	1:D:245:ILE:CG2	2.92	0.48
1:D:282:TYR:CE2	1:D:432:VAL:CA	2.93	0.48
1:D:282:TYR:O	1:D:285:GLN:HG2	2.12	0.48
1:D:282:TYR:CD2	1:D:432:VAL:O	2.66	0.48
1:D:450:GLY:O	1:D:454:GLU:HG3	2.14	0.48
1:D:456:GLN:O	1:D:460:LYS:CD	2.62	0.48
1:D:484:TRP:NE1	1:D:488:LEU:CD1	2.77	0.48
1:D:607:TRP:O	1:D:608:ASP:CG	2.51	0.48
1:D:741:ASP:OD1	1:D:741:ASP:N	2.47	0.48
1:D:743:ASP:HB2	1:D:744:ILE:CD1	2.43	0.48
1:D:73:PRO:HG2	1:D:90:GLN:HG2	1.90	0.48
2:E:131:LEU:HD23	2:E:132:HIS:H	1.79	0.48
2:E:297:LYS:O	2:E:298:GLU:CB	2.62	0.48
2:F:183:GLU:HA	2:F:183:GLU:OE2	2.14	0.48
2:F:243:THR:CB	2:F:334:CYS:HB2	2.44	0.48
2:F:368:HIS:CG	2:F:368:HIS:O	2.65	0.48
2:F:86:LEU:O	2:F:87:SER:C	2.51	0.48
1:A:1054:GLU:O	1:A:1058:PHE:CD2	2.65	0.48
1:A:121:PRO:O	1:A:1175:GLN:NE2	2.35	0.48
1:A:210:THR:HB	1:A:224:CYS:SG	2.53	0.48
1:A:301:ILE:HD11	1:A:414:PHE:HB3	1.96	0.48
1:A:346:ASP:HB3	1:A:1040:LYS:HZ2	1.67	0.48
1:A:445:LEU:C	1:A:448:ALA:HB3	2.33	0.48
1:A:483:PRO:HB3	1:A:712:VAL:CG1	2.42	0.48
1:A:605:LEU:O	1:A:781:ALA:HB3	2.14	0.48
1:A:658:LYS:O	1:A:660:CYS:N	2.47	0.48
1:A:624:VAL:HG21	1:A:747:CYS:SG	2.54	0.48
1:A:764:SER:O	1:A:767:ALA:HB2	2.13	0.48
1:A:916:PHE:O	1:A:920:THR:CA	2.62	0.48
2:B:213:VAL:HA	2:B:235:THR:CA	2.43	0.48
2:B:313:HIS:O	2:B:316:PRO:CD	2.55	0.48
2:B:315:TYR:CB	2:B:322:LEU:HD11	2.44	0.48
2:B:437:ILE:O	2:B:458:ARG:CZ	2.62	0.48
2:B:74:GLN:HE21	2:B:77:HIS:CD2	2.32	0.48
2:C:306:LEU:H	2:C:335:VAL:HG21	1.77	0.48
2:C:74:GLN:CG	2:C:75:ARG:N	2.76	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:78:PHE:CD2	2:C:79:LEU:CD2	2.97	0.48
2:B:450:ASN:ND2	2:B:452:LEU:HD23	2.24	0.48
1:D:101:ARG:O	1:D:102:ARG:C	2.52	0.48
1:D:308:GLN:NE2	1:D:1027:LYS:HZ1	2.10	0.47
1:D:1028:VAL:CG2	1:D:1096:ARG:CZ	2.89	0.47
1:D:1111:LEU:CD1	1:D:1111:LEU:H	2.27	0.47
1:D:301:ILE:CG2	1:D:302:SER:N	2.77	0.47
1:D:904:ASP:OD1	1:D:904:ASP:N	2.34	0.47
2:E:94:GLY:CA	2:E:234:LYS:HE2	2.43	0.47
2:E:440:THR:HG22	2:E:441:VAL:N	2.29	0.47
2:E:448:LEU:CD2	2:E:448:LEU:C	2.82	0.47
2:E:86:LEU:CG	2:E:97:PRO:HD2	2.43	0.47
2:F:236:GLU:HB3	2:F:341:ASP:HB3	1.95	0.47
2:F:331:VAL:HG13	2:F:332:VAL:N	2.28	0.47
2:F:454:HIS:CA	2:F:466:MET:O	2.62	0.47
1:A:1006:ARG:O	1:A:1010:LEU:HB2	2.14	0.47
1:A:313:ILE:HG13	1:A:314:ALA:H	1.79	0.47
1:A:463:LEU:HD22	1:A:464:MET:N	2.29	0.47
1:A:620:TRP:C	1:A:748:TRP:HB2	2.35	0.47
1:A:756:ASP:OD1	1:A:756:ASP:N	2.47	0.47
1:A:795:ASN:HD21	1:A:796:LYS:HG3	1.78	0.47
1:A:900:ALA:CA	1:A:915:ALA:CB	2.80	0.47
1:A:972:ARG:C	1:A:974:THR:N	2.66	0.47
2:B:196:CYS:O	2:B:197:LEU:C	2.53	0.47
2:B:261:GLN:O	2:B:261:GLN:CG	2.61	0.47
2:B:383:VAL:CG2	2:B:384:ALA:N	2.76	0.47
2:B:435:MET:O	2:B:436:SER:C	2.51	0.47
2:C:384:ALA:HB2	2:C:437:ILE:CD1	2.38	0.47
1:D:497:GLN:OE1	1:D:516:GLU:O	2.32	0.47
1:A:526:ASP:OD2	1:A:526:ASP:C	2.52	0.47
2:E:270:PRO:CA	2:E:273:PHE:CD2	2.88	0.47
2:B:82:SER:H	2:B:85:GLN:HG3	1.78	0.47
2:F:90:SER:O	2:F:93:SER:OG	2.27	0.47
1:D:1008:LEU:CD2	1:D:1008:LEU:C	2.82	0.47
1:D:1044:VAL:O	1:D:1047:ARG:CB	2.62	0.47
1:D:1122:GLU:O	1:D:1123:PHE:CB	2.63	0.47
1:D:78:MET:HB2	1:D:1175:GLN:HB3	1.96	0.47
1:D:877:MET:HE1	1:D:1203:PRO:HA	1.96	0.47
1:D:1221:TYR:CA	1:D:1224:ILE:CG2	2.73	0.47
1:D:257:THR:O	1:D:258:GLN:C	2.53	0.47
1:D:293:ASP:HB3	1:D:296:SER:CB	2.42	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:357:ALA:O	1:D:359:VAL:N	2.47	0.47
1:D:389:PHE:HD1	1:D:389:PHE:C	2.16	0.47
1:D:420:HIS:ND1	1:D:421:PRO:CD	2.47	0.47
1:D:451:THR:O	1:D:454:GLU:HB2	2.13	0.47
1:D:459:MET:CB	1:D:460:LYS:NZ	2.77	0.47
1:D:485:LEU:HD12	1:D:489:GLU:CD	2.34	0.47
1:D:486:TRP:CZ3	1:D:598:VAL:CG2	2.97	0.47
1:D:607:TRP:HA	1:D:607:TRP:HE3	1.80	0.47
1:D:771:LEU:O	1:D:775:GLU:N	2.47	0.47
1:D:774:MET:HB3	1:D:779:LEU:CD1	2.44	0.47
1:D:992:LEU:O	1:D:996:ARG:CZ	2.62	0.47
2:E:431:LYS:CG	2:E:432:TYR:H	2.27	0.47
2:F:193:TYR:CZ	2:F:333:PRO:HG3	2.49	0.47
2:F:311:LEU:CD2	2:F:311:LEU:H	2.26	0.47
2:F:99:PHE:N	2:F:99:PHE:CD2	2.82	0.47
1:A:1014:ARG:HB2	1:A:1026:ARG:NE	2.27	0.47
1:A:1136:GLU:N	1:A:1136:GLU:OE2	2.47	0.47
1:A:81:ARG:NH2	1:A:126:ARG:HA	2.29	0.47
1:A:226:GLN:CD	1:A:226:GLN:C	2.73	0.47
1:A:302:SER:HB2	1:A:351:SER:HB3	1.96	0.47
1:A:389:PHE:CB	1:A:393:MET:HG2	2.44	0.47
1:A:406:VAL:HG22	1:A:407:PHE:CD2	2.48	0.47
1:A:601:LYS:CE	1:A:616:GLU:OE2	2.61	0.47
1:A:869:ARG:N	1:A:872:SER:HB2	2.28	0.47
2:B:72:ILE:C	2:B:75:ARG:HG3	2.33	0.47
2:C:375:HIS:HD2	2:C:377:CYS:HB2	1.78	0.47
1:A:484:TRP:HH2	2:C:461:THR:O	1.97	0.47
1:A:503:VAL:HG21	1:A:521:PRO:C	2.35	0.47
2:E:389:ARG:HH21	2:E:390:GLY:C	2.17	0.47
1:A:774:MET:CA	1:A:777:GLY:HA3	2.32	0.47
2:F:219:PHE:O	2:F:220:ASP:O	2.31	0.47
1:D:1026:ARG:O	1:D:1026:ARG:HG2	2.14	0.47
1:D:1040:LYS:HE3	1:D:1040:LYS:HA	1.97	0.47
1:D:172:ALA:O	1:D:173:GLU:CB	2.62	0.47
1:D:275:ARG:HH11	1:D:843:GLN:CG	2.20	0.47
1:D:384:ASP:N	1:D:384:ASP:OD1	2.42	0.47
1:D:459:MET:CA	1:D:460:LYS:HZ1	2.23	0.47
1:D:550:GLN:H	1:D:551:LYS:CE	2.21	0.47
1:D:633:LYS:HB3	1:D:638:THR:CB	2.44	0.47
1:D:656:TYR:HD1	1:D:744:ILE:CD1	2.24	0.47
1:D:247:LEU:HA	1:D:815:TRP:HB2	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:900:ALA:O	1:D:915:ALA:HB2	2.14	0.47
1:D:923:GLY:HA2	1:D:931:LEU:CD1	2.44	0.47
2:E:108:LYS:HG3	2:E:109:ASN:N	2.29	0.47
2:E:289:LEU:O	2:E:290:TYR:CG	2.68	0.47
2:E:425:LEU:C	2:E:428:LEU:HD11	2.34	0.47
2:F:200:VAL:CG1	2:F:204:LEU:HD22	2.43	0.47
2:F:300:ILE:CG2	2:F:301:GLU:HG3	2.31	0.47
2:F:82:SER:O	2:F:83:LYS:HB2	2.12	0.47
1:A:1016:GLU:N	1:A:1016:GLU:OE2	2.46	0.47
1:A:373:PRO:HD2	1:A:374:ARG:HG2	1.94	0.47
1:A:372:GLU:C	1:A:374:ARG:O	2.52	0.47
1:A:374:ARG:C	1:A:376:LEU:H	2.17	0.47
1:A:383:LYS:O	1:A:386:ARG:NH1	2.48	0.47
1:A:462:SER:O	1:A:463:LEU:C	2.52	0.47
1:A:587:PRO:O	1:A:588:GLY:O	2.31	0.47
1:A:596:MET:HB3	1:A:596:MET:HE2	1.59	0.47
1:A:670:PRO:HA	1:A:673:ALA:CB	2.44	0.47
1:A:753:PRO:HD2	1:A:770:PHE:CD2	2.49	0.47
1:A:275:ARG:HG2	1:A:843:GLN:HB2	1.95	0.47
1:A:958:GLY:HA2	1:A:986:TYR:CZ	2.49	0.47
2:B:305:ASN:C	2:B:307:GLY:H	2.17	0.47
2:B:386:ASP:OD1	2:B:387:VAL:HG22	2.14	0.47
2:C:239:LEU:C	2:C:338:VAL:CG1	2.82	0.47
2:C:364:LYS:N	2:C:364:LYS:CD	2.76	0.47
2:C:388:GLY:O	2:C:444:THR:CA	2.54	0.47
1:A:503:VAL:CG2	1:A:521:PRO:CA	2.82	0.47
2:E:186:LEU:O	2:E:187:HIS:C	2.51	0.47
2:B:469:SER:OG	2:B:470:LYS:N	2.46	0.47
1:D:97:GLU:C	1:D:101:ARG:HG3	2.35	0.47
1:D:162:LEU:HD12	1:D:162:LEU:C	2.34	0.47
1:A:141:LEU:O	1:A:144:GLN:CB	2.63	0.47
1:D:306:SER:HB2	1:D:1058:PHE:CE2	2.49	0.47
1:D:197:PHE:CD1	1:D:213:VAL:O	2.67	0.47
1:D:273:PHE:O	1:D:276:ALA:HB3	2.14	0.47
1:D:312:TRP:HB3	1:D:316:LYS:CD	2.44	0.47
1:D:348:LEU:CD2	1:D:350:ILE:CG1	2.85	0.47
1:D:891:VAL:CG2	1:D:891:VAL:O	2.62	0.47
2:E:193:TYR:O	2:E:197:LEU:HB2	2.14	0.47
2:E:322:LEU:HB3	2:E:332:VAL:HG13	1.96	0.47
2:E:382:LYS:HE2	2:E:412:SER:C	2.23	0.47
2:E:468:ILE:O	2:E:469:SER:OG	2.25	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1209:ARG:HD2	2:F:249:ASN:CA	2.45	0.47
2:F:344:ARG:HG2	2:F:347:LEU:HD22	1.96	0.47
2:F:351:TYR:HD1	2:F:355:GLN:HA	1.78	0.47
1:D:472:GLN:CB	2:F:460:THR:O	2.56	0.47
1:A:1137:VAL:HG12	1:A:1139:TYR:CZ	2.48	0.47
1:A:885:THR:CA	1:A:1146:ARG:HE	2.23	0.47
1:A:420:HIS:O	1:A:424:LEU:HD11	2.14	0.47
1:A:441:TRP:CE3	1:A:445:LEU:CD1	2.96	0.47
1:A:468:ASN:O	1:A:469:ASP:C	2.52	0.47
1:A:664:GLY:HA2	1:A:667:GLN:NE2	2.28	0.47
1:A:792:LEU:C	1:A:796:LYS:HD3	2.35	0.47
1:A:914:THR:HA	1:A:918:TRP:CB	2.29	0.47
1:A:951:TYR:O	1:A:955:TYR:CD1	2.67	0.47
2:B:336:LEU:HD23	2:B:337:SER:N	2.29	0.47
2:C:193:TYR:CD1	2:C:242:PHE:CE2	3.02	0.47
2:C:291:TYR:CG	2:C:348:ALA:HB1	2.50	0.47
2:C:314:MET:O	2:C:316:PRO:HD3	2.14	0.47
2:C:342:LEU:HD23	2:C:342:LEU:C	2.35	0.47
2:C:371:VAL:HG13	2:C:372:LEU:O	2.14	0.47
2:C:405:GLU:OE1	2:C:405:GLU:C	2.53	0.47
2:C:428:LEU:HB3	2:C:432:TYR:CE1	2.50	0.47
2:C:455:LEU:CB	2:C:456:ARG:HE	2.26	0.47
2:C:456:ARG:HG3	2:C:463:LYS:HB3	1.96	0.47
1:D:164:PRO:CD	1:D:165:LYS:N	2.73	0.47
2:F:219:PHE:HE2	2:F:228:VAL:C	2.17	0.47
2:F:219:PHE:O	2:F:220:ASP:C	2.50	0.47
2:F:365:LYS:HB3	2:F:367:LEU:HD21	1.96	0.47
2:C:59:ALA:O	2:C:62:SER:OG	2.28	0.47
1:D:1041:TRP:CD1	1:D:1044:VAL:HG11	2.50	0.47
1:D:1132:SER:O	1:D:1132:SER:OG	2.33	0.47
1:D:1208:ARG:CG	1:D:1209:ARG:N	2.77	0.47
1:D:127:LEU:HD23	1:D:128:PRO:N	2.29	0.47
1:D:224:CYS:SG	1:D:228:LEU:HD23	2.54	0.47
1:D:556:THR:HG23	1:D:559:LEU:HD23	1.95	0.47
1:D:724:GLY:HA2	1:D:725:PRO:HA	1.64	0.47
1:D:801:TRP:CH2	1:D:805:HIS:CD2	3.02	0.47
1:D:846:THR:O	1:D:847:ALA:C	2.52	0.47
1:D:115:GLN:NE2	1:D:924:ARG:NH1	2.61	0.47
2:E:131:LEU:CD2	2:E:132:HIS:H	2.26	0.47
2:F:199:LEU:HD12	2:F:200:VAL:N	2.30	0.47
2:F:204:LEU:O	2:F:205:PRO:C	2.53	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:350:LEU:H	2:F:350:LEU:CD1	2.16	0.47
2:F:458:ARG:HH11	2:F:458:ARG:CG	2.27	0.47
1:A:1154:GLN:HE21	1:A:1220:ILE:HG13	1.77	0.47
1:A:420:HIS:O	1:A:424:LEU:CD1	2.62	0.47
1:A:454:GLU:C	1:A:458:GLU:HG3	2.34	0.47
1:A:784:GLY:CA	2:C:363:ARG:HD3	2.44	0.47
1:A:801:TRP:O	1:A:802:ARG:C	2.50	0.47
1:A:907:PHE:CZ	1:A:1001:GLY:HA3	2.50	0.47
2:B:289:LEU:HB2	2:B:301:GLU:H	1.80	0.47
2:B:338:VAL:O	2:B:338:VAL:HG12	2.14	0.47
2:B:379:ALA:HB3	2:B:438:LEU:HD11	1.96	0.47
2:C:120:VAL:CG2	2:C:121:PHE:H	2.28	0.47
2:C:290:TYR:HD1	2:C:299:LEU:HA	1.80	0.47
2:C:403:PHE:CD2	2:C:407:LEU:HB2	2.50	0.47
2:C:403:PHE:O	2:C:406:LEU:HB2	2.14	0.47
1:D:525:MET:HG3	1:D:526:ASP:OD1	2.15	0.47
1:A:1056:GLU:HG3	1:A:1059:ASN:HD21	1.79	0.47
2:B:475:LEU:O	2:B:476:ILE:C	2.51	0.47
1:A:741:ASP:OD2	1:A:742:VAL:HG23	2.14	0.47
2:E:272:ASN:HB3	2:E:292:ASN:HB2	1.97	0.47
1:D:1024:ASP:C	1:D:1026:ARG:H	2.18	0.47
1:D:253:ALA:O	1:D:254:SER:HB3	2.14	0.47
1:D:361:ARG:HH22	1:D:362:LEU:HD21	1.80	0.47
1:D:458:GLU:N	1:D:460:LYS:HE3	2.29	0.47
1:D:888:GLY:O	1:D:889:ALA:HB2	2.13	0.47
2:F:110:LEU:O	2:F:111:ALA:C	2.51	0.47
2:F:207:GLY:CA	2:F:239:LEU:HD11	2.43	0.47
2:F:305:ASN:C	2:F:307:GLY:N	2.68	0.47
1:A:996:ARG:HH12	1:A:1004:LEU:N	2.12	0.47
1:A:1131:ILE:CG1	1:A:1132:SER:N	2.77	0.47
1:A:81:ARG:CD	1:A:125:LEU:O	2.57	0.47
1:A:247:LEU:HD12	1:A:251:THR:HG23	1.93	0.47
1:A:258:GLN:CD	1:A:258:GLN:N	2.67	0.47
1:A:583:PRO:C	1:A:585:TRP:H	2.14	0.47
1:A:821:LEU:CD1	1:A:825:VAL:CG1	2.92	0.47
2:C:99:PHE:CB	2:C:103:GLY:HA3	2.44	0.47
2:C:206:TYR:N	2:C:206:TYR:CD2	2.52	0.47
2:C:236:GLU:CA	2:C:341:ASP:CG	2.82	0.47
2:C:389:ARG:HG3	2:C:390:GLY:N	2.27	0.47
2:C:385:LEU:CD2	2:C:402:LEU:HD23	2.38	0.47
2:C:431:LYS:HZ3	2:C:435:MET:HG2	1.72	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:506:GLU:HG3	1:A:507:PRO:N	2.29	0.47
1:D:1002:GLU:CG	1:D:1003:TRP:H	2.28	0.47
1:D:1107:ASP:C	1:D:1111:LEU:HD13	2.35	0.47
1:D:121:PRO:O	1:D:122:ASP:OD2	2.32	0.47
1:D:255:SER:HB3	1:D:256:PRO:CA	2.43	0.47
1:D:195:LEU:O	1:D:267:VAL:CG1	2.63	0.47
1:D:484:TRP:HE1	1:D:488:LEU:CD1	2.28	0.47
1:D:550:GLN:CG	1:D:551:LYS:HE3	2.45	0.47
1:D:808:ILE:HG22	1:D:811:GLN:OE1	2.14	0.47
1:D:821:LEU:HB3	1:D:822:PRO:HA	1.95	0.47
1:D:851:THR:OG1	1:D:1102:GLN:HG2	2.15	0.47
1:D:848:GLY:HA3	1:D:855:VAL:HG21	1.97	0.47
1:D:921:LEU:CD1	1:D:921:LEU:H	2.27	0.47
1:D:945:HIS:HE1	1:D:969:PHE:HB2	1.79	0.47
2:E:129:ASP:OD2	2:F:97:PRO:O	2.33	0.47
2:E:78:PHE:CZ	2:E:350:LEU:HD12	2.49	0.47
2:F:356:LEU:HB3	2:F:359:ASN:HB2	1.96	0.47
2:F:432:TYR:O	2:F:435:MET:N	2.45	0.47
2:F:459:ASP:OD1	2:F:461:THR:N	2.40	0.47
1:D:1109:LEU:HD21	1:D:1137:VAL:CG2	2.44	0.47
1:D:1136:GLU:HG3	1:D:1138:ARG:NH1	2.29	0.47
1:D:1203:PRO:O	1:D:1204:THR:C	2.52	0.47
1:D:1204:THR:O	1:D:1205:GLY:O	2.32	0.47
1:D:188:ALA:O	1:D:189:ILE:C	2.52	0.47
1:D:733:HIS:CE1	1:D:750:PHE:CE1	3.02	0.47
1:D:787:SER:O	1:D:789:PRO:CD	2.63	0.47
1:D:807:ARG:NH1	1:D:1062:GLU:HG2	2.30	0.47
1:D:865:ALA:CA	1:D:872:SER:O	2.63	0.47
1:D:967:MET:SD	1:D:975:GLN:HA	2.55	0.47
1:D:987:ALA:HA	1:D:991:GLY:HA2	1.96	0.47
2:E:246:ARG:HD2	2:E:326:ASP:OD1	2.13	0.47
2:E:193:TYR:HE2	2:E:322:LEU:CB	2.28	0.47
2:F:208:LEU:HD22	2:F:242:PHE:CD2	2.48	0.47
2:F:311:LEU:O	2:F:312:LEU:C	2.52	0.47
2:F:456:ARG:N	2:F:456:ARG:CD	2.78	0.47
2:F:458:ARG:CD	2:F:459:ASP:N	2.78	0.47
2:F:468:ILE:O	2:F:471:LEU:HD13	2.15	0.47
1:A:1011:PRO:CD	1:A:1012:VAL:H	2.24	0.47
1:A:1141:VAL:HB	1:A:1146:ARG:HD3	1.94	0.47
1:A:1196:ASP:C	1:A:1196:ASP:OD2	2.52	0.47
1:A:1223:ILE:HA	1:A:1226:LEU:CD2	2.43	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:157:LEU:HG	1:A:194:ALA:CB	2.44	0.47
1:A:386:ARG:HD3	1:A:387:GLU:N	2.27	0.47
1:A:213:VAL:HG11	1:A:401:TRP:NE1	2.26	0.47
1:A:515:ILE:CG1	1:A:565:HIS:O	2.56	0.47
1:A:655:LEU:CD1	1:A:719:LEU:O	2.62	0.47
1:A:727:ASP:N	1:A:727:ASP:OD2	2.33	0.47
1:A:451:THR:CB	1:A:869:ARG:HH22	2.28	0.47
1:A:884:TYR:OH	1:A:1142:ARG:NH1	2.48	0.47
1:A:916:PHE:O	1:A:920:THR:N	2.48	0.47
2:B:186:LEU:O	2:B:190:LEU:HD23	2.15	0.47
2:B:320:SER:O	2:B:323:HIS:NE2	2.47	0.47
2:B:461:THR:OG1	2:B:463:LYS:CE	2.62	0.47
2:B:81:GLY:HA3	2:B:86:LEU:HD11	1.96	0.47
2:C:186:LEU:O	2:C:190:LEU:HG	2.15	0.47
2:B:419:GLU:OE2	2:C:201:ASN:ND2	2.48	0.47
1:A:1068:ASP:O	1:A:1069:ILE:CB	2.63	0.47
1:A:119:PRO:C	1:A:121:PRO:HD3	2.35	0.47
1:A:293:ASP:CB	1:A:296:SER:OG	2.63	0.47
1:A:305:SER:O	1:A:309:ARG:HG2	2.13	0.47
1:A:557:GLU:C	1:A:559:LEU:N	2.67	0.47
1:A:572:TRP:CA	1:A:572:TRP:CE3	2.98	0.47
1:A:581:ASP:OD1	1:A:581:ASP:O	2.32	0.47
1:A:793:GLU:N	1:A:796:LYS:HD3	2.30	0.47
1:A:824:ALA:HA	1:A:827:ARG:HB3	1.96	0.47
1:A:848:GLY:O	1:A:849:THR:CB	2.59	0.47
1:A:76:ILE:HG12	1:A:911:HIS:NE2	2.30	0.47
1:A:950:ASN:O	1:A:954:ILE:N	2.31	0.47
1:A:948:ILE:O	1:A:951:TYR:HB2	2.14	0.47
1:A:992:LEU:HB2	1:A:997:LEU:HB2	1.97	0.47
2:B:246:ARG:C	2:B:246:ARG:NE	2.68	0.47
2:B:254:PHE:O	2:B:257:ARG:N	2.46	0.47
2:C:441:VAL:CG1	2:C:454:HIS:CE1	2.88	0.47
2:C:466:MET:HE2	2:C:471:LEU:HB3	1.96	0.47
2:C:79:LEU:HD12	2:C:98:GLY:O	2.15	0.47
1:D:515:ILE:HG22	1:D:565:HIS:ND1	2.30	0.47
2:B:389:ARG:HG2	2:B:444:THR:HG22	1.96	0.47
2:B:445:GLU:C	2:B:447:THR:N	2.68	0.47
1:D:499:LYS:NZ	1:D:499:LYS:CB	2.76	0.47
2:E:272:ASN:CA	2:E:292:ASN:HD22	2.26	0.47
2:E:481:SER:HA	2:E:484:ASN:OD1	2.14	0.47
2:C:479:ILE:H	2:C:479:ILE:HG13	1.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:216:HIS:O	2:C:232:GLY:N	2.40	0.47
1:D:1027:LYS:HZ3	1:D:1091:GLU:CG	2.20	0.47
1:D:1055:SER:O	1:D:1058:PHE:HD2	1.97	0.47
1:D:271:VAL:HG22	1:D:295:MET:CG	2.44	0.47
1:D:275:ARG:HG3	1:D:279:ARG:HG3	1.96	0.47
1:D:307:PHE:O	1:D:310:SER:OG	2.23	0.47
1:D:308:GLN:HE22	1:D:1027:LYS:HZ2	1.63	0.47
1:D:455:LEU:HD22	1:D:794:ILE:HB	1.96	0.47
1:D:80:SER:O	1:D:81:ARG:C	2.52	0.47
2:E:254:PHE:HD1	2:E:257:ARG:HE	1.63	0.47
2:F:347:LEU:HD12	2:F:347:LEU:N	2.30	0.47
2:F:360:SER:HB3	2:F:364:LYS:CE	2.44	0.47
2:F:406:LEU:O	2:F:407:LEU:C	2.54	0.47
1:A:114:GLY:N	1:A:116:PRO:CD	2.73	0.47
1:A:81:ARG:HH12	1:A:127:LEU:H	1.61	0.47
1:A:371:LYS:O	1:A:372:GLU:CG	2.63	0.47
1:A:392:LEU:HD22	1:A:393:MET:CE	2.44	0.47
1:A:824:ALA:CA	1:A:827:ARG:HB3	2.44	0.47
1:A:830:ASP:C	1:A:831:TYR:CD1	2.83	0.47
1:A:860:LEU:CD2	1:A:1133:ILE:HA	2.44	0.47
1:A:925:LYS:HG2	1:A:926:SER:H	1.78	0.47
1:A:959:GLN:N	1:A:986:TYR:CE1	2.83	0.47
2:B:236:GLU:CG	2:B:341:ASP:HB2	2.43	0.47
2:B:81:GLY:O	2:B:86:LEU:HD11	2.14	0.47
2:C:452:LEU:HD12	2:C:453:ILE:CA	2.44	0.47
1:D:515:ILE:O	1:D:565:HIS:ND1	2.48	0.47
1:A:500:ALA:HA	1:A:521:PRO:HD3	1.97	0.47
1:A:633:LYS:HZ2	1:A:634:LEU:HB2	1.80	0.47
1:D:966:LEU:CA	1:D:970:ASN:HD22	2.28	0.47
1:A:230:GLU:CG	1:A:231:GLU:N	2.78	0.47
2:B:251:TRP:HA	2:B:251:TRP:CE3	2.49	0.47
1:D:1002:GLU:HG3	1:D:1003:TRP:N	2.28	0.47
1:D:948:ILE:CG2	1:D:1071:ARG:O	2.63	0.47
1:D:114:GLY:O	1:D:116:PRO:HD3	2.13	0.47
1:D:249:VAL:CG1	1:D:251:THR:HG22	2.45	0.47
1:D:627:ARG:O	1:D:627:ARG:HG3	2.15	0.47
1:D:597:ARG:NH1	1:D:725:PRO:CD	2.78	0.47
2:E:109:ASN:CB	2:E:378:LEU:HB3	2.44	0.47
2:E:182:ARG:N	2:E:215:PHE:CZ	2.83	0.47
2:E:319:VAL:CG1	2:E:320:SER:N	2.78	0.47
2:F:303:LEU:CA	2:F:337:SER:CB	2.92	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:413:VAL:HG12	2:F:414:TRP:H	1.80	0.47
1:A:1019:TRP:O	1:A:1020:ILE:HB	2.15	0.47
1:A:867:PRO:HG3	1:A:1067:SER:OG	2.14	0.47
1:A:1144:GLU:H	1:A:1144:GLU:HG2	1.32	0.47
1:A:1150:ALA:CB	1:A:1223:ILE:HG13	2.44	0.47
1:A:132:GLY:CA	1:A:138:HIS:CE1	2.98	0.47
1:A:175:TRP:HB3	1:A:186:PRO:CG	2.45	0.47
1:A:256:PRO:HD2	1:A:257:THR:N	2.19	0.47
1:A:197:PHE:CB	1:A:277:HIS:CD2	2.98	0.47
1:A:295:MET:O	1:A:299:MET:HG2	2.15	0.47
1:A:369:PRO:O	1:A:370:PRO:C	2.50	0.47
1:A:441:TRP:CD1	1:A:877:MET:HB3	2.49	0.47
1:A:481:GLU:C	1:A:482:ASP:CG	2.73	0.47
1:A:79:LEU:HB3	1:A:83:LEU:CD2	2.45	0.47
1:A:866:ARG:CB	1:A:867:PRO:HD3	2.35	0.47
2:B:400:GLN:CA	2:B:400:GLN:OE1	2.63	0.47
2:C:188:GLY:O	2:C:190:LEU:N	2.48	0.47
2:C:241:TRP:CZ3	2:C:255:TRP:CE2	2.99	0.47
2:C:394:GLU:OE1	2:C:394:GLU:N	2.48	0.47
2:C:398:VAL:HA	2:C:401:GLY:CA	2.43	0.47
2:C:469:SER:O	2:C:472:LYS:HB2	2.15	0.47
1:D:497:GLN:O	1:D:517:GLY:HA2	2.15	0.47
1:D:1000:GLU:C	1:D:1004:LEU:HD13	2.35	0.47
1:D:1019:TRP:HD1	1:D:1020:ILE:H	1.63	0.47
1:D:1107:ASP:O	1:D:1108:TYR:C	2.53	0.47
1:D:1111:LEU:HD22	1:D:1164:PHE:CD1	2.49	0.47
1:D:1133:ILE:N	1:D:1133:ILE:CD1	2.78	0.47
1:D:196:VAL:CG1	1:D:268:GLY:H	2.27	0.47
1:D:306:SER:CB	1:D:1058:PHE:CZ	2.98	0.47
1:D:616:GLU:O	1:D:617:ARG:HG3	2.14	0.47
1:D:646:VAL:HG12	1:D:647:CYS:H	1.79	0.47
1:D:909:GLY:O	1:D:912:GLY:N	2.48	0.47
1:D:923:GLY:HA2	1:D:931:LEU:HD11	1.96	0.47
1:D:922:GLN:HG3	1:D:931:LEU:HD22	1.96	0.47
2:E:86:LEU:HB2	2:E:91:LEU:HD21	1.96	0.47
2:F:406:LEU:CA	2:F:409:ASN:HD21	2.19	0.47
2:F:470:LYS:CE	2:F:471:LEU:HD21	2.45	0.47
1:A:290:ARG:O	1:A:291:PHE:HD2	1.98	0.47
1:A:439:GLN:HG2	1:A:835:GLY:C	2.34	0.47
1:A:456:GLN:HE21	1:A:798:ILE:CD1	2.27	0.47
1:A:465:ASP:CG	1:A:466:LEU:N	2.67	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:576:LEU:CD1	1:A:577:CYS:HB2	2.41	0.47
1:A:719:LEU:CD2	1:A:719:LEU:N	2.73	0.47
1:A:843:GLN:O	1:A:843:GLN:CG	2.55	0.47
2:B:257:ARG:O	2:B:258:HIS:O	2.33	0.47
2:C:247:THR:HG1	2:C:251:TRP:HD1	1.58	0.47
2:C:308:ASP:OD1	2:C:333:PRO:O	2.33	0.47
2:C:397:GLN:HG2	2:C:398:VAL:N	2.29	0.47
1:A:503:VAL:CG1	1:A:520:ALA:C	2.65	0.47
1:A:739:TYR:O	1:A:741:ASP:N	2.25	0.47
1:D:498:LYS:HD3	1:D:521:PRO:HG3	1.97	0.47
2:C:94:GLY:HA2	2:C:96:HIS:CE1	2.49	0.47
1:A:736:ASN:ND2	1:A:736:ASN:N	2.63	0.47
1:D:1019:TRP:C	1:D:1019:TRP:CD1	2.87	0.46
1:D:1025:LEU:O	1:D:1027:LYS:CD	2.57	0.46
1:D:1040:LYS:C	1:D:1042:GLU:H	2.17	0.46
1:D:1209:ARG:HE	2:F:253:ASP:HB2	1.80	0.46
1:D:78:MET:SD	1:D:121:PRO:HB2	2.55	0.46
1:D:1147:TYR:HB3	1:D:1227:THR:HG22	1.97	0.46
1:D:175:TRP:H	1:D:220:TRP:HB2	1.80	0.46
1:D:173:GLU:OE2	1:D:223:TRP:NE1	2.48	0.46
1:D:359:VAL:CG1	1:D:363:TYR:HB3	2.44	0.46
1:D:930:ASP:OD2	1:D:930:ASP:O	2.34	0.46
1:D:963:GLU:HB3	1:D:978:ALA:O	2.15	0.46
2:E:323:HIS:C	2:E:331:VAL:O	2.52	0.46
2:E:403:PHE:CE1	2:E:407:LEU:HD11	2.50	0.46
2:E:461:THR:C	2:E:462:MET:HE2	2.36	0.46
2:F:343:ASP:O	2:F:346:MET:HB3	2.15	0.46
2:F:86:LEU:N	2:F:86:LEU:HD12	2.30	0.46
1:A:1057:MET:CA	1:A:1064:ILE:HG21	2.45	0.46
1:A:386:ARG:CZ	1:A:387:GLU:CG	2.93	0.46
1:A:407:PHE:O	1:A:411:LEU:HB3	2.15	0.46
1:A:465:ASP:C	1:A:465:ASP:OD1	2.54	0.46
1:A:484:TRP:C	1:A:488:LEU:HG	2.33	0.46
1:A:601:LYS:HZ3	1:A:616:GLU:CG	2.24	0.46
1:A:752:LEU:CB	1:A:753:PRO:CA	2.93	0.46
1:A:879:GLN:CA	1:A:879:GLN:NE2	2.77	0.46
1:A:935:THR:HG21	1:A:950:ASN:HD21	1.78	0.46
1:A:966:LEU:O	1:A:970:ASN:N	2.46	0.46
2:B:283:GLY:HA2	2:B:306:LEU:HD23	1.97	0.46
2:C:440:THR:O	2:C:457:SER:OG	2.24	0.46
1:D:235:TRP:O	1:D:236:THR:HG23	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:228:VAL:O	2:F:228:VAL:HG23	2.15	0.46
1:D:1042:GLU:C	1:D:1044:VAL:H	2.19	0.46
1:D:1064:ILE:O	1:D:1064:ILE:CG2	2.62	0.46
1:D:944:GLU:OE1	1:D:1071:ARG:NH1	2.49	0.46
1:D:1122:GLU:O	1:D:1123:PHE:CD1	2.68	0.46
1:D:1187:ARG:HG3	1:D:1212:ILE:HG12	1.96	0.46
1:D:307:PHE:CD2	1:D:1048:ALA:HB1	2.50	0.46
1:D:417:ARG:CG	1:D:417:ARG:HH11	2.28	0.46
1:D:648:PRO:CG	1:D:649:TYR:N	2.78	0.46
1:D:735:GLY:CA	1:D:749:PHE:CB	2.79	0.46
2:E:303:LEU:N	2:E:303:LEU:HD12	2.29	0.46
2:E:96:HIS:C	2:E:98:GLY:N	2.68	0.46
2:F:349:TYR:O	2:F:352:ASP:HB3	2.14	0.46
2:F:447:THR:HA	2:F:452:LEU:C	2.35	0.46
1:A:1158:LEU:C	1:A:1162:CYS:SG	2.93	0.46
1:A:403:THR:O	1:A:404:HIS:C	2.53	0.46
1:A:612:LEU:HD23	1:A:614:TYR:HA	1.96	0.46
1:A:657:ARG:O	1:A:660:CYS:CB	2.51	0.46
1:A:757:GLY:O	1:A:758:ASN:CB	2.62	0.46
1:A:618:HIS:CA	1:A:762:VAL:HG21	2.42	0.46
2:C:109:ASN:O	2:C:113:GLU:N	2.31	0.46
2:C:130:ALA:HB1	2:C:182:ARG:NH1	2.24	0.46
2:C:78:PHE:CE2	2:C:350:LEU:HD23	2.50	0.46
1:D:1020:ILE:HD13	1:D:1021:SER:HA	1.94	0.46
1:D:1030:ARG:C	1:D:1032:THR:N	2.68	0.46
1:D:1060:LYS:HD2	1:D:1060:LYS:H	1.80	0.46
1:D:118:VAL:CG2	1:D:121:PRO:HD3	2.45	0.46
1:D:1224:ILE:O	1:D:1227:THR:C	2.54	0.46
1:D:184:ALA:C	1:D:185:VAL:CG1	2.84	0.46
1:D:486:TRP:CA	1:D:489:GLU:OE2	2.56	0.46
1:D:656:TYR:CG	1:D:744:ILE:HD12	2.49	0.46
1:D:73:PRO:O	1:D:74:LEU:HD22	2.15	0.46
1:D:768:LYS:CA	1:D:772:PRO:CG	2.93	0.46
1:D:886:LEU:HA	1:D:1146:ARG:HH22	1.81	0.46
2:F:256:LEU:HD12	2:F:256:LEU:HA	1.82	0.46
2:F:314:MET:CE	2:F:315:TYR:CE2	2.94	0.46
2:F:242:PHE:CB	2:F:333:PRO:HB2	2.25	0.46
1:D:546:ARG:HH22	2:F:408:GLU:CD	2.19	0.46
2:F:421:MET:HE3	2:F:421:MET:HB2	1.85	0.46
2:F:454:HIS:NE2	2:F:472:LYS:CE	2.78	0.46
2:F:470:LYS:HB2	2:F:471:LEU:HG	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:84:GLN:C	2:F:86:LEU:N	2.68	0.46
1:A:1019:TRP:C	1:A:1020:ILE:HG22	2.35	0.46
1:A:1048:ALA:HB3	1:A:1092:PHE:CE1	2.48	0.46
1:A:1094:THR:O	1:A:1095:SER:C	2.53	0.46
1:A:1113:LEU:HD23	1:A:1113:LEU:H	1.78	0.46
1:A:1124:ALA:CA	1:A:1148:ARG:NH1	2.78	0.46
1:A:1124:ALA:O	1:A:1126:ASP:OD1	2.34	0.46
1:A:1131:ILE:HD13	1:A:1137:VAL:CA	2.46	0.46
1:A:237:SER:O	1:A:238:GLN:HB2	2.15	0.46
1:A:272:SER:CA	1:A:275:ARG:HB2	2.35	0.46
1:A:294:THR:HG22	1:A:411:LEU:CD2	2.46	0.46
1:A:887:VAL:HG23	1:A:1146:ARG:NH1	2.20	0.46
2:B:212:GLY:O	2:B:235:THR:O	2.34	0.46
2:B:99:PHE:CE2	2:C:131:LEU:HD11	2.44	0.46
2:C:208:LEU:HD11	2:C:240:VAL:CA	2.45	0.46
2:C:344:ARG:CB	2:C:347:LEU:HD22	2.46	0.46
2:C:402:LEU:O	2:C:403:PHE:C	2.54	0.46
2:C:464:GLU:OE2	2:C:466:MET:SD	2.74	0.46
1:D:524:PRO:O	1:D:525:MET:CB	2.53	0.46
2:C:88:ARG:NH1	2:C:88:ARG:CB	2.78	0.46
2:C:90:SER:O	2:C:93:SER:OG	2.26	0.46
1:D:668:LEU:HD23	1:D:668:LEU:N	2.30	0.46
2:C:272:ASN:CB	2:C:292:ASN:HB2	2.43	0.46
1:D:1019:TRP:CZ2	1:D:1167:LYS:HG3	2.51	0.46
1:D:1030:ARG:HA	1:D:1030:ARG:HD3	1.54	0.46
1:D:1157:ASN:O	1:D:1158:LEU:C	2.54	0.46
1:D:1209:ARG:NH1	2:F:285:LYS:NZ	2.63	0.46
1:D:253:ALA:HB2	1:D:281:GLN:N	2.30	0.46
1:D:271:VAL:HG22	1:D:295:MET:CB	2.46	0.46
1:D:307:PHE:C	1:D:311:LEU:HD12	2.34	0.46
1:D:309:ARG:O	1:D:310:SER:C	2.49	0.46
1:D:720:THR:C	1:D:721:ALA:O	2.32	0.46
1:D:794:ILE:O	1:D:798:ILE:HG23	2.15	0.46
1:D:945:HIS:CE1	1:D:969:PHE:CB	2.99	0.46
2:E:293:PHE:O	2:E:295:TRP:N	2.49	0.46
2:E:353:SER:OG	2:E:372:LEU:HD21	2.15	0.46
2:E:460:THR:C	2:E:461:THR:HG22	2.35	0.46
2:F:197:LEU:O	2:F:202:LYS:CA	2.60	0.46
2:F:213:VAL:CG2	2:F:235:THR:OG1	2.63	0.46
2:F:414:TRP:O	2:F:415:PRO:O	2.34	0.46
1:D:472:GLN:C	2:F:461:THR:HG23	2.36	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1068:ASP:O	1:A:1069:ILE:CG2	2.64	0.46
1:A:430:MET:HA	1:A:1130:CYS:SG	2.55	0.46
1:A:77:GLN:HB3	1:A:1174:PRO:HB3	1.96	0.46
1:A:134:ASN:O	1:A:137:GLN:N	2.47	0.46
1:A:196:VAL:HG11	1:A:404:HIS:CB	2.45	0.46
1:A:259:ARG:CA	1:A:259:ARG:HE	2.27	0.46
1:A:273:PHE:C	1:A:276:ALA:CB	2.83	0.46
1:A:752:LEU:HB3	1:A:753:PRO:HA	1.98	0.46
1:A:827:ARG:CG	1:A:828:HIS:N	2.78	0.46
1:A:873:GLU:OE1	1:A:875:LYS:HG2	2.15	0.46
1:A:949:PHE:HD2	1:A:949:PHE:H	1.62	0.46
1:A:960:PRO:HD2	1:A:961:PHE:HE1	1.79	0.46
2:B:117:SER:CB	2:B:262:TRP:HE1	2.21	0.46
2:C:184:ASN:C	2:C:186:LEU:N	2.67	0.46
2:C:201:ASN:O	2:C:202:LYS:CB	2.64	0.46
2:C:400:GLN:CD	2:C:400:GLN:C	2.74	0.46
2:C:428:LEU:O	2:C:429:TYR:C	2.52	0.46
2:C:448:LEU:O	2:C:449:GLU:HG3	2.16	0.46
2:C:474:PHE:CE1	2:C:478:TYR:CD2	2.94	0.46
2:C:134:LYS:HB2	2:C:180:LYS:HE2	1.96	0.46
2:E:184:ASN:C	2:E:186:LEU:N	2.68	0.46
2:F:317:GLY:C	2:F:319:VAL:N	2.56	0.46
2:F:145:PHE:O	2:F:146:ARG:CG	2.64	0.46
2:C:67:GLU:O	2:C:71:GLU:N	2.46	0.46
2:E:345:GLY:O	2:E:348:ALA:HB3	2.15	0.46
1:D:1042:GLU:C	1:D:1044:VAL:N	2.69	0.46
1:D:1058:PHE:CG	1:D:1059:ASN:N	2.82	0.46
1:D:1124:ALA:CB	1:D:1148:ARG:HH12	2.26	0.46
1:D:192:GLU:HG2	1:D:193:ARG:N	2.31	0.46
1:D:272:SER:HB3	1:D:843:GLN:OE1	2.15	0.46
1:D:429:GLU:HG2	1:D:1117:LYS:CE	2.42	0.46
1:D:460:LYS:HG2	1:D:461:LYS:H	1.80	0.46
1:D:599:THR:C	1:D:602:LEU:HD12	2.36	0.46
1:D:751:LYS:HE2	1:D:751:LYS:HB3	1.55	0.46
1:D:865:ALA:HB1	1:D:872:SER:CB	2.45	0.46
1:D:995:TYR:H	1:D:996:ARG:HH11	1.58	0.46
2:E:116:THR:C	2:E:118:VAL:N	2.69	0.46
2:F:233:GLU:O	2:F:234:LYS:CG	2.61	0.46
2:F:454:HIS:HB2	2:F:466:MET:CA	2.45	0.46
1:A:1017:GLY:C	1:A:1019:TRP:N	2.69	0.46
1:A:1020:ILE:CG1	1:A:1021:SER:H	2.28	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1165:ALA:HB2	1:A:1173:LEU:HD11	1.97	0.46
1:A:1222:GLN:O	1:A:1225:GLU:HB2	2.15	0.46
1:A:192:GLU:OE2	1:A:193:ARG:N	2.48	0.46
1:A:213:VAL:N	1:A:221:TYR:HE2	2.11	0.46
1:A:313:ILE:HD12	1:A:313:ILE:C	2.36	0.46
1:A:452:TYR:HA	1:A:869:ARG:NH2	2.19	0.46
1:A:666:GLN:HA	1:A:669:MET:CE	2.45	0.46
1:A:711:ALA:C	1:A:713:PRO:CD	2.65	0.46
1:A:720:THR:CG2	1:A:721:ALA:H	2.28	0.46
1:A:959:GLN:CB	1:A:960:PRO:CD	2.60	0.46
1:A:967:MET:O	1:A:970:ASN:O	2.33	0.46
2:B:102:LEU:HD12	2:B:102:LEU:N	2.19	0.46
2:B:306:LEU:O	2:B:335:VAL:HB	2.16	0.46
2:B:336:LEU:HG	2:B:337:SER:N	2.30	0.46
2:C:236:GLU:CB	2:C:341:ASP:HB3	2.45	0.46
2:C:360:SER:HB3	2:C:364:LYS:HE2	1.97	0.46
1:A:634:LEU:CB	1:A:635:PRO:CA	2.78	0.46
2:E:315:TYR:O	2:E:316:PRO:O	2.34	0.46
2:F:367:LEU:HD12	2:F:367:LEU:C	2.36	0.46
2:E:84:GLN:O	2:E:84:GLN:NE2	2.49	0.46
1:D:1014:ARG:NE	1:D:1022:LEU:HD22	2.30	0.46
1:D:1019:TRP:CE3	1:D:1168:LEU:CG	2.75	0.46
1:D:1151:LEU:HD11	1:D:1227:THR:OG1	2.15	0.46
1:D:157:LEU:N	1:D:157:LEU:HD23	2.30	0.46
1:D:179:GLY:C	1:D:182:GLY:N	2.69	0.46
1:D:345:TRP:N	1:D:1040:LYS:HZ2	2.13	0.46
1:D:636:THR:CG2	1:D:637:GLY:N	2.74	0.46
1:D:719:LEU:O	1:D:722:ARG:HD3	2.15	0.46
1:D:720:THR:HA	1:D:721:ALA:O	2.15	0.46
1:D:827:ARG:CG	1:D:828:HIS:N	2.79	0.46
2:E:283:GLY:HA3	2:E:306:LEU:HD23	1.98	0.46
2:E:326:ASP:O	2:E:327:GLY:C	2.54	0.46
2:F:322:LEU:HD22	2:F:323:HIS:N	2.31	0.46
2:F:431:LYS:O	2:F:434:GLU:HB2	2.15	0.46
2:F:477:LYS:O	2:F:478:TYR:C	2.53	0.46
1:A:1030:ARG:CD	1:A:1040:LYS:HZ2	2.29	0.46
1:A:1043:VAL:HG22	1:A:1046:GLU:HB3	1.98	0.46
1:A:285:GLN:HA	1:A:1128:ARG:NH2	2.31	0.46
1:A:860:LEU:HD23	1:A:1133:ILE:HG12	1.96	0.46
1:A:1190:ARG:CZ	1:A:1190:ARG:HB2	2.45	0.46
1:A:132:GLY:HA2	1:A:138:HIS:CE1	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:195:LEU:CD2	1:A:217:PRO:CD	2.87	0.46
1:A:252:GLY:O	1:A:253:ALA:CB	2.64	0.46
1:A:309:ARG:HA	1:A:312:TRP:HB2	1.98	0.46
1:A:313:ILE:HG13	1:A:314:ALA:N	2.31	0.46
1:A:441:TRP:HE1	1:A:878:VAL:N	2.13	0.46
1:A:569:HIS:CB	1:A:570:PRO:HD2	2.45	0.46
1:A:570:PRO:HG3	1:A:572:TRP:CZ3	2.51	0.46
1:A:649:TYR:HA	1:A:652:ILE:HB	1.97	0.46
1:A:817:PRO:HG2	1:A:818:ARG:N	2.30	0.46
2:B:187:HIS:O	2:B:190:LEU:HB2	2.16	0.46
2:B:423:SER:HG	2:B:428:LEU:HD22	1.79	0.46
2:B:74:GLN:O	2:B:75:ARG:C	2.53	0.46
2:C:185:LEU:CG	2:C:214:CYS:SG	3.03	0.46
2:C:300:ILE:CA	2:C:342:LEU:HD12	2.42	0.46
2:C:375:HIS:NE2	2:C:377:CYS:SG	2.87	0.46
1:A:503:VAL:CB	1:A:521:PRO:CA	2.76	0.46
1:D:503:VAL:O	1:D:505:LYS:O	2.33	0.46
2:F:61:GLY:HA2	2:F:64:GLU:HG2	1.97	0.46
1:D:163:PRO:HB2	1:D:165:LYS:CG	2.36	0.46
2:E:311:LEU:N	2:E:314:MET:HG3	2.31	0.46
1:A:973:LEU:CB	1:A:977:GLU:HB2	2.45	0.46
2:F:280:ASP:C	2:F:282:GLU:H	2.19	0.46
1:D:1192:GLU:O	1:D:1192:GLU:CG	2.59	0.46
1:D:1206:MET:SD	1:D:1206:MET:C	2.93	0.46
1:D:1206:MET:SD	2:F:257:ARG:NH1	2.88	0.46
1:D:201:VAL:HG13	1:D:202:CYS:N	2.31	0.46
1:D:271:VAL:CG2	1:D:295:MET:HG3	2.46	0.46
1:D:375:GLU:HG3	1:D:379:LYS:HG3	1.93	0.46
1:D:658:LYS:HE3	1:D:713:PRO:HG3	1.98	0.46
1:D:748:TRP:CE3	1:D:748:TRP:O	2.69	0.46
1:D:444:TYR:CZ	1:D:877:MET:HE1	2.51	0.46
1:D:943:ARG:O	1:D:946:ALA:N	2.49	0.46
2:E:116:THR:C	2:E:120:VAL:HG12	2.36	0.46
2:E:325:ARG:HG2	2:E:330:ASN:H	1.80	0.46
2:E:373:LYS:O	2:E:374:LEU:HG	2.16	0.46
2:E:99:PHE:O	2:F:128:VAL:CG1	2.63	0.46
2:F:115:TRP:CD1	2:F:119:VAL:CG2	2.98	0.46
2:F:295:TRP:O	2:F:295:TRP:CD1	2.69	0.46
2:F:300:ILE:HA	2:F:342:LEU:HD12	1.97	0.46
2:F:344:ARG:O	2:F:345:GLY:C	2.52	0.46
2:F:354:PHE:HD1	2:F:371:VAL:O	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:455:LEU:HB3	2:F:456:ARG:NE	2.29	0.46
2:F:459:ASP:O	2:F:460:THR:HB	2.16	0.46
2:F:70:LEU:HD13	2:F:70:LEU:C	2.36	0.46
1:A:773:LYS:HZ1	1:A:1057:MET:HE3	1.81	0.46
1:A:1123:PHE:C	1:A:1125:ILE:H	2.19	0.46
1:A:1171:ASN:O	1:A:1172:ASP:OD1	2.33	0.46
1:A:265:LEU:HD12	1:A:265:LEU:C	2.36	0.46
1:A:297:MET:HE2	1:A:297:MET:HB3	1.62	0.46
1:A:311:LEU:HD22	1:A:1047:ARG:HD2	1.97	0.46
1:A:363:ARG:HH21	1:A:410:GLN:HG2	1.81	0.46
1:A:439:GLN:NE2	1:A:837:TYR:CE2	2.79	0.46
1:A:599:THR:HB	1:A:600:PRO:HD3	1.96	0.46
1:A:657:ARG:HD3	1:A:744:ILE:CG2	2.45	0.46
1:A:731:SER:O	1:A:732:TYR:C	2.52	0.46
1:A:70:ARG:HD3	1:A:81:ARG:CB	2.44	0.46
2:B:132:HIS:HA	2:B:182:ARG:CZ	2.46	0.46
2:C:115:TRP:NE1	2:C:119:VAL:CG2	2.79	0.46
2:C:243:THR:HG21	2:C:247:THR:HG1	1.80	0.46
1:D:623:LEU:HB3	1:D:667:GLN:HE22	1.71	0.46
2:B:475:LEU:CA	2:B:479:ILE:HD13	2.46	0.46
1:A:751:LYS:HB3	1:A:751:LYS:HE2	1.57	0.46
1:D:1050:LYS:O	1:D:1053:THR:HB	2.16	0.46
1:D:877:MET:HE3	1:D:1207:GLU:HB2	1.97	0.46
1:D:1207:GLU:CB	1:D:1210:TYR:HD2	2.28	0.46
1:D:270:ASN:N	1:D:294:THR:HG23	2.30	0.46
1:D:374:ARG:NE	1:D:392:LEU:HD21	2.31	0.46
1:D:559:LEU:CD1	1:D:560:PRO:HD3	2.39	0.46
1:D:597:ARG:HG2	1:D:617:ARG:HD2	1.98	0.46
1:D:622:TYR:O	1:D:746:GLY:HA2	2.14	0.46
1:D:649:TYR:O	1:D:653:GLU:N	2.45	0.46
1:D:729:GLN:HB3	1:D:731:SER:OG	2.16	0.46
2:E:344:ARG:O	2:E:346:MET:N	2.49	0.46
2:E:428:LEU:O	2:E:431:LYS:CB	2.57	0.46
2:F:389:ARG:CD	2:F:390:GLY:H	2.28	0.46
1:D:457:ARG:NH2	2:F:485:VAL:HG23	2.20	0.46
1:A:1141:VAL:HG12	1:A:1142:ARG:N	2.31	0.46
1:A:282:TYR:CE2	1:A:291:PHE:CE1	3.04	0.46
1:A:302:SER:HB2	1:A:351:SER:CB	2.45	0.46
1:A:369:PRO:O	1:A:371:LYS:N	2.49	0.46
1:A:630:ASN:ND2	1:A:631:LEU:N	2.62	0.46
1:A:658:LYS:O	1:A:662:GLU:CD	2.53	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:732:TYR:HD1	1:A:755:LYS:NZ	2.08	0.46
1:A:657:ARG:HD3	1:A:744:ILE:HG22	1.98	0.46
1:A:829:PRO:O	1:A:830:ASP:HB2	2.15	0.46
1:A:933:SER:CB	1:A:943:ARG:CZ	2.91	0.46
1:A:993:ARG:HG3	1:A:994:TRP:HD1	1.81	0.46
2:B:126:PHE:HB3	2:C:101:PRO:HB3	1.98	0.46
2:B:261:GLN:O	2:B:262:TRP:O	2.34	0.46
2:B:289:LEU:HB3	2:B:300:ILE:HG13	1.97	0.46
2:B:378:LEU:CD1	2:B:378:LEU:N	2.78	0.46
2:C:115:TRP:NE1	2:C:119:VAL:HG21	2.31	0.46
2:C:291:TYR:CD1	2:C:348:ALA:HB1	2.51	0.46
2:C:348:ALA:O	2:C:351:TYR:CB	2.63	0.46
2:C:361:PHE:CD2	2:C:362:THR:HG23	2.48	0.46
2:C:455:LEU:HD12	2:C:455:LEU:H	1.81	0.46
1:D:664:GLY:CA	1:D:667:GLN:HG3	2.45	0.46
2:E:390:GLY:HA3	2:E:395:LEU:HD23	1.98	0.46
1:A:539:PHE:O	1:A:540:GLN:HB2	2.16	0.46
2:E:230:SER:O	2:E:231:ILE:HB	2.15	0.46
2:C:221:THR:CG2	2:C:222:LYS:H	2.28	0.46
2:B:454:HIS:N	2:B:454:HIS:ND1	2.63	0.46
1:D:1024:ASP:O	1:D:1026:ARG:N	2.48	0.46
1:D:1162:CYS:O	1:D:1166:TYR:CE1	2.69	0.46
1:D:1172:ASP:O	1:D:1173:LEU:CG	2.44	0.46
1:D:1198:LYS:O	1:D:1201:SER:O	2.34	0.46
1:D:868:ASP:OD1	1:D:1199:THR:CB	2.63	0.46
1:D:140:ARG:NH1	1:D:140:ARG:CG	2.74	0.46
1:D:277:HIS:C	1:D:291:PHE:CE1	2.89	0.46
1:D:293:ASP:CB	1:D:296:SER:CB	2.94	0.46
1:D:400:VAL:O	1:D:400:VAL:HG12	2.16	0.46
1:D:550:GLN:OE1	1:D:551:LYS:HE3	2.16	0.46
1:D:794:ILE:HA	1:D:797:MET:HG3	1.96	0.46
1:D:822:PRO:N	1:D:825:VAL:HB	2.31	0.46
1:D:865:ALA:CB	1:D:872:SER:O	2.63	0.46
1:D:88:PHE:O	1:D:1172:ASP:OD1	2.33	0.46
1:D:116:PRO:CA	1:D:921:LEU:HD13	2.40	0.46
2:E:114:TRP:CZ2	2:E:238:SER:O	2.69	0.46
2:E:386:ASP:N	2:E:414:TRP:O	2.49	0.46
2:F:259:ARG:HE	2:F:339:ASN:ND2	2.14	0.46
2:F:92:LEU:CD2	2:F:344:ARG:HE	2.27	0.46
1:A:1067:SER:O	1:A:1068:ASP:CB	2.62	0.46
1:A:1068:ASP:H	1:A:1071:ARG:CD	2.28	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1111:LEU:N	1:A:1111:LEU:CD1	2.79	0.46
1:A:1198:LYS:O	1:A:1199:THR:OG1	2.33	0.46
1:A:598:VAL:N	1:A:616:GLU:OE1	2.49	0.46
1:A:801:TRP:O	1:A:805:HIS:CB	2.64	0.46
1:A:879:GLN:HG3	1:A:886:LEU:CD1	2.46	0.46
1:A:76:ILE:CG2	1:A:911:HIS:CE1	2.98	0.46
