



Full wwPDB EM Validation Report ⓘ

Jan 1, 2025 – 01:40 AM EST

PDB ID : 8RRS
EMDB ID : EMD-19463
Title : Structure of mouse RyR2 solubilised in detergent in open state in complex with Ca²⁺, ATP, caffeine and Nb9657.
Authors : Li, C.; Efremov, R.G.
Deposited on : 2024-01-23
Resolution : 3.40 Å(reported)

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

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<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>
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The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev113
Mogul : 2022.3.0, CSD as543be (2022)
MolProbity : 4.02b-467
buster-report : 1.1.7 (2018)
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.40

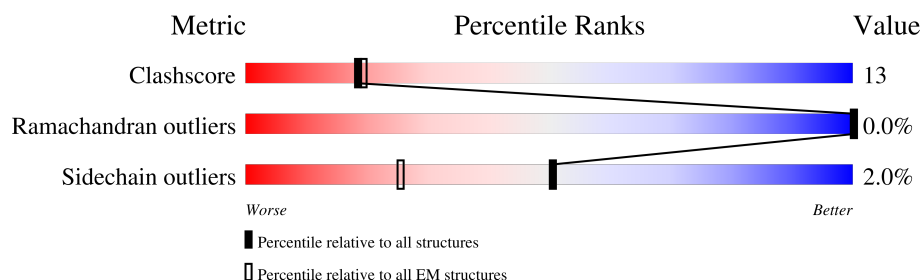
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.40 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	210492	15764
Ramachandran outliers	207382	16835
Sidechain outliers	206894	16415

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

Mol	Chain	Length	Quality of chain
1	A	4966	
1	C	4966	
1	E	4966	
1	F	4966	
2	B	137	
2	D	137	
2	G	137	
2	I	137	

2 Entry composition

There are 6 unique types of molecules in this entry. The entry contains 136400 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called Ryanodine receptor 2.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	4142	Total	C	N	O	S	0	0
			33088	21113	5650	6106	219		
1	C	4142	Total	C	N	O	S	0	0
			33088	21113	5650	6106	219		
1	E	4142	Total	C	N	O	S	0	0
			33088	21113	5650	6106	219		
1	F	4142	Total	C	N	O	S	0	0
			33088	21113	5650	6106	219		

- Molecule 2 is a protein called Nanobody 9657.

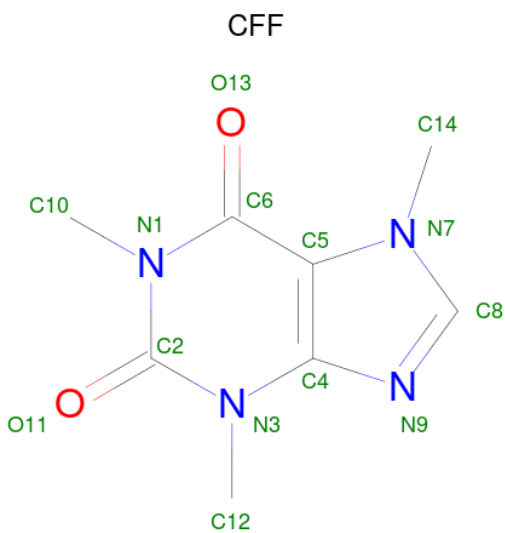
Mol	Chain	Residues	Atoms					AltConf	Trace
2	B	126	Total	C	N	O	S	0	0
			965	595	170	195	5		
2	D	126	Total	C	N	O	S	0	0
			965	595	170	195	5		
2	G	126	Total	C	N	O	S	0	0
			965	595	170	195	5		
2	I	126	Total	C	N	O	S	0	0
			965	595	170	195	5		

- Molecule 3 is ADENOSINE-5'-TRIPHOSPHATE (three-letter code: ATP) (formula: $C_{10}H_{16}N_5O_{13}P_3$).



Mol	Chain	Residues	Atoms					AltConf
3	A	1	Total 31	C 10	N 5	O 13	P 3	0
3	C	1	Total 31	C 10	N 5	O 13	P 3	0
3	E	1	Total 31	C 10	N 5	O 13	P 3	0
3	F	1	Total 31	C 10	N 5	O 13	P 3	0

- Molecule 4 is CAFFEINE (three-letter code: CFF) (formula: $\text{C}_8\text{H}_{10}\text{N}_4\text{O}_2$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms				AltConf
4	A	1	Total	C	N	O	0
			14	8	4	2	
4	C	1	Total	C	N	O	0
			14	8	4	2	
4	E	1	Total	C	N	O	0
			14	8	4	2	
4	F	1	Total	C	N	O	0
			14	8	4	2	

- Molecule 5 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		AltConf
5	A	1	Total	Zn	0
			1	1	
5	C	1	Total	Zn	0
			1	1	
5	E	1	Total	Zn	0
			1	1	
5	F	1	Total	Zn	0
			1	1	

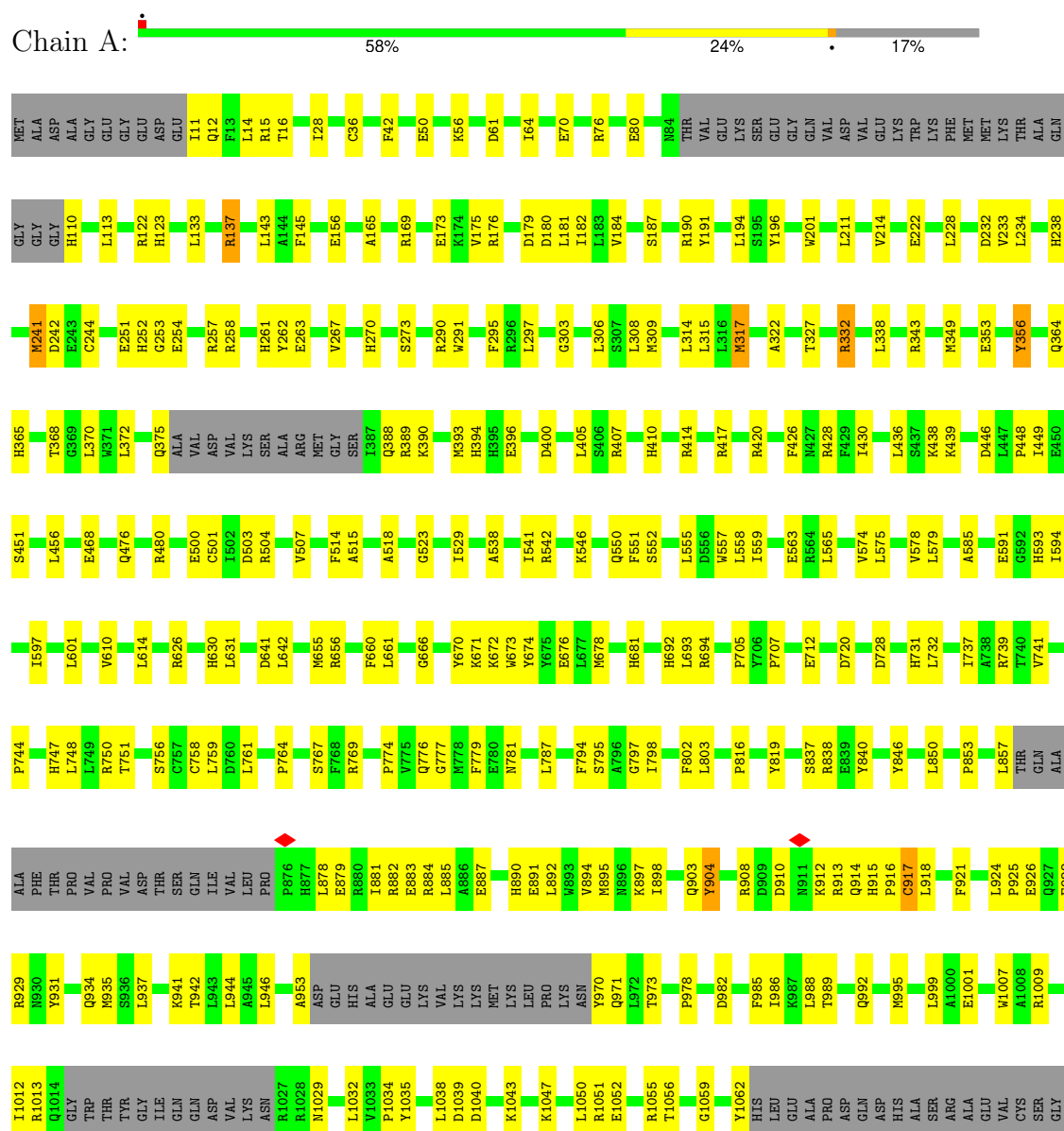
- Molecule 6 is CALCIUM ION (three-letter code: CA) (formula: Ca).

Mol	Chain	Residues	Atoms		AltConf
6	A	1	Total	Ca	0
			1	1	
6	C	1	Total	Ca	0
			1	1	
6	E	1	Total	Ca	0
			1	1	
6	F	1	Total	Ca	0
			1	1	

3 Residue-property plots

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

• Molecule 1: Ryanodine receptor 2



Y2518	P2442	Q2210	L2079	G2007	L1907	L1840	E1733	P1584	Q1436	PRO	R1303	L1207	THR
L2519	THR	Q2210	L2079	I2008	L1908	K1841	L1739	P1585	E1437	GLU	R1304	G1208	GLY
C2520	ALA	Y2219	M2083	LEU	L1909	H1842	L1739	H1588	P1438	PHE	S1305	R1209	GLU
L2526	LYS	M2233	L2086	ASP	C1913	L1844	L1749	Q1589	A1439	ASN	C1310	A1210	F1085
L2527	ASP	M2233	L2086	GLU	D1914	Q1845	P1750	Q1590	N1440	HIS	C1310	Q1211	R1086
T2528	GLY	T2237	R2089	ASP	C1915	L1846	L1754	V1595	G1444	LYS	S1315	N1216	E1091
R2529	LYS	P2238	Q2090	GLY	R1920	I1847	L1758	L1596	V1445	ASP	S1315	F1217	K1092
C2530	VAL	L2239	L2094	LEU	S1928	F1852	L1758	W1597	I1446	ALA	VAL	G1218	Y1094
A2531	THR	A2244	V2098	ASP	D1929	GLU	M1762	N1602	H1451	LYS	GLY	D1220	K1097
F2534	ASP	E2258	R2089	SER	V1932	ALA	M1762	V1609	D1458	PRO	GLY	V1221	A1098
A2535	ILE	P2259	A2100	ASN	V1932	VAL	P1767	S1610	L1459	SER	GLY	Y1236	G1099
G2536	GLU	L2101	GLU	ASP	L1935	PRO	SER	P1611	D1460	ARG	PRO	R1100	R1090
T2537	GLU	E2262	P2102	THR	Q1939	GLU	VAL	G1617	R1461	LEU	GLY	N1242	W1101
H2540	GLU	K2263	K2103	ILE	R1942	GLY	VAL	G1617	T1466	LYS	ALA	Y1102	F1103
A2541	ASP	V2264	T2104	ARG	R1942	GLY	ILE	V1620	T1467	GLN	PHE	D1245	E1104
S2542	ASP	R2265	Y2105	GLY	R1942	GLY	SER	Q1621	T1468	ARG	TYR	D1247	
L2543	THR	R2266		LEU	S1953	THR	ASN	C1622	L1469	PHE	GLY	T1248	M1113
I2544	LEU	C2276	S2121	LEU	A1954	PRO	ASP	L1623	G1470	LEU	PRO	L1251	R1114
D2545	SER	M2278	I2125	SER	A1955	GLU	C1776	E1636	D1471	ARG	LYS	L1251	W1117
S2546	LEU	L2279	V2131	LEU	L1956	GLU	D1786	E1637	H1477	THR	ASN	R1254	
L2548	VAL	R2289	R2132	VAL	R1959	GLU	K1789	R1638	I1480	LYS	ASP	Q1257	P1120
H2549	GLU	M2296	M2133	LYS	K1960	SER	I1791	D1641	R1481	PRO	ASP	F1258	P1124
Y2552	THR	L2145	L2145	THR	R1962	ILE	K1791	E1644	S1483	TYR	PHE	P1262	E1127
R2553	TYR	M2149	M2149	TYR	R1965	ALA	M1795	E1649	M1484	THR	VAL	L1128	L1128
L2554	LEU	F2300	L2154	LEU	S1966	LYS	K1801	E1649	C1485	GLY	ASP	H1265	A1136
S2555	LYS	L2302	F2154	LYS	P1967	LEU	K1802	C1666	Y1486	HIS	SER	H1267	F1137
T2561	LYS	P2158	P2158	LYS	P1968	GLU	E1802	C1666	M1487	SER	ASP	I1268	D1138
Q2564	GLN	N2169	N2169	GLN	Q1971	GLY	L1805	L1677	M1494	ALA	PHE	E1269	
R2565	ALA	L2160	L2160	ALA	I1972	GLU	L1805	L1677	M1494	ARG	GLU	R1272	R1144
D2566	GLU	M2161	M2161	GLU	I1972	GLU	L1805	L1677	M1494	LEU	VAL	I1273	W1156
I2568	PRO	L2164	L2164	PRO	L1975	ALA	R1808	V1681	E1507	THR	LEU	GLY	C1164
E2569	VAL	G2165	G2165	VAL	L1975	LYS	D1809	L1686	C1510	GLU	MET	D1274	W1156
V2570	ALA	M2166	M2166	ALA	F1978	GLY	V1811	Y1694	E1535	VAL	THR	THR	THR
L2573	SER	F2330	V2175	SER	C1985	LYS	T1814	Y1694	E1535	LEU	ALA	THR	THR
Q2578	SER	L2342	K2184	SER	P1988	ARG	T1815	Y1704	T1538	ALA	HIS	ILE	ALA
L2579	ARG	L2342	K2184	ARG	P1988	PRO	E1816	D1705	V1553	ASP	GLY	ASP	V1166
R2580	LYS	K2184	K2184	LYS	P1988	LYS	F1817	L1706	V1553	SER	HIS	SER	M1168
P2581	CYS	E1989	E1989	CYS	E1989	GLU	L1818	L1707	F1554	ASP	VAL	PRO	M1173
S2582	SER	S2056	S2056	SER	E1990	GLU	P1821	D1709	E1587	PRO	PRO	CYS	M1174
M2583	LEU	L2057	L2057	LEU	I1991	R1992	P1821	D1709	E1587	ASP	ASP	LEU	D1177
M2584	VAL	M2065	M2065	VAL	L1995	M1895	Y1827	I1710	R1560	ARG	ARG	K1284	L1177
Q2585	ASP	R2067	R2067	ASP	L1995	E1899	Y1827	I1710	R1560	ILE	ILE	V1285	L1182
H2586	LYS	K2351	K2351	LYS	L1995	P1900	I1831	S1713	I1561	ASP	ASP	Q1287	L1182
L2587	LYS	R2197	R2197	LYS	F1998	V1901	I1831	M1722	N1563	LYS	LYS	F1290	L1190
L2588	GLY	Y2201	Y2201	GLY	E2000	L1903	I1834	I1727	H1576	ASP	LYS	F1290	L1190
L2591	ASP	F2202	F2202	ASP	D2001	Q1904	M1837	V1728	K1577	GLU	GLU	F1192	F1192
V2592	THR	R2358	R2358	THR	C2006	M1905	E1838	P1729	C1583	THR	THR	Q1293	F1201
						C1906	D1839			PRO	LYS	N1295	I1202



- Molecule 1: Ryanodine receptor 2

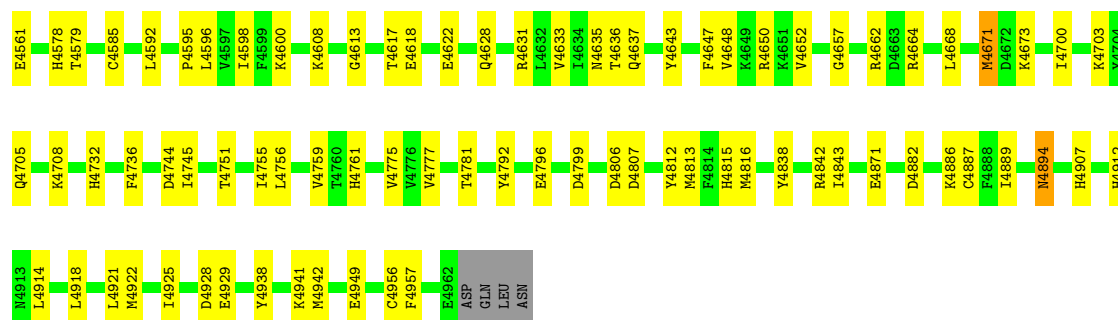
Frequency	Percentage
Often	59%
Sometimes	24%
Rarely	17%



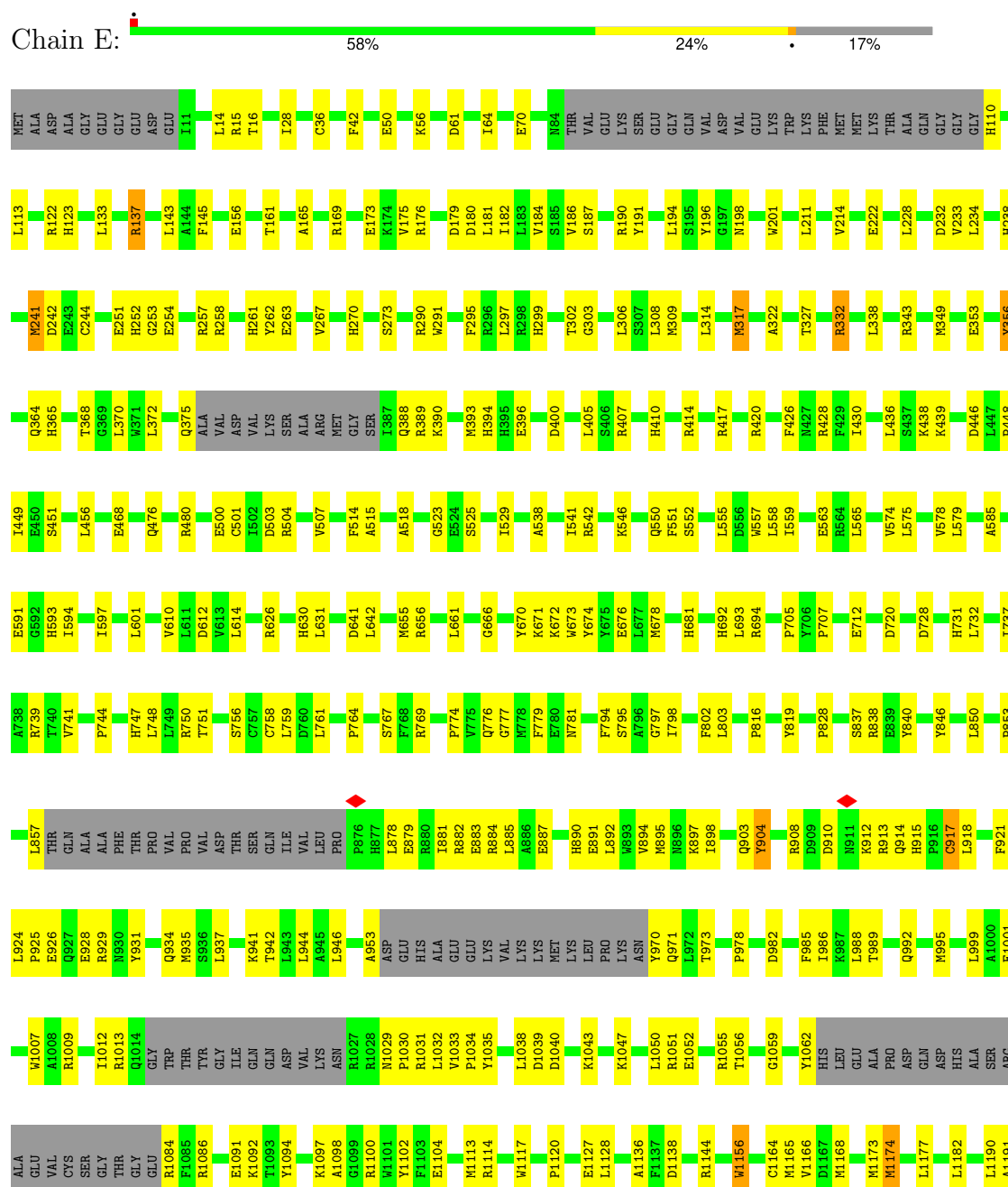
K1853	L1768	Q1590	H1440	HIS	C1310	Q1211	R1084	GLY	Y931	THR	H747	I594	L456	T388
GLU	R1759	V1595	G1444	LYS	S1315	M1216	F1085	TRP	Q934	VAL	L748	I597	VAL	G369
ALA		V1595	W1445	ALA	SER	G1218	E1091	TYR	M935	VAL	R750	L601	VAL	L370
VAL	M1762	N1602	I1446	ALA	SER	K1219	K1092	ILE	S936	ASP	T751	L610	ASP	W371
PRO				GLN	VAL	D1220	T1093	GLN	L937			V610	ASP	L372
GLU	P1767	V1609	H1451	GLU	ALA	V1221	Y1094	GLN	K941	SER	S756	V610	THR	Q375
GLU	S1610	S1610		LYS	GLY			GLN	T942	THR	C757	R480	THR	ALA
GLU	PHE	R1611	D1458	PRO	GLY	Y1236	K1097	ASP	T942	GLN	C768	L614	GLN	ALA
GLY	VAL	G1617	L1459	SER	LEU	Y1236	A1098	VAL	T943	ILE	L759	E500	ILE	VAL
GLY	SER	G1617	D1460	ARG	PRO	Y1236	A1098	LYS	L944	VAL	D760	R626	C501	ASP
THR	ILE		R1461	LEU	GLY	M1242	G1099	ASN	L944	VAL	L761	E502	VAL	VAL
PRO		V1620		LYS	ALA	N1242	G1099	ASN	A945	LEU	L761	H630	LYS	LYS
GLU	SER	V1620	T1466	GLN	GLY	R1245	R1100	R1027	L946	PRO	P764	L631	R504	SER
GLU	ASN	Q1621	V1467	GLN	GLY	D1246	W1101	R1028					R504	ALA
LYS	C1622	L1623	T1468	PHE	PHE	F1102	Y1102	M1029	A953	H877	S767	D641	V507	ARG
ILE	C1776		T1469	TYR	TYR	T1247	F1103	L1032	ASP	L878	R768	L642	GLU	MET
ILE			L1469	GLY	GLY	T1248	E1104	V1033	HIS	R879	R769	H613	GLY	GLY
ILE	D1786	E1636	G1470	LEU	LYS	L1251	M1113	P1034	ALA	R880	P774	M655	F514	SER
LYS		N1637	D1471	ARG	ASN	R1254	R1114	Y1035	GLU	I881	V775	R656	A515	Q388
GLU	K1789	R1638	H1477	ARG	ASP				GLU	R882	G776	L661	D516	R388
ASP	A1790		I1480	THR	ASP	Q1257	W1117	L1038	GLU	R883	G777	L661	V517	R389
ALA	K1791		K1481	PRO	GLU	F1258	P1120	D1039	LYS	R884	G777	L661	A518	K390
LYS		E1644	S1483	ASP	ASP			D1040	VAL	L885	G777	L666		
GLU	M1795		M1487	PHE	PHE				LYS	A886	M778		G523	
GLY		E1649	M1494	SER	ASP	P1262	E1127	K1043	MET	E887	F779	Y670	M393	
GLU	K1801		E1507	LEU	VAL		L1128		LYS	H890	E780	Y670	H394	
GLU	E1802		C1510	GLU	ASP	H1265	L1128	K1047	LYS	E891	N781	K671	H395	
GLU		C1666	T1486	GLY	ASP	E1266	A1136		PRO	R891	F794	K672	E396	
ALA	L1805	L1677	M1487	HIS	SER	H1267	F1137	L1050	PRO	R892	W673	W673	A538	
LYS	R1808		M1487	SER	PHE	I1268	D1138	R1051	LYS	R893	S795	Y675		D400
GLY	D1809		M1494	ALA	GLY	E1269		E1052	ASN	M894	A796	V675	I541	
GLY	P1810			VAL	VAL		R1144		Y970	R895	G797	R695	R542	L405
LYS	V1811			LEU	GLY					R896	L798	L677	S406	
ARG		L1686		THR	LEU	R1272		R1055	T973	K897	F802	H678	K546	R407
PRO	T1814	Y1694	C1510	GLU	MET	I1273	W1156	T1056		I897	F802	H681	Q550	H410
LYS	T1815			VAL	LYS	D1274			P978	I898	L803		F551	
GLU	E1816	Y1704	E1535	THR	THR	GLY	C1164	G1059	D932	Q903	P816	H692	S552	R414
GLY	F1817	D1705	LEU	ALA	ALA	THR	M1165			Y904	Y819	L693		
	L1818	L1706	T1538	ALA	HIS	ILE	V1166	Y1062				R694	L555	
		L1707		ASP	GLY	ASP	D1167	HIS	F985	R908	Y819		D556	R417
	Y1827	L1708		ASP	HIS	SER	M1168	LEU	R986	D909	P828		W557	R420
		F1553	V1554	ARG	LEU	PRO	M1173	ALA	R987	D910	L988		L558	F426
	I1831	I1710		ASP	VAL	CYS	M1174	PRO	T989	N911	E712		I559	M427
			E1557	ASP	PRO	LEU		ASP	Q982	K912	R838		F429	R428
	I1834	S1713		ASP	ASP		L1177	GLN		R913	E839		E563	F429
			R1560	ARG	ILE	K1284		ASP		Q914	Y840		R394	R428
	N1837	M1722	I1561	ASP	ASP	V1285		ASP		H915			L565	F429
	E1838		K1562	LYS	LYS	Q1287	L1182	HIS	M995	P916	Y846		L565	I430
	C1906	I1727	M1563	ASP	ASP			ALA		C917				
	L1907	V1728		GLY	GLY	F1290	L1190	SER	L999	C917	L850	H731	V574	L436
	K1841		H1576	LYS	VAL		A1191	ARG	E1000	L918	L732	L732	L575	S437
			K1577	GLY	THR		F1192	ALA	E1001	F921	P853	I737	L578	K438
	H1842	E1733		THR	THR	Q1293	F1201	GLU	W1007				V578	K439
	I1843		C1583	PRO	PRO	M1294	I1202	VAL	A1008	L924	L857	R739	L579	D446
	C1913	L1739		LYS	LYS	M1295	I1202	CYS	A1008	P925	THR	A738	A585	
	D1914	P1584		PRO	PRO			SER	R1009	E926	GLN			P448
	C1915	P1585		GLY	GLY	R1303	L1207	GLY		Q927	ALA	V741	E591	I449
	I1847	P1750		PHE	PHE	L1304	G1208	THR	I1012	R928	ALA		G592	E450
			H1583	ASN	ASN	S1305	V1209	GLY	R1013	R929	ALA	P744	E592	
				ASN	ASN		A1210	GLU	Q1014	N930	PHE		H593	S451

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<div><div></div><div>A2618</div></div>	<div><div></div><div>A2619</div></div>	<div><div></div><div>A2634</div></div>	<div><div></div><div>A2635</div></div>	<div><div></div><div>A2636</div></div>	<div><div></div><div>A2637</div></div>	<div><div></div><div>A2638</div></div>	<div><div></div><div>A2639</div></div>	<div><div></div><div>A2640</div></div>	<div><div></div><div>A2641</div></div>	<div><div></div><div>A2642</div></div>	<div><div></div><div>A2643</div></div>	<div><div></div><div>A2644</div></div>	<div><div></div><div>A2645</div></div>	<div><div></div><div>A2646</div></div>	<div><div></div><div>A2647</div></div>	<div><div></div><div>A2648</div></div>	<div><div></div><div>A2649</div></div>	<div><div></div><div>A2650</div></div>	<div><div></div><div>A2651</div></div>	<div><div></div><div>A2652</div></div>	<div><div></div><div>A2653</div></div>	<div><div></div><div>A2654</div></div>	<div><div></div><div>A2655</div></div>	<div><div></div><div>A2656</div></div>	<div><div></div><div>A2657</div></div>	<div><div></div><div>A2658</div></div>	<div><div></div><div>A2659</div></div>	<div><div></div><div>A2660</div></div>	<div><div></div><div>A2661</div></div>	<div><div></div><div>A2662</div></div>	<div><div></div><div>A2663</div></div>	<div><div></div><div>A2664</div></div>	<div><div></div><div>A2665</div></div>	<div><div></div><div>A2666</div></div>	<div><div></div><div>A2667</div></div>	<div><div></div><div>A2668</div></div>	<div><div></div><div>A2669</div></div>	<div><div></div><div>A2670</div></div>	<div><div></div><div>A2671</div></div>	<div><div></div><div>A2672</div></div>	<div><div></div><div>A2673</div></div>	<div><div></div><div>A2674</div></div>	<div><div></div><div>A2675</div></div>	<div><div></div><div>A2676</div></div>	<div><div></div><div>A2677</div></div>	<div><div></div><div>A2678</div></div>	<div><div></div><div>A2679</div></div>	<div><div></div><div>A2680</div></div>	<div><div></div><div>A2681</div></div>	<div><div></div><div>A2682</div></div>	<div><div></div><div>A2683</div></div>	<div><div></div><div>A2684</div></div>	<div><div></div><div>A2685</div></div>	<div><div></div><div>A2686</div></div>	<div><div></div><div>A2687</div></div>	<div><div></div><div>A2688</div></div>	<div><div></div><div>A2689</div></div>	<div><div></div><div>A2690</div></div>	<div><div></div><div>A2691</div></div>	<div><div></div><div>A2692</div></div>	<div><div></div><div>A2693</div></div>	<div><div></div><div>A2694</div></div>	<div><div></div><div>A2695</div></div>	<div><div></div><div>A2696</div></div>	<div><div></div><div>A2697</div></div>	<div><div></div><div>A2698</div></div>	<div><div></div><div>A2699</div></div>	<div><div></div><div>A2700</div></div>	<div><div></div><div>A2701</div></div>	<div><div></div><div>A2702</div></div>	<div><div></div><div>A2703</div></div>	<div><div></div><div>A2704</div></div>	<div><div></div><div>A2705</div></div>	<div><div></div><div>A2706</div></div>	<div><div></div><div>A2707</div></div>	<div><div></div><div>A2708</div></div>	<div><div></div><div>A2709</div></div>	<div><div></div><div>A2710</div></div>	<div><div></div><div>A2711</div></div>	<div><div></div><div>A2712</div></div>	<div><div></div><div>A2713</div></div>	<div><div></div><div>A2714</div></div>	<div><div></div><div>A2715</div></div>	<div><div></div><div>A2716</div></div>	<div><div></div><div>A2717</div></div>	<div><div></div><div>A2718</div></div>	<div><div></div><div>A2719</div></div>	<div><div></div><div>A2720</div></div>	<div><div></div><div>A2721</div></div>	<div><div></div><div>A2722</div></div>	<div><div></div><div>A2723</div></div>	<div><div></div><div>A2724</div></div>	<div><div></div><div>A2725</div></div>	<div><div></div><div>A2726</div></div>	<div><div></div><div>A2727</div></div>	<div><div></div><div>A2728</div></div>	<div><div></div><div>A2729</div></div>	<div><div></div><div>A2730</div></div>	<div><div></div><div>A2731</div></div>	<div><div></div><div>A2732</div></div>	<div><div></div><div>A2733</div></div>	<div><div></div><div>A2734</div></div>	<div><div></div><div>A2735</div></div>	<div><div></div><div>A2736</div></div>	<div><div></div><div>A2737</div></div>	<div><div></div><div>A2738</div></div>	<div><div></div><div>A2739</div></div>	<div><div></div><div>A2740</div></div>	<div><div></div><div>A2741</div></div>	<div><div></div><div>A2742</div></div>	<div><div></div><div>A2743</div></div>	<div><div></div><div>A2744</div></div>	<div><div></div><div>A2745</div></div>	<div><div></div><div>A2746</div></div>	<div><div></div><div>A2747</div></div>	<div><div></div><div>A2748</div></div>	<div><div></div><div>A2749</div></div>	<div><div></div><div>A2750</div></div>	<div><div></div><div>A2751</div></div>	<div><div></div><div>A2752</div></div>	<div><div></div><div>A2753</div></div>	<div><div></div><div>A2754</div></div>	<div><div></div><div>A2755</div></div>	<div><div></div><div>A2756</div></div>	<div><div></div><div>A2757</div></div>	<div><div></div><div>A2758</div></div>	<div><div></div><div>A2759</div></div>	<div><di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● Molecule 1: Ryanodine receptor 2





E3717	R3614	E3548	R3448	I3354	I3283	R3184	L3074	PRO	ASP	SER	K2680
Q3721	A3615	R3549	K3449	L3361	N3286	T3185	L3078	LEU	LEU	ILE	GLU
Q3727	N3617	V3550	K3450	A3362	L3287	R3186	Q3078	CYS	ASP	ASP	SER
A3728	L3618	G3552	K3454	R3363	G3288	R3189	K3083	THR	LEU	ALA	ASN
R3729	F3619	G3553	R3457	D3364	I3289	P3197	S3084	GLY	THR	HIS	TYR
L3730	L3620	I3556	R3465	L3365	E3291	P3197	Q3085	HIS	PRO	GLY	VAL
H3731	Y3623	L3557	S3463	F3368	G3292	P3197	V3089	ALA	SER	TYR	SER
R3733	V3627	S3564	L3464	Y3369	A3293	E3201	V3089	N2997	LEU	PRO	MET
A3736	E3636	L3572	V3466	L3372	N3295	D3202	I3093	K2918	ARG	ALA	GLU
I3764	A3644	V3376	K3470	V3376	L3298	P3208	I3094	E2999	ALA	ILE	LYS
N3769	G3647	D3377	R3471	D3377	F3301	E3211	I3095	K3000	ASP	SER	GLN
A3648	G3648	N3378	L3472	N3378	S3302	K3212	T3096	E3001	MET	ASP	SER
F3649	F3649	R3380	P3474	R3380	Q3303	L3213	T3097	M3002	SER	SER	SER
V3772	LEU	A3381	I3475	A3381	Q3303	M3214	S3105	L3006	ASN	ASN	ASP
K3775	PRO	F3382	I3479	F3382	I3306	I3217	F3108	F3007	THR	THR	GLU
M3776	VAL	N3383	C3480	L3384	R3307	I3217	E3109	L3010	LEU	LEU	GLY
L3802	GLU	K3385	A3481	K3385	K3308	H3110	H3110	L3013	SER	SER	ASN
N3805	ASP	E3386	P3482	E3386	V3309	L3220	F3116	V3014	ARG	PHE	ASN
R3809	GLU	F3387	G3483	F3387	K3310	I3225	G3117	I3018	L2835	L2836	PRO
Q3810	ALA	N3388	D3484	N3388	L3313	R3226	G3117	A3025	M2839	R2771	GLN
R3811	SER	F3389	Q3485	F3389	L3314	R3226	L3120	A2995	I2774	I2774	PRO
V3814	LYS	A3391	L3487	A3391	K3315	M3233	L3121	H2996	K2767	K2767	VAL
E3814	LYS	E3392	I3488	E3392	H3316	M3234	I3122	V2938	A2852	L2778	ASP
M3818	ALA	E3393	L3488	E3393	F3318	V3236	E3123	N3030	K2855	M2781	THR
V3828	VAL	L3394	L3490	L3394	L3319	V3237	D3124	K3030	K2856	K2784	ASN
D3831	TRP	F3395	L3490	F3395	F3320	L3238	V3125	C3031	L2857	V2784	THR
D3832	LYS	K3396	R3505	K3396	L3321	R3239	Q3126	I3034	E2858	TRP	ILE
E3833	LYS	M3397	R3508	M3397	M3322	M3240	G3129	I3034	L2859	ARG	THR
F3834	ARG	V3398	I3508	V3398	E3323	L3241	L3133	L3035	E2860	ARG	THR
R3840	ARG	A3399	T3512	A3399	L3324	Y3244	T3134	T3038	S2861	ILE	THR
Q3843	ALA	E3400	G3516	E3400	L3325	M3245	S3135	L3039	K2862	GLY	THR
L3844	VAL	F3401	K3517	F3401	K3326	S3246	S3135	D3040	G2863	ARG	THR
N3850	VAL	I3403	L3518	I3403	K3328	R3247	V3146	A3041	G2864	ARG	THR
S3851	GLY	F3411	A3522	F3411	M3331	W3249	V3147	R3042	G2865	GLY	THR
D3852	GLY	N3425	A3522	N3425	V3332	E3250	R3151	T3043	G2865	GLY	THR
F3853	ASP	M3426	W3525	M3426	E3335	H3256	G3155	K3046	N2866	ASP	THR
Q3854	CYS	M3527	Q3526	M3527	E3335	R3256	E3156	T3047	H2867	MET	THR
R3858	PHE	Y3530	F3516	Y3530	L3339	M3262	A3159	G3048	L2877	ALA	THR
T3859	ARG	K3435	K3517	K3435	K3340	H3271	A3160	L3049	A2878	LEU	THR
I3870	ARG	S3436	L3533	S3436	A3341	M3272	F3161	D3050	K2881	TYR	THR
V3874	PRO	P3534	P3533	P3534	E3342	N3273	P3166	V3052	Q2889	ASN	THR
D3875	LEU	R3535	P3534	R3535	A3343	T3274	P3166	K3053	D2890	ARG	THR
	TRP	R3536	R3535	R3536	G3345	L3275	T3167	R3057	I2891	ARG	THR
	ASN	T3537	T3537	T3537	D3346	L3276	T3172	E3065	L2982	ILE	THR
	LEU	P3540	P3540	P3540	S3347	I3279	N3178	K3069	S2983	GLN	THR
	LYS	V3547	V3547	V3547	L3352	K3281	V3179	T3070	K2893	SER	THR
					L3353	I3282	X3180	M3071	G2905	PHE	THR

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• Molecule 1: Ryanodine receptor 2

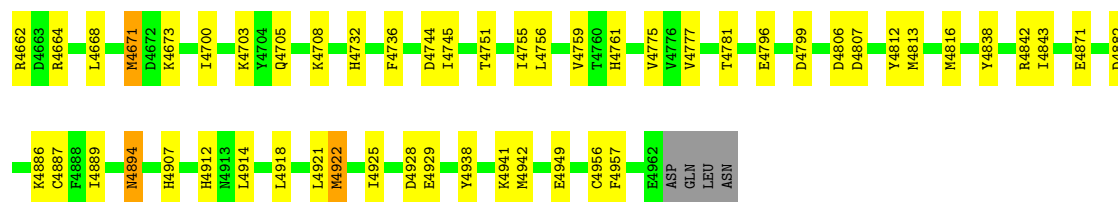
Chain F:  58% 24% 17%

MET	ALA	ASP	ALA	GLY	GLU	GLY	GLY	ASP	GLU	I11	L14	R15	T16	I28	C36	F42	E50	D61	I64	E70	R84	THR	VAL	GLU	LYS	SER	GLU	GLY	GLN	VAL	ASP	VAL	VAL	LYS	LYS	THR	ALA	GLN	GLY	GLY	H110	L113					
R122	H123	L133	L133	R137	L143	A144	F145	E156	T161	A165	R169	E173	R174	V175	R176	D179	D180	L181	I182	I183	V184	S185	V186	S187	R190	Y191	L194	S195	Y196	W201	L211	V214	E222	L228	D232	V233	L234	H238	M241	D242	E243						
C244	E251	H252	C253	E254	R257	R258	H261	Y262	E263	V267	H270	S273	R290	W291	F295	R296	L297	R298	H299	T302	G303	L306	S307	L308	M309	L314	M317	A322	T327	R332	L338	R343	M349	F353	Y356	Q364	K365										
T368	G369	L370	W371	L372	Q375	ALA	VAL	ASP	VAL	LYS	SER	ALA	ARG	MET	GLY	SER	T387	Q388	R389	K390	K393	H394	H395	E396	D400	L405	S406	R407	H410	R414	R417	R420	F426	N427	R428	F429	I430	L436	S437	K438	K439	D446	L447	P448	I449	E450	S451