2:B:79:LEU:HA	2:B:100:GLY:HA3	1.98	0.46
2:B:259:ARG:O	2:B:260:LEU:O	2.34	0.46
2:B:283:GLY:O	2:B:284:ARG:CB	2.64	0.46
1:A:604:ALA:HB3	2:C:363:ARG:NH2	2.30	0.46
2:B:447:THR:HG23	2:B:451:GLY:C	2.35	0.46
1:D:1118:TRP:C	1:D:1122:GLU:HG2	2.35	0.46
1:D:150:TYR:CE1	1:D:266:VAL:HG21	2.51	0.46
1:D:279:ARG:O	1:D:841:LEU:CG	2.54	0.46
1:D:368:PRO:CB	1:D:369:LEU:CD2	2.94	0.46
1:D:627:ARG:O	1:D:627:ARG:CG	2.64	0.46
1:D:652:ILE:HA	1:D:655:LEU:CB	2.46	0.46
1:D:770:PHE:O	1:D:773:LYS:CG	2.64	0.46
1:D:801:TRP:CD2	1:D:870:VAL:CG1	2.99	0.46
1:D:904:ASP:OD1	1:D:915:ALA:HB3	2.16	0.46
1:D:935:THR:O	1:D:938:THR:OG1	2.33	0.46
1:D:932:HIS:CE1	1:D:947:LYS:HG2	2.51	0.46
1:D:986:TYR:CD1	1:D:990:LYS:HD2	2.51	0.46
2:E:254:PHE:CZ	2:E:258:HIS:HB2	2.51	0.46
2:E:263:TRP:CH2	2:E:300:ILE:HD12	2.51	0.46
2:E:419:GLU:OE1	2:E:420:THR:HG22	2.16	0.46
2:F:246:ARG:C	2:F:248:SER:N	2.56	0.46
2:F:306:LEU:HB3	2:F:335:VAL:HG21	1.98	0.46
2:F:423:SER:OG	2:F:424:SER:N	2.47	0.46
2:F:63:GLY:O	2:F:67:GLU:HG3	2.16	0.46
2:F:76:ARG:CG	2:F:76:ARG:NH1	2.79	0.46
1:A:176:THR:O	1:A:177:ARG:HG2	2.16	0.46
1:A:811:GLN:HB2	1:A:840:ILE:CG2	2.33	0.46
1:A:821:LEU:CG	1:A:826:ILE:HG23	2.43	0.46
1:A:845:VAL:CG2	1:A:855:VAL:HB	2.45	0.46
1:A:952:GLY:HA2	1:A:955:TYR:HD1	1.81	0.46
2:B:182:ARG:O	2:B:215:PHE:HB2	2.15	0.46
2:B:245:PRO:O	2:B:248:SER:CB	2.64	0.46
2:B:400:GLN:O	2:B:403:PHE:HB3	2.15	0.46
2:B:374:LEU:N	2:B:458:ARG:NH1	2.64	0.46
2:C:130:ALA:CA	2:C:182:ARG:NH1	2.79	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:350:LEU:N	2:C:350:LEU:CD1	2.58	0.46
2:C:444:THR:H	2:C:447:THR:CG2	2.23	0.46
2:C:452:LEU:HD12	2:C:453:ILE:HG13	1.98	0.46
2:C:74:GLN:NE2	2:C:75:ARG:HH21	2.14	0.46
1:D:665:LYS:O	1:D:669:MET:N	2.38	0.46
1:A:987:ALA:HA	1:A:991:GLY:HA3	1.98	0.46
1:D:1034:ARG:O	1:D:1035:LYS:HB2	2.16	0.46
1:D:1028:VAL:HA	1:D:1031:GLU:HG2	1.98	0.45
1:D:1206:MET:O	1:D:1210:TYR:CA	2.64	0.45
1:D:375:GLU:O	1:D:376:LEU:C	2.54	0.45
1:D:407:PHE:HD1	1:D:411:LEU:CG	2.29	0.45
1:D:461:LYS:HA	1:D:464:MET:HE2	1.96	0.45
1:D:608:ASP:OD1	1:D:611:PRO:C	2.55	0.45
1:D:614:TYR:HB3	1:D:615:SER:O	2.16	0.45
1:D:645:VAL:HG13	1:D:650:ARG:CD	2.45	0.45
1:D:768:LYS:O	1:D:769:ASP:CB	2.63	0.45
2:E:211:ILE:N	2:E:211:ILE:CD1	2.79	0.45
2:F:128:VAL:CG1	2:F:129:ASP:N	2.79	0.45
2:F:131:LEU:O	2:F:132:HIS:C	2.54	0.45
2:F:72:ILE:HD12	2:F:76:ARG:HE	1.81	0.45
1:A:1069:ILE:O	1:A:1069:ILE:HD13	2.16	0.45
1:A:1109:LEU:HD13	1:A:1109:LEU:HA	1.60	0.45
1:A:1198:LYS:HA	1:A:1204:THR:HG23	1.98	0.45
1:A:1227:THR:O	1:A:1228:LYS:O	2.35	0.45
1:A:185:VAL:N	1:A:186:PRO:CD	2.80	0.45
1:A:238:GLN:CB	1:A:239:LEU:HG	2.42	0.45
1:A:407:PHE:N	1:A:407:PHE:CD2	2.83	0.45
1:A:438:ASN:O	1:A:439:GLN:C	2.55	0.45
1:A:600:PRO:HG3	1:A:765:PRO:HG3	1.98	0.45
1:A:608:ASP:CA	1:A:620:TRP:HH2	2.29	0.45
1:A:77:GLN:O	1:A:79:LEU:HG	2.16	0.45
1:A:788:GLY:O	1:A:791:ALA:HB3	2.16	0.45
1:A:808:ILE:O	1:A:811:GLN:NE2	2.49	0.45
1:A:857:PRO:O	1:A:861:THR:CB	2.64	0.45
1:A:943:ARG:O	1:A:946:ALA:N	2.49	0.45
2:B:246:ARG:O	2:B:248:SER:N	2.49	0.45
2:B:439:PHE:HD1	2:B:456:ARG:O	1.98	0.45
2:C:310:GLU:CD	2:C:310:GLU:N	2.70	0.45
2:C:394:GLU:CA	2:C:397:GLN:HB3	2.45	0.45
2:C:432:TYR:O	2:C:436:SER:N	2.48	0.45
2:C:395:LEU:CD2	2:C:443:VAL:CG1	2.89	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:471:LEU:O	2:C:472:LYS:C	2.55	0.45
2:C:470:LYS:O	2:C:474:PHE:N	2.49	0.45
2:F:138:LEU:HD22	2:F:138:LEU:N	2.31	0.45
1:A:94:MET:SD	1:A:909:GLY:HA3	2.56	0.45
1:D:1131:ILE:CD1	1:D:1138:ARG:O	2.63	0.45
1:D:1207:GLU:C	1:D:1210:TYR:H	2.19	0.45
1:D:271:VAL:CG1	1:D:272:SER:N	2.49	0.45
1:D:370:GLU:HB2	1:D:399:ASP:CG	2.37	0.45
1:D:608:ASP:O	1:D:609:GLY:O	2.34	0.45
1:D:739:TYR:O	1:D:740:ASN:ND2	2.49	0.45
1:D:801:TRP:CH2	1:D:805:HIS:CB	2.99	0.45
2:E:106:LEU:HD12	2:E:109:ASN:HD21	1.80	0.45
2:F:129:ASP:OD1	2:F:131:LEU:HD11	2.17	0.45
2:F:351:TYR:CA	2:F:355:GLN:HB2	2.44	0.45
2:F:389:ARG:C	2:F:395:LEU:HD11	2.36	0.45
1:A:1102:GLN:O	1:A:1103:SER:C	2.53	0.45
1:A:1173:LEU:HA	1:A:1174:PRO:HD3	1.65	0.45
1:A:1190:ARG:HD2	1:A:1195:MET:CE	2.45	0.45
1:A:1231:LEU:CG	1:A:1232:GLU:H	2.16	0.45
1:A:278:ILE:HG23	1:A:291:PHE:HE1	1.80	0.45
1:A:350:ILE:O	1:A:351:SER:C	2.54	0.45
1:A:386:ARG:NH1	1:A:387:GLU:CG	2.79	0.45
1:A:623:LEU:CG	1:A:623:LEU:O	2.45	0.45
1:A:630:ASN:HB2	1:A:744:ILE:C	2.37	0.45
1:A:651:ALA:O	1:A:655:LEU:HD23	2.16	0.45
1:A:604:ALA:HB2	1:A:717:LEU:HD13	1.99	0.45
1:A:788:GLY:O	1:A:789:PRO:C	2.51	0.45
1:A:439:GLN:CG	1:A:835:GLY:CA	2.75	0.45
1:A:944:GLU:N	1:A:947:LYS:HE3	2.31	0.45
2:B:186:LEU:CD1	2:B:337:SER:HB2	2.37	0.45
2:C:131:LEU:O	2:C:132:HIS:C	2.54	0.45
2:C:314:MET:HE3	2:C:315:TYR:CE2	2.52	0.45
2:C:432:TYR:HB3	2:C:437:ILE:CG1	2.46	0.45
2:C:458:ARG:CG	2:C:458:ARG:HH11	2.20	0.45
2:C:86:LEU:O	2:C:91:LEU:HB2	2.16	0.45
1:D:1070:PRO:HD3	1:D:1071:ARG:NH1	2.30	0.45
1:D:1136:GLU:HG3	1:D:1138:ARG:HH12	1.81	0.45
1:D:1193:VAL:O	1:D:1194:THR:C	2.53	0.45
1:D:1195:MET:O	1:D:1196:ASP:CB	2.63	0.45
1:D:157:LEU:CG	1:D:194:ALA:CB	2.95	0.45
1:D:179:GLY:O	1:D:180:PRO:C	2.53	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:201:VAL:CG1	1:D:202:CYS:N	2.80	0.45
1:D:479:TYR:HE1	1:D:480:LYS:HE2	1.82	0.45
1:D:575:LYS:HE3	1:D:581:ASP:HB3	1.97	0.45
1:D:649:TYR:CD2	1:D:652:ILE:CD1	2.99	0.45
1:D:632:ALA:HB2	1:D:657:ARG:HD2	1.98	0.45
1:D:810:SER:O	1:D:811:GLN:C	2.54	0.45
1:D:922:GLN:CD	1:D:931:LEU:HD13	2.35	0.45
2:E:387:VAL:O	2:E:442:LEU:CD2	2.62	0.45
2:E:422:GLN:HG2	2:E:424:SER:CA	2.47	0.45
2:F:234:LYS:HZ1	2:F:344:ARG:CG	2.28	0.45
1:A:1020:ILE:O	1:A:1021:SER:CB	2.64	0.45
1:A:1200:PRO:HB2	1:A:1203:PRO:HD2	1.98	0.45
1:A:268:GLY:O	1:A:269:HIS:CD2	2.69	0.45
1:A:313:ILE:CG1	1:A:314:ALA:N	2.79	0.45
1:A:297:MET:HE3	1:A:411:LEU:HD22	1.98	0.45
1:A:712:VAL:N	1:A:713:PRO:CD	2.80	0.45
1:A:879:GLN:CB	1:A:886:LEU:CD1	2.92	0.45
1:A:79:LEU:CB	1:A:89:GLY:O	2.64	0.45
1:A:918:TRP:CD1	1:A:918:TRP:O	2.69	0.45
1:A:996:ARG:NH2	1:A:1003:TRP:CB	2.68	0.45
2:B:192:HIS:O	2:B:195:ASN:HB3	2.17	0.45
2:C:194:VAL:O	2:C:197:LEU:CB	2.63	0.45
1:A:783:PRO:CG	2:C:363:ARG:NH1	2.79	0.45
2:C:416:GLY:HA2	2:C:419:GLU:HB3	1.98	0.45
2:C:443:VAL:CB	2:C:447:THR:HG21	2.44	0.45
2:C:86:LEU:HD12	2:C:86:LEU:N	2.31	0.45
1:D:665:LYS:O	1:D:669:MET:O	2.34	0.45
1:D:502:LYS:HD2	1:D:506:GLU:HB3	1.98	0.45
2:B:122:ARG:HD3	2:B:124:GLN:HE22	1.81	0.45
1:D:1099:TRP:CE3	1:D:1099:TRP:HA	2.51	0.45
1:D:1023:GLN:OE1	1:D:1100:VAL:HG22	2.17	0.45
1:D:258:GLN:O	1:D:259:ARG:HB3	2.16	0.45
1:D:316:LYS:HZ1	1:D:348:LEU:CD2	2.29	0.45
1:D:461:LYS:O	1:D:465:ASP:OD2	2.34	0.45
1:D:606:THR:HG1	1:D:614:TYR:HH	1.58	0.45
1:D:652:ILE:O	1:D:655:LEU:HB2	2.17	0.45
1:D:279:ARG:NH1	1:D:841:LEU:HD13	2.31	0.45
1:D:893:SER:OG	1:D:894:GLN:N	2.48	0.45
2:E:329:LYS:O	2:E:330:ASN:O	2.34	0.45
2:E:450:ASN:ND2	2:E:452:LEU:HD21	2.27	0.45
2:E:478:TYR:HD1	2:E:478:TYR:O	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:199:LEU:O	2:F:200:VAL:HB	2.17	0.45
1:D:784:GLY:C	2:F:363:ARG:HG2	2.37	0.45
1:A:1005:VAL:O	1:A:1006:ARG:NH1	2.50	0.45
1:A:857:PRO:HB3	1:A:1062:GLU:CD	2.37	0.45
1:A:1069:ILE:HG23	1:A:1070:PRO:HG3	1.98	0.45
1:A:271:VAL:HG22	1:A:295:MET:CE	2.46	0.45
1:A:515:ILE:HG12	1:A:565:HIS:C	2.37	0.45
1:A:720:THR:HG22	1:A:721:ALA:H	1.82	0.45
1:A:79:LEU:HD13	1:A:1172:ASP:OD2	2.16	0.45
1:A:850:ILE:HD11	1:A:1099:TRP:CE2	2.51	0.45
1:A:869:ARG:HG2	1:A:870:VAL:CG2	2.37	0.45
2:B:115:TRP:CZ2	2:C:108:LYS:CD	2.98	0.45
2:B:196:CYS:O	2:B:199:LEU:N	2.43	0.45
2:B:243:THR:OG1	2:B:244:PRO:HD2	2.16	0.45
2:B:425:LEU:HD23	2:B:425:LEU:C	2.36	0.45
2:B:127:PRO:HG2	2:C:104:VAL:HB	1.97	0.45
2:C:264:ARG:HG2	2:C:270:PRO:HB2	1.98	0.45
2:C:308:ASP:CG	2:C:334:CYS:HA	2.37	0.45
2:C:78:PHE:CE2	2:C:350:LEU:CD2	2.99	0.45
2:C:372:LEU:CG	2:C:436:SER:HB2	2.46	0.45
2:C:456:ARG:HB3	2:C:463:LYS:O	2.16	0.45
2:C:476:ILE:HD13	2:C:477:LYS:H	1.79	0.45
2:E:285:LYS:O	2:E:304:TRP:CE3	2.70	0.45
1:D:586:THR:O	1:D:586:THR:CG2	2.65	0.45
2:B:447:THR:HA	2:B:450:ASN:CG	2.37	0.45
2:C:317:GLY:HA2	2:C:320:SER:CB	2.42	0.45
2:E:272:ASN:OD1	2:E:292:ASN:ND2	2.46	0.45
1:D:1126:ASP:HB2	1:D:1141:VAL:HG13	1.98	0.45
1:D:1140:LEU:HD23	1:D:1141:VAL:H	1.81	0.45
1:D:1231:LEU:HD21	1:D:1236:GLN:HE22	1.81	0.45
1:D:214:ALA:HB3	1:D:219:ALA:N	2.31	0.45
1:D:248:GLU:HB3	1:D:249:VAL:CA	2.07	0.45
1:D:279:ARG:HD2	1:D:841:LEU:CD1	2.46	0.45
1:D:271:VAL:HG22	1:D:295:MET:HB2	1.98	0.45
1:D:365:GLY:C	1:D:367:PRO:HD3	2.36	0.45
1:D:424:LEU:O	1:D:428:LEU:HD12	2.15	0.45
1:D:652:ILE:CG2	1:D:656:TYR:CE2	2.99	0.45
1:D:76:ILE:CG1	1:D:77:GLN:N	2.79	0.45
1:D:897:TRP:CZ3	1:D:1177:VAL:CG1	2.98	0.45
1:D:916:PHE:CD1	1:D:920:THR:O	2.69	0.45
2:E:125:VAL:HG22	2:E:241:TRP:CZ3	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:254:PHE:HA	2:E:257:ARG:NE	2.30	0.45
2:E:259:ARG:NH1	2:E:263:TRP:NE1	2.64	0.45
2:E:110:LEU:HA	2:E:378:LEU:HD23	1.97	0.45
2:F:76:ARG:O	2:F:102:LEU:HD13	2.16	0.45
2:F:131:LEU:O	2:F:182:ARG:HD3	2.15	0.45
2:F:399:CYS:SG	2:F:417:TYR:CE2	3.09	0.45
2:F:384:ALA:HA	2:F:414:TRP:HB3	1.98	0.45
2:F:413:VAL:HG12	2:F:415:PRO:CD	2.46	0.45
2:F:446:THR:C	2:F:450:ASN:H	2.19	0.45
2:F:78:PHE:O	2:F:99:PHE:HB3	2.17	0.45
1:A:1038:TRP:CZ3	1:A:1041:TRP:CG	3.04	0.45
1:A:1207:GLU:C	1:A:1210:TYR:CE1	2.90	0.45
1:A:242:ALA:O	1:A:245:ILE:HG23	2.17	0.45
1:A:269:HIS:C	1:A:294:THR:OG1	2.55	0.45
1:A:441:TRP:CZ2	1:A:874:LEU:HB3	2.51	0.45
1:A:577:CYS:HB3	1:A:578:PRO:HD3	1.87	0.45
1:A:591:LEU:CD1	1:A:594:LEU:HD11	2.47	0.45
1:A:652:ILE:O	1:A:656:TYR:CD2	2.70	0.45
2:B:184:ASN:C	2:B:186:LEU:N	2.69	0.45
2:B:285:LYS:O	2:B:305:ASN:N	2.49	0.45
2:B:69:LEU:HD21	2:B:351:TYR:CE1	2.51	0.45
2:C:454:HIS:HB3	2:C:466:MET:C	2.36	0.45
2:C:471:LEU:HA	2:C:474:PHE:CB	2.34	0.45
2:C:88:ARG:CB	2:C:88:ARG:CZ	2.94	0.45
2:E:280:ASP:N	2:E:304:TRP:CZ3	2.84	0.45
1:D:1103:SER:O	1:D:1106:VAL:N	2.50	0.45
1:D:887:VAL:N	1:D:1139:TYR:O	2.48	0.45
1:D:897:TRP:CH2	1:D:1173:LEU:HB3	2.52	0.45
1:D:1198:LYS:N	1:D:1204:THR:CG2	2.78	0.45
1:D:190:PRO:O	1:D:192:GLU:N	2.49	0.45
1:D:277:HIS:HB3	1:D:291:PHE:CD2	2.52	0.45
1:D:253:ALA:HB2	1:D:280:GLU:C	2.37	0.45
1:D:375:GLU:CD	1:D:379:LYS:HA	2.37	0.45
1:D:389:PHE:CA	1:D:393:MET:HG2	2.45	0.45
1:D:576:LEU:CD1	1:D:577:CYS:SG	3.05	0.45
1:D:597:ARG:HH12	1:D:725:PRO:N	2.14	0.45
1:D:793:GLU:O	1:D:797:MET:CG	2.64	0.45
1:D:992:LEU:HB2	1:D:997:LEU:HB2	1.98	0.45
2:E:109:ASN:HB2	2:E:378:LEU:C	2.37	0.45
2:E:322:LEU:O	2:E:323:HIS:C	2.52	0.45
2:E:372:LEU:O	2:E:458:ARG:CD	2.63	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:111:ALA:O	2:F:115:TRP:HB2	2.15	0.45
2:F:236:GLU:HA	2:F:341:ASP:CG	2.35	0.45
2:F:353:SER:HG	2:F:354:PHE:H	1.65	0.45
2:F:375:HIS:HE2	2:F:377:CYS:HG	1.64	0.45
2:F:455:LEU:CB	2:F:456:ARG:NE	2.74	0.45
1:A:107:LEU:HD13	1:A:107:LEU:N	2.32	0.45
1:A:240:SER:O	1:A:243:ASP:CG	2.54	0.45
1:A:288:ARG:CZ	1:A:290:ARG:HH21	2.29	0.45
1:A:386:ARG:CG	1:A:390:GLN:HE22	2.29	0.45
1:A:597:ARG:HE	1:A:597:ARG:HB2	1.28	0.45
1:A:788:GLY:O	1:A:792:LEU:N	2.49	0.45
2:B:78:PHE:CE1	2:B:103:GLY:N	2.85	0.45
2:B:301:GLU:HG2	2:B:303:LEU:HD12	1.99	0.45
2:B:336:LEU:CG	2:B:337:SER:N	2.79	0.45
2:B:77:HIS:CG	2:C:195:ASN:OD1	2.69	0.45
2:C:241:TRP:CG	2:C:243:THR:HG23	2.51	0.45
2:C:342:LEU:HD23	2:C:345:GLY:N	2.17	0.45
2:B:130:ALA:HB1	2:B:188:GLY:HA3	1.99	0.45
1:D:1040:LYS:HE3	1:D:1040:LYS:CA	2.46	0.45
1:D:944:GLU:CB	1:D:1071:ARG:HD3	2.44	0.45
1:D:1157:ASN:C	1:D:1161:ARG:HD2	2.36	0.45
1:D:1209:ARG:NH1	2:F:252:LEU:HD22	2.32	0.45
1:D:179:GLY:N	1:D:180:PRO:CD	2.79	0.45
1:D:251:THR:O	1:D:254:SER:N	2.49	0.45
1:D:466:LEU:O	1:D:470:ALA:HB3	2.17	0.45
1:D:635:PRO:HA	1:D:639:THR:HG1	1.77	0.45
1:D:652:ILE:HG12	1:D:655:LEU:HG	1.98	0.45
1:D:653:GLU:HA	1:D:656:TYR:CD2	2.51	0.45
1:D:439:GLN:HG3	1:D:836:LEU:H	1.82	0.45
1:D:877:MET:CE	1:D:1203:PRO:C	2.84	0.45
2:E:382:LYS:HE3	2:E:383:VAL:C	2.37	0.45
2:E:69:LEU:O	2:E:72:ILE:HB	2.17	0.45
2:F:193:TYR:CD1	2:F:242:PHE:CE2	3.04	0.45
2:F:322:LEU:HD22	2:F:324:GLY:N	2.32	0.45
2:F:78:PHE:CE2	2:F:350:LEU:CD2	2.99	0.45
2:F:363:ARG:HB3	2:F:363:ARG:HH11	1.77	0.45
2:F:399:CYS:O	2:F:400:GLN:C	2.55	0.45
1:A:1006:ARG:N	1:A:1006:ARG:HD2	2.31	0.45
1:A:1042:GLU:OE1	1:A:1042:GLU:HA	2.17	0.45
1:A:1109:LEU:CD1	1:A:1113:LEU:HD21	2.38	0.45
1:A:1131:ILE:CG2	1:A:1137:VAL:CA	2.84	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1158:LEU:HD11	1:A:1179:PHE:CE1	2.51	0.45
1:A:1188:CYS:C	1:A:1189:LEU:CD2	2.75	0.45
1:A:170:ALA:CB	1:A:175:TRP:CZ2	2.67	0.45
1:A:460:LYS:NZ	1:A:594:LEU:HD21	2.32	0.45
1:A:462:SER:HG	1:A:463:LEU:H	1.61	0.45
1:A:577:CYS:HB3	1:A:578:PRO:HD2	1.97	0.45
1:A:586:THR:O	1:A:587:PRO:C	2.54	0.45
1:A:612:LEU:HD21	1:A:614:TYR:C	2.36	0.45
1:A:606:THR:OG1	1:A:613:HIS:CA	2.65	0.45
1:A:607:TRP:HZ2	1:A:670:PRO:C	2.20	0.45
1:A:784:GLY:HA3	2:C:362:THR:C	2.36	0.45
1:A:304:LEU:O	1:A:850:ILE:HG21	2.17	0.45
1:A:892:ASP:CB	1:A:927:ARG:HH22	2.30	0.45
1:A:931:LEU:O	1:A:934:LYS:N	2.49	0.45
1:A:972:ARG:C	1:A:974:THR:H	2.20	0.45
2:B:261:GLN:N	2:B:264:ARG:HD3	2.29	0.45
2:B:320:SER:O	2:B:323:HIS:CD2	2.70	0.45
2:B:385:LEU:N	2:B:414:TRP:O	2.49	0.45
2:C:187:HIS:HA	2:C:190:LEU:CD1	2.46	0.45
2:C:197:LEU:O	2:C:202:LYS:HA	2.17	0.45
2:C:200:VAL:HG11	2:C:204:LEU:CD1	2.44	0.45
2:C:320:SER:HB3	2:C:321:LYS:CG	2.38	0.45
1:D:103:SER:OG	1:D:104:VAL:N	2.49	0.45
2:B:122:ARG:CB	2:B:124:GLN:HE21	2.26	0.45
1:A:984:GLN:O	1:A:987:ALA:HB3	2.17	0.45
1:D:1158:LEU:CG	1:D:1159:LEU:H	2.27	0.45
1:D:1200:PRO:CG	1:D:1200:PRO:O	2.65	0.45
1:D:877:MET:HE1	1:D:1203:PRO:CA	2.47	0.45
1:D:225:SER:HA	1:D:228:LEU:CD2	2.44	0.45
1:D:228:LEU:HG	1:D:229:VAL:H	1.82	0.45
1:D:196:VAL:HA	1:D:267:VAL:HG12	1.99	0.45
1:D:580:LEU:C	1:D:580:LEU:CD1	2.75	0.45
1:D:630:ASN:C	1:D:631:LEU:HD23	2.37	0.45
1:D:655:LEU:C	1:D:659:HIS:NE2	2.70	0.45
1:D:805:HIS:O	1:D:806:LYS:C	2.53	0.45
2:E:104:VAL:CG2	2:F:128:VAL:O	2.65	0.45
2:E:199:LEU:HG	2:E:200:VAL:H	1.80	0.45
2:E:407:LEU:HD21	2:E:412:SER:HA	1.93	0.45
2:E:425:LEU:C	2:E:426:GLU:OE1	2.55	0.45
2:F:124:GLN:C	2:F:125:VAL:CG1	2.85	0.45
2:F:452:LEU:O	2:F:453:ILE:CB	2.64	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1005:VAL:O	1:A:1006:ARG:C	2.54	0.45
1:A:1074:VAL:CG2	1:A:1098:ASN:HD21	2.30	0.45
1:A:145:LYS:CG	1:A:1118:TRP:CH2	2.92	0.45
1:A:298:HIS:O	1:A:299:MET:C	2.54	0.45
1:A:408:GLN:C	1:A:411:LEU:HG	2.36	0.45
1:A:436:PRO:HB3	1:A:880:ALA:H	1.81	0.45
1:A:800:PHE:CE2	1:A:867:PRO:CB	3.00	0.45
1:A:296:SER:CB	1:A:846:THR:OG1	2.64	0.45
1:A:851:THR:O	1:A:852:ARG:HG3	2.16	0.45
1:A:808:ILE:HG21	1:A:874:LEU:HD13	1.99	0.45
1:A:995:TYR:C	1:A:996:ARG:HG3	2.37	0.45
2:B:301:GLU:HA	2:B:340:GLY:CA	2.33	0.45
2:B:312:LEU:HA	2:B:312:LEU:HD12	1.71	0.45
2:B:338:VAL:O	2:B:339:ASN:ND2	2.40	0.45
2:C:200:VAL:HG11	2:C:204:LEU:CB	2.46	0.45
1:A:784:GLY:HA3	2:C:363:ARG:CD	2.47	0.45
1:A:633:LYS:CD	1:A:634:LEU:N	2.78	0.45
1:D:163:PRO:C	1:D:165:LYS:N	2.64	0.45
2:E:230:SER:OG	2:E:231:ILE:N	2.49	0.45
1:D:1154:GLN:HG2	1:D:1179:PHE:CD2	2.52	0.45
1:D:294:THR:CB	1:D:407:PHE:CE2	2.99	0.45
1:D:346:ASP:H	1:D:1040:LYS:CD	2.24	0.45
1:D:361:ARG:NH2	1:D:362:LEU:CG	2.79	0.45
1:D:461:LYS:H	1:D:461:LYS:HD2	1.82	0.45
1:D:482:ASP:CB	1:D:483:PRO:CD	2.95	0.45
1:D:815:TRP:CZ3	1:D:836:LEU:CB	2.99	0.45
2:E:196:CYS:O	2:E:197:LEU:C	2.55	0.45
2:E:214:CYS:SG	2:E:236:GLU:CB	3.05	0.45
2:E:258:HIS:O	2:E:261:GLN:HB2	2.17	0.45
2:E:322:LEU:HD12	2:E:332:VAL:HG11	1.98	0.45
2:E:91:LEU:HD22	2:E:91:LEU:N	2.32	0.45
2:F:389:ARG:NH2	2:F:390:GLY:HA3	2.32	0.45
2:F:431:LYS:CE	2:F:435:MET:HG2	2.47	0.45
2:F:69:LEU:HD12	2:F:69:LEU:O	2.17	0.45
2:F:72:ILE:O	2:F:75:ARG:O	2.35	0.45
1:A:1108:TYR:CE2	1:A:1112:MET:CE	2.99	0.45
1:A:1131:ILE:HG23	1:A:1137:VAL:HA	1.94	0.45
1:A:1108:TYR:CA	1:A:1164:PHE:HZ	2.29	0.45
1:A:197:PHE:HB3	1:A:277:HIS:HD2	1.79	0.45
1:A:296:SER:OG	1:A:297:MET:N	2.50	0.45
1:A:312:TRP:CH2	1:A:346:ASP:N	2.85	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:316:LYS:HB2	1:A:348:LEU:HD12	1.97	0.45
1:A:424:LEU:N	1:A:424:LEU:CD2	2.64	0.45
1:A:452:TYR:CD1	1:A:801:TRP:NE1	2.83	0.45
1:A:462:SER:C	1:A:466:LEU:CD2	2.81	0.45
1:A:559:LEU:CG	1:A:560:PRO:HD3	2.46	0.45
1:A:611:PRO:HG3	1:A:671:GLN:HG3	1.99	0.45
1:A:821:LEU:CD1	1:A:825:VAL:CB	2.92	0.45
1:A:869:ARG:O	1:A:870:VAL:C	2.55	0.45
1:A:875:LYS:HA	1:A:878:VAL:CG2	2.47	0.45
2:B:348:ALA:C	2:B:350:LEU:H	2.20	0.45
2:B:391:PRO:CD	2:B:392:THR:H	2.29	0.45
2:B:431:LYS:O	2:B:431:LYS:HG2	2.16	0.45
2:C:121:PHE:C	2:C:123:GLU:N	2.69	0.45
2:B:99:PHE:O	2:C:128:VAL:CG1	2.65	0.45
2:C:185:LEU:HD12	2:C:236:GLU:HB2	1.99	0.45
2:C:325:ARG:NH2	2:C:327:GLY:CA	2.80	0.45
2:C:484:ASN:O	2:C:485:VAL:C	2.55	0.45
1:D:104:VAL:CG1	1:D:105:GLU:N	2.79	0.45
2:F:146:ARG:HD2	2:F:228:VAL:HG21	1.98	0.45
2:B:405:GLU:HA	2:B:405:GLU:OE1	2.16	0.45
1:D:138:HIS:HA	1:D:141:LEU:CD1	2.47	0.45
1:D:208:CYS:SG	1:D:245:ILE:HG22	2.57	0.45
1:D:173:GLU:C	1:D:220:TRP:HB3	2.38	0.45
1:D:457:ARG:O	1:D:461:LYS:HD2	2.17	0.45
1:D:807:ARG:HH22	1:D:1061:LEU:CD1	2.30	0.45
1:D:822:PRO:CD	1:D:825:VAL:CG2	2.78	0.45
1:D:80:SER:C	1:D:83:LEU:HD22	2.37	0.45
1:D:438:ASN:HD21	1:D:882:PRO:HD3	1.82	0.45
1:D:972:ARG:C	1:D:974:THR:H	2.20	0.45
2:E:193:TYR:CE2	2:E:322:LEU:HB3	2.49	0.45
2:E:431:LYS:HD3	2:E:434:GLU:HG3	1.99	0.45
2:F:126:PHE:HB2	2:F:128:VAL:HG23	1.98	0.45
1:A:131:TYR:CD2	1:A:131:TYR:N	2.79	0.45
1:A:406:VAL:CG2	1:A:407:PHE:HD2	2.30	0.45
1:A:614:TYR:HE2	1:A:620:TRP:CE3	2.35	0.45
1:A:657:ARG:O	1:A:658:LYS:C	2.53	0.45
1:A:828:HIS:O	1:A:830:ASP:OD2	2.36	0.45
1:A:875:LYS:CA	1:A:878:VAL:HG23	2.47	0.45
2:B:106:LEU:O	2:B:109:ASN:CB	2.65	0.45
2:B:250:GLN:O	2:B:252:LEU:N	2.42	0.45
2:B:114:TRP:HD1	2:B:262:TRP:CE3	2.35	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:114:TRP:HB2	2:C:262:TRP:CZ2	2.51	0.45
2:C:275:SER:CB	2:C:287:ASN:ND2	2.75	0.45
2:C:306:LEU:N	2:C:335:VAL:CG1	2.69	0.45
2:C:353:SER:HB3	2:C:373:LYS:O	2.17	0.45
2:C:476:ILE:HD13	2:C:477:LYS:HG3	1.99	0.45
2:F:317:GLY:O	2:F:318:ASN:C	2.55	0.45
1:A:98:ALA:O	1:A:102:ARG:CG	2.57	0.45
1:A:979:ALA:O	1:A:980:GLU:C	2.55	0.45
1:D:539:PHE:O	1:D:540:GLN:HB2	2.17	0.45
1:D:1184:ASP:OD1	1:D:1184:ASP:N	2.48	0.45
1:D:429:GLU:CG	1:D:1117:LYS:HE2	2.43	0.44
1:D:1139:TYR:CD1	1:D:1153:LEU:HD13	2.53	0.44
1:D:189:ILE:CG2	1:D:258:GLN:CB	2.89	0.44
1:D:282:TYR:CD1	1:D:282:TYR:N	2.84	0.44
1:D:295:MET:CE	1:D:354:ASN:HA	2.46	0.44
1:D:374:ARG:CD	1:D:392:LEU:CD2	2.87	0.44
1:D:375:GLU:C	1:D:378:VAL:H	2.19	0.44
1:D:416:GLU:C	1:D:418:CYS:N	2.69	0.44
1:D:612:LEU:HD23	1:D:620:TRP:O	2.17	0.44
1:D:743:ASP:CB	1:D:744:ILE:HD13	2.46	0.44
1:D:733:HIS:O	1:D:750:PHE:O	2.35	0.44
1:D:829:PRO:O	1:D:829:PRO:CD	2.66	0.44
1:D:770:PHE:CE1	1:D:968:GLN:C	2.91	0.44
2:F:307:GLY:N	2:F:335:VAL:CG1	2.80	0.44
2:F:344:ARG:CG	2:F:347:LEU:HD22	2.47	0.44
2:F:375:HIS:NE2	2:F:377:CYS:SG	2.89	0.44
2:F:429:TYR:O	2:F:432:TYR:HB2	2.17	0.44
2:F:471:LEU:O	2:F:472:LYS:C	2.56	0.44
2:F:474:PHE:CD2	2:F:475:LEU:HD23	2.53	0.44
1:A:1030:ARG:CG	1:A:1040:LYS:NZ	2.62	0.44
1:A:252:GLY:O	1:A:256:PRO:HB3	2.17	0.44
1:A:348:LEU:HD13	1:A:350:ILE:CD1	2.45	0.44
1:A:371:LYS:C	1:A:372:GLU:CG	2.79	0.44
1:A:412:PRO:O	1:A:416:GLU:HG3	2.17	0.44
1:A:444:TYR:O	1:A:448:ALA:HB2	2.17	0.44
1:A:498:LYS:CG	1:A:518:ALA:HB3	2.47	0.44
1:A:83:LEU:O	1:A:86:GLN:N	2.50	0.44
1:A:897:TRP:O	1:A:900:ALA:CB	2.53	0.44
1:A:938:THR:O	1:A:939:VAL:CG2	2.65	0.44
2:B:214:CYS:HB3	2:B:236:GLU:CD	2.38	0.44
2:B:243:THR:O	2:B:334:CYS:HB2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:76:ARG:O	2:B:77:HIS:O	2.35	0.44
2:C:251:TRP:CZ3	2:C:254:PHE:CG	3.04	0.44
2:C:431:LYS:CE	2:C:435:MET:HG2	2.44	0.44
2:C:70:LEU:O	2:C:73:CYS:N	2.46	0.44
2:C:83:LYS:O	2:C:86:LEU:HB2	2.17	0.44
2:C:382:LYS:HD3	2:C:411:ILE:CD1	2.46	0.44
2:B:453:ILE:HD13	2:B:466:MET:O	2.17	0.44
2:F:326:ASP:CG	2:F:329:LYS:CB	2.86	0.44
2:F:229:LYS:O	2:F:230:SER:HB2	2.17	0.44
1:D:1067:SER:HB3	1:D:1073:PRO:HD2	1.98	0.44
1:D:1099:TRP:HE3	1:D:1099:TRP:HA	1.82	0.44
1:D:1108:TYR:O	1:D:1109:LEU:C	2.54	0.44
1:D:1131:ILE:HG13	1:D:1138:ARG:HB2	1.99	0.44
1:D:1155:ILE:O	1:D:1157:ASN:N	2.50	0.44
1:D:1180:PHE:O	1:D:1181:SER:C	2.56	0.44
1:D:887:VAL:CG1	1:D:1183:VAL:HG11	2.47	0.44
1:D:223:TRP:CZ3	1:D:250:PRO:HB3	2.49	0.44
1:D:267:VAL:CG2	1:D:291:PHE:CE2	2.99	0.44
1:D:486:TRP:CE3	1:D:598:VAL:HG21	2.53	0.44
1:D:542:ASP:O	1:D:546:ARG:CB	2.65	0.44
1:D:576:LEU:H	1:D:579:ARG:HD2	1.83	0.44
1:D:840:ILE:C	1:D:841:LEU:HD23	2.37	0.44
2:E:318:ASN:C	2:E:320:SER:N	2.70	0.44
2:E:380:PRO:CG	2:E:438:LEU:CD1	2.96	0.44
2:F:389:ARG:CD	2:F:390:GLY:N	2.81	0.44
2:F:484:ASN:O	2:F:485:VAL:C	2.56	0.44
2:E:199:LEU:CB	2:F:77:HIS:NE2	2.74	0.44
1:A:773:LYS:HZ2	1:A:1057:MET:HE1	1.83	0.44
1:A:1048:ALA:CB	1:A:1092:PHE:CE1	3.00	0.44
1:A:273:PHE:N	1:A:273:PHE:CD2	2.85	0.44
1:A:316:LYS:HZ2	1:A:350:ILE:HG21	1.82	0.44
1:A:392:LEU:CG	1:A:395:TYR:HB2	2.47	0.44
1:A:490:TRP:HH2	1:A:597:ARG:O	1.99	0.44
1:A:630:ASN:HD21	1:A:632:ALA:CB	2.30	0.44
1:A:659:HIS:CD2	1:A:720:THR:CG2	3.00	0.44
1:A:724:GLY:HA2	1:A:725:PRO:HA	1.55	0.44
2:B:289:LEU:N	2:B:301:GLU:O	2.48	0.44
2:B:105:GLU:HG3	2:B:380:PRO:O	2.17	0.44
2:C:214:CYS:SG	2:C:236:GLU:CG	3.05	0.44
2:C:243:THR:HB	2:C:247:THR:HG21	1.99	0.44
2:C:348:ALA:O	2:C:351:TYR:HB2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:83:LYS:N	2:C:86:LEU:HD22	2.32	0.44
2:C:320:SER:CB	2:C:321:LYS:HG3	2.35	0.44
1:D:529:ASP:O	1:D:530:LEU:CB	2.57	0.44
2:E:311:LEU:H	2:E:313:HIS:HB3	1.82	0.44
1:A:1035:LYS:HG3	1:A:1039:LYS:HE2	0.80	0.44
2:C:280:ASP:C	2:C:282:GLU:H	2.21	0.44
1:D:1015:THR:HB	1:D:1016:GLU:H	1.19	0.44
1:D:1046:GLU:O	1:D:1049:TRP:CD1	2.70	0.44
1:D:886:LEU:N	1:D:1146:ARG:NH1	2.66	0.44
1:D:127:LEU:HD23	1:D:128:PRO:HD2	1.98	0.44
1:D:220:TRP:NE1	1:D:221:TYR:HD2	2.14	0.44
1:D:221:TYR:O	1:D:222:SER:CB	2.65	0.44
1:D:257:THR:C	1:D:258:GLN:CG	2.85	0.44
1:D:711:ALA:O	1:D:712:VAL:HB	2.17	0.44
1:D:823:ARG:NH1	1:D:824:ALA:CB	2.80	0.44
1:D:992:LEU:C	1:D:996:ARG:CD	2.86	0.44
2:E:293:PHE:O	2:E:294:PRO:C	2.56	0.44
2:F:196:CYS:HB2	2:F:206:TYR:OH	2.18	0.44
2:F:239:LEU:N	2:F:338:VAL:HG13	2.22	0.44
2:F:385:LEU:HA	2:F:385:LEU:HD23	1.75	0.44
2:F:407:LEU:HD13	2:F:408:GLU:OE2	2.17	0.44
1:D:546:ARG:NH2	2:F:408:GLU:CD	2.71	0.44
1:A:907:PHE:CZ	1:A:1001:GLY:CA	2.99	0.44
1:A:1068:ASP:N	1:A:1071:ARG:HG2	2.02	0.44
1:A:129:PRO:O	1:A:131:TYR:CE2	2.70	0.44
1:A:196:VAL:O	1:A:401:TRP:CH2	2.61	0.44
1:A:210:THR:HA	1:A:224:CYS:HB2	1.95	0.44
1:A:266:VAL:HG22	1:A:267:VAL:H	1.82	0.44
1:A:470:ALA:O	1:A:474:LEU:HD13	2.17	0.44
1:A:549:LEU:HA	1:A:549:LEU:HD23	1.54	0.44
1:A:656:TYR:HA	1:A:659:HIS:CD2	2.51	0.44
1:A:439:GLN:HG2	1:A:836:LEU:H	1.78	0.44
2:B:385:LEU:O	2:B:415:PRO:HA	2.17	0.44
2:C:450:ASN:OD1	2:C:453:ILE:HD12	2.17	0.44
1:D:515:ILE:CG2	1:D:565:HIS:CG	3.00	0.44
2:C:87:SER:O	2:C:90:SER:N	2.49	0.44
2:F:59:ALA:N	2:F:60:PRO:HD2	2.32	0.44
1:D:1111:LEU:HD12	1:D:1111:LEU:N	2.32	0.44
1:D:1111:LEU:CD1	1:D:1111:LEU:N	2.80	0.44
1:D:1166:TYR:CE2	1:D:1171:ASN:HA	2.53	0.44
1:D:249:VAL:HG11	1:D:251:THR:CG2	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:305:SER:HB2	1:D:308:GLN:CD	2.38	0.44
1:D:803:ASN:OD1	1:D:803:ASN:N	2.50	0.44
1:D:958:GLY:O	1:D:959:GLN:C	2.56	0.44
2:E:94:GLY:CA	2:E:96:HIS:CE1	2.97	0.44
2:F:405:GLU:OE1	2:F:406:LEU:HD12	2.16	0.44
2:F:407:LEU:HD23	2:F:407:LEU:O	2.17	0.44
1:A:1133:ILE:HG23	1:A:1133:ILE:O	2.18	0.44
1:A:1146:ARG:HH11	1:A:1146:ARG:HG3	1.82	0.44
1:A:1209:ARG:HH22	2:C:256:LEU:HG	1.82	0.44
1:A:215:ILE:HD12	1:A:401:TRP:CE3	2.51	0.44
1:A:248:GLU:O	1:A:249:VAL:C	2.54	0.44
1:A:316:LYS:CD	1:A:350:ILE:CG1	2.85	0.44
1:A:477:GLU:CG	1:A:478:ARG:N	2.32	0.44
1:A:497:GLN:O	1:A:497:GLN:OE1	2.35	0.44
1:A:515:ILE:HG13	1:A:568:GLY:O	2.17	0.44
1:A:464:MET:SD	1:A:578:PRO:O	2.76	0.44
1:A:579:ARG:NH1	1:A:579:ARG:HG2	2.33	0.44
1:A:580:LEU:HD23	1:A:596:MET:SD	2.58	0.44
1:A:612:LEU:HD22	1:A:620:TRP:CA	2.45	0.44
1:A:801:TRP:CE2	1:A:805:HIS:HB2	2.53	0.44
1:A:862:ALA:HB1	1:A:875:LYS:CE	2.46	0.44
1:A:945:HIS:NE2	1:A:969:PHE:CG	2.85	0.44
1:A:993:ARG:HG3	1:A:994:TRP:CD1	2.52	0.44
2:C:185:LEU:CD2	2:C:214:CYS:SG	3.05	0.44
2:C:241:TRP:CZ3	2:C:255:TRP:CZ2	3.06	0.44
2:C:92:LEU:CD2	2:C:344:ARG:HE	2.28	0.44
2:C:375:HIS:CD2	2:C:377:CYS:H	2.36	0.44
1:A:472:GLN:CG	2:C:460:THR:HB	2.41	0.44
1:A:504:LYS:O	1:A:505:LYS:HG3	2.17	0.44
1:A:646:VAL:O	1:A:647:CYS:HB2	2.17	0.44
1:A:737:GLY:N	1:A:750:PHE:HE1	2.16	0.44
1:A:980:GLU:O	1:A:984:GLN:HG3	2.18	0.44
2:B:251:TRP:HA	2:B:251:TRP:HE3	1.83	0.44
1:D:1027:LYS:CB	1:D:1091:GLU:CA	2.85	0.44
1:D:1129:PHE:HB2	1:D:1138:ARG:O	2.17	0.44
1:D:897:TRP:CH2	1:D:1173:LEU:HD22	2.53	0.44
1:D:876:ALA:O	1:D:1189:LEU:HD23	2.18	0.44
1:D:1209:ARG:NH1	2:F:252:LEU:CD1	2.64	0.44
1:D:1224:ILE:HD11	1:D:1228:LYS:HE3	2.00	0.44
1:D:192:GLU:HG2	1:D:193:ARG:C	2.38	0.44
1:D:301:ILE:HG13	1:D:415:LEU:HD12	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:384:ASP:O	1:D:388:ASN:OD1	2.36	0.44
1:D:549:LEU:C	1:D:551:LYS:HD2	2.38	0.44
1:D:726:LYS:O	1:D:727:ASP:HB3	2.17	0.44
1:D:786:ALA:O	1:D:787:SER:HB3	2.17	0.44
1:D:826:ILE:O	1:D:829:PRO:HB3	2.18	0.44
2:E:109:ASN:CB	2:E:378:LEU:O	2.66	0.44
2:E:382:LYS:CD	2:E:383:VAL:H	2.31	0.44
2:E:382:LYS:HG3	2:E:383:VAL:H	1.68	0.44
2:E:384:ALA:HB1	2:E:385:LEU:H	1.13	0.44
2:E:441:VAL:HG22	2:E:455:LEU:HD13	2.00	0.44
2:F:129:ASP:OD1	2:F:131:LEU:HD13	2.18	0.44
2:F:263:TRP:CZ3	2:F:349:TYR:CE2	3.06	0.44
2:F:394:GLU:N	2:F:394:GLU:OE1	2.51	0.44
2:F:426:GLU:O	2:F:429:TYR:HB2	2.17	0.44
2:F:444:THR:O	2:F:447:THR:HG21	2.12	0.44
2:F:454:HIS:CD2	2:F:468:ILE:HG13	2.53	0.44
1:A:1014:ARG:O	1:A:1014:ARG:HG2	2.17	0.44
1:A:247:LEU:HD11	1:A:251:THR:HG21	1.96	0.44
1:A:484:TRP:O	1:A:488:LEU:CG	2.57	0.44
1:A:515:ILE:HG12	1:A:565:HIS:HA	1.99	0.44
1:A:467:ALA:HB2	1:A:602:LEU:HD13	2.00	0.44
1:A:756:ASP:OD1	1:A:770:PHE:HZ	2.00	0.44
1:A:762:VAL:HG23	1:A:763:GLY:N	2.32	0.44
1:A:893:SER:O	1:A:894:GLN:C	2.55	0.44
2:B:79:LEU:C	2:B:100:GLY:HA3	2.38	0.44
2:B:461:THR:CG2	2:B:463:LYS:NZ	2.80	0.44
2:B:100:GLY:CA	2:C:128:VAL:HG11	2.47	0.44
2:C:395:LEU:CD1	2:C:444:THR:CA	2.90	0.44
2:C:449:GLU:H	2:C:451:GLY:N	2.15	0.44
2:C:459:ASP:O	2:C:460:THR:OG1	2.32	0.44
2:F:138:LEU:HB3	2:F:139:LEU:H	1.57	0.44
1:D:1027:LYS:HD3	1:D:1027:LYS:N	2.33	0.44
1:D:1118:TRP:NE1	1:D:1122:GLU:OE1	2.51	0.44
1:D:1163:MET:CE	1:D:1167:LYS:CG	2.95	0.44
1:D:368:PRO:HA	1:D:369:LEU:HA	1.41	0.44
1:D:597:ARG:O	1:D:600:PRO:HD2	2.17	0.44
1:D:598:VAL:N	1:D:616:GLU:OE1	2.49	0.44
1:D:801:TRP:CD2	1:D:870:VAL:HG12	2.53	0.44
1:D:83:LEU:C	1:D:87:ILE:O	2.56	0.44
1:D:985:MET:O	1:D:986:TYR:C	2.56	0.44
2:E:386:ASP:HB3	2:E:441:VAL:O	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:241:TRP:HE3	2:F:336:LEU:CD1	2.30	0.44
2:F:259:ARG:CZ	2:F:339:ASN:HD21	2.31	0.44
2:F:79:LEU:HD13	2:F:79:LEU:HA	1.72	0.44
1:A:1026:ARG:CA	1:A:1029:GLN:NE2	2.80	0.44
1:A:1041:TRP:CD1	1:A:1043:VAL:CB	2.97	0.44
1:A:1146:ARG:HH22	1:A:1186:ASP:H	1.65	0.44
1:A:652:ILE:O	1:A:655:LEU:HD12	2.17	0.44
1:A:655:LEU:HA	1:A:713:PRO:HB3	2.00	0.44
1:A:905:ALA:CA	1:A:910:MET:HB3	2.47	0.44
1:A:933:SER:HB2	1:A:943:ARG:NH1	2.33	0.44
1:A:934:LYS:HE3	1:A:934:LYS:HB2	1.77	0.44
2:B:132:HIS:CD2	2:B:182:ARG:NH1	2.85	0.44
2:B:259:ARG:HD2	2:B:262:TRP:CB	2.48	0.44
2:C:199:LEU:HD12	2:C:200:VAL:N	2.32	0.44
1:A:1208:ARG:CZ	2:C:277:ASP:OD1	2.65	0.44
2:C:483:LYS:O	2:C:485:VAL:CG1	2.66	0.44
1:D:518:ALA:O	1:D:521:PRO:HD2	2.17	0.44
2:B:217:PRO:HB2	2:B:231:ILE:HB	1.97	0.44
1:A:973:LEU:HD13	1:A:977:GLU:OE1	2.18	0.44
1:D:1190:ARG:NH1	1:D:1195:MET:SD	2.90	0.44
1:D:1200:PRO:O	1:D:1203:PRO:CD	2.40	0.44
1:D:1204:THR:C	1:D:1207:GLU:H	2.21	0.44
1:D:120:LEU:N	1:D:121:PRO:HD2	2.33	0.44
1:D:127:LEU:HA	1:D:128:PRO:HD3	1.56	0.44
1:D:143:ALA:HA	1:D:146:GLN:HG3	1.99	0.44
1:D:225:SER:O	1:D:229:VAL:HG22	2.18	0.44
1:D:416:GLU:HG3	1:D:416:GLU:H	1.66	0.44
1:D:448:ALA:HB1	1:D:870:VAL:CB	2.37	0.44
1:D:728:THR:HG22	1:D:729:GLN:N	2.32	0.44
2:E:203:ARG:NH2	2:E:203:ARG:CG	2.79	0.44
2:E:299:LEU:HD23	2:E:299:LEU:C	2.38	0.44
2:E:325:ARG:HA	2:E:331:VAL:HG12	1.98	0.44
2:E:437:ILE:CG2	2:E:439:PHE:O	2.66	0.44
2:E:90:SER:OG	2:E:96:HIS:N	2.51	0.44
2:F:106:LEU:O	2:F:107:ARG:C	2.53	0.44
2:F:128:VAL:HG12	2:F:129:ASP:H	1.83	0.44
2:F:78:PHE:HZ	2:F:346:MET:CE	2.29	0.44
1:D:784:GLY:O	2:F:363:ARG:HG2	2.17	0.44
1:A:1064:ILE:C	1:A:1066:THR:H	2.21	0.44
1:A:1179:PHE:O	1:A:1220:ILE:CD1	2.65	0.44
1:A:184:ALA:O	1:A:185:VAL:CG1	2.58	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:188:ALA:O	1:A:189:ILE:CG1	2.65	0.44
1:A:192:GLU:HB3	1:A:195:LEU:HD21	1.95	0.44
1:A:316:LYS:NZ	1:A:350:ILE:CG1	2.80	0.44
1:A:316:LYS:CB	1:A:348:LEU:HD11	2.47	0.44
1:A:515:ILE:HG12	1:A:565:HIS:CA	2.48	0.44
1:A:623:LEU:H	1:A:623:LEU:CD2	2.29	0.44
1:A:630:ASN:O	1:A:631:LEU:HD23	2.18	0.44
1:A:792:LEU:O	1:A:796:LYS:CG	2.66	0.44
1:A:70:ARG:HD3	1:A:81:ARG:HB3	1.99	0.44
1:A:901:VAL:HA	1:A:904:ASP:OD2	2.18	0.44
1:A:992:LEU:HD12	1:A:993:ARG:N	2.28	0.44
2:B:132:HIS:CG	2:B:182:ARG:CZ	3.01	0.44
2:B:259:ARG:HH11	2:B:259:ARG:CB	2.29	0.44
2:B:343:ASP:CG	2:B:344:ARG:HH21	2.18	0.44
2:B:423:SER:O	2:B:424:SER:HB3	2.17	0.44
2:C:200:VAL:HB	2:C:204:LEU:HD22	1.99	0.44
2:C:254:PHE:HE2	2:C:255:TRP:CE2	2.36	0.44
2:C:293:PHE:HB3	2:C:294:PRO:CD	2.47	0.44
2:C:363:ARG:NH1	2:C:363:ARG:CB	2.81	0.44
2:C:387:VAL:CG1	2:C:395:LEU:O	2.66	0.44
1:D:515:ILE:O	1:D:565:HIS:CE1	2.70	0.44
2:B:277:ASP:HA	2:B:287:ASN:ND2	2.33	0.44
1:A:741:ASP:O	1:A:742:VAL:CB	2.50	0.44
1:D:1019:TRP:HE3	1:D:1168:LEU:CD1	2.30	0.44
1:D:175:TRP:CZ2	1:D:183:GLU:CG	3.01	0.44
1:D:195:LEU:C	1:D:267:VAL:HG12	2.38	0.44
1:D:242:ALA:O	1:D:246:PRO:CD	2.57	0.44
1:D:269:HIS:C	1:D:294:THR:CG2	2.79	0.44
1:D:279:ARG:NH1	1:D:841:LEU:HA	2.33	0.44
1:D:293:ASP:OD2	1:D:294:THR:N	2.51	0.44
1:D:482:ASP:HB2	1:D:483:PRO:CD	2.46	0.44
1:D:547:ALA:C	1:D:551:LYS:CD	2.85	0.44
1:D:728:THR:CG2	1:D:741:ASP:OD2	2.64	0.44
1:D:790:ARG:C	1:D:792:LEU:N	2.71	0.44
1:D:866:ARG:CB	1:D:867:PRO:HD3	2.29	0.44
1:D:870:VAL:HG22	1:D:874:LEU:HD22	2.00	0.44
1:D:927:ARG:HH11	1:D:927:ARG:HG3	1.83	0.44
2:E:125:VAL:HG22	2:E:241:TRP:HZ3	1.82	0.44
2:E:185:LEU:HD12	2:E:237:ALA:O	2.17	0.44
2:E:128:VAL:CB	2:E:208:LEU:HB3	2.35	0.44
2:E:215:PHE:C	2:E:216:HIS:HD2	2.21	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:335:VAL:CG2	2:E:336:LEU:H	2.31	0.44
2:E:404:ASN:C	2:E:406:LEU:H	2.21	0.44
2:F:270:PRO:O	2:F:273:PHE:HD2	2.01	0.44
2:F:451:GLY:O	2:F:452:LEU:CB	2.55	0.44
2:F:476:ILE:HD13	2:F:477:LYS:H	1.82	0.44
1:A:1065:ALA:O	1:A:1071:ARG:HB3	2.17	0.44
1:A:106:HIS:C	1:A:108:GLN:H	2.20	0.44
1:A:79:LEU:CD1	1:A:1172:ASP:OD2	2.66	0.44
1:A:170:ALA:HB1	1:A:175:TRP:CH2	2.47	0.44
1:A:617:ARG:HB2	1:A:725:PRO:HD3	2.00	0.44
1:A:869:ARG:HB3	1:A:872:SER:OG	2.18	0.44
1:A:930:ASP:C	1:A:933:SER:HB3	2.39	0.44
1:A:967:MET:HG2	1:A:978:ALA:HB2	1.99	0.44
1:A:967:MET:SD	1:A:978:ALA:HB2	2.57	0.44
2:B:114:TRP:C	2:B:116:THR:N	2.69	0.44
2:B:236:GLU:CG	2:B:341:ASP:HB3	2.48	0.44
2:C:331:VAL:CG1	2:C:332:VAL:N	2.79	0.44
2:C:348:ALA:O	2:C:352:ASP:N	2.48	0.44
2:C:351:TYR:O	2:C:355:GLN:HB2	2.09	0.44
2:C:361:PHE:HD2	2:C:362:THR:CG2	2.30	0.44
1:D:585:TRP:N	1:D:585:TRP:CE3	2.86	0.44
2:B:468:ILE:CG2	2:B:468:ILE:O	2.62	0.44
1:A:634:LEU:CD1	1:A:638:THR:HG21	2.47	0.44
2:C:367:LEU:C	2:C:367:LEU:HD12	2.38	0.44
2:B:229:LYS:N	2:B:229:LYS:CD	2.78	0.44
1:D:873:GLU:HB2	1:D:1203:PRO:HB3	2.00	0.44
1:D:404:HIS:O	1:D:408:GLN:CB	2.66	0.44
1:D:601:LYS:CE	1:D:721:ALA:HB1	2.32	0.44
1:D:768:LYS:CA	1:D:772:PRO:CD	2.96	0.44
1:D:904:ASP:OD2	1:D:915:ALA:CB	2.63	0.44
1:D:914:THR:HA	1:D:918:TRP:CB	2.48	0.44
2:F:342:LEU:HD23	2:F:342:LEU:C	2.38	0.44
1:A:1030:ARG:NH2	1:A:1044:VAL:CG1	2.81	0.44
1:A:1179:PHE:HB3	1:A:1220:ILE:CG1	2.31	0.44
1:A:195:LEU:HA	1:A:216:SER:OG	2.18	0.44
1:A:241:PRO:C	1:A:244:LEU:HB2	2.38	0.44
1:A:493:GLN:O	1:A:495:PHE:CD1	2.71	0.44
1:A:494:GLU:HB2	1:A:572:TRP:CE2	2.52	0.44
1:A:612:LEU:HD12	1:A:612:LEU:C	2.38	0.44
1:A:730:PRO:CB	1:A:760:CYS:O	2.66	0.44
1:A:789:PRO:O	1:A:793:GLU:CB	2.65	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:950:ASN:O	1:A:951:TYR:O	2.35	0.44
1:A:963:GLU:HA	1:A:966:LEU:HG	1.99	0.44
2:B:79:LEU:HA	2:B:100:GLY:CA	2.48	0.44
2:B:206:TYR:CE1	2:B:242:PHE:HZ	2.36	0.44
2:B:418:LEU:O	2:B:418:LEU:HD23	2.18	0.44
2:C:115:TRP:CD1	2:C:119:VAL:HG21	2.53	0.44
2:C:204:LEU:HG	2:C:206:TYR:CG	2.53	0.44
1:A:635:PRO:C	1:A:637:GLY:N	2.71	0.44
1:A:97:GLU:O	1:A:98:ALA:C	2.54	0.44
2:F:279:GLN:NE2	2:F:279:GLN:HA	2.33	0.44
1:D:119:PRO:HG2	1:D:120:LEU:HD22	1.99	0.43
1:D:166:PRO:HA	1:D:167:PRO:HD3	1.47	0.43
1:D:247:LEU:N	1:D:815:TRP:HD1	2.14	0.43
1:D:553:LYS:HD3	2:F:468:ILE:HG21	1.99	0.43
1:D:588:GLY:N	1:D:806:LYS:HZ1	2.16	0.43
1:D:776:ASP:C	1:D:776:ASP:OD1	2.57	0.43
1:D:79:LEU:HD11	1:D:1173:LEU:C	2.33	0.43
1:D:80:SER:N	1:D:83:LEU:HD11	2.33	0.43
1:D:770:PHE:HE1	1:D:968:GLN:CA	2.30	0.43
1:D:990:LYS:HE2	1:D:1000:GLU:OE2	2.14	0.43
2:E:235:THR:H	2:E:343:ASP:CG	2.22	0.43
2:E:72:ILE:CD1	2:E:72:ILE:N	2.80	0.43
2:F:191:GLU:CD	2:F:315:TYR:HH	2.21	0.43
2:F:251:TRP:HZ3	2:F:254:PHE:CG	2.35	0.43
2:F:253:ASP:OD1	2:F:256:LEU:HD23	2.18	0.43
2:F:387:VAL:HG11	2:F:399:CYS:HB2	1.98	0.43
2:F:415:PRO:HA	2:F:417:TYR:HE2	1.82	0.43
2:F:471:LEU:CA	2:F:474:PHE:CD2	2.91	0.43
2:F:70:LEU:O	2:F:73:CYS:N	2.47	0.43
1:A:1195:MET:O	1:A:1196:ASP:CG	2.56	0.43
1:A:1218:LEU:O	1:A:1219:ASP:CB	2.46	0.43
1:A:480:LYS:HA	1:A:484:TRP:HD1	1.82	0.43
1:A:486:TRP:CZ3	1:A:598:VAL:CG2	2.96	0.43
1:A:498:LYS:HG2	1:A:518:ALA:H	1.83	0.43
1:A:560:PRO:CB	2:C:452:LEU:HD21	2.47	0.43
1:A:80:SER:H	1:A:83:LEU:CD1	2.08	0.43
1:A:877:MET:HA	1:A:877:MET:HE3	2.00	0.43
2:B:182:ARG:CD	2:B:214:CYS:HA	2.48	0.43
2:B:461:THR:HG23	2:B:461:THR:O	2.18	0.43
2:C:262:TRP:HZ3	2:C:263:TRP:CZ2	2.35	0.43
2:C:294:PRO:HG3	2:C:355:GLN:NE2	2.32	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:346:MET:CE	2:C:347:LEU:HD12	2.48	0.43
2:C:444:THR:N	2:C:447:THR:HG21	2.33	0.43
2:C:90:SER:HB2	2:C:95:CYS:SG	2.59	0.43
2:B:471:LEU:HD12	2:B:474:PHE:HD2	1.83	0.43
2:F:326:ASP:OD1	2:F:329:LYS:CG	2.56	0.43
2:F:186:LEU:O	2:F:187:HIS:C	2.56	0.43
1:D:1010:LEU:H	1:D:1011:PRO:CD	2.28	0.43
1:D:768:LYS:HZ2	1:D:1064:ILE:HG22	1.83	0.43
1:D:76:ILE:HD11	1:D:1174:PRO:N	2.33	0.43
1:D:1195:MET:C	1:D:1196:ASP:O	2.49	0.43
1:D:1151:LEU:CG	1:D:1223:ILE:HD11	2.48	0.43
1:D:180:PRO:O	1:D:181:GLU:OE2	2.36	0.43
1:D:175:TRP:CB	1:D:185:VAL:HG22	2.47	0.43
1:D:265:LEU:HD21	1:D:267:VAL:CB	2.48	0.43
1:D:353:VAL:HG23	1:D:354:ASN:OD1	2.19	0.43
1:D:363:TYR:O	1:D:364:VAL:CB	2.65	0.43
1:D:424:LEU:HD23	1:D:424:LEU:H	1.82	0.43
1:D:437:VAL:HG13	1:D:441:TRP:HB2	1.99	0.43
1:D:592:LEU:HA	1:D:595:GLN:HG2	1.93	0.43
1:D:779:LEU:HD22	1:D:780:GLN:CA	2.46	0.43
1:D:826:ILE:CB	1:D:831:TYR:HE2	2.26	0.43
1:D:444:TYR:CZ	1:D:877:MET:CE	3.01	0.43
1:D:878:VAL:O	1:D:879:GLN:HB3	2.18	0.43
1:D:950:ASN:O	1:D:953:ARG:HB2	2.18	0.43
1:D:954:ILE:HG22	1:D:955:TYR:N	2.33	0.43
1:D:989:THR:C	1:D:998:SER:HB3	2.38	0.43
2:E:100:GLY:O	2:E:104:VAL:N	2.48	0.43
2:E:131:LEU:O	2:E:182:ARG:NH2	2.47	0.43
2:E:289:LEU:C	2:E:290:TYR:CG	2.91	0.43
2:E:344:ARG:C	2:E:346:MET:N	2.70	0.43
2:E:406:LEU:HB2	2:E:413:VAL:HG23	2.00	0.43
2:E:406:LEU:HB2	2:E:413:VAL:CG2	2.48	0.43
2:E:448:LEU:HD23	2:E:449:GLU:CA	2.48	0.43
2:F:245:PRO:CG	2:F:246:ARG:N	2.81	0.43
2:F:331:VAL:CG1	2:F:332:VAL:N	2.78	0.43
2:F:431:LYS:HZ3	2:F:435:MET:CG	2.32	0.43
2:F:386:ASP:OD2	2:F:441:VAL:N	2.51	0.43
1:A:1046:GLU:CD	1:A:1049:TRP:CD1	2.91	0.43
1:A:1181:SER:O	1:A:1182:ALA:HB2	2.18	0.43
1:A:1227:THR:O	1:A:1235:SER:OG	2.34	0.43
1:A:267:VAL:CG2	1:A:291:PHE:CE2	3.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:402:ALA:O	1:A:403:THR:C	2.56	0.43
1:A:585:TRP:N	1:A:585:TRP:CE3	2.83	0.43
1:A:601:LYS:HG2	1:A:721:ALA:HB3	1.98	0.43
1:A:70:ARG:N	1:A:81:ARG:HD2	2.33	0.43
1:A:749:PHE:N	1:A:749:PHE:HD1	2.15	0.43
1:A:790:ARG:O	1:A:794:ILE:HG13	2.18	0.43
1:A:836:LEU:O	1:A:837:TYR:HD2	2.02	0.43
1:A:892:ASP:O	1:A:893:SER:OG	2.27	0.43
1:A:931:LEU:O	1:A:932:HIS:C	2.56	0.43
1:A:995:TYR:O	1:A:996:ARG:CG	2.66	0.43
2:B:117:SER:OG	2:B:259:ARG:HD3	2.18	0.43
2:B:132:HIS:CE1	2:B:182:ARG:HH22	2.36	0.43
2:B:194:VAL:HG13	2:B:195:ASN:H	1.83	0.43
2:B:246:ARG:C	2:B:248:SER:N	2.71	0.43
2:B:259:ARG:NH1	2:B:259:ARG:CB	2.81	0.43
2:B:261:GLN:O	2:B:262:TRP:C	2.55	0.43
2:C:207:GLY:CA	2:C:239:LEU:HD11	2.45	0.43
2:C:386:ASP:OD2	2:C:441:VAL:N	2.51	0.43
2:C:76:ARG:NE	2:C:434:GLU:HB3	2.32	0.43
2:E:252:LEU:HD13	2:E:252:LEU:O	2.18	0.43
1:D:113:TRP:CE3	1:D:114:GLY:N	2.58	0.43
1:D:352:SER:HB3	1:D:352:SER:C	2.36	0.43
1:D:440:ASN:CB	1:D:441:TRP:CE3	3.02	0.43
1:D:441:TRP:CE3	1:D:441:TRP:N	2.86	0.43
1:D:802:ARG:CG	1:D:803:ASN:N	2.81	0.43
1:D:874:LEU:HG	1:D:878:VAL:HG13	2.01	0.43
1:D:926:SER:O	1:D:929:THR:HG22	2.18	0.43
2:E:371:VAL:O	2:E:372:LEU:HB2	2.18	0.43
2:F:247:THR:OG1	2:F:251:TRP:HB2	2.18	0.43
2:F:396:ARG:CZ	2:F:397:GLN:HA	2.48	0.43
2:F:433:ASP:C	2:F:435:MET:H	2.21	0.43
2:F:76:ARG:HD3	2:F:434:GLU:HB2	2.00	0.43
2:F:449:GLU:O	2:F:451:GLY:N	2.46	0.43
2:F:69:LEU:HD23	2:F:351:TYR:CD1	2.53	0.43
2:F:70:LEU:O	2:F:71:GLU:C	2.57	0.43
1:A:1151:LEU:O	1:A:1152:ALA:C	2.56	0.43
1:A:175:TRP:O	1:A:176:THR:HB	2.18	0.43
1:A:282:TYR:CZ	1:A:291:PHE:CZ	3.06	0.43
1:A:346:ASP:HB2	1:A:1040:LYS:HG3	2.00	0.43
1:A:570:PRO:CG	1:A:572:TRP:CZ3	3.02	0.43
1:A:601:LYS:C	1:A:602:LEU:O	2.45	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:610:PHE:HB2	1:A:611:PRO:HD3	2.00	0.43
1:A:588:GLY:N	1:A:806:LYS:HZ1	2.15	0.43
1:A:813:VAL:HG13	1:A:815:TRP:CZ2	2.52	0.43
1:A:79:LEU:HB2	1:A:89:GLY:O	2.19	0.43
1:A:904:ASP:OD1	1:A:915:ALA:HB3	2.19	0.43
1:A:916:PHE:CE2	1:A:920:THR:O	2.72	0.43
2:B:213:VAL:O	2:B:213:VAL:HG13	2.18	0.43
2:B:78:PHE:CE1	2:B:103:GLY:CA	3.01	0.43
2:C:121:PHE:C	2:C:123:GLU:H	2.21	0.43
2:C:322:LEU:C	2:C:322:LEU:HD13	2.36	0.43
2:C:372:LEU:HD21	2:C:436:SER:CB	2.46	0.43
1:A:560:PRO:HG2	2:C:452:LEU:HD21	2.00	0.43
2:E:389:ARG:C	2:E:389:ARG:NE	2.70	0.43
2:E:312:LEU:CD1	2:E:312:LEU:H	2.32	0.43
1:A:542:ASP:O	1:A:546:ARG:HG3	2.18	0.43
2:C:280:ASP:HB3	2:C:282:GLU:O	2.18	0.43
2:F:479:ILE:HG13	2:F:479:ILE:H	1.47	0.43
2:E:476:ILE:O	2:E:480:SER:CB	2.66	0.43
1:D:1008:LEU:O	1:D:1012:VAL:CG2	2.63	0.43
1:D:1091:GLU:O	1:D:1092:PHE:CD2	2.71	0.43
1:D:887:VAL:N	1:D:1146:ARG:HH22	2.09	0.43
1:D:1232:GLU:HB2	1:D:1234:ARG:N	2.33	0.43
1:D:136:ASP:C	1:D:136:ASP:OD1	2.57	0.43
1:D:370:GLU:O	1:D:371:LYS:CB	2.60	0.43
1:D:169:TRP:CD1	1:D:390:GLN:O	2.71	0.43
1:D:545:ALA:O	1:D:548:CYS:SG	2.75	0.43
1:D:576:LEU:HB3	1:D:579:ARG:HD2	2.00	0.43
1:D:604:ALA:O	1:D:782:GLY:N	2.48	0.43
1:D:876:ALA:O	1:D:878:VAL:N	2.51	0.43
1:D:878:VAL:O	1:D:879:GLN:CB	2.61	0.43
2:E:205:PRO:CG	2:E:241:TRP:NE1	2.79	0.43
2:E:215:PHE:CD1	2:E:215:PHE:O	2.71	0.43
2:E:374:LEU:HB3	2:E:375:HIS:H	1.59	0.43
2:E:422:GLN:O	2:E:424:SER:N	2.51	0.43
2:E:379:ALA:CB	2:E:438:LEU:HD21	2.49	0.43
2:E:462:MET:HA	2:E:462:MET:HE1	2.01	0.43
2:F:262:TRP:O	2:F:266:PHE:HD2	2.02	0.43
2:F:403:PHE:HD2	2:F:404:ASN:N	2.15	0.43
2:F:427:GLN:C	2:F:430:SER:HG	2.21	0.43
1:A:1040:LYS:C	1:A:1042:GLU:H	2.20	0.43
1:A:1108:TYR:HE2	1:A:1112:MET:HE1	1.81	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1118:TRP:CZ3	1:A:1121:GLU:OE1	2.72	0.43
1:A:401:TRP:O	1:A:402:ALA:C	2.56	0.43
1:A:359:ALA:CB	1:A:407:PHE:HE2	2.22	0.43
1:A:407:PHE:N	1:A:407:PHE:HD2	2.16	0.43
1:A:459:MET:O	1:A:462:SER:OG	2.29	0.43
1:A:588:GLY:HA3	1:A:589:PRO:HD2	1.78	0.43
1:A:610:PHE:H	1:A:610:PHE:HD2	1.59	0.43
1:A:769:ASP:O	1:A:772:PRO:CD	2.66	0.43
1:A:79:LEU:HA	1:A:79:LEU:HD23	1.65	0.43
1:A:81:ARG:HA	1:A:84:HIS:CD2	2.52	0.43
1:A:83:LEU:HD22	1:A:89:GLY:CA	2.39	0.43
1:A:960:PRO:HA	1:A:963:GLU:HG3	1.99	0.43
2:B:382:LYS:N	2:B:412:SER:OG	2.50	0.43
2:B:95:CYS:O	2:B:96:HIS:C	2.56	0.43
1:A:564:GLN:N	1:A:564:GLN:NE2	2.61	0.43
1:A:531:GLY:HA2	1:A:532:PRO:HD3	1.77	0.43
2:E:339:ASN:N	2:E:339:ASN:HD22	2.17	0.43
1:D:1047:ARG:O	1:D:1048:ALA:C	2.56	0.43
1:D:1051:GLY:HA2	1:D:1054:GLU:CG	2.36	0.43
1:D:1056:GLU:HA	1:D:1059:ASN:ND2	2.22	0.43
1:D:1126:ASP:O	1:D:1142:ARG:HG3	2.19	0.43
1:D:177:ARG:HE	1:D:216:SER:N	2.13	0.43
1:D:214:ALA:CB	1:D:219:ALA:HB3	2.48	0.43
1:D:281:GLN:NE2	1:D:287:SER:OG	2.51	0.43
1:D:359:VAL:CG1	1:D:406:VAL:HG21	2.48	0.43
1:D:559:LEU:CD1	1:D:560:PRO:CD	2.96	0.43
1:D:597:ARG:O	1:D:598:VAL:C	2.56	0.43
1:D:598:VAL:HG22	1:D:601:LYS:HE2	2.00	0.43
1:D:774:MET:HB3	1:D:779:LEU:HD13	2.01	0.43
1:D:783:PRO:CG	1:D:784:GLY:N	2.81	0.43
1:D:921:LEU:H	1:D:921:LEU:HD12	1.84	0.43
1:D:954:ILE:CG2	1:D:955:TYR:N	2.82	0.43
1:D:770:PHE:HE1	1:D:968:GLN:C	2.21	0.43
2:F:203:ARG:CG	2:F:325:ARG:HD3	2.49	0.43
2:F:259:ARG:HE	2:F:339:ASN:HD21	1.66	0.43
2:F:235:THR:O	2:F:341:ASP:CG	2.57	0.43
2:F:351:TYR:O	2:F:355:GLN:CB	2.65	0.43
2:F:446:THR:CB	2:F:450:ASN:CB	2.80	0.43
1:A:1218:LEU:O	1:A:1222:GLN:CD	2.57	0.43
1:A:172:ALA:O	1:A:173:GLU:CB	2.67	0.43
1:A:249:VAL:HB	1:A:250:PRO:HD3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:350:ILE:HA	1:A:350:ILE:HD12	1.67	0.43
1:A:371:LYS:HG2	1:A:372:GLU:HG2	1.99	0.43
1:A:415:LEU:C	1:A:417:ARG:H	2.21	0.43
1:A:420:HIS:ND1	1:A:422:VAL:HG23	2.33	0.43
1:A:498:LYS:HG2	1:A:518:ALA:N	2.33	0.43
1:A:761:ASN:O	1:A:762:VAL:C	2.55	0.43
1:A:77:GLN:HB2	1:A:1174:PRO:CG	2.48	0.43
1:A:811:GLN:HG3	1:A:840:ILE:HD13	2.00	0.43
1:A:878:VAL:C	1:A:879:GLN:HE21	2.21	0.43
1:A:876:ALA:O	1:A:878:VAL:N	2.52	0.43
1:A:905:ALA:HA	1:A:910:MET:CB	2.48	0.43
1:A:968:GLN:O	1:A:970:ASN:O	2.37	0.43
2:B:101:PRO:HB2	2:B:102:LEU:HD12	2.01	0.43
2:B:206:TYR:CD1	2:B:207:GLY:O	2.70	0.43
2:B:343:ASP:OD2	2:B:344:ARG:NH2	2.33	0.43
2:B:433:ASP:O	2:B:436:SER:N	2.51	0.43
2:B:87:SER:O	2:B:91:LEU:HG	2.18	0.43
2:C:204:LEU:HG	2:C:206:TYR:CD2	2.54	0.43
2:C:241:TRP:HZ3	2:C:255:TRP:CZ2	2.35	0.43
2:C:338:VAL:C	2:C:339:ASN:O	2.52	0.43
2:C:341:ASP:O	2:C:342:LEU:CB	2.66	0.43
2:C:344:ARG:O	2:C:347:LEU:N	2.52	0.43
2:C:456:ARG:NE	2:C:456:ARG:N	2.64	0.43
1:A:307:PHE:CD1	1:A:307:PHE:N	2.81	0.43
2:B:133:HIS:CG	2:C:231:ILE:HG21	2.52	0.43
1:A:539:PHE:C	1:A:539:PHE:CD2	2.92	0.43
1:A:543:VAL:HG13	1:A:544:MET:H	1.82	0.43
2:F:278:CYS:SG	2:F:279:GLN:N	2.92	0.43
2:E:401:GLY:O	2:E:405:GLU:HB2	2.18	0.43
2:C:136:GLY:HA2	2:C:137:PRO:HD3	1.74	0.43
1:D:907:PHE:CE1	1:D:1002:GLU:HB3	2.52	0.43
1:D:1024:ASP:C	1:D:1026:ARG:N	2.71	0.43
1:D:897:TRP:CZ2	1:D:1177:VAL:HG11	2.50	0.43
1:D:177:ARG:NH2	1:D:216:SER:HA	2.34	0.43
1:D:256:PRO:HB3	1:D:281:GLN:HA	2.00	0.43
1:D:405:GLU:OE2	1:D:405:GLU:CA	2.66	0.43
1:D:458:GLU:N	1:D:460:LYS:CE	2.82	0.43
1:D:629:ASP:OD2	1:D:744:ILE:O	2.37	0.43
1:D:916:PHE:CD1	1:D:916:PHE:O	2.70	0.43
1:D:989:THR:C	1:D:998:SER:CB	2.87	0.43
2:E:214:CYS:SG	2:E:236:GLU:OE1	2.77	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:326:ASP:N	2:E:331:VAL:HG12	2.29	0.43
2:E:79:LEU:C	2:E:100:GLY:CA	2.86	0.43
2:F:78:PHE:HD1	2:F:102:LEU:CD2	2.30	0.43
2:F:193:TYR:CB	2:F:242:PHE:CE2	2.97	0.43
1:D:786:ALA:CB	2:F:358:GLU:OE2	2.67	0.43
2:F:454:HIS:HD2	2:F:468:ILE:CG1	2.31	0.43
1:A:1123:PHE:N	1:A:1123:PHE:CD1	2.86	0.43
1:A:1131:ILE:HD13	1:A:1137:VAL:C	2.38	0.43
1:A:1161:ARG:O	1:A:1162:CYS:C	2.55	0.43
1:A:1170:LEU:N	1:A:1170:LEU:CD1	2.82	0.43
1:A:1231:LEU:CD1	1:A:1235:SER:CB	2.97	0.43
1:A:165:LYS:HG3	1:A:178:TYR:HB2	2.00	0.43
1:A:167:PRO:CB	1:A:177:ARG:NE	2.81	0.43
1:A:168:ALA:C	1:A:169:TRP:CG	2.91	0.43
1:A:448:ALA:HA	1:A:1202:ASN:HD21	1.84	0.43
1:A:583:PRO:C	1:A:585:TRP:N	2.71	0.43
1:A:845:VAL:HG11	1:A:848:GLY:CA	2.48	0.43
1:A:847:ALA:C	1:A:848:GLY:O	2.55	0.43
1:A:891:VAL:HG22	1:A:894:GLN:CB	2.48	0.43
1:A:893:SER:OG	1:A:893:SER:O	2.36	0.43
1:A:897:TRP:HE1	1:A:901:VAL:HG11	1.81	0.43
1:A:904:ASP:HB2	1:A:910:MET:C	2.37	0.43
1:A:76:ILE:HD13	1:A:911:HIS:CD2	2.54	0.43
2:B:419:GLU:CD	2:C:201:ASN:ND2	2.71	0.43
2:C:102:LEU:HG	2:C:380:PRO:HG3	1.99	0.43
2:B:108:LYS:HE3	2:C:115:TRP:CZ2	2.54	0.43
2:C:256:LEU:O	2:C:257:ARG:C	2.57	0.43
2:C:364:LYS:N	2:C:364:LYS:HD3	2.18	0.43
2:C:475:LEU:H	2:C:475:LEU:HG	1.50	0.43
2:E:260:LEU:O	2:E:264:ARG:HG3	2.19	0.43
1:D:231:GLU:HA	1:D:231:GLU:OE1	2.19	0.43
1:D:1046:GLU:HG2	1:D:1049:TRP:NE1	2.34	0.43
1:D:285:GLN:HE21	1:D:1128:ARG:HG2	1.74	0.43
1:D:1212:ILE:HA	1:D:1212:ILE:HD12	1.80	0.43
1:D:420:HIS:O	1:D:424:LEU:CD1	2.66	0.43
1:D:421:PRO:HG2	1:D:422:VAL:HG23	2.00	0.43
1:D:486:TRP:CZ2	1:D:722:ARG:CA	2.81	0.43
1:D:740:ASN:HB2	1:D:741:ASP:H	1.56	0.43
1:D:788:GLY:O	1:D:791:ALA:CA	2.67	0.43
1:D:922:GLN:HB2	1:D:924:ARG:HD3	2.01	0.43
1:D:955:TYR:O	1:D:956:GLY:O	2.37	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:993:ARG:NH2	1:D:994:TRP:CD1	2.87	0.43
1:D:994:TRP:O	1:D:994:TRP:CE3	2.71	0.43
2:E:109:ASN:CG	2:E:378:LEU:HB3	2.39	0.43
2:E:205:PRO:HG2	2:E:241:TRP:NE1	2.33	0.43
2:E:437:ILE:CG2	2:E:439:PHE:C	2.86	0.43
2:E:447:THR:HG22	2:E:451:GLY:HA2	2.00	0.43
2:E:456:ARG:CZ	2:E:463:LYS:CE	2.83	0.43
2:E:475:LEU:O	2:E:479:ILE:HG13	2.19	0.43
2:F:374:LEU:O	2:F:375:HIS:C	2.57	0.43
2:F:454:HIS:CD2	2:F:472:LYS:HE3	2.54	0.43
2:F:474:PHE:HA	2:F:477:LYS:CG	2.49	0.43
1:A:1030:ARG:NE	1:A:1040:LYS:NZ	2.66	0.43
1:A:139:PHE:CE2	1:A:1111:LEU:HG	2.54	0.43
1:A:1190:ARG:HD2	1:A:1195:MET:HG2	2.00	0.43
1:A:166:PRO:CG	1:A:398:GLN:NE2	2.48	0.43
1:A:197:PHE:CZ	1:A:214:ALA:HB2	2.54	0.43
1:A:461:LYS:HE2	1:A:461:LYS:HB3	1.80	0.43
1:A:513:LEU:HB2	1:A:568:GLY:HA3	1.84	0.43
1:A:606:THR:CG2	1:A:620:TRP:CE3	2.78	0.43
1:A:784:GLY:HA3	2:C:363:ARG:CG	2.46	0.43
1:A:964:ARG:O	1:A:968:GLN:HG3	2.18	0.43
1:A:914:THR:CB	1:A:999:ASP:HB3	2.48	0.43
2:B:257:ARG:HG2	2:B:258:HIS:N	2.34	0.43
2:B:269:SER:O	2:B:272:ASN:CG	2.57	0.43
2:C:78:PHE:HD1	2:C:102:LEU:HD22	1.79	0.43
2:C:374:LEU:HB3	2:C:378:LEU:CB	2.36	0.43
1:D:509:THR:O	1:D:510:ALA:O	2.36	0.43
1:A:739:TYR:HB3	1:A:740:ASN:H	1.51	0.43
1:D:1007:GLU:C	1:D:1011:PRO:HD3	2.34	0.43
1:D:1073:PRO:O	1:D:1074:VAL:C	2.57	0.43
1:D:1131:ILE:CG1	1:D:1138:ARG:HB2	2.48	0.43
1:D:1136:GLU:CG	1:D:1138:ARG:HH12	2.32	0.43
1:D:220:TRP:HD1	1:D:221:TYR:HD2	1.62	0.43
1:D:253:ALA:CA	1:D:280:GLU:HB2	2.49	0.43
1:D:275:ARG:O	1:D:279:ARG:HG2	2.18	0.43
1:D:401:TRP:CD1	1:D:404:HIS:CD2	2.96	0.43
1:D:477:GLU:HG2	1:D:478:ARG:N	2.34	0.43
1:D:482:ASP:OD1	1:D:484:TRP:HB3	2.19	0.43
1:D:549:LEU:O	1:D:552:LEU:N	2.51	0.43
1:D:559:LEU:CB	1:D:560:PRO:CD	2.97	0.43
1:D:715:GLN:O	1:D:716:PRO:C	2.57	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:788:GLY:O	1:D:791:ALA:CB	2.64	0.43
1:D:427:MET:C	1:D:846:THR:HG21	2.38	0.43
1:D:851:THR:O	1:D:852:ARG:CB	2.66	0.43
1:D:932:HIS:HE1	1:D:943:ARG:HB3	1.83	0.43
2:E:471:LEU:O	2:E:474:PHE:HB3	2.18	0.43
2:F:248:SER:OG	2:F:249:ASN:N	2.47	0.43
2:F:302:THR:H	2:F:339:ASN:HA	1.82	0.43
2:F:338:VAL:HG12	2:F:339:ASN:N	2.34	0.43
2:F:472:LYS:O	2:F:475:LEU:CG	2.67	0.43
1:A:1060:LYS:HE2	1:A:1065:ALA:H	1.76	0.43
1:A:177:ARG:O	1:A:178:TYR:C	2.56	0.43
1:A:276:ALA:O	1:A:279:ARG:N	2.44	0.43
1:A:275:ARG:C	1:A:278:ILE:HB	2.39	0.43
1:A:313:ILE:HA	1:A:316:LYS:CE	2.45	0.43
1:A:618:HIS:NE2	1:A:726:LYS:O	2.52	0.43
1:A:653:GLU:OE2	1:A:657:ARG:CZ	2.67	0.43
1:A:608:ASP:C	1:A:779:LEU:HD22	2.39	0.43
2:C:115:TRP:CH2	2:C:210:GLN:HA	2.54	0.43
2:C:293:PHE:HB3	2:C:294:PRO:HD2	2.01	0.43
2:C:432:TYR:C	2:C:435:MET:H	2.21	0.43
2:C:464:GLU:OE2	2:C:466:MET:CG	2.64	0.43
2:C:468:ILE:C	2:C:471:LEU:HD13	2.39	0.43
2:C:76:ARG:NH1	2:C:76:ARG:CG	2.81	0.43
1:D:525:MET:C	1:D:526:ASP:CG	2.77	0.43
2:B:280:ASP:HB3	2:B:281:GLU:CB	2.49	0.43
2:F:438:LEU:CD2	2:F:438:LEU:N	2.81	0.43
1:D:501:LYS:HG2	1:D:502:LYS:N	2.33	0.43
2:E:271:SER:C	2:E:273:PHE:N	2.71	0.43
1:A:737:GLY:N	1:A:750:PHE:CE1	2.86	0.43
1:D:248:GLU:OE1	1:D:250:PRO:CD	2.67	0.43
1:D:368:PRO:HB3	1:D:369:LEU:HD23	2.00	0.43
1:D:376:LEU:C	1:D:378:VAL:H	2.22	0.43
1:D:440:ASN:HB2	1:D:441:TRP:CE3	2.53	0.43
1:D:469:ASP:O	1:D:473:LEU:HG	2.18	0.43
1:D:771:LEU:CD1	1:D:771:LEU:H	2.32	0.43
1:D:433:SER:HB3	1:D:844:VAL:HG21	2.00	0.43
1:D:844:VAL:HG13	1:D:856:GLU:HG3	2.00	0.43
1:D:909:GLY:O	1:D:912:GLY:CA	2.66	0.43
1:D:961:PHE:CD1	1:D:961:PHE:N	2.83	0.43
2:E:183:GLU:O	2:E:214:CYS:SG	2.70	0.43
1:A:1133:ILE:HD13	1:A:1134:HIS:N	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:225:SER:HA	1:A:228:LEU:CD1	2.31	0.43
1:A:255:SER:N	1:A:256:PRO:CA	2.81	0.43
1:A:262:GLN:HB3	1:A:263:GLU:H	1.54	0.43
1:A:386:ARG:CA	1:A:389:PHE:HB3	2.46	0.43
1:A:444:TYR:CD2	1:A:874:LEU:CD2	2.85	0.43
1:A:616:GLU:OE2	1:A:724:GLY:N	2.48	0.43
1:A:618:HIS:N	1:A:725:PRO:CG	2.57	0.43
1:A:600:PRO:CG	1:A:765:PRO:HG3	2.49	0.43
1:A:754:HIS:HA	1:A:770:PHE:HE2	1.83	0.43
1:A:798:ILE:O	1:A:799:SER:C	2.57	0.43
1:A:824:ALA:C	1:A:827:ARG:H	2.22	0.43
1:A:944:GLU:HG2	1:A:947:LYS:HZ2	1.83	0.43
2:B:204:LEU:HG	2:B:325:ARG:CZ	2.48	0.43
2:B:234:LYS:O	2:B:236:GLU:CG	2.63	0.43
2:B:285:LYS:CG	2:B:286:GLY:N	2.37	0.43
2:B:311:LEU:HD13	2:B:322:LEU:HD13	2.00	0.43
2:B:375:HIS:HA	2:B:376:PRO:HD2	1.76	0.43
2:B:76:ARG:HH21	2:B:431:LYS:HB2	1.84	0.43
2:B:72:ILE:HA	2:B:75:ARG:CD	2.42	0.43
2:C:108:LYS:O	2:C:109:ASN:C	2.56	0.43
2:C:341:ASP:C	2:C:342:LEU:HD22	2.39	0.43
2:C:234:LYS:HZ1	2:C:344:ARG:CD	2.28	0.43
2:C:375:HIS:O	2:C:376:PRO:C	2.55	0.43
2:C:406:LEU:N	2:C:406:LEU:HD13	2.34	0.43
2:C:466:MET:CE	2:C:471:LEU:HD23	2.49	0.43
1:D:162:LEU:CG	1:D:162:LEU:O	2.67	0.43
2:F:108:LYS:O	2:F:109:ASN:C	2.55	0.43
1:D:1119:LEU:N	1:D:1119:LEU:HD23	2.34	0.43
1:D:1157:ASN:HB3	1:D:1161:ARG:CZ	2.48	0.43
1:D:444:TYR:HH	1:D:1206:MET:CG	2.31	0.43
1:D:441:TRP:HZ3	1:D:1210:TYR:CE1	2.36	0.43
1:D:80:SER:CB	1:D:125:LEU:HG	2.43	0.43
1:D:919:MET:O	1:D:920:THR:O	2.36	0.43
1:D:955:TYR:HB2	1:D:956:GLY:H	1.43	0.43
2:E:78:PHE:HD2	2:E:102:LEU:CD1	2.29	0.43
2:E:113:GLU:HG3	2:E:377:CYS:SG	2.59	0.43
2:E:212:GLY:O	2:E:236:GLU:N	2.52	0.43
2:E:398:VAL:O	2:E:402:LEU:N	2.49	0.43
2:F:300:ILE:CG2	2:F:301:GLU:N	2.63	0.43
2:F:342:LEU:HD21	2:F:345:GLY:H	1.70	0.43
2:F:372:LEU:CD2	2:F:436:SER:HB2	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:446:THR:O	2:F:449:GLU:N	2.47	0.43
1:A:1208:ARG:CZ	2:C:277:ASP:CG	2.87	0.43
1:A:293:ASP:CB	1:A:296:SER:CB	2.97	0.43
1:A:348:LEU:CD1	1:A:350:ILE:HD13	2.45	0.43
1:A:599:THR:O	1:A:603:MET:HB2	2.19	0.43
1:A:624:VAL:CG2	1:A:747:CYS:SG	3.06	0.43
1:A:716:PRO:O	1:A:717:LEU:C	2.56	0.43
1:A:601:LYS:CA	1:A:717:LEU:HB3	2.41	0.43
1:A:817:PRO:CG	1:A:818:ARG:H	2.32	0.43
1:A:444:TYR:CB	1:A:877:MET:HG3	2.49	0.43
1:A:986:TYR:O	1:A:990:LYS:HG3	2.18	0.43
2:B:235:THR:OG1	2:B:343:ASP:CA	2.66	0.43
2:B:373:LYS:HE2	2:B:459:ASP:O	2.18	0.43
2:B:81:GLY:C	2:B:86:LEU:HD11	2.38	0.43
2:C:123:GLU:O	2:C:124:GLN:HB3	2.19	0.43
2:C:128:VAL:HG12	2:C:192:HIS:NE2	2.34	0.43
1:D:525:MET:HE2	1:D:525:MET:HB3	1.90	0.43
1:D:666:GLN:HA	1:D:669:MET:CG	2.48	0.43
1:A:1036:SER:O	1:A:1039:LYS:HA	2.19	0.43
2:B:180:LYS:HG3	2:B:181:LEU:N	2.34	0.43
1:D:1025:LEU:C	1:D:1026:ARG:HD3	2.37	0.42
1:D:1027:LYS:CD	1:D:1027:LYS:N	2.78	0.42
1:D:1065:ALA:HA	1:D:1073:PRO:HG3	2.00	0.42
1:D:1193:VAL:C	1:D:1195:MET:HE2	2.40	0.42
1:D:204:ALA:O	1:D:205:GLU:CB	2.66	0.42
1:D:457:ARG:N	1:D:460:LYS:NZ	2.46	0.42
1:D:576:LEU:HG	1:D:579:ARG:NE	2.32	0.42
1:D:598:VAL:C	1:D:600:PRO:HD2	2.39	0.42
1:D:613:HIS:CE1	1:D:717:LEU:HD22	2.53	0.42
1:D:648:PRO:HG2	1:D:649:TYR:N	2.34	0.42
1:D:851:THR:C	1:D:852:ARG:HG3	2.39	0.42
1:D:85:GLU:OE1	1:D:130:LEU:CD2	2.63	0.42
1:D:959:GLN:O	1:D:962:ALA:HB3	2.19	0.42
2:E:128:VAL:CG1	2:E:208:LEU:O	2.66	0.42
2:E:372:LEU:O	2:E:458:ARG:CG	2.67	0.42
2:E:385:LEU:O	2:E:386:ASP:HB3	2.17	0.42
2:E:456:ARG:HG3	2:E:462:MET:O	2.18	0.42
2:E:86:LEU:HB3	2:E:97:PRO:HD2	2.01	0.42
2:F:389:ARG:CZ	2:F:390:GLY:N	2.82	0.42
1:A:285:GLN:C	1:A:1128:ARG:HH22	2.12	0.42
1:A:382:MET:HE3	1:A:383:LYS:HA	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:468:ASN:HB3	2:C:460:THR:HG22	2.01	0.42
1:A:497:GLN:OE1	1:A:516:GLU:HB3	2.19	0.42
1:A:659:HIS:O	1:A:662:GLU:HG2	2.18	0.42
1:A:656:TYR:HD2	1:A:720:THR:HG1	1.55	0.42
1:A:728:THR:C	1:A:729:GLN:NE2	2.72	0.42
1:A:772:PRO:HG2	1:A:773:LYS:N	2.33	0.42
1:A:788:GLY:N	1:A:789:PRO:CD	2.82	0.42
1:A:824:ALA:O	1:A:827:ARG:N	2.51	0.42
1:A:856:GLU:O	1:A:857:PRO:C	2.56	0.42
1:A:448:ALA:C	1:A:870:VAL:CG1	2.86	0.42
1:A:996:ARG:NH2	1:A:1000:GLU:OE2	2.52	0.42
2:B:116:THR:HG23	2:B:120:VAL:HB	2.00	0.42
2:B:386:ASP:OD1	2:B:387:VAL:N	2.42	0.42
2:C:100:GLY:O	2:C:103:GLY:N	2.52	0.42
2:C:267:ALA:CB	2:C:270:PRO:HB3	2.47	0.42
2:E:315:TYR:CD1	2:E:316:PRO:HD2	2.51	0.42
2:F:136:GLY:HA2	2:F:137:PRO:HD3	1.71	0.42
1:D:1108:TYR:CE1	1:D:1160:THR:CG2	3.02	0.42
1:D:1148:ARG:CZ	1:D:1231:LEU:CD2	2.97	0.42
1:D:916:PHE:CD2	1:D:1174:PRO:HG2	2.53	0.42
1:D:1208:ARG:NH1	2:F:285:LYS:CE	2.81	0.42
1:D:249:VAL:HG22	1:D:250:PRO:HD2	1.78	0.42
1:D:267:VAL:CB	1:D:269:HIS:NE2	2.81	0.42
1:D:304:LEU:H	1:D:309:ARG:HH21	1.59	0.42
1:D:458:GLU:OE2	2:F:270:PRO:HG3	2.18	0.42
1:D:479:TYR:C	1:D:481:GLU:H	2.22	0.42
1:D:547:ALA:HA	1:D:551:LYS:CE	2.39	0.42
1:D:577:CYS:HB3	1:D:578:PRO:CD	2.46	0.42
1:D:614:TYR:HE2	1:D:620:TRP:CD2	2.35	0.42
1:D:718:ALA:O	1:D:722:ARG:NE	2.52	0.42
1:D:618:HIS:CA	1:D:762:VAL:HG21	2.46	0.42
1:D:776:ASP:N	1:D:796:LYS:HZ3	2.16	0.42
1:D:83:LEU:HB3	1:D:88:PHE:C	2.39	0.42
1:D:967:MET:SD	1:D:974:THR:C	2.97	0.42
1:D:996:ARG:CA	1:D:996:ARG:NE	2.82	0.42
2:E:120:VAL:HG12	2:E:121:PHE:CD1	2.53	0.42
2:E:130:ALA:CB	2:E:210:GLN:NE2	2.82	0.42
2:F:262:TRP:O	2:F:265:LYS:HB3	2.19	0.42
2:F:197:LEU:CD1	2:F:322:LEU:HD11	2.50	0.42
2:F:435:MET:CE	2:F:435:MET:CA	2.90	0.42
1:A:1049:TRP:CG	1:A:1050:LYS:N	2.87	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:756:ASP:CG	1:A:1057:MET:HG2	2.38	0.42
1:A:1007:GLU:CD	1:A:1093:MET:HE2	2.39	0.42
1:A:1157:ASN:HD22	1:A:1157:ASN:N	2.17	0.42
1:A:304:LEU:O	1:A:850:ILE:CG2	2.67	0.42
1:A:369:PRO:O	1:A:369:PRO:HG2	2.20	0.42
1:A:477:GLU:CD	1:A:478:ARG:HG2	2.39	0.42
1:A:499:LYS:HB2	1:A:499:LYS:HE3	1.85	0.42
1:A:612:LEU:HB2	1:A:621:GLY:N	2.34	0.42
1:A:656:TYR:CA	1:A:720:THR:HB	2.49	0.42
2:B:241:TRP:CD2	2:B:255:TRP:HH2	2.36	0.42
2:B:301:GLU:HG2	2:B:303:LEU:HD11	2.01	0.42
2:B:303:LEU:CD1	2:B:303:LEU:N	2.79	0.42
2:B:306:LEU:O	2:B:308:ASP:O	2.37	0.42
2:C:443:VAL:HG13	2:C:447:THR:HG21	2.00	0.42
2:C:451:GLY:O	2:C:452:LEU:CG	2.67	0.42
2:C:87:SER:O	2:C:88:ARG:C	2.58	0.42
2:E:264:ARG:NH2	2:E:273:PHE:O	2.52	0.42
2:F:221:THR:CG2	2:F:222:LYS:H	2.32	0.42
2:F:145:PHE:CD1	2:F:145:PHE:N	2.87	0.42
1:A:120:LEU:HD13	1:A:120:LEU:N	2.34	0.42
1:D:1020:ILE:O	1:D:1021:SER:HB2	2.20	0.42
1:D:1150:ALA:O	1:D:1153:LEU:HB2	2.19	0.42
1:D:274:ASP:HA	1:D:277:HIS:CD2	2.54	0.42
1:D:292:LEU:HD23	1:D:297:MET:HE3	1.98	0.42
1:D:395:TYR:N	1:D:395:TYR:CD1	2.87	0.42
1:D:421:PRO:CG	1:D:422:VAL:N	2.81	0.42
1:D:439:GLN:O	1:D:440:ASN:C	2.57	0.42
1:D:549:LEU:C	1:D:551:LYS:HZ2	2.22	0.42
1:D:581:ASP:OD2	1:D:593:SER:HB2	2.18	0.42
1:D:475:SER:C	1:D:712:VAL:HG23	2.40	0.42
2:E:254:PHE:HA	2:E:257:ARG:CB	2.27	0.42
2:E:350:LEU:C	2:E:350:LEU:HD23	2.40	0.42
2:E:382:LYS:O	2:E:383:VAL:CG2	2.67	0.42
2:E:404:ASN:O	2:E:407:LEU:N	2.53	0.42
2:E:419:GLU:HA	2:F:203:ARG:HH21	1.75	0.42
2:E:474:PHE:CG	2:E:475:LEU:N	2.88	0.42
2:F:118:VAL:HG12	2:F:119:VAL:HG23	2.01	0.42
2:F:251:TRP:HE3	2:F:254:PHE:HB3	1.85	0.42
2:F:357:THR:C	2:F:359:ASN:N	2.72	0.42
1:A:1129:PHE:CG	1:A:1130:CYS:N	2.87	0.42
1:A:202:CYS:C	1:A:204:ALA:N	2.71	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:209:PRO:C	1:A:210:THR:HG23	2.39	0.42
1:A:304:LEU:CB	1:A:309:ARG:HB3	2.50	0.42
1:A:513:LEU:O	1:A:568:GLY:C	2.58	0.42
1:A:627:ARG:O	1:A:627:ARG:CG	2.67	0.42
1:A:824:ALA:O	1:A:825:VAL:C	2.58	0.42
1:A:829:PRO:HG3	1:A:883:GLY:CA	2.49	0.42
1:A:905:ALA:CA	1:A:910:MET:HA	2.49	0.42
2:B:262:TRP:O	2:B:265:LYS:CA	2.66	0.42
2:C:124:GLN:C	2:C:125:VAL:CG1	2.88	0.42
2:C:355:GLN:OE1	2:C:356:LEU:HD23	2.19	0.42
2:C:358:GLU:CA	2:C:361:PHE:HB2	2.47	0.42
2:C:372:LEU:HD21	2:C:436:SER:N	2.32	0.42
2:C:419:GLU:OE2	2:C:428:LEU:CD1	2.50	0.42
2:C:80:SER:H	2:C:99:PHE:HA	1.82	0.42
1:D:586:THR:N	1:D:587:PRO:HD2	2.29	0.42
1:D:1090:GLU:HB3	1:D:1092:PHE:CE1	2.54	0.42
1:D:1173:LEU:HA	1:D:1174:PRO:HD3	1.75	0.42
1:D:375:GLU:O	1:D:378:VAL:C	2.54	0.42
1:D:658:LYS:CE	1:D:713:PRO:CB	2.85	0.42
1:D:860:LEU:HA	1:D:860:LEU:HD23	1.74	0.42
1:D:954:ILE:O	1:D:955:TYR:CD1	2.73	0.42
2:E:379:ALA:N	2:E:380:PRO:HD3	2.35	0.42
2:F:113:GLU:HG2	2:F:266:PHE:CE1	2.54	0.42
2:F:293:PHE:CE2	2:F:298:GLU:HB2	2.55	0.42
2:F:305:ASN:HA	2:F:335:VAL:HG13	2.00	0.42
2:F:389:ARG:HD2	2:F:389:ARG:HA	1.74	0.42
2:F:414:TRP:O	2:F:415:PRO:C	2.57	0.42
1:A:1026:ARG:CG	1:A:1029:GLN:NE2	2.82	0.42
1:A:1065:ALA:C	1:A:1066:THR:HG1	2.18	0.42
1:A:1134:HIS:HB2	1:A:1136:GLU:OE2	2.18	0.42
1:A:1180:PHE:HB3	1:A:1181:SER:H	1.26	0.42
1:A:377:PHE:HD2	1:A:377:PHE:HA	1.41	0.42
1:A:512:LYS:C	1:A:513:LEU:HD23	2.38	0.42
1:A:582:ASP:CG	1:A:592:LEU:HD22	2.40	0.42
1:A:613:HIS:O	1:A:614:TYR:CG	2.72	0.42
1:A:614:TYR:HD2	1:A:620:TRP:HB3	1.85	0.42
1:A:607:TRP:H	1:A:620:TRP:HZ3	1.67	0.42
1:A:601:LYS:CE	1:A:721:ALA:HB1	2.50	0.42
1:A:765:PRO:HG2	1:A:766:PHE:H	1.84	0.42
1:A:275:ARG:NE	1:A:843:GLN:HB2	2.34	0.42
1:A:842:PRO:CB	1:A:856:GLU:OE1	2.57	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:868:ASP:HB3	1:A:1199:THR:CB	2.45	0.42
1:A:914:THR:OG1	1:A:999:ASP:CG	2.58	0.42
2:B:189:ALA:O	2:B:190:LEU:C	2.58	0.42
2:B:187:HIS:HA	2:B:190:LEU:HD23	2.02	0.42
2:B:267:ALA:HB1	2:B:270:PRO:CD	2.49	0.42
2:B:344:ARG:N	2:B:344:ARG:NE	2.67	0.42
2:B:69:LEU:C	2:B:73:CYS:HG	2.22	0.42
2:B:90:SER:C	2:B:93:SER:HG	2.21	0.42
2:C:197:LEU:CD1	2:C:322:LEU:HD11	2.49	0.42
2:C:213:VAL:HG23	2:C:235:THR:OG1	2.19	0.42
1:A:783:PRO:HG2	2:C:363:ARG:CD	2.49	0.42
2:C:385:LEU:HA	2:C:385:LEU:HD23	1.74	0.42
2:C:395:LEU:C	2:C:397:GLN:N	2.72	0.42
1:D:121:PRO:HB3	1:D:1175:GLN:NE2	2.35	0.42
1:D:1212:ILE:HA	1:D:1213:PRO:HD3	1.29	0.42
1:D:258:GLN:O	1:D:259:ARG:CB	2.67	0.42
1:D:278:ILE:N	1:D:291:PHE:CD1	2.88	0.42
1:D:349:ASP:O	1:D:361:ARG:CD	2.56	0.42
1:D:434:TYR:OH	1:D:436:PRO:HB3	2.20	0.42
1:D:481:GLU:O	1:D:482:ASP:CB	2.68	0.42
1:D:487:ASP:O	1:D:491:ASP:OD1	2.36	0.42
1:D:612:LEU:HD11	1:D:622:TYR:HA	2.00	0.42
1:D:620:TRP:H	1:D:620:TRP:HD1	1.64	0.42
1:D:649:TYR:CD2	1:D:652:ILE:HD13	2.54	0.42
1:D:792:LEU:O	1:D:793:GLU:C	2.57	0.42
1:D:873:GLU:HB3	1:D:1200:PRO:CG	2.47	0.42
1:D:967:MET:HE1	1:D:974:THR:C	2.40	0.42
2:E:461:THR:HG23	2:E:461:THR:O	2.19	0.42
2:F:245:PRO:CD	2:F:246:ARG:H	2.32	0.42
2:F:102:LEU:HG	2:F:380:PRO:HG3	2.02	0.42
2:F:389:ARG:CZ	2:F:390:GLY:CA	2.96	0.42
1:A:1030:ARG:HD2	1:A:1040:LYS:NZ	2.31	0.42
1:A:1108:TYR:HE2	1:A:1112:MET:HE3	1.84	0.42
1:A:1157:ASN:C	1:A:1161:ARG:HD2	2.40	0.42
1:A:1205:GLY:O	1:A:1208:ARG:CB	2.62	0.42
1:A:1232:GLU:CD	1:A:1234:ARG:HH21	2.22	0.42
1:A:849:THR:HB	1:A:852:ARG:N	2.34	0.42
1:A:895:GLU:O	1:A:898:ILE:HG12	2.20	0.42
1:A:900:ALA:CB	1:A:915:ALA:HB1	2.49	0.42
1:A:967:MET:CG	1:A:978:ALA:CB	2.98	0.42
2:B:241:TRP:CE3	2:B:255:TRP:HH2	2.38	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:190:LEU:O	2:C:193:TYR:N	2.47	0.42
2:C:243:THR:O	2:C:334:CYS:HB2	2.20	0.42
2:C:306:LEU:H	2:C:335:VAL:CB	2.33	0.42
2:C:328:ARG:C	2:C:329:LYS:HD2	2.40	0.42
2:C:344:ARG:O	2:C:348:ALA:N	2.53	0.42
2:C:69:LEU:HD13	2:C:354:PHE:CD2	2.53	0.42
2:C:389:ARG:HA	2:C:389:ARG:HD2	1.60	0.42
2:C:78:PHE:CD1	2:C:102:LEU:CD2	2.99	0.42
2:B:466:MET:HE3	2:B:471:LEU:HD13	2.02	0.42
2:F:145:PHE:HD1	2:F:145:PHE:N	2.17	0.42
2:F:216:HIS:O	2:F:232:GLY:N	2.31	0.42
1:D:1137:VAL:CG1	1:D:1139:TYR:CE1	2.96	0.42
1:D:134:ASN:O	1:D:135:LEU:C	2.57	0.42
1:D:278:ILE:HG12	1:D:291:PHE:CD1	2.54	0.42
1:D:716:PRO:HB2	1:D:717:LEU:HD23	2.02	0.42
1:D:762:VAL:HG23	1:D:763:GLY:N	2.33	0.42
1:D:275:ARG:NH2	1:D:842:PRO:CA	2.79	0.42
1:D:879:GLN:O	1:D:880:ALA:HB2	2.20	0.42
1:D:898:ILE:HG12	1:D:899:ALA:H	1.85	0.42
1:D:943:ARG:O	1:D:946:ALA:CB	2.64	0.42
2:E:293:PHE:CE1	2:E:351:TYR:CE2	3.08	0.42
2:E:78:PHE:CE1	2:E:346:MET:CE	3.02	0.42
2:F:190:LEU:O	2:F:193:TYR:N	2.51	0.42
2:F:88:ARG:HD3	2:F:295:TRP:CE2	2.54	0.42
2:F:236:GLU:CA	2:F:341:ASP:CB	2.73	0.42
2:F:469:SER:O	2:F:472:LYS:N	2.52	0.42
1:A:1028:VAL:O	1:A:1031:GLU:C	2.58	0.42
1:A:955:TYR:HE2	1:A:1102:GLN:NE2	2.13	0.42
1:A:1102:GLN:O	1:A:1106:VAL:HG23	2.18	0.42
1:A:398:GLN:O	1:A:399:ASP:C	2.56	0.42
1:A:452:TYR:CD2	1:A:453:GLU:N	2.87	0.42
1:A:782:GLY:HA3	1:A:789:PRO:HD3	2.02	0.42
1:A:827:ARG:O	1:A:831:TYR:HE1	2.02	0.42
1:A:904:ASP:HB3	1:A:912:GLY:CA	2.49	0.42
2:B:197:LEU:HD11	2:B:325:ARG:HH12	1.84	0.42
2:B:185:LEU:HD11	2:B:238:SER:HB2	2.02	0.42
2:B:204:LEU:HG	2:B:325:ARG:HE	1.82	0.42
2:C:110:LEU:O	2:C:113:GLU:N	2.53	0.42
2:C:247:THR:OG1	2:C:251:TRP:HB2	2.20	0.42
2:C:308:ASP:O	2:C:312:LEU:HD11	2.19	0.42
2:C:424:SER:O	2:C:425:LEU:C	2.58	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:80:SER:HG	2:C:99:PHE:C	2.22	0.42
1:D:623:LEU:CG	1:D:623:LEU:O	2.55	0.42
2:E:389:ARG:CZ	2:E:390:GLY:O	2.68	0.42
2:B:122:ARG:CD	2:B:124:GLN:NE2	2.81	0.42
2:C:225:ARG:HB3	2:C:226:ASN:H	1.57	0.42
1:D:1197:CYS:C	1:D:1204:THR:CG2	2.88	0.42
1:D:1220:ILE:O	1:D:1223:ILE:N	2.52	0.42
1:D:242:ALA:CA	1:D:245:ILE:HG23	2.42	0.42
1:D:309:ARG:CA	1:D:312:TRP:HB2	2.39	0.42
1:D:348:LEU:CD1	1:D:350:ILE:CD1	2.94	0.42
1:D:369:LEU:O	1:D:370:GLU:C	2.52	0.42
1:D:618:HIS:N	1:D:725:PRO:HG2	2.35	0.42
1:D:829:PRO:CG	1:D:831:TYR:CE1	3.02	0.42
2:E:403:PHE:HA	2:E:413:VAL:HG21	2.01	0.42
2:E:456:ARG:HG3	2:E:457:SER:H	1.85	0.42
2:E:457:SER:OG	2:E:458:ARG:N	2.52	0.42
2:E:67:GLU:HG2	2:E:70:LEU:HD12	1.97	0.42
2:F:183:GLU:O	2:F:184:ASN:OD1	2.38	0.42
2:F:185:LEU:CD1	2:F:236:GLU:HB2	2.46	0.42
2:F:357:THR:O	2:F:358:GLU:CB	2.66	0.42
2:E:203:ARG:NH1	2:F:418:LEU:HD13	2.34	0.42
1:A:1090:GLU:O	1:A:1091:GLU:CB	2.68	0.42
1:A:1113:LEU:CB	1:A:1117:LYS:HZ2	2.28	0.42
1:A:107:LEU:O	1:A:111:GLY:CA	2.67	0.42
1:A:209:PRO:C	1:A:210:THR:CG2	2.87	0.42
1:A:554:GLY:O	1:A:557:GLU:CG	2.68	0.42
1:A:581:ASP:CA	1:A:583:PRO:HD2	2.49	0.42
1:A:604:ALA:CB	1:A:717:LEU:HD13	2.49	0.42
1:A:608:ASP:HA	1:A:779:LEU:HD22	1.99	0.42
1:A:823:ARG:O	1:A:824:ALA:CB	2.68	0.42
1:A:864:ASN:HB2	1:A:865:ALA:H	1.58	0.42
1:A:884:TYR:O	1:A:885:THR:HG23	2.19	0.42
1:A:892:ASP:OD1	1:A:892:ASP:N	2.42	0.42
1:A:951:TYR:C	1:A:955:TYR:CD1	2.93	0.42
2:B:93:SER:C	2:B:234:LYS:HZ1	2.23	0.42
2:B:238:SER:CA	2:B:339:ASN:ND2	2.83	0.42
2:B:392:THR:HG22	2:B:393:LEU:N	2.35	0.42
2:B:396:ARG:CD	2:B:418:LEU:H	2.25	0.42
2:C:127:PRO:HD2	2:C:128:VAL:H	1.81	0.42
2:C:322:LEU:HD22	2:C:324:GLY:N	2.34	0.42
2:C:347:LEU:N	2:C:347:LEU:HD12	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:419:GLU:HG3	2:C:432:TYR:OH	2.19	0.42
2:B:201:ASN:HD22	2:C:421:MET:CE	2.33	0.42
2:E:395:LEU:HD13	2:E:395:LEU:O	2.20	0.42
2:B:471:LEU:CD1	2:B:474:PHE:HD2	2.32	0.42
2:E:310:GLU:C	2:E:314:MET:HG3	2.40	0.42
2:F:272:ASN:CA	2:F:292:ASN:ND2	2.83	0.42
1:D:818:ARG:HE	1:D:818:ARG:HB2	1.62	0.42
1:D:1015:THR:O	1:D:1016:GLU:HB2	2.19	0.42
1:D:285:GLN:HE22	1:D:1128:ARG:CB	2.32	0.42
1:D:1124:ALA:HB2	1:D:1231:LEU:HB3	2.02	0.42
1:D:239:LEU:O	1:D:242:ALA:N	2.47	0.42
1:D:285:GLN:NE2	1:D:1128:ARG:CB	2.83	0.42
1:D:289:MET:CE	1:D:290:ARG:N	2.83	0.42
1:D:348:LEU:HD13	1:D:350:ILE:CD1	2.48	0.42
1:D:387:GLU:O	1:D:390:GLN:CG	2.67	0.42
1:D:462:SER:O	1:D:466:LEU:HD13	2.20	0.42
1:D:538:GLU:CD	1:D:542:ASP:CG	2.78	0.42
1:D:576:LEU:HD22	2:F:481:SER:CB	2.49	0.42
1:D:801:TRP:HE3	1:D:804:ALA:HB3	1.85	0.42
1:D:871:GLY:O	1:D:872:SER:C	2.57	0.42
1:D:955:TYR:HB3	1:D:1098:ASN:OD1	2.19	0.42
2:E:250:GLN:CD	2:E:250:GLN:C	2.78	0.42
2:E:109:ASN:HB2	2:E:378:LEU:O	2.20	0.42
2:E:398:VAL:HG22	2:E:402:LEU:HG	2.02	0.42
2:E:406:LEU:HG	2:E:413:VAL:HG21	2.01	0.42
2:E:95:CYS:O	2:E:96:HIS:C	2.58	0.42
2:F:263:TRP:O	2:F:264:ARG:C	2.58	0.42
2:F:442:LEU:CB	2:F:455:LEU:HD22	2.49	0.42
1:A:1022:LEU:HB3	1:A:1026:ARG:HH22	1.85	0.42
1:A:267:VAL:HB	1:A:269:HIS:CE1	2.55	0.42
1:A:269:HIS:O	1:A:293:ASP:HA	2.19	0.42
1:A:474:LEU:HB3	1:A:712:VAL:H	1.85	0.42
1:A:805:HIS:CG	1:A:806:LYS:N	2.88	0.42
1:A:275:ARG:HH22	1:A:845:VAL:CA	2.23	0.42
1:A:962:ALA:CB	1:A:986:TYR:CZ	2.98	0.42
2:B:289:LEU:CB	2:B:300:ILE:HG13	2.50	0.42
2:B:374:LEU:H	2:B:458:ARG:CZ	2.20	0.42
2:B:424:SER:OG	2:B:425:LEU:N	2.52	0.42
2:B:435:MET:O	2:B:437:ILE:N	2.53	0.42
2:B:461:THR:OG1	2:B:463:LYS:HE3	2.20	0.42
2:C:241:TRP:CD1	2:C:243:THR:CG2	3.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:306:LEU:C	2:C:335:VAL:HB	2.40	0.42
2:C:86:LEU:O	2:C:87:SER:C	2.58	0.42
1:D:666:GLN:HA	1:D:669:MET:HG3	2.00	0.42
2:B:466:MET:CG	2:B:467:HIS:N	2.80	0.42
2:C:59:ALA:N	2:C:60:PRO:HD2	2.35	0.42
1:A:284:ILE:O	1:A:284:ILE:HG22	2.19	0.42
2:E:472:LYS:NZ	2:E:473:ASP:OD1	2.53	0.42
1:D:1015:THR:C	1:D:1017:GLY:H	2.23	0.42