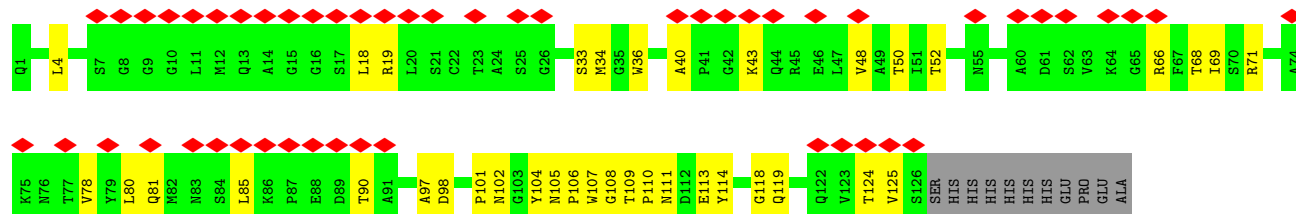
L1907	L1908	L1909	C1913	C1914	C1915	R1920	S1928	D1929	V1932	L1935	Q1939	R1942	S1953	A1954	A1955	L1956	R1959	K1960	T1961	R1962	R1965	S1966	P1967	P1968	Q1971	I1972	L1975	F1978	K1979	C1985	P1988	E1989	E1990	I1991	R1992	L1995	F1998	K1999	L1903	Q1904	D2001	C2006																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																
L1840	K1841	H1842	L1843	L1844	Q1845	L1846	I1847	F1852	K1853	GLU	ALA	ALA	VAL	PRO	GLU	GLU	GLY	GLY	THR	THR	PRO	GLU	LYS	GLU	ILE	SER	ILE	GLY	GLU	ALA	LYS	F1967	K1801	E1802	L1805	R1808	D1809	P1810	L1811	T1814	T1815	F1816	L1818	Y1827	I1831	I1834	N1837	E1838	D1839																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																									
V1728	P1729	E1733	L1739	P1750	L1754	L1758	R1759	M1762	P1767	SER	PHE	VAL	THR	ILE	ASN	ASP	C1776	D1786	K1791	M1795	K1801	E1802	L1805	R1808	D1809	P1810	L1811	T1814	T1815	F1816	L1818	Y1827	I1831	I1834	N1837	E1838	D1839	Y1840	L1846	L1847	L1848	L1849	L1850	L1851	L1852	L1853	L1854	L1855	L1856	L1857	L1858	L1859	L1860	L1861	L1862	L1863	L1864	L1865	L1866	L1867	L1868	L1869	L1870	L1871	L1872	L1873	L1874	L1875	L1876	L1877	L1878	L1879	L1880	L1881	L1882	L1883	L1884	L1885	L1886	L1887	L1888	L1889	L1890	L1891	L1892	L1893	L1894	L1895	L1896	L1897	L1898	L1899	L1900	L1901	L1902	L1903	L1904	L1905	L1906	L1907	L1908	L1909	L1910	L1911	L1912	L1913	L1914	L1915	L1916	L1917	L1918	L1919	L1920	L1921	L1922	L1923	L1924	L1925	L1926	L1927	L1928	L1929	L1930	L1931	L1932	L1933	L1934	L1935	L1936	L1937	L1938	L1939	L1940	L1941	L1942	L1943	L1944	L1945	L1946	L1947	L1948	L1949	L1950	L1951	L1952	L1953	L1954	L1955	L1956	L1957	L1958	L1959	L1960	L1961	L1962	L1963	L1964	L1965	L1966	L1967	L1968	L1969	L1970	L1971	L1972	L1973	L1974	L1975	L1976	L1977	L1978	L1979	L1980	L1981	L1982	L1983	L1984	L1985	L1986	L1987	L1988	L1989	L1990	L1991	L1992	L1993	L1994	L1995	L1996	L1997	L1998	L1999	L2000	L2001	L2002	L2003	L2004	L2005	L2006	L2007	L2008	L2009	L2010	L2011	L2012	L2013	L2014	L2015	L2016	L2017	L2018	L2019	L2020	L2021	L2022	L2023	L2024	L2025	L2026	L2027	L2028	L2029	L2030	L2031	L2032	L2033	L2034	L2035	L2036	L2037	L2038	L2039	L2040	L2041	L2042	L2043	L2044	L2045	L2046	L2047	L2048	L2049	L2050	L2051	L2052	L2053	L2054	L2055	L2056	L2057	L2058	L2059	L2060	L2061	L2062	L2063	L2064	L2065	L2066	L2067	L2068	L2069	L2070	L2071	L2072	L2073	L2074	L2075	L2076	L2077	L2078	L2079	L2080	L2081	L2082	L2083	L2084	L2085	L2086	L2087	L2088	L2089	L2090	L2091	L2092	L2093	L2094	L2095	L2096	L2097	L2098	L2099	L2100	L2101	L2102	L2103	L2104	L2105	L2106	L2107	L2108	L2109	L2110	L2111	L2112	L2113	L2114	L2115	L2116	L2117	L2118	L2119	L2120	L2121	L2122	L2123	L2124	L2125	L2126	L2127	L2128	L2129	L2130	L2131	L2132	L2133	L2134	L2135	L2136	L2137	L2138	L2139	L2140	L2141	L2142	L2143	L2144	L2145	L2146	L2147	L2148	L2149	L2150	L2151	L2152	L2153	L2154	L2155	L2156	L2157	L2158	L2159	L2160	L2161	L2162	L2163	L2164	L2165	L2166	L2167	L2168	L2169	L2170	L2171	L2172	L2173	L2174	L2175	L2176	L2177	L2178	L2179	L2180	L2181	L2182	L2183	L2184	L2185	L2186	L2187	L2188	L2189	L2190	L2191	L2192	L2193	L2194	L2195	L2196	L2197	L2198	L2199	L2200	L2201	L2202	L2203	L2204	L2205	L2206	L2207	L2208	L2209	L2210	L2211	L2212	L2213	L2214	L2215	L2216	L2217	L2218	L2219	L2220	L2221	L2222	L2223	L2224	L2225	L2226	L2227	L2228	L2229	L2230	L2231	L2232	L2233	L2234	L2235	L2236	L2237	L2238	L2239	L2240	L2241	L2242	L2243	L2244	L2245	L2246	L2247	L2248	L2249	L2250	L2251	L2252	L2253	L2254	L2255	L2256	L2257	L2258	L2259	L2260	L2261	L2262	L2263	L2264	L2265	L2266	L2267	L2268	L2269	L2270	L2271	L2272	L2273	L2274	L2275	L2276	L2277	L2278	L2279	L2280	L2281	L2282	L2283	L2284	L2285	L2286	L2287	L2288	L2289	L2290	L2291	L2292	L2293	L2294	L2295	L2296	L2297	L2298	L2299	L2300	L2301	L2302	L2303	L2304	L2305	L2306	L2307	L2308	L2309	L2310	L2311	L2312	L2313	L2314	L2315	L2316	L2317	L2318	L2319	L2320	L2321	L2322	L2323	L2324	L2325	L2326	L2327	L2328	L2329	L2330	L2331	L2332	L2333	L2334	L2335	L2336	L2337	L2338	L2339	L2340	L2341	L2342	L2343	L2344	L2345	L2346	L2347	L2348	L2349	L2350	L2351	L2352	L2353	L2354	L2355	L2356	L2357	L2358	L2359	L2360	L2361	L2362	L2363	L2364	L2365	L2366	L2367	L2368	L2369	L2370	L2371	L2372	L2373	L2374	L2375	L2376	L2377	L2378	L2379	L2380	L2381	L2382	L2383	L2384	L2385	L2386	L2387	L2388	L2389	L2390	L2391	L2392	L2393	L2394	L2395	L2396	L2397	L2398	L2399	L2400	L2401	L2402	L2403	L2404	L2405	L2406	L2407	L2408	L2409	L2410	L2411	L2412	L2413	L2414	L2415	L2416	L2417	L2418	L2419	L2420	L2421	L2422	L2423	L2424	L2425	L2426	L2427	L2428	L2429	L2430	L2431	L2432	L2433	L2434	L2435	L2436	L2437	L2438	L2439	L2440	L2441	L2442	L2443	L2444	L2445	L2446	L2447	L2448	L2449	L2450	L2451	L2452	L2453	L2454	L2455	L2456	L2457	L2458	L2459	L2460	L2461	L2462	L2463	L2464	L2465	L2466	L2467	L2468	L2469	L2470	L2471	L2472	L2473	L2474	L2475	L2476	L2477	L2478	L2479	L2480	L2481	L2482	L2483	L2484	L2485	L2486	L2487	L2488	L2489	L2490	L2491	L2492	L2493	L2494	L2495	L2496	L2497	L2498	L2499	L2500	L2501	L2502	L2503	L2504	L2505	L2506	L2507	L2508	L2509	L2510	L2511	L2512	L2513	L2514	L2515	L2516	L2517	L2518	L2519	L2520	L2521	L2522	L2523	L2524	L2525	L2526	L2527	L2528	L2529	L2530	L2531	L2532	L2533	L2534	L2535	L2536	L2537	L2538	L2539	L2540	L2541	L2542	L2543	L2544	L2545	L2546	L2547	L2548	L2549	L2550	L2551	L2552	L2553	L2554	L2555	L2556	L2557	L2558	L2559	L2560	L2561	L2562	L2563	L2564	L2565	L2566	L2567	L2568	L2569	L2570	L2571	L2572	L2573	L2574	L2575	L2576	L2577	L2578	L2579	L2580	L2581	L2582	L2583	L2584	L2585	L2586	L2587	L2588	L2589	L2590	L2591	L2592	L2593	L2594	L2595	L2596	L2597	L2598	L2599	L2600	L2601	L2602	L2603	L2604	L2605	L2606	L2607	L2608	L2609	L2610	L2611	L2612	L2613	L2614	L2615	L2616	L2617	L2618	L2619	L2620	L2621	L2622	L2623	L2624	L2625	L2626	L2627	L2628	L2629	L2630	L2631	L2632	L2633	L2634	L2635	L2636	L2637	L2638	L2639	L2640	L2641	L2642	L2643	L2644	L2645	L2646	L2647	L2648	L2649	L2650	L2651	L2652	L2653	L2654	L2655	L2656	L2657	L2658	L2659	L2660	L2661	L2662	L2663	L2664	L2665	L2666	L2667	L2668	L2669	L2670	L2671	L2672	L2673	L2674	L2675	L2676	L2677	L2678	L2679	L2680	L2681	L2682	L2683	L2684	L2685	L2686	L2687	L2688	L2689	L2690	L2691	L2692	L2693	L2694	L2695	L2696	L2697	L2698	L2699	L2700	L2701	L2702	L2703	L2704	L2705	L2706	L2707	L2708	L2709	L2710	L2711	L2712	L2713	L2714	L2715	L2716	L2717	L2718	L2719	L2720	L2721	L2722	L2723	L2724	L2725	L2726	L2727	L2728	L2729	L2730	L2731	L2732	L2733	L2734	L2735	L2736	L2737	L2738	L2739	L2740	L2741	L2742	L2743	L2744	L2745	L2746	L2747	L2748	L2749	L2750	L2751	L2752	L2753	L2754	L2755	L2756	L2757	L2758	L2759	L2760	L2761	L2762	L2763	L2764	L2765	L2766	L2767	L2768	L2769	L2770	L2771	L2772	L2773	L2774	L2775	L2776	L2777	L2778	L2779	L2780	L2781	L2782	L2783	L2784	L2785	L2786	L2787	L2788	L2789	L2790	L2791	L2792	L2793	L2794	L2795	L2796	L2797	L2798	L2799	L2800	L2801	L2802	L2803	L2804	L2805	L2806	L2807	L2808	L2809	L2810	L2811	L2812	L2813	L2814	L2815	L2816	L2817	L2818	L2819	L2820	L2821	L2822	L2823	L2824	L2825	L2826	L2827	L2828	L2829	L2830	L2831	L2832	L2833	L2834	L2835	L2836	L2837	L2838	L2839	L2840	L2841	L2842	L2843	L2844	L2845	L2846	L2847	L2848	L2849	L2850	L2851	L2852	L2853	L2854	L2855	L2856	L2857	L2858	L2859	L2860	L2861	L2862	L2863	L2864	L2865	L2866	L2867	L2868	L2869	L2870	L2871	L2872	L2873	L2874	L2875	L2876	L2877	L2878	L2879	L2880	L2881	L2882	L2883	L2884	L2885	L2886	L2887	L2888	L2889	L2890	L2891	L2892	L2893	L2894	L2895	L2896	L2897	L2898	L2899	L2900	L2901	L2902	L2903	L2904	L2905	L2906	L2907	L2908	L2909	L2910	L2911	L2912	L2913	L2914	L2915	L2916	L2917	L2918	L2919	L2920	L2921	L2922	L2923	L2924	L2925	L2926	L2927	L2928	L2929	L2930	L2931	L2932	L2933	L2934	L2935	L2936	L2937	L2938	L2939	L2940	L2941	L2942	L2943	L2944	L2945	L2946	L2947	L2948	L2949	L2950	L2951	L2952	L2953	L2954	L2955	L2956	L2957	L2958	L2959	L2960	L2961	L2962	L2963	L2964	L2965	L2966	L2967	L2968	L2969	L2970	L2971	L2972	L2973	L2974	L2975	L2976	L2977	L2978	L2979	L2980	L2981	L2982	L2983	L2984	L2985	L2986	L2987	L2988	L2989	L2990	L2991	L2992	L2993	L2994	L2995	L2996	L2997	L2998	L2999	L3000	L3001	L3002	L3003	L3004	L3005	L3006	L3007	L3008	L3009	L3010	L3011	L3012	L3013	L3014	L3015	L3016	L3017	L3018	L3019	L3020	L3021	L3022	L3023	L3024	L3025	L3026	L3027	L3028	L3029	L3030	L3031	L3032	L3033	L3034	L3035	L3036	L3037	L3038	L3039	L3040	L3041	L3042	L3043	L3044	L3045	L3046	L3047	L3048	L3049	L3050	L3051	L3052	L3053	L3054	L3055	L3056	L3057	L3058	L3059	L3060	L3061	L3062	L3063	L3064	L3065	L3066	L3067	L3068	L3069	L3070	L3071	L3072	L3073	L3074	L3075	L3076	L3077	L3078	L3079	L3080	L30



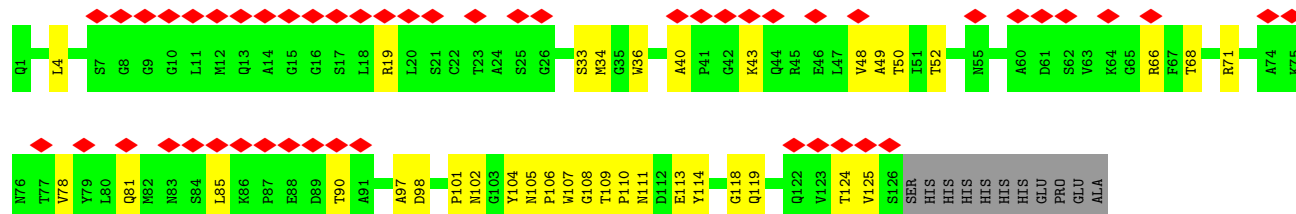




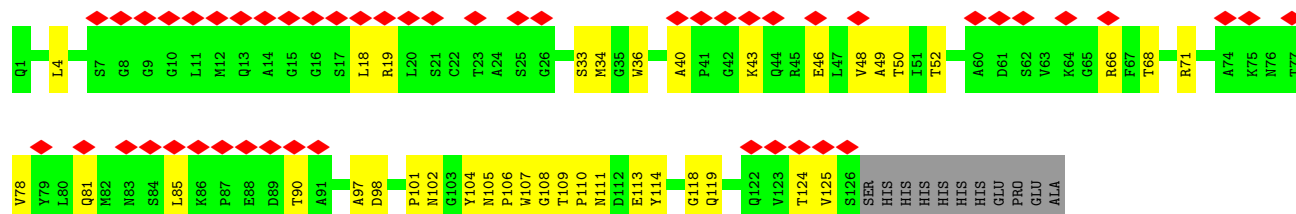
• Molecule 2: Nanobody 9657



• Molecule 2: Nanobody 9657

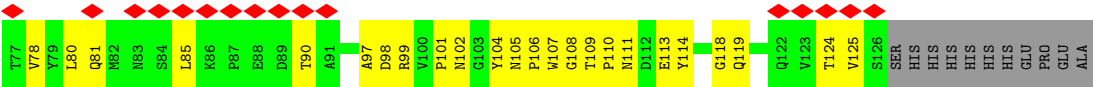


• Molecule 2: Nanobody 9657



• Molecule 2: Nanobody 9657





4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	76852	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	JEOL CRYO ARM 300	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	60	Depositor
Minimum defocus (nm)	1500	Depositor
Maximum defocus (nm)	2500	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor
Maximum map value	5.181	Depositor
Minimum map value	-0.204	Depositor
Average map value	0.074	Depositor
Map value standard deviation	0.151	Depositor
Recommended contour level	0.35	Depositor
Map size (Å)	490.56, 490.56, 490.56	wwPDB
Map dimensions	336, 336, 336	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.46, 1.46, 1.46	Depositor

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: ATP, ZN, CFF, CA

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	0.26	0/33802	0.50	1/45653 (0.0%)
1	C	0.26	0/33802	0.50	1/45653 (0.0%)
1	E	0.26	0/33802	0.50	1/45653 (0.0%)
1	F	0.26	0/33802	0.50	1/45653 (0.0%)
2	B	0.26	0/984	0.51	0/1335
2	D	0.26	0/984	0.51	0/1335
2	G	0.26	0/984	0.51	0/1335
2	I	0.26	0/984	0.51	0/1335
All	All	0.26	0/139144	0.50	4/187952 (0.0%)

There are no bond length outliers.

All (4) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	E	1494	MET	CA-CB-CG	5.38	122.45	113.30
1	F	1494	MET	CA-CB-CG	5.38	122.44	113.30
1	C	1494	MET	CA-CB-CG	5.37	122.43	113.30
1	A	1494	MET	CA-CB-CG	5.36	122.41	113.30

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	33088	0	32662	852	0
1	C	33088	0	32662	837	0
1	E	33088	0	32662	853	0
1	F	33088	0	32662	854	0
2	B	965	0	910	28	0
2	D	965	0	910	27	0
2	G	965	0	910	28	0
2	I	965	0	910	27	0
3	A	31	0	12	3	0
3	C	31	0	12	3	0
3	E	31	0	12	3	0
3	F	31	0	12	3	0
4	A	14	0	10	0	0
4	C	14	0	10	0	0
4	E	14	0	10	0	0
4	F	14	0	10	0	0
5	A	1	0	0	0	0
5	C	1	0	0	0	0
5	E	1	0	0	0	0
5	F	1	0	0	0	0
6	A	1	0	0	0	0
6	C	1	0	0	0	0
6	E	1	0	0	0	0
6	F	1	0	0	0	0
All	All	136400	0	134376	3465	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 13.

All (3465) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2641:ARG:HH12	1:A:2680:MET:HE3	1.42	0.85
1:C:2641:ARG:HH12	1:C:2680:MET:HE3	1.41	0.84
1:E:2641:ARG:HH12	1:E:2680:MET:HE3	1.41	0.84
1:A:4040:GLY:HA3	1:A:4078:ASP:HA	1.60	0.83
1:F:2641:ARG:HH12	1:F:2680:MET:HE3	1.41	0.83
1:C:4040:GLY:HA3	1:C:4078:ASP:HA	1.60	0.83
1:E:4040:GLY:HA3	1:E:4078:ASP:HA	1.60	0.82
1:F:4040:GLY:HA3	1:F:4078:ASP:HA	1.60	0.81
1:F:3392:GLU:HG2	1:F:3479:ILE:HG12	1.63	0.81
1:C:3392:GLU:HG2	1:C:3479:ILE:HG12	1.63	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3376:VAL:O	1:A:3380:ARG:HB2	1.82	0.80
1:C:3376:VAL:O	1:C:3380:ARG:HB2	1.82	0.80
1:E:3376:VAL:O	1:E:3380:ARG:HB2	1.82	0.79
1:A:3392:GLU:HG2	1:A:3479:ILE:HG12	1.63	0.79
1:C:306:LEU:HD11	1:C:314:LEU:HB3	1.65	0.78
1:F:3376:VAL:O	1:F:3380:ARG:HB2	1.82	0.78
1:E:3392:GLU:HG2	1:E:3479:ILE:HG12	1.63	0.78
1:A:306:LEU:HD11	1:A:314:LEU:HB3	1.65	0.77
1:E:306:LEU:HD11	1:E:314:LEU:HB3	1.65	0.77
1:F:306:LEU:HD11	1:F:314:LEU:HB3	1.65	0.77
1:C:2916:ILE:O	1:C:2919:ARG:N	2.20	0.75
1:E:2916:ILE:O	1:E:2919:ARG:N	2.20	0.75
1:F:2916:ILE:O	1:F:2919:ARG:N	2.20	0.75
1:A:2916:ILE:O	1:A:2919:ARG:N	2.20	0.74
1:E:2609:LEU:HD12	1:E:2613:TYR:HE1	1.54	0.73
1:F:891:GLU:HB2	1:F:978:PRO:HB3	1.71	0.72
1:E:891:GLU:HB2	1:E:978:PRO:HB3	1.71	0.72
1:F:2545:ASP:O	1:F:2549:HIS:ND1	2.20	0.72
1:F:794:PHE:HB2	1:F:798:ILE:HG21	1.72	0.72
1:C:2609:LEU:HD12	1:C:2613:TYR:HE1	1.54	0.72
1:E:3049:LEU:HD23	1:E:3052:VAL:HG23	1.72	0.72
1:A:3049:LEU:HD23	1:A:3052:VAL:HG23	1.72	0.71
1:A:2609:LEU:HD12	1:A:2613:TYR:HE1	1.54	0.71
1:C:794:PHE:HB2	1:C:798:ILE:HG21	1.72	0.71
1:C:3238:LEU:HD12	1:C:3241:LEU:HD11	1.73	0.71
1:E:3238:LEU:HD12	1:E:3241:LEU:HD11	1.73	0.71
1:F:2609:LEU:HD12	1:F:2613:TYR:HE1	1.54	0.71
1:F:3238:LEU:HD12	1:F:3241:LEU:HD11	1.73	0.71
1:A:794:PHE:HB2	1:A:798:ILE:HG21	1.72	0.71
1:E:794:PHE:HB2	1:E:798:ILE:HG21	1.72	0.71
1:A:2545:ASP:O	1:A:2549:HIS:ND1	2.21	0.71
1:E:2545:ASP:O	1:E:2549:HIS:ND1	2.21	0.71
1:E:2922:TYR:HB2	1:E:3002:MET:HE1	1.71	0.70
2:B:34:MET:HE1	2:B:78:VAL:HG11	1.74	0.70
1:E:3843:GLN:HG3	1:E:3921:GLU:HG3	1.72	0.70
1:A:3238:LEU:HD12	1:A:3241:LEU:HD11	1.73	0.70
1:A:891:GLU:HB2	1:A:978:PRO:HB3	1.71	0.70
1:A:3843:GLN:HG3	1:A:3921:GLU:HG3	1.72	0.70
1:E:2501:LEU:HD13	1:E:2512:ALA:HA	1.74	0.70
1:F:3843:GLN:HG3	1:F:3921:GLU:HG3	1.72	0.70
1:A:924:LEU:HD12	1:A:925:PRO:HD2	1.74	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:924:LEU:HD12	1:C:925:PRO:HD2	1.74	0.70
1:F:2922:TYR:HB2	1:F:3002:MET:HE1	1.73	0.70
1:C:891:GLU:HB2	1:C:978:PRO:HB3	1.71	0.70
1:A:2922:TYR:HB2	1:A:3002:MET:HE1	1.73	0.70
1:C:2545:ASP:O	1:C:2549:HIS:ND1	2.20	0.70
1:E:3399:ALA:HB1	1:E:3556:VAL:HG21	1.74	0.70
1:F:3049:LEU:HD23	1:F:3052:VAL:HG23	1.72	0.70
1:C:3226:ARG:HA	1:C:3234:MET:SD	2.32	0.70
1:C:3399:ALA:HB1	1:C:3556:VAL:HG21	1.74	0.70
1:F:3399:ALA:HB1	1:F:3556:VAL:HG21	1.74	0.70
1:C:3049:LEU:HD23	1:C:3052:VAL:HG23	1.72	0.69
1:A:3399:ALA:HB1	1:A:3556:VAL:HG21	1.74	0.69
1:C:2501:LEU:HD13	1:C:2512:ALA:HA	1.74	0.69
1:F:3226:ARG:HA	1:F:3234:MET:SD	2.32	0.69
1:A:3273:ASN:HD21	1:A:3310:LYS:HB2	1.56	0.69
1:E:924:LEU:HD12	1:E:925:PRO:HD2	1.74	0.69
1:C:3843:GLN:HG3	1:C:3921:GLU:HG3	1.71	0.69
1:E:3226:ARG:HA	1:E:3234:MET:SD	2.32	0.69
2:G:34:MET:HE1	2:G:78:VAL:HG11	1.74	0.69
1:A:3226:ARG:HA	1:A:3234:MET:SD	2.32	0.69
1:C:3273:ASN:HD21	1:C:3310:LYS:HB2	1.57	0.69
1:E:3273:ASN:HD21	1:E:3310:LYS:HB2	1.57	0.69
1:A:2501:LEU:HD13	1:A:2512:ALA:HA	1.74	0.69
1:A:4013:LEU:HD13	1:A:4121:ALA:HB2	1.75	0.69
1:E:946:LEU:HD13	1:E:995:MET:HG2	1.75	0.69
1:F:764:PRO:HB2	1:F:781:ASN:H	1.58	0.69
1:F:946:LEU:HD13	1:F:995:MET:HG2	1.75	0.69
1:F:2501:LEU:HD13	1:F:2512:ALA:HA	1.74	0.69
1:A:764:PRO:HB2	1:A:781:ASN:H	1.58	0.69
1:E:764:PRO:HB2	1:E:781:ASN:H	1.58	0.69
1:E:3326:LYS:HD3	1:E:3397:MET:HB3	1.75	0.69
1:F:924:LEU:HD12	1:F:925:PRO:HD2	1.74	0.69
1:F:4013:LEU:HD13	1:F:4121:ALA:HB2	1.75	0.69
1:C:764:PRO:HB2	1:C:781:ASN:H	1.58	0.69
1:A:2620:TYR:HB2	1:A:2674:ALA:HB1	1.76	0.69
1:A:3326:LYS:HD3	1:A:3397:MET:HB3	1.75	0.69
1:C:2620:TYR:HB2	1:C:2674:ALA:HB1	1.75	0.69
1:F:3727:GLN:O	1:F:3731:HIS:HB3	1.93	0.69
1:C:234:LEU:HD12	1:C:405:LEU:HB3	1.76	0.68
1:A:3727:GLN:O	1:A:3731:HIS:HB3	1.93	0.68
1:E:3727:GLN:O	1:E:3731:HIS:HB3	1.93	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3273:ASN:HD21	1:F:3310:LYS:HB2	1.57	0.68
1:C:4013:LEU:HD13	1:C:4121:ALA:HB2	1.75	0.68
1:F:3326:LYS:HD3	1:F:3397:MET:HB3	1.75	0.68
1:F:2620:TYR:HB2	1:F:2674:ALA:HB1	1.75	0.68
2:I:34:MET:HE1	2:I:78:VAL:HG11	1.75	0.68
1:C:3326:LYS:HD3	1:C:3397:MET:HB3	1.75	0.68
1:E:4013:LEU:HD13	1:E:4121:ALA:HB2	1.75	0.68
1:F:1303:ARG:NH2	1:F:1590:GLN:OE1	2.27	0.68
1:F:3167:ILE:O	1:F:3247:ARG:NH2	2.27	0.68
1:C:1303:ARG:NH2	1:C:1590:GLN:OE1	2.27	0.68
2:G:36:TRP:HB2	2:G:48:VAL:HB	1.76	0.68
1:E:3000:LYS:HD2	1:E:3043:THR:HG21	1.76	0.68
1:E:3167:ILE:O	1:E:3247:ARG:NH2	2.27	0.68
1:C:3000:LYS:HD2	1:C:3043:THR:HG21	1.76	0.67
1:C:3727:GLN:O	1:C:3731:HIS:HB3	1.93	0.67
1:E:2620:TYR:HB2	1:E:2674:ALA:HB1	1.75	0.67
1:A:1303:ARG:NH2	1:A:1590:GLN:OE1	2.27	0.67
1:A:3993:THR:HA	1:A:3996:LYS:HE2	1.76	0.67
1:E:234:LEU:HD12	1:E:405:LEU:HB3	1.76	0.67
1:F:234:LEU:HD12	1:F:405:LEU:HB3	1.76	0.67
1:C:946:LEU:HD13	1:C:995:MET:HG2	1.75	0.67
1:F:3000:LYS:HD2	1:F:3043:THR:HG21	1.76	0.67
1:A:982:ASP:HB3	1:A:985:PHE:HB2	1.76	0.67
1:A:3000:LYS:HD2	1:A:3043:THR:HG21	1.77	0.67
1:C:3993:THR:HA	1:C:3996:LYS:HE2	1.77	0.67
1:F:3041:ALA:HB3	1:F:3116:PHE:HB3	1.76	0.67
2:I:36:TRP:HB2	2:I:48:VAL:HB	1.76	0.67
1:C:982:ASP:HB3	1:C:985:PHE:HB2	1.76	0.67
1:A:234:LEU:HD12	1:A:405:LEU:HB3	1.76	0.67
1:E:3993:THR:HA	1:E:3996:LYS:HE2	1.77	0.67
1:E:4158:GLN:HB3	1:E:4199:MET:HB3	1.76	0.67
1:F:3993:THR:HA	1:F:3996:LYS:HE2	1.77	0.67
1:A:946:LEU:HD13	1:A:995:MET:HG2	1.75	0.67
1:A:4942:MET:HE1	1:A:4949:GLU:HG3	1.76	0.67
1:E:1303:ARG:NH2	1:E:1590:GLN:OE1	2.27	0.67
1:A:2918:LYS:HD2	1:A:2919:ARG:N	2.10	0.67
1:F:3237:VAL:HA	1:F:3240:MET:HG2	1.77	0.67
1:A:3041:ALA:HB3	1:A:3116:PHE:HB3	1.76	0.66
1:A:3167:ILE:O	1:A:3247:ARG:NH2	2.27	0.66
1:E:3041:ALA:HB3	1:E:3116:PHE:HB3	1.76	0.66
1:A:1560:ARG:NH1	1:A:1561:ILE:O	2.29	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1560:ARG:NH1	1:C:1561:ILE:O	2.29	0.66
1:C:3466:VAL:HG22	1:C:3470:LYS:HE3	1.78	0.66
1:E:2918:LYS:HD2	1:E:2919:ARG:N	2.10	0.66
1:A:3237:VAL:HA	1:A:3240:MET:HG2	1.77	0.66
1:C:4158:GLN:HB3	1:C:4199:MET:HB3	1.76	0.66
1:F:1267:HIS:HB3	1:F:1295:ASN:H	1.61	0.66
1:A:3211:GLU:HA	1:A:3214:MET:HG2	1.77	0.66
1:A:3466:VAL:HG22	1:A:3470:LYS:HE3	1.78	0.66
1:A:4158:GLN:HB3	1:A:4199:MET:HB3	1.76	0.66
2:D:105:ASN:ND2	2:D:108:GLY:O	2.29	0.66
1:E:982:ASP:HB3	1:E:985:PHE:HB2	1.76	0.66
2:I:105:ASN:ND2	2:I:108:GLY:O	2.29	0.66
1:C:3041:ALA:HB3	1:C:3116:PHE:HB3	1.76	0.66
1:E:3211:GLU:HA	1:E:3214:MET:HG2	1.77	0.66
1:C:1437:GLU:O	1:C:1440:ASN:ND2	2.29	0.66
1:F:1560:ARG:NH1	1:F:1561:ILE:O	2.29	0.66
1:F:4942:MET:HE1	1:F:4949:GLU:HG3	1.78	0.66
1:E:3818:MET:N	1:E:3818:MET:SD	2.70	0.66
1:F:982:ASP:HB3	1:F:985:PHE:HB2	1.76	0.66
1:F:3211:GLU:HA	1:F:3214:MET:HG2	1.77	0.66
1:F:3466:VAL:HG22	1:F:3470:LYS:HE3	1.78	0.66
2:B:36:TRP:HB2	2:B:48:VAL:HB	1.76	0.65
2:D:36:TRP:HB2	2:D:48:VAL:HB	1.76	0.65
2:B:105:ASN:ND2	2:B:108:GLY:O	2.29	0.65
1:C:3167:ILE:O	1:C:3247:ARG:NH2	2.27	0.65
1:E:1560:ARG:NH1	1:E:1561:ILE:O	2.29	0.65
1:E:3039:LEU:O	1:E:3110:HIS:NE2	2.30	0.65
1:A:1267:HIS:HB3	1:A:1295:ASN:H	1.61	0.65
1:C:3237:VAL:HA	1:C:3240:MET:HG2	1.77	0.65
1:E:3237:VAL:HA	1:E:3240:MET:HG2	1.77	0.65
1:A:3039:LEU:O	1:A:3110:HIS:NE2	2.29	0.65
1:E:2439:PHE:CZ	1:E:2464:LYS:HD2	2.32	0.65
1:F:3818:MET:N	1:F:3818:MET:SD	2.70	0.65
1:A:2439:PHE:CZ	1:A:2464:LYS:HD2	2.32	0.65
1:C:2918:LYS:HD2	1:C:2919:ARG:N	2.10	0.65
1:F:4158:GLN:HB3	1:F:4199:MET:HB3	1.76	0.65
1:A:317:MET:N	1:A:317:MET:SD	2.70	0.65
1:E:1267:HIS:HB3	1:E:1295:ASN:H	1.61	0.65
1:E:1913:CYS:SG	1:E:2090:GLN:NE2	2.61	0.65
2:G:105:ASN:ND2	2:G:108:GLY:O	2.29	0.65
1:C:3039:LEU:O	1:C:3110:HIS:NE2	2.30	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:222:GLU:HB2	1:E:349:MET:HG3	1.79	0.65
1:E:1437:GLU:O	1:E:1440:ASN:ND2	2.29	0.65
1:F:222:GLU:HB2	1:F:349:MET:HG3	1.79	0.65
1:F:1437:GLU:O	1:F:1440:ASN:ND2	2.29	0.65
1:C:317:MET:N	1:C:317:MET:SD	2.70	0.65
1:E:2355:ASP:HB3	1:E:2358:ARG:HB2	1.79	0.65
1:E:2579:LEU:O	1:E:2615:ARG:NH1	2.30	0.65
1:E:3466:VAL:HG22	1:E:3470:LYS:HE3	1.78	0.65
1:F:3039:LEU:O	1:F:3110:HIS:NE2	2.29	0.65
1:A:946:LEU:HD11	1:A:999:LEU:HB2	1.79	0.65
1:C:3211:GLU:HA	1:C:3214:MET:HG2	1.77	0.65
1:E:317:MET:N	1:E:317:MET:SD	2.70	0.65
1:F:2439:PHE:CZ	1:F:2464:LYS:HD2	2.32	0.65
1:F:2579:LEU:O	1:F:2615:ARG:NH1	2.30	0.65
1:F:2918:LYS:HD2	1:F:2919:ARG:N	2.10	0.65
1:C:2439:PHE:CZ	1:C:2464:LYS:HD2	2.32	0.64
1:C:3614:ARG:HH22	1:C:3618:LEU:HD21	1.62	0.64
1:C:3818:MET:SD	1:C:3818:MET:N	2.70	0.64
1:E:555:LEU:HD21	1:E:578:VAL:HG11	1.79	0.64
1:E:946:LEU:HD11	1:E:999:LEU:HB2	1.79	0.64
1:F:317:MET:SD	1:F:317:MET:N	2.70	0.64
1:A:2579:LEU:O	1:A:2615:ARG:NH1	2.30	0.64
1:C:1267:HIS:HB3	1:C:1295:ASN:H	1.61	0.64
1:C:3151:ARG:HD3	1:C:3236:VAL:HG21	1.79	0.64
1:A:3614:ARG:HH22	1:A:3618:LEU:HD21	1.62	0.64
1:C:2579:LEU:O	1:C:2615:ARG:NH1	2.30	0.64
1:F:671:LYS:HB3	1:F:761:LEU:HB3	1.80	0.64
1:A:222:GLU:HB2	1:A:349:MET:HG3	1.79	0.64
1:A:3818:MET:N	1:A:3818:MET:SD	2.70	0.64
1:F:555:LEU:HD21	1:F:578:VAL:HG11	1.79	0.64
1:A:1437:GLU:O	1:A:1440:ASN:ND2	2.29	0.64
1:C:222:GLU:HB2	1:C:349:MET:HG3	1.79	0.64
1:C:2355:ASP:HB3	1:C:2358:ARG:HB2	1.78	0.64
1:F:3614:ARG:HH22	1:F:3618:LEU:HD21	1.62	0.64
1:C:2938:TYR:HB3	1:C:2955:TYR:HB3	1.79	0.64
1:C:4942:MET:HE1	1:C:4949:GLU:HG3	1.80	0.64
1:E:2265:VAL:HG12	1:E:2326:ARG:HH21	1.63	0.64
1:E:2938:TYR:HB3	1:E:2955:TYR:HB3	1.79	0.64
1:E:3614:ARG:HH22	1:E:3618:LEU:HD21	1.62	0.64
1:F:946:LEU:HD11	1:F:999:LEU:HB2	1.79	0.64
1:F:2265:VAL:HG12	1:F:2326:ARG:HH21	1.63	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2355:ASP:HB3	1:F:2358:ARG:HB2	1.79	0.64
1:E:3151:ARG:HD3	1:E:3236:VAL:HG21	1.79	0.64
1:A:3151:ARG:HD3	1:A:3236:VAL:HG21	1.79	0.64
1:F:4617:THR:OG1	1:F:4618:GLU:OE1	2.16	0.64
1:C:555:LEU:HD21	1:C:578:VAL:HG11	1.79	0.63
1:C:946:LEU:HD11	1:C:999:LEU:HB2	1.79	0.63
1:F:891:GLU:HA	1:F:894:VAL:HG22	1.79	0.63
1:F:2938:TYR:HB3	1:F:2955:TYR:HB3	1.79	0.63
1:C:2265:VAL:HG12	1:C:2326:ARG:HH21	1.63	0.63
1:C:2778:LEU:HA	1:C:2781:MET:HE3	1.81	0.63
2:D:34:MET:HE1	2:D:78:VAL:HG11	1.80	0.63
1:F:3151:ARG:HD3	1:F:3236:VAL:HG21	1.79	0.63
1:A:671:LYS:HB3	1:A:761:LEU:HB3	1.79	0.63
1:A:2938:TYR:HB3	1:A:2955:TYR:HB3	1.79	0.63
1:E:671:LYS:HB3	1:E:761:LEU:HB3	1.79	0.63
1:A:1913:CYS:SG	1:A:2090:GLN:NE2	2.61	0.63
1:F:1845:GLN:NE2	1:F:1852:PHE:O	2.32	0.63
1:A:891:GLU:HA	1:A:894:VAL:HG22	1.79	0.63
1:A:2265:VAL:HG12	1:A:2326:ARG:HH21	1.63	0.63
1:A:2355:ASP:HB3	1:A:2358:ARG:HB2	1.79	0.63
1:C:3217:ILE:HA	1:C:3220:LEU:HG	1.81	0.63
1:C:4617:THR:OG1	1:C:4618:GLU:OE1	2.16	0.63
1:E:4617:THR:OG1	1:E:4618:GLU:OE1	2.16	0.63
1:F:3129:CYS:HB3	1:F:3161:PHE:HE1	1.64	0.63
1:E:891:GLU:HA	1:E:894:VAL:HG22	1.79	0.63
1:F:594:ILE:HD12	1:F:631:LEU:HD13	1.81	0.63
1:C:3129:CYS:HB3	1:C:3161:PHE:HE1	1.64	0.63
1:A:555:LEU:HD21	1:A:578:VAL:HG11	1.79	0.62
1:A:3129:CYS:HB3	1:A:3161:PHE:HE1	1.64	0.62
1:A:4051:MET:HA	1:A:4054:HIS:HB2	1.81	0.62
1:C:1845:GLN:NE2	1:C:1852:PHE:O	2.32	0.62
1:E:1845:GLN:NE2	1:E:1852:PHE:O	2.32	0.62
1:E:4051:MET:HA	1:E:4054:HIS:HB2	1.81	0.62
1:F:1913:CYS:SG	1:F:2090:GLN:NE2	2.61	0.62
1:F:3217:ILE:HA	1:F:3220:LEU:HG	1.81	0.62
1:C:891:GLU:HA	1:C:894:VAL:HG22	1.79	0.62
1:C:2587:LEU:O	1:C:2591:LEU:HG	2.00	0.62
1:E:3129:CYS:HB3	1:E:3161:PHE:HE1	1.64	0.62
1:F:2713:PRO:HD2	1:F:2716:LEU:HD12	1.81	0.62
1:A:1935:LEU:HD11	1:A:1975:LEU:HD22	1.82	0.62
1:C:671:LYS:HB3	1:C:761:LEU:HB3	1.79	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2713:PRO:HD2	1:A:2716:LEU:HD12	1.81	0.62
1:A:3217:ILE:HA	1:A:3220:LEU:HG	1.81	0.62
1:E:2587:LEU:O	1:E:2591:LEU:HG	2.00	0.62
1:A:594:ILE:HD12	1:A:631:LEU:HD13	1.81	0.62
1:C:1935:LEU:HD11	1:C:1975:LEU:HD22	1.82	0.62
1:C:4051:MET:HA	1:C:4054:HIS:HB2	1.81	0.62
1:F:2981:PHE:HB3	1:F:3000:LYS:HE2	1.82	0.62
1:A:2585:GLN:OE1	1:A:2585:GLN:N	2.33	0.62
1:C:594:ILE:HD12	1:C:631:LEU:HD13	1.81	0.62
1:E:1935:LEU:HD11	1:E:1975:LEU:HD22	1.82	0.62
1:F:1789:LYS:HE3	1:F:1834:ILE:HG22	1.82	0.62
1:F:1935:LEU:HD11	1:F:1975:LEU:HD22	1.82	0.62
1:A:1953:SER:H	1:A:1956:LEU:HB2	1.65	0.62
1:A:4622:GLU:HA	1:A:4628:GLN:HE21	1.65	0.62
1:C:671:LYS:HG3	1:C:761:LEU:HD22	1.81	0.62
1:C:1953:SER:H	1:C:1956:LEU:HB2	1.65	0.62
1:C:2981:PHE:HB3	1:C:3000:LYS:HE2	1.82	0.62
1:E:671:LYS:HG3	1:E:761:LEU:HD22	1.81	0.62
1:E:3331:MET:O	1:E:3335:GLU:HG2	2.00	0.62
1:E:4942:MET:HE1	1:E:4949:GLU:HG3	1.81	0.62
1:A:1845:GLN:NE2	1:A:1852:PHE:O	2.32	0.62
1:A:4617:THR:OG1	1:A:4618:GLU:OE1	2.16	0.62
1:C:2585:GLN:N	1:C:2585:GLN:OE1	2.33	0.62
1:E:2974:PHE:HB3	1:E:3038:THR:HG21	1.82	0.62
1:F:1953:SER:H	1:F:1956:LEU:HB2	1.65	0.62
1:F:2585:GLN:N	1:F:2585:GLN:OE1	2.33	0.62
1:F:2587:LEU:O	1:F:2591:LEU:HG	2.00	0.62
1:C:2086:LEU:HD12	1:C:2089:ARG:HD3	1.82	0.61
1:E:2585:GLN:OE1	1:E:2585:GLN:N	2.33	0.61
1:E:2973:TYR:O	1:E:2977:HIS:ND1	2.33	0.61
1:F:3900:GLN:OE1	1:F:3903:ARG:NH1	2.33	0.61
1:A:2412:LYS:HG3	1:A:2415:ALA:H	1.65	0.61
1:A:2974:PHE:HB3	1:A:3038:THR:HG21	1.82	0.61
1:A:3331:MET:O	1:A:3335:GLU:HG2	2.00	0.61
1:C:176:ARG:HD3	1:C:179:ASP:HB3	1.82	0.61
1:C:2973:TYR:O	1:C:2977:HIS:ND1	2.33	0.61
1:E:165:ALA:HB2	1:E:182:ILE:HG12	1.82	0.61
1:E:594:ILE:HD12	1:E:631:LEU:HD13	1.81	0.61
1:E:747:HIS:ND1	1:E:748:LEU:O	2.31	0.61
1:E:2086:LEU:HD12	1:E:2089:ARG:HD3	1.81	0.61
1:F:2973:TYR:O	1:F:2977:HIS:ND1	2.33	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:4051:MET:HA	1:F:4054:HIS:HB2	1.81	0.61
1:F:4132:LEU:HD11	1:F:4148:TYR:HB3	1.83	0.61
1:A:2587:LEU:O	1:A:2591:LEU:HG	2.00	0.61
1:A:2778:LEU:HA	1:A:2781:MET:HE3	1.83	0.61
1:A:3382:LYS:HA	1:A:3385:LYS:HD2	1.82	0.61
1:E:1953:SER:H	1:E:1956:LEU:HB2	1.65	0.61
1:E:2412:LYS:HG3	1:E:2415:ALA:H	1.65	0.61
1:F:165:ALA:HB2	1:F:182:ILE:HG12	1.83	0.61
1:F:3764:ILE:O	1:F:3769:ASN:ND2	2.31	0.61
1:C:2713:PRO:HD2	1:C:2716:LEU:HD12	1.81	0.61
1:C:3331:MET:O	1:C:3335:GLU:HG2	2.00	0.61
1:C:3900:GLN:OE1	1:C:3903:ARG:NH1	2.33	0.61
1:E:953:ALA:H	1:E:1062:TYR:HD1	1.49	0.61
1:E:3382:LYS:HA	1:E:3385:LYS:HD2	1.83	0.61
1:F:671:LYS:HG3	1:F:761:LEU:HD22	1.81	0.61
1:A:2086:LEU:HD12	1:A:2089:ARG:HD3	1.82	0.61
1:A:2669:SER:O	1:A:2972:GLN:NE2	2.34	0.61
1:A:3440:LYS:HA	1:A:3443:ILE:HD11	1.82	0.61
1:A:4132:LEU:HD11	1:A:4148:TYR:HB3	1.82	0.61
1:C:2716:LEU:HD13	1:C:2778:LEU:HB3	1.83	0.61
1:C:3382:LYS:HA	1:C:3385:LYS:HD2	1.83	0.61
1:E:2981:PHE:HB3	1:E:3000:LYS:HE2	1.82	0.61
1:E:3217:ILE:HA	1:E:3220:LEU:HG	1.81	0.61
1:E:4132:LEU:HD11	1:E:4148:TYR:HB3	1.82	0.61
1:F:953:ALA:H	1:F:1062:TYR:HD1	1.49	0.61
1:F:2716:LEU:HD13	1:F:2778:LEU:HB3	1.83	0.61
1:F:3331:MET:O	1:F:3335:GLU:HG2	2.00	0.61
1:F:3382:LYS:HA	1:F:3385:LYS:HD2	1.83	0.61
1:A:176:ARG:HD3	1:A:179:ASP:HB3	1.82	0.61
1:A:671:LYS:HG3	1:A:761:LEU:HD22	1.81	0.61
1:A:2973:TYR:O	1:A:2977:HIS:ND1	2.33	0.61
1:C:4132:LEU:HD11	1:C:4148:TYR:HB3	1.82	0.61
1:E:3440:LYS:HA	1:E:3443:ILE:HD11	1.82	0.61
1:E:3900:GLN:OE1	1:E:3903:ARG:NH1	2.33	0.61
1:F:2922:TYR:O	1:F:2926:GLN:HB2	2.01	0.61
2:I:68:THR:HB	2:I:81:GLN:HB2	1.83	0.61
1:A:3233:MET:HA	1:A:3237:VAL:HB	1.83	0.61
1:A:3900:GLN:OE1	1:A:3903:ARG:NH1	2.33	0.61
1:C:953:ALA:H	1:C:1062:TYR:HD1	1.49	0.61
1:C:4079:TYR:O	1:C:4083:VAL:HG12	2.01	0.61
1:E:476:GLN:NE2	1:E:3677:GLU:OE1	2.34	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1789:LYS:HE3	1:E:1834:ILE:HG22	1.82	0.61
1:E:2922:TYR:O	1:E:2926:GLN:HB2	2.01	0.61
1:E:3233:MET:HA	1:E:3237:VAL:HB	1.83	0.61
1:E:4079:TYR:O	1:E:4083:VAL:HG12	2.01	0.61
1:F:4195:THR:O	1:F:4199:MET:HG3	2.01	0.61
1:A:2343:LEU:HD23	1:A:2430:ASP:HA	1.82	0.61
1:A:2716:LEU:HD13	1:A:2778:LEU:HB3	1.83	0.61
1:C:1086:ARG:HH21	1:C:1251:LEU:HD13	1.66	0.61
1:C:2343:LEU:HD23	1:C:2430:ASP:HA	1.82	0.61
1:E:2713:PRO:HD2	1:E:2716:LEU:HD12	1.81	0.61
1:E:3384:LEU:HD21	1:E:3474:PRO:HB2	1.83	0.61
1:F:2086:LEU:HD12	1:F:2089:ARG:HD3	1.82	0.61
1:F:4079:TYR:O	1:F:4083:VAL:HG12	2.01	0.61
1:A:476:GLN:NE2	1:A:3677:GLU:OE1	2.34	0.61
1:A:747:HIS:ND1	1:A:748:LEU:O	2.31	0.61
1:E:1961:THR:HA	1:E:1965:ARG:HB2	1.83	0.61
1:E:2343:LEU:HD23	1:E:2430:ASP:HA	1.82	0.61
1:F:2974:PHE:HB3	1:F:3038:THR:HG21	1.82	0.61
2:B:68:THR:HB	2:B:81:GLN:HB2	1.83	0.60
1:C:777:GLY:HA3	1:C:1469:LEU:HD12	1.83	0.60
1:C:2669:SER:O	1:C:2972:GLN:NE2	2.34	0.60
1:C:4195:THR:O	1:C:4199:MET:HG3	2.01	0.60
1:E:2716:LEU:HD13	1:E:2778:LEU:HB3	1.83	0.60
1:E:4622:GLU:HA	1:E:4628:GLN:HE21	1.65	0.60
1:F:3233:MET:HA	1:F:3237:VAL:HB	1.83	0.60
1:A:165:ALA:HB2	1:A:182:ILE:HG12	1.83	0.60
1:A:953:ALA:H	1:A:1062:TYR:HD1	1.49	0.60
1:A:4195:THR:O	1:A:4199:MET:HG3	2.01	0.60
1:C:2661:PHE:HD2	1:C:2965:VAL:HG21	1.66	0.60
1:E:2519:LEU:HD21	1:E:2554:LEU:HD21	1.83	0.60
1:F:2669:SER:O	1:F:2972:GLN:NE2	2.34	0.60
1:A:2981:PHE:HB3	1:A:3000:LYS:HE2	1.82	0.60
1:C:476:GLN:NE2	1:C:3677:GLU:OE1	2.34	0.60
1:C:2922:TYR:O	1:C:2926:GLN:HB2	2.01	0.60
1:C:3233:MET:HA	1:C:3237:VAL:HB	1.83	0.60
1:E:1827:TYR:CZ	1:E:1831:ILE:HD11	2.37	0.60
1:F:176:ARG:HD3	1:F:179:ASP:HB3	1.82	0.60
1:F:332:ARG:NH1	1:F:364:GLN:OE1	2.34	0.60
1:F:887:GLU:HA	1:F:890:HIS:CD2	2.37	0.60
1:F:2237:THR:HG22	1:F:2239:LEU:H	1.67	0.60
1:F:4622:GLU:HA	1:F:4628:GLN:HE21	1.65	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:68:THR:HB	2:G:81:GLN:HB2	1.83	0.60
1:A:1086:ARG:HH21	1:A:1251:LEU:HD13	1.66	0.60
1:A:1144:ARG:HH11	1:A:1191:ALA:HA	1.66	0.60
1:A:1789:LYS:HE3	1:A:1834:ILE:HG22	1.82	0.60
1:A:2519:LEU:HD21	1:A:2554:LEU:HD21	1.83	0.60
1:C:165:ALA:HB2	1:C:182:ILE:HG12	1.82	0.60
1:C:2412:LYS:HG3	1:C:2415:ALA:H	1.65	0.60
1:C:2974:PHE:HB3	1:C:3038:THR:HG21	1.82	0.60
1:E:1086:ARG:HH21	1:E:1251:LEU:HD13	1.66	0.60
1:E:3089:VAL:O	1:E:3093:ILE:HG12	2.01	0.60
1:F:1144:ARG:HH11	1:F:1191:ALA:HA	1.66	0.60
1:C:1827:TYR:CZ	1:C:1831:ILE:HD11	2.37	0.60
1:E:2669:SER:O	1:E:2972:GLN:NE2	2.34	0.60
1:F:476:GLN:NE2	1:F:3677:GLU:OE1	2.34	0.60
1:F:2343:LEU:HD23	1:F:2430:ASP:HA	1.82	0.60
1:A:4079:TYR:O	1:A:4083:VAL:HG12	2.01	0.60
1:C:1789:LYS:HE3	1:C:1834:ILE:HG22	1.82	0.60
1:C:2435:ILE:HD12	1:C:2464:LYS:HG2	1.84	0.60
1:C:4622:GLU:HA	1:C:4628:GLN:HE21	1.65	0.60
1:E:2661:PHE:HD2	1:E:2965:VAL:HG21	1.66	0.60
1:F:3089:VAL:O	1:F:3093:ILE:HG12	2.01	0.60
1:A:1827:TYR:CZ	1:A:1831:ILE:HD11	2.37	0.60
1:C:3089:VAL:O	1:C:3093:ILE:HG12	2.01	0.60
1:E:332:ARG:NH1	1:E:364:GLN:OE1	2.34	0.60
1:E:887:GLU:HA	1:E:890:HIS:CD2	2.37	0.60
1:E:2237:THR:HG22	1:E:2239:LEU:H	1.67	0.60
1:E:2928:LEU:HD13	1:E:2970:ILE:HG22	1.84	0.60
1:F:3440:LYS:HA	1:F:3443:ILE:HD11	1.82	0.60
1:A:1174:MET:HB3	1:A:1190:LEU:HA	1.84	0.60
1:A:3089:VAL:O	1:A:3093:ILE:HG12	2.01	0.60
1:C:887:GLU:HA	1:C:890:HIS:CD2	2.37	0.60
1:C:2237:THR:HG22	1:C:2239:LEU:H	1.67	0.60
1:E:1174:MET:HB3	1:E:1190:LEU:HA	1.84	0.60
1:A:707:PRO:HG2	1:A:838:ARG:HB2	1.84	0.60
1:A:777:GLY:HA3	1:A:1469:LEU:HD12	1.83	0.60
2:D:68:THR:HB	2:D:81:GLN:HB2	1.83	0.60
1:E:1144:ARG:HH11	1:E:1191:ALA:HA	1.66	0.60
1:E:4761:HIS:NE2	1:E:4871:GLU:OE2	2.35	0.60
1:F:2412:LYS:HG3	1:F:2415:ALA:H	1.65	0.60
1:A:2435:ILE:HD12	1:A:2464:LYS:HG2	1.84	0.60
1:C:3440:LYS:HA	1:C:3443:ILE:HD11	1.82	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4761:HIS:NE2	1:C:4871:GLU:OE2	2.35	0.60
1:E:176:ARG:HD3	1:E:179:ASP:HB3	1.82	0.60
1:F:332:ARG:NH2	1:F:338:LEU:O	2.35	0.60
1:F:1086:ARG:HH21	1:F:1251:LEU:HD13	1.66	0.60
1:F:4761:HIS:NE2	1:F:4871:GLU:OE2	2.35	0.60
1:A:113:LEU:HD12	1:A:175:VAL:HG21	1.84	0.59
1:A:332:ARG:NH1	1:A:364:GLN:OE1	2.34	0.59
1:E:332:ARG:NH2	1:E:338:LEU:O	2.35	0.59
1:F:2661:PHE:HD2	1:F:2965:VAL:HG21	1.66	0.59
1:A:1929:ASP:OD1	1:A:3612:ARG:NH2	2.35	0.59
1:A:1961:THR:HA	1:A:1965:ARG:HB2	1.83	0.59
1:A:4761:HIS:NE2	1:A:4871:GLU:OE2	2.35	0.59
1:C:332:ARG:NH1	1:C:364:GLN:OE1	2.34	0.59
1:C:707:PRO:HG2	1:C:838:ARG:HB2	1.84	0.59
1:E:1929:ASP:OD1	1:E:3612:ARG:NH2	2.35	0.59
1:F:1827:TYR:CZ	1:F:1831:ILE:HD11	2.37	0.59
1:A:2928:LEU:HD13	1:A:2970:ILE:HG22	1.84	0.59
1:C:2519:LEU:HD21	1:C:2554:LEU:HD21	1.83	0.59
1:E:707:PRO:HG2	1:E:838:ARG:HB2	1.84	0.59
1:F:707:PRO:HG2	1:F:838:ARG:HB2	1.84	0.59
1:F:777:GLY:HA3	1:F:1469:LEU:HD12	1.83	0.59
1:F:1009:ARG:HH21	1:F:1013:ARG:HH12	1.50	0.59
1:A:887:GLU:HA	1:A:890:HIS:CD2	2.37	0.59
1:C:2922:TYR:HB2	1:C:3002:MET:HE1	1.83	0.59
1:F:3384:LEU:HD21	1:F:3474:PRO:HB2	1.83	0.59
1:C:1009:ARG:HH21	1:C:1013:ARG:HH12	1.50	0.59
1:C:1961:THR:HA	1:C:1965:ARG:HB2	1.83	0.59
1:E:777:GLY:HA3	1:E:1469:LEU:HD12	1.83	0.59
1:F:1929:ASP:OD1	1:F:3612:ARG:NH2	2.35	0.59
1:F:1961:THR:HA	1:F:1965:ARG:HB2	1.83	0.59
1:A:3384:LEU:HD21	1:A:3474:PRO:HB2	1.83	0.59
1:C:262:TYR:HB2	1:C:389:ARG:HB2	1.85	0.59
1:F:2519:LEU:HD21	1:F:2554:LEU:HD21	1.83	0.59
1:F:3530:TYR:HA	1:F:3533:LEU:HD12	1.85	0.59
1:A:2463:HIS:O	1:A:2467:MET:HG2	2.03	0.59
1:A:2669:SER:HB3	1:A:2972:GLN:HG3	1.85	0.59
1:C:2145:LEU:O	1:C:2149:MET:HG2	2.03	0.59
1:C:2463:HIS:O	1:C:2467:MET:HG2	2.03	0.59
1:C:3384:LEU:HD21	1:C:3474:PRO:HB2	1.83	0.59
1:E:173:GLU:HA	1:F:3938:ARG:HH12	1.68	0.59
1:E:3530:TYR:HA	1:E:3533:LEU:HD12	1.85	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:4195:THR:O	1:E:4199:MET:HG3	2.01	0.59
1:F:113:LEU:HD12	1:F:175:VAL:HG21	1.84	0.59
1:A:1009:ARG:HH21	1:A:1013:ARG:HH12	1.50	0.59
1:A:1285:VAL:O	1:A:1287:GLN:NE2	2.36	0.59
1:C:1038:LEU:HD23	1:C:1043:LYS:HG2	1.85	0.59
1:C:1144:ARG:HH11	1:C:1191:ALA:HA	1.66	0.59
1:C:2928:LEU:HD13	1:C:2970:ILE:HG22	1.84	0.59
1:E:2435:ILE:HD12	1:E:2464:LYS:HG2	1.84	0.59
1:E:3487:LEU:HD11	1:E:3512:ILE:HA	1.85	0.59
1:F:747:HIS:ND1	1:F:748:LEU:O	2.31	0.59
1:F:3276:LEU:O	1:F:3280:LEU:HG	2.03	0.59
1:A:2661:PHE:HD2	1:A:2965:VAL:HG21	1.66	0.59
1:A:3530:TYR:HA	1:A:3533:LEU:HD12	1.85	0.59
1:A:4956:CYS:SG	1:A:4957:PHE:N	2.76	0.59
1:C:3487:LEU:HD11	1:C:3512:ILE:HA	1.85	0.59
1:E:1285:VAL:O	1:E:1287:GLN:NE2	2.36	0.59
1:F:262:TYR:HB2	1:F:389:ARG:HB2	1.85	0.59
1:A:332:ARG:NH2	1:A:338:LEU:O	2.35	0.59
1:A:1038:LEU:HD23	1:A:1043:LYS:HG2	1.85	0.59
1:A:2145:LEU:O	1:A:2149:MET:HG2	2.03	0.59
1:C:113:LEU:HD12	1:C:175:VAL:HG21	1.84	0.59
1:C:1174:MET:HB3	1:C:1190:LEU:HA	1.84	0.59
1:C:1704:TYR:O	1:C:1708:ILE:HG12	2.03	0.59
1:C:3530:TYR:HA	1:C:3533:LEU:HD12	1.85	0.59
1:C:4956:CYS:SG	1:C:4957:PHE:N	2.76	0.59
1:F:1100:ARG:HB3	1:F:1236:TYR:HA	1.84	0.59
1:F:1704:TYR:O	1:F:1708:ILE:HG12	2.03	0.59
1:A:1704:TYR:O	1:A:1708:ILE:HG12	2.03	0.58
1:C:1929:ASP:OD1	1:C:3612:ARG:NH2	2.35	0.58
1:E:2480:GLN:NE2	1:E:2537:THR:OG1	2.36	0.58
1:E:2669:SER:HB3	1:E:2972:GLN:HG3	1.85	0.58
1:F:194:LEU:HD11	1:F:201:TRP:HB3	1.85	0.58
1:F:1174:MET:HB3	1:F:1190:LEU:HA	1.84	0.58
1:F:2086:LEU:O	1:F:2090:GLN:HG2	2.03	0.58
1:C:194:LEU:HD11	1:C:201:TRP:HB3	1.85	0.58
1:C:3053:LYS:HG3	1:C:3057:ARG:HH12	1.68	0.58
1:E:1038:LEU:HD23	1:E:1043:LYS:HG2	1.85	0.58
1:F:4956:CYS:SG	1:F:4957:PHE:N	2.76	0.58
1:A:3938:ARG:HH12	1:C:173:GLU:HA	1.68	0.58
1:C:332:ARG:NH2	1:C:338:LEU:O	2.35	0.58
1:C:3276:LEU:O	1:C:3280:LEU:HG	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:3053:LYS:HG3	1:E:3057:ARG:HH12	1.68	0.58
1:F:1038:LEU:HD23	1:F:1043:LYS:HG2	1.85	0.58
1:F:3487:LEU:HD11	1:F:3512:ILE:HA	1.85	0.58
1:A:262:TYR:HB2	1:A:389:ARG:HB2	1.85	0.58
1:A:3053:LYS:HG3	1:A:3057:ARG:HH12	1.68	0.58
1:A:3487:LEU:HD11	1:A:3512:ILE:HA	1.85	0.58
1:E:113:LEU:HD12	1:E:175:VAL:HG21	1.84	0.58
1:E:1100:ARG:HB3	1:E:1236:TYR:HA	1.84	0.58
1:E:3276:LEU:O	1:E:3280:LEU:HG	2.03	0.58
1:A:2922:TYR:O	1:A:2926:GLN:HB2	2.01	0.58
1:C:2086:LEU:O	1:C:2090:GLN:HG2	2.03	0.58
1:E:2145:LEU:O	1:E:2149:MET:HG2	2.03	0.58
1:E:3508:ILE:HG22	1:E:3547:VAL:HG22	1.85	0.58
1:A:194:LEU:HD11	1:A:201:TRP:HB3	1.85	0.58
1:C:1285:VAL:O	1:C:1287:GLN:NE2	2.36	0.58
1:C:3508:ILE:HG22	1:C:3547:VAL:HG22	1.85	0.58
1:E:156:GLU:HB2	1:E:187:SER:HB3	1.84	0.58
1:E:194:LEU:HD11	1:E:201:TRP:HB3	1.85	0.58
1:F:2145:LEU:O	1:F:2149:MET:HG2	2.03	0.58
1:A:156:GLU:HB2	1:A:187:SER:HB3	1.84	0.58
1:A:2086:LEU:O	1:A:2090:GLN:HG2	2.03	0.58
1:A:2237:THR:HG22	1:A:2239:LEU:H	1.67	0.58
1:A:3508:ILE:HG22	1:A:3547:VAL:HG22	1.85	0.58
1:C:1007:TRP:HH2	2:D:113:GLU:HB2	1.69	0.58
1:F:156:GLU:HB2	1:F:187:SER:HB3	1.84	0.58
1:F:1733:GLU:HB2	1:F:1754:LEU:HD21	1.85	0.58
1:F:2435:ILE:HD12	1:F:2464:LYS:HG2	1.84	0.58
2:I:4:LEU:HB2	2:I:118:GLY:HA3	1.86	0.58
1:A:3276:LEU:O	1:A:3280:LEU:HG	2.03	0.58
1:A:4843:ILE:HD13	1:E:4813:MET:SD	2.44	0.58
1:C:1100:ARG:HB3	1:C:1236:TYR:HA	1.85	0.58
1:C:3938:ARG:HH12	1:F:173:GLU:HA	1.69	0.58
1:E:2585:GLN:HA	1:E:2588:LEU:HD12	1.86	0.58
1:E:4956:CYS:SG	1:E:4957:PHE:N	2.76	0.58
1:F:2463:HIS:O	1:F:2467:MET:HG2	2.03	0.58
2:G:4:LEU:HB2	2:G:118:GLY:HA3	1.86	0.58