1:D:1056:GLU:CA	1:D:1059:ASN:HD21	2.24	0.42
1:D:1102:GLN:O	1:D:1103:SER:C	2.56	0.42
1:D:1151:LEU:HD21	1:D:1224:ILE:HB	2.02	0.42
1:D:175:TRP:HZ2	1:D:183:GLU:HG3	1.79	0.42
1:D:278:ILE:CA	1:D:291:PHE:CD1	3.03	0.42
1:D:380:GLY:O	1:D:381:THR:CG2	2.58	0.42
1:D:472:GLN:O	2:F:461:THR:CG2	2.68	0.42
1:D:544:MET:O	1:D:548:CYS:N	2.52	0.42
1:D:631:LEU:O	1:D:633:LYS:N	2.53	0.42
1:D:634:LEU:HA	1:D:635:PRO:HD3	1.78	0.42
2:F:203:ARG:HA	2:F:325:ARG:CG	2.47	0.42
2:F:300:ILE:N	2:F:344:ARG:HH12	2.18	0.42
2:F:374:LEU:HB2	2:F:379:ALA:HB2	2.02	0.42
2:F:408:GLU:O	2:F:409:ASN:C	2.58	0.42
2:F:427:GLN:O	2:F:430:SER:OG	2.28	0.42
2:F:436:SER:O	2:F:437:ILE:C	2.58	0.42
2:F:473:ASP:O	2:F:476:ILE:HD12	2.20	0.42
1:A:1013:ASP:CG	1:A:1014:ARG:N	2.73	0.42
1:A:1219:ASP:O	1:A:1223:ILE:CG2	2.65	0.42
1:A:204:ALA:C	1:A:206:GLY:N	2.72	0.42
1:A:290:ARG:O	1:A:291:PHE:CD2	2.73	0.42
1:A:316:LYS:HB2	1:A:348:LEU:HD11	2.02	0.42
1:A:312:TRP:CZ2	1:A:346:ASP:O	2.54	0.42
1:A:357:SER:O	1:A:361:VAL:HG12	2.20	0.42
1:A:593:SER:C	1:A:594:LEU:HD23	2.40	0.42
1:A:869:ARG:NE	1:A:870:VAL:HG23	2.35	0.42
1:A:996:ARG:HD2	1:A:1004:LEU:HD13	2.01	0.42
2:B:261:GLN:OE1	2:B:266:PHE:CE1	2.73	0.42
2:C:300:ILE:CG2	2:C:342:LEU:CD1	2.83	0.42
2:C:322:LEU:HD22	2:C:323:HIS:N	2.35	0.42
2:C:325:ARG:NH2	2:C:327:GLY:HA2	2.35	0.42
2:C:361:PHE:C	2:C:362:THR:HG23	2.40	0.42
2:C:433:ASP:C	2:C:435:MET:N	2.72	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:72:ILE:HG23	2:C:76:ARG:HE	1.84	0.42
1:D:525:MET:O	1:D:526:ASP:CG	2.57	0.42
1:A:500:ALA:HB1	1:A:520:ALA:HB2	2.02	0.42
2:E:308:ASP:O	2:E:309:HIS:O	2.38	0.42
1:A:941:ILE:HG22	1:A:942:SER:O	2.20	0.42
2:B:118:VAL:HG21	2:B:239:LEU:HD22	2.02	0.42
2:B:107:ARG:HB2	2:B:346:MET:SD	2.60	0.42
1:D:1057:MET:CB	1:D:1064:ILE:HD11	2.29	0.42
1:D:1103:SER:O	1:D:1104:SER:C	2.57	0.42
1:D:289:MET:HE2	1:D:290:ARG:N	2.35	0.42
1:D:739:TYR:C	1:D:740:ASN:CG	2.79	0.42
1:D:770:PHE:CE1	1:D:968:GLN:CA	3.03	0.42
1:D:797:MET:H	1:D:797:MET:HG3	1.49	0.42
1:D:813:VAL:HG21	1:D:815:TRP:CZ2	2.55	0.42
1:D:841:LEU:HA	1:D:842:PRO:HD3	1.72	0.42
1:D:967:MET:SD	1:D:978:ALA:CB	3.07	0.42
2:E:121:PHE:HA	2:F:407:LEU:CD1	2.49	0.42
2:E:471:LEU:HD12	2:E:471:LEU:O	2.20	0.42
2:E:80:SER:O	2:E:86:LEU:HD21	2.20	0.42
2:F:184:ASN:O	2:F:185:LEU:C	2.58	0.42
1:D:546:ARG:NH2	2:F:408:GLU:OE1	2.53	0.42
2:F:468:ILE:C	2:F:471:LEU:HD13	2.41	0.42
1:A:1025:LEU:O	1:A:1026:ARG:CG	2.63	0.42
1:A:955:TYR:CB	1:A:1098:ASN:OD1	2.65	0.42
1:A:1135:ASP:N	1:A:1136:GLU:OE2	2.45	0.42
1:A:1207:GLU:H	1:A:1210:TYR:H	1.66	0.42
1:A:199:VAL:HG21	1:A:277:HIS:CB	2.50	0.42
1:A:213:VAL:HG13	1:A:397:ALA:CB	2.44	0.42
1:A:394:GLN:OE1	1:A:395:TYR:HA	2.20	0.42
1:A:607:TRP:HZ2	1:A:671:GLN:N	2.18	0.42
1:A:608:ASP:OD2	1:A:614:TYR:OH	2.29	0.42
2:B:182:ARG:HD3	2:B:214:CYS:HA	2.02	0.42
2:B:128:VAL:HG12	2:B:209:ALA:O	2.20	0.42
2:C:198:ASP:OD1	2:C:198:ASP:O	2.38	0.42
2:C:252:LEU:CD2	2:C:252:LEU:C	2.89	0.42
2:C:456:ARG:CG	2:C:457:SER:N	2.79	0.42
2:F:275:SER:HA	2:F:288:LYS:O	2.20	0.42
1:D:1037:GLN:H	1:D:1037:GLN:HG2	1.38	0.42
1:D:873:GLU:N	1:D:1200:PRO:HD3	2.35	0.41
1:D:172:ALA:HB1	1:D:221:TYR:HB2	2.02	0.41
1:D:206:GLY:C	1:D:208:CYS:H	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:245:ILE:HG13	1:D:246:PRO:CG	2.47	0.41
1:D:288:ARG:NH1	1:D:290:ARG:HH22	2.18	0.41
1:D:420:HIS:CD2	1:D:1020:ILE:CG2	3.02	0.41
1:D:434:TYR:CD1	1:D:841:LEU:HD21	2.55	0.41
1:D:457:ARG:C	1:D:460:LYS:NZ	2.73	0.41
1:D:461:LYS:HE3	1:D:461:LYS:HB3	1.84	0.41
1:D:488:LEU:HD23	1:D:488:LEU:N	2.34	0.41
1:D:808:ILE:HD13	1:D:874:LEU:CD2	2.50	0.41
2:E:379:ALA:HB3	2:E:438:LEU:HD11	2.02	0.41
2:E:456:ARG:NH1	2:E:463:LYS:HE3	2.32	0.41
2:F:286:GLY:CA	2:F:303:LEU:O	2.68	0.41
2:F:293:PHE:C	2:F:295:TRP:N	2.74	0.41
2:F:197:LEU:HD12	2:F:322:LEU:HD21	2.00	0.41
2:F:335:VAL:HG22	2:F:336:LEU:N	2.35	0.41
2:F:239:LEU:HG	2:F:336:LEU:CD1	2.50	0.41
2:F:383:VAL:CG2	2:F:384:ALA:H	2.33	0.41
2:F:394:GLU:HA	2:F:397:GLN:CD	2.39	0.41
2:E:203:ARG:NH2	2:F:418:LEU:CD1	2.83	0.41
2:F:372:LEU:CG	2:F:436:SER:HB2	2.49	0.41
2:F:474:PHE:CZ	2:F:475:LEU:HD23	2.55	0.41
1:A:293:ASP:HB2	1:A:296:SER:OG	2.20	0.41
1:A:373:PRO:CD	1:A:374:ARG:H	2.32	0.41
1:A:415:LEU:C	1:A:417:ARG:N	2.73	0.41
1:A:454:GLU:O	1:A:455:LEU:C	2.57	0.41
1:A:612:LEU:CD2	1:A:614:TYR:CA	2.98	0.41
1:A:730:PRO:HG2	1:A:731:SER:H	1.85	0.41
1:A:886:LEU:CD1	1:A:1189:LEU:HD21	2.50	0.41
2:B:196:CYS:O	2:B:199:LEU:HB3	2.20	0.41
2:B:269:SER:C	2:B:271:SER:N	2.68	0.41
1:D:236:THR:O	1:D:237:SER:HB3	2.20	0.41
1:A:643:ALA:O	1:A:644:GLY:C	2.58	0.41
1:D:669:MET:HA	1:D:670:PRO:HA	1.58	0.41
1:A:1193:VAL:HG22	1:A:1194:THR:H	1.85	0.41
2:E:476:ILE:O	2:E:480:SER:N	2.43	0.41
1:D:1019:TRP:NE1	1:D:1020:ILE:HG22	2.32	0.41
1:D:1122:GLU:OE2	1:D:1122:GLU:CA	2.68	0.41
1:D:1220:ILE:HB	1:D:1221:TYR:H	1.41	0.41
1:D:209:PRO:O	1:D:210:THR:OG1	2.38	0.41
1:D:345:TRP:O	1:D:345:TRP:CE3	2.73	0.41
1:D:444:TYR:CE2	1:D:877:MET:SD	3.13	0.41
1:D:546:ARG:HH21	2:F:404:ASN:ND2	2.18	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:611:PRO:HG2	1:D:622:TYR:CE2	2.55	0.41
1:D:631:LEU:O	1:D:633:LYS:HG3	2.20	0.41
1:D:797:MET:HB2	1:D:869:ARG:HH11	1.85	0.41
1:D:896:LEU:HD13	1:D:896:LEU:HA	1.85	0.41
2:E:213:VAL:HG13	2:E:235:THR:HG23	1.96	0.41
2:F:201:ASN:O	2:F:202:LYS:CB	2.68	0.41
2:F:238:SER:HA	2:F:338:VAL:CB	2.49	0.41
2:F:299:LEU:CD2	2:F:299:LEU:H	2.09	0.41
2:F:308:ASP:OD2	2:F:334:CYS:SG	2.76	0.41
1:A:955:TYR:HE2	1:A:1102:GLN:HE21	1.68	0.41
1:A:1230:SER:HB3	1:A:1231:LEU:H	1.58	0.41
1:A:177:ARG:O	1:A:179:GLY:N	2.53	0.41
1:A:196:VAL:HA	1:A:267:VAL:HG12	2.02	0.41
1:A:449:GLN:HA	1:A:452:TYR:CD1	2.53	0.41
1:A:485:LEU:O	1:A:489:GLU:HG3	2.20	0.41
1:A:630:ASN:C	1:A:630:ASN:ND2	2.74	0.41
1:A:784:GLY:N	2:C:362:THR:O	2.54	0.41
1:A:792:LEU:O	1:A:796:LYS:CD	2.67	0.41
1:A:906:HIS:C	1:A:908:ALA:H	2.23	0.41
2:B:259:ARG:HD2	2:B:262:TRP:HB2	2.01	0.41
2:B:76:ARG:NH2	2:B:431:LYS:HG3	2.35	0.41
2:C:102:LEU:HD11	2:C:435:MET:SD	2.60	0.41
2:C:121:PHE:CE1	2:C:122:ARG:HD2	2.55	0.41
2:C:125:VAL:CG1	2:C:206:TYR:HD1	2.33	0.41
2:C:208:LEU:HG	2:C:208:LEU:H	0.84	0.41
2:C:208:LEU:CD1	2:C:240:VAL:C	2.80	0.41
2:C:473:ASP:O	2:C:477:LYS:HG3	2.20	0.41
2:E:389:ARG:HH21	2:E:391:PRO:C	2.20	0.41
2:E:415:PRO:HB2	2:E:417:TYR:CZ	2.55	0.41
2:C:272:ASN:HB3	2:C:292:ASN:CB	2.47	0.41
1:D:765:PRO:HG2	1:D:766:PHE:CG	2.55	0.41
2:E:476:ILE:O	2:E:480:SER:HB3	2.19	0.41
1:D:1129:PHE:CE2	1:D:1139:TYR:CD2	3.09	0.41
1:D:1154:GLN:HG2	1:D:1179:PHE:HE2	1.82	0.41
1:D:444:TYR:CE2	1:D:1210:TYR:HB2	2.55	0.41
1:D:180:PRO:O	1:D:181:GLU:CD	2.59	0.41
1:D:256:PRO:HB3	1:D:281:GLN:CA	2.49	0.41
1:D:267:VAL:CB	1:D:269:HIS:HD2	2.14	0.41
1:D:275:ARG:NE	1:D:842:PRO:C	2.72	0.41
1:D:451:THR:HA	1:D:454:GLU:HG3	2.01	0.41
1:D:468:ASN:O	1:D:472:GLN:HB2	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:722:ARG:CG	1:D:722:ARG:NH1	2.83	0.41
1:D:601:LYS:NZ	1:D:722:ARG:H	2.17	0.41
1:D:74:LEU:CD1	1:D:76:ILE:O	2.67	0.41
1:D:812:MET:O	1:D:840:ILE:CG2	2.58	0.41
1:D:79:LEU:CB	1:D:83:LEU:HD11	2.46	0.41
2:E:203:ARG:CZ	2:E:203:ARG:CB	2.97	0.41
2:E:259:ARG:NH2	2:E:301:GLU:OE1	2.53	0.41
2:E:328:ARG:O	2:E:329:LYS:HG3	2.20	0.41
2:E:87:SER:C	2:E:89:ASP:H	2.24	0.41
2:F:254:PHE:HA	2:F:257:ARG:HD2	2.02	0.41
2:F:375:HIS:N	2:F:378:LEU:HB2	2.35	0.41
2:F:386:ASP:O	2:F:443:VAL:N	2.36	0.41
2:F:403:PHE:CD2	2:F:407:LEU:HB2	2.54	0.41
2:F:426:GLU:OE2	2:F:426:GLU:HA	2.20	0.41
2:F:468:ILE:CG2	2:F:469:SER:N	2.83	0.41
2:F:470:LYS:O	2:F:474:PHE:N	2.54	0.41
2:F:475:LEU:H	2:F:475:LEU:HG	1.58	0.41
1:A:1010:LEU:C	1:A:1012:VAL:H	2.19	0.41
1:A:1028:VAL:HG23	1:A:1029:GLN:HG3	2.01	0.41
1:A:1108:TYR:CE2	1:A:1112:MET:HE1	2.55	0.41
1:A:1128:ARG:O	1:A:1139:TYR:HA	2.21	0.41
1:A:110:HIS:C	1:A:112:LEU:HD12	2.41	0.41
1:A:1195:MET:O	1:A:1196:ASP:CB	2.67	0.41
1:A:196:VAL:HA	1:A:269:HIS:NE2	2.35	0.41
1:A:239:LEU:H	1:A:239:LEU:HG	1.55	0.41
1:A:253:ALA:CB	1:A:256:PRO:HB3	2.37	0.41
1:A:272:SER:O	1:A:274:ASP:N	2.52	0.41
1:A:481:GLU:C	1:A:482:ASP:OD1	2.59	0.41
1:A:620:TRP:HD1	1:A:749:PHE:O	2.03	0.41
1:A:726:LYS:C	1:A:727:ASP:O	2.57	0.41
1:A:829:PRO:HB3	1:A:883:GLY:N	2.35	0.41
1:A:913:CYS:O	1:A:917:GLY:N	2.54	0.41
2:C:188:GLY:C	2:C:190:LEU:N	2.70	0.41
2:C:208:LEU:CD1	2:C:208:LEU:H	1.98	0.41
2:C:413:VAL:CG1	2:C:414:TRP:N	2.83	0.41
2:C:433:ASP:C	2:C:435:MET:H	2.23	0.41
2:C:134:LYS:HD3	2:C:180:LYS:HZ2	1.86	0.41
2:E:264:ARG:HG2	2:E:273:PHE:CG	2.55	0.41
1:D:500:ALA:HB2	1:D:521:PRO:CD	2.49	0.41
2:C:222:LYS:HB2	2:C:223:GLN:H	1.56	0.41
1:A:105:GLU:O	1:A:109:LYS:HE2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1043:VAL:HG23	1:D:1043:VAL:O	2.20	0.41
1:D:869:ARG:HA	1:D:1200:PRO:HB2	2.02	0.41
1:D:247:LEU:N	1:D:247:LEU:HD12	2.35	0.41
1:D:265:LEU:HG	1:D:267:VAL:N	2.35	0.41
1:D:387:GLU:O	1:D:390:GLN:HG2	2.20	0.41
1:D:863:SER:O	1:D:875:LYS:HE2	2.20	0.41
2:E:212:GLY:O	2:E:235:THR:C	2.58	0.41
2:E:302:THR:O	2:E:338:VAL:HA	2.20	0.41
2:E:447:THR:CG2	2:E:452:LEU:H	2.32	0.41
2:E:455:LEU:O	2:E:463:LYS:CA	2.59	0.41
2:F:114:TRP:HB2	2:F:262:TRP:HZ2	1.77	0.41
2:F:193:TYR:O	2:F:193:TYR:CD1	2.74	0.41
2:F:204:LEU:H	2:F:204:LEU:HD12	1.84	0.41
2:F:289:LEU:O	2:F:299:LEU:C	2.58	0.41
2:F:360:SER:OG	2:F:364:LYS:HD2	2.21	0.41
2:F:432:TYR:O	2:F:436:SER:N	2.53	0.41
2:F:82:SER:CA	2:F:86:LEU:CD2	2.77	0.41
1:A:1041:TRP:NE1	1:A:1043:VAL:CG2	2.83	0.41
1:A:1093:MET:HG3	1:A:1093:MET:H	1.68	0.41
1:A:1122:GLU:C	1:A:1123:PHE:HD1	2.20	0.41
1:A:1158:LEU:HD23	1:A:1158:LEU:N	2.34	0.41
1:A:1233:LYS:C	1:A:1236:GLN:NE2	2.73	0.41
1:A:175:TRP:O	1:A:176:THR:CB	2.68	0.41
1:A:259:ARG:O	1:A:260:ASP:CB	2.68	0.41
1:A:371:LYS:HB2	1:A:374:ARG:HD2	2.03	0.41
1:A:417:ARG:C	1:A:419:PRO:HD3	2.41	0.41
1:A:425:ALA:O	1:A:428:LEU:HB2	2.21	0.41
1:A:497:GLN:O	1:A:498:LYS:CG	2.68	0.41
1:A:560:PRO:O	1:A:561:LYS:HB2	2.20	0.41
1:A:606:THR:HG22	1:A:614:TYR:CE2	2.55	0.41
1:A:617:ARG:HB3	1:A:763:GLY:C	2.41	0.41
1:A:441:TRP:HD1	1:A:877:MET:SD	2.44	0.41
1:A:944:GLU:O	1:A:948:ILE:HG12	2.21	0.41
2:B:183:GLU:C	2:B:214:CYS:SG	2.99	0.41
2:C:197:LEU:C	2:C:199:LEU:N	2.72	0.41
2:C:238:SER:HB2	2:C:338:VAL:HG11	2.01	0.41
2:C:432:TYR:HB3	2:C:437:ILE:CD1	2.50	0.41
1:A:556:THR:CG2	2:C:467:HIS:NE2	2.84	0.41
1:A:645:VAL:CG1	1:A:647:CYS:O	2.68	0.41
1:D:97:GLU:CB	1:D:101:ARG:NH1	2.83	0.41
2:C:282:GLU:C	2:C:284:ARG:N	2.71	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:282:GLU:C	2:F:284:ARG:H	2.22	0.41
1:D:531:GLY:HA2	1:D:532:PRO:HD3	1.68	0.41
1:D:1005:VAL:O	1:D:1006:ARG:C	2.55	0.41
1:D:1132:SER:CA	1:D:1136:GLU:O	2.62	0.41
1:D:1200:PRO:HD2	1:D:1203:PRO:CB	2.51	0.41
1:D:255:SER:HB3	1:D:256:PRO:O	2.18	0.41
1:D:269:HIS:O	1:D:274:ASP:OD2	2.38	0.41
1:D:823:ARG:CD	1:D:824:ALA:H	2.32	0.41
1:D:902:LEU:HA	1:D:902:LEU:HD12	1.80	0.41
2:E:323:HIS:O	2:E:330:ASN:HA	2.20	0.41
2:E:325:ARG:CG	2:E:330:ASN:H	2.34	0.41
2:E:354:PHE:HB2	2:E:372:LEU:CD1	2.50	0.41
2:F:78:PHE:HD1	2:F:102:LEU:HD22	1.78	0.41
2:F:131:LEU:HD12	2:F:131:LEU:HA	1.88	0.41
2:F:247:THR:HG1	2:F:248:SER:N	2.19	0.41
2:F:460:THR:C	2:F:461:THR:HG23	2.41	0.41
1:A:139:PHE:CD2	1:A:1111:LEU:HG	2.56	0.41
1:A:1131:ILE:HG23	1:A:1137:VAL:CG1	2.49	0.41
1:A:1179:PHE:CD1	1:A:1220:ILE:HG12	2.55	0.41
1:A:255:SER:N	1:A:256:PRO:HA	2.35	0.41
1:A:606:THR:HG21	1:A:613:HIS:N	2.25	0.41
1:A:597:ARG:HB2	1:A:616:GLU:OE1	2.20	0.41
1:A:618:HIS:O	1:A:762:VAL:HG21	2.20	0.41
1:A:659:HIS:HD2	1:A:720:THR:H	1.67	0.41
1:A:667:GLN:CG	1:A:668:LEU:N	2.82	0.41
1:A:765:PRO:HG2	1:A:766:PHE:N	2.36	0.41
1:A:769:ASP:OD1	1:A:1057:MET:HE2	2.20	0.41
2:B:125:VAL:CG1	2:B:126:PHE:N	2.82	0.41
2:B:431:LYS:HG2	2:B:435:MET:HG3	2.02	0.41
2:B:436:SER:C	2:B:458:ARG:HH11	2.23	0.41
2:C:115:TRP:CD1	2:C:119:VAL:CG2	3.03	0.41
2:C:208:LEU:CD1	2:C:208:LEU:N	2.57	0.41
2:C:413:VAL:HG12	2:C:414:TRP:H	1.84	0.41
1:D:163:PRO:O	1:D:165:LYS:HE3	2.21	0.41
1:A:739:TYR:CA	1:A:741:ASP:O	2.67	0.41
1:A:156:LEU:CD2	1:A:156:LEU:N	2.83	0.41
1:A:1183:VAL:O	1:A:1217:ALA:CB	2.68	0.41
1:A:1184:ASP:N	1:A:1184:ASP:OD1	2.54	0.41
2:E:341:ASP:C	2:E:341:ASP:OD1	2.59	0.41
1:A:345:TRP:CD1	1:A:345:TRP:O	2.74	0.41
1:D:1165:ALA:HB1	1:D:1173:LEU:HD12	1.91	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:140:ARG:O	1:D:144:GLN:CB	2.61	0.41
1:D:146:GLN:H	1:D:146:GLN:HG2	1.44	0.41
1:D:295:MET:O	1:D:298:HIS:N	2.54	0.41
1:D:203:LEU:CB	1:D:379:LYS:O	2.69	0.41
1:D:391:ASP:O	1:D:395:TYR:CD1	2.74	0.41
1:D:495:PHE:O	1:D:573:TYR:CZ	2.73	0.41
1:D:611:PRO:CB	1:D:612:LEU:HD13	2.37	0.41
1:D:738:PRO:C	1:D:739:TYR:O	2.57	0.41
1:D:622:TYR:HB2	1:D:747:CYS:O	2.21	0.41
1:D:81:ARG:HD3	1:D:126:ARG:HA	2.01	0.41
1:D:444:TYR:CD2	1:D:877:MET:SD	3.14	0.41
1:D:950:ASN:O	1:D:953:ARG:N	2.53	0.41
2:E:106:LEU:HD12	2:E:109:ASN:ND2	2.36	0.41
2:E:123:GLU:O	2:E:125:VAL:O	2.39	0.41
2:E:201:ASN:OD1	2:F:422:GLN:CD	2.59	0.41
2:E:205:PRO:HB3	2:E:242:PHE:C	2.41	0.41
2:E:353:SER:HA	2:E:373:LYS:O	2.20	0.41
2:E:370:LYS:O	2:E:433:ASP:CG	2.58	0.41
2:E:406:LEU:HG	2:E:413:VAL:HG22	2.01	0.41
2:F:336:LEU:HD21	2:F:337:SER:O	2.21	0.41
1:A:756:ASP:OD2	1:A:1057:MET:HG2	2.20	0.41
1:A:879:GLN:CG	1:A:1189:LEU:CD2	2.79	0.41
1:A:1198:LYS:O	1:A:1199:THR:CB	2.68	0.41
1:A:174:GLY:O	1:A:220:TRP:HB2	2.21	0.41
1:A:256:PRO:CB	1:A:281:GLN:HA	2.46	0.41
1:A:439:GLN:HG3	1:A:836:LEU:O	2.20	0.41
1:A:670:PRO:O	1:A:671:GLN:C	2.59	0.41
1:A:792:LEU:O	1:A:796:LYS:HD3	2.21	0.41
1:A:904:ASP:OD1	1:A:904:ASP:N	2.51	0.41
2:B:261:GLN:CA	2:B:264:ARG:NE	2.70	0.41
2:C:393:LEU:O	2:C:397:GLN:HB3	2.21	0.41
2:C:452:LEU:HD12	2:C:453:ILE:HG12	2.02	0.41
1:D:522:GLY:C	1:D:524:PRO:N	2.74	0.41
2:E:389:ARG:N	2:E:395:LEU:CD1	2.77	0.41
2:E:418:LEU:HD23	2:E:418:LEU:HA	1.88	0.41
2:B:471:LEU:HA	2:B:474:PHE:HB3	2.01	0.41
2:F:61:GLY:CA	2:F:64:GLU:HG2	2.51	0.41
1:A:985:MET:H	1:A:985:MET:HG3	1.73	0.41
1:D:1045:ALA:HB1	1:D:1092:PHE:CE1	2.55	0.41
1:D:1097:VAL:O	1:D:1098:ASN:C	2.57	0.41
1:D:1193:VAL:O	1:D:1195:MET:CE	2.69	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:138:HIS:O	1:D:142:LEU:CG	2.52	0.41
1:D:270:ASN:C	1:D:274:ASP:CG	2.79	0.41
1:D:350:ILE:HG22	1:D:352:SER:CB	2.40	0.41
1:D:374:ARG:NE	1:D:392:LEU:CD2	2.83	0.41
1:D:466:LEU:N	1:D:466:LEU:CD1	2.82	0.41
1:D:490:TRP:O	1:D:580:LEU:CA	2.68	0.41
1:D:637:GLY:O	1:D:641:GLU:HG3	2.21	0.41
1:D:752:LEU:HB3	1:D:753:PRO:CB	2.49	0.41
1:D:770:PHE:O	1:D:773:LYS:HG2	2.20	0.41
1:D:897:TRP:HH2	1:D:1173:LEU:CD1	2.26	0.41
1:D:989:THR:C	1:D:998:SER:OG	2.59	0.41
2:E:259:ARG:NE	2:E:301:GLU:OE1	2.54	0.41
2:E:76:ARG:HG3	2:E:435:MET:HG3	2.03	0.41
2:E:441:VAL:HG22	2:E:455:LEU:CB	2.51	0.41
2:F:114:TRP:CD1	2:F:259:ARG:NH1	2.81	0.41
2:F:197:LEU:C	2:F:199:LEU:N	2.74	0.41
2:F:350:LEU:CD1	2:F:351:TYR:H	2.32	0.41
2:F:446:THR:CB	2:F:450:ASN:OD1	2.69	0.41
2:F:482:ALA:O	2:F:483:LYS:C	2.59	0.41
2:E:129:ASP:OD1	2:F:98:GLY:HA3	2.21	0.41
1:A:1125:ILE:C	1:A:1126:ASP:OD1	2.59	0.41
1:A:211:LEU:HA	1:A:222:SER:H	1.86	0.41
1:A:276:ALA:CB	1:A:279:ARG:HG2	2.51	0.41
1:A:292:LEU:HD11	1:A:428:LEU:CD2	2.51	0.41
1:A:412:PRO:O	1:A:415:LEU:N	2.53	0.41
1:A:456:GLN:HE21	1:A:798:ILE:HD11	1.86	0.41
1:A:465:ASP:OD1	1:A:466:LEU:N	2.54	0.41
1:A:745:PRO:HD2	1:A:746:GLY:H	1.85	0.41
1:A:78:MET:O	1:A:79:LEU:HD23	2.21	0.41
1:A:817:PRO:CG	1:A:818:ARG:N	2.84	0.41
1:A:900:ALA:HB1	1:A:915:ALA:HB1	2.02	0.41
1:A:928:GLY:C	1:A:930:ASP:N	2.74	0.41
1:A:931:LEU:CD2	1:A:932:HIS:N	2.77	0.41
2:C:113:GLU:O	2:C:116:THR:CG2	2.59	0.41
2:C:290:TYR:CD1	2:C:299:LEU:HD22	2.56	0.41
2:C:240:VAL:C	2:C:336:LEU:CB	2.83	0.41
2:C:344:ARG:N	2:C:347:LEU:HD22	2.35	0.41
2:C:362:THR:O	2:C:363:ARG:HD2	2.21	0.41
2:B:473:ASP:O	2:B:477:LYS:HE3	2.21	0.41
2:F:138:LEU:HA	2:F:138:LEU:HD13	1.81	0.41
1:D:1067:SER:HB3	1:D:1073:PRO:CG	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1144:GLU:HG3	1:D:1145:ASP:OD1	2.20	0.41
1:D:1200:PRO:HD2	1:D:1203:PRO:HB2	2.01	0.41
1:D:213:VAL:C	1:D:401:TRP:CZ2	2.71	0.41
1:D:385:ILE:CG2	1:D:386:ARG:N	2.83	0.41
1:D:399:ASP:O	1:D:403:THR:HG22	2.20	0.41
1:D:434:TYR:O	1:D:1130:CYS:CB	2.58	0.41
1:D:647:CYS:HB3	1:D:649:TYR:H	1.85	0.41
1:D:618:HIS:CB	1:D:725:PRO:CB	2.98	0.41
1:D:944:GLU:HG2	1:D:947:LYS:HE3	2.03	0.41
2:E:439:PHE:CZ	2:E:478:TYR:CZ	3.08	0.41
2:E:456:ARG:HG3	2:E:462:MET:CA	2.51	0.41
2:F:125:VAL:C	2:F:126:PHE:CG	2.92	0.41
2:F:447:THR:HG23	2:F:448:LEU:N	2.35	0.41
2:F:78:PHE:C	2:F:79:LEU:HD22	2.41	0.41
1:A:1032:THR:C	1:A:1034:ARG:H	2.23	0.41
1:A:851:THR:OG1	1:A:1099:TRP:CD1	2.69	0.41
1:A:1134:HIS:CB	1:A:1136:GLU:OE2	2.69	0.41
1:A:1179:PHE:CD1	1:A:1220:ILE:CG2	2.99	0.41
1:A:1200:PRO:O	1:A:1202:ASN:N	2.53	0.41
1:A:1206:MET:O	1:A:1207:GLU:CB	2.69	0.41
1:A:1208:ARG:C	1:A:1210:TYR:N	2.70	0.41
1:A:1218:LEU:O	1:A:1222:GLN:NE2	2.53	0.41
1:A:169:TRP:CD1	1:A:169:TRP:N	2.88	0.41
1:A:226:GLN:OE1	1:A:229:VAL:CG2	2.69	0.41
1:A:289:MET:O	1:A:290:ARG:HG3	2.20	0.41
1:A:143:ALA:CB	1:A:421:PRO:HB2	2.50	0.41
1:A:441:TRP:CZ2	1:A:874:LEU:CB	3.04	0.41
1:A:656:TYR:HA	1:A:659:HIS:CB	2.50	0.41
1:A:811:GLN:O	1:A:812:MET:C	2.59	0.41
1:A:848:GLY:O	1:A:849:THR:O	2.38	0.41
2:B:104:VAL:HG21	2:C:128:VAL:C	2.40	0.41
2:B:250:GLN:C	2:B:252:LEU:N	2.74	0.41
2:B:238:SER:CA	2:B:339:ASN:HD21	2.33	0.41
2:C:247:THR:HG1	2:C:251:TRP:HB2	1.86	0.41
2:C:254:PHE:O	2:C:257:ARG:HB2	2.21	0.41
2:C:289:LEU:C	2:C:290:TYR:CG	2.94	0.41
2:C:346:MET:O	2:C:350:LEU:HD11	2.20	0.41
2:C:397:GLN:C	2:C:399:CYS:N	2.73	0.41
2:C:413:VAL:HG12	2:C:415:PRO:CD	2.51	0.41
1:A:503:VAL:HG11	1:A:520:ALA:O	2.17	0.41
2:B:474:PHE:CG	2:B:475:LEU:N	2.89	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:741:ASP:CG	1:A:742:VAL:N	2.72	0.41
1:D:498:LYS:HZ2	1:D:521:PRO:HB3	1.85	0.41
1:A:539:PHE:O	1:A:539:PHE:CD2	2.73	0.41
1:A:443:ARG:HG3	1:A:447:GLU:OE1	2.21	0.41
2:F:218:VAL:HG13	2:F:218:VAL:O	2.21	0.41
1:D:1000:GLU:O	1:D:1003:TRP:CB	2.69	0.41
1:D:1067:SER:CB	1:D:1071:ARG:HB3	2.50	0.41
1:D:1163:MET:HE2	1:D:1167:LYS:HG3	2.03	0.41
1:D:1108:TYR:CA	1:D:1164:PHE:CE1	3.01	0.41
1:D:1206:MET:HG2	2:F:257:ARG:CZ	2.45	0.41
1:D:143:ALA:HB1	1:D:425:ALA:HB3	2.03	0.41
1:D:297:MET:CG	1:D:411:LEU:HD13	2.50	0.41
1:D:432:VAL:HG11	1:D:1128:ARG:NE	2.06	0.41
1:D:482:ASP:CG	1:D:483:PRO:HD2	2.42	0.41
1:D:482:ASP:C	1:D:482:ASP:OD1	2.60	0.41
1:D:598:VAL:N	1:D:616:GLU:OE2	2.53	0.41
1:D:720:THR:CA	1:D:721:ALA:O	2.69	0.41
1:D:992:LEU:C	1:D:996:ARG:NE	2.75	0.41
2:E:201:ASN:CB	2:F:421:MET:SD	3.09	0.41
2:F:401:GLY:O	2:F:402:LEU:C	2.59	0.41
2:F:481:SER:O	2:F:484:ASN:HB2	2.21	0.41
2:F:86:LEU:C	2:F:87:SER:O	2.59	0.41
1:D:1003:TRP:CH2	1:D:1101:VAL:CG2	3.04	0.41
1:D:1129:PHE:CZ	1:D:1139:TYR:CE2	3.09	0.41
1:D:1166:TYR:HA	1:D:1169:GLY:C	2.41	0.41
1:D:1171:ASN:O	1:D:1172:ASP:OD1	2.39	0.41
1:D:282:TYR:HB2	1:D:841:LEU:HD11	2.01	0.41
1:D:289:MET:C	1:D:290:ARG:HD2	2.41	0.41
1:D:546:ARG:O	1:D:550:GLN:CD	2.59	0.41
1:D:592:LEU:O	1:D:593:SER:C	2.56	0.41
1:D:612:LEU:CD1	1:D:622:TYR:HA	2.51	0.41
1:D:647:CYS:CB	1:D:648:PRO:HD2	2.51	0.41
1:D:840:ILE:HG22	1:D:841:LEU:N	2.36	0.41
1:D:913:CYS:O	1:D:918:TRP:N	2.54	0.41
1:D:914:THR:O	1:D:914:THR:CG2	2.69	0.41
1:D:923:GLY:CA	1:D:931:LEU:CD1	2.98	0.41
2:E:201:ASN:HB3	2:F:421:MET:HE1	2.02	0.41
2:E:442:LEU:HD13	2:E:443:VAL:CA	2.49	0.41
2:F:181:LEU:O	2:F:182:ARG:C	2.59	0.41
2:F:289:LEU:HA	2:F:289:LEU:HD23	1.85	0.41
2:F:303:LEU:C	2:F:304:TRP:CD1	2.93	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:336:LEU:HD23	2:F:337:SER:O	2.20	0.41
1:D:546:ARG:NH1	2:F:408:GLU:OE1	2.54	0.41
2:F:454:HIS:O	2:F:455:LEU:CB	2.49	0.41
2:F:458:ARG:CG	2:F:459:ASP:N	2.83	0.41
2:F:80:SER:HB2	2:F:98:GLY:C	2.41	0.41
2:F:86:LEU:N	2:F:86:LEU:CD1	2.84	0.41
1:A:1024:ASP:O	1:A:1026:ARG:HG3	2.21	0.41
1:A:1144:GLU:C	1:A:1146:ARG:N	2.74	0.41
1:A:1198:LYS:O	1:A:1199:THR:HG23	2.20	0.41
1:A:1231:LEU:CD1	1:A:1235:SER:HB3	2.51	0.41
1:A:312:TRP:HH2	1:A:346:ASP:N	2.18	0.41
1:A:371:LYS:HG3	1:A:372:GLU:HG2	2.02	0.41
1:A:411:LEU:C	1:A:411:LEU:CD1	2.89	0.41
1:A:814:VAL:HG22	1:A:839:ALA:O	2.21	0.41
1:A:875:LYS:HB3	1:A:1189:LEU:O	2.21	0.41
1:A:880:ALA:H	1:A:881:PRO:CD	2.32	0.41
1:A:886:LEU:CD2	1:A:886:LEU:N	2.68	0.41
1:A:993:ARG:O	1:A:994:TRP:C	2.58	0.41
2:B:186:LEU:HD23	2:B:186:LEU:O	2.21	0.41
2:B:185:LEU:CD1	2:B:238:SER:HB2	2.50	0.41
2:C:184:ASN:HB3	2:C:187:HIS:H	1.85	0.41
2:C:247:THR:HG1	2:C:251:TRP:CB	2.32	0.41
2:C:253:ASP:HA	2:C:256:LEU:CD2	2.49	0.41
2:C:237:ALA:N	2:C:341:ASP:HB3	2.36	0.41
2:C:344:ARG:HG3	2:C:344:ARG:H	1.58	0.41
2:C:354:PHE:O	2:C:356:LEU:N	2.54	0.41
2:C:358:GLU:O	2:C:361:PHE:N	2.45	0.41
2:C:72:ILE:HG23	2:C:76:ARG:CD	2.51	0.41
1:A:1038:TRP:CZ3	1:A:1041:TRP:CD2	3.09	0.41
1:A:1155:ILE:HD13	1:A:1158:LEU:HD12	2.02	0.41
1:A:136:ASP:CB	1:A:1163:MET:HE3	2.51	0.41
1:A:136:ASP:HB2	1:A:1163:MET:HE3	2.03	0.41
1:A:1166:TYR:CD2	1:A:1171:ASN:HA	2.56	0.41
1:A:197:PHE:CB	1:A:277:HIS:HD2	2.33	0.41
1:A:313:ILE:HA	1:A:316:LYS:CG	2.51	0.41
1:A:406:VAL:HG22	1:A:407:PHE:HD2	1.85	0.41
1:A:444:TYR:OH	1:A:1200:PRO:CG	2.66	0.41
1:A:601:LYS:HZ1	1:A:721:ALA:CA	2.34	0.41
1:A:607:TRP:O	1:A:610:PHE:HA	2.21	0.41
1:A:74:LEU:CD1	1:A:78:MET:N	2.84	0.41
1:A:617:ARG:CD	1:A:763:GLY:O	2.69	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:76:ILE:HG13	1:A:77:GLN:N	2.34	0.41
2:B:199:LEU:HD13	2:B:200:VAL:N	2.35	0.41
2:B:428:LEU:HD12	2:B:431:LYS:NZ	2.35	0.41
2:C:193:TYR:O	2:C:194:VAL:C	2.59	0.41
2:C:256:LEU:HD13	2:C:303:LEU:HD22	2.02	0.41
1:A:556:THR:CG2	2:C:452:LEU:HB3	2.33	0.41
1:D:515:ILE:HG12	1:D:568:GLY:HA3	2.02	0.41
1:A:501:LYS:HB3	1:A:502:LYS:H	1.44	0.41
1:A:503:VAL:C	1:A:505:LYS:H	2.23	0.41
2:E:393:LEU:O	2:E:396:ARG:CG	2.68	0.41
1:D:586:THR:O	1:D:586:THR:HG22	2.19	0.41
1:A:645:VAL:C	1:A:650:ARG:HD2	2.33	0.41
1:D:666:GLN:HA	1:D:669:MET:O	2.19	0.41
2:B:389:ARG:N	2:B:389:ARG:NE	2.68	0.41
1:A:634:LEU:HA	1:A:638:THR:OG1	2.21	0.41
2:F:326:ASP:OD2	2:F:329:LYS:O	2.39	0.41
1:A:102:ARG:HG2	1:A:102:ARG:H	1.51	0.41
1:D:500:ALA:CB	1:D:521:PRO:CD	2.98	0.41
1:A:1184:ASP:HA	1:A:1217:ALA:HB2	2.02	0.41
1:D:1212:ILE:HG13	1:D:1213:PRO:HD2	2.03	0.41
1:D:157:LEU:HD22	1:D:157:LEU:HA	1.83	0.41
1:D:274:ASP:HA	1:D:277:HIS:HD2	1.85	0.41
1:D:420:HIS:O	1:D:424:LEU:CG	2.68	0.41
1:D:444:TYR:C	1:D:446:ALA:N	2.72	0.41
1:D:477:GLU:HG2	1:D:478:ARG:H	1.86	0.41
1:D:654:SER:O	1:D:655:LEU:C	2.58	0.41
1:D:656:TYR:HA	1:D:659:HIS:HE1	1.85	0.41
1:D:72:ASN:OD1	1:D:80:SER:OG	2.31	0.41
1:D:865:ALA:HB2	1:D:872:SER:C	2.40	0.41
2:F:244:PRO:HB3	2:F:245:PRO:HD2	2.02	0.41
2:F:251:TRP:HZ3	2:F:254:PHE:CD1	2.38	0.41
2:F:291:TYR:O	2:F:297:LYS:HA	2.21	0.41
2:F:342:LEU:C	2:F:344:ARG:N	2.62	0.41
2:F:348:ALA:HA	2:F:351:TYR:HB2	2.02	0.41
2:F:453:ILE:CD1	2:F:465:MET:HB3	2.36	0.41
1:A:1113:LEU:HD22	1:A:1139:TYR:OH	2.20	0.41
1:A:1149:ALA:O	1:A:1152:ALA:N	2.54	0.41
1:A:181:GLU:CG	1:A:182:GLY:N	2.83	0.41
1:A:456:GLN:O	1:A:457:ARG:C	2.58	0.41
1:A:798:ILE:HD11	1:A:802:ARG:NE	2.30	0.41
2:C:347:LEU:CD1	2:C:347:LEU:N	2.82	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:502:LYS:CD	1:A:506:GLU:OE1	2.41	0.41
2:B:450:ASN:ND2	2:B:452:LEU:HB2	2.36	0.41
1:A:566:LEU:O	1:A:567:PRO:C	2.59	0.41
1:D:500:ALA:HB1	1:D:520:ALA:HB3	2.03	0.41
2:F:365:LYS:HG3	2:F:366:ASN:N	2.35	0.41
2:C:146:ARG:NE	2:C:228:VAL:HG11	2.36	0.41
1:A:148:LEU:HA	1:A:148:LEU:HD23	1.88	0.41
1:D:1066:THR:C	1:D:1073:PRO:HD3	2.41	0.40
1:D:174:GLY:HA2	1:D:220:TRP:N	2.36	0.40
1:D:189:ILE:HG21	1:D:258:GLN:CB	2.45	0.40
1:D:198:ASP:CB	1:D:400:VAL:CG1	2.87	0.40
1:D:408:GLN:HA	1:D:411:LEU:CD1	2.46	0.40
1:D:455:LEU:HD11	1:D:797:MET:SD	2.61	0.40
1:D:453:GLU:C	1:D:457:ARG:HG2	2.40	0.40
1:D:849:THR:HG22	1:D:850:ILE:HG12	2.03	0.40
1:D:441:TRP:CG	1:D:878:VAL:O	2.74	0.40
1:D:900:ALA:HB1	1:D:915:ALA:CA	2.51	0.40
2:E:123:GLU:O	2:E:125:VAL:N	2.54	0.40
2:E:259:ARG:HE	2:E:301:GLU:CD	2.23	0.40
2:E:319:VAL:HG23	2:E:332:VAL:HG21	2.03	0.40
2:E:380:PRO:O	2:E:382:LYS:N	2.54	0.40
2:F:239:LEU:O	2:F:336:LEU:CD2	2.46	0.40
2:F:252:LEU:HD12	2:F:305:ASN:HB2	2.01	0.40
2:F:347:LEU:H	2:F:347:LEU:HD13	1.86	0.40
2:F:69:LEU:HD21	2:F:351:TYR:HA	2.03	0.40
2:F:418:LEU:O	2:F:418:LEU:HD12	2.21	0.40
2:F:419:GLU:CG	2:F:432:TYR:OH	2.68	0.40
2:F:455:LEU:HG	2:F:456:ARG:NH2	2.36	0.40
1:A:1066:THR:CB	1:A:1071:ARG:HB3	2.50	0.40
1:A:107:LEU:HA	1:A:110:HIS:HB2	2.02	0.40
1:A:1144:GLU:O	1:A:1145:ASP:C	2.57	0.40
1:A:279:ARG:HB3	1:A:280:GLU:OE2	2.20	0.40
1:A:301:ILE:CD1	1:A:414:PHE:C	2.90	0.40
1:A:513:LEU:HA	1:A:514:PRO:HD2	1.61	0.40
1:A:576:LEU:HD12	1:A:577:CYS:H	1.77	0.40
1:A:812:MET:O	1:A:813:VAL:CB	2.69	0.40
1:A:920:THR:O	1:A:921:LEU:C	2.58	0.40
2:B:264:ARG:O	2:B:267:ALA:N	2.54	0.40
2:C:125:VAL:HG22	2:C:208:LEU:CB	2.48	0.40
2:C:241:TRP:CE3	2:C:336:LEU:CD1	3.02	0.40
2:C:291:TYR:OH	2:C:351:TYR:HB3	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:393:LEU:O	2:C:397:GLN:CB	2.69	0.40
2:C:411:ILE:HG23	2:C:411:ILE:O	2.20	0.40
1:D:585:TRP:N	1:D:585:TRP:HE3	2.18	0.40
2:B:474:PHE:CD2	2:B:475:LEU:HG	2.56	0.40
2:F:146:ARG:NE	2:F:228:VAL:HG11	2.36	0.40
2:F:187:HIS:O	2:F:188:GLY:C	2.59	0.40
1:D:1025:LEU:HD22	1:D:1091:GLU:OE2	2.21	0.40
1:D:1068:ASP:O	1:D:1071:ARG:HG2	2.21	0.40
1:D:308:GLN:HE21	1:D:1091:GLU:N	2.18	0.40
1:D:223:TRP:C	1:D:227:ARG:HD2	2.41	0.40
1:D:369:LEU:HD22	1:D:372:GLU:HG3	2.03	0.40
1:D:453:GLU:C	1:D:457:ARG:CG	2.89	0.40
1:D:483:PRO:HA	1:D:486:TRP:CD1	2.55	0.40
1:D:602:LEU:O	1:D:604:ALA:N	2.54	0.40
1:D:620:TRP:HE1	1:D:751:LYS:HB2	1.84	0.40
1:D:605:LEU:HA	1:D:782:GLY:H	1.86	0.40
1:D:894:GLN:HE22	1:D:1180:PHE:HE2	1.69	0.40
2:E:115:TRP:CD1	2:E:115:TRP:O	2.75	0.40
2:E:116:THR:O	2:E:120:VAL:N	2.47	0.40
2:E:375:HIS:CG	2:E:376:PRO:CA	3.04	0.40
2:E:380:PRO:O	2:E:382:LYS:CA	2.68	0.40
2:E:386:ASP:O	2:E:387:VAL:CG1	2.70	0.40
2:E:445:GLU:O	2:E:446:THR:C	2.58	0.40
2:E:453:ILE:O	2:E:465:MET:HA	2.21	0.40
2:F:106:LEU:HG	2:F:346:MET:CG	2.51	0.40
2:F:298:GLU:CG	2:F:344:ARG:NH2	2.84	0.40
2:F:400:GLN:C	2:F:400:GLN:NE2	2.75	0.40
2:F:403:PHE:CD2	2:F:404:ASN:N	2.89	0.40
2:F:406:LEU:N	2:F:406:LEU:HD13	2.36	0.40
2:E:126:PHE:CE2	2:F:414:TRP:CZ3	3.09	0.40
2:F:422:GLN:O	2:F:423:SER:OG	2.24	0.40
2:F:384:ALA:HB3	2:F:440:THR:HG22	2.04	0.40
2:F:441:VAL:HB	2:F:454:HIS:O	2.21	0.40
2:F:455:LEU:CA	2:F:456:ARG:NE	2.73	0.40
1:A:995:TYR:HB2	1:A:1004:LEU:HD11	2.02	0.40
1:A:1163:MET:O	1:A:1164:PHE:C	2.59	0.40
1:A:1177:VAL:O	1:A:1178:ALA:C	2.59	0.40
1:A:191:GLU:O	1:A:193:ARG:NH1	2.54	0.40
1:A:260:ASP:O	1:A:260:ASP:OD1	2.39	0.40
1:A:260:ASP:CA	1:A:264:GLN:NE2	2.84	0.40
1:A:313:ILE:CD1	1:A:314:ALA:N	2.85	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:423:THR:HB	1:A:424:LEU:HD23	2.03	0.40
1:A:486:TRP:HA	1:A:489:GLU:OE2	2.21	0.40
1:A:610:PHE:CB	1:A:611:PRO:HD3	2.51	0.40
1:A:78:MET:SD	1:A:123:VAL:HG11	2.61	0.40
1:A:812:MET:CG	1:A:813:VAL:N	2.84	0.40
1:A:813:VAL:HB	1:A:840:ILE:CG1	2.43	0.40
1:A:79:LEU:HB2	1:A:89:GLY:CA	2.50	0.40
2:B:261:GLN:O	2:B:265:LYS:HA	2.21	0.40
2:C:254:PHE:N	2:C:257:ARG:HH21	2.19	0.40
2:C:299:LEU:O	2:C:300:ILE:C	2.59	0.40
2:C:343:ASP:HB2	2:C:347:LEU:HD11	2.02	0.40
2:C:69:LEU:HD21	2:C:351:TYR:HA	2.03	0.40
2:C:389:ARG:O	2:C:392:THR:HA	2.21	0.40
2:C:437:ILE:O	2:C:437:ILE:HG13	2.21	0.40
1:A:633:LYS:O	1:A:634:LEU:CD1	2.70	0.40
2:B:481:SER:O	2:B:484:ASN:HB2	2.21	0.40
1:D:1066:THR:CG2	1:D:1066:THR:O	2.69	0.40
1:D:851:THR:HG21	1:D:1102:GLN:CG	2.51	0.40
1:D:1163:MET:HE2	1:D:1167:LYS:CG	2.51	0.40
1:D:1224:ILE:HG12	1:D:1228:LYS:HA	2.03	0.40
1:D:175:TRP:HA	1:D:185:VAL:CG2	2.48	0.40
1:D:193:ARG:HD2	1:D:263:GLU:OE2	2.21	0.40
1:D:266:VAL:HA	1:D:289:MET:CE	2.52	0.40
1:D:350:ILE:CA	1:D:352:SER:H	2.33	0.40
1:D:484:TRP:NE1	1:D:488:LEU:CG	2.84	0.40
1:D:597:ARG:CB	1:D:616:GLU:CD	2.89	0.40
1:D:734:HIS:HA	1:D:752:LEU:HD11	2.03	0.40
2:E:370:LYS:O	2:E:433:ASP:C	2.60	0.40
2:F:180:LYS:O	2:F:180:LYS:HE3	2.17	0.40
2:F:315:TYR:N	2:F:315:TYR:CD2	2.89	0.40
2:F:350:LEU:CD1	2:F:351:TYR:N	2.85	0.40
2:F:431:LYS:HD3	2:F:432:TYR:N	2.36	0.40
2:F:473:ASP:O	2:F:476:ILE:CD1	2.69	0.40
1:A:1170:LEU:N	1:A:1170:LEU:HD12	2.37	0.40
1:A:174:GLY:CA	1:A:220:TRP:HB2	2.50	0.40
1:A:258:GLN:O	1:A:259:ARG:NH2	2.54	0.40
1:A:261:TRP:CA	1:A:261:TRP:CE3	3.00	0.40
1:A:273:PHE:CA	1:A:276:ALA:H	2.33	0.40
1:A:352:SER:C	1:A:358:LEU:CD2	2.88	0.40
1:A:822:PRO:O	1:A:825:VAL:HB	2.22	0.40
1:A:851:THR:O	1:A:852:ARG:CB	2.69	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:755:LYS:CD	1:A:964:ARG:NH1	2.67	0.40
2:B:192:HIS:O	2:B:193:TYR:C	2.58	0.40
2:B:204:LEU:CD1	2:B:324:GLY:HA3	2.51	0.40
2:B:69:LEU:CA	2:B:72:ILE:HG12	2.52	0.40
2:C:190:LEU:O	2:C:191:GLU:C	2.58	0.40
2:B:77:HIS:CB	2:C:195:ASN:OD1	2.67	0.40
2:C:197:LEU:HD11	2:C:322:LEU:CD1	2.51	0.40
2:C:200:VAL:O	2:C:201:ASN:CB	2.69	0.40
2:C:406:LEU:HD11	2:C:472:LYS:HZ2	1.85	0.40
2:C:403:PHE:N	2:C:406:LEU:HD22	2.36	0.40
2:C:422:GLN:O	2:C:423:SER:OG	2.22	0.40
2:C:464:GLU:O	2:C:464:GLU:OE2	2.39	0.40
2:C:99:PHE:N	2:C:99:PHE:CD2	2.89	0.40
1:D:238:GLN:OE1	1:D:238:GLN:HA	2.19	0.40
2:C:64:GLU:CG	2:C:65:GLY:N	2.84	0.40
1:D:1071:ARG:C	1:D:1073:PRO:HD2	2.42	0.40
1:D:148:LEU:HA	1:D:148:LEU:HD23	1.88	0.40
1:D:307:PHE:CA	1:D:310:SER:OG	2.67	0.40
1:D:421:PRO:CG	1:D:422:VAL:H	2.32	0.40
1:D:458:GLU:C	1:D:460:LYS:CE	2.90	0.40
1:D:484:TRP:CE2	1:D:488:LEU:HD11	2.56	0.40
1:D:603:MET:HB3	1:D:605:LEU:HD23	2.00	0.40
1:D:606:THR:CG2	1:D:611:PRO:O	2.69	0.40
1:D:614:TYR:CE2	1:D:620:TRP:CD2	3.08	0.40
1:D:630:ASN:ND2	1:D:632:ALA:N	2.69	0.40
1:D:788:GLY:C	1:D:792:LEU:HD13	2.42	0.40
1:D:950:ASN:O	1:D:951:TYR:C	2.58	0.40
2:E:370:LYS:HE3	2:E:370:LYS:HB3	1.65	0.40
2:F:111:ALA:O	2:F:115:TRP:CB	2.70	0.40
2:F:180:LYS:CA	2:F:180:LYS:HE3	2.47	0.40
2:F:254:PHE:HA	2:F:257:ARG:HD3	2.03	0.40
2:F:341:ASP:OD1	2:F:342:LEU:C	2.60	0.40
2:F:76:ARG:CZ	2:F:434:GLU:CD	2.89	0.40
1:A:422:VAL:HG11	1:A:1111:LEU:HD11	2.03	0.40
1:A:175:TRP:CB	1:A:184:ALA:C	2.86	0.40
1:A:174:GLY:HA2	1:A:220:TRP:CB	2.51	0.40
1:A:659:HIS:HD1	1:A:663:GLN:CG	2.35	0.40
1:A:761:ASN:N	1:A:761:ASN:OD1	2.54	0.40
1:A:851:THR:HG21	1:A:853:ARG:HB2	2.00	0.40
1:A:83:LEU:O	1:A:87:ILE:O	2.40	0.40
1:A:905:ALA:HA	1:A:910:MET:HA	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:905:ALA:HB1	1:A:1169:GLY:HA2	2.03	0.40
2:B:256:LEU:O	2:B:257:ARG:C	2.60	0.40
2:B:265:LYS:O	2:B:266:PHE:CB	2.69	0.40
2:B:456:ARG:HB3	2:B:457:SER:H	1.70	0.40
2:B:87:SER:C	2:B:89:ASP:N	2.73	0.40
2:C:335:VAL:O	2:C:335:VAL:HG13	2.20	0.40
2:C:240:VAL:O	2:C:336:LEU:HD13	2.21	0.40
2:C:385:LEU:HD21	2:C:402:LEU:CD2	2.39	0.40
2:C:399:CYS:C	2:C:401:GLY:N	2.69	0.40
2:C:476:ILE:HD13	2:C:477:LYS:CG	2.50	0.40
1:D:513:LEU:H	1:D:570:PRO:HD3	1.85	0.40
2:C:82:SER:C	2:C:86:LEU:CD1	2.89	0.40
2:B:420:THR:OG1	2:B:420:THR:O	2.32	0.40
1:D:1067:SER:CA	1:D:1071:ARG:CA	2.82	0.40
1:D:1200:PRO:O	1:D:1201:SER:C	2.59	0.40
1:D:197:PHE:CZ	1:D:219:ALA:CB	3.04	0.40
1:D:282:TYR:HE2	1:D:432:VAL:CB	2.34	0.40
1:D:371:LYS:CD	1:D:392:LEU:CD2	2.98	0.40
1:D:624:VAL:HB	1:D:747:CYS:SG	2.62	0.40
1:D:780:GLN:OE1	1:D:781:ALA:N	2.49	0.40
1:D:848:GLY:O	1:D:855:VAL:HG22	2.21	0.40
1:D:851:THR:C	1:D:853:ARG:H	2.23	0.40
1:D:910:MET:C	1:D:910:MET:SD	3.00	0.40
1:D:933:SER:O	1:D:935:THR:N	2.54	0.40
2:E:403:PHE:HA	2:E:413:VAL:HG11	2.04	0.40
2:E:428:LEU:C	2:E:431:LYS:H	2.25	0.40
2:E:474:PHE:O	2:E:475:LEU:C	2.60	0.40
2:E:78:PHE:CE1	2:E:350:LEU:CD1	3.04	0.40
2:F:406:LEU:H	2:F:406:LEU:HD13	1.87	0.40
2:F:381:ILE:CB	2:F:437:ILE:HG22	2.52	0.40
1:A:903:GLY:N	1:A:1002:GLU:HB3	2.32	0.40
1:A:1111:LEU:HB3	1:A:1160:THR:HG23	2.03	0.40
1:A:181:GLU:CD	1:A:181:GLU:C	2.80	0.40
1:A:206:GLY:C	1:A:208:CYS:H	2.24	0.40
1:A:244:LEU:HD23	1:A:244:LEU:HA	1.81	0.40
1:A:203:LEU:HD11	1:A:379:LYS:C	2.34	0.40
1:A:606:THR:O	1:A:607:TRP:CD1	2.75	0.40
1:A:612:LEU:CD2	1:A:614:TYR:N	2.83	0.40
1:A:617:ARG:O	1:A:762:VAL:CG2	2.60	0.40
1:A:659:HIS:CD2	1:A:716:PRO:O	2.75	0.40
2:B:185:LEU:HB3	2:B:214:CYS:SG	2.61	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:306:LEU:O	2:B:307:GLY:C	2.58	0.40
2:B:375:HIS:ND1	2:B:378:LEU:HD13	2.37	0.40
2:B:393:LEU:O	2:B:396:ARG:HB3	2.21	0.40
2:C:368:HIS:O	2:C:368:HIS:CG	2.75	0.40
2:C:368:HIS:O	2:C:368:HIS:ND1	2.55	0.40
2:C:470:LYS:N	2:C:471:LEU:HD12	2.36	0.40
2:E:280:ASP:N	2:E:304:TRP:CH2	2.90	0.40
1:D:584:ALA:HB1	1:D:585:TRP:CZ3	2.57	0.40
1:A:526:ASP:CG	1:A:527:GLN:N	2.74	0.40
1:D:536:GLU:H	1:D:536:GLU:HG2	1.78	0.40