1:A:1733:GLU:HB2	1:A:1754:LEU:HD21	1.85	0.58
1:C:2480:GLN:HE22	1:C:2534:PHE:HA	1.69	0.58
1:C:2669:SER:HB3	1:C:2972:GLN:HG3	1.85	0.58
1:E:2086:LEU:O	1:E:2090:GLN:HG2	2.03	0.58
1:A:1007:TRP:HH2	2:B:113:GLU:HB2	1.69	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:4:LEU:HB2	2:B:118:GLY:HA3	1.86	0.58
1:C:156:GLU:HB2	1:C:187:SER:HB3	1.84	0.58
1:C:2330:PHE:HE2	1:C:2425:LEU:HD21	1.69	0.58
1:C:2585:GLN:HA	1:C:2588:LEU:HD12	1.86	0.58
1:C:3122:LEU:HA	1:C:3126:GLN:HE21	1.69	0.58
1:E:2330:PHE:HE2	1:E:2425:LEU:HD21	1.69	0.58
1:F:2669:SER:HB3	1:F:2972:GLN:HG3	1.85	0.58
1:F:3053:LYS:HG3	1:F:3057:ARG:HH12	1.68	0.58
1:F:3389:PRO:HG3	1:F:3537:THR:HG21	1.86	0.58
1:A:1100:ARG:HB3	1:A:1236:TYR:HA	1.85	0.57
1:A:2480:GLN:NE2	1:A:2537:THR:OG1	2.37	0.57
1:A:3764:ILE:O	1:A:3769:ASN:ND2	2.31	0.57
1:C:552:SER:HA	1:C:555:LEU:HG	1.86	0.57
1:C:3831:ASP:HB2	1:C:3834:PHE:HB3	1.86	0.57
1:E:1009:ARG:HH21	1:E:1013:ARG:HH12	1.50	0.57
1:E:2480:GLN:HE22	1:E:2534:PHE:HA	1.69	0.57
1:F:1310:CYS:SG	1:F:1538:THR:N	2.76	0.57
1:F:2480:GLN:NE2	1:F:2537:THR:OG1	2.36	0.57
1:F:3122:LEU:HA	1:F:3126:GLN:HE21	1.69	0.57
1:A:3389:PRO:HG3	1:A:3537:THR:HG21	1.86	0.57
1:C:1733:GLU:HB2	1:C:1754:LEU:HD21	1.86	0.57
1:C:2480:GLN:NE2	1:C:2537:THR:OG1	2.36	0.57
1:E:262:TYR:HB2	1:E:389:ARG:HB2	1.85	0.57
1:F:2928:LEU:HD13	1:F:2970:ILE:HG22	1.84	0.57
1:A:2480:GLN:HE22	1:A:2534:PHE:HA	1.69	0.57
1:A:3292:GLY:HA3	1:A:3295:MET:CE	2.34	0.57
1:C:3339:LEU:HD22	1:C:3354:ILE:HD12	1.87	0.57
1:E:2520:CYS:SG	1:E:2564:GLN:HG2	2.45	0.57
1:F:3339:LEU:HD22	1:F:3354:ILE:HD12	1.87	0.57
1:F:3831:ASP:HB2	1:F:3834:PHE:HB3	1.86	0.57
1:A:2520:CYS:SG	1:A:2564:GLN:HG2	2.45	0.57
1:C:3292:GLY:HA3	1:C:3295:MET:CE	2.34	0.57
1:C:4813:MET:SD	1:F:4843:ILE:HD13	2.44	0.57
1:E:1704:TYR:O	1:E:1708:ILE:HG12	2.03	0.57
1:E:1733:GLU:HB2	1:E:1754:LEU:HD21	1.85	0.57
1:E:3001:GLU:HG2	1:E:3048:GLY:HA2	1.86	0.57
1:E:4843:ILE:HD13	1:F:4813:MET:SD	2.45	0.57
1:F:448:PRO:HB2	1:F:451:SER:HB3	1.87	0.57
1:F:2503:THR:OG1	1:F:2506:LEU:HB2	2.04	0.57
1:F:3272:MET:HG3	1:F:3309:VAL:HG12	1.87	0.57
1:C:515:ALA:HB2	1:C:523:GLY:HA3	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3001:GLU:HG2	1:C:3048:GLY:HA2	1.86	0.57
1:E:903:GLN:O	1:E:915:HIS:N	2.35	0.57
1:E:3122:LEU:HA	1:E:3126:GLN:HE21	1.69	0.57
1:F:2480:GLN:HE22	1:F:2534:PHE:HA	1.69	0.57
1:F:3393:GLU:HG3	1:F:3397:MET:HE2	1.86	0.57
1:F:3508:ILE:HG22	1:F:3547:VAL:HG22	1.85	0.57
1:A:173:GLU:HA	1:E:3938:ARG:HH12	1.69	0.57
1:A:515:ALA:HB2	1:A:523:GLY:HA3	1.87	0.57
1:A:1901:VAL:O	1:A:1905:MET:HG3	2.05	0.57
1:A:2585:GLN:HA	1:A:2588:LEU:HD12	1.86	0.57
1:A:3272:MET:HE2	1:A:3308:LYS:HB2	1.85	0.57
1:C:448:PRO:HB2	1:C:451:SER:HB3	1.87	0.57
1:C:3389:PRO:HG3	1:C:3537:THR:HG21	1.85	0.57
1:E:1007:TRP:HH2	2:I:113:GLU:HB2	1.69	0.57
1:E:3831:ASP:HB2	1:E:3834:PHE:HB3	1.86	0.57
1:F:2330:PHE:HE2	1:F:2425:LEU:HD21	1.69	0.57
1:A:2350:ILE:O	1:A:2354:GLU:HG2	2.05	0.57
1:A:3122:LEU:HA	1:A:3126:GLN:HE21	1.69	0.57
1:E:2463:HIS:O	1:E:2467:MET:HG2	2.03	0.57
1:F:2350:ILE:O	1:F:2354:GLU:HG2	2.05	0.57
1:A:3001:GLU:HG2	1:A:3048:GLY:HA2	1.86	0.57
1:C:2350:ILE:O	1:C:2354:GLU:HG2	2.05	0.57
1:C:2612:HIS:CE1	1:C:2620:TYR:HH	2.23	0.57
1:E:3292:GLY:HA3	1:E:3295:MET:CE	2.34	0.57
1:E:3328:LYS:O	1:E:3332:VAL:HG23	2.05	0.57
1:E:3389:PRO:HG3	1:E:3537:THR:HG21	1.86	0.57
1:F:908:ARG:HE	2:G:104:TYR:HB3	1.70	0.57
1:F:1285:VAL:O	1:F:1287:GLN:NE2	2.36	0.57
1:F:2585:GLN:HA	1:F:2588:LEU:HD12	1.86	0.57
1:F:3328:LYS:O	1:F:3332:VAL:HG23	2.05	0.57
1:C:2520:CYS:SG	1:C:2564:GLN:HG2	2.45	0.57
1:C:2592:VAL:HA	1:C:2643:LEU:HD13	1.87	0.57
1:C:3272:MET:HG3	1:C:3309:VAL:HG12	1.87	0.57
2:D:4:LEU:HB2	2:D:118:GLY:HA3	1.86	0.57
1:F:2520:CYS:SG	1:F:2564:GLN:HG2	2.45	0.57
1:A:538:ALA:O	1:A:542:ARG:HB3	2.05	0.57
1:A:552:SER:HA	1:A:555:LEU:HG	1.86	0.57
1:A:3339:LEU:HD22	1:A:3354:ILE:HD12	1.87	0.57
1:C:1098:ALA:HA	1:C:1168:MET:HB3	1.87	0.57
1:C:1913:CYS:SG	1:C:2090:GLN:NE2	2.61	0.57
1:C:2503:THR:OG1	1:C:2506:LEU:HB2	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2350:ILE:O	1:E:2354:GLU:HG2	2.05	0.57
1:F:3001:GLU:HG2	1:F:3048:GLY:HA2	1.86	0.57
1:A:448:PRO:HB2	1:A:451:SER:HB3	1.87	0.56
1:A:2385:ASN:HD21	1:A:2457:ALA:HA	1.70	0.56
1:A:2503:THR:OG1	1:A:2506:LEU:HB2	2.05	0.56
1:A:3831:ASP:HB2	1:A:3834:PHE:HB3	1.86	0.56
1:C:538:ALA:O	1:C:542:ARG:HB3	2.05	0.56
1:C:3272:MET:HE2	1:C:3308:LYS:HB2	1.85	0.56
1:E:538:ALA:O	1:E:542:ARG:HB3	2.05	0.56
1:E:1262:PRO:HG2	1:E:1265:HIS:HB2	1.87	0.56
1:E:1310:CYS:SG	1:E:1538:THR:N	2.76	0.56
1:E:2592:VAL:HA	1:E:2643:LEU:HD13	1.87	0.56
1:E:3172:THR:HB	1:E:3201:GLU:HG3	1.87	0.56
1:F:538:ALA:O	1:F:542:ARG:HB3	2.05	0.56
1:F:552:SER:HA	1:F:555:LEU:HG	1.86	0.56
1:F:2385:ASN:HD21	1:F:2457:ALA:HA	1.70	0.56
1:C:1686:LEU:HD11	1:C:1710:ILE:HD11	1.87	0.56
1:E:1246:ASP:OD1	1:E:1694:TYR:OH	2.23	0.56
1:E:3393:GLU:HG3	1:E:3397:MET:HE2	1.86	0.56
1:F:1007:TRP:HH2	2:G:113:GLU:HB2	1.68	0.56
1:F:1901:VAL:O	1:F:1905:MET:HG3	2.05	0.56
1:F:3292:GLY:HA3	1:F:3295:MET:CE	2.34	0.56
1:F:3447:GLU:HA	1:F:3450:LYS:HD2	1.87	0.56
1:A:1262:PRO:HG2	1:A:1265:HIS:HB2	1.87	0.56
1:A:1686:LEU:HD11	1:A:1710:ILE:HD11	1.87	0.56
1:A:2330:PHE:HE2	1:A:2425:LEU:HD21	1.69	0.56
1:A:3272:MET:HG3	1:A:3309:VAL:HG12	1.87	0.56
1:C:137:ARG:HA	1:C:137:ARG:HE	1.71	0.56
1:C:2166:MET:N	1:C:2166:MET:SD	2.78	0.56
1:E:552:SER:HA	1:E:555:LEU:HG	1.86	0.56
1:E:2385:ASN:HD21	1:E:2457:ALA:HA	1.70	0.56
1:E:2778:LEU:HA	1:E:2781:MET:HE3	1.86	0.56
1:E:3272:MET:HG3	1:E:3309:VAL:HG12	1.87	0.56
1:F:137:ARG:HA	1:F:137:ARG:HE	1.71	0.56
1:F:559:ILE:HD13	1:F:593:HIS:HB3	1.87	0.56
1:F:1104:GLU:HB2	1:F:1216:ASN:HB3	1.88	0.56
1:F:1120:PRO:HG3	1:F:1202:ILE:HD11	1.88	0.56
1:F:2342:LEU:O	1:F:2346:MET:HG2	2.05	0.56
1:F:4003:VAL:HG11	1:F:4113:ARG:HD2	1.88	0.56
1:F:4494:ALA:HB1	1:F:4592:LEU:HD13	1.88	0.56
1:A:2967:LEU:HD11	1:A:3028:ILE:HA	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3447:GLU:HA	1:C:3450:LYS:HD2	1.87	0.56
1:E:1120:PRO:HG3	1:E:1202:ILE:HD11	1.87	0.56
1:F:4181:GLU:HA	1:F:4184:LYS:HB2	1.88	0.56
1:A:3647:GLY:O	1:A:3658:LYS:NZ	2.37	0.56
1:C:1901:VAL:O	1:C:1905:MET:HG3	2.05	0.56
1:C:4494:ALA:HB1	1:C:4592:LEU:HD13	1.88	0.56
1:E:1901:VAL:O	1:E:1905:MET:HG3	2.05	0.56
1:E:3272:MET:HE2	1:E:3308:LYS:HB2	1.88	0.56
1:E:3339:LEU:HD22	1:E:3354:ILE:HD12	1.87	0.56
1:F:1098:ALA:HA	1:F:1168:MET:HB3	1.87	0.56
1:A:1310:CYS:SG	1:A:1538:THR:N	2.76	0.56
1:C:2278:MET:N	1:C:2278:MET:SD	2.79	0.56
1:C:3393:GLU:HA	1:C:3396:ARG:HD2	1.88	0.56
1:C:4003:VAL:HG11	1:C:4113:ARG:HD2	1.88	0.56
1:E:448:PRO:HB2	1:E:451:SER:HB3	1.87	0.56
1:E:1811:VAL:HB	1:E:1818:LEU:HD13	1.87	0.56
1:A:3172:THR:HB	1:A:3201:GLU:HG3	1.87	0.56
1:A:3920:THR:O	1:A:3924:GLN:HG2	2.06	0.56
1:C:1104:GLU:HB2	1:C:1216:ASN:HB3	1.88	0.56
1:C:1811:VAL:HB	1:C:1818:LEU:HD13	1.87	0.56
1:C:4628:GLN:OE1	1:C:4631:ARG:NH2	2.39	0.56
1:E:515:ALA:HB2	1:E:523:GLY:HA3	1.87	0.56
1:E:2641:ARG:NH1	1:E:2680:MET:HE3	2.18	0.56
1:E:3393:GLU:HA	1:E:3396:ARG:HD2	1.88	0.56
1:E:3396:ARG:O	1:E:3400:GLU:HG3	2.06	0.56
1:E:4494:ALA:HB1	1:E:4592:LEU:HD13	1.88	0.56
1:A:1727:ILE:HD11	1:A:2164:LEU:HD21	1.87	0.56
1:C:908:ARG:HE	2:D:104:TYR:HB3	1.70	0.56
1:C:1262:PRO:HG2	1:C:1265:HIS:HB2	1.87	0.56
1:C:2342:LEU:O	1:C:2346:MET:HG2	2.05	0.56
1:C:3920:THR:O	1:C:3924:GLN:HG2	2.06	0.56
1:E:137:ARG:HA	1:E:137:ARG:HE	1.71	0.56
1:E:769:ARG:HH22	1:E:816:PRO:HD3	1.71	0.56
1:E:4628:GLN:OE1	1:E:4631:ARG:NH2	2.39	0.56
1:F:1262:PRO:HG2	1:F:1265:HIS:HB2	1.87	0.56
1:A:3328:LYS:O	1:A:3332:VAL:HG23	2.05	0.56
1:A:4628:GLN:OE1	1:A:4631:ARG:NH2	2.39	0.56
1:C:4181:GLU:HA	1:C:4184:LYS:HB2	1.88	0.56
1:E:559:ILE:HD13	1:E:593:HIS:HB3	1.87	0.56
1:E:2967:LEU:HD11	1:E:3028:ILE:HA	1.87	0.56
1:E:3731:HIS:CE1	1:E:3775:LYS:HD2	2.41	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:4181:GLU:HA	1:E:4184:LYS:HB2	1.88	0.56
1:F:676:GLU:HG3	1:F:803:LEU:HB2	1.88	0.56
1:F:1641:ASP:HB3	1:F:1644:GLU:HG3	1.88	0.56
1:A:137:ARG:HA	1:A:137:ARG:HE	1.71	0.56
1:A:559:ILE:HD13	1:A:593:HIS:HB3	1.87	0.56
1:A:2592:VAL:HA	1:A:2643:LEU:HD13	1.87	0.56
1:C:2967:LEU:HD11	1:C:3028:ILE:HA	1.87	0.56
1:C:3172:THR:HB	1:C:3201:GLU:HG3	1.87	0.56
1:C:3764:ILE:O	1:C:3769:ASN:ND2	2.31	0.56
1:E:1686:LEU:HD11	1:E:1710:ILE:HD11	1.87	0.56
1:E:3764:ILE:O	1:E:3769:ASN:ND2	2.31	0.56
1:F:515:ALA:HB2	1:F:523:GLY:HA3	1.87	0.56
1:F:1727:ILE:HD11	1:F:2164:LEU:HD21	1.87	0.56
1:F:3172:THR:HB	1:F:3201:GLU:HG3	1.87	0.56
1:A:908:ARG:HE	2:B:104:TYR:HB3	1.70	0.55
1:A:1098:ALA:HA	1:A:1168:MET:HB3	1.87	0.55
1:A:1104:GLU:HB2	1:A:1216:ASN:HB3	1.88	0.55
1:A:1120:PRO:HG3	1:A:1202:ILE:HD11	1.88	0.55
1:C:1641:ASP:HB3	1:C:1644:GLU:HG3	1.88	0.55
1:C:2385:ASN:HD21	1:C:2457:ALA:HA	1.70	0.55
1:E:4003:VAL:HG11	1:E:4113:ARG:HD2	1.88	0.55
1:A:2342:LEU:O	1:A:2346:MET:HG2	2.05	0.55
1:A:4494:ALA:HB1	1:A:4592:LEU:HD13	1.88	0.55
1:C:2981:PHE:CG	1:C:2996:SER:HB3	2.42	0.55
1:C:3348:SER:O	1:C:3352:LEU:HG	2.06	0.55
1:E:676:GLU:HG3	1:E:803:LEU:HB2	1.88	0.55
1:E:2503:THR:OG1	1:E:2506:LEU:HB2	2.04	0.55
1:F:3348:SER:O	1:F:3352:LEU:HG	2.06	0.55
1:F:3731:HIS:CE1	1:F:3775:LYS:HD2	2.41	0.55
1:A:3393:GLU:HA	1:A:3396:ARG:HD2	1.88	0.55
1:A:3396:ARG:O	1:A:3400:GLU:HG3	2.06	0.55
1:A:4181:GLU:HA	1:A:4184:LYS:HB2	1.88	0.55
1:C:676:GLU:HG3	1:C:803:LEU:HB2	1.88	0.55
1:C:1120:PRO:HG3	1:C:1202:ILE:HD11	1.87	0.55
1:C:3328:LYS:O	1:C:3332:VAL:HG23	2.05	0.55
1:E:2932:VAL:HG21	1:E:3006:LEU:HD11	1.89	0.55
1:F:1686:LEU:HD11	1:F:1710:ILE:HD11	1.87	0.55
1:F:3396:ARG:O	1:F:3400:GLU:HG3	2.06	0.55
1:F:3920:THR:O	1:F:3924:GLN:HG2	2.06	0.55
1:A:989:THR:HG23	1:A:992:GLN:H	1.71	0.55
1:A:2542:SER:HB2	1:A:2877:THR:HB	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1480:ILE:H	1:C:1480:ILE:HD12	1.72	0.55
1:C:3393:GLU:O	1:C:3397:MET:HE2	2.07	0.55
1:E:891:GLU:O	1:E:895:MET:HG3	2.07	0.55
1:E:2342:LEU:O	1:E:2346:MET:HG2	2.05	0.55
1:F:1811:VAL:HB	1:F:1818:LEU:HD13	1.87	0.55
1:F:2542:SER:HB2	1:F:2877:THR:HB	1.88	0.55
1:F:3393:GLU:HA	1:F:3396:ARG:HD2	1.88	0.55
1:A:1165:MET:HB2	1:A:1174:MET:SD	2.46	0.55
1:A:4813:MET:SD	1:C:4843:ILE:HD13	2.47	0.55
1:C:559:ILE:HD13	1:C:593:HIS:HB3	1.87	0.55
1:C:891:GLU:O	1:C:895:MET:HG3	2.07	0.55
1:C:1248:THR:HG22	1:C:1602:ASN:HD21	1.72	0.55
1:E:838:ARG:HH21	1:E:1254:ARG:HD2	1.71	0.55
1:E:1098:ALA:HA	1:E:1168:MET:HB3	1.87	0.55
1:E:2278:MET:N	1:E:2278:MET:SD	2.79	0.55
1:E:2580:ARG:HD2	1:E:2581:PRO:HD2	1.89	0.55
1:E:3647:GLY:O	1:E:3658:LYS:NZ	2.37	0.55
1:E:3920:THR:O	1:E:3924:GLN:HG2	2.06	0.55
1:F:4628:GLN:OE1	1:F:4631:ARG:NH2	2.39	0.55
1:A:676:GLU:HG3	1:A:803:LEU:HB2	1.88	0.55
1:A:1641:ASP:HB3	1:A:1644:GLU:HG3	1.88	0.55
1:A:3348:SER:O	1:A:3352:LEU:HG	2.06	0.55
1:A:3731:HIS:CE1	1:A:3775:LYS:HD2	2.41	0.55
1:E:989:THR:HG23	1:E:992:GLN:H	1.71	0.55
1:E:1104:GLU:HB2	1:E:1216:ASN:HB3	1.88	0.55
1:E:1641:ASP:HB3	1:E:1644:GLU:HG3	1.88	0.55
1:F:2580:ARG:HD2	1:F:2581:PRO:HD2	1.89	0.55
1:A:769:ARG:HH22	1:A:816:PRO:HD3	1.71	0.55
1:A:838:ARG:HH21	1:A:1254:ARG:HD2	1.71	0.55
1:A:1480:ILE:H	1:A:1480:ILE:HD12	1.72	0.55
1:C:1165:MET:HB2	1:C:1174:MET:SD	2.47	0.55
1:C:1310:CYS:SG	1:C:1538:THR:N	2.76	0.55
1:C:2191:MET:HG3	1:C:2191:MET:O	2.07	0.55
1:E:1248:THR:HG22	1:E:1602:ASN:HD21	1.72	0.55
1:E:1727:ILE:HD11	1:E:2164:LEU:HD21	1.87	0.55
1:E:1843:ILE:HD13	1:E:1846:LEU:HD21	1.89	0.55
1:E:3348:SER:O	1:E:3352:LEU:HG	2.06	0.55
1:F:2317:ASN:O	1:F:2321:ARG:HG2	2.07	0.55
1:A:1811:VAL:HB	1:A:1818:LEU:HD13	1.87	0.55
1:A:1843:ILE:HD13	1:A:1846:LEU:HD21	1.89	0.55
1:C:1727:ILE:HD11	1:C:2164:LEU:HD21	1.87	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3396:ARG:O	1:C:3400:GLU:HG3	2.06	0.55
1:E:1480:ILE:HD12	1:E:1480:ILE:H	1.72	0.55
1:F:769:ARG:HH22	1:F:816:PRO:HD3	1.71	0.55
1:F:1165:MET:HB2	1:F:1174:MET:SD	2.47	0.55
1:F:1480:ILE:H	1:F:1480:ILE:HD12	1.72	0.55
1:F:2278:MET:N	1:F:2278:MET:SD	2.79	0.55
1:F:2932:VAL:HG21	1:F:3006:LEU:HD11	1.89	0.55
1:A:113:LEU:HB2	1:A:175:VAL:HB	1.89	0.55
1:A:3447:GLU:HA	1:A:3450:LYS:HD2	1.87	0.55
1:A:4003:VAL:HG11	1:A:4113:ARG:HD2	1.88	0.55
1:C:3379:ASN:HB2	1:C:3382:LYS:HD3	1.89	0.55
1:C:3731:HIS:CE1	1:C:3775:LYS:HD2	2.41	0.55
1:F:2592:VAL:HA	1:F:2643:LEU:HD13	1.87	0.55
1:A:2981:PHE:CG	1:A:2996:SER:HB3	2.42	0.55
1:E:853:PRO:HD3	1:E:1086:ARG:HG3	1.89	0.55
1:E:3472:LEU:HD23	1:E:3475:ILE:HD12	1.88	0.55
1:A:2580:ARG:HD2	1:A:2581:PRO:HD2	1.89	0.54
1:A:3379:ASN:HB2	1:A:3382:LYS:HD3	1.89	0.54
1:A:3802:LEU:HD22	1:A:3828:VAL:HG22	1.89	0.54
1:C:838:ARG:HH21	1:C:1254:ARG:HD2	1.71	0.54
1:C:853:PRO:HD3	1:C:1086:ARG:HG3	1.89	0.54
1:C:3472:LEU:HD23	1:C:3475:ILE:HD12	1.88	0.54
1:C:3802:LEU:HD22	1:C:3828:VAL:HG22	1.89	0.54
1:C:3840:ARG:HH21	1:C:3844:LEU:HD21	1.72	0.54
1:E:908:ARG:HE	2:I:104:TYR:HB3	1.71	0.54
1:E:1165:MET:HB2	1:E:1174:MET:SD	2.47	0.54
1:E:3802:LEU:HD22	1:E:3828:VAL:HG22	1.89	0.54
1:F:2967:LEU:HD11	1:F:3028:ILE:HA	1.87	0.54
1:F:3684:ASP:O	1:F:3688:MET:HG2	2.07	0.54
1:F:4046:ASP:O	1:F:4049:LYS:HG2	2.08	0.54
1:A:3472:LEU:HD23	1:A:3475:ILE:HD12	1.88	0.54
1:A:4046:ASP:O	1:A:4049:LYS:HG2	2.07	0.54
1:C:143:LEU:O	1:C:190:ARG:NE	2.34	0.54
1:C:1246:ASP:OD1	1:C:1694:TYR:OH	2.22	0.54
1:E:2981:PHE:CG	1:E:2996:SER:HB3	2.42	0.54
1:F:891:GLU:O	1:F:895:MET:HG3	2.07	0.54
1:F:2981:PHE:CG	1:F:2996:SER:HB3	2.41	0.54
1:A:891:GLU:O	1:A:895:MET:HG3	2.07	0.54
1:A:2278:MET:N	1:A:2278:MET:SD	2.79	0.54
1:A:3208:PRO:HB3	1:A:3212:LYS:HD3	1.89	0.54
1:A:3237:VAL:HA	1:A:3240:MET:CG	2.37	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3840:ARG:HH21	1:A:3844:LEU:HD21	1.72	0.54
1:A:4099:VAL:HG12	1:A:4132:LEU:HD13	1.90	0.54
1:C:113:LEU:HB2	1:C:175:VAL:HB	1.89	0.54
1:E:3447:GLU:HA	1:E:3450:LYS:HD2	1.87	0.54
1:E:3684:ASP:O	1:E:3688:MET:HG2	2.07	0.54
1:F:3802:LEU:HD22	1:F:3828:VAL:HG22	1.89	0.54
1:A:1248:THR:HG22	1:A:1602:ASN:HD21	1.72	0.54
1:A:1942:ARG:HH22	1:A:1975:LEU:HD21	1.73	0.54
1:A:2612:HIS:CE1	1:A:2620:TYR:HH	2.24	0.54
1:A:2932:VAL:HG21	1:A:3006:LEU:HD11	1.89	0.54
1:A:3684:ASP:O	1:A:3688:MET:HG2	2.07	0.54
1:C:769:ARG:HH22	1:C:816:PRO:HD3	1.71	0.54
1:C:2542:SER:HB2	1:C:2877:THR:HB	1.88	0.54
1:C:2580:ARG:HD2	1:C:2581:PRO:HD2	1.89	0.54
1:C:4099:VAL:HG12	1:C:4132:LEU:HD13	1.90	0.54
1:E:113:LEU:HB2	1:E:175:VAL:HB	1.89	0.54
1:E:2166:MET:N	1:E:2166:MET:SD	2.78	0.54
1:A:1791:LYS:O	1:A:1795:MET:HG3	2.08	0.54
1:A:2592:VAL:HG22	1:A:2643:LEU:HB2	1.89	0.54
1:A:3388:ASN:ND2	1:A:3390:GLU:OE1	2.41	0.54
1:E:143:LEU:O	1:E:190:ARG:NE	2.34	0.54
1:E:3237:VAL:HA	1:E:3240:MET:CG	2.37	0.54
1:E:4099:VAL:HG12	1:E:4132:LEU:HD13	1.89	0.54
1:F:1843:ILE:HD13	1:F:1846:LEU:HD21	1.89	0.54
1:F:3379:ASN:HB2	1:F:3382:LYS:HD3	1.89	0.54
1:A:2658:GLN:O	1:A:2662:LYS:HB2	2.08	0.54
1:C:3178:ASN:OD1	1:C:3180:TYR:N	2.40	0.54
1:E:1942:ARG:HH22	1:E:1975:LEU:HD21	1.73	0.54
1:C:2932:VAL:HG21	1:C:3006:LEU:HD11	1.89	0.54
1:C:2998:LYS:HG3	1:C:3002:MET:HE2	1.88	0.54
1:E:4046:ASP:O	1:E:4049:LYS:HG2	2.08	0.54
1:F:3472:LEU:HD23	1:F:3475:ILE:HD12	1.88	0.54
1:A:1012:ILE:HG22	1:A:1032:LEU:HG	1.90	0.54
1:A:1246:ASP:OD1	1:A:1694:TYR:OH	2.22	0.54
1:C:747:HIS:ND1	1:C:748:LEU:O	2.31	0.54
1:C:2317:ASN:O	1:C:2321:ARG:HG2	2.07	0.54
1:C:2658:GLN:O	1:C:2662:LYS:HB2	2.08	0.54
1:C:2672:ALA:O	1:C:2977:HIS:NE2	2.36	0.54
1:E:2317:ASN:O	1:E:2321:ARG:HG2	2.07	0.54
1:F:113:LEU:HB2	1:F:175:VAL:HB	1.89	0.54
1:F:3178:ASN:OD1	1:F:3180:TYR:N	2.40	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3208:PRO:HB3	1:F:3212:LYS:HD3	1.89	0.54
1:C:1942:ARG:HH22	1:C:1975:LEU:HD21	1.73	0.54
1:C:3684:ASP:O	1:C:3688:MET:HG2	2.07	0.54
1:E:3840:ARG:HH21	1:E:3844:LEU:HD21	1.72	0.54
1:F:853:PRO:HD3	1:F:1086:ARG:HG3	1.89	0.54
1:F:2592:VAL:HG22	1:F:2643:LEU:HB2	1.89	0.54
1:F:2641:ARG:NH1	1:F:2680:MET:HE3	2.18	0.54
1:A:853:PRO:HD3	1:A:1086:ARG:HG3	1.89	0.54
1:A:3178:ASN:OD1	1:A:3180:TYR:N	2.40	0.54
1:C:989:THR:HG23	1:C:992:GLN:H	1.71	0.54
1:C:2419:ARG:NE	1:C:2474:VAL:O	2.36	0.54
1:C:4046:ASP:O	1:C:4049:LYS:HG2	2.08	0.54
1:E:2612:HIS:CE1	1:E:2620:TYR:HH	2.25	0.54
1:E:3388:ASN:ND2	1:E:3390:GLU:OE1	2.41	0.54
1:F:989:THR:HG23	1:F:992:GLN:H	1.71	0.54
1:F:4099:VAL:HG12	1:F:4132:LEU:HD13	1.90	0.54
1:A:2460:CYS:HB2	1:A:2463:HIS:ND1	2.23	0.53
1:A:2673:GLY:HA2	1:A:2977:HIS:CE1	2.43	0.53
1:C:1012:ILE:HG22	1:C:1032:LEU:HG	1.90	0.53
1:C:1843:ILE:HD13	1:C:1846:LEU:HD21	1.89	0.53
1:E:2542:SER:HB2	1:E:2877:THR:HB	1.88	0.53
1:E:2658:GLN:O	1:E:2662:LYS:HB2	2.08	0.53
1:E:3475:ILE:O	1:E:3479:ILE:HG13	2.08	0.53
1:F:3475:ILE:O	1:F:3479:ILE:HG13	2.08	0.53
1:A:2166:MET:N	1:A:2166:MET:SD	2.78	0.53
1:C:2460:CYS:HB2	1:C:2463:HIS:ND1	2.23	0.53
1:C:2544:ILE:O	1:C:2548:LEU:HG	2.08	0.53
1:E:2460:CYS:HB2	1:E:2463:HIS:ND1	2.23	0.53
1:E:2673:GLY:HA2	1:E:2977:HIS:CE1	2.43	0.53
1:F:838:ARG:HH21	1:F:1254:ARG:HD2	1.71	0.53
1:F:942:THR:O	1:F:946:LEU:HG	2.08	0.53
1:F:1248:THR:HG22	1:F:1602:ASN:HD21	1.72	0.53
1:F:1891:GLY:O	1:F:1895:MET:HG3	2.09	0.53
1:F:1942:ARG:HH22	1:F:1975:LEU:HD21	1.73	0.53
1:A:2317:ASN:O	1:A:2321:ARG:HG2	2.07	0.53
1:C:1791:LYS:O	1:C:1795:MET:HG3	2.08	0.53
1:C:3388:ASN:ND2	1:C:3390:GLU:OE1	2.41	0.53
1:E:2592:VAL:HG22	1:E:2643:LEU:HB2	1.89	0.53
1:E:3208:PRO:HB3	1:E:3212:LYS:HD3	1.89	0.53
1:F:1791:LYS:O	1:F:1795:MET:HG3	2.08	0.53
1:F:2544:ILE:O	1:F:2548:LEU:HG	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2889:GLN:HG2	1:F:2893:LYS:HE3	1.91	0.53
1:F:3237:VAL:HA	1:F:3240:MET:CG	2.37	0.53
1:A:895:MET:HG2	1:A:978:PRO:HG2	1.91	0.53
1:A:2544:ILE:O	1:A:2548:LEU:HG	2.08	0.53
1:C:394:HIS:ND1	1:C:396:GLU:HG3	2.24	0.53
1:C:3208:PRO:HB3	1:C:3212:LYS:HD3	1.89	0.53
1:C:3392:GLU:O	1:C:3396:ARG:HG3	2.08	0.53
1:E:2544:ILE:O	1:E:2548:LEU:HG	2.08	0.53
1:F:1242:ASN:HB3	1:F:1808:ARG:HD2	1.90	0.53
1:F:1902:LYS:HD2	1:F:2079:LEU:HD21	1.90	0.53
1:F:2438:ALA:HA	1:F:2464:LYS:HZ1	1.73	0.53
1:F:3647:GLY:O	1:F:3658:LYS:NZ	2.37	0.53
1:F:3840:ARG:HH21	1:F:3844:LEU:HD21	1.72	0.53
1:A:2889:GLN:HG2	1:A:2893:LYS:HE3	1.91	0.53
1:C:4806:ASP:OD1	1:C:4807:ASP:N	2.42	0.53
1:E:1012:ILE:HG22	1:E:1032:LEU:HG	1.90	0.53
1:E:1902:LYS:HD2	1:E:2079:LEU:HD21	1.90	0.53
1:E:2438:ALA:HA	1:E:2464:LYS:HZ1	1.72	0.53
1:F:2658:GLN:O	1:F:2662:LYS:HB2	2.08	0.53
1:F:4806:ASP:OD1	1:F:4807:ASP:N	2.42	0.53
1:A:394:HIS:ND1	1:A:396:GLU:HG3	2.24	0.53
1:A:942:THR:O	1:A:946:LEU:HG	2.08	0.53
1:A:1242:ASN:HB3	1:A:1808:ARG:HD2	1.90	0.53
1:A:1891:GLY:O	1:A:1895:MET:HG3	2.09	0.53
1:A:2931:TYR:HB3	1:A:2962:PHE:HE1	1.74	0.53
1:E:575:LEU:O	1:E:579:LEU:HG	2.09	0.53
1:E:895:MET:HG2	1:E:978:PRO:HG2	1.91	0.53
1:E:1791:LYS:O	1:E:1795:MET:HG3	2.08	0.53
1:E:4806:ASP:OD1	1:E:4807:ASP:N	2.42	0.53
1:F:2540:HIS:O	1:F:2543:LEU:N	2.42	0.53
1:F:2778:LEU:HA	1:F:2781:MET:HE3	1.89	0.53
1:F:2931:TYR:HB3	1:F:2962:PHE:HE1	1.74	0.53
1:A:4781:THR:HG21	1:A:4812:TYR:HB2	1.91	0.53
1:C:895:MET:HG2	1:C:978:PRO:HG2	1.91	0.53
1:C:942:THR:O	1:C:946:LEU:HG	2.08	0.53
1:C:2931:TYR:HB3	1:C:2962:PHE:CE1	2.44	0.53
1:E:2191:MET:O	1:E:2191:MET:HG3	2.07	0.53
1:F:3388:ASN:ND2	1:F:3390:GLU:OE1	2.41	0.53
1:F:3392:GLU:O	1:F:3396:ARG:HG3	2.08	0.53
2:G:19:ARG:HG3	2:G:81:GLN:HE22	1.74	0.53
2:G:33:SER:HB2	2:G:98:ASP:O	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2931:TYR:HB3	1:A:2962:PHE:CE1	2.44	0.53
1:A:3392:GLU:O	1:A:3396:ARG:HG3	2.08	0.53
1:C:2673:GLY:HA2	1:C:2977:HIS:CE1	2.43	0.53
1:E:942:THR:O	1:E:946:LEU:HG	2.08	0.53
1:E:2889:GLN:HG2	1:E:2893:LYS:HE3	1.91	0.53
1:E:3379:ASN:HB2	1:E:3382:LYS:HD3	1.89	0.53
1:E:4058:THR:O	1:E:4062:THR:HG23	2.09	0.53
1:F:1012:ILE:HG22	1:F:1032:LEU:HG	1.90	0.53
1:A:2419:ARG:NE	1:A:2474:VAL:O	2.35	0.53
1:C:1891:GLY:O	1:C:1895:MET:HG3	2.09	0.53
1:C:2931:TYR:HB3	1:C:2962:PHE:HE1	1.74	0.53
1:E:1891:GLY:O	1:E:1895:MET:HG3	2.09	0.53
1:E:2925:LEU:HD23	1:E:2928:LEU:HD12	1.91	0.53
1:E:3878:LEU:HD22	1:E:3938:ARG:HG2	1.91	0.53
1:F:2460:CYS:HB2	1:F:2463:HIS:ND1	2.23	0.53
1:F:2931:TYR:HB3	1:F:2962:PHE:CE1	2.44	0.53
1:A:1609:VAL:HG13	1:A:1611:ARG:HG3	1.91	0.53
2:D:19:ARG:HG3	2:D:81:GLN:HE22	1.74	0.53
1:E:2931:TYR:HB3	1:E:2962:PHE:CE1	2.44	0.53
1:F:575:LEU:O	1:F:579:LEU:HG	2.09	0.53
1:F:1649:GLU:OE1	1:F:1649:GLU:N	2.36	0.53
1:F:2673:GLY:HA2	1:F:2977:HIS:CE1	2.43	0.53
1:F:2925:LEU:HD23	1:F:2928:LEU:HD12	1.91	0.53
2:I:19:ARG:HG3	2:I:81:GLN:HE22	1.74	0.53
1:A:3878:LEU:HD22	1:A:3938:ARG:HG2	1.91	0.52
1:C:3237:VAL:HA	1:C:3240:MET:CG	2.37	0.52
1:C:4781:THR:HG21	1:C:4812:TYR:HB2	1.91	0.52
1:C:4889:ILE:HD13	1:C:4912:HIS:HB3	1.91	0.52
1:F:394:HIS:ND1	1:F:396:GLU:HG3	2.24	0.52
1:F:3878:LEU:HD22	1:F:3938:ARG:HG2	1.91	0.52
2:I:33:SER:HB2	2:I:98:ASP:O	2.09	0.52
1:A:575:LEU:O	1:A:579:LEU:HG	2.09	0.52
1:A:1419:TYR:HE2	1:A:1563:ASN:HB3	1.74	0.52
1:A:2925:LEU:HD23	1:A:2928:LEU:HD12	1.91	0.52
1:A:3475:ILE:O	1:A:3479:ILE:HG13	2.08	0.52
1:A:4058:THR:O	1:A:4062:THR:HG23	2.09	0.52
1:C:2889:GLN:HG2	1:C:2893:LYS:HE3	1.91	0.52
1:C:2922:TYR:O	1:C:2926:GLN:CB	2.57	0.52
1:C:3878:LEU:HD22	1:C:3938:ARG:HG2	1.91	0.52
1:C:4058:THR:O	1:C:4062:THR:HG23	2.09	0.52
2:D:4:LEU:HD21	2:D:97:ALA:HB2	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:394:HIS:ND1	1:E:396:GLU:HG3	2.24	0.52
1:E:3178:ASN:OD1	1:E:3180:TYR:N	2.41	0.52
1:E:4907:HIS:CE1	1:E:4912:HIS:ND1	2.78	0.52
1:F:261:HIS:CD2	1:F:263:GLU:HG3	2.45	0.52
1:A:2540:HIS:O	1:A:2543:LEU:N	2.42	0.52
2:B:19:ARG:HG3	2:B:81:GLN:HE22	1.74	0.52
1:C:2540:HIS:O	1:C:2543:LEU:N	2.42	0.52
1:C:2592:VAL:HG22	1:C:2643:LEU:HB2	1.89	0.52
1:C:2925:LEU:HD23	1:C:2928:LEU:HD12	1.91	0.52
1:C:3475:ILE:O	1:C:3479:ILE:HG13	2.08	0.52
1:E:931:TYR:CD2	2:I:101:PRO:HG3	2.44	0.52
1:E:2931:TYR:HB3	1:E:2962:PHE:HE1	1.74	0.52
1:E:4889:ILE:HD13	1:E:4912:HIS:HB3	1.91	0.52
1:C:892:LEU:HA	1:C:895:MET:HE3	1.92	0.52
1:C:1242:ASN:HB3	1:C:1808:ARG:HD2	1.90	0.52
1:C:1902:LYS:HD2	1:C:2079:LEU:HD21	1.90	0.52
1:E:1221:VAL:HG21	1:F:3525:TRP:CG	2.45	0.52
1:E:1419:TYR:HE2	1:E:1563:ASN:HB3	1.74	0.52
1:E:3392:GLU:O	1:E:3396:ARG:HG3	2.08	0.52
1:F:1609:VAL:HG13	1:F:1611:ARG:HG3	1.92	0.52
1:A:3108:PHE:HB2	1:A:3160:ALA:HB1	1.92	0.52
1:C:1419:TYR:HE2	1:C:1563:ASN:HB3	1.74	0.52
1:C:2189:PRO:HA	1:C:2192:VAL:HG12	1.92	0.52
1:E:1955:ALA:O	1:E:1959:ARG:HG2	2.10	0.52
1:F:895:MET:HG2	1:F:978:PRO:HG2	1.91	0.52
1:A:4806:ASP:OD1	1:A:4807:ASP:N	2.42	0.52
1:A:4889:ILE:HD13	1:A:4912:HIS:HB3	1.90	0.52
1:E:1609:VAL:HG13	1:E:1611:ARG:HG3	1.91	0.52
1:E:2201:TYR:O	1:E:2205:ILE:HG23	2.10	0.52
1:E:2540:HIS:O	1:E:2543:LEU:N	2.42	0.52
1:E:4781:THR:HG21	1:E:4812:TYR:HB2	1.91	0.52
1:F:928:GLU:OE2	2:G:102:ASN:ND2	2.43	0.52
1:F:1900:PRO:O	1:F:1904:GLN:HG2	2.10	0.52
1:F:2201:TYR:O	1:F:2205:ILE:HG23	2.10	0.52
1:F:2922:TYR:O	1:F:2926:GLN:CB	2.57	0.52
1:F:3916:PHE:O	1:F:3920:THR:HG23	2.10	0.52
1:F:4058:THR:O	1:F:4062:THR:HG23	2.09	0.52
1:A:892:LEU:HA	1:A:895:MET:HE3	1.91	0.52
1:A:2191:MET:HG3	1:A:2191:MET:O	2.07	0.52
1:C:238:HIS:HB2	1:C:242:ASP:H	1.75	0.52
1:C:666:GLY:HA3	1:C:1034:PRO:HG3	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1609:VAL:HG13	1:C:1611:ARG:HG3	1.92	0.52
1:C:2997:ASN:O	1:C:3001:GLU:HG3	2.10	0.52
1:C:3525:TRP:CG	1:F:1221:VAL:HG21	2.45	0.52
1:E:3108:PHE:HB2	1:E:3160:ALA:HB1	1.92	0.52
1:F:1092:LYS:HE3	1:F:1120:PRO:HB3	1.92	0.52
1:F:1419:TYR:HE2	1:F:1563:ASN:HB3	1.74	0.52
1:C:261:HIS:CD2	1:C:263:GLU:HG3	2.44	0.52
1:C:3916:PHE:O	1:C:3920:THR:HG23	2.10	0.52
1:E:1242:ASN:HB3	1:E:1808:ARG:HD2	1.90	0.52
1:E:2067:ARG:O	1:E:2071:GLU:HG2	2.10	0.52
1:E:2922:TYR:O	1:E:2926:GLN:CB	2.57	0.52
1:E:4796:GLU:OE1	1:E:4796:GLU:N	2.37	0.52
1:F:666:GLY:HA3	1:F:1034:PRO:HG3	1.92	0.52
1:F:1246:ASP:OD1	1:F:1694:TYR:OH	2.23	0.52
1:F:4781:THR:HG21	1:F:4812:TYR:HB2	1.91	0.52
1:A:1902:LYS:HD2	1:A:2079:LEU:HD21	1.90	0.52
1:A:2201:TYR:O	1:A:2205:ILE:HG23	2.10	0.52
1:A:2922:TYR:O	1:A:2926:GLN:CB	2.57	0.52
1:A:4907:HIS:CE1	1:A:4912:HIS:ND1	2.78	0.52
1:C:575:LEU:O	1:C:579:LEU:HG	2.09	0.52
1:C:1900:PRO:O	1:C:1904:GLN:HG2	2.10	0.52
1:C:2075:GLU:OE1	1:C:2075:GLU:N	2.43	0.52
1:C:2657:GLU:HB2	1:C:2660:LEU:HB3	1.92	0.52
1:A:546:LYS:O	1:A:550:GLN:HG2	2.10	0.52
1:A:551:PHE:HE2	1:A:558:LEU:HD11	1.75	0.52
1:C:1092:LYS:HE3	1:C:1120:PRO:HB3	1.92	0.52
1:C:1471:ASP:HB2	1:C:1477:HIS:NE2	2.25	0.52
1:C:2441:MET:HE2	1:C:2506:LEU:HD11	1.92	0.52
1:E:238:HIS:HB2	1:E:242:ASP:H	1.75	0.52
1:E:2657:GLU:HB2	1:E:2660:LEU:HB3	1.92	0.52
1:E:2997:ASN:O	1:E:3001:GLU:HG3	2.10	0.52
1:F:143:LEU:O	1:F:190:ARG:NE	2.34	0.52
1:F:251:GLU:HG3	1:F:252:HIS:CE1	2.45	0.52
1:F:887:GLU:O	1:F:891:GLU:HG2	2.10	0.52
1:F:1955:ALA:O	1:F:1959:ARG:HG2	2.10	0.52
1:F:2405:MET:HE2	1:F:2407:LEU:H	1.74	0.52
1:F:4907:HIS:CE1	1:F:4912:HIS:ND1	2.78	0.52
1:A:251:GLU:HG3	1:A:252:HIS:CE1	2.45	0.51
1:A:2552:TYR:CE2	1:A:2556:LYS:HE3	2.45	0.51
2:B:4:LEU:HD21	2:B:97:ALA:HB2	1.91	0.51
1:C:2552:TYR:CE2	1:C:2556:LYS:HE3	2.45	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:551:PHE:HE2	1:E:558:LEU:HD11	1.75	0.51
1:E:3729:ARG:O	1:E:3733:ARG:NH1	2.43	0.51
1:F:2067:ARG:O	1:F:2071:GLU:HG2	2.10	0.51
2:G:4:LEU:HD21	2:G:97:ALA:HB2	1.91	0.51
1:A:261:HIS:CD2	1:A:263:GLU:HG3	2.45	0.51
1:A:1221:VAL:HG21	1:E:3525:TRP:CG	2.45	0.51
1:A:1272:ARG:NH2	1:A:1583:CYS:SG	2.83	0.51
1:A:2189:PRO:HA	1:A:2192:VAL:HG12	1.91	0.51
1:A:2438:ALA:HA	1:A:2464:LYS:HZ1	1.74	0.51
1:A:2997:ASN:O	1:A:3001:GLU:HG3	2.10	0.51
1:A:3273:ASN:ND2	1:A:3310:LYS:HB2	2.23	0.51
2:B:33:SER:HB2	2:B:98:ASP:O	2.09	0.51
1:C:251:GLU:HG3	1:C:252:HIS:CE1	2.45	0.51
1:C:546:LYS:O	1:C:550:GLN:HG2	2.10	0.51
1:E:3916:PHE:O	1:E:3920:THR:HG23	2.10	0.51
1:F:2997:ASN:O	1:F:3001:GLU:HG3	2.10	0.51
1:A:238:HIS:HB2	1:A:242:ASP:H	1.75	0.51
1:E:666:GLY:HA3	1:E:1034:PRO:HG3	1.92	0.51
1:E:1257:GLN:HG2	1:E:1451:HIS:HE1	1.76	0.51
1:E:1900:PRO:O	1:E:1904:GLN:HG2	2.10	0.51
1:F:2657:GLU:HB2	1:F:2660:LEU:HB3	1.92	0.51
1:F:3273:ASN:ND2	1:F:3310:LYS:HB2	2.23	0.51
1:F:3280:LEU:HD12	1:F:3317:HIS:HB2	1.93	0.51
1:A:1959:ARG:HA	1:A:1962:ARG:HB3	1.92	0.51
1:A:2067:ARG:O	1:A:2071:GLU:HG2	2.10	0.51
1:A:2657:GLU:HB2	1:A:2660:LEU:HB3	1.92	0.51
1:C:887:GLU:O	1:C:891:GLU:HG2	2.10	0.51
1:C:2527:LEU:HA	1:C:2531:ALA:HB2	1.93	0.51
1:C:3636:GLU:HG2	1:C:3696:LYS:HB2	1.93	0.51
1:E:601:LEU:HB2	1:E:610:VAL:HG11	1.93	0.51
1:E:2189:PRO:HA	1:E:2192:VAL:HG12	1.91	0.51
1:E:2441:MET:HE2	1:E:2506:LEU:HD11	1.92	0.51
1:E:4156:ARG:O	1:E:4160:GLU:HG2	2.10	0.51
1:F:238:HIS:HB2	1:F:242:ASP:H	1.75	0.51
1:F:1959:ARG:HA	1:F:1962:ARG:HB3	1.92	0.51
1:A:666:GLY:HA3	1:A:1034:PRO:HG3	1.92	0.51
1:A:1052:GLU:O	1:A:1056:THR:HG23	2.11	0.51
1:A:1092:LYS:HE3	1:A:1120:PRO:HB3	1.92	0.51
1:A:1955:ALA:O	1:A:1959:ARG:HG2	2.10	0.51
1:C:903:GLN:O	1:C:915:HIS:N	2.35	0.51
1:E:244:CYS:SG	1:E:273:SER:HB2	2.51	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1471:ASP:HB2	1:E:1477:HIS:NE2	2.25	0.51
1:E:1636:GLU:HB2	1:E:1638:ARG:HG2	1.93	0.51
1:E:2405:MET:HE2	1:E:2407:LEU:H	1.75	0.51
1:E:3031:CYS:HA	1:E:3034:ILE:HG22	1.93	0.51
1:E:3280:LEU:HD12	1:E:3317:HIS:HB2	1.93	0.51
1:F:1636:GLU:HB2	1:F:1638:ARG:HG2	1.93	0.51
1:F:3031:CYS:HA	1:F:3034:ILE:HG22	1.93	0.51
1:F:3729:ARG:O	1:F:3733:ARG:NH1	2.43	0.51
1:A:887:GLU:O	1:A:891:GLU:HG2	2.10	0.51
1:A:1900:PRO:O	1:A:1904:GLN:HG2	2.10	0.51
1:A:3293:ALA:HA	1:A:3363:ARG:HD2	1.92	0.51
1:C:1955:ALA:O	1:C:1959:ARG:HG2	2.10	0.51
1:C:3108:PHE:HB2	1:C:3160:ALA:HB1	1.92	0.51
1:C:3280:LEU:HD12	1:C:3317:HIS:HB2	1.93	0.51
2:D:33:SER:HB2	2:D:98:ASP:O	2.09	0.51
1:E:1814:THR:HB	1:E:1817:PHE:HD2	1.76	0.51
1:E:2075:GLU:OE1	1:E:2075:GLU:N	2.43	0.51
1:E:2352:ILE:HA	1:E:2358:ARG:HB3	1.92	0.51
1:F:3293:ALA:HA	1:F:3363:ARG:HD2	1.92	0.51
1:A:2075:GLU:OE1	1:A:2075:GLU:N	2.43	0.51
1:C:928:GLU:OE2	2:D:102:ASN:ND2	2.44	0.51
1:C:1959:ARG:HA	1:C:1962:ARG:HB3	1.92	0.51
1:C:2201:TYR:O	1:C:2205:ILE:HG23	2.10	0.51
1:C:2641:ARG:NH1	1:C:2680:MET:HE3	2.19	0.51
1:E:261:HIS:CD2	1:E:263:GLU:HG3	2.45	0.51
1:E:2382:HIS:HB2	1:E:2385:ASN:HB2	1.92	0.51
1:E:3273:ASN:ND2	1:E:3310:LYS:HB2	2.24	0.51
1:F:551:PHE:HE2	1:F:558:LEU:HD11	1.75	0.51
1:F:892:LEU:HD23	1:F:895:MET:HE1	1.91	0.51
1:F:1269:GLU:OE2	1:F:1293:GLN:NE2	2.30	0.51
1:F:1471:ASP:HB2	1:F:1477:HIS:NE2	2.25	0.51
1:F:1814:THR:HB	1:F:1817:PHE:HD2	1.76	0.51
1:A:3805:ASN:O	1:A:3809:ARG:HG3	2.11	0.51
1:A:3916:PHE:O	1:A:3920:THR:HG23	2.10	0.51
1:C:2065:MET:HE1	1:C:2086:LEU:HD23	1.93	0.51
1:C:4907:HIS:CE1	1:C:4912:HIS:ND1	2.78	0.51
1:E:546:LYS:O	1:E:550:GLN:HG2	2.10	0.51
1:F:2552:TYR:CE2	1:F:2556:LYS:HE3	2.45	0.51
2:I:4:LEU:HD21	2:I:97:ALA:HB2	1.91	0.51
1:A:3729:ARG:O	1:A:3733:ARG:NH1	2.43	0.51
1:A:4048:HIS:ND1	1:A:4066:LEU:HD11	2.26	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4156:ARG:O	1:A:4160:GLU:HG2	2.10	0.51
1:C:244:CYS:SG	1:C:273:SER:HB2	2.51	0.51
1:C:2067:ARG:O	1:C:2071:GLU:HG2	2.10	0.51
1:E:251:GLU:HG3	1:E:252:HIS:CE1	2.45	0.51
1:E:1052:GLU:O	1:E:1056:THR:HG23	2.11	0.51
1:E:1092:LYS:HE3	1:E:1120:PRO:HB3	1.92	0.51
1:F:546:LYS:O	1:F:550:GLN:HG2	2.10	0.51
1:F:601:LEU:HB2	1:F:610:VAL:HG11	1.93	0.51
1:A:244:CYS:SG	1:A:273:SER:HB2	2.51	0.51
1:A:928:GLU:OE2	2:B:102:ASN:ND2	2.44	0.51
1:A:3393:GLU:HG3	1:A:3397:MET:HE2	1.92	0.51
1:A:3636:GLU:HG2	1:A:3696:LYS:HB2	1.93	0.51
1:C:737:ILE:HB	1:C:1482:ARG:HH21	1.76	0.51
1:C:2352:ILE:HA	1:C:2358:ARG:HB3	1.92	0.51
1:C:3159:ALA:HB2	1:C:3240:MET:HE2	1.93	0.51
1:C:3273:ASN:ND2	1:C:3310:LYS:HB2	2.24	0.51
1:C:3729:ARG:O	1:C:3733:ARG:NH1	2.43	0.51
1:C:4796:GLU:OE1	1:C:4796:GLU:N	2.37	0.51
1:E:2527:LEU:HA	1:E:2531:ALA:HB2	1.93	0.51
1:F:244:CYS:SG	1:F:273:SER:HB2	2.51	0.51
1:F:1257:GLN:HG2	1:F:1451:HIS:HE1	1.76	0.51
1:F:2189:PRO:HA	1:F:2192:VAL:HG12	1.91	0.51
1:F:2382:HIS:HB2	1:F:2385:ASN:HB2	1.92	0.51
1:F:3108:PHE:HB2	1:F:3160:ALA:HB1	1.92	0.51
1:F:4889:ILE:HD13	1:F:4912:HIS:HB3	1.90	0.51
1:A:3393:GLU:O	1:A:3397:MET:HE2	2.10	0.50
1:C:551:PHE:HE2	1:C:558:LEU:HD11	1.75	0.50
1:C:3155:GLY:O	1:C:3240:MET:HE1	2.12	0.50
1:F:2075:GLU:OE1	1:F:2075:GLU:N	2.43	0.50
1:F:4156:ARG:O	1:F:4160:GLU:HG2	2.10	0.50
1:A:1257:GLN:HG2	1:A:1451:HIS:HE1	1.76	0.50
1:A:1269:GLU:OE2	1:A:1293:GLN:NE2	2.30	0.50
1:A:1471:ASP:HB2	1:A:1477:HIS:NE2	2.25	0.50
1:A:1636:GLU:HB2	1:A:1638:ARG:HG2	1.93	0.50
1:A:2527:LEU:HA	1:A:2531:ALA:HB2	1.93	0.50
1:A:3031:CYS:HA	1:A:3034:ILE:HG22	1.93	0.50
1:A:3280:LEU:HD12	1:A:3317:HIS:HB2	1.93	0.50
1:A:3484:ASP:O	1:A:3488:ILE:HG13	2.11	0.50
1:C:2382:HIS:HB2	1:C:2385:ASN:HB2	1.92	0.50
1:C:3647:GLY:O	1:C:3658:LYS:NZ	2.37	0.50
1:E:737:ILE:HB	1:E:1482:ARG:HH21	1.76	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1269:GLU:OE2	1:E:1293:GLN:NE2	2.30	0.50
1:F:263:GLU:OE2	1:F:388:GLN:NE2	2.40	0.50
1:F:737:ILE:HB	1:F:1482:ARG:HH21	1.76	0.50
1:F:3166:PRO:HD2	1:F:3167:ILE:HD12	1.93	0.50
1:F:4045:ARG:HA	1:F:4045:ARG:HE	1.76	0.50
1:A:4045:ARG:HA	1:A:4045:ARG:HE	1.76	0.50
1:C:1257:GLN:HG2	1:C:1451:HIS:HE1	1.76	0.50
1:C:1636:GLU:HB2	1:C:1638:ARG:HG2	1.93	0.50
1:C:3031:CYS:HA	1:C:3034:ILE:HG22	1.93	0.50
1:C:3484:ASP:O	1:C:3488:ILE:HG13	2.11	0.50
1:C:4156:ARG:O	1:C:4160:GLU:HG2	2.10	0.50
1:E:739:ARG:HH12	1:E:1480:ILE:HG21	1.77	0.50
1:E:2672:ALA:HB3	1:E:2972:GLN:HE22	1.77	0.50
1:F:4796:GLU:OE1	1:F:4796:GLU:N	2.37	0.50
1:A:2352:ILE:HA	1:A:2358:ARG:HB3	1.92	0.50
1:C:2101:LEU:O	1:C:2104:THR:HG22	2.12	0.50
1:C:2482:PHE:CZ	1:C:2486:LEU:HD11	2.47	0.50
1:C:2672:ALA:HB3	1:C:2972:GLN:HE22	1.77	0.50
1:E:1959:ARG:HA	1:E:1962:ARG:HB3	1.92	0.50
1:F:1128:LEU:HG	1:F:1136:ALA:HB2	1.93	0.50
1:F:2672:ALA:HB3	1:F:2972:GLN:HE22	1.77	0.50
1:F:3155:GLY:O	1:F:3240:MET:HE1	2.12	0.50
1:F:4111:ASP:O	1:F:4115:GLN:HG2	2.12	0.50
1:A:1446:ILE:HG22	1:A:1485:CYS:HA	1.94	0.50
1:C:1084:ARG:HG3	1:C:1084:ARG:O	2.12	0.50
1:C:2438:ALA:HA	1:C:2464:LYS:HZ1	1.75	0.50
1:E:887:GLU:O	1:E:891:GLU:HG2	2.10	0.50
1:E:3636:GLU:HG2	1:E:3696:LYS:HB2	1.93	0.50
1:F:1052:GLU:O	1:F:1056:THR:HG23	2.11	0.50
1:F:2482:PHE:CZ	1:F:2486:LEU:HD11	2.47	0.50
1:F:3380:ARG:HH21	1:F:3474:PRO:HG2	1.77	0.50
1:F:4048:HIS:ND1	1:F:4066:LEU:HD11	2.26	0.50
1:A:253:GLY:O	1:A:257:ARG:HG3	2.12	0.50
1:A:2101:LEU:O	1:A:2104:THR:HG22	2.12	0.50
1:C:3178:ASN:O	1:C:3184:ASN:ND2	2.45	0.50
1:C:4045:ARG:HA	1:C:4045:ARG:HE	1.76	0.50
1:C:4048:HIS:ND1	1:C:4066:LEU:HD11	2.26	0.50
1:E:656:ARG:HG2	1:E:837:SER:HA	1.93	0.50
1:E:3380:ARG:HH21	1:E:3474:PRO:HG2	1.77	0.50
1:E:4657:GLY:HA3	1:E:4662:ARG:HG2	1.94	0.50
1:F:656:ARG:HG2	1:F:837:SER:HA	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2527:LEU:HA	1:F:2531:ALA:HB2	1.93	0.50
1:F:3522:ALA:HA	1:F:3525:TRP:CD1	2.47	0.50
1:C:3166:PRO:HD2	1:C:3167:ILE:HD12	1.93	0.50
1:C:3293:ALA:HA	1:C:3363:ARG:HD2	1.93	0.50
1:E:1939:GLN:HA	1:E:1942:ARG:HG2	1.94	0.50
1:E:1988:PRO:HB2	1:E:1991:ILE:HG12	1.93	0.50
1:E:2552:TYR:CE2	1:E:2556:LYS:HE3	2.45	0.50
1:E:3155:GLY:O	1:E:3240:MET:HE1	2.12	0.50
1:E:3314:LEU:HD12	1:E:3315:LYS:N	2.27	0.50
1:E:3805:ASN:O	1:E:3809:ARG:HG3	2.11	0.50
1:E:4111:ASP:O	1:E:4115:GLN:HG2	2.12	0.50
1:F:739:ARG:HH12	1:F:1480:ILE:HG21	1.77	0.50
1:F:3178:ASN:O	1:F:3184:ASN:ND2	2.45	0.50
1:F:3636:GLU:HG2	1:F:3696:LYS:HB2	1.93	0.50
1:A:601:LEU:HB2	1:A:610:VAL:HG11	1.93	0.50
1:A:737:ILE:HB	1:A:1482:ARG:HH21	1.76	0.50
1:A:1814:THR:HB	1:A:1817:PHE:HD2	1.76	0.50
1:A:3525:TRP:CG	1:C:1221:VAL:HG21	2.46	0.50
1:C:253:GLY:O	1:C:257:ARG:HG3	2.12	0.50
1:C:1052:GLU:O	1:C:1056:THR:HG23	2.11	0.50
1:C:1446:ILE:HG22	1:C:1485:CYS:HA	1.93	0.50
1:C:2480:GLN:HG2	1:C:2537:THR:HG21	1.94	0.50
1:C:3380:ARG:HH21	1:C:3474:PRO:HG2	1.77	0.50
1:C:3522:ALA:HA	1:C:3525:TRP:CD1	2.47	0.50
1:C:3805:ASN:O	1:C:3809:ARG:HG3	2.11	0.50
1:C:4647:PHE:HD1	1:C:4650:ARG:HD3	1.76	0.50
1:E:892:LEU:HA	1:E:895:MET:HE3	1.93	0.50
1:E:1729:PRO:HD3	1:E:1758:LEU:HD22	1.94	0.50
1:E:3484:ASP:O	1:E:3488:ILE:HG13	2.11	0.50
1:E:4635:ASN:OD1	1:E:4703:LYS:NZ	2.38	0.50
1:F:2330:PHE:CE2	1:F:2425:LEU:HD21	2.47	0.50
1:F:2352:ILE:HA	1:F:2358:ARG:HB3	1.92	0.50
1:F:4116:THR:O	1:F:4119:GLU:HG2	2.12	0.50
1:F:4478:PHE:HA	1:F:4481:LYS:HE2	1.94	0.50
1:A:1113:MET:HB2	1:A:1156:TRP:HZ2	1.77	0.50
1:A:1128:LEU:HG	1:A:1136:ALA:HB2	1.93	0.50
1:A:1666:CYS:HB3	1:A:1677:LEU:HD12	1.94	0.50
1:C:2330:PHE:CE2	1:C:2425:LEU:HD21	2.47	0.50
1:C:3018:ILE:HG21	1:C:3095:TYR:HB2	1.94	0.50
1:E:449:ILE:HG23	1:E:529:ILE:HD11	1.94	0.50
1:E:3178:ASN:O	1:E:3184:ASN:ND2	2.45	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3050:ASP:HA	1:F:3053:LYS:HG2	1.94	0.50
1:A:557:TRP:CE3	1:A:558:LEU:HD23	2.47	0.49
1:A:924:LEU:HD23	1:A:929:ARG:HA	1.94	0.49
1:A:1979:LYS:HE2	1:A:3627:TRP:CD1	2.47	0.49
1:C:1814:THR:HB	1:C:1817:PHE:HD2	1.76	0.49
1:E:1113:MET:HB2	1:E:1156:TRP:HZ2	1.77	0.49
1:E:3293:ALA:HA	1:E:3363:ARG:HD2	1.92	0.49
1:F:2480:GLN:HG2	1:F:2537:THR:HG21	1.94	0.49
1:F:4657:GLY:HA3	1:F:4662:ARG:HG2	1.94	0.49
1:A:449:ILE:HG23	1:A:529:ILE:HD11	1.94	0.49
1:A:739:ARG:HH12	1:A:1480:ILE:HG21	1.77	0.49
1:A:3050:ASP:HA	1:A:3053:LYS:HG2	1.94	0.49
1:A:3155:GLY:O	1:A:3240:MET:HE1	2.12	0.49
1:A:3314:LEU:HD12	1:A:3315:LYS:N	2.27	0.49
1:A:3380:ARG:HH21	1:A:3474:PRO:HG2	1.77	0.49
1:A:4478:PHE:HA	1:A:4481:LYS:HE2	1.94	0.49
1:C:449:ILE:HG23	1:C:529:ILE:HD11	1.94	0.49
1:C:601:LEU:HB2	1:C:610:VAL:HG11	1.93	0.49
1:C:1649:GLU:OE1	1:C:1649:GLU:N	2.36	0.49
1:C:2999:GLU:HA	1:C:3002:MET:SD	2.52	0.49
1:C:3050:ASP:HA	1:C:3053:LYS:HG2	1.94	0.49
1:E:303:GLY:HA2	1:E:420:ARG:HH11	1.77	0.49
1:F:1666:CYS:HB3	1:F:1677:LEU:HD12	1.94	0.49
1:F:3314:LEU:HD12	1:F:3315:LYS:N	2.27	0.49
1:F:3805:ASN:O	1:F:3809:ARG:HG3	2.11	0.49
1:A:1553:VAL:HG23	1:A:1554:PHE:H	1.78	0.49
1:A:1988:PRO:HB2	1:A:1991:ILE:HG12	1.93	0.49
1:C:1666:CYS:HB3	1:C:1677:LEU:HD12	1.94	0.49
1:C:3314:LEU:HD12	1:C:3315:LYS:N	2.27	0.49
1:C:4478:PHE:HA	1:C:4481:LYS:HE2	1.94	0.49
1:E:557:TRP:CE3	1:E:558:LEU:HD23	2.47	0.49
1:E:2643:LEU:O	1:E:2647:ILE:HG13	2.12	0.49
1:E:2999:GLU:HA	1:E:3002:MET:SD	2.53	0.49
1:E:3050:ASP:HA	1:E:3053:LYS:HG2	1.94	0.49
1:E:4048:HIS:ND1	1:E:4066:LEU:HD11	2.26	0.49
1:E:4647:PHE:HD1	1:E:4650:ARG:HD3	1.76	0.49
1:F:1988:PRO:HB2	1:F:1991:ILE:HG12	1.93	0.49
1:F:2419:ARG:NE	1:F:2474:VAL:O	2.35	0.49
1:F:2643:LEU:O	1:F:2647:ILE:HG13	2.12	0.49
1:F:3802:LEU:HB2	1:F:3883:SER:HB2	1.95	0.49
1:A:2330:PHE:CE2	1:A:2425:LEU:HD21	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2382:HIS:HB2	1:A:2385:ASN:HB2	1.92	0.49
1:A:2482:PHE:CZ	1:A:2486:LEU:HD11	2.47	0.49
1:A:3905:PHE:O	1:A:3909:ILE:HG12	2.12	0.49
1:A:4116:THR:O	1:A:4119:GLU:HG2	2.12	0.49
1:A:4647:PHE:HD1	1:A:4650:ARG:HD3	1.76	0.49
1:C:303:GLY:HA2	1:C:420:ARG:HH11	1.77	0.49
1:C:924:LEU:HD23	1:C:929:ARG:HA	1.94	0.49
1:C:1939:GLN:HA	1:C:1942:ARG:HG2	1.94	0.49
1:E:2482:PHE:CZ	1:E:2486:LEU:HD11	2.47	0.49
1:E:2672:ALA:O	1:E:2977:HIS:NE2	2.36	0.49
1:F:557:TRP:CE3	1:F:558:LEU:HD23	2.47	0.49
1:F:924:LEU:HD23	1:F:929:ARG:HA	1.94	0.49
1:F:3484:ASP:O	1:F:3488:ILE:HG13	2.11	0.49
1:A:503:ASP:O	1:A:507:VAL:HG23	2.13	0.49
1:A:2999:GLU:HA	1:A:3002:MET:SD	2.53	0.49
1:A:4057:TYR:HB3	1:A:4061:GLU:HB2	1.95	0.49
1:C:2503:THR:O	1:C:2507:SER:N	2.35	0.49
1:C:3646:PRO:HB2	1:C:3658:LYS:HZ2	1.78	0.49
1:E:253:GLY:O	1:E:257:ARG:HG3	2.12	0.49
1:E:892:LEU:HD23	1:E:895:MET:HE1	1.93	0.49
1:E:1128:LEU:HG	1:E:1136:ALA:HB2	1.93	0.49
1:E:1553:VAL:HG23	1:E:1554:PHE:H	1.78	0.49
1:E:3392:GLU:OE1	1:E:3537:THR:OG1	2.31	0.49
1:E:3522:ALA:HA	1:E:3525:TRP:CD1	2.47	0.49
1:F:303:GLY:HA2	1:F:420:ARG:HH11	1.77	0.49
1:F:449:ILE:HG23	1:F:529:ILE:HD11	1.94	0.49
1:F:1729:PRO:HD3	1:F:1758:LEU:HD22	1.94	0.49
1:F:3272:MET:HE2	1:F:3308:LYS:HB2	1.93	0.49
1:F:4636:THR:OG1	1:F:4637:GLN:N	2.46	0.49
1:A:656:ARG:HG2	1:A:837:SER:HA	1.94	0.49
1:A:894:VAL:HA	1:A:897:LYS:HB2	1.94	0.49
1:A:2643:LEU:O	1:A:2647:ILE:HG13	2.12	0.49
1:A:3324:LYS:HA	1:A:3327:LYS:HD2	1.95	0.49
1:C:557:TRP:CE3	1:C:558:LEU:HD23	2.47	0.49
1:C:4116:THR:O	1:C:4119:GLU:HG2	2.12	0.49
1:E:1609:VAL:HG23	1:E:1620:VAL:HG22	1.95	0.49
1:E:1666:CYS:HB3	1:E:1677:LEU:HD12	1.94	0.49
1:F:563:GLU:OE1	1:F:563:GLU:N	2.39	0.49
1:F:2166:MET:N	1:F:2166:MET:SD	2.78	0.49
1:F:2191:MET:O	1:F:2191:MET:HG3	2.07	0.49
1:F:3018:ILE:HG21	1:F:3095:TYR:HB2	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3935:ALA:O	1:F:3940:TRP:NE1	2.44	0.49
1:A:1084:ARG:HG3	1:A:1084:ARG:O	2.11	0.49
1:A:4796:GLU:OE1	1:A:4796:GLU:N	2.37	0.49
1:C:739:ARG:HH12	1:C:1480:ILE:HG21	1.77	0.49
1:C:4111:ASP:O	1:C:4115:GLN:HG2	2.12	0.49
1:C:4112:THR:O	1:C:4116:THR:HG23	2.13	0.49
1:C:4195:THR:HB	1:C:4918:LEU:HD11	1.95	0.49
1:E:2998:LYS:O	1:E:3002:MET:HG3	2.13	0.49
1:E:4636:THR:OG1	1:E:4637:GLN:N	2.46	0.49
1:F:253:GLY:O	1:F:257:ARG:HG3	2.12	0.49
1:F:894:VAL:HA	1:F:897:LYS:HB2	1.94	0.49
1:F:925:PRO:HG2	1:F:928:GLU:HB2	1.94	0.49
1:F:1084:ARG:HG3	1:F:1084:ARG:O	2.12	0.49
1:F:1979:LYS:HE2	1:F:3627:TRP:CD1	2.47	0.49
1:F:4112:THR:O	1:F:4116:THR:HG23	2.13	0.49
1:A:670:TYR:HD2	1:A:672:LYS:HB2	1.78	0.49
1:A:846:TYR:CZ	1:A:1219:LYS:HB2	2.48	0.49
1:A:1939:GLN:HA	1:A:1942:ARG:HG2	1.94	0.49
1:A:2569:GLU:O	1:A:2573:LEU:HG	2.13	0.49
1:A:3041:ALA:HB1	1:A:3120:LEU:HB2	1.94	0.49
1:C:656:ARG:HG2	1:C:837:SER:HA	1.94	0.49
1:C:892:LEU:HD23	1:C:895:MET:HE1	1.94	0.49
1:C:1553:VAL:HG23	1:C:1554:PHE:H	1.78	0.49
1:C:2569:GLU:O	1:C:2573:LEU:HG	2.13	0.49
1:C:3905:PHE:O	1:C:3909:ILE:HG12	2.12	0.49
1:C:4057:TYR:HB3	1:C:4061:GLU:HB2	1.95	0.49
1:E:180:ASP:HB3	1:E:211:LEU:HD12	1.94	0.49
1:E:503:ASP:O	1:E:507:VAL:HG23	2.13	0.49
1:E:925:PRO:HG2	1:E:928:GLU:HB2	1.94	0.49
1:E:1979:LYS:HE2	1:E:3627:TRP:CD1	2.47	0.49
1:E:3802:LEU:HB2	1:E:3883:SER:HB2	1.95	0.49
1:E:4057:TYR:HB3	1:E:4061:GLU:HB2	1.95	0.49
1:F:816:PRO:HG2	1:F:819:TYR:CG	2.48	0.49
1:F:931:TYR:CD2	2:G:101:PRO:HG3	2.47	0.49
1:F:1446:ILE:HG22	1:F:1485:CYS:HA	1.94	0.49
1:F:1706:LEU:O	1:F:1710:ILE:HG12	2.13	0.49
1:F:2778:LEU:HD23	1:F:2781:MET:HE3	1.95	0.49
1:F:3905:PHE:O	1:F:3909:ILE:HG12	2.12	0.49
1:F:4647:PHE:HD1	1:F:4650:ARG:HD3	1.76	0.49
1:A:180:ASP:HB3	1:A:211:LEU:HD12	1.94	0.49
1:A:2480:GLN:HG2	1:A:2537:THR:HG21	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2839:MET:HG3	1:A:2892:PHE:HZ	1.78	0.49
1:A:2998:LYS:O	1:A:3002:MET:HG3	2.13	0.49
1:A:3018:ILE:HG21	1:A:3095:TYR:HB2	1.94	0.49
1:A:3159:ALA:HB2	1:A:3240:MET:HE2	1.94	0.49
1:A:3802:LEU:HB2	1:A:3883:SER:HB2	1.95	0.49
1:A:4111:ASP:O	1:A:4115:GLN:HG2	2.12	0.49
1:C:1128:LEU:HG	1:C:1136:ALA:HB2	1.93	0.49
1:C:1190:LEU:HD23	1:C:1190:LEU:H	1.78	0.49
1:C:1979:LYS:HE2	1:C:3627:TRP:CD1	2.47	0.49
1:C:3397:MET:O	1:C:3401:VAL:HG23	2.13	0.49
1:C:3935:ALA:O	1:C:3940:TRP:NE1	2.44	0.49
1:C:4657:GLY:HA3	1:C:4662:ARG:HG2	1.94	0.49
1:E:1681:VAL:HG21	1:E:1706:LEU:HD21	1.95	0.49
1:E:2065:MET:HE1	1:E:2086:LEU:HD23	1.95	0.49
1:E:2503:THR:O	1:E:2507:SER:N	2.35	0.49
1:E:4045:ARG:HA	1:E:4045:ARG:HE	1.76	0.49
1:F:1553:VAL:HG23	1:F:1554:PHE:H	1.78	0.49
1:F:2569:GLU:O	1:F:2573:LEU:HG	2.13	0.49
1:F:2999:GLU:HA	1:F:3002:MET:SD	2.53	0.49
1:F:3393:GLU:O	1:F:3397:MET:HE2	2.13	0.49
1:A:1649:GLU:OE1	1:A:1649:GLU:N	2.36	0.49
1:A:2383:MET:O	1:A:2387:ILE:HG12	2.13	0.49
1:A:3178:ASN:O	1:A:3184:ASN:ND2	2.45	0.49
1:A:4664:ARG:HD2	1:A:4668:LEU:HD23	1.95	0.49
1:C:846:TYR:CZ	1:C:1219:LYS:HB2	2.48	0.49
1:C:1706:LEU:O	1:C:1710:ILE:HG12	2.13	0.49
1:E:816:PRO:HG2	1:E:819:TYR:CG	2.48	0.49
1:E:846:TYR:CZ	1:E:1219:LYS:HB2	2.48	0.49
1:E:924:LEU:HD23	1:E:929:ARG:HA	1.94	0.49
1:E:1272:ARG:NH2	1:E:1583:CYS:SG	2.83	0.49
1:E:1446:ILE:HG22	1:E:1485:CYS:HA	1.94	0.49
1:F:1113:MET:SD	1:F:1211:GLN:HB3	2.53	0.49
1:F:1272:ARG:NH2	1:F:1583:CYS:SG	2.83	0.49
1:F:2101:LEU:O	1:F:2104:THR:HG22	2.12	0.49
1:F:3239:PRO:HB3	1:F:3301:PHE:CG	2.48	0.49
1:F:4664:ARG:HD2	1:F:4668:LEU:HD23	1.95	0.49
1:A:4195:THR:HB	1:A:4918:LEU:HD11	1.95	0.48
1:A:4657:GLY:HA3	1:A:4662:ARG:HG2	1.94	0.48
1:C:816:PRO:HG2	1:C:819:TYR:CG	2.48	0.48
1:C:1113:MET:HB2	1:C:1156:TRP:HZ2	1.77	0.48
1:C:1729:PRO:HD3	1:C:1758:LEU:HD22	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1988:PRO:HB2	1:C:1991:ILE:HG12	1.93	0.48
1:C:3071:MET:HE3	1:C:3135:SER:HA	1.95	0.48
1:E:2419:ARG:NE	1:E:2474:VAL:O	2.35	0.48
1:E:4043:SER:HA	1:E:4076:THR:HA	1.95	0.48
1:E:4478:PHE:HA	1:E:4481:LYS:HE2	1.94	0.48
1:F:180:ASP:HB3	1:F:211:LEU:HD12	1.94	0.48
1:F:503:ASP:O	1:F:507:VAL:HG23	2.13	0.48
1:F:846:TYR:CZ	1:F:1219:LYS:HB2	2.48	0.48
1:F:892:LEU:HA	1:F:895:MET:HE3	1.95	0.48
1:F:1113:MET:HB2	1:F:1156:TRP:HZ2	1.77	0.48
1:F:1681:VAL:HG21	1:F:1706:LEU:HD21	1.95	0.48
1:F:2065:MET:HE1	1:F:2086:LEU:HD23	1.95	0.48
1:F:2423:ARG:NH2	1:F:2475:TYR:O	2.44	0.48
1:A:1190:LEU:HD23	1:A:1190:LEU:H	1.78	0.48
1:A:3397:MET:O	1:A:3401:VAL:HG23	2.13	0.48
1:A:4636:THR:OG1	1:A:4637:GLN:N	2.46	0.48
1:C:3505:ARG:HB2	1:C:3551:LEU:HD22	1.95	0.48
1:E:417:ARG:HG3	1:E:420:ARG:HH21	1.77	0.48
1:E:670:TYR:HD2	1:E:672:LYS:HB2	1.78	0.48
1:E:2330:PHE:CE2	1:E:2425:LEU:HD21	2.47	0.48
1:E:2839:MET:HG3	1:E:2892:PHE:HZ	1.78	0.48
1:E:3239:PRO:HB3	1:E:3301:PHE:CG	2.48	0.48
1:F:4195:THR:HB	1:F:4918:LEU:HD11	1.95	0.48
1:A:892:LEU:HD23	1:A:895:MET:HE1	1.94	0.48
1:A:1609:VAL:HG23	1:A:1620:VAL:HG22	1.95	0.48
1:A:1729:PRO:HD3	1:A:1758:LEU:HD22	1.94	0.48
1:C:180:ASP:HB3	1:C:211:LEU:HD12	1.94	0.48
1:C:693:LEU:HD22	1:C:798:ILE:HD11	1.96	0.48
1:E:1113:MET:SD	1:E:1211:GLN:HB3	2.53	0.48
1:F:417:ARG:HG3	1:F:420:ARG:HH21	1.77	0.48
1:F:670:TYR:HD2	1:F:672:LYS:HB2	1.78	0.48
1:F:1708:ILE:O	1:F:1713:SER:HB3	2.14	0.48
1:F:2878:ALA:HA	1:F:2881:LYS:HE3	1.95	0.48
1:F:4043:SER:HA	1:F:4076:THR:HA	1.95	0.48
1:A:1708:ILE:O	1:A:1713:SER:HB3	2.14	0.48
1:A:2618:LYS:O	1:A:2625:GLY:HA2	2.14	0.48
1:A:3522:ALA:HA	1:A:3525:TRP:CD1	2.47	0.48
1:A:4112:THR:O	1:A:4116:THR:HG23	2.13	0.48
1:A:4700:ILE:HD12	1:A:4705:GLN:HG3	1.96	0.48
1:C:670:TYR:HD2	1:C:672:LYS:HB2	1.78	0.48
1:C:894:VAL:HA	1:C:897:LYS:HB2	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:904:TYR:HB3	1:C:918:LEU:HD23	1.95	0.48
1:C:2643:LEU:O	1:C:2647:ILE:HG13	2.12	0.48
1:C:2839:MET:HG3	1:C:2892:PHE:HZ	1.78	0.48
1:C:3041:ALA:HB1	1:C:3120:LEU:HB2	1.94	0.48
1:E:1708:ILE:O	1:E:1713:SER:HB3	2.14	0.48
1:E:2101:LEU:O	1:E:2104:THR:HG22	2.12	0.48
1:E:2878:ALA:HA	1:E:2881:LYS:HE3	1.95	0.48
1:E:2925:LEU:O	1:E:2929:ILE:HG13	2.14	0.48
1:E:3041:ALA:HB1	1:E:3120:LEU:HB2	1.94	0.48
1:E:3166:PRO:HD2	1:E:3167:ILE:HD12	1.93	0.48
1:E:3772:VAL:O	1:E:3776:MET:HG3	2.14	0.48
1:F:903:GLN:O	1:F:915:HIS:N	2.35	0.48
1:F:1939:GLN:HA	1:F:1942:ARG:HG2	1.94	0.48
1:F:3071:MET:HE3	1:F:3135:SER:HA	1.94	0.48
1:F:3159:ALA:HB2	1:F:3240:MET:HE2	1.94	0.48
1:A:263:GLU:OE2	1:A:388:GLN:NE2	2.40	0.48
1:A:1681:VAL:HG21	1:A:1706:LEU:HD21	1.95	0.48
1:A:3166:PRO:HD2	1:A:3167:ILE:HD12	1.93	0.48
1:C:555:LEU:HD11	1:C:585:ALA:HB1	1.96	0.48
1:C:1681:VAL:HG21	1:C:1706:LEU:HD21	1.95	0.48
1:C:2618:LYS:O	1:C:2625:GLY:HA2	2.14	0.48
1:C:2925:LEU:O	1:C:2929:ILE:HG13	2.14	0.48
1:C:3239:PRO:HB3	1:C:3301:PHE:CG	2.49	0.48
1:E:894:VAL:HA	1:E:897:LYS:HB2	1.94	0.48
1:E:1436:GLN:NE2	1:E:1440:ASN:OD1	2.46	0.48
1:F:1190:LEU:HD23	1:F:1190:LEU:H	1.78	0.48
1:F:1968:PRO:O	1:F:1972:ILE:HG12	2.13	0.48
1:F:2925:LEU:O	1:F:2929:ILE:HG13	2.14	0.48
1:A:143:LEU:O	1:A:190:ARG:NE	2.34	0.48
1:C:503:ASP:O	1:C:507:VAL:HG23	2.13	0.48
1:C:925:PRO:HG2	1:C:928:GLU:HB2	1.95	0.48
1:C:3078:GLN:HB3	1:C:3085:GLN:HG3	1.96	0.48
1:C:4636:THR:OG1	1:C:4637:GLN:N	2.46	0.48
1:C:4664:ARG:HD2	1:C:4668:LEU:HD23	1.95	0.48
1:E:2480:GLN:HG2	1:E:2537:THR:HG21	1.94	0.48
1:E:3018:ILE:HG21	1:E:3095:TYR:HB2	1.94	0.48
1:E:3905:PHE:O	1:E:3909:ILE:HG12	2.12	0.48
1:E:4664:ARG:HD2	1:E:4668:LEU:HD23	1.95	0.48
1:F:3505:ARG:HB2	1:F:3551:LEU:HD22	1.95	0.48
1:A:692:HIS:HB3	1:A:795:SER:HB2	1.96	0.48
1:A:925:PRO:HG2	1:A:928:GLU:HB2	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2672:ALA:HB3	1:A:2972:GLN:HE22	1.77	0.48
1:C:1113:MET:SD	1:C:1211:GLN:HB3	2.53	0.48
1:E:1429:SER:OG	1:E:1557:GLU:HB2	2.14	0.48
1:E:1972:ILE:HG13	1:E:3618:LEU:HB3	1.96	0.48
1:E:2569:GLU:O	1:E:2573:LEU:HG	2.13	0.48
1:E:3393:GLU:O	1:E:3397:MET:HE2	2.13	0.48
1:E:4044:LYS:HZ1	1:E:4071:THR:HA	1.79	0.48
1:F:1609:VAL:HG23	1:F:1620:VAL:HG22	1.95	0.48
1:F:2321:ARG:O	1:F:2325:ARG:HG2	2.14	0.48
1:F:2503:THR:O	1:F:2507:SER:N	2.35	0.48
1:F:2618:LYS:O	1:F:2625:GLY:HA2	2.14	0.48
1:F:4057:TYR:HB3	1:F:4061:GLU:HB2	1.95	0.48
1:A:303:GLY:HA2	1:A:420:ARG:HH11	1.77	0.48
1:A:904:TYR:HB3	1:A:918:LEU:HD23	1.95	0.48
1:A:931:TYR:CD2	2:B:101:PRO:HG3	2.49	0.48
1:A:1051:ARG:HG2	1:A:1055:ARG:HE	1.79	0.48
1:C:417:ARG:HG3	1:C:420:ARG:HH21	1.77	0.48
1:C:692:HIS:HB3	1:C:795:SER:HB2	1.96	0.48
1:C:712:GLU:HG2	1:C:838:ARG:HG2	1.96	0.48
1:C:1968:PRO:O	1:C:1972:ILE:HG12	2.13	0.48
1:E:1084:ARG:HG3	1:E:1084:ARG:O	2.12	0.48
1:E:1928:SER:HG	1:E:3619:PHE:HD1	1.60	0.48
1:E:2383:MET:O	1:E:2387:ILE:HG12	2.13	0.48
1:E:3324:LYS:HA	1:E:3327:LYS:HD2	1.95	0.48
1:E:4116:THR:O	1:E:4119:GLU:HG2	2.12	0.48
1:F:1429:SER:OG	1:F:1557:GLU:HB2	2.14	0.48
1:F:1722:MET:HE2	1:F:1759:ARG:HG2	1.96	0.48
1:F:3772:VAL:O	1:F:3776:MET:HG3	2.14	0.48
1:F:4635:ASN:OD1	1:F:4703:LYS:NZ	2.38	0.48
1:A:712:GLU:HG2	1:A:838:ARG:HG2	1.96	0.48
1:A:816:PRO:HG2	1:A:819:TYR:CG	2.48	0.48
1:A:1047:LYS:HB3	1:A:1051:ARG:HH12	1.79	0.48
1:A:1968:PRO:O	1:A:1972:ILE:HG12	2.13	0.48
1:A:1972:ILE:HG13	1:A:3618:LEU:HB3	1.96	0.48
1:A:2472:ASP:OD2	1:A:2529:ARG:NE	2.47	0.48
1:A:2641:ARG:NH1	1:A:2680:MET:HE3	2.19	0.48
1:C:3772:VAL:O	1:C:3776:MET:HG3	2.14	0.48
1:E:1047:LYS:HB3	1:E:1051:ARG:HH12	1.79	0.48
1:E:1144:ARG:HB2	1:E:1192:PHE:CZ	2.49	0.48
1:E:1190:LEU:HD23	1:E:1190:LEU:H	1.78	0.48
1:E:2430:ASP:O	1:E:2434:VAL:HG23	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:3053:LYS:HG3	1:E:3057:ARG:NH1	2.29	0.48
1:E:3505:ARG:HB2	1:E:3551:LEU:HD22	1.95	0.48
1:F:1431:ARG:NH2	1:F:1507:GLU:OE1	2.47	0.48
1:F:2998:LYS:O	1:F:3002:MET:HG3	2.13	0.48
1:F:4044:LYS:HZ1	1:F:4071:THR:HA	1.78	0.48
1:A:781:ASN:HB3	1:A:1466:THR:HG22	1.96	0.48
1:A:1144:ARG:HB2	1:A:1192:PHE:CZ	2.49	0.48
1:A:1431:ARG:NH2	1:A:1507:GLU:OE1	2.47	0.48
1:A:1436:GLN:NE2	1:A:1440:ASN:OD1	2.46	0.48
1:C:781:ASN:HB3	1:C:1466:THR:HG22	1.96	0.48
1:C:931:TYR:CD2	2:D:101:PRO:HG3	2.48	0.48
1:C:2175:VAL:HG12	1:C:2219:TYR:CZ	2.49	0.48
1:C:2998:LYS:O	1:C:3002:MET:HG3	2.13	0.48
1:C:3802:LEU:HB2	1:C:3883:SER:HB2	1.95	0.48
1:E:263:GLU:OE2	1:E:388:GLN:NE2	2.40	0.48
1:E:1444:GLY:HA3	1:E:1487:MET:HA	1.96	0.48
1:E:1968:PRO:O	1:E:1972:ILE:HG12	2.13	0.48
1:E:4195:THR:HB	1:E:4918:LEU:HD11	1.95	0.48
1:E:4700:ILE:HD12	1:E:4705:GLN:HG3	1.96	0.48
1:F:693:LEU:HD22	1:F:798:ILE:HD11	1.96	0.48
1:F:802:PHE:O	1:F:1617:GLY:HA3	2.14	0.48
1:F:2441:MET:HE2	1:F:2506:LEU:HD11	1.95	0.48
1:F:3053:LYS:HG3	1:F:3057:ARG:NH1	2.29	0.48
1:F:4104:LEU:O	1:F:4108:MET:HB2	2.14	0.48
1:A:417:ARG:HG3	1:A:420:ARG:HH21	1.77	0.47
1:A:693:LEU:HD22	1:A:798:ILE:HD11	1.96	0.47
1:A:3122:LEU:H	1:A:3125:VAL:HB	1.79	0.47
1:A:3239:PRO:HB3	1:A:3301:PHE:CG	2.48	0.47
1:A:3772:VAL:O	1:A:3776:MET:HG3	2.14	0.47
1:C:1609:VAL:HG23	1:C:1620:VAL:HG22	1.95	0.47
1:E:802:PHE:O	1:E:1617:GLY:HA3	2.14	0.47
1:E:2774:ILE:HD12	1:E:2891:ILE:HD12	1.96	0.47
1:E:4112:THR:O	1:E:4116:THR:HG23	2.13	0.47
1:F:555:LEU:HD11	1:F:585:ALA:HB1	1.96	0.47
1:F:781:ASN:HB3	1:F:1466:THR:HG22	1.96	0.47
1:F:2472:ASP:OD2	1:F:2529:ARG:NE	2.47	0.47
1:F:2774:ILE:HD12	1:F:2891:ILE:HD12	1.96	0.47
1:F:2839:MET:HG3	1:F:2892:PHE:HZ	1.78	0.47
1:A:1429:SER:OG	1:A:1557:GLU:HB2	2.14	0.47
1:A:1706:LEU:O	1:A:1710:ILE:HG12	2.13	0.47
1:A:2405:MET:HE2	1:A:2407:LEU:H	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2727:SER:OG	1:A:2767:LYS:NZ	2.47	0.47
1:A:3392:GLU:OE1	1:A:3537:THR:OG1	2.31	0.47
1:A:3505:ARG:HB2	1:A:3551:LEU:HD22	1.95	0.47
1:A:4887:CYS:HA	3:A:5101:ATP:H2	1.79	0.47
1:C:626:ARG:O	1:C:630:HIS:ND1	2.47	0.47
1:C:1269:GLU:OE2	1:C:1293:GLN:NE2	2.30	0.47
1:C:2472:ASP:OD2	1:C:2529:ARG:NE	2.47	0.47
1:C:3324:LYS:HA	1:C:3327:LYS:HD2	1.95	0.47
1:C:4777:VAL:HG11	1:C:4816:MET:HG2	1.97	0.47
1:E:693:LEU:HD22	1:E:798:ILE:HD11	1.96	0.47
1:E:2618:LYS:O	1:E:2625:GLY:HA2	2.14	0.47
1:E:3397:MET:O	1:E:3401:VAL:HG23	2.13	0.47