All (10) 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:A:673:ALA:O	2:F:318:ASN:O[2_734]	1.91	0.29
1:A:673:ALA:C	2:F:318:ASN:O[2_734]	1.98	0.22
1:A:1034:ARG:NE	2:F:450:ASN:ND2[1_545]	2.04	0.16
1:A:994:TRP:NE1	1:D:522:GLY:N[1_545]	2.06	0.14
1:D:1232:GLU:OE2	2:F:223:GLN:OE1[2_844]	2.11	0.09
2:C:450:ASN:ND2	1:D:1034:ARG:NH2[1_445]	2.12	0.08
1:A:518:ALA:O	1:D:993:ARG:O[1_445]	2.13	0.07
1:A:727:ASP:OD1	1:D:133:ASP:O[1_445]	2.16	0.04
1:A:133:ASP:OD1	1:D:727:ASP:OD2[1_545]	2.19	0.01
1:A:993:ARG:NH2	1:D:520:ALA:O[1_545]	2.19	0.01

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all 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	1085/1172 (93%)	629 (58%)	254 (23%)	202 (19%)	0 0

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	D	1086/1172 (93%)	645 (59%)	234 (22%)	207 (19%)	0	0
2	B	349/427 (82%)	205 (59%)	94 (27%)	50 (14%)	0	1
2	C	394/427 (92%)	254 (64%)	84 (21%)	56 (14%)	0	1
2	E	349/427 (82%)	219 (63%)	79 (23%)	51 (15%)	0	1
2	F	394/427 (92%)	251 (64%)	93 (24%)	50 (13%)	0	2
All	All	3657/4052 (90%)	2203 (60%)	838 (23%)	616 (17%)	0	1