1:E:4777:VAL:HG11	1:E:4816:MET:HG2	1.97	0.47
1:F:1436:GLN:NE2	1:F:1440:ASN:OD1	2.46	0.47
1:F:2727:SER:OG	1:F:2767:LYS:NZ	2.47	0.47
1:F:3041:ALA:HB1	1:F:3120:LEU:HB2	1.94	0.47
1:A:1113:MET:SD	1:A:1211:GLN:HB3	2.53	0.47
1:A:2175:VAL:HG12	1:A:2219:TYR:CZ	2.49	0.47
1:C:15:ARG:HB3	1:C:110:HIS:HB3	1.96	0.47
1:C:1436:GLN:NE2	1:C:1440:ASN:OD1	2.46	0.47
1:C:1920:ARG:NH1	1:C:2006:CYS:SG	2.87	0.47
1:C:2383:MET:O	1:C:2387:ILE:HG12	2.13	0.47
1:C:2878:ALA:HA	1:C:2881:LYS:HE3	1.95	0.47
1:E:15:ARG:HB3	1:E:110:HIS:HB3	1.97	0.47
1:E:70:GLU:OE1	1:E:122:ARG:NH2	2.48	0.47
1:E:781:ASN:HB3	1:E:1466:THR:HG22	1.96	0.47
1:E:2472:ASP:OD2	1:E:2529:ARG:NE	2.47	0.47
1:E:3071:MET:HE3	1:E:3135:SER:HA	1.96	0.47
1:E:4887:CYS:HA	3:E:5101:ATP:H2	1.79	0.47
1:F:626:ARG:O	1:F:630:HIS:ND1	2.47	0.47
1:F:1144:ARG:HB2	1:F:1192:PHE:CZ	2.49	0.47
1:F:3078:GLN:HB3	1:F:3085:GLN:HG3	1.96	0.47
1:A:1920:ARG:NH1	1:A:2006:CYS:SG	2.87	0.47
1:A:3425:ASN:ND2	1:A:3428:SER:HB3	2.30	0.47
1:A:4043:SER:HA	1:A:4076:THR:HA	1.95	0.47
1:A:4635:ASN:OD1	1:A:4703:LYS:NZ	2.38	0.47
1:C:1458:ASP:OD1	1:C:1459:LEU:N	2.48	0.47
1:C:3122:LEU:H	1:C:3125:VAL:HB	1.80	0.47
1:C:3425:ASN:ND2	1:C:3428:SER:HB3	2.30	0.47
1:E:4104:LEU:O	1:E:4108:MET:HB2	2.14	0.47
1:F:70:GLU:OE1	1:F:122:ARG:NH2	2.48	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:1444:GLY:HA3	1:F:1487:MET:HA	1.96	0.47
1:F:1972:ILE:HG13	1:F:3618:LEU:HB3	1.96	0.47
1:F:2383:MET:O	1:F:2387:ILE:HG12	2.13	0.47
1:A:626:ARG:O	1:A:630:HIS:ND1	2.47	0.47
1:A:2320:VAL:HG11	1:A:2418:ILE:HD12	1.97	0.47
1:A:2503:THR:O	1:A:2507:SER:N	2.35	0.47
1:A:3053:LYS:HG3	1:A:3057:ARG:NH1	2.29	0.47
1:C:802:PHE:O	1:C:1617:GLY:HA3	2.14	0.47
1:C:1429:SER:OG	1:C:1557:GLU:HB2	2.14	0.47
1:C:2430:ASP:O	1:C:2434:VAL:HG23	2.14	0.47
1:C:2566:ASP:O	1:C:2570:VAL:HG23	2.15	0.47
1:C:2774:ILE:HD12	1:C:2891:ILE:HD12	1.96	0.47
1:C:3852:ASP:OD1	1:C:3853:PHE:N	2.48	0.47
1:C:4887:CYS:HA	3:C:5101:ATP:H2	1.79	0.47
1:E:1932:VAL:HG13	1:E:3612:ARG:HH12	1.80	0.47
1:E:2321:ARG:O	1:E:2325:ARG:HG2	2.14	0.47
1:E:2506:LEU:HA	1:E:2506:LEU:HD12	1.62	0.47
1:E:2727:SER:OG	1:E:2767:LYS:NZ	2.47	0.47
1:E:3324:LYS:O	1:E:3328:LYS:HG2	2.15	0.47
1:F:712:GLU:HG2	1:F:838:ARG:HG2	1.96	0.47
1:F:1051:ARG:HG2	1:F:1055:ARG:HE	1.79	0.47
1:F:1920:ARG:NH1	1:F:2006:CYS:SG	2.87	0.47
1:A:802:PHE:O	1:A:1617:GLY:HA3	2.14	0.47
1:A:853:PRO:HG2	1:A:1209:VAL:HA	1.96	0.47
1:A:903:GLN:O	1:A:915:HIS:N	2.35	0.47
1:A:2566:ASP:O	1:A:2570:VAL:HG23	2.15	0.47
1:A:2767:LYS:HG2	1:A:2771:ARG:HH21	1.80	0.47
1:C:1708:ILE:O	1:C:1713:SER:HB3	2.13	0.47
1:C:2973:TYR:CE1	1:C:2979:LEU:HD21	2.50	0.47
1:C:4043:SER:HA	1:C:4076:THR:HA	1.95	0.47
1:C:4700:ILE:HD12	1:C:4705:GLN:HG3	1.96	0.47
1:E:1706:LEU:O	1:E:1710:ILE:HG12	2.13	0.47
1:E:2566:ASP:O	1:E:2570:VAL:HG23	2.15	0.47
1:F:2320:VAL:HG11	1:F:2418:ILE:HD12	1.97	0.47
1:F:2430:ASP:O	1:F:2434:VAL:HG23	2.14	0.47
1:F:2609:LEU:O	1:F:2613:TYR:HD1	1.98	0.47
1:F:3273:ASN:HA	1:F:3313:LEU:HD11	1.96	0.47
1:A:70:GLU:OE1	1:A:122:ARG:NH2	2.48	0.47
1:A:2321:ARG:O	1:A:2325:ARG:HG2	2.14	0.47
1:A:2672:ALA:O	1:A:2977:HIS:NE2	2.36	0.47
1:A:2878:ALA:HA	1:A:2881:LYS:HE3	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2925:LEU:O	1:A:2929:ILE:HG13	2.14	0.47
2:B:85:LEU:HD13	2:B:125:VAL:HG22	1.97	0.47
1:C:70:GLU:OE1	1:C:122:ARG:NH2	2.48	0.47
1:C:137:ARG:HA	1:C:137:ARG:NE	2.30	0.47
1:C:1051:ARG:HG2	1:C:1055:ARG:HE	1.79	0.47
1:C:2320:VAL:HG11	1:C:2418:ILE:HD12	1.97	0.47
1:C:2767:LYS:HG2	1:C:2771:ARG:HH21	1.80	0.47
1:C:3939:LEU:HD21	1:C:3980:MET:HE1	1.97	0.47
1:E:681:HIS:HA	1:E:751:THR:HG22	1.97	0.47
1:E:1051:ARG:HG2	1:E:1055:ARG:HE	1.79	0.47
1:E:1431:ARG:NH2	1:E:1507:GLU:OE1	2.47	0.47
1:E:1722:MET:HE2	1:E:1759:ARG:HG2	1.97	0.47
1:E:2320:VAL:HG11	1:E:2418:ILE:HD12	1.97	0.47
1:E:2767:LYS:HG2	1:E:2771:ARG:HH21	1.80	0.47
1:E:3852:ASP:OD1	1:E:3853:PHE:N	2.48	0.47
1:F:290:ARG:NH2	1:F:343:ARG:O	2.48	0.47
1:F:934:GLN:OE1	1:F:935:MET:HG2	2.15	0.47
1:F:1458:ASP:OD1	1:F:1459:LEU:N	2.48	0.47
1:F:1932:VAL:HG13	1:F:3612:ARG:HH12	1.80	0.47
1:F:2767:LYS:HG2	1:F:2771:ARG:HH21	1.80	0.47
1:F:2973:TYR:CE1	1:F:2979:LEU:HD21	2.50	0.47
1:F:3309:VAL:H	1:F:3378:TYR:HE2	1.62	0.47
1:F:3324:LYS:HA	1:F:3327:LYS:HD2	1.95	0.47
1:F:3392:GLU:OE1	1:F:3537:THR:OG1	2.31	0.47
1:F:3425:ASN:ND2	1:F:3428:SER:HB3	2.30	0.47
1:F:4777:VAL:HG11	1:F:4816:MET:HG2	1.96	0.47
1:A:2973:TYR:CE1	1:A:2979:LEU:HD21	2.50	0.47
1:A:3944:VAL:HG13	1:A:4005:SER:HB3	1.96	0.47
1:C:2727:SER:OG	1:C:2767:LYS:NZ	2.47	0.47
1:C:3241:LEU:HA	1:C:3244:TYR:HB2	1.97	0.47
1:C:3273:ASN:HA	1:C:3313:LEU:HD11	1.96	0.47
1:C:3392:GLU:OE1	1:C:3537:THR:OG1	2.31	0.47
1:C:3944:VAL:HG13	1:C:4005:SER:HB3	1.96	0.47
1:E:626:ARG:O	1:E:630:HIS:ND1	2.47	0.47
1:E:692:HIS:HB3	1:E:795:SER:HB2	1.96	0.47
1:E:3159:ALA:HB2	1:E:3240:MET:CE	2.45	0.47
1:E:3234:MET:O	1:E:3238:LEU:HB3	2.15	0.47
1:E:3944:VAL:HG13	1:E:4005:SER:HB3	1.96	0.47
1:F:137:ARG:HA	1:F:137:ARG:NE	2.30	0.47
1:F:2455:MET:SD	1:F:2456:SER:N	2.88	0.47
1:F:3241:LEU:HA	1:F:3244:TYR:HB2	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3964:ILE:O	1:F:3968:LYS:HG3	2.15	0.47
2:G:85:LEU:HD13	2:G:125:VAL:HG22	1.97	0.47
2:I:85:LEU:HD13	2:I:125:VAL:HG22	1.96	0.47
1:A:915:HIS:CE1	1:A:917:CYS:HB3	2.50	0.47
1:A:2455:MET:SD	1:A:2456:SER:N	2.88	0.47
1:A:2998:LYS:HG3	1:A:3002:MET:CE	2.45	0.47
1:A:3071:MET:HE3	1:A:3135:SER:HA	1.97	0.47
1:A:3852:ASP:OD1	1:A:3853:PHE:N	2.48	0.47
1:A:4777:VAL:HG11	1:A:4816:MET:HG2	1.97	0.47
1:C:915:HIS:CE1	1:C:917:CYS:HB3	2.50	0.47
1:C:1047:LYS:HB3	1:C:1051:ARG:HH12	1.79	0.47
1:C:1932:VAL:HG13	1:C:3612:ARG:HH12	1.80	0.47
1:C:3250:GLU:O	1:C:3256:HIS:NE2	2.47	0.47
1:C:3393:GLU:HG3	1:C:3397:MET:CE	2.45	0.47
1:E:1649:GLU:OE1	1:E:1649:GLU:N	2.36	0.47
1:E:1920:ARG:NH1	1:E:2006:CYS:SG	2.87	0.47
1:E:4882:ASP:O	1:E:4886:LYS:HG2	2.15	0.47
1:F:904:TYR:HB3	1:F:918:LEU:HD23	1.95	0.47
1:F:1047:LYS:HB3	1:F:1051:ARG:HH12	1.79	0.47
1:F:3234:MET:O	1:F:3238:LEU:HB3	2.15	0.47
1:A:290:ARG:NH2	1:A:343:ARG:O	2.48	0.47
1:A:2774:ILE:HD12	1:A:2891:ILE:HD12	1.96	0.47
1:A:3273:ASN:HA	1:A:3313:LEU:HD11	1.96	0.47
1:A:3536:ARG:HH22	1:A:3540:PRO:HG3	1.80	0.47
1:C:468:GLU:OE1	1:C:468:GLU:N	2.41	0.47
1:C:853:PRO:HG2	1:C:1209:VAL:HA	1.96	0.47
1:C:1245:ARG:HD2	1:C:1694:TYR:CZ	2.50	0.47
1:C:2526:LEU:HB3	1:C:2534:PHE:CE1	2.50	0.47
1:C:4745:ILE:HD11	1:F:4775:VAL:HG21	1.97	0.47
1:E:290:ARG:NH2	1:E:343:ARG:O	2.48	0.47
1:E:2175:VAL:HG12	1:E:2219:TYR:CZ	2.49	0.47
1:E:3309:VAL:H	1:E:3378:TYR:HE2	1.62	0.47
1:E:3536:ARG:HH22	1:E:3540:PRO:HG3	1.80	0.47
1:E:3964:ILE:O	1:E:3968:LYS:HG3	2.15	0.47
1:F:692:HIS:HB3	1:F:795:SER:HB2	1.96	0.47
1:F:2175:VAL:HG12	1:F:2219:TYR:CZ	2.50	0.47
1:F:3250:GLU:O	1:F:3256:HIS:NE2	2.47	0.47
1:F:3536:ARG:HH22	1:F:3540:PRO:HG3	1.80	0.47
1:F:3944:VAL:HG13	1:F:4005:SER:HB3	1.96	0.47
1:A:681:HIS:HA	1:A:751:THR:HG22	1.97	0.46
1:A:1928:SER:HG	1:A:3619:PHE:HD1	1.60	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2430:ASP:O	1:A:2434:VAL:HG23	2.14	0.46
1:A:2433:GLY:O	1:A:2437:ILE:HG12	2.16	0.46
1:A:3159:ALA:HB2	1:A:3240:MET:CE	2.45	0.46
1:A:4104:LEU:O	1:A:4108:MET:HB2	2.14	0.46
1:C:1444:GLY:HA3	1:C:1487:MET:HA	1.96	0.46
1:C:2998:LYS:HG3	1:C:3002:MET:CE	2.45	0.46
1:C:3309:VAL:H	1:C:3378:TYR:HE2	1.63	0.46
1:C:3964:ILE:O	1:C:3968:LYS:HG3	2.15	0.46
1:E:712:GLU:HG2	1:E:838:ARG:HG2	1.96	0.46
1:E:2998:LYS:HG3	1:E:3002:MET:CE	2.45	0.46
1:E:3078:GLN:HB3	1:E:3085:GLN:HG3	1.96	0.46
1:F:853:PRO:HG2	1:F:1209:VAL:HA	1.96	0.46
1:F:4700:ILE:HD12	1:F:4705:GLN:HG3	1.96	0.46
1:A:2526:LEU:HB3	1:A:2534:PHE:CE1	2.50	0.46
1:A:2609:LEU:O	1:A:2613:TYR:HD1	1.98	0.46
1:A:3250:GLU:O	1:A:3256:HIS:NE2	2.47	0.46
1:A:3324:LYS:O	1:A:3328:LYS:HG2	2.15	0.46
1:A:3964:ILE:O	1:A:3968:LYS:HG3	2.15	0.46
1:C:308:LEU:HD21	1:C:370:LEU:HD12	1.97	0.46
1:C:1117:TRP:HB3	1:C:1201:PHE:HB3	1.98	0.46
1:C:1431:ARG:NH2	1:C:1507:GLU:OE1	2.47	0.46
1:C:1801:LYS:HE3	1:C:1801:LYS:HB3	1.71	0.46
1:C:1972:ILE:HG13	1:C:3618:LEU:HB3	1.96	0.46
1:E:555:LEU:HD11	1:E:585:ALA:HB1	1.96	0.46
1:E:563:GLU:OE1	1:E:563:GLU:N	2.39	0.46
1:E:2433:GLY:O	1:E:2437:ILE:HG12	2.16	0.46
1:E:3273:ASN:HA	1:E:3313:LEU:HD11	1.96	0.46
1:F:1086:ARG:NH1	1:F:1254:ARG:HH21	2.13	0.46
1:F:3245:MET:O	1:F:3249:TRP:HB2	2.15	0.46
1:A:15:ARG:HB3	1:A:110:HIS:HB3	1.97	0.46
1:A:563:GLU:OE1	1:A:563:GLU:N	2.39	0.46
1:A:1117:TRP:HB3	1:A:1201:PHE:HB3	1.97	0.46
1:A:1444:GLY:HA3	1:A:1487:MET:HA	1.96	0.46
1:A:3555:ASN:O	1:A:3559:HIS:ND1	2.41	0.46
1:C:290:ARG:NH2	1:C:343:ARG:O	2.48	0.46
1:C:3324:LYS:O	1:C:3328:LYS:HG2	2.15	0.46
2:D:34:MET:O	2:D:50:THR:HG23	2.16	0.46
1:E:1086:ARG:NH1	1:E:1254:ARG:HH21	2.13	0.46
1:E:2325:ARG:HA	1:E:2325:ARG:HD3	1.73	0.46
1:E:2455:MET:SD	1:E:2456:SER:N	2.88	0.46
1:E:2973:TYR:CE1	1:E:2979:LEU:HD21	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:3245:MET:O	1:E:3249:TRP:HB2	2.16	0.46
1:F:681:HIS:HA	1:F:751:THR:HG22	1.97	0.46
1:F:1117:TRP:HB3	1:F:1201:PHE:HB3	1.97	0.46
1:F:2998:LYS:HG3	1:F:3002:MET:CE	2.45	0.46
1:F:3319:LEU:HB2	1:F:3320:PRO:HD3	1.97	0.46
1:F:3324:LYS:O	1:F:3328:LYS:HG2	2.15	0.46
1:F:3397:MET:O	1:F:3401:VAL:HG23	2.13	0.46
1:F:3555:ASN:O	1:F:3559:HIS:ND1	2.41	0.46
1:A:1458:ASP:OD1	1:A:1459:LEU:N	2.48	0.46
1:A:3234:MET:O	1:A:3238:LEU:HB3	2.15	0.46
1:A:3393:GLU:HG3	1:A:3397:MET:CE	2.45	0.46
1:A:4775:VAL:HG21	1:E:4745:ILE:HD11	1.97	0.46
2:B:34:MET:O	2:B:50:THR:HG23	2.16	0.46
1:C:375:GLN:NE2	1:C:390:LYS:HB2	2.31	0.46
1:C:1722:MET:HE2	1:C:1759:ARG:HG2	1.97	0.46
1:C:2455:MET:SD	1:C:2456:SER:N	2.88	0.46
1:C:3289:ILE:HD12	1:C:3291:GLU:H	1.81	0.46
1:C:4882:ASP:O	1:C:4886:LYS:HG2	2.15	0.46
1:E:673:TRP:HD1	1:E:759:LEU:HD12	1.81	0.46
1:E:904:TYR:HB3	1:E:918:LEU:HD23	1.95	0.46
1:E:934:GLN:OE1	1:E:935:MET:HG2	2.15	0.46
1:E:2497:ALA:O	1:E:2500:SER:OG	2.29	0.46
1:E:2526:LEU:HB3	1:E:2534:PHE:HE1	1.80	0.46
1:E:2609:LEU:O	1:E:2613:TYR:HD1	1.98	0.46
1:E:4151:ILE:HD12	1:E:4156:ARG:HH21	1.81	0.46
1:F:2526:LEU:HB3	1:F:2534:PHE:CE1	2.50	0.46
1:F:3159:ALA:HB2	1:F:3240:MET:CE	2.45	0.46
2:G:52:THR:HG21	2:G:102:ASN:HA	1.98	0.46
1:A:308:LEU:HD21	1:A:370:LEU:HD12	1.97	0.46
1:A:3241:LEU:HA	1:A:3244:TYR:HB2	1.97	0.46
1:A:4882:ASP:O	1:A:4886:LYS:HG2	2.15	0.46
1:C:887:GLU:OE1	1:C:890:HIS:NE2	2.49	0.46
1:C:934:GLN:OE1	1:C:935:MET:HG2	2.15	0.46
1:C:1272:ARG:NH2	1:C:1583:CYS:SG	2.83	0.46
1:C:2321:ARG:O	1:C:2325:ARG:HG2	2.14	0.46
1:C:3536:ARG:HH22	1:C:3540:PRO:HG3	1.80	0.46
1:C:3717:GLU:O	1:C:3721:GLN:HG2	2.16	0.46
1:C:3920:THR:HG22	1:C:3980:MET:HA	1.98	0.46
1:E:16:THR:HB	1:E:110:HIS:HA	1.98	0.46
1:E:853:PRO:HG2	1:E:1209:VAL:HA	1.96	0.46
1:E:1458:ASP:OD1	1:E:1459:LEU:N	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:308:LEU:HD21	1:F:370:LEU:HD12	1.97	0.46
1:F:887:GLU:OE1	1:F:890:HIS:NE2	2.49	0.46
1:F:1458:ASP:HB3	1:F:1461:ARG:HE	1.80	0.46
1:F:3393:GLU:HG3	1:F:3397:MET:CE	2.45	0.46
1:F:3852:ASP:OD1	1:F:3853:PHE:N	2.48	0.46
1:F:4887:CYS:HA	3:F:5101:ATP:H2	1.79	0.46
2:I:52:THR:HG21	2:I:102:ASN:HA	1.98	0.46
1:C:681:HIS:HA	1:C:751:THR:HG22	1.97	0.46
1:C:846:TYR:HE2	1:C:1218:GLY:H	1.64	0.46
1:C:2432:VAL:O	1:C:2435:ILE:HG22	2.16	0.46
1:C:2526:LEU:HB3	1:C:2534:PHE:HE1	1.80	0.46
1:C:2609:LEU:O	1:C:2613:TYR:HD1	1.98	0.46
1:C:4104:LEU:O	1:C:4108:MET:HB2	2.15	0.46
2:D:85:LEU:HD13	2:D:125:VAL:HG22	1.97	0.46
1:E:228:LEU:HA	1:E:356:TYR:CD2	2.50	0.46
1:E:375:GLN:NE2	1:E:390:LYS:HB2	2.31	0.46
1:E:2526:LEU:HB3	1:E:2534:PHE:CE1	2.50	0.46
1:E:3108:PHE:N	1:E:3108:PHE:CD1	2.83	0.46
1:E:3220:LEU:HD22	1:E:3225:ILE:HD13	1.98	0.46
1:F:228:LEU:HA	1:F:356:TYR:CD2	2.50	0.46
1:F:846:TYR:HE2	1:F:1218:GLY:H	1.64	0.46
1:F:915:HIS:CE1	1:F:917:CYS:HB3	2.50	0.46
1:F:3122:LEU:H	1:F:3125:VAL:HB	1.80	0.46
1:F:4894:ASN:HD22	1:F:4894:ASN:C	2.19	0.46
1:A:228:LEU:HA	1:A:356:TYR:CD2	2.50	0.46
1:A:375:GLN:NE2	1:A:390:LYS:HB2	2.31	0.46
1:A:1932:VAL:HG13	1:A:3612:ARG:HH12	1.80	0.46
1:A:2441:MET:HE2	1:A:2506:LEU:HD11	1.97	0.46
1:C:3159:ALA:HB2	1:C:3240:MET:CE	2.45	0.46
1:E:137:ARG:HA	1:E:137:ARG:NE	2.30	0.46
1:E:915:HIS:CE1	1:E:917:CYS:HB3	2.50	0.46
1:E:1482:ARG:HG2	1:E:1535:GLU:OE2	2.16	0.46
1:E:2778:LEU:HD23	1:E:2781:MET:HE3	1.98	0.46
1:E:3393:GLU:HG3	1:E:3397:MET:CE	2.45	0.46
1:E:4914:LEU:H	3:E:5101:ATP:HN61	1.64	0.46
1:F:15:ARG:HB3	1:F:110:HIS:HB3	1.97	0.46
1:F:1245:ARG:HD2	1:F:1694:TYR:CZ	2.50	0.46
1:F:2731:TRP:O	1:F:2735:LYS:HG2	2.16	0.46
1:F:2996:SER:O	1:F:3000:LYS:HG3	2.16	0.46
1:A:555:LEU:HD11	1:A:585:ALA:HB1	1.96	0.46
1:A:1086:ARG:NH1	1:A:1254:ARG:HH21	2.13	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1245:ARG:HD2	1:A:1694:TYR:CZ	2.50	0.46
1:A:3220:LEU:HD22	1:A:3225:ILE:HD13	1.98	0.46
1:A:3319:LEU:HB2	1:A:3320:PRO:HD3	1.97	0.46
1:C:322:ALA:HB1	1:C:327:THR:HG21	1.98	0.46
1:C:1458:ASP:HB3	1:C:1461:ARG:HE	1.80	0.46
1:C:3053:LYS:HG3	1:C:3057:ARG:NH1	2.29	0.46
1:C:3117:GLY:HA2	1:C:3121:ILE:HD13	1.98	0.46
1:E:1117:TRP:HB3	1:E:1201:PHE:HB3	1.98	0.46
1:E:2158:PRO:HA	1:E:2161:MET:HB2	1.98	0.46
1:E:3122:LEU:H	1:E:3125:VAL:HB	1.80	0.46
1:E:3250:GLU:O	1:E:3256:HIS:NE2	2.47	0.46
1:F:2566:ASP:O	1:F:2570:VAL:HG23	2.14	0.46
1:F:3220:LEU:HD22	1:F:3225:ILE:HD13	1.98	0.46
1:F:3717:GLU:O	1:F:3721:GLN:HG2	2.16	0.46
2:G:34:MET:O	2:G:50:THR:HG23	2.16	0.46
1:A:137:ARG:HA	1:A:137:ARG:NE	2.30	0.46
1:A:375:GLN:HE21	1:A:390:LYS:HB2	1.80	0.46
1:A:468:GLU:OE1	1:A:468:GLU:N	2.41	0.46
1:A:2441:MET:CE	1:A:2506:LEU:HD11	2.46	0.46
1:A:2731:TRP:O	1:A:2735:LYS:HG2	2.16	0.46
1:A:3078:GLN:HB3	1:A:3085:GLN:HG3	1.96	0.46
1:A:3920:THR:HG22	1:A:3980:MET:HA	1.98	0.46
1:C:16:THR:HB	1:C:110:HIS:HA	1.98	0.46
1:C:1144:ARG:HB2	1:C:1192:PHE:CZ	2.49	0.46
1:C:2433:GLY:O	1:C:2437:ILE:HG12	2.16	0.46
1:C:2996:SER:O	1:C:3000:LYS:HG3	2.16	0.46
1:C:4151:ILE:HD12	1:C:4156:ARG:HH21	1.81	0.46
1:E:375:GLN:HE21	1:E:390:LYS:HB2	1.80	0.46
1:E:1483:SER:HB3	1:E:1486:TYR:CE2	2.51	0.46
1:E:3241:LEU:HA	1:E:3244:TYR:HB2	1.97	0.46
1:E:4000:ASP:O	1:E:4004:GLU:HG2	2.16	0.46
1:F:2526:LEU:HB3	1:F:2534:PHE:HE1	1.80	0.46
1:F:2621:CYS:SG	1:F:2622:LEU:N	2.89	0.46
1:A:254:GLU:O	1:A:258:ARG:HG3	2.16	0.46
1:A:2158:PRO:HA	1:A:2161:MET:HB2	1.98	0.46
1:A:2484:LEU:HD13	1:A:2540:HIS:ND1	2.32	0.46
1:A:2526:LEU:HB3	1:A:2534:PHE:HE1	1.80	0.46
1:A:2614:GLU:HG3	1:A:2670:ALA:HB2	1.98	0.46
1:A:3122:LEU:HA	1:A:3126:GLN:HG3	1.98	0.46
1:A:3249:TRP:CZ3	1:A:3308:LYS:HD3	2.51	0.46
1:A:3717:GLU:O	1:A:3721:GLN:HG2	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:309:MET:N	1:C:309:MET:SD	2.89	0.46
1:C:1482:ARG:HG2	1:C:1535:GLU:OE2	2.16	0.46
1:C:3234:MET:O	1:C:3238:LEU:HB3	2.15	0.46
1:C:4044:LYS:HZ1	1:C:4071:THR:HA	1.81	0.46
1:C:4914:LEU:H	3:C:5101:ATP:HN61	1.64	0.46
1:E:887:GLU:OE1	1:E:890:HIS:NE2	2.49	0.46
1:E:986:ILE:HG21	1:E:1059:GLY:HA2	1.98	0.46
1:E:2405:MET:CE	1:E:2407:LEU:H	2.29	0.46
1:E:3831:ASP:OD1	1:E:3831:ASP:N	2.49	0.46
1:F:1482:ARG:HG2	1:F:1535:GLU:OE2	2.16	0.46
1:F:2405:MET:CE	1:F:2407:LEU:H	2.29	0.46
1:F:3854:GLN:NE2	1:F:3921:GLU:O	2.49	0.46
2:G:105:ASN:ND2	2:G:111:ASN:HB2	2.31	0.46
1:A:1458:ASP:HB3	1:A:1461:ARG:HE	1.80	0.45
1:A:1577:LYS:HE2	1:A:1577:LYS:HA	1.98	0.45
1:A:1903:LEU:O	1:A:1907:LEU:HG	2.16	0.45
1:A:2496:ARG:NH2	1:A:2546:SER:OG	2.49	0.45
1:A:2506:LEU:HD12	1:A:2506:LEU:HA	1.62	0.45
1:A:3117:GLY:HA2	1:A:3121:ILE:HD13	1.98	0.45
1:A:3934:LEU:HD23	1:A:3939:LEU:HD22	1.98	0.45
1:A:4000:ASP:O	1:A:4004:GLU:HG2	2.16	0.45
1:A:4151:ILE:HD12	1:A:4156:ARG:HH21	1.81	0.45
1:C:1483:SER:HB3	1:C:1486:TYR:CE2	2.51	0.45
1:C:1903:LEU:O	1:C:1907:LEU:HG	2.16	0.45
1:C:2614:GLU:HG3	1:C:2670:ALA:HB2	1.98	0.45
1:C:2621:CYS:SG	1:C:2622:LEU:N	2.89	0.45
1:C:2731:TRP:O	1:C:2735:LYS:HG2	2.16	0.45
1:C:2977:HIS:O	1:C:3440:LYS:HD3	2.16	0.45
1:C:3122:LEU:HA	1:C:3126:GLN:HG3	1.98	0.45
1:C:3220:LEU:HD22	1:C:3225:ILE:HD13	1.98	0.45
1:E:1245:ARG:HD2	1:E:1694:TYR:CZ	2.50	0.45
1:E:1272:ARG:HH12	1:E:1585:PRO:HA	1.82	0.45
1:E:1903:LEU:O	1:E:1907:LEU:HG	2.16	0.45
1:F:1577:LYS:HE2	1:F:1577:LYS:HA	1.98	0.45
1:F:2496:ARG:NH2	1:F:2546:SER:OG	2.49	0.45
1:F:4151:ILE:HD12	1:F:4156:ARG:HH21	1.81	0.45
1:F:4882:ASP:O	1:F:4886:LYS:HG2	2.15	0.45
1:F:4914:LEU:H	3:F:5101:ATP:HN61	1.64	0.45
2:I:34:MET:O	2:I:50:THR:HG23	2.16	0.45
1:A:887:GLU:OE1	1:A:890:HIS:NE2	2.49	0.45
1:A:934:GLN:OE1	1:A:935:MET:HG2	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2621:CYS:SG	1:A:2622:LEU:N	2.89	0.45
1:A:3000:LYS:HB2	1:A:3043:THR:HG21	1.99	0.45
1:C:1086:ARG:NH1	1:C:1254:ARG:HH21	2.13	0.45
1:C:2065:MET:HG3	1:C:2083:MET:HG2	1.98	0.45
1:E:309:MET:N	1:E:309:MET:SD	2.89	0.45
1:E:846:TYR:HE2	1:E:1218:GLY:H	1.64	0.45
1:E:1801:LYS:HE3	1:E:1801:LYS:HB3	1.71	0.45
1:E:3319:LEU:HB2	1:E:3320:PRO:HD3	1.97	0.45
1:E:3425:ASN:ND2	1:E:3428:SER:HB3	2.30	0.45
1:E:4017:ASP:OD1	1:E:4124:VAL:HG13	2.17	0.45
1:F:673:TRP:HD1	1:F:759:LEU:HD12	1.81	0.45
1:F:1903:LEU:O	1:F:1907:LEU:HG	2.16	0.45
1:F:2433:GLY:O	1:F:2437:ILE:HG12	2.16	0.45
1:F:3010:LEU:O	1:F:3014:VAL:HG23	2.17	0.45
1:F:3122:LEU:HA	1:F:3126:GLN:HG3	1.98	0.45
1:A:322:ALA:HB1	1:A:327:THR:HG21	1.98	0.45
1:A:912:LYS:HB2	1:A:914:GLN:HG2	1.99	0.45
1:A:986:ILE:HG21	1:A:1059:GLY:HA2	1.98	0.45
1:A:1272:ARG:HH12	1:A:1585:PRO:HA	1.81	0.45
1:A:3309:VAL:H	1:A:3378:TYR:HE2	1.62	0.45
1:A:4017:ASP:OD1	1:A:4124:VAL:HG13	2.17	0.45
1:C:228:LEU:HA	1:C:356:TYR:CD2	2.50	0.45
1:C:2680:MET:HA	1:C:2680:MET:CE	2.46	0.45
1:C:3736:ALA:HB1	1:C:3776:MET:HG2	1.98	0.45
1:C:3854:GLN:NE2	1:C:3921:GLU:O	2.49	0.45
1:E:2614:GLU:HG3	1:E:2670:ALA:HB2	1.98	0.45
1:E:2731:TRP:O	1:E:2735:LYS:HG2	2.16	0.45
1:E:3000:LYS:HB2	1:E:3043:THR:HG21	1.99	0.45
1:E:3030:ASN:HA	1:E:3033:HIS:HD2	1.81	0.45
1:F:16:THR:HB	1:F:110:HIS:HA	1.98	0.45
1:F:2352:ILE:HD13	1:F:2358:ARG:HB3	1.98	0.45
1:F:2441:MET:CE	1:F:2506:LEU:HD11	2.46	0.45
1:F:3289:ILE:HD12	1:F:3291:GLU:H	1.81	0.45
1:F:3736:ALA:HB1	1:F:3776:MET:HG2	1.98	0.45
1:A:673:TRP:HD1	1:A:759:LEU:HD12	1.81	0.45
1:A:2233:MET:O	1:A:2296:ARG:NH2	2.48	0.45
1:C:169:ARG:HD2	1:C:176:ARG:HH12	1.82	0.45
1:C:986:ILE:HG21	1:C:1059:GLY:HA2	1.98	0.45
1:C:1576:HIS:CD2	1:C:1577:LYS:HG2	2.51	0.45
1:E:308:LEU:HD21	1:E:370:LEU:HD12	1.97	0.45
1:E:912:LYS:HB2	1:E:914:GLN:HG2	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2352:ILE:HD13	1:E:2358:ARG:HB3	1.98	0.45
1:F:322:ALA:HB1	1:F:327:THR:HG21	1.98	0.45
1:F:1174:MET:CB	1:F:1190:LEU:HA	2.47	0.45
1:F:3000:LYS:HB2	1:F:3043:THR:HG21	1.99	0.45
1:A:2432:VAL:O	1:A:2435:ILE:HG22	2.16	0.45
1:A:2977:HIS:O	1:A:3440:LYS:HD3	2.16	0.45
1:A:3010:LEU:O	1:A:3014:VAL:HG23	2.16	0.45
1:A:3108:PHE:N	1:A:3108:PHE:CD1	2.83	0.45
1:C:2441:MET:CE	1:C:2506:LEU:HD11	2.46	0.45
1:C:2496:ARG:NH2	1:C:2546:SER:OG	2.49	0.45
1:C:3383:TRP:CH2	1:C:3394:LEU:HD23	2.52	0.45
1:C:4183:GLU:HG2	1:C:4186:GLU:HB3	1.98	0.45
1:E:436:LEU:HD21	1:E:518:ALA:HB2	1.99	0.45
1:E:1576:HIS:CD2	1:E:1577:LYS:HG2	2.51	0.45
1:E:2065:MET:HG3	1:E:2083:MET:HG2	1.98	0.45
1:E:2432:VAL:O	1:E:2435:ILE:HG22	2.16	0.45
1:E:2680:MET:CE	1:E:2680:MET:HA	2.46	0.45
1:E:3010:LEU:O	1:E:3014:VAL:HG23	2.17	0.45
1:E:3249:TRP:CZ3	1:E:3308:LYS:HD3	2.51	0.45
1:E:3289:ILE:HD12	1:E:3291:GLU:H	1.81	0.45
1:E:3717:GLU:O	1:E:3721:GLN:HG2	2.16	0.45
1:F:1272:ARG:HH12	1:F:1585:PRO:HA	1.81	0.45
1:F:1483:SER:HB3	1:F:1486:TYR:CE2	2.51	0.45
1:F:2735:LYS:HB3	1:F:2740:TRP:HB2	1.99	0.45
1:F:3327:LYS:O	1:F:3331:MET:HG3	2.16	0.45
1:F:3920:THR:HG22	1:F:3980:MET:HA	1.98	0.45
1:A:241:MET:HE2	1:A:241:MET:HB2	1.78	0.45
1:A:1482:ARG:HG2	1:A:1535:GLU:OE2	2.16	0.45
1:A:1576:HIS:CD2	1:A:1577:LYS:HG2	2.51	0.45
1:A:2680:MET:HA	1:A:2680:MET:CE	2.46	0.45
2:B:119:GLN:OE1	2:B:119:GLN:N	2.50	0.45
1:C:254:GLU:O	1:C:258:ARG:HG3	2.16	0.45
1:C:563:GLU:OE1	1:C:563:GLU:N	2.39	0.45
1:C:912:LYS:HB2	1:C:914:GLN:HG2	1.98	0.45
1:C:1303:ARG:O	1:C:1590:GLN:N	2.38	0.45
1:C:2423:ARG:NH2	1:C:2475:TYR:O	2.44	0.45
1:C:3065:GLU:HG2	1:C:3069:LYS:HE3	1.99	0.45
1:C:3549:ARG:O	1:C:3553:ILE:HG12	2.17	0.45
1:C:3831:ASP:N	1:C:3831:ASP:OD1	2.49	0.45
1:E:1458:ASP:HB3	1:E:1461:ARG:HE	1.80	0.45
1:E:2441:MET:CE	1:E:2506:LEU:HD11	2.46	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2484:LEU:HD13	1:E:2540:HIS:ND1	2.32	0.45
1:E:2977:HIS:O	1:E:3440:LYS:HD3	2.16	0.45
1:E:4579:THR:HG1	1:E:4732:HIS:CD2	2.34	0.45
1:E:4775:VAL:HG21	1:F:4745:ILE:HD11	1.98	0.45
1:F:375:GLN:HE21	1:F:390:LYS:HB2	1.80	0.45
1:F:2432:VAL:O	1:F:2435:ILE:HG22	2.16	0.45
1:F:2540:HIS:HB3	1:F:2543:LEU:HD23	1.99	0.45
1:F:2680:MET:HA	1:F:2680:MET:CE	2.46	0.45
1:F:3108:PHE:N	1:F:3108:PHE:CD1	2.83	0.45
1:F:3831:ASP:N	1:F:3831:ASP:OD1	2.49	0.45
1:F:3934:LEU:HD23	1:F:3939:LEU:HD22	1.98	0.45
1:F:4000:ASP:O	1:F:4004:GLU:HG2	2.16	0.45
1:A:356:TYR:CD1	1:A:407:ARG:HG2	2.52	0.45
1:A:898:ILE:HG21	1:A:973:THR:HA	1.99	0.45
1:A:1483:SER:HB3	1:A:1486:TYR:CE2	2.51	0.45
1:A:2065:MET:HE1	1:A:2086:LEU:HD23	1.99	0.45
1:A:2264:VAL:HG21	1:A:2300:PHE:HE2	1.82	0.45
1:A:2464:LYS:HB3	1:A:2518:TYR:CE1	2.52	0.45
1:A:3030:ASN:HA	1:A:3033:HIS:HD2	1.81	0.45
1:A:3736:ALA:HB1	1:A:3776:MET:HG2	1.98	0.45
1:A:4169:ARG:NH2	3:A:5101:ATP:O1G	2.50	0.45
1:A:4183:GLU:HG2	1:A:4186:GLU:HB3	1.99	0.45
1:C:295:PHE:HE1	1:C:297:LEU:HG	1.82	0.45
1:C:1471:ASP:HB2	1:C:1477:HIS:CE1	2.52	0.45
1:C:2405:MET:CE	1:C:2407:LEU:H	2.29	0.45
1:C:2484:LEU:HD13	1:C:2540:HIS:ND1	2.32	0.45
1:C:2735:LYS:HB3	1:C:2740:TRP:HB2	1.99	0.45
1:C:3097:THR:HG21	1:C:3146:TYR:CE2	2.52	0.45
1:C:3108:PHE:N	1:C:3108:PHE:CD1	2.83	0.45
1:C:3934:LEU:HD23	1:C:3939:LEU:HD22	1.97	0.45
1:C:4000:ASP:O	1:C:4004:GLU:HG2	2.16	0.45
1:C:4894:ASN:HD22	1:C:4894:ASN:C	2.19	0.45
2:D:52:THR:HG21	2:D:102:ASN:HA	1.98	0.45
1:E:898:ILE:HG21	1:E:973:THR:HA	1.99	0.45
1:E:3122:LEU:HA	1:E:3126:GLN:HG3	1.98	0.45
1:E:3327:LYS:O	1:E:3331:MET:HG3	2.16	0.45
1:E:3854:GLN:NE2	1:E:3921:GLU:O	2.49	0.45
1:E:3920:THR:HG22	1:E:3980:MET:HA	1.98	0.45
1:E:4169:ARG:NH2	3:E:5101:ATP:O1G	2.50	0.45
1:E:4941:LYS:HE2	1:E:4941:LYS:HB3	1.82	0.45
1:F:375:GLN:NE2	1:F:390:LYS:HB2	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2158:PRO:HA	1:F:2161:MET:HB2	1.98	0.45
1:F:2614:GLU:HG3	1:F:2670:ALA:HB2	1.98	0.45
1:F:3097:THR:HG21	1:F:3146:TYR:CE2	2.52	0.45
1:A:593:HIS:O	1:A:597:ILE:HG13	2.17	0.45
1:A:3245:MET:O	1:A:3249:TRP:HB2	2.16	0.45
1:A:3289:ILE:HD12	1:A:3291:GLU:H	1.81	0.45
1:A:3327:LYS:O	1:A:3331:MET:HG3	2.16	0.45
1:C:2352:ILE:HD13	1:C:2358:ARG:HB3	1.98	0.45
1:C:3000:LYS:HB2	1:C:3043:THR:HG21	1.99	0.45
1:E:2442:PRO:HA	1:E:2454:ASP:CG	2.37	0.45
1:E:2621:CYS:SG	1:E:2622:LEU:N	2.89	0.45
1:E:2996:SER:O	1:E:3000:LYS:HG3	2.16	0.45
1:E:3549:ARG:O	1:E:3553:ILE:HG12	2.17	0.45
1:E:3736:ALA:HB1	1:E:3776:MET:HG2	1.98	0.45
1:E:3934:LEU:HD23	1:E:3939:LEU:HD22	1.98	0.45
1:F:169:ARG:HD2	1:F:176:ARG:HH12	1.82	0.45
1:F:309:MET:N	1:F:309:MET:SD	2.89	0.45
1:F:1576:HIS:CD2	1:F:1577:LYS:HG2	2.51	0.45
1:F:2264:VAL:HG21	1:F:2300:PHE:HE2	1.82	0.45
1:F:3850:ASN:ND2	1:F:3853:PHE:HB2	2.32	0.45
2:I:105:ASN:ND2	2:I:111:ASN:HB2	2.31	0.45
1:A:16:THR:HB	1:A:110:HIS:HA	1.98	0.45
1:A:436:LEU:HD21	1:A:518:ALA:HB2	1.99	0.45
1:A:846:TYR:HE2	1:A:1218:GLY:H	1.64	0.45
1:A:1722:MET:HE2	1:A:1759:ARG:HG2	1.99	0.45
1:A:1786:ASP:OD1	1:A:1786:ASP:N	2.50	0.45
1:A:2405:MET:CE	1:A:2407:LEU:H	2.29	0.45
1:A:2459:PHE:HE1	1:A:2464:LYS:HG3	1.82	0.45
1:A:2996:SER:O	1:A:3000:LYS:HG3	2.16	0.45
1:A:4914:LEU:H	3:A:5101:ATP:HN61	1.64	0.45
2:B:52:THR:HG21	2:B:102:ASN:HA	1.98	0.45
2:B:105:ASN:ND2	2:B:111:ASN:HB2	2.31	0.45
1:C:673:TRP:HD1	1:C:759:LEU:HD12	1.81	0.45
1:C:1786:ASP:OD1	1:C:1786:ASP:N	2.50	0.45
1:C:2464:LYS:HB3	1:C:2518:TYR:CE1	2.52	0.45
1:C:3249:TRP:CZ3	1:C:3308:LYS:HD3	2.51	0.45
1:C:3858:ARG:HG2	1:C:3859:THR:HG23	1.99	0.45
1:E:254:GLU:O	1:E:258:ARG:HG3	2.16	0.45
1:E:2496:ARG:NH2	1:E:2546:SER:OG	2.49	0.45
1:E:3025:ALA:O	1:E:3029:VAL:HG23	2.17	0.45
1:E:3850:ASN:ND2	1:E:3853:PHE:HB2	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:4894:ASN:C	1:E:4894:ASN:HD22	2.19	0.45
1:F:375:GLN:NE2	1:F:390:LYS:O	2.50	0.45
1:F:593:HIS:O	1:F:597:ILE:HG13	2.17	0.45
1:F:917:CYS:HA	1:F:924:LEU:HD11	1.99	0.45
1:F:1786:ASP:OD1	1:F:1786:ASP:N	2.50	0.45
1:F:3025:ALA:O	1:F:3029:VAL:HG23	2.17	0.45
1:F:3858:ARG:HG2	1:F:3859:THR:HG23	1.99	0.45
1:F:4169:ARG:NH2	3:F:5101:ATP:O1G	2.50	0.45
1:A:372:LEU:HD23	1:A:372:LEU:H	1.81	0.45
1:A:1471:ASP:HB2	1:A:1477:HIS:CE1	2.52	0.45
1:A:3065:GLU:HG2	1:A:3069:LYS:HE3	1.99	0.45
1:A:3549:ARG:O	1:A:3553:ILE:HG12	2.17	0.45
1:A:3850:ASN:ND2	1:A:3853:PHE:HB2	2.32	0.45
1:C:375:GLN:HE21	1:C:390:LYS:HB2	1.80	0.45
1:C:2276:CYS:SG	1:C:2279:LEU:HG	2.57	0.45
1:C:3025:ALA:O	1:C:3029:VAL:HG23	2.17	0.45
1:C:3245:MET:O	1:C:3249:TRP:HB2	2.16	0.45
1:C:3319:LEU:HB2	1:C:3320:PRO:HD3	1.97	0.45
1:C:3327:LYS:O	1:C:3331:MET:HG3	2.16	0.45
1:C:4635:ASN:OD1	1:C:4703:LYS:NZ	2.38	0.45
2:D:105:ASN:ND2	2:D:111:ASN:HB2	2.31	0.45
1:E:2464:LYS:HB3	1:E:2518:TYR:CE1	2.52	0.45
1:E:2934:GLU:HB3	1:E:2938:TYR:CZ	2.52	0.45
1:F:11:ILE:HD12	1:F:11:ILE:HA	1.85	0.45
1:F:898:ILE:HG21	1:F:973:THR:HA	1.99	0.45
1:F:1419:TYR:CE2	1:F:1563:ASN:HB3	2.52	0.45
1:F:2065:MET:HG3	1:F:2083:MET:HG2	1.98	0.45
1:F:3249:TRP:CZ3	1:F:3308:LYS:HD3	2.51	0.45
1:F:3292:GLY:HA3	1:F:3295:MET:HE3	1.98	0.45
1:F:4017:ASP:OD1	1:F:4124:VAL:HG13	2.17	0.45
1:A:1091:GLU:HB3	1:A:1094:TYR:HD2	1.82	0.44
1:A:2065:MET:HG3	1:A:2083:MET:HG2	1.98	0.44
1:A:2540:HIS:HB3	1:A:2543:LEU:HD23	1.99	0.44
1:C:356:TYR:CD1	1:C:407:ARG:HG2	2.52	0.44
1:C:1029:ASN:O	1:C:1032:LEU:HB2	2.17	0.44
1:C:1174:MET:CB	1:C:1190:LEU:HA	2.47	0.44
1:C:3030:ASN:HA	1:C:3033:HIS:HD2	1.81	0.44
1:C:3463:SER:HB3	1:C:3466:VAL:HG12	1.99	0.44
1:C:3850:ASN:ND2	1:C:3853:PHE:HB2	2.32	0.44
1:C:4169:ARG:NH2	3:C:5101:ATP:O1G	2.50	0.44
1:C:4928:ASP:OD1	1:C:4929:GLU:N	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1577:LYS:HE2	1:E:1577:LYS:HA	1.98	0.44
1:E:3117:GLY:HA2	1:E:3121:ILE:HD13	1.98	0.44
1:E:4928:ASP:OD1	1:E:4929:GLU:N	2.50	0.44
1:F:436:LEU:HD21	1:F:518:ALA:HB2	1.99	0.44
1:F:986:ILE:HG21	1:F:1059:GLY:HA2	1.98	0.44
1:F:2484:LEU:HD13	1:F:2540:HIS:ND1	2.32	0.44
1:F:3117:GLY:HA2	1:F:3121:ILE:HD13	1.98	0.44
1:F:3939:LEU:HD21	1:F:3980:MET:HE1	1.99	0.44
1:A:1029:ASN:O	1:A:1032:LEU:HB2	2.18	0.44
1:A:3226:ARG:HH12	1:A:3286:ASN:HA	1.82	0.44
1:A:3831:ASP:N	1:A:3831:ASP:OD1	2.49	0.44
1:A:3967:LEU:O	1:A:3971:MET:HG3	2.18	0.44
1:C:908:ARG:NH1	1:C:910:ASP:OD1	2.51	0.44
1:C:1577:LYS:HA	1:C:1577:LYS:HE2	1.98	0.44
1:C:2094:ILE:O	1:C:2098:VAL:HG22	2.18	0.44
1:C:2264:VAL:HG21	1:C:2300:PHE:HE2	1.82	0.44
1:C:2934:GLU:HB3	1:C:2938:TYR:CZ	2.52	0.44
2:D:119:GLN:N	2:D:119:GLN:OE1	2.50	0.44
1:E:295:PHE:HE1	1:E:297:LEU:HG	1.82	0.44
1:E:908:ARG:NH1	1:E:910:ASP:OD1	2.51	0.44
1:E:1029:ASN:O	1:E:1032:LEU:HB2	2.18	0.44
1:E:1786:ASP:N	1:E:1786:ASP:OD1	2.50	0.44
1:E:2459:PHE:HE1	1:E:2464:LYS:HG3	1.82	0.44
1:E:2753:GLN:HG2	1:E:2755:LEU:H	1.83	0.44
1:E:3133:LEU:HB2	1:E:3161:PHE:CE2	2.53	0.44
1:E:3383:TRP:CH2	1:E:3394:LEU:HD23	2.52	0.44
1:E:4183:GLU:HG2	1:E:4186:GLU:HB3	1.99	0.44
1:F:254:GLU:O	1:F:258:ARG:HG3	2.16	0.44
1:F:1471:ASP:HB2	1:F:1477:HIS:CE1	2.52	0.44
1:F:2464:LYS:HB3	1:F:2518:TYR:CE1	2.52	0.44
1:F:2506:LEU:HD12	1:F:2506:LEU:HA	1.62	0.44
1:F:2753:GLN:HG2	1:F:2755:LEU:H	1.83	0.44
1:A:1844:LEU:HA	1:A:1847:ILE:HG22	1.99	0.44
1:A:2442:PRO:HA	1:A:2454:ASP:CG	2.37	0.44
1:A:3097:THR:HG21	1:A:3146:TYR:CE2	2.52	0.44
1:A:4042:ILE:HD11	1:A:4079:TYR:HB3	2.00	0.44
1:C:375:GLN:NE2	1:C:390:LYS:O	2.50	0.44
1:C:1114:ARG:NH1	1:C:1127:GLU:OE1	2.51	0.44
1:C:2158:PRO:HA	1:C:2161:MET:HB2	1.98	0.44
1:C:2405:MET:HE2	1:C:2407:LEU:H	1.83	0.44
1:C:2981:PHE:HB3	1:C:3000:LYS:CE	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4088:GLU:OE1	1:C:4088:GLU:N	2.51	0.44
1:E:372:LEU:HD23	1:E:372:LEU:H	1.81	0.44
1:E:426:PHE:HZ	1:E:456:LEU:HD21	1.82	0.44
1:E:593:HIS:O	1:E:597:ILE:HG13	2.17	0.44
1:E:1762:MET:HE2	1:E:1762:MET:HB2	1.81	0.44
1:E:1844:LEU:HA	1:E:1847:ILE:HG22	1.99	0.44
1:E:2735:LYS:HB3	1:E:2740:TRP:HB2	1.99	0.44
1:E:3486:GLU:O	1:E:3490:LEU:HG	2.17	0.44
1:E:3660:VAL:HG13	1:E:3664:HIS:ND1	2.33	0.44
1:E:4608:LYS:O	1:E:4613:GLY:N	2.50	0.44
1:E:4838:TYR:O	1:E:4842:ARG:HB2	2.18	0.44
1:F:1905:MET:O	1:F:1909:LEU:HG	2.17	0.44
1:F:3133:LEU:HB2	1:F:3161:PHE:CE2	2.53	0.44
2:G:119:GLN:N	2:G:119:GLN:OE1	2.50	0.44
2:I:66:ARG:HA	2:I:66:ARG:NE	2.33	0.44
2:I:119:GLN:OE1	2:I:119:GLN:N	2.50	0.44
1:A:309:MET:N	1:A:309:MET:SD	2.89	0.44
1:A:375:GLN:NE2	1:A:390:LYS:O	2.50	0.44
1:A:1039:ASP:O	1:A:1043:LYS:HG3	2.18	0.44
1:A:1837:ASN:HA	1:A:1840:LEU:HD12	1.99	0.44
1:A:2352:ILE:HD13	1:A:2358:ARG:HB3	1.98	0.44
1:A:2423:ARG:NH2	1:A:2475:TYR:O	2.44	0.44
1:A:3854:GLN:NE2	1:A:3921:GLU:O	2.49	0.44
1:A:4838:TYR:O	1:A:4842:ARG:HB2	2.18	0.44
1:C:2775:LYS:HE3	1:C:2775:LYS:HB3	1.89	0.44
1:C:2851:TRP:HH2	1:C:2869:LEU:HD13	1.82	0.44
1:C:3123:GLU:OE2	1:C:3186:ARG:NH2	2.51	0.44
1:C:3555:ASN:O	1:C:3559:HIS:ND1	2.41	0.44
1:E:375:GLN:NE2	1:E:390:LYS:O	2.50	0.44
1:F:372:LEU:HD23	1:F:372:LEU:H	1.81	0.44
1:F:393:MET:HE2	1:F:393:MET:HA	1.99	0.44
1:F:739:ARG:HE	1:F:1467:VAL:HG11	1.82	0.44
1:F:1114:ARG:NH1	1:F:1127:GLU:OE1	2.50	0.44
1:F:1267:HIS:HB3	1:F:1295:ASN:N	2.31	0.44
1:F:3030:ASN:HA	1:F:3033:HIS:HD2	1.81	0.44
1:F:3065:GLU:HG2	1:F:3069:LYS:HE3	1.99	0.44
1:F:3226:ARG:HH12	1:F:3286:ASN:HA	1.82	0.44
1:F:4662:ARG:HE	1:F:4662:ARG:HB3	1.68	0.44
2:G:66:ARG:NE	2:G:66:ARG:HA	2.33	0.44
1:A:12:GLN:H	1:A:12:GLN:HG3	1.61	0.44
1:A:291:TRP:CD1	1:A:353:GLU:HB3	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:739:ARG:HE	1:A:1467:VAL:HG11	1.82	0.44
1:A:908:ARG:NH1	1:A:910:ASP:OD1	2.51	0.44
1:A:999:LEU:HD21	1:A:1050:LEU:HD12	2.00	0.44
1:A:2680:MET:HE3	1:A:2680:MET:HA	1.99	0.44
1:A:4894:ASN:C	1:A:4894:ASN:HD22	2.19	0.44
1:C:732:LEU:HG	1:C:741:VAL:HB	2.00	0.44
1:C:2561:THR:O	1:C:2565:ARG:HG3	2.18	0.44
1:C:3010:LEU:O	1:C:3014:VAL:HG23	2.17	0.44
1:C:3226:ARG:HH12	1:C:3286:ASN:HA	1.82	0.44
1:C:4017:ASP:OD1	1:C:4124:VAL:HG13	2.17	0.44
1:E:322:ALA:HB1	1:E:327:THR:HG21	1.98	0.44
1:E:393:MET:HA	1:E:393:MET:HE2	1.99	0.44
1:E:428:ARG:HH21	1:E:446:ASP:HB2	1.83	0.44
1:E:1516:SER:O	1:E:1533:GLN:NE2	2.38	0.44
1:E:1727:ILE:HG22	1:E:1758:LEU:HD23	2.00	0.44
1:E:1905:MET:O	1:E:1909:LEU:HG	2.17	0.44
1:E:2264:VAL:HG21	1:E:2300:PHE:HE2	1.82	0.44
1:E:2435:ILE:HD12	1:E:2435:ILE:HA	1.86	0.44
1:E:2981:PHE:HB3	1:E:3000:LYS:CE	2.47	0.44
1:E:4042:ILE:HD11	1:E:4079:TYR:HB3	2.00	0.44
1:F:356:TYR:CD1	1:F:407:ARG:HG2	2.52	0.44
1:F:1928:SER:HG	1:F:3619:PHE:HD1	1.65	0.44
1:F:2094:ILE:O	1:F:2098:VAL:HG22	2.18	0.44
1:F:2262:GLU:O	1:F:2266:ARG:HG3	2.18	0.44
1:F:2977:HIS:O	1:F:3440:LYS:HD3	2.16	0.44
1:F:2981:PHE:HB3	1:F:3000:LYS:CE	2.47	0.44
1:A:169:ARG:HD2	1:A:176:ARG:HH12	1.82	0.44
1:A:2723:TYR:CE2	1:A:2774:ILE:HG13	2.53	0.44
1:A:2981:PHE:HB3	1:A:3000:LYS:CE	2.47	0.44
1:A:3383:TRP:CH2	1:A:3394:LEU:HD23	2.52	0.44
1:A:4044:LYS:HZ1	1:A:4071:THR:HA	1.82	0.44
1:A:4921:LEU:O	1:A:4925:ILE:HG13	2.18	0.44
2:B:66:ARG:NE	2:B:66:ARG:HA	2.33	0.44
1:C:12:GLN:H	1:C:12:GLN:HG3	1.61	0.44
1:C:400:ASP:OD1	1:C:400:ASP:N	2.51	0.44
1:C:797:GLY:HA2	1:C:1623:LEU:HA	2.00	0.44
1:C:898:ILE:HG21	1:C:973:THR:HA	1.99	0.44
1:C:2591:LEU:HD11	1:C:2608:LEU:HD23	2.00	0.44
1:C:2753:GLN:HG2	1:C:2755:LEU:H	1.83	0.44
1:C:3133:LEU:HB2	1:C:3161:PHE:CE2	2.53	0.44
1:E:2405:MET:SD	1:E:2408:ILE:HG12	2.58	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:4927:LYS:HE2	1:E:4927:LYS:HB3	1.84	0.44
1:F:661:LEU:HD13	1:F:673:TRP:NE1	2.33	0.44
1:F:797:GLY:HA2	1:F:1623:LEU:HA	2.00	0.44
1:F:1029:ASN:O	1:F:1032:LEU:HB2	2.18	0.44
1:F:2276:CYS:SG	1:F:2279:LEU:HG	2.57	0.44
1:F:2459:PHE:HE1	1:F:2464:LYS:HG3	1.82	0.44
1:F:2497:ALA:O	1:F:2500:SER:OG	2.29	0.44
1:F:2672:ALA:O	1:F:2977:HIS:NE2	2.36	0.44
1:F:2851:TRP:HH2	1:F:2869:LEU:HD13	1.82	0.44
1:F:2934:GLU:HB3	1:F:2938:TYR:CZ	2.52	0.44
1:F:3395:PHE:HD1	1:F:3472:LEU:HB3	1.82	0.44
1:F:3967:LEU:O	1:F:3971:MET:HG3	2.18	0.44
1:F:4183:GLU:HG2	1:F:4186:GLU:HB3	1.98	0.44
1:A:2276:CYS:SG	1:A:2279:LEU:HG	2.58	0.44
1:A:3262:MET:SD	1:A:3262:MET:N	2.91	0.44
1:A:3361:LEU:O	1:A:3365:LEU:HG	2.18	0.44
1:A:3399:ALA:O	1:A:3403:ILE:HG12	2.18	0.44
1:A:3858:ARG:HG2	1:A:3859:THR:HG23	1.99	0.44
1:A:4596:LEU:HG	1:A:4600:LYS:HE3	2.00	0.44
1:A:4598:ILE:HD13	1:A:4708:LYS:HE3	2.00	0.44
1:C:393:MET:HE2	1:C:393:MET:HA	1.99	0.44
1:C:426:PHE:HZ	1:C:456:LEU:HD21	1.82	0.44
1:C:1039:ASP:O	1:C:1043:LYS:HG3	2.18	0.44
1:C:2262:GLU:O	1:C:2266:ARG:HG3	2.18	0.44
1:C:2436:SER:HB3	1:C:2489:VAL:HG12	2.00	0.44
1:E:1174:MET:CB	1:E:1190:LEU:HA	2.47	0.44
1:E:2500:SER:O	1:E:2506:LEU:HD23	2.18	0.44
1:F:916:PRO:HG2	2:G:104:TYR:CE2	2.53	0.44
1:F:1102:TYR:HA	1:F:1164:CYS:O	2.18	0.44
1:F:2442:PRO:HA	1:F:2454:ASP:CG	2.37	0.44
1:F:3383:TRP:CH2	1:F:3394:LEU:HD23	2.52	0.44
1:F:3660:VAL:HG13	1:F:3664:HIS:ND1	2.33	0.44
1:F:4608:LYS:O	1:F:4613:GLY:N	2.50	0.44
1:A:232:ASP:OD1	1:A:233:VAL:N	2.51	0.44
1:A:917:CYS:HA	1:A:924:LEU:HD11	1.99	0.44
1:A:2405:MET:SD	1:A:2408:ILE:HG12	2.58	0.44
1:A:2735:LYS:HB3	1:A:2740:TRP:HB2	1.99	0.44
1:A:3463:SER:HB3	1:A:3466:VAL:HG12	1.99	0.44
1:A:3480:CYS:HB2	1:A:3485:GLN:NE2	2.33	0.44
1:A:4008:ASN:O	1:A:4012:ILE:HG13	2.18	0.44
1:A:4608:LYS:O	1:A:4613:GLY:N	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4928:ASP:OD1	1:A:4929:GLU:N	2.50	0.44
1:C:372:LEU:H	1:C:372:LEU:HD23	1.81	0.44
1:C:728:ASP:OD1	1:C:731:HIS:N	2.41	0.44
1:C:917:CYS:HA	1:C:924:LEU:HD11	1.99	0.44
1:C:1102:TYR:HA	1:C:1164:CYS:O	2.18	0.44
1:C:1272:ARG:HH12	1:C:1585:PRO:HA	1.81	0.44
1:C:1998:PHE:HA	1:C:2001:ASP:OD2	2.18	0.44
1:C:3399:ALA:O	1:C:3403:ILE:HG12	2.18	0.44
1:C:3967:LEU:O	1:C:3971:MET:HG3	2.18	0.44
1:C:4921:LEU:O	1:C:4925:ILE:HG13	2.18	0.44
2:D:66:ARG:NE	2:D:66:ARG:HA	2.33	0.44
1:E:661:LEU:HD13	1:E:673:TRP:NE1	2.33	0.44
1:E:732:LEU:HG	1:E:741:VAL:HB	2.00	0.44
1:E:1040:ASP:HA	1:E:1043:LYS:HE2	2.00	0.44
1:E:2437:ILE:O	1:E:2464:LYS:HE3	2.18	0.44
1:E:2517:ARG:O	1:E:2521:THR:HG23	2.18	0.44
1:E:2540:HIS:HB3	1:E:2543:LEU:HD23	1.99	0.44
1:E:3097:THR:HG21	1:E:3146:TYR:CE2	2.52	0.44
1:E:3123:GLU:OE2	1:E:3186:ARG:NH2	2.51	0.44
1:E:3262:MET:N	1:E:3262:MET:SD	2.91	0.44
1:F:1727:ILE:HG22	1:F:1758:LEU:HD23	2.00	0.44
1:F:2983:SER:HA	1:F:3439:SER:HB3	2.00	0.44
1:F:3197:PRO:HD2	1:F:3203:VAL:HG22	2.00	0.44
1:F:3399:ALA:O	1:F:3403:ILE:HG12	2.18	0.44
1:F:3549:ARG:O	1:F:3553:ILE:HG12	2.17	0.44
1:F:4928:ASP:OD1	1:F:4929:GLU:N	2.50	0.44
1:A:1905:MET:O	1:A:1909:LEU:HG	2.17	0.44
1:A:3074:LEU:HD22	1:A:3147:VAL:HG23	2.00	0.44
1:A:3486:GLU:O	1:A:3490:LEU:HG	2.17	0.44
1:A:4069:ALA:HA	1:A:4082:PHE:CE1	2.53	0.44
1:A:4745:ILE:HD11	1:C:4775:VAL:HG21	1.99	0.44
1:C:739:ARG:HE	1:C:1467:VAL:HG11	1.82	0.44
1:C:1040:ASP:HA	1:C:1043:LYS:HE2	2.00	0.44
1:C:2517:ARG:O	1:C:2521:THR:HG23	2.18	0.44
1:C:2723:TYR:CE2	1:C:2774:ILE:HG13	2.53	0.44
1:C:3480:CYS:HB2	1:C:3485:GLN:NE2	2.33	0.44
1:C:4008:ASN:O	1:C:4012:ILE:HG13	2.18	0.44
1:C:4598:ILE:HD13	1:C:4708:LYS:HE3	2.00	0.44
1:C:4608:LYS:O	1:C:4613:GLY:N	2.50	0.44
1:E:356:TYR:CD1	1:E:407:ARG:HG2	2.52	0.44
1:E:739:ARG:HE	1:E:1467:VAL:HG11	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:917:CYS:HA	1:E:924:LEU:HD11	1.99	0.44
1:E:1039:ASP:O	1:E:1043:LYS:HG3	2.18	0.44
1:E:1419:TYR:CE2	1:E:1563:ASN:HB3	2.52	0.44
1:E:2276:CYS:SG	1:E:2279:LEU:HG	2.57	0.44
1:E:2859:LEU:HD11	1:E:2867:HIS:CD2	2.53	0.44
1:E:3226:ARG:HH12	1:E:3286:ASN:HA	1.82	0.44
1:E:3399:ALA:O	1:E:3403:ILE:HG12	2.18	0.44
1:E:4088:GLU:N	1:E:4088:GLU:OE1	2.51	0.44
1:F:426:PHE:HZ	1:F:456:LEU:HD21	1.82	0.44
1:F:732:LEU:HG	1:F:741:VAL:HB	2.00	0.44
1:F:1091:GLU:HB3	1:F:1094:TYR:HD2	1.82	0.44
1:F:1750:PRO:HG3	1:F:2057:LEU:HD22	2.00	0.44
1:F:2561:THR:O	1:F:2565:ARG:HG3	2.18	0.44
1:F:3123:GLU:OE2	1:F:3186:ARG:NH2	2.51	0.44
1:F:4008:ASN:O	1:F:4012:ILE:HG13	2.18	0.44
1:F:4009:VAL:O	1:F:4013:LEU:HG	2.18	0.44
1:A:267:VAL:HA	1:A:270:HIS:ND1	2.33	0.43
1:A:393:MET:HE2	1:A:393:MET:HA	1.99	0.43
1:A:884:ARG:HG3	1:A:885:LEU:N	2.33	0.43
1:A:916:PRO:HG2	2:B:104:TYR:CE2	2.53	0.43
1:A:1303:ARG:O	1:A:1590:GLN:N	2.38	0.43
1:A:1762:MET:HE2	1:A:1762:MET:HB2	1.84	0.43
1:A:2851:TRP:HH2	1:A:2869:LEU:HD13	1.82	0.43
1:A:3133:LEU:HB2	1:A:3161:PHE:CE2	2.53	0.43
1:A:3395:PHE:HD1	1:A:3472:LEU:HB3	1.82	0.43
1:A:4088:GLU:OE1	1:A:4088:GLU:N	2.51	0.43
2:B:40:ALA:HB3	2:B:43:LYS:HB2	2.00	0.43
1:C:291:TRP:CD1	1:C:353:GLU:HB3	2.53	0.43
1:C:436:LEU:HD21	1:C:518:ALA:HB2	1.99	0.43
1:C:1837:ASN:HA	1:C:1840:LEU:HD12	2.00	0.43
1:C:3361:LEU:O	1:C:3365:LEU:HG	2.18	0.43
1:C:4838:TYR:O	1:C:4842:ARG:HB2	2.18	0.43
1:E:884:ARG:HG3	1:E:885:LEU:N	2.33	0.43
1:E:3553:ILE:O	1:E:3557:LEU:HG	2.18	0.43
1:E:3858:ARG:HG2	1:E:3859:THR:HG23	2.00	0.43
1:E:4030:THR:HA	1:E:4033:GLU:HG3	2.00	0.43
1:E:4596:LEU:HG	1:E:4600:LYS:HE3	2.00	0.43
1:F:468:GLU:OE1	1:F:468:GLU:N	2.41	0.43
1:F:884:ARG:HG3	1:F:885:LEU:N	2.33	0.43
1:F:2500:SER:O	1:F:2506:LEU:HD23	2.18	0.43
1:F:3463:SER:HB3	1:F:3466:VAL:HG12	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:3488:ILE:HG12	1:F:3550:VAL:HA	2.00	0.43
1:F:4069:ALA:HA	1:F:4082:PHE:CE1	2.53	0.43
1:A:400:ASP:OD1	1:A:400:ASP:N	2.51	0.43
1:A:428:ARG:HH21	1:A:446:ASP:HB2	1.83	0.43
1:A:1114:ARG:NH1	1:A:1127:GLU:OE1	2.51	0.43
1:A:2500:SER:O	1:A:2506:LEU:HD23	2.18	0.43
1:A:2936:HIS:CE1	1:A:3013:LEU:HD13	2.53	0.43
1:A:3403:ILE:HD11	1:A:3556:VAL:HA	2.00	0.43
1:A:3553:ILE:O	1:A:3557:LEU:HG	2.18	0.43
1:C:593:HIS:O	1:C:597:ILE:HG13	2.17	0.43
1:C:1113:MET:HB2	1:C:1156:TRP:CZ2	2.54	0.43
1:C:1419:TYR:CE2	1:C:1563:ASN:HB3	2.52	0.43
1:C:2086:LEU:O	1:C:2089:ARG:HG2	2.18	0.43
1:C:2498:ALA:HB2	1:C:2515:LEU:HD21	2.00	0.43
1:C:4009:VAL:O	1:C:4013:LEU:HG	2.19	0.43
1:C:4579:THR:HG1	1:C:4732:HIS:CD2	2.34	0.43
1:E:169:ARG:HD2	1:E:176:ARG:HH12	1.82	0.43
1:E:610:VAL:O	1:E:614:LEU:HG	2.18	0.43
1:E:1809:ASP:N	1:E:1809:ASP:OD1	2.51	0.43
1:E:2414:GLU:O	1:E:2418:ILE:HG12	2.18	0.43
1:E:2936:HIS:CE1	1:E:3013:LEU:HD13	2.53	0.43
1:F:133:LEU:O	1:F:145:PHE:HB3	2.19	0.43
1:F:291:TRP:CD1	1:F:353:GLU:HB3	2.53	0.43
1:F:295:PHE:HE1	1:F:297:LEU:HG	1.82	0.43
1:F:912:LYS:HB2	1:F:914:GLN:HG2	1.99	0.43
1:F:1001:GLU:HG2	1:F:1035:TYR:CD1	2.53	0.43
1:F:1040:ASP:HA	1:F:1043:LYS:HE2	2.00	0.43
1:F:2591:LEU:HD11	1:F:2608:LEU:HD23	2.00	0.43
1:F:3262:MET:SD	1:F:3262:MET:N	2.91	0.43
1:F:3361:LEU:O	1:F:3365:LEU:HG	2.18	0.43
1:A:295:PHE:HE1	1:A:297:LEU:HG	1.82	0.43
1:A:2094:ILE:O	1:A:2098:VAL:HG22	2.18	0.43
1:A:2561:THR:O	1:A:2565:ARG:HG3	2.18	0.43
1:A:2859:LEU:HD11	1:A:2867:HIS:CD2	2.53	0.43
1:A:3197:PRO:HD2	1:A:3203:VAL:HG22	2.00	0.43
1:A:3488:ILE:HG12	1:A:3550:VAL:HA	2.00	0.43
1:A:4595:PRO:HA	1:A:4598:ILE:HG12	2.01	0.43
1:C:1905:MET:O	1:C:1909:LEU:HG	2.17	0.43
1:C:2442:PRO:HA	1:C:2454:ASP:CG	2.38	0.43
1:C:2540:HIS:HB3	1:C:2543:LEU:HD23	1.99	0.43
1:C:2980:TYR:O	1:C:3000:LYS:HE2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3395:PHE:HD1	1:C:3472:LEU:HB3	1.82	0.43
1:E:232:ASP:OD1	1:E:233:VAL:N	2.51	0.43
1:E:267:VAL:HA	1:E:270:HIS:ND1	2.33	0.43
1:E:291:TRP:CD1	1:E:353:GLU:HB3	2.53	0.43
1:E:365:HIS:HD2	1:E:368:THR:HG22	1.83	0.43
1:E:400:ASP:OD1	1:E:400:ASP:N	2.51	0.43
1:E:879:GLU:O	1:E:883:GLU:HG2	2.19	0.43
1:E:2094:ILE:O	1:E:2098:VAL:HG22	2.18	0.43
1:E:2561:THR:O	1:E:2565:ARG:HG3	2.18	0.43
1:E:3967:LEU:O	1:E:3971:MET:HG3	2.18	0.43
1:E:4069:ALA:HA	1:E:4082:PHE:CE1	2.53	0.43
1:E:4595:PRO:HA	1:E:4598:ILE:HG12	2.01	0.43
1:F:610:VAL:O	1:F:614:LEU:HG	2.18	0.43
1:F:881:ILE:HA	1:F:884:ARG:HG2	2.00	0.43
1:F:1113:MET:HB2	1:F:1156:TRP:CZ2	2.54	0.43
1:F:2086:LEU:O	1:F:2089:ARG:HG2	2.18	0.43
1:F:2723:TYR:CE2	1:F:2774:ILE:HG13	2.53	0.43
1:F:4042:ILE:HD11	1:F:4079:TYR:HB3	2.00	0.43
1:F:4922:MET:HE2	1:F:4922:MET:HB2	1.92	0.43
1:A:610:VAL:O	1:A:614:LEU:HG	2.18	0.43
1:A:732:LEU:HG	1:A:741:VAL:HB	2.00	0.43
1:A:797:GLY:HA2	1:A:1623:LEU:HA	2.00	0.43
1:A:2262:GLU:O	1:A:2266:ARG:HG3	2.18	0.43
1:A:2924:PHE:O	1:A:2928:LEU:HG	2.19	0.43
1:A:3025:ALA:O	1:A:3029:VAL:HG23	2.17	0.43
1:C:1437:GLU:OE2	1:C:1439:ALA:HB3	2.19	0.43
1:C:2459:PHE:HE1	1:C:2464:LYS:HG3	1.82	0.43
1:C:2500:SER:O	1:C:2506:LEU:HD23	2.18	0.43
1:C:3014:VAL:HG12	1:C:3095:TYR:HE1	1.83	0.43
1:C:4941:LYS:HE2	1:C:4941:LYS:HB3	1.82	0.43
1:E:1001:GLU:HG2	1:E:1035:TYR:CD1	2.53	0.43
1:E:1471:ASP:HB2	1:E:1477:HIS:CE1	2.52	0.43
1:E:1998:PHE:HA	1:E:2001:ASP:OD2	2.18	0.43
1:E:3065:GLU:HG2	1:E:3069:LYS:HE3	1.99	0.43
1:E:3395:PHE:HD1	1:E:3472:LEU:HB3	1.82	0.43
1:E:4009:VAL:O	1:E:4013:LEU:HG	2.18	0.43
1:F:232:ASP:OD1	1:F:233:VAL:N	2.51	0.43
1:F:879:GLU:O	1:F:883:GLU:HG2	2.19	0.43
1:F:1809:ASP:OD1	1:F:1809:ASP:N	2.51	0.43
1:F:1837:ASN:HA	1:F:1840:LEU:HD12	1.99	0.43
1:F:1998:PHE:HA	1:F:2001:ASP:OD2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2325:ARG:HA	1:F:2325:ARG:HD3	1.73	0.43
1:F:2498:ALA:HB2	1:F:2515:LEU:HD21	2.00	0.43
1:F:2517:ARG:O	1:F:2521:THR:HG23	2.18	0.43
1:F:3480:CYS:HB2	1:F:3485:GLN:NE2	2.33	0.43
1:F:4030:THR:HA	1:F:4033:GLU:HG3	2.00	0.43
1:F:4088:GLU:OE1	1:F:4088:GLU:N	2.51	0.43
1:A:3123:GLU:OE2	1:A:3186:ARG:NH2	2.51	0.43
1:C:50:GLU:OE2	1:C:61:ASP:N	2.52	0.43
1:C:241:MET:HB2	1:C:241:MET:HE2	1.78	0.43
1:C:916:PRO:HG2	2:D:104:TYR:CE2	2.54	0.43
1:C:999:LEU:HD21	1:C:1050:LEU:HD12	2.00	0.43
1:C:1091:GLU:HB3	1:C:1094:TYR:HD2	1.82	0.43
1:C:2859:LEU:HD11	1:C:2867:HIS:CD2	2.53	0.43
1:C:3074:LEU:HD22	1:C:3147:VAL:HG23	2.00	0.43
1:C:3660:VAL:HG13	1:C:3664:HIS:ND1	2.33	0.43
1:C:3731:HIS:NE2	1:C:3775:LYS:HD2	2.34	0.43
1:C:4042:ILE:HD11	1:C:4079:TYR:HB3	2.00	0.43
1:E:797:GLY:HA2	1:E:1623:LEU:HA	2.00	0.43
1:E:881:ILE:HA	1:E:884:ARG:HG2	2.00	0.43
1:E:928:GLU:OE2	2:I:102:ASN:ND2	2.49	0.43
1:E:1102:TYR:HA	1:E:1164:CYS:O	2.18	0.43
1:E:1750:PRO:HG3	1:E:2057:LEU:HD22	2.00	0.43
1:E:2262:GLU:O	1:E:2266:ARG:HG3	2.18	0.43
1:E:3463:SER:HB3	1:E:3466:VAL:HG12	1.99	0.43
1:E:3488:ILE:HG12	1:E:3550:VAL:HA	2.00	0.43
1:E:4598:ILE:HD13	1:E:4708:LYS:HE3	2.00	0.43
1:F:890:HIS:O	1:F:894:VAL:HG13	2.19	0.43
1:F:2980:TYR:O	1:F:3000:LYS:HE2	2.18	0.43
1:F:3365:LEU:O	1:F:3369:TYR:HB2	2.19	0.43
1:A:426:PHE:HZ	1:A:456:LEU:HD21	1.82	0.43
1:A:591:GLU:HA	1:A:631:LEU:HD11	2.01	0.43
1:A:728:ASP:OD1	1:A:731:HIS:N	2.41	0.43
1:A:1040:ASP:HA	1:A:1043:LYS:HE2	2.00	0.43
1:A:1102:TYR:HA	1:A:1164:CYS:O	2.18	0.43
1:A:1113:MET:HB2	1:A:1156:TRP:CZ2	2.54	0.43
1:A:2437:ILE:O	1:A:2464:LYS:HE3	2.18	0.43
1:A:2765:LYS:O	1:A:2768:GLU:HG3	2.19	0.43
1:A:2980:TYR:O	1:A:3000:LYS:HE2	2.18	0.43
1:A:4662:ARG:HH21	1:A:4673:LYS:HD2	1.84	0.43
1:C:267:VAL:HA	1:C:270:HIS:ND1	2.33	0.43
1:C:591:GLU:HA	1:C:631:LEU:HD11	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:661:LEU:HD13	1:C:673:TRP:NE1	2.33	0.43
1:C:1001:GLU:HG2	1:C:1035:TYR:CD1	2.53	0.43
1:C:1928:SER:HG	1:C:3619:PHE:HD1	1.66	0.43
1:C:2405:MET:SD	1:C:2408:ILE:HG12	2.58	0.43
1:C:3486:GLU:O	1:C:3490:LEU:HG	2.17	0.43
1:C:3488:ILE:HG12	1:C:3550:VAL:HA	2.00	0.43
1:C:4030:THR:HA	1:C:4033:GLU:HG3	2.00	0.43
1:C:4508:ALA:HB2	1:C:4578:HIS:HE1	1.83	0.43
1:C:4662:ARG:HH21	1:C:4673:LYS:HD2	1.84	0.43
1:E:894:VAL:O	1:E:898:ILE:HG12	2.18	0.43
1:E:1091:GLU:HB3	1:E:1094:TYR:HD2	1.82	0.43
1:E:2723:TYR:CE2	1:E:2774:ILE:HG13	2.53	0.43
1:E:2983:SER:HA	1:E:3439:SER:HB3	2.00	0.43
1:E:3361:LEU:O	1:E:3365:LEU:HG	2.18	0.43
1:E:4559:VAL:HG22	1:E:4561:GLU:H	1.84	0.43
1:F:267:VAL:HA	1:F:270:HIS:ND1	2.33	0.43
1:F:365:HIS:HD2	1:F:368:THR:HG22	1.83	0.43
1:F:2405:MET:SD	1:F:2408:ILE:HG12	2.58	0.43
1:F:2414:GLU:O	1:F:2418:ILE:HG12	2.18	0.43
1:F:2435:ILE:HD12	1:F:2435:ILE:HA	1.87	0.43
1:F:3014:VAL:HG12	1:F:3095:TYR:HE1	1.83	0.43
1:F:3486:GLU:O	1:F:3490:LEU:HG	2.18	0.43
1:F:4595:PRO:HA	1:F:4598:ILE:HG12	2.00	0.43
1:F:4838:TYR:O	1:F:4842:ARG:HB2	2.18	0.43
1:A:2591:LEU:HD11	1:A:2608:LEU:HD23	2.00	0.43
1:A:2753:GLN:HG2	1:A:2755:LEU:H	1.83	0.43
1:A:4671:MET:HG2	1:A:4671:MET:O	2.19	0.43
1:C:894:VAL:O	1:C:898:ILE:HG12	2.18	0.43
1:C:2936:HIS:CE1	1:C:3013:LEU:HD13	2.53	0.43
1:C:3262:MET:N	1:C:3262:MET:SD	2.91	0.43
1:C:3303:GLN:O	1:C:3306:ILE:HG22	2.19	0.43
1:C:4035:ASP:OD1	1:C:4042:ILE:HG23	2.19	0.43
1:C:4069:ALA:HA	1:C:4082:PHE:CE1	2.53	0.43
1:C:4559:VAL:HG22	1:C:4561:GLU:H	1.83	0.43
1:E:241:MET:HB2	1:E:241:MET:HE2	1.78	0.43
1:E:999:LEU:HD21	1:E:1050:LEU:HD12	2.00	0.43
1:E:2143:ARG:HE	1:E:2143:ARG:HB3	1.67	0.43
1:E:2851:TRP:HH2	1:E:2869:LEU:HD13	1.82	0.43
1:F:400:ASP:OD1	1:F:400:ASP:N	2.51	0.43
1:F:428:ARG:HH21	1:F:446:ASP:HB2	1.83	0.43
1:F:999:LEU:HD21	1:F:1050:LEU:HD12	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:1039:ASP:OD1	1:F:1039:ASP:N	2.52	0.43
1:F:1039:ASP:O	1:F:1043:LYS:HG3	2.18	0.43
1:F:2437:ILE:O	1:F:2464:LYS:HE3	2.18	0.43
1:F:2765:LYS:O	1:F:2768:GLU:HG3	2.19	0.43
1:F:2859:LEU:HD11	1:F:2867:HIS:CD2	2.53	0.43
1:F:3394:LEU:HA	1:F:3397:MET:HE3	2.01	0.43
1:F:4042:ILE:HB	1:F:4047:PHE:HB2	2.00	0.43
1:F:4508:ALA:HB2	1:F:4578:HIS:HE1	1.83	0.43
1:F:4671:MET:HG2	1:F:4671:MET:O	2.19	0.43
1:A:11:ILE:HD12	1:A:11:ILE:HA	1.85	0.43
1:A:390:LYS:HA	1:A:390:LYS:HD3	1.89	0.43
1:A:797:GLY:N	1:A:1622:CYS:O	2.52	0.43
1:A:941:LYS:NZ	1:A:944:LEU:HD11	2.34	0.43
1:A:1001:GLU:HG2	1:A:1035:TYR:CD1	2.53	0.43
1:A:1727:ILE:HG22	1:A:1758:LEU:HD23	2.00	0.43
1:A:1801:LYS:HB3	1:A:1801:LYS:HE3	1.71	0.43
1:A:2086:LEU:O	1:A:2089:ARG:HG2	2.18	0.43
1:A:2414:GLU:O	1:A:2418:ILE:HG12	2.19	0.43
1:A:3303:GLN:O	1:A:3306:ILE:HG22	2.19	0.43
1:A:3660:VAL:HG13	1:A:3664:HIS:ND1	2.33	0.43
1:A:4166:GLU:O	1:A:4169:ARG:HG2	2.19	0.43
1:A:4508:ALA:HB2	1:A:4578:HIS:HE1	1.83	0.43
1:C:232:ASP:OD1	1:C:233:VAL:N	2.51	0.43
1:C:2289:TRP:CZ2	1:C:2387:ILE:HD12	2.54	0.43
1:C:2414:GLU:O	1:C:2418:ILE:HG12	2.19	0.43
1:C:3553:ILE:O	1:C:3557:LEU:HG	2.18	0.43
1:E:758:CYS:SG	1:E:767:SER:HB3	2.59	0.43
1:E:1837:ASN:HA	1:E:1840:LEU:HD12	1.99	0.43
1:E:2210:GLN:NE2	1:E:2244:ALA:O	2.45	0.43
1:E:3394:LEU:HA	1:E:3397:MET:HE3	2.01	0.43
1:E:3731:HIS:NE2	1:E:3775:LYS:HD2	2.34	0.43
1:F:908:ARG:NH1	1:F:910:ASP:OD1	2.51	0.43
1:F:970:TYR:HB2	1:F:971:GLN:H	1.70	0.43
1:F:2936:HIS:CE1	1:F:3013:LEU:HD13	2.53	0.43
1:F:4166:GLU:O	1:F:4169:ARG:HG2	2.19	0.43
1:A:890:HIS:O	1:A:894:VAL:HG13	2.18	0.43
1:A:1174:MET:CB	1:A:1190:LEU:HA	2.47	0.43
1:A:1998:PHE:HA	1:A:2001:ASP:OD2	2.18	0.43
1:A:4135:ILE:HG12	1:A:4149:PHE:HE1	1.84	0.43
2:B:90:THR:HG23	2:B:124:THR:HA	2.01	0.43
1:C:133:LEU:O	1:C:145:PHE:HB3	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:430:ILE:HD13	1:C:500:GLU:HG3	2.01	0.43
1:C:565:LEU:HD23	1:C:565:LEU:O	2.19	0.43
1:C:1727:ILE:HG22	1:C:1758:LEU:HD23	2.00	0.43
1:C:2514:ALA:HA	1:C:2517:ARG:HD2	2.01	0.43
1:C:2765:LYS:O	1:C:2768:GLU:HG3	2.19	0.43
1:C:2852:ALA:O	1:C:2855:LYS:HG3	2.19	0.43
1:C:4199:MET:HE2	1:C:4199:MET:HB2	1.81	0.43
1:C:4596:LEU:HG	1:C:4600:LYS:HE3	2.00	0.43
1:E:890:HIS:O	1:E:894:VAL:HG13	2.19	0.43
1:E:1113:MET:HB2	1:E:1156:TRP:CZ2	2.54	0.43
1:E:1114:ARG:NH1	1:E:1127:GLU:OE1	2.51	0.43
1:E:1258:PHE:HB2	1:E:1303:ARG:HH21	1.84	0.43
1:E:2580:ARG:HG2	1:E:2583:MET:HE3	2.00	0.43
1:E:3159:ALA:HB2	1:E:3240:MET:HE2	2.00	0.43
1:E:3365:LEU:O	1:E:3369:TYR:HB2	2.19	0.43
1:E:4008:ASN:O	1:E:4012:ILE:HG13	2.18	0.43
1:F:1762:MET:HE2	1:F:1762:MET:HB2	1.85	0.43
1:F:3731:HIS:NE2	1:F:3775:LYS:HD2	2.34	0.43
1:F:4559:VAL:HG22	1:F:4561:GLU:H	1.84	0.43
2:I:40:ALA:HB3	2:I:43:LYS:HB2	2.00	0.43
1:A:438:LYS:HG3	1:A:439:LYS:HG2	2.00	0.43
1:A:882:ARG:HH11	1:A:937:LEU:HA	1.84	0.43
1:A:2076:ASP:HB3	1:A:2079:LEU:HB3	2.01	0.43
1:A:2517:ARG:O	1:A:2521:THR:HG23	2.18	0.43
1:A:2929:ILE:HG12	1:A:3006:LEU:HD13	2.01	0.43
1:C:428:ARG:HH21	1:C:446:ASP:HB2	1.83	0.43
1:C:610:VAL:O	1:C:614:LEU:HG	2.18	0.43
1:C:1809:ASP:N	1:C:1809:ASP:OD1	2.51	0.43
1:C:1844:LEU:HA	1:C:1847:ILE:HG22	1.99	0.43
1:C:2924:PHE:O	1:C:2928:LEU:HG	2.19	0.43
1:C:2983:SER:HA	1:C:3439:SER:HB3	2.00	0.43
2:D:40:ALA:HB3	2:D:43:LYS:HB2	2.01	0.43
1:E:591:GLU:HA	1:E:631:LEU:HD11	2.01	0.43
1:E:732:LEU:HB3	1:E:779:PHE:CZ	2.54	0.43
1:E:878:LEU:HD23	1:E:878:LEU:H	1.84	0.43
1:E:1437:GLU:OE2	1:E:1439:ALA:HB3	2.19	0.43
1:E:2498:ALA:HB2	1:E:2515:LEU:HD21	2.00	0.43
1:E:2929:ILE:HG12	1:E:3006:LEU:HD13	2.01	0.43
1:E:3240:MET:HE2	1:E:3240:MET:HA	2.00	0.43
1:E:4035:ASP:OD1	1:E:4042:ILE:HG23	2.18	0.43
1:E:4135:ILE:HG12	1:E:4149:PHE:HE1	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:4166:GLU:O	1:E:4169:ARG:HG2	2.19	0.43
1:E:4648:VAL:O	1:E:4652:VAL:HG12	2.19	0.43
1:F:430:ILE:HD13	1:F:500:GLU:HG3	2.01	0.43
1:F:2193:ALA:O	1:F:2197:ARG:HG3	2.19	0.43
1:F:2436:SER:HB3	1:F:2489:VAL:HG12	2.00	0.43
1:F:3007:PHE:HD1	1:F:3007:PHE:O	2.02	0.43
1:F:4035:ASP:OD1	1:F:4042:ILE:HG23	2.19	0.43
1:A:14:LEU:HD11	1:A:214:VAL:HG21	2.01	0.42
1:A:365:HIS:HD2	1:A:368:THR:HG22	1.83	0.42
1:A:410:HIS:O	1:A:414:ARG:HG2	2.19	0.42
1:A:565:LEU:O	1:A:565:LEU:HD23	2.19	0.42
1:A:2289:TRP:CZ2	1:A:2387:ILE:HD12	2.54	0.42
1:A:2436:SER:HB3	1:A:2489:VAL:HG12	2.00	0.42
1:A:2514:ALA:HA	1:A:2517:ARG:HD2	2.01	0.42
1:A:2852:ALA:O	1:A:2855:LYS:HG3	2.19	0.42
1:C:557:TRP:HE3	1:C:558:LEU:HD23	1.84	0.42
1:C:878:LEU:HD23	1:C:878:LEU:H	1.84	0.42
1:C:941:LYS:NZ	1:C:944:LEU:HD11	2.34	0.42
1:C:2497:ALA:O	1:C:2500:SER:OG	2.29	0.42
1:C:4671:MET:O	1:C:4671:MET:HG2	2.19	0.42
2:D:90:THR:HG23	2:D:124:THR:HA	2.01	0.42
1:E:133:LEU:O	1:E:145:PHE:HB3	2.19	0.42
1:E:430:ILE:HD13	1:E:500:GLU:HG3	2.01	0.42
1:E:797:GLY:N	1:E:1622:CYS:O	2.52	0.42
1:E:2591:LEU:HD11	1:E:2608:LEU:HD23	2.00	0.42
1:E:2852:ALA:O	1:E:2855:LYS:HG3	2.19	0.42
1:E:3197:PRO:HD2	1:E:3203:VAL:HG22	2.00	0.42
1:E:4662:ARG:HH21	1:E:4673:LYS:HD2	1.84	0.42
1:F:184:VAL:HG22	1:F:191:TYR:CD1	2.54	0.42
1:F:797:GLY:N	1:F:1622:CYS:O	2.52	0.42
1:F:1437:GLU:OE2	1:F:1439:ALA:HB3	2.19	0.42
1:F:1844:LEU:HA	1:F:1847:ILE:HG22	1.99	0.42
1:F:2580:ARG:HG2	1:F:2583:MET:HE3	2.01	0.42
1:F:4598:ILE:HD13	1:F:4708:LYS:HE3	2.00	0.42
1:A:661:LEU:HD13	1:A:673:TRP:NE1	2.33	0.42