All (616) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	72	ASN
1	A	76	ILE
1	A	96	GLY
1	A	163	PRO
1	A	169	TRP
1	A	170	ALA
1	A	176	THR
1	A	184	ALA
1	A	185	VAL
1	A	189	ILE
1	A	205	GLU
1	A	211	LEU
1	A	216	SER
1	A	217	PRO
1	A	231	GLU
1	A	232	ARG
1	A	234	SER
1	A	240	SER
1	A	253	ALA
1	A	255	SER
1	A	271	VAL
1	A	273	PHE
1	A	287	SER
1	A	348	LEU
1	A	353	VAL
1	A	354	ASN
1	A	355	SER
1	A	369	PRO
1	A	372	GLU
1	A	378	VAL

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Mol	Chain	Res	Type
1	A	419	PRO
1	A	436	PRO
1	A	437	VAL
1	A	463	LEU
1	A	482	ASP
1	A	499	LYS
1	A	500	ALA
1	A	503	VAL
1	A	527	GLN
1	A	540	GLN
1	A	559	LEU
1	A	573	TYR
1	A	583	PRO
1	A	588	GLY
1	A	589	PRO
1	A	598	VAL
1	A	610	PHE
1	A	611	PRO
1	A	612	LEU
1	A	614	TYR
1	A	626	GLY
1	A	628	ARG
1	A	634	LEU
1	A	635	PRO
1	A	712	VAL
1	A	727	ASP
1	A	728	THR
1	A	731	SER
1	A	738	PRO
1	A	744	ILE
1	A	758	ASN
1	A	768	LYS
1	A	869	ARG
1	A	880	ALA
1	A	900	ALA
1	A	915	ALA
1	A	920	THR
1	A	927	ARG
1	A	939	VAL
1	A	959	GLN
1	A	1020	ILE
1	A	1032	THR

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Mol	Chain	Res	Type
1	A	1034	ARG
1	A	1059	ASN
1	A	1069	ILE
1	A	1070	PRO
1	A	1125	ILE
1	A	1131	ILE
1	A	1170	LEU
1	A	1171	ASN
1	A	1192	GLU
1	A	1193	VAL
1	A	1200	PRO
1	A	1206	MET
1	A	1208	ARG
1	A	1213	PRO
1	A	1214	GLN
1	A	1219	ASP
1	A	1236	GLN
2	B	77	HIS
2	B	79	LEU
2	B	101	PRO
2	B	123	GLU
2	B	205	PRO
2	B	216	HIS
2	B	217	PRO
2	B	246	ARG
2	B	258	HIS
2	B	262	TRP
2	B	280	ASP
2	B	281	GLU
2	B	284	ARG
2	B	285	LYS
2	B	323	HIS
2	B	376	PRO
2	B	391	PRO
2	B	437	ILE
2	C	83	LYS
2	C	122	ARG
2	C	134	LYS
2	C	145	PHE
2	C	178	SER
2	C	204	LEU
2	C	205	PRO

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Mol	Chain	Res	Type
2	C	223	GLN
2	C	257	ARG
2	C	319	VAL
2	C	342	LEU
2	C	352	ASP
2	C	353	SER
2	C	356	LEU
2	C	398	VAL
2	C	415	PRO
2	C	438	LEU
2	C	449	GLU
2	C	452	LEU
2	C	456	ARG
2	C	460	THR
1	D	73	PRO
1	D	81	ARG
1	D	116	PRO
1	D	163	PRO
1	D	167	PRO
1	D	168	ALA
1	D	171	TRP
1	D	173	GLU
1	D	193	ARG
1	D	205	GLU
1	D	211	LEU
1	D	212	ALA
1	D	213	VAL
1	D	219	ALA
1	D	221	TYR
1	D	234	SER
1	D	248	GLU
1	D	249	VAL
1	D	255	SER
1	D	303	GLY
1	D	350	ILE
1	D	364	VAL
1	D	368	PRO
1	D	372	GLU
1	D	378	VAL
1	D	419	PRO
1	D	433	SER
1	D	482	ASP

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Mol	Chain	Res	Type
1	D	492	LEU
1	D	504	LYS
1	D	511	SER
1	D	517	GLY
1	D	521	PRO
1	D	523	ASP
1	D	530	LEU
1	D	540	GLN
1	D	559	LEU
1	D	577	CYS
1	D	589	PRO
1	D	597	ARG
1	D	598	VAL
1	D	599	THR
1	D	610	PHE
1	D	611	PRO
1	D	612	LEU
1	D	614	TYR
1	D	628	ARG
1	D	643	ALA
1	D	646	VAL
1	D	647	CYS
1	D	710	ALA
1	D	712	VAL
1	D	713	PRO
1	D	727	ASP
1	D	728	THR
1	D	735	GLY
1	D	736	ASN
1	D	742	VAL
1	D	744	ILE
1	D	753	PRO
1	D	762	VAL
1	D	776	ASP
1	D	781	ALA
1	D	783	PRO
1	D	787	SER
1	D	829	PRO
1	D	869	ARG
1	D	890	ASP
1	D	891	VAL
1	D	913	CYS

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Mol	Chain	Res	Type
1	D	925	LYS
1	D	939	VAL
1	D	956	GLY
1	D	959	GLN
1	D	974	THR
1	D	992	LEU
1	D	1027	LYS
1	D	1069	ILE
1	D	1073	PRO
1	D	1174	PRO
1	D	1176	SER
1	D	1180	PHE
1	D	1203	PRO
1	D	1232	GLU
1	D	1236	GLN
2	E	70	LEU
2	E	87	SER
2	E	239	LEU
2	E	248	SER
2	E	281	GLU
2	E	294	PRO
2	E	309	HIS
2	E	314	MET
2	E	316	PRO
2	E	319	VAL
2	E	323	HIS
2	E	330	ASN
2	E	331	VAL
2	E	375	HIS
2	E	376	PRO
2	E	384	ALA
2	E	385	LEU
2	E	420	THR
2	E	432	TYR
2	E	449	GLU
2	F	83	LYS
2	F	87	SER
2	F	134	LYS
2	F	139	LEU
2	F	145	PHE
2	F	178	SER
2	F	223	GLN

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Mol	Chain	Res	Type
2	F	319	VAL
2	F	341	ASP
2	F	352	ASP
2	F	356	LEU
2	F	360	SER
2	F	365	LYS
2	F	389	ARG
2	F	415	PRO
2	F	453	ILE
2	F	456	ARG
2	F	460	THR
1	A	73	PRO
1	A	84	HIS
1	A	132	GLY
1	A	171	TRP
1	A	238	GLN
1	A	262	GLN
1	A	283	LEU
1	A	292	LEU
1	A	371	LYS
1	A	383	LYS
1	A	470	ALA
1	A	507	PRO
1	A	508	ALA
1	A	517	GLY
1	A	561	LYS
1	A	576	LEU
1	A	596	MET
1	A	607	TRP
1	A	613	HIS
1	A	617	ARG
1	A	639	THR
1	A	659	HIS
1	A	719	LEU
1	A	735	GLY
1	A	762	VAL
1	A	813	VAL
1	A	864	ASN
1	A	923	GLY
1	A	940	GLY
1	A	1018	GLY
1	A	1021	SER

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Mol	Chain	Res	Type
1	A	1052	GLY
1	A	1094	THR
1	A	1122	GLU
1	A	1145	ASP
1	A	1228	LYS
1	A	1232	GLU
2	B	88	ARG
2	B	117	SER
2	B	179	GLY
2	B	265	LYS
2	B	282	GLU
2	B	309	HIS
2	B	312	LEU
2	B	343	ASP
2	B	353	SER
2	B	354	PHE
2	B	377	CYS
2	B	410	GLY
2	B	436	SER
2	B	441	VAL
2	B	469	SER
2	C	88	ARG
2	C	248	SER
2	C	300	ILE
2	C	321	LYS
2	C	328	ARG
2	C	340	GLY
2	C	351	TYR
2	C	445	GLU
2	C	451	GLY
2	C	453	ILE
2	C	458	ARG
1	D	78	MET
1	D	130	LEU
1	D	174	GLY
1	D	218	SER
1	D	220	TRP
1	D	223	TRP
1	D	292	LEU
1	D	371	LYS
1	D	392	LEU
1	D	421	PRO

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Mol	Chain	Res	Type
1	D	469	ASP
1	D	510	ALA
1	D	519	GLY
1	D	525	MET
1	D	529	ASP
1	D	588	GLY
1	D	609	GLY
1	D	624	VAL
1	D	656	TYR
1	D	758	ASN
1	D	802	ARG
1	D	810	SER
1	D	855	VAL
1	D	868	ASP
1	D	877	MET
1	D	880	ALA
1	D	920	THR
1	D	940	GLY
1	D	1000	GLU
1	D	1021	SER
1	D	1025	LEU
1	D	1034	ARG
1	D	1042	GLU
1	D	1062	GLU
1	D	1066	THR
1	D	1068	ASP
1	D	1171	ASN
1	D	1172	ASP
1	D	1193	VAL
1	D	1220	ILE
1	D	1238	GLY
2	E	82	SER
2	E	85	GLN
2	E	94	GLY
2	E	116	THR
2	E	117	SER
2	E	231	ILE
2	E	264	ARG
2	E	298	GLU
2	E	311	LEU
2	E	401	GLY
2	E	405	GLU

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Mol	Chain	Res	Type
2	E	408	GLU
2	E	469	SER
2	F	122	ARG
2	F	125	VAL
2	F	182	ARG
2	F	200	VAL
2	F	220	ASP
2	F	257	ARG
2	F	300	ILE
2	F	342	LEU
2	F	353	SER
2	F	438	LEU
2	F	455	LEU
1	A	123	VAL
1	A	233	TYR
1	A	239	LEU
1	A	393	MET
1	A	570	PRO
1	A	587	PRO
1	A	716	PRO
1	A	726	LYS
1	A	763	GLY
1	A	777	GLY
1	A	787	SER
1	A	805	HIS
1	A	922	GLN
1	A	990	LYS
1	A	1000	GLU
1	A	1037	GLN
1	A	1038	TRP
1	A	1042	GLU
1	A	1071	ARG
1	A	1095	SER
1	A	1195	MET
1	A	1212	ILE
2	B	75	ARG
2	B	199	LEU
2	B	247	THR
2	B	267	ALA
2	B	283	GLY
2	B	349	TYR
2	C	137	PRO

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Mol	Chain	Res	Type
2	C	220	ASP
2	C	368	HIS
2	C	423	SER
2	C	425	LEU
2	C	448	LEU
1	D	84	HIS
1	D	183	GLU
1	D	254	SER
1	D	278	ILE
1	D	306	SER
1	D	470	ALA
1	D	527	GLN
1	D	560	PRO
1	D	578	PRO
1	D	579	ARG
1	D	627	ARG
1	D	719	LEU
1	D	731	SER
1	D	811	GLN
1	D	876	ALA
1	D	1020	ILE
1	D	1026	ARG
1	D	1037	GLN
1	D	1040	LYS
1	D	1094	THR
1	D	1128	ARG
1	D	1162	CYS
1	D	1181	SER
1	D	1191	LYS
1	D	1196	ASP
2	E	203	ARG
2	E	328	ARG
2	E	382	LYS
2	F	137	PRO
2	F	328	ARG
2	F	403	PHE
2	F	436	SER
2	F	452	LEU
2	F	484	ASN
1	A	162	LEU
1	A	397	ALA
1	A	412	PRO

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Mol	Chain	Res	Type
1	A	501	LYS
1	A	577	CYS
1	A	633	LYS
1	A	722	ARG
1	A	742	VAL
1	A	802	ARG
1	A	817	PRO
1	A	868	ASP
1	A	973	LEU
1	A	1043	VAL
1	A	1068	ASP
1	A	1129	PHE
1	A	1167	LYS
1	A	1199	THR
2	B	214	CYS
2	C	140	PRO
2	C	182	ARG
2	C	189	ALA
2	C	200	VAL
2	C	226	ASN
2	C	270	PRO
2	C	299	LEU
2	C	392	THR
2	C	419	GLU
2	C	478	TYR
2	C	484	ASN
1	D	97	GLU
1	D	119	PRO
1	D	165	LYS
1	D	179	GLY
1	D	210	THR
1	D	259	ARG
1	D	271	VAL
1	D	287	SER
1	D	500	ALA
1	D	520	ALA
1	D	625	PRO
1	D	632	ALA
1	D	655	LEU
1	D	672	GLU
1	D	715	GLN
1	D	723	GLY

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Mol	Chain	Res	Type
1	D	730	PRO
1	D	922	GLN
1	D	995	TYR
1	D	1035	LYS
1	D	1108	TYR
1	D	1134	HIS
1	D	1192	GLU
2	E	77	HIS
2	E	88	ARG
2	E	121	PHE
2	E	215	PHE
2	E	240	VAL
2	E	285	LYS
2	E	292	ASN
2	F	77	HIS
2	F	140	PRO
2	F	217	PRO
2	F	248	SER
2	F	280	ASP
2	F	423	SER
2	F	451	GLY
2	F	478	TYR
1	A	71	HIS
1	A	248	GLU
1	A	380	GLY
1	A	421	PRO
1	A	462	SER
1	A	464	MET
1	A	506	GLU
1	A	568	GLY
1	A	625	PRO
1	A	646	VAL
1	A	660	CYS
1	A	717	LEU
1	A	830	ASP
1	A	908	ALA
1	A	910	MET
1	A	1062	GLU
1	A	1092	PHE
1	A	1182	ALA
2	B	201	ASN
2	B	316	PRO

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Mol	Chain	Res	Type
2	B	413	VAL
2	B	484	ASN
2	C	110	LEU
2	C	124	GLN
2	C	280	ASP
2	C	389	ARG
1	D	169	TRP
1	D	177	ARG
1	D	184	ALA
1	D	191	GLU
1	D	279	ARG
1	D	405	GLU
1	D	477	GLU
1	D	562	ARG
1	D	633	LYS
1	D	780	GLN
1	D	882	PRO
1	D	941	ILE
1	D	1118	TRP
1	D	1122	GLU
2	E	210	GLN
2	E	272	ASN
2	E	280	ASP
2	E	372	LEU
2	E	423	SER
2	F	121	PHE
2	F	130	ALA
2	F	138	LEU
2	F	351	TYR
1	A	857	PRO
1	A	1093	MET
1	A	1207	GLU
2	B	270	PRO
2	B	297	LYS
2	C	125	VAL
1	D	128	PRO
1	D	569	HIS
1	D	572	TRP
1	D	853	ARG
1	D	1103	SER
1	A	128	PRO
1	A	215	ILE

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Mol	Chain	Res	Type
1	A	645	VAL
1	D	166	PRO
1	D	411	LEU
1	D	881	PRO
1	D	1205	GLY
2	F	204	LEU
1	A	87	ILE
1	A	871	GLY
1	A	878	VAL
1	A	1010	LEU
1	A	1127	GLY
2	B	327	GLY
1	D	267	VAL
1	D	406	VAL
1	D	436	PRO
1	D	563	PRO
1	D	1155	ILE
1	D	1211	GLY
2	F	128	VAL
1	A	366	VAL
1	A	432	VAL
1	A	515	ILE
1	A	715	GLN
2	B	307	GLY
2	B	380	PRO
2	C	120	VAL
2	C	414	TRP
1	D	1137	VAL
2	F	414	TRP
1	A	560	PRO
1	A	621	GLY
1	A	855	VAL
2	B	213	VAL
1	D	435	LEU
1	D	817	PRO
2	E	135	PRO
2	E	179	GLY
2	E	200	VAL
1	D	648	PRO
1	A	753	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	918/988 (93%)	611 (67%)	307 (33%)	0	0
1	D	919/988 (93%)	584 (64%)	335 (36%)	0	0
2	B	317/380 (83%)	259 (82%)	58 (18%)	2	10
2	C	350/380 (92%)	260 (74%)	90 (26%)	0	2
2	E	317/380 (83%)	270 (85%)	47 (15%)	4	17
2	F	350/380 (92%)	245 (70%)	105 (30%)	0	1
All	All	3171/3496 (91%)	2229 (70%)	942 (30%)	0	1

All (942) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	70	ARG
1	A	78	MET
1	A	83	LEU
1	A	90	GLN
1	A	93	GLU
1	A	94	MET
1	A	97	GLU
1	A	101	ARG
1	A	102	ARG
1	A	107	LEU
1	A	109	LYS
1	A	120	LEU
1	A	125	LEU
1	A	128	PRO
1	A	131	TYR
1	A	135	LEU
1	A	138	HIS
1	A	141	LEU
1	A	146	GLN
1	A	156	LEU
1	A	157	LEU
1	A	161	GLN

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Mol	Chain	Res	Type
1	A	169	TRP
1	A	177	ARG
1	A	183	GLU
1	A	187	VAL
1	A	190	PRO
1	A	193	ARG
1	A	195	LEU
1	A	197	PHE
1	A	202	CYS
1	A	208	CYS
1	A	216	SER
1	A	217	PRO
1	A	220	TRP
1	A	223	TRP
1	A	224	CYS
1	A	226	GLN
1	A	227	ARG
1	A	228	LEU
1	A	235	TRP
1	A	236	THR
1	A	246	PRO
1	A	247	LEU
1	A	248	GLU
1	A	259	ARG
1	A	260	ASP
1	A	261	TRP
1	A	262	GLN
1	A	263	GLU
1	A	265	LEU
1	A	267	VAL
1	A	270	ASN
1	A	272	SER
1	A	275	ARG
1	A	279	ARG
1	A	288	ARG
1	A	293	ASP
1	A	294	THR
1	A	298	HIS
1	A	307	PHE
1	A	308	GLN
1	A	311	LEU
1	A	312	TRP

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Mol	Chain	Res	Type
1	A	346	ASP
1	A	350	ILE
1	A	351	SER
1	A	352	SER
1	A	354	ASN
1	A	356	ASN
1	A	358	LEU
1	A	361	VAL
1	A	364	LEU
1	A	365	TYR
1	A	369	PRO
1	A	372	GLU
1	A	374	ARG
1	A	376	LEU
1	A	377	PHE
1	A	379	LYS
1	A	384	ASP
1	A	386	ARG
1	A	388	ASN
1	A	389	PHE
1	A	393	MET
1	A	394	GLN
1	A	398	GLN
1	A	403	THR
1	A	406	VAL
1	A	415	LEU
1	A	416	GLU
1	A	417	ARG
1	A	424	LEU
1	A	429	GLU
1	A	430	MET
1	A	436	PRO
1	A	441	TRP
1	A	442	GLU
1	A	445	LEU
1	A	451	THR
1	A	453	GLU
1	A	457	ARG
1	A	460	LYS
1	A	463	LEU
1	A	465	ASP
1	A	466	LEU

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Mol	Chain	Res	Type
1	A	473	LEU
1	A	474	LEU
1	A	477	GLU
1	A	481	GLU
1	A	482	ASP
1	A	484	TRP
1	A	485	LEU
1	A	487	ASP
1	A	490	TRP
1	A	491	ASP
1	A	496	LYS
1	A	498	LYS
1	A	502	LYS
1	A	512	LYS
1	A	524	PRO
1	A	526	ASP
1	A	527	GLN
1	A	528	GLU
1	A	535	GLU
1	A	536	GLU
1	A	549	LEU
1	A	557	GLU
1	A	558	LEU
1	A	559	LEU
1	A	560	PRO
1	A	561	LYS
1	A	564	GLN
1	A	567	PRO
1	A	569	HIS
1	A	572	TRP
1	A	576	LEU
1	A	577	CYS
1	A	579	ARG
1	A	583	PRO
1	A	585	TRP
1	A	586	THR
1	A	587	PRO
1	A	591	LEU
1	A	592	LEU
1	A	594	LEU
1	A	595	GLN
1	A	596	MET

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Mol	Chain	Res	Type
1	A	597	ARG
1	A	601	LYS
1	A	607	TRP
1	A	608	ASP
1	A	610	PHE
1	A	611	PRO
1	A	612	LEU
1	A	618	HIS
1	A	620	TRP
1	A	624	VAL
1	A	627	ARG
1	A	628	ARG
1	A	630	ASN
1	A	631	LEU
1	A	633	LYS
1	A	636	THR
1	A	639	THR
1	A	640	LEU
1	A	655	LEU
1	A	659	HIS
1	A	660	CYS
1	A	661	LEU
1	A	662	GLU
1	A	666	GLN
1	A	668	LEU
1	A	672	GLU
1	A	712	VAL
1	A	719	LEU
1	A	727	ASP
1	A	733	HIS
1	A	734	HIS
1	A	736	ASN
1	A	738	PRO
1	A	741	ASP
1	A	744	ILE
1	A	748	TRP
1	A	752	LEU
1	A	754	HIS
1	A	755	LYS
1	A	756	ASP
1	A	758	ASN
1	A	760	CYS

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Mol	Chain	Res	Type
1	A	771	LEU
1	A	776	ASP
1	A	778	THR
1	A	779	LEU
1	A	792	LEU
1	A	795	ASN
1	A	798	ILE
1	A	803	ASN
1	A	805	HIS
1	A	810	SER
1	A	812	MET
1	A	813	VAL
1	A	817	PRO
1	A	823	ARG
1	A	826	ILE
1	A	827	ARG
1	A	831	TYR
1	A	833	GLU
1	A	840	ILE
1	A	850	ILE
1	A	851	THR
1	A	853	ARG
1	A	858	THR
1	A	864	ASN
1	A	866	ARG
1	A	877	MET
1	A	879	GLN
1	A	884	TYR
1	A	885	THR
1	A	886	LEU
1	A	891	VAL
1	A	892	ASP
1	A	894	GLN
1	A	898	ILE
1	A	904	ASP
1	A	906	HIS
1	A	910	MET
1	A	911	HIS
1	A	913	CYS
1	A	916	PHE
1	A	919	MET
1	A	921	LEU

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Mol	Chain	Res	Type
1	A	922	GLN
1	A	927	ARG
1	A	930	ASP
1	A	931	LEU
1	A	933	SER
1	A	941	ILE
1	A	948	ILE
1	A	949	PHE
1	A	954	ILE
1	A	961	PHE
1	A	966	LEU
1	A	989	THR
1	A	992	LEU
1	A	997	LEU
1	A	1003	TRP
1	A	1006	ARG
1	A	1008	LEU
1	A	1011	PRO
1	A	1014	ARG
1	A	1015	THR
1	A	1016	GLU
1	A	1028	VAL
1	A	1029	GLN
1	A	1031	GLU
1	A	1041	TRP
1	A	1044	VAL
1	A	1047	ARG
1	A	1056	GLU
1	A	1066	THR
1	A	1069	ILE
1	A	1102	GLN
1	A	1109	LEU
1	A	1112	MET
1	A	1113	LEU
1	A	1116	MET
1	A	1125	ILE
1	A	1126	ASP
1	A	1130	CYS
1	A	1133	ILE
1	A	1138	ARG
1	A	1140	LEU
1	A	1145	ASP

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Mol	Chain	Res	Type
1	A	1148	ARG
1	A	1153	LEU
1	A	1154	GLN
1	A	1156	THR
1	A	1160	THR
1	A	1166	TYR
1	A	1170	LEU
1	A	1171	ASN
1	A	1172	ASP
1	A	1177	VAL
1	A	1180	PHE
1	A	1191	LYS
1	A	1193	VAL
1	A	1197	CYS
1	A	1198	LYS
1	A	1202	ASN
1	A	1206	MET
1	A	1209	ARG
1	A	1210	TYR
1	A	1213	PRO
1	A	1214	GLN
1	A	1218	LEU
1	A	1219	ASP
1	A	1220	ILE
1	A	1223	ILE
1	A	1224	ILE
1	A	1225	GLU
1	A	1226	LEU
1	A	1227	THR
1	A	1231	LEU
1	A	1233	LYS
1	A	1236	GLN
1	A	1237	PRO
2	B	74	GLN
2	B	77	HIS
2	B	78	PHE
2	B	83	LYS
2	B	84	GLN
2	B	86	LEU
2	B	89	ASP
2	B	96	HIS
2	B	101	PRO

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Mol	Chain	Res	Type
2	B	105	GLU
2	B	109	ASN
2	B	122	ARG
2	B	123	GLU
2	B	184	ASN
2	B	203	ARG
2	B	205	PRO
2	B	208	LEU
2	B	210	GLN
2	B	230	SER
2	B	231	ILE
2	B	234	LYS
2	B	236	GLU
2	B	242	PHE
2	B	246	ARG
2	B	251	TRP
2	B	254	PHE
2	B	260	LEU
2	B	262	TRP
2	B	264	ARG
2	B	272	ASN
2	B	281	GLU
2	B	284	ARG
2	B	298	GLU
2	B	304	TRP
2	B	309	HIS
2	B	311	LEU
2	B	312	LEU
2	B	313	HIS
2	B	316	PRO
2	B	318	ASN
2	B	325	ARG
2	B	331	VAL
2	B	344	ARG
2	B	346	MET
2	B	347	LEU
2	B	349	TYR
2	B	351	TYR
2	B	352	ASP
2	B	377	CYS
2	B	380	PRO
2	B	414	TRP

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Mol	Chain	Res	Type
2	B	417	TYR
2	B	429	TYR
2	B	431	LYS
2	B	441	VAL
2	B	453	ILE
2	B	454	HIS
2	B	472	LYS
2	C	88	ARG
2	C	89	ASP
2	C	91	LEU
2	C	92	LEU
2	C	99	PHE
2	C	116	THR
2	C	117	SER
2	C	118	VAL
2	C	120	VAL
2	C	124	GLN
2	C	125	VAL
2	C	128	VAL
2	C	131	LEU
2	C	132	HIS
2	C	133	HIS
2	C	180	LYS
2	C	186	LEU
2	C	193	TYR
2	C	197	LEU
2	C	198	ASP
2	C	199	LEU
2	C	205	PRO
2	C	206	TYR
2	C	208	LEU
2	C	210	GLN
2	C	214	CYS
2	C	215	PHE
2	C	224	ILE
2	C	226	ASN
2	C	230	SER
2	C	236	GLU
2	C	244	PRO
2	C	245	PRO
2	C	246	ARG
2	C	247	THR

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Mol	Chain	Res	Type
2	C	251	TRP
2	C	254	PHE
2	C	256	LEU
2	C	280	ASP
2	C	285	LYS
2	C	298	GLU
2	C	299	LEU
2	C	308	ASP
2	C	309	HIS
2	C	310	GLU
2	C	312	LEU
2	C	320	SER
2	C	321	LYS
2	C	322	LEU
2	C	325	ARG
2	C	328	ARG
2	C	330	ASN
2	C	337	SER
2	C	341	ASP
2	C	342	LEU
2	C	350	LEU
2	C	351	TYR
2	C	357	THR
2	C	363	ARG
2	C	364	LYS
2	C	367	LEU
2	C	369	ARG
2	C	385	LEU
2	C	387	VAL
2	C	389	ARG
2	C	394	GLU
2	C	395	LEU
2	C	396	ARG
2	C	399	CYS
2	C	400	GLN
2	C	402	LEU
2	C	403	PHE
2	C	406	LEU
2	C	407	LEU
2	C	417	TYR
2	C	431	LYS
2	C	435	MET

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Mol	Chain	Res	Type
2	C	439	PHE
2	C	448	LEU
2	C	456	ARG
2	C	457	SER
2	C	458	ARG
2	C	464	GLU
2	C	468	ILE
2	C	471	LEU
2	C	474	PHE
2	C	475	LEU
2	C	476	ILE
2	C	483	LYS
2	C	485	VAL
1	D	70	ARG
1	D	77	GLN
1	D	78	MET
1	D	79	LEU
1	D	81	ARG
1	D	83	LEU
1	D	85	GLU
1	D	86	GLN
1	D	87	ILE
1	D	97	GLU
1	D	101	ARG
1	D	104	VAL
1	D	107	LEU
1	D	113	TRP
1	D	120	LEU
1	D	122	ASP
1	D	125	LEU
1	D	127	LEU
1	D	130	LEU
1	D	133	ASP
1	D	135	LEU
1	D	140	ARG
1	D	144	GLN
1	D	145	LYS
1	D	146	GLN
1	D	157	LEU
1	D	161	GLN
1	D	162	LEU
1	D	169	TRP

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Mol	Chain	Res	Type
1	D	175	TRP
1	D	176	THR
1	D	177	ARG
1	D	181	GLU
1	D	185	VAL
1	D	189	ILE
1	D	193	ARG
1	D	196	VAL
1	D	201	VAL
1	D	203	LEU
1	D	205	GLU
1	D	208	CYS
1	D	216	SER
1	D	217	PRO
1	D	218	SER
1	D	220	TRP
1	D	223	TRP
1	D	224	CYS
1	D	225	SER
1	D	227	ARG
1	D	228	LEU
1	D	232	ARG
1	D	236	THR
1	D	238	GLN
1	D	239	LEU
1	D	243	ASP
1	D	247	LEU
1	D	256	PRO
1	D	258	GLN
1	D	259	ARG
1	D	260	ASP
1	D	265	LEU
1	D	267	VAL
1	D	269	HIS
1	D	284	ILE
1	D	285	GLN
1	D	288	ARG
1	D	289	MET
1	D	293	ASP
1	D	298	HIS
1	D	301	ILE
1	D	305	SER

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Mol	Chain	Res	Type
1	D	307	PHE
1	D	308	GLN
1	D	313	ILE
1	D	316	LYS
1	D	317	GLN
1	D	347	TRP
1	D	348	LEU
1	D	352	SER
1	D	362	LEU
1	D	371	LYS
1	D	374	ARG
1	D	379	LYS
1	D	384	ASP
1	D	386	ARG
1	D	389	PHE
1	D	390	GLN
1	D	391	ASP
1	D	394	GLN
1	D	395	TYR
1	D	403	THR
1	D	405	GLU
1	D	407	PHE
1	D	411	LEU
1	D	415	LEU
1	D	416	GLU
1	D	417	ARG
1	D	419	PRO
1	D	420	HIS
1	D	424	LEU
1	D	428	LEU
1	D	429	GLU
1	D	430	MET
1	D	435	LEU
1	D	436	PRO
1	D	438	ASN
1	D	439	GLN
1	D	441	TRP
1	D	442	GLU
1	D	445	LEU
1	D	447	GLU
1	D	457	ARG
1	D	460	LYS

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Mol	Chain	Res	Type
1	D	461	LYS
1	D	466	LEU
1	D	469	ASP
1	D	471	CYS
1	D	474	LEU
1	D	479	TYR
1	D	484	TRP
1	D	491	ASP
1	D	493	GLN
1	D	494	GLU
1	D	496	LYS
1	D	499	LYS
1	D	501	LYS
1	D	503	VAL
1	D	504	LYS
1	D	524	PRO
1	D	527	GLN
1	D	530	LEU
1	D	542	ASP
1	D	543	VAL
1	D	550	GLN
1	D	551	LYS
1	D	557	GLU
1	D	558	LEU
1	D	560	PRO
1	D	564	GLN
1	D	565	HIS
1	D	576	LEU
1	D	581	ASP
1	D	585	TRP
1	D	592	LEU
1	D	594	LEU
1	D	595	GLN
1	D	597	ARG
1	D	601	LYS
1	D	602	LEU
1	D	606	THR
1	D	608	ASP
1	D	612	LEU
1	D	615	SER
1	D	617	ARG
1	D	618	HIS

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Mol	Chain	Res	Type
1	D	620	TRP
1	D	627	ARG
1	D	630	ASN
1	D	631	LEU
1	D	638	THR
1	D	645	VAL
1	D	647	CYS
1	D	649	TYR
1	D	653	GLU
1	D	654	SER
1	D	660	CYS
1	D	661	LEU
1	D	666	GLN
1	D	667	GLN
1	D	668	LEU
1	D	713	PRO
1	D	716	PRO
1	D	717	LEU
1	D	719	LEU
1	D	722	ARG
1	D	733	HIS
1	D	736	ASN
1	D	740	ASN
1	D	742	VAL
1	D	744	ILE
1	D	745	PRO
1	D	748	TRP
1	D	753	PRO
1	D	754	HIS
1	D	755	LYS
1	D	756	ASP
1	D	758	ASN
1	D	760	CYS
1	D	761	ASN
1	D	768	LYS
1	D	773	LYS
1	D	776	ASP
1	D	778	THR
1	D	779	LEU
1	D	780	GLN
1	D	795	ASN
1	D	797	MET

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Mol	Chain	Res	Type
1	D	801	TRP
1	D	802	ARG
1	D	805	HIS
1	D	807	ARG
1	D	809	SER
1	D	812	MET
1	D	813	VAL
1	D	819	SER
1	D	821	LEU
1	D	823	ARG
1	D	826	ILE
1	D	827	ARG
1	D	829	PRO
1	D	831	TYR
1	D	841	LEU
1	D	843	GLN
1	D	844	VAL
1	D	846	THR
1	D	850	ILE
1	D	858	THR
1	D	863	SER
1	D	864	ASN
1	D	872	SER
1	D	874	LEU
1	D	878	VAL
1	D	879	GLN
1	D	885	THR
1	D	886	LEU
1	D	892	ASP
1	D	895	GLU
1	D	896	LEU
1	D	898	ILE
1	D	904	ASP
1	D	906	HIS
1	D	907	PHE
1	D	910	MET
1	D	916	PHE
1	D	919	MET
1	D	921	LEU
1	D	922	GLN
1	D	929	THR
1	D	930	ASP

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Mol	Chain	Res	Type
1	D	932	HIS
1	D	933	SER
1	D	934	LYS
1	D	948	ILE
1	D	955	TYR
1	D	961	PHE
1	D	963	GLU
1	D	967	MET
1	D	983	GLN
1	D	989	THR
1	D	992	LEU
1	D	993	ARG
1	D	994	TRP
1	D	995	TYR
1	D	996	ARG
1	D	997	LEU
1	D	1006	ARG
1	D	1008	LEU
1	D	1014	ARG
1	D	1020	ILE
1	D	1022	LEU
1	D	1023	GLN
1	D	1025	LEU
1	D	1026	ARG
1	D	1030	ARG
1	D	1034	ARG
1	D	1036	SER
1	D	1037	GLN
1	D	1038	TRP
1	D	1041	TRP
1	D	1043	VAL
1	D	1047	ARG
1	D	1055	SER
1	D	1056	GLU
1	D	1057	MET
1	D	1058	PHE
1	D	1060	LYS
1	D	1061	LEU
1	D	1066	THR
1	D	1067	SER
1	D	1068	ASP
1	D	1070	PRO

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Mol	Chain	Res	Type
1	D	1071	ARG
1	D	1072	THR
1	D	1073	PRO
1	D	1099	TRP
1	D	1104	SER
1	D	1109	LEU
1	D	1113	LEU
1	D	1121	GLU
1	D	1122	GLU
1	D	1126	ASP
1	D	1130	CYS
1	D	1131	ILE
1	D	1132	SER
1	D	1133	ILE
1	D	1134	HIS
1	D	1135	ASP
1	D	1136	GLU
1	D	1138	ARG
1	D	1140	LEU
1	D	1146	ARG
1	D	1148	ARG
1	D	1153	LEU
1	D	1158	LEU
1	D	1163	MET
1	D	1164	PHE
1	D	1166	TYR
1	D	1171	ASN
1	D	1172	ASP
1	D	1175	GLN
1	D	1180	PHE
1	D	1181	SER
1	D	1184	ASP
1	D	1187	ARG
1	D	1191	LYS
1	D	1192	GLU
1	D	1195	MET
1	D	1197	CYS
1	D	1200	PRO
1	D	1203	PRO
1	D	1204	THR
1	D	1206	MET
1	D	1207	GLU

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Mol	Chain	Res	Type
1	D	1208	ARG
1	D	1210	TYR
1	D	1212	ILE
1	D	1218	LEU
1	D	1219	ASP
1	D	1221	TYR
1	D	1223	ILE
1	D	1224	ILE
1	D	1225	GLU
1	D	1226	LEU
1	D	1227	THR
1	D	1230	SER
2	E	67	GLU
2	E	86	LEU
2	E	88	ARG
2	E	107	ARG
2	E	108	LYS
2	E	116	THR
2	E	120	VAL
2	E	122	ARG
2	E	123	GLU
2	E	131	LEU
2	E	191	GLU
2	E	202	LYS
2	E	211	ILE
2	E	213	VAL
2	E	250	GLN
2	E	255	TRP
2	E	276	SER
2	E	277	ASP
2	E	278	CYS
2	E	293	PHE
2	E	294	PRO
2	E	295	TRP
2	E	315	TYR
2	E	328	ARG
2	E	329	LYS
2	E	336	LEU
2	E	352	ASP
2	E	353	SER
2	E	378	LEU
2	E	385	LEU

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Mol	Chain	Res	Type
2	E	389	ARG
2	E	394	GLU
2	E	397	GLN
2	E	417	TYR
2	E	422	GLN
2	E	425	LEU
2	E	426	GLU
2	E	428	LEU
2	E	432	TYR
2	E	437	ILE
2	E	445	GLU
2	E	446	THR
2	E	447	THR
2	E	449	GLU
2	E	455	LEU
2	E	467	HIS
2	E	483	LYS
2	F	86	LEU
2	F	88	ARG
2	F	89	ASP
2	F	91	LEU
2	F	92	LEU
2	F	95	CYS
2	F	99	PHE
2	F	116	THR
2	F	117	SER
2	F	118	VAL
2	F	120	VAL
2	F	125	VAL
2	F	128	VAL
2	F	131	LEU
2	F	132	HIS
2	F	133	HIS
2	F	142	ASP
2	F	145	PHE
2	F	180	LYS
2	F	186	LEU
2	F	193	TYR
2	F	195	ASN
2	F	197	LEU
2	F	198	ASP
2	F	199	LEU

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Mol	Chain	Res	Type
2	F	204	LEU
2	F	208	LEU
2	F	210	GLN
2	F	214	CYS
2	F	215	PHE
2	F	223	GLN
2	F	226	ASN
2	F	230	SER
2	F	231	ILE
2	F	236	GLU
2	F	238	SER
2	F	244	PRO
2	F	245	PRO
2	F	246	ARG
2	F	247	THR
2	F	251	TRP
2	F	253	ASP
2	F	254	PHE
2	F	256	LEU
2	F	279	GLN
2	F	280	ASP
2	F	281	GLU
2	F	285	LYS
2	F	298	GLU
2	F	299	LEU
2	F	308	ASP
2	F	309	HIS
2	F	310	GLU
2	F	320	SER
2	F	321	LYS
2	F	322	LEU
2	F	325	ARG
2	F	328	ARG
2	F	330	ASN
2	F	331	VAL
2	F	337	SER
2	F	339	ASN
2	F	341	ASP
2	F	342	LEU
2	F	347	LEU
2	F	350	LEU
2	F	351	TYR

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Mol	Chain	Res	Type
2	F	357	THR
2	F	364	LYS
2	F	367	LEU
2	F	369	ARG
2	F	387	VAL
2	F	389	ARG
2	F	394	GLU
2	F	395	LEU
2	F	396	ARG
2	F	400	GLN
2	F	402	LEU
2	F	403	PHE
2	F	406	LEU
2	F	407	LEU
2	F	412	SER
2	F	417	TYR
2	F	421	MET
2	F	427	GLN
2	F	431	LYS
2	F	435	MET
2	F	439	PHE
2	F	446	THR
2	F	450	ASN
2	F	452	LEU
2	F	453	ILE
2	F	455	LEU
2	F	456	ARG
2	F	458	ARG
2	F	459	ASP
2	F	461	THR
2	F	462	MET
2	F	464	GLU
2	F	471	LEU
2	F	474	PHE
2	F	475	LEU
2	F	476	ILE
2	F	483	LYS
2	F	485	VAL

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (119) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	110	HIS
1	A	155	ASN
1	A	238	GLN
1	A	264	GLN
1	A	308	GLN
1	A	388	ASN
1	A	390	GLN
1	A	438	ASN
1	A	468	ASN
1	A	564	GLN
1	A	565	HIS
1	A	630	ASN
1	A	663	GLN
1	A	666	GLN
1	A	729	GLN
1	A	736	ASN
1	A	758	ASN
1	A	795	ASN
1	A	879	GLN
1	A	922	GLN
1	A	970	ASN
1	A	976	GLN
1	A	1023	GLN
1	A	1029	GLN
1	A	1037	GLN
1	A	1059	ASN
1	A	1202	ASN
1	A	1214	GLN
1	A	1222	GLN
1	A	1236	GLN
2	B	74	GLN
2	B	84	GLN
2	B	96	HIS
2	B	109	ASN
2	B	124	GLN
2	B	195	ASN
2	B	201	ASN
2	B	250	GLN
2	B	261	GLN
2	B	287	ASN
2	B	292	ASN
2	B	305	ASN
2	B	313	HIS

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Mol	Chain	Res	Type
2	B	318	ASN
2	B	323	HIS
2	B	422	GLN
2	B	427	GLN
2	C	74	GLN
2	C	109	ASN
2	C	124	GLN
2	C	187	HIS
2	C	279	GLN
2	C	292	ASN
2	C	454	HIS
1	D	71	HIS
1	D	86	GLN
1	D	106	HIS
1	D	110	HIS
1	D	115	GLN
1	D	138	HIS
1	D	155	ASN
1	D	258	GLN
1	D	277	HIS
1	D	281	GLN
1	D	298	HIS
1	D	317	GLN
1	D	394	GLN
1	D	404	HIS
1	D	438	ASN
1	D	439	GLN
1	D	440	ASN
1	D	456	GLN
1	D	472	GLN
1	D	493	GLN
1	D	527	GLN
1	D	630	ASN
1	D	663	GLN
1	D	671	GLN
1	D	733	HIS
1	D	736	ASN
1	D	795	ASN
1	D	811	GLN
1	D	970	ASN
1	D	976	GLN
1	D	983	GLN

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Mol	Chain	Res	Type
1	D	1037	GLN
1	D	1059	ASN
1	D	1134	HIS
1	D	1175	GLN
1	D	1236	GLN
2	E	84	GLN
2	E	96	HIS
2	E	124	GLN
2	E	184	ASN
2	E	210	GLN
2	E	216	HIS
2	E	249	ASN
2	E	250	GLN
2	E	272	ASN
2	E	292	ASN
2	E	318	ASN
2	E	323	HIS
2	E	339	ASN
2	E	400	GLN
2	E	422	GLN
2	E	427	GLN
2	E	450	ASN
2	E	454	HIS
2	F	74	GLN
2	F	84	GLN
2	F	187	HIS
2	F	192	HIS
2	F	279	GLN
2	F	292	ASN
2	F	339	ASN
2	F	400	GLN
2	F	404	ASN
2	F	409	ASN
2	F	454	HIS

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no carbohydrates in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 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	1093/1172 (93%)	0.23	61 (5%)	28	18	27, 80, 112, 124	0
1	D	1094/1172 (93%)	0.15	53 (4%)	34	23	27, 81, 113, 124	0
2	B	355/427 (83%)	0.44	31 (8%)	13	8	81, 111, 129, 141	0
2	C	396/427 (92%)	0.31	20 (5%)	32	21	42, 85, 113, 127	1 (0%)
2	E	355/427 (83%)	0.41	30 (8%)	13	9	64, 108, 129, 144	0
2	F	396/427 (92%)	0.20	22 (5%)	28	18	42, 87, 114, 128	0
All	All	3689/4052 (91%)	0.25	217 (5%)	26	16	27, 88, 120, 144	1 (0%)

All (217) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
2	B	267	ALA	7.6
1	D	583	PRO	7.0
2	F	327	GLY	6.9
1	D	277	HIS	6.2
1	A	563	PRO	6.2
2	E	334	CYS	6.2
2	E	335	VAL	6.2
2	E	305	ASN	6.0
1	A	748	TRP	5.9
1	D	1003	TRP	5.7
1	A	750	PHE	5.6
1	A	272	SER	5.2
2	B	179	GLY	5.1
1	D	582	ASP	5.1
1	D	866	ARG	5.0
1	A	281	GLN	4.7
1	A	723	GLY	4.6
2	C	386	ASP	4.6
2	E	299	LEU	4.5

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Mol	Chain	Res	Type	RSRZ
2	B	241	TRP	4.5
1	A	582	ASP	4.5
1	D	918	TRP	4.5
1	A	654	SER	4.4
1	D	1063	SER	4.1
1	A	91	GLY	4.0
2	F	178	SER	4.0
1	D	723	GLY	4.0
1	A	152	GLU	4.0
1	D	591	LEU	3.9
1	D	1239	PRO	3.9
1	D	221	TYR	3.9
2	B	180	LYS	3.8
2	E	336	LEU	3.8
1	A	233	TYR	3.7
2	E	284	ARG	3.7
1	A	241	PRO	3.7
2	F	423	SER	3.6
1	A	592	LEU	3.6
2	B	335	VAL	3.6
1	D	590	SER	3.5
2	C	137	PRO	3.5
2	E	136	GLY	3.5
2	E	204	LEU	3.5
1	A	277	HIS	3.5
1	A	235	TRP	3.5
1	A	1063	SER	3.5
2	B	243	THR	3.4
1	A	265	LEU	3.4
1	D	178	TYR	3.4
1	A	538	GLU	3.4
1	D	1090	GLU	3.4
1	A	590	SER	3.3
1	D	90	GLN	3.3
2	B	131	LEU	3.3
1	D	219	ALA	3.3
2	C	449	GLU	3.3
1	D	89	GLY	3.3
2	B	200	VAL	3.3
2	B	428	LEU	3.3
2	E	438	LEU	3.2
2	B	462	MET	3.2

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Mol	Chain	Res	Type	RSRZ
1	A	558	LEU	3.2
2	F	419	GLU	3.2
1	A	219	ALA	3.2
1	A	537	GLU	3.2
2	E	245	PRO	3.1
1	A	1054	GLU	3.1
1	D	195	LEU	3.1
1	D	721	ALA	3.0
1	D	617	ARG	3.0
1	A	1185	ILE	3.0
2	B	250	GLN	3.0
2	E	454	HIS	3.0
1	D	758	ASN	3.0
1	A	94	MET	3.0
2	E	133	HIS	3.0
2	E	388	GLY	3.0
2	E	243	THR	3.0
2	E	137	PRO	3.0
2	C	324	GLY	3.0
2	B	205	PRO	2.9
1	D	520	ALA	2.9
2	B	456	ARG	2.9
2	E	275	SER	2.9
2	F	422	GLN	2.9
2	F	341	ASP	2.9
1	D	753	PRO	2.9
1	A	1064	ILE	2.9
1	D	1045	ALA	2.9
1	D	1212	ILE	2.8
1	D	519	GLY	2.8
2	B	334	CYS	2.8
1	A	164	PRO	2.8
1	A	753	PRO	2.8
2	E	179	GLY	2.8
1	D	929	THR	2.8
2	B	130	ALA	2.8
1	D	197	PHE	2.8
2	C	178	SER	2.8
1	A	197	PHE	2.8
1	A	710	ALA	2.8
1	D	235	TRP	2.8
2	F	78	PHE	2.7

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Mol	Chain	Res	Type	RSRZ
1	A	1093	MET	2.7
1	D	368	PRO	2.7
2	B	445	GLU	2.7
2	B	328	ARG	2.7
2	B	336	LEU	2.7
2	C	442	LEU	2.7
1	A	264	GLN	2.7
2	B	427	GLN	2.7
1	A	758	ASN	2.7
1	D	283	LEU	2.6
2	B	133	HIS	2.6
2	F	244	PRO	2.6
1	D	500	ALA	2.6
1	D	1012	VAL	2.6
1	A	867	PRO	2.6
2	E	267	ALA	2.6
2	B	300	ILE	2.6
2	C	385	LEU	2.6
2	C	454	HIS	2.6
1	A	841	LEU	2.6
2	F	390	GLY	2.6
2	C	333	PRO	2.6
1	D	641	GLU	2.6
1	D	234	SER	2.5
2	F	224	ILE	2.5
2	F	85	GLN	2.5
1	A	199	VAL	2.5
1	D	585	TRP	2.5
1	D	1008	LEU	2.5
1	A	918	TRP	2.5
2	E	306	LEU	2.5
1	A	614	TYR	2.5
2	C	77	HIS	2.5
1	A	956	GLY	2.5
1	A	1126	ASP	2.5
2	E	198	ASP	2.5
1	D	363	TYR	2.4
1	A	237	SER	2.4
1	A	273	PHE	2.4
2	B	372	LEU	2.4
2	C	219	PHE	2.4
2	B	291	TYR	2.4

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Mol	Chain	Res	Type	RSRZ
1	A	1090	GLU	2.4
2	F	204	LEU	2.4
1	A	559	LEU	2.4
2	E	203	ARG	2.4
1	A	591	LEU	2.4
2	E	349	TYR	2.4
1	D	150	TYR	2.4
2	C	439	PHE	2.4
2	E	429	TYR	2.4
1	D	615	SER	2.3
1	A	1048	ALA	2.3
2	C	59	ALA	2.3
1	A	1003	TRP	2.3
1	A	266	VAL	2.3
2	C	444	THR	2.3
2	E	483	LYS	2.3
2	F	411	ILE	2.3
2	E	117	SER	2.3
1	A	208	CYS	2.3
1	D	673	ALA	2.3
2	F	386	ASP	2.3
2	B	258	HIS	2.3
1	A	866	ARG	2.3
2	C	278	CYS	2.3
1	D	94	MET	2.3
2	F	129	ASP	2.2
1	A	239	LEU	2.2
1	A	560	PRO	2.2
2	E	417	TYR	2.2
1	D	228	LEU	2.2
2	C	264	ARG	2.2
2	B	444	THR	2.2
1	D	748	TRP	2.2
2	C	289	LEU	2.2
1	A	943	ARG	2.2
2	C	244	PRO	2.2
2	F	333	PRO	2.2
1	A	1233	LYS	2.2
2	E	485	VAL	2.2
1	D	1126	ASP	2.2
2	E	121	PHE	2.2
1	A	206	GLY	2.2

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Mol	Chain	Res	Type	RSRZ
1	D	587	PRO	2.2
1	A	583	PRO	2.1
2	F	308	ASP	2.1
1	A	155	ASN	2.1
2	F	278	CYS	2.1
2	F	382	LYS	2.1
1	D	995	TYR	2.1
2	B	110	LEU	2.1
2	F	134	LYS	2.1
2	B	137	PRO	2.1
2	E	396	ARG	2.1
2	F	122	ARG	2.1
2	E	415	PRO	2.1
2	C	298	GLU	2.1
1	A	501	LYS	2.1
1	D	1093	MET	2.1
2	B	292	ASN	2.1
2	B	404	ASN	2.1
2	B	104	VAL	2.1
1	A	127	LEU	2.1
2	F	123	GLU	2.1
1	D	867	PRO	2.0
2	C	421	MET	2.0
1	A	292	LEU	2.0
1	A	627	ARG	2.0
1	D	579	ARG	2.0
1	D	751	LYS	2.0
1	D	566	LEU	2.0
1	D	152	GLU	2.0
2	B	201	ASN	2.0

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

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

6.3 Carbohydrates ⓘ

There are no carbohydrates in this entry.

6.4 Ligands

There are no ligands in this entry.

6.5 Other polymers

There are no such residues in this entry.