1:A:1437:GLU:OE2	1:A:1439:ALA:HB3	2.19	0.42
1:A:1739:LEU:HD23	1:A:1739:LEU:H	1.84	0.42
1:A:1750:PRO:HG3	1:A:2057:LEU:HD22	2.00	0.42
1:A:1809:ASP:OD1	1:A:1809:ASP:N	2.51	0.42
1:A:3454:LYS:HA	1:A:3457:ARG:HG3	2.01	0.42
1:A:4009:VAL:O	1:A:4013:LEU:HG	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4648:VAL:O	1:A:4652:VAL:HG12	2.19	0.42
1:C:881:ILE:HA	1:C:884:ARG:HG2	2.00	0.42
1:C:3294:TRP:O	1:C:3298:LEU:HG	2.20	0.42
1:C:4595:PRO:HA	1:C:4598:ILE:HG12	2.01	0.42
1:E:1039:ASP:OD1	1:E:1039:ASP:N	2.52	0.42
1:E:2436:SER:HB3	1:E:2489:VAL:HG12	2.00	0.42
1:F:438:LYS:HG3	1:F:439:LYS:HG2	2.01	0.42
1:F:2998:LYS:HG3	1:F:3002:MET:HE2	2.00	0.42
1:F:3303:GLN:O	1:F:3306:ILE:HG22	2.19	0.42
1:F:3403:ILE:HD11	1:F:3556:VAL:HA	2.00	0.42
1:A:2498:ALA:HB2	1:A:2515:LEU:HD21	2.00	0.42
1:A:2934:GLU:HB3	1:A:2938:TYR:CZ	2.52	0.42
1:A:4042:ILE:HB	1:A:4047:PHE:HB2	2.00	0.42
2:B:18:LEU:HD23	2:B:18:LEU:HA	1.87	0.42
1:C:365:HIS:HD2	1:C:368:THR:HG22	1.83	0.42
1:C:758:CYS:SG	1:C:767:SER:HB3	2.59	0.42
1:C:884:ARG:HG3	1:C:885:LEU:N	2.33	0.42
1:C:890:HIS:O	1:C:894:VAL:HG13	2.18	0.42
1:C:1039:ASP:OD1	1:C:1039:ASP:N	2.52	0.42
1:C:2233:MET:O	1:C:2296:ARG:NH2	2.48	0.42
1:C:2943:ASP:OD2	1:C:3017:ARG:NE	2.48	0.42
1:C:4010:GLU:HG2	1:C:4120:LEU:HD13	2.02	0.42
1:E:480:ARG:NH2	1:E:3677:GLU:OE2	2.52	0.42
1:E:2980:TYR:O	1:E:3000:LYS:HE2	2.18	0.42
1:E:3074:LEU:HD22	1:E:3147:VAL:HG23	2.00	0.42
1:E:3085:GLN:HE21	1:E:3089:VAL:HG12	1.84	0.42
1:E:3480:CYS:HB2	1:E:3485:GLN:NE2	2.33	0.42
1:E:4500:MET:HG2	1:E:4585:CYS:SG	2.60	0.42
1:E:4921:LEU:O	1:E:4925:ILE:HG13	2.18	0.42
1:F:769:ARG:HA	1:F:774:PRO:HA	2.01	0.42
1:F:894:VAL:O	1:F:898:ILE:HG12	2.18	0.42
1:F:1428:TYR:CE1	1:F:1510:CYS:HB2	2.54	0.42
1:F:2852:ALA:O	1:F:2855:LYS:HG3	2.19	0.42
1:F:3553:ILE:O	1:F:3557:LEU:HG	2.18	0.42
1:F:3909:ILE:HG21	1:F:3969:GLU:HB3	2.01	0.42
1:F:4921:LEU:O	1:F:4925:ILE:HG13	2.18	0.42
2:G:107:TRP:HD1	2:G:110:PRO:HB2	1.84	0.42
1:A:732:LEU:HB3	1:A:779:PHE:CZ	2.54	0.42
1:A:894:VAL:O	1:A:898:ILE:HG12	2.18	0.42
1:A:1791:LYS:HG3	1:A:1795:MET:SD	2.59	0.42
1:A:1995:LEU:HD11	1:A:3623:TYR:CE1	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3085:GLN:HE21	1:A:3089:VAL:HG12	1.84	0.42
1:A:4030:THR:HA	1:A:4033:GLU:HG3	2.00	0.42
1:C:480:ARG:NH2	1:C:3677:GLU:OE2	2.53	0.42
1:C:2325:ARG:HD3	1:C:2325:ARG:HA	1.73	0.42
1:C:2584:MET:HE2	1:C:2588:LEU:HG	2.02	0.42
1:C:3197:PRO:HD2	1:C:3203:VAL:HG22	2.00	0.42
1:C:3217:ILE:HD11	1:C:3241:LEU:HD22	2.02	0.42
1:C:3403:ILE:HD11	1:C:3556:VAL:HA	2.00	0.42
1:C:3454:LYS:HA	1:C:3457:ARG:HG3	2.02	0.42
1:C:4042:ILE:HB	1:C:4047:PHE:HB2	2.00	0.42
1:E:14:LEU:HD11	1:E:214:VAL:HG21	2.01	0.42
1:E:430:ILE:HG23	1:E:504:ARG:HE	1.84	0.42
1:E:850:LEU:O	1:E:1207:LEU:HD12	2.20	0.42
1:E:1428:TYR:CE1	1:E:1510:CYS:HB2	2.54	0.42
1:E:2086:LEU:O	1:E:2089:ARG:HG2	2.18	0.42
1:E:2121:SER:O	1:E:2125:ILE:HG12	2.19	0.42
1:E:2175:VAL:HG12	1:E:2219:TYR:CE2	2.55	0.42
1:E:2514:ALA:HA	1:E:2517:ARG:HD2	2.01	0.42
1:E:3303:GLN:O	1:E:3306:ILE:HG22	2.19	0.42
1:E:4508:ALA:HB2	1:E:4578:HIS:HE1	1.83	0.42
1:E:4671:MET:O	1:E:4671:MET:HG2	2.19	0.42
1:F:557:TRP:HE3	1:F:558:LEU:HD23	1.84	0.42
1:F:2121:SER:O	1:F:2125:ILE:HG12	2.19	0.42
1:F:2289:TRP:CZ2	1:F:2387:ILE:HD12	2.54	0.42
1:F:2929:ILE:HG12	1:F:3006:LEU:HD13	2.01	0.42
1:F:3294:TRP:O	1:F:3298:LEU:HG	2.20	0.42
2:I:107:TRP:HD1	2:I:110:PRO:HB2	1.84	0.42
1:A:878:LEU:H	1:A:878:LEU:HD23	1.84	0.42
1:A:881:ILE:HA	1:A:884:ARG:HG2	2.00	0.42
1:A:2175:VAL:HG12	1:A:2219:TYR:CE2	2.55	0.42
1:A:4010:GLU:HG2	1:A:4120:LEU:HD13	2.02	0.42
1:A:4199:MET:HE2	1:A:4199:MET:HB2	1.77	0.42
1:C:430:ILE:HG23	1:C:504:ARG:HE	1.85	0.42
1:C:769:ARG:HA	1:C:774:PRO:HA	2.01	0.42
1:C:797:GLY:N	1:C:1622:CYS:O	2.52	0.42
1:C:850:LEU:O	1:C:1207:LEU:HD12	2.20	0.42
1:C:1750:PRO:HG3	1:C:2057:LEU:HD22	2.00	0.42
1:C:2193:ALA:O	1:C:2197:ARG:HG3	2.19	0.42
1:C:3636:GLU:HG3	1:C:3693:ILE:HG23	2.01	0.42
1:E:184:VAL:HG22	1:E:191:TYR:CD1	2.54	0.42
1:E:882:ARG:HH11	1:E:937:LEU:HA	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1990:GLU:CD	1:E:1990:GLU:H	2.22	0.42
1:E:1995:LEU:HD11	1:E:3623:TYR:CE1	2.54	0.42
1:E:2076:ASP:HB3	1:E:2079:LEU:HB3	2.01	0.42
1:E:3014:VAL:HG12	1:E:3095:TYR:HE1	1.83	0.42
1:E:3275:LEU:O	1:E:3279:ILE:HG13	2.19	0.42
1:E:3403:ILE:HD11	1:E:3556:VAL:HA	2.00	0.42
1:F:758:CYS:SG	1:F:767:SER:HB3	2.59	0.42
1:F:1595:VAL:HG23	1:F:1595:VAL:O	2.20	0.42
1:F:2175:VAL:HG12	1:F:2219:TYR:CE2	2.55	0.42
1:F:2514:ALA:HA	1:F:2517:ARG:HD2	2.01	0.42
1:F:3074:LEU:HD22	1:F:3147:VAL:HG23	2.00	0.42
1:F:3454:LYS:HA	1:F:3457:ARG:HG3	2.02	0.42
1:F:4596:LEU:HG	1:F:4600:LYS:HE3	2.00	0.42
1:A:430:ILE:HD13	1:A:500:GLU:HG3	2.01	0.42
1:A:480:ARG:NH2	1:A:3677:GLU:OE2	2.52	0.42
1:A:758:CYS:SG	1:A:767:SER:HB3	2.59	0.42
1:A:1113:MET:SD	1:A:1207:LEU:HD22	2.60	0.42
1:A:1419:TYR:CE2	1:A:1563:ASN:HB3	2.52	0.42
1:A:1990:GLU:H	1:A:1990:GLU:CD	2.22	0.42
1:A:2121:SER:O	1:A:2125:ILE:HG12	2.19	0.42
1:A:3007:PHE:HD1	1:A:3007:PHE:O	2.02	0.42
1:A:3014:VAL:HG12	1:A:3095:TYR:HE1	1.83	0.42
1:A:3275:LEU:O	1:A:3279:ILE:HG13	2.19	0.42
1:A:3769:ASN:OD1	1:A:3769:ASN:N	2.53	0.42
1:A:4035:ASP:OD1	1:A:4042:ILE:HG23	2.18	0.42
1:A:4483:ILE:O	1:A:4486:GLN:HG3	2.19	0.42
1:A:4559:VAL:HG22	1:A:4561:GLU:H	1.84	0.42
1:C:176:ARG:HE	1:C:181:LEU:HB3	1.85	0.42
1:C:879:GLU:O	1:C:883:GLU:HG2	2.19	0.42
1:C:1739:LEU:HD23	1:C:1739:LEU:H	1.84	0.42
1:C:1762:MET:HE2	1:C:1762:MET:HB2	1.87	0.42
1:C:2480:GLN:HE21	1:C:2484:LEU:HD11	1.84	0.42
1:E:4616:ILE:HD13	1:E:4616:ILE:HA	1.88	0.42
1:F:50:GLU:OE2	1:F:61:ASP:N	2.52	0.42
1:F:565:LEU:HD23	1:F:565:LEU:O	2.19	0.42
1:F:591:GLU:HA	1:F:631:LEU:HD11	2.01	0.42
1:F:1739:LEU:HD23	1:F:1739:LEU:H	1.84	0.42
1:F:2076:ASP:HB3	1:F:2079:LEU:HB3	2.01	0.42
1:F:2924:PHE:O	1:F:2928:LEU:HG	2.19	0.42
1:F:3217:ILE:HD11	1:F:3241:LEU:HD22	2.02	0.42
1:F:4662:ARG:HH21	1:F:4673:LYS:HD2	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:28:ILE:HD12	1:A:201:TRP:HE1	1.85	0.42
1:A:1177:LEU:HB2	1:A:1182:LEU:HD21	2.02	0.42
1:A:2983:SER:HA	1:A:3439:SER:HB3	2.00	0.42
1:A:3217:ILE:HD11	1:A:3241:LEU:HD22	2.02	0.42
1:A:3939:LEU:HD21	1:A:3980:MET:HE1	2.01	0.42
1:C:184:VAL:HG22	1:C:191:TYR:CD1	2.54	0.42
1:C:840:TYR:HE2	1:C:1086:ARG:HH12	1.68	0.42
1:C:1258:PHE:HB2	1:C:1303:ARG:HH21	1.84	0.42
1:C:2105:TYR:CZ	1:C:2160:LEU:HB2	2.55	0.42
1:C:2175:VAL:HG12	1:C:2219:TYR:CE2	2.55	0.42
1:C:3769:ASN:OD1	1:C:3769:ASN:N	2.53	0.42
1:C:4166:GLU:O	1:C:4169:ARG:HG2	2.19	0.42
1:E:941:LYS:NZ	1:E:944:LEU:HD11	2.34	0.42
1:E:1267:HIS:HB3	1:E:1295:ASN:N	2.31	0.42
1:E:1595:VAL:HG23	1:E:1595:VAL:O	2.20	0.42
1:E:1841:LYS:O	1:E:1845:GLN:HG2	2.20	0.42
1:E:2765:LYS:O	1:E:2768:GLU:HG3	2.19	0.42
1:E:3698:CYS:SG	1:E:3730:LEU:HD21	2.60	0.42
1:E:3935:ALA:O	1:E:3940:TRP:NE1	2.44	0.42
1:E:4042:ILE:HB	1:E:4047:PHE:HB2	2.00	0.42
1:E:4483:ILE:O	1:E:4486:GLN:HG3	2.19	0.42
1:F:430:ILE:HG23	1:F:504:ARG:HE	1.85	0.42
1:F:941:LYS:NZ	1:F:944:LEU:HD11	2.34	0.42
1:A:879:GLU:O	1:A:883:GLU:HG2	2.19	0.42
1:A:3286:ASN:HB3	1:A:3295:MET:HE3	2.01	0.42
1:A:3731:HIS:NE2	1:A:3775:LYS:HD2	2.34	0.42
2:B:107:TRP:HD1	2:B:110:PRO:HB2	1.84	0.42
1:C:1553:VAL:HG23	1:C:1554:PHE:N	2.35	0.42
1:C:1595:VAL:HG23	1:C:1595:VAL:O	2.20	0.42
1:C:2437:ILE:O	1:C:2464:LYS:HE3	2.18	0.42
1:C:3275:LEU:O	1:C:3279:ILE:HG13	2.19	0.42
1:C:4519:LYS:HE2	1:C:4519:LYS:HB3	1.88	0.42
1:E:438:LYS:HG3	1:E:439:LYS:HG2	2.01	0.42
1:E:565:LEU:O	1:E:565:LEU:HD23	2.19	0.42
1:E:2193:ALA:O	1:E:2197:ARG:HG3	2.19	0.42
1:E:2233:MET:O	1:E:2296:ARG:NH2	2.48	0.42
1:E:2455:MET:HE3	1:E:2456:SER:HB2	2.02	0.42
1:E:2579:LEU:N	1:E:2615:ARG:HH12	2.18	0.42
1:E:2924:PHE:O	1:E:2928:LEU:HG	2.19	0.42
1:F:694:ARG:NH1	1:F:720:ASP:OD1	2.53	0.42
1:F:840:TYR:HE2	1:F:1086:ARG:HH12	1.68	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:1553:VAL:HG23	1:F:1554:PHE:N	2.35	0.42
1:F:1995:LEU:HD11	1:F:3623:TYR:CE1	2.54	0.42
1:F:4135:ILE:HG12	1:F:4149:PHE:HE1	1.84	0.42
1:F:4500:MET:HG2	1:F:4585:CYS:SG	2.60	0.42
1:F:4941:LYS:HB3	1:F:4941:LYS:HE2	1.82	0.42
2:G:40:ALA:HB3	2:G:43:LYS:HB2	2.00	0.42
1:A:840:TYR:HE2	1:A:1086:ARG:HH12	1.68	0.42
1:A:1267:HIS:HB3	1:A:1295:ASN:N	2.31	0.42
1:A:3018:ILE:HD12	1:A:3095:TYR:HD1	1.85	0.42
1:A:3294:TRP:O	1:A:3298:LEU:HG	2.20	0.42
1:A:3698:CYS:SG	1:A:3730:LEU:HD21	2.60	0.42
1:C:694:ARG:NH1	1:C:720:ASP:OD1	2.53	0.42
1:C:1791:LYS:HG3	1:C:1795:MET:SD	2.59	0.42
1:C:2121:SER:O	1:C:2125:ILE:HG12	2.19	0.42
1:C:3085:GLN:HE21	1:C:3089:VAL:HG12	1.84	0.42
1:C:3365:LEU:O	1:C:3369:TYR:HB2	2.19	0.42
1:C:3909:ILE:HG21	1:C:3969:GLU:HB3	2.01	0.42
1:C:3993:THR:O	1:C:3997:GLN:HG3	2.20	0.42
1:E:176:ARG:HE	1:E:181:LEU:HB3	1.85	0.42
1:E:468:GLU:OE1	1:E:468:GLU:N	2.41	0.42
1:E:1739:LEU:HD23	1:E:1739:LEU:H	1.84	0.42
1:E:2099:ARG:O	1:E:2103:LYS:NZ	2.34	0.42
1:E:2480:GLN:HE21	1:E:2484:LEU:HD11	1.84	0.42
1:E:3007:PHE:O	1:E:3007:PHE:HD1	2.02	0.42
1:E:3482:PRO:HD2	1:E:3527:MET:SD	2.60	0.42
1:E:3939:LEU:HD21	1:E:3980:MET:HE1	2.01	0.42
1:E:4010:GLU:HG2	1:E:4120:LEU:HD13	2.02	0.42
1:E:4508:ALA:O	1:E:4511:ILE:HG22	2.20	0.42
1:E:4633:VAL:O	1:E:4636:THR:HG22	2.20	0.42
1:F:238:HIS:HB2	1:F:242:ASP:N	2.35	0.42
1:F:626:ARG:HH21	1:F:2131:VAL:HG11	1.84	0.42
1:F:3698:CYS:SG	1:F:3730:LEU:HD21	2.60	0.42
1:F:4648:VAL:O	1:F:4652:VAL:HG12	2.19	0.42
1:A:674:TYR:HE1	1:A:756:SER:HB2	1.85	0.42
1:A:850:LEU:O	1:A:1207:LEU:HD12	2.20	0.42
1:A:1841:LYS:O	1:A:1845:GLN:HG2	2.20	0.42
1:A:2193:ALA:O	1:A:2197:ARG:HG3	2.19	0.42
1:A:3482:PRO:HD2	1:A:3527:MET:SD	2.60	0.42
1:A:3727:GLN:O	1:A:3731:HIS:CB	2.66	0.42
1:A:3909:ILE:HG21	1:A:3969:GLU:HB3	2.01	0.42
1:A:4500:MET:HG2	1:A:4585:CYS:SG	2.60	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1113:MET:SD	1:C:1207:LEU:HD22	2.60	0.42
1:C:1802:GLU:HA	1:C:1805:LEU:HG	2.02	0.42
1:C:2580:ARG:HG2	1:C:2583:MET:HE3	2.01	0.42
1:C:3226:ARG:NH1	1:C:3286:ASN:HA	2.35	0.42
1:C:3322:MET:HB3	1:C:3368:PHE:CE2	2.55	0.42
1:C:3482:PRO:HD2	1:C:3527:MET:SD	2.60	0.42
2:D:109:THR:OG1	2:D:110:PRO:HD3	2.20	0.42
1:E:50:GLU:OE2	1:E:61:ASP:N	2.52	0.42
1:E:238:HIS:HB2	1:E:242:ASP:N	2.35	0.42
1:E:410:HIS:O	1:E:414:ARG:HG2	2.20	0.42
1:E:747:HIS:CE1	1:E:750:ARG:HG2	2.55	0.42
1:E:769:ARG:HA	1:E:774:PRO:HA	2.01	0.42
1:E:840:TYR:HE2	1:E:1086:ARG:HH12	1.68	0.42
1:E:934:GLN:CD	2:I:99:ARG:HH22	2.22	0.42
1:E:1553:VAL:HG23	1:E:1554:PHE:N	2.35	0.42
1:E:1791:LYS:HG3	1:E:1795:MET:SD	2.59	0.42
1:E:2258:GLU:N	1:E:2259:PRO:HD2	2.35	0.42
1:E:2289:TRP:CZ2	1:E:2387:ILE:HD12	2.54	0.42
1:E:3018:ILE:HD12	1:E:3095:TYR:HD1	1.85	0.42
1:E:3322:MET:HB3	1:E:3368:PHE:CE2	2.55	0.42
1:F:410:HIS:O	1:F:414:ARG:HG2	2.20	0.42
1:F:480:ARG:NH2	1:F:3677:GLU:OE2	2.52	0.42
1:F:850:LEU:O	1:F:1207:LEU:HD12	2.20	0.42
1:F:1166:VAL:HG22	1:F:1173:MET:HB2	2.02	0.42
1:F:3283:ILE:O	1:F:3287:LEU:HG	2.20	0.42
1:F:3636:GLU:HG3	1:F:3693:ILE:HG23	2.01	0.42
1:F:4168:LYS:HE2	1:F:4168:LYS:HB3	1.86	0.42
1:A:184:VAL:HG22	1:A:191:TYR:CD1	2.54	0.41
1:A:2132:ARG:HG2	1:A:2133:MET:H	1.85	0.41
1:A:2258:GLU:N	1:A:2259:PRO:HD2	2.35	0.41
1:A:4508:ALA:O	1:A:4511:ILE:HG22	2.20	0.41
1:C:238:HIS:HB2	1:C:242:ASP:N	2.35	0.41
1:C:410:HIS:O	1:C:414:ARG:HG2	2.20	0.41
1:C:882:ARG:HH11	1:C:937:LEU:HA	1.84	0.41
1:C:1990:GLU:H	1:C:1990:GLU:CD	2.22	0.41
1:C:2426:ILE:O	1:C:2475:TYR:OH	2.37	0.41
1:C:4500:MET:HG2	1:C:4585:CYS:SG	2.60	0.41
1:C:4792:TYR:HH	1:C:4815:HIS:CE1	2.38	0.41
2:D:107:TRP:HD1	2:D:110:PRO:HB2	1.84	0.41
1:E:64:ILE:HA	1:E:123:HIS:CE1	2.55	0.41
1:E:626:ARG:HH21	1:E:2131:VAL:HG11	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1113:MET:SD	1:E:1207:LEU:HD22	2.60	0.41
1:E:1814:THR:HG22	1:E:1816:GLU:H	1.85	0.41
1:E:2581:PRO:HD2	1:E:2629:PHE:HD2	1.85	0.41
1:E:3226:ARG:NH1	1:E:3286:ASN:HA	2.35	0.41
1:E:3454:LYS:HA	1:E:3457:ARG:HG3	2.02	0.41
1:E:3636:GLU:HG3	1:E:3693:ILE:HG23	2.02	0.41
1:E:3993:THR:O	1:E:3997:GLN:HG3	2.20	0.41
1:F:390:LYS:HA	1:F:390:LYS:HD3	1.89	0.41
1:F:1114:ARG:HG2	1:F:1138:ASP:HB2	2.01	0.41
1:F:1791:LYS:HG3	1:F:1795:MET:SD	2.59	0.41
1:F:1841:LYS:O	1:F:1845:GLN:HG2	2.20	0.41
1:F:4505:LEU:HD13	1:F:4744:ASP:HB3	2.02	0.41
1:F:4633:VAL:O	1:F:4636:THR:HG22	2.20	0.41
1:A:133:LEU:O	1:A:145:PHE:HB3	2.19	0.41
1:A:626:ARG:HH21	1:A:2131:VAL:HG11	1.84	0.41
1:A:2753:GLN:NE2	1:A:2762:LEU:O	2.49	0.41
1:A:2974:PHE:HD1	1:A:2979:LEU:HD12	1.86	0.41
1:A:3638:LYS:HE2	1:A:3638:LYS:HB3	1.89	0.41
1:A:3935:ALA:O	1:A:3940:TRP:NE1	2.44	0.41
2:B:104:TYR:HD1	2:B:106:PRO:HD3	1.84	0.41
1:C:64:ILE:HA	1:C:123:HIS:CE1	2.55	0.41
1:C:1114:ARG:HG2	1:C:1138:ASP:HB2	2.01	0.41
1:C:1166:VAL:HG22	1:C:1173:MET:HB2	2.02	0.41
1:C:4135:ILE:HG12	1:C:4149:PHE:HE1	1.84	0.41
1:C:4648:VAL:O	1:C:4652:VAL:HG12	2.19	0.41
1:E:28:ILE:HD12	1:E:201:TRP:HE1	1.85	0.41
1:E:728:ASP:OD1	1:E:731:HIS:N	2.41	0.41
1:E:1114:ARG:HG2	1:E:1138:ASP:HB2	2.01	0.41
1:E:1144:ARG:H	1:E:1144:ARG:HG2	1.73	0.41
1:E:2132:ARG:HG2	1:E:2133:MET:H	1.85	0.41
1:E:3909:ILE:HG21	1:E:3969:GLU:HB3	2.01	0.41
1:F:241:MET:HE2	1:F:241:MET:HB2	1.78	0.41
1:F:674:TYR:HE1	1:F:756:SER:HB2	1.85	0.41
1:F:828:PRO:HG2	1:F:1033:VAL:HG21	2.02	0.41
1:F:1113:MET:SD	1:F:1207:LEU:HD22	2.60	0.41
1:F:1985:CYS:SG	1:F:1992:ARG:NH1	2.94	0.41
1:F:3482:PRO:HD2	1:F:3527:MET:SD	2.60	0.41
1:F:3769:ASN:OD1	1:F:3769:ASN:N	2.53	0.41
2:G:18:LEU:HD23	2:G:18:LEU:HA	1.87	0.41
2:I:90:THR:HG23	2:I:124:THR:HA	2.01	0.41
2:I:109:THR:OG1	2:I:110:PRO:HD3	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:557:TRP:HE3	1:A:558:LEU:HD23	1.84	0.41
1:A:694:ARG:NH1	1:A:720:ASP:OD1	2.53	0.41
1:A:769:ARG:HA	1:A:774:PRO:HA	2.01	0.41
1:A:1039:ASP:OD1	1:A:1039:ASP:N	2.52	0.41
1:A:1595:VAL:O	1:A:1595:VAL:HG23	2.20	0.41
1:A:1985:CYS:SG	1:A:1992:ARG:NH1	2.94	0.41
1:A:2459:PHE:CE1	1:A:2464:LYS:HG3	2.56	0.41
1:A:2998:LYS:HG3	1:A:3002:MET:HE2	2.01	0.41
1:A:3365:LEU:O	1:A:3369:TYR:HB2	2.19	0.41
1:A:3993:THR:O	1:A:3997:GLN:HG3	2.20	0.41
1:C:14:LEU:HD11	1:C:214:VAL:HG21	2.01	0.41
1:C:1177:LEU:HB2	1:C:1182:LEU:HD21	2.02	0.41
1:C:1428:TYR:CE1	1:C:1510:CYS:HB2	2.54	0.41
1:C:3369:TYR:HE2	1:C:3465:ILE:HA	1.85	0.41
1:C:4751:THR:O	1:C:4755:ILE:HG13	2.21	0.41
1:E:674:TYR:HE1	1:E:756:SER:HB2	1.85	0.41
1:E:1165:MET:HA	1:E:1165:MET:HE3	2.03	0.41
1:E:1999:HIS:CG	1:E:3627:TRP:HD1	2.39	0.41
1:E:3217:ILE:HD11	1:E:3241:LEU:HD22	2.02	0.41
1:E:3426:ASN:O	1:E:3430:LEU:HG	2.21	0.41
1:E:3832:ASP:N	1:E:3832:ASP:OD1	2.53	0.41
1:F:882:ARG:HH11	1:F:937:LEU:HA	1.84	0.41
1:F:2233:MET:O	1:F:2296:ARG:NH2	2.48	0.41
1:F:3085:GLN:HE21	1:F:3089:VAL:HG12	1.84	0.41
1:F:3275:LEU:O	1:F:3279:ILE:HG13	2.19	0.41
1:F:3322:MET:HB3	1:F:3368:PHE:CE2	2.55	0.41
1:F:3369:TYR:HE2	1:F:3465:ILE:HA	1.85	0.41
2:G:90:THR:HG23	2:G:124:THR:HA	2.01	0.41
1:A:64:ILE:HA	1:A:123:HIS:CE1	2.55	0.41
1:A:1303:ARG:HD3	1:A:1446:ILE:HD11	2.03	0.41
1:A:1553:VAL:HG23	1:A:1554:PHE:N	2.35	0.41
1:A:1899:GLU:HA	1:A:1902:LYS:HG2	2.03	0.41
1:A:2579:LEU:N	1:A:2615:ARG:HH12	2.18	0.41
1:A:3283:ILE:O	1:A:3287:LEU:HG	2.20	0.41
1:A:3482:PRO:O	1:A:3486:GLU:HG3	2.20	0.41
1:A:4030:THR:HG23	1:A:4054:HIS:HE1	1.86	0.41
1:C:747:HIS:CE1	1:C:750:ARG:HG2	2.55	0.41
1:C:2929:ILE:HG12	1:C:3006:LEU:HD13	2.01	0.41
1:C:3007:PHE:O	1:C:3007:PHE:HD1	2.02	0.41
1:C:3018:ILE:HD12	1:C:3095:TYR:HD1	1.85	0.41
1:C:3254:GLU:OE1	1:C:3254:GLU:N	2.47	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3426:ASN:O	1:C:3430:LEU:HG	2.21	0.41
1:C:3875:ASP:O	1:C:3879:ARG:HG3	2.21	0.41
1:C:4483:ILE:O	1:C:4486:GLN:HG3	2.19	0.41
1:C:4505:LEU:HD13	1:C:4744:ASP:HB3	2.02	0.41
1:E:1303:ARG:O	1:E:1590:GLN:N	2.38	0.41
1:E:1899:GLU:HA	1:E:1902:LYS:HG2	2.03	0.41
1:E:2459:PHE:CE1	1:E:2464:LYS:HG3	2.56	0.41
1:E:2918:LYS:HA	1:E:2999:GLU:OE2	2.21	0.41
1:E:4751:THR:O	1:E:4755:ILE:HG13	2.21	0.41
1:F:176:ARG:HE	1:F:181:LEU:HB3	1.85	0.41
1:F:732:LEU:HB3	1:F:779:PHE:CZ	2.54	0.41
1:F:878:LEU:HD23	1:F:878:LEU:H	1.84	0.41
1:F:1258:PHE:HB2	1:F:1303:ARG:HH21	1.84	0.41
1:F:1516:SER:O	1:F:1533:GLN:NE2	2.38	0.41
1:F:1801:LYS:HB3	1:F:1801:LYS:HE3	1.71	0.41
1:F:1990:GLU:H	1:F:1990:GLU:CD	2.22	0.41
1:F:2105:TYR:CZ	1:F:2160:LEU:HB2	2.55	0.41
1:F:2202:PHE:O	1:F:2205:ILE:HG12	2.21	0.41
1:F:2581:PRO:HD2	1:F:2629:PHE:HD2	1.85	0.41
1:F:3226:ARG:NH1	1:F:3286:ASN:HA	2.35	0.41
1:F:3875:ASP:O	1:F:3879:ARG:HG3	2.21	0.41
1:A:1814:THR:HG22	1:A:1816:GLU:H	1.85	0.41
1:A:2480:GLN:HE21	1:A:2484:LEU:HD11	1.84	0.41
1:A:2581:PRO:HD2	1:A:2629:PHE:HD2	1.85	0.41
1:A:3129:CYS:HB3	1:A:3161:PHE:CE1	2.51	0.41
1:A:3226:ARG:NH1	1:A:3286:ASN:HA	2.35	0.41
1:A:3254:GLU:OE1	1:A:3254:GLU:N	2.47	0.41
1:A:4633:VAL:O	1:A:4636:THR:HG22	2.20	0.41
2:B:109:THR:OG1	2:B:110:PRO:HD3	2.20	0.41
1:C:744:PRO:HD3	1:C:776:GLN:HE21	1.85	0.41
2:D:52:THR:HA	2:D:71:ARG:HH11	1.86	0.41
1:E:2105:TYR:CZ	1:E:2160:LEU:HB2	2.55	0.41
1:E:3769:ASN:OD1	1:E:3769:ASN:N	2.53	0.41
1:F:251:GLU:HG3	1:F:252:HIS:ND1	2.36	0.41
1:F:299:HIS:HB3	1:F:302:THR:HG22	2.02	0.41
1:F:908:ARG:NE	2:G:104:TYR:HB3	2.36	0.41
1:F:3046:LYS:HD2	1:F:3046:LYS:O	2.21	0.41
1:F:3488:ILE:HA	1:F:3550:VAL:HG13	2.03	0.41
1:F:4010:GLU:HG2	1:F:4120:LEU:HD13	2.02	0.41
1:F:4483:ILE:O	1:F:4486:GLN:HG3	2.19	0.41
1:F:4508:ALA:O	1:F:4511:ILE:HG22	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:104:TYR:HD1	2:G:106:PRO:HD3	1.84	0.41
1:A:176:ARG:HE	1:A:181:LEU:HB3	1.85	0.41
1:A:921:PHE:HA	1:A:924:LEU:HB2	2.02	0.41
1:A:970:TYR:HB2	1:A:971:GLN:H	1.70	0.41
1:A:2455:MET:HE3	1:A:2456:SER:HB2	2.03	0.41
1:A:2996:SER:O	1:A:2999:GLU:HB2	2.21	0.41
1:A:3240:MET:HE2	1:A:3240:MET:HA	2.03	0.41
1:A:3426:ASN:O	1:A:3430:LEU:HG	2.21	0.41
1:A:3636:GLU:HG3	1:A:3693:ILE:HG23	2.01	0.41
1:A:4140:SER:O	1:A:4140:SER:OG	2.39	0.41
1:C:28:ILE:HD12	1:C:201:TRP:HE1	1.85	0.41
1:C:732:LEU:HB3	1:C:779:PHE:CZ	2.54	0.41
1:C:1841:LYS:O	1:C:1845:GLN:HG2	2.20	0.41
1:C:3383:TRP:HH2	1:C:3394:LEU:HD23	1.85	0.41
1:C:3832:ASP:N	1:C:3832:ASP:OD1	2.54	0.41
1:C:3870:ILE:O	1:C:3874:VAL:HG23	2.20	0.41
1:C:4164:VAL:HG21	1:C:4199:MET:HG2	2.03	0.41
1:E:3369:TYR:HE2	1:E:3465:ILE:HA	1.85	0.41
1:E:3875:ASP:O	1:E:3879:ARG:HG3	2.21	0.41
1:F:161:THR:HG23	1:F:186:VAL:HG22	2.03	0.41
1:F:747:HIS:CE1	1:F:750:ARG:HG2	2.55	0.41
1:F:1588:HIS:CE1	1:F:1590:GLN:HE21	2.39	0.41
1:F:1814:THR:HG22	1:F:1816:GLU:H	1.85	0.41
1:F:1899:GLU:HA	1:F:1902:LYS:HG2	2.03	0.41
1:F:2480:GLN:HE21	1:F:2484:LEU:HD11	1.84	0.41
1:F:2859:LEU:HD13	1:F:2866:ASN:HA	2.02	0.41
1:F:3240:MET:HE2	1:F:3240:MET:HA	2.03	0.41
1:F:4164:VAL:HG21	1:F:4199:MET:HG2	2.03	0.41
1:A:56:LYS:HE2	1:A:56:LYS:HB2	1.88	0.41
1:A:251:GLU:HG3	1:A:252:HIS:ND1	2.36	0.41
1:A:430:ILE:HG23	1:A:504:ARG:HE	1.84	0.41
1:A:1258:PHE:HB2	1:A:1303:ARG:HH21	1.84	0.41
1:A:3369:TYR:HE2	1:A:3465:ILE:HA	1.85	0.41
1:A:4164:VAL:HG21	1:A:4199:MET:HG2	2.03	0.41
1:A:4505:LEU:HD13	1:A:4744:ASP:HB3	2.02	0.41
1:C:263:GLU:OE2	1:C:388:GLN:NE2	2.40	0.41
1:C:299:HIS:HB3	1:C:302:THR:HG22	2.02	0.41
1:C:1420:LEU:O	1:C:1423:THR:HG22	2.21	0.41
1:C:1686:LEU:HD13	1:C:1707:LEU:HD13	2.02	0.41
1:C:1995:LEU:HD11	1:C:3623:TYR:CE1	2.54	0.41
1:C:2996:SER:O	1:C:2999:GLU:HB2	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3283:ILE:O	1:C:3287:LEU:HG	2.20	0.41
1:C:3943:VAL:HG23	1:C:3977:MET:SD	2.61	0.41
1:C:3998:MET:O	1:C:4002:LEU:HG	2.21	0.41
1:C:4188:PHE:CE1	1:C:4914:LEU:HD22	2.56	0.41
1:C:4508:ALA:O	1:C:4511:ILE:HG22	2.20	0.41
1:E:557:TRP:HE3	1:E:558:LEU:HD23	1.84	0.41
1:E:1097:LYS:HD2	1:E:1097:LYS:N	2.36	0.41
1:E:2423:ARG:NH2	1:E:2475:TYR:O	2.44	0.41
1:E:4140:SER:O	1:E:4140:SER:OG	2.39	0.41
1:E:4164:VAL:HG21	1:E:4199:MET:HG2	2.03	0.41
1:F:28:ILE:HD12	1:F:201:TRP:HE1	1.85	0.41
1:F:2258:GLU:N	1:F:2259:PRO:HD2	2.35	0.41
1:F:2276:CYS:HB2	1:F:2290:ASN:ND2	2.36	0.41
1:F:2621:CYS:HA	1:F:2676:PRO:HG3	2.02	0.41
1:F:3383:TRP:HH2	1:F:3394:LEU:HD23	1.85	0.41
1:F:4501:ARG:HA	1:F:4501:ARG:HD2	1.93	0.41
1:A:50:GLU:OE2	1:A:61:ASP:N	2.52	0.41
1:A:744:PRO:HD3	1:A:776:GLN:HE21	1.85	0.41
1:A:928:GLU:OE1	1:A:928:GLU:N	2.49	0.41
1:A:1097:LYS:HD2	1:A:1097:LYS:N	2.36	0.41
1:A:1114:ARG:HG2	1:A:1138:ASP:HB2	2.01	0.41
1:A:1305:SER:N	1:A:1588:HIS:O	2.54	0.41
1:A:3943:VAL:HG23	1:A:3977:MET:SD	2.61	0.41
1:A:4044:LYS:HB2	1:A:4075:GLU:OE2	2.21	0.41
1:C:674:TYR:HE1	1:C:756:SER:HB2	1.85	0.41
1:C:1267:HIS:HB3	1:C:1295:ASN:N	2.31	0.41
1:C:1814:THR:HG22	1:C:1816:GLU:H	1.85	0.41
1:C:1899:GLU:HA	1:C:1902:LYS:HG2	2.03	0.41
1:C:1967:PRO:O	1:C:1971:GLN:HG3	2.21	0.41
1:C:2076:ASP:HB3	1:C:2079:LEU:HB3	2.01	0.41
1:C:2258:GLU:N	1:C:2259:PRO:HD2	2.35	0.41
1:C:3029:VAL:HG12	1:C:3033:HIS:NE2	2.36	0.41
1:C:3482:PRO:O	1:C:3486:GLU:HG3	2.20	0.41
1:C:3811:ASN:O	1:C:3814:GLU:HG2	2.21	0.41
1:C:4020:LEU:HD12	1:C:4124:VAL:HG12	2.03	0.41
1:C:4633:VAL:O	1:C:4636:THR:HG22	2.20	0.41
1:E:299:HIS:HB3	1:E:302:THR:HG22	2.02	0.41
1:E:641:ASP:O	1:E:642:LEU:HB3	2.21	0.41
1:E:705:PRO:HG3	1:E:857:LEU:HD12	2.03	0.41
1:E:828:PRO:HG2	1:E:1033:VAL:HG21	2.02	0.41
1:E:921:PHE:HB2	1:E:929:ARG:NH1	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:1166:VAL:HG22	1:E:1173:MET:HB2	2.02	0.41
1:E:1802:GLU:HA	1:E:1805:LEU:HG	2.02	0.41
1:E:1985:CYS:SG	1:E:1992:ARG:NH1	2.94	0.41
1:E:2065:MET:HE2	1:E:2065:MET:HB2	1.99	0.41
1:E:2279:LEU:HB3	1:E:2284:TYR:HB2	2.03	0.41
1:E:3283:ILE:O	1:E:3287:LEU:HG	2.20	0.41
1:E:3294:TRP:O	1:E:3298:LEU:HG	2.19	0.41
1:E:4044:LYS:HB2	1:E:4075:GLU:OE2	2.21	0.41
1:E:4199:MET:HE2	1:E:4199:MET:HB2	1.76	0.41
1:E:4521:SER:O	1:E:4556:VAL:N	2.54	0.41
1:F:261:HIS:HD2	1:F:263:GLU:HG3	1.86	0.41
1:F:1097:LYS:HD2	1:F:1097:LYS:N	2.36	0.41
1:F:2348:GLU:O	1:F:2352:ILE:HG12	2.21	0.41
1:F:3029:VAL:HG12	1:F:3033:HIS:NE2	2.36	0.41
1:F:3832:ASP:N	1:F:3832:ASP:OD1	2.54	0.41
1:F:3998:MET:O	1:F:4002:LEU:HG	2.21	0.41
1:A:641:ASP:O	1:A:642:LEU:HB3	2.21	0.41
1:A:1428:TYR:CE1	1:A:1510:CYS:HB2	2.54	0.41
1:A:2105:TYR:CZ	1:A:2160:LEU:HB2	2.55	0.41
1:A:2210:GLN:NE2	1:A:2244:ALA:O	2.45	0.41
1:A:3322:MET:HB3	1:A:3368:PHE:CE2	2.55	0.41
1:A:4645:ASP:OD1	1:A:4645:ASP:N	2.44	0.41
1:A:4751:THR:O	1:A:4755:ILE:HG13	2.21	0.41
1:A:4756:LEU:O	1:A:4759:VAL:HG12	2.21	0.41
2:B:52:THR:HA	2:B:71:ARG:HH11	1.85	0.41
1:C:251:GLU:HG3	1:C:252:HIS:ND1	2.36	0.41
1:C:438:LYS:HG3	1:C:439:LYS:HG2	2.01	0.41
1:C:556:ASP:OD1	1:C:556:ASP:N	2.54	0.41
1:C:601:LEU:HD13	1:C:610:VAL:HB	2.03	0.41
1:C:1097:LYS:N	1:C:1097:LYS:HD2	2.36	0.41
1:C:1985:CYS:SG	1:C:1992:ARG:NH1	2.94	0.41
1:C:2154:PHE:CE1	1:C:2205:ILE:HD13	2.56	0.41
1:C:2202:PHE:O	1:C:2205:ILE:HG12	2.21	0.41
1:C:2276:CYS:HB2	1:C:2290:ASN:ND2	2.36	0.41
1:C:2348:GLU:O	1:C:2352:ILE:HG12	2.21	0.41
1:C:2579:LEU:N	1:C:2615:ARG:HH12	2.18	0.41
1:C:2974:PHE:HD1	1:C:2979:LEU:HD12	1.86	0.41
1:C:3124:ASP:N	1:C:3124:ASP:OD1	2.54	0.41
1:C:3698:CYS:SG	1:C:3730:LEU:HD21	2.60	0.41
1:C:4662:ARG:HE	1:C:4662:ARG:HB3	1.68	0.41
1:C:4756:LEU:O	1:C:4759:VAL:HG12	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:D:104:TYR:HD1	2:D:106:PRO:HD3	1.84	0.41
1:E:56:LYS:HE2	1:E:56:LYS:HB2	1.88	0.41
1:E:161:THR:HG23	1:E:186:VAL:HG22	2.03	0.41
1:E:173:GLU:HA	1:F:3938:ARG:NH1	2.35	0.41
1:E:198:ASN:N	1:E:198:ASN:OD1	2.54	0.41
1:E:430:ILE:HD11	1:E:501:CYS:HB3	2.02	0.41
1:E:612:ASP:OD1	1:E:1657:HIS:ND1	2.50	0.41
1:E:694:ARG:NH1	1:E:720:ASP:OD1	2.53	0.41
1:E:1177:LEU:HB2	1:E:1182:LEU:HD21	2.02	0.41
1:E:2154:PHE:CE1	1:E:2205:ILE:HD13	2.56	0.41
1:E:2584:MET:HE2	1:E:2588:LEU:HG	2.03	0.41
1:E:3129:CYS:HB3	1:E:3161:PHE:CE1	2.51	0.41
1:E:3214:MET:SD	1:E:3271:HIS:NE2	2.94	0.41
1:E:3369:TYR:HA	1:E:3372:LEU:HB3	2.03	0.41
1:E:3386:GLU:HB3	1:E:3535:ASN:ND2	2.36	0.41
1:E:3482:PRO:O	1:E:3486:GLU:HG3	2.21	0.41
1:E:3998:MET:O	1:E:4002:LEU:HG	2.21	0.41
1:E:4030:THR:HG23	1:E:4054:HIS:HE1	1.86	0.41
1:E:4756:LEU:O	1:E:4759:VAL:HG12	2.21	0.41
1:F:744:PRO:HD3	1:F:776:GLN:HE21	1.85	0.41
1:F:921:PHE:HB2	1:F:929:ARG:NH1	2.36	0.41
1:F:928:GLU:OE1	1:F:928:GLU:N	2.49	0.41
1:F:1273:ILE:HB	1:F:1287:GLN:OE1	2.21	0.41
1:F:1303:ARG:HD3	1:F:1446:ILE:HD11	2.03	0.41
1:F:1802:GLU:HA	1:F:1805:LEU:HG	2.02	0.41
1:F:2143:ARG:HE	1:F:2143:ARG:HB3	1.67	0.41
1:F:2392:ALA:HB2	1:F:2463:HIS:CD2	2.56	0.41
1:F:2459:PHE:CE1	1:F:2464:LYS:HG3	2.56	0.41
1:F:2584:MET:HE2	1:F:2588:LEU:HG	2.03	0.41
1:F:2974:PHE:HD1	1:F:2979:LEU:HD12	1.86	0.41
1:F:3018:ILE:HD12	1:F:3095:TYR:HD1	1.85	0.41
1:F:3482:PRO:O	1:F:3486:GLU:HG3	2.21	0.41
1:F:3638:LYS:HE2	1:F:3638:LYS:HB3	1.89	0.41
1:F:3870:ILE:O	1:F:3874:VAL:HG23	2.21	0.41
1:F:3993:THR:O	1:F:3997:GLN:HG3	2.20	0.41
1:F:4030:THR:HG23	1:F:4054:HIS:HE1	1.86	0.41
1:F:4188:PHE:CE1	1:F:4914:LEU:HD22	2.56	0.41
2:G:109:THR:OG1	2:G:110:PRO:HD3	2.20	0.41
1:A:747:HIS:CE1	1:A:750:ARG:HG2	2.55	0.41
1:A:2392:ALA:HB2	1:A:2463:HIS:CD2	2.56	0.41
1:A:2426:ILE:O	1:A:2475:TYR:OH	2.37	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2554:LEU:HG	1:A:2568:ILE:HD13	2.03	0.41
1:A:2918:LYS:HA	1:A:2999:GLU:OE2	2.21	0.41
1:A:3046:LYS:HD2	1:A:3046:LYS:O	2.21	0.41
1:A:3214:MET:SD	1:A:3271:HIS:NE2	2.94	0.41
1:A:3616:VAL:O	1:A:3620:LEU:HG	2.21	0.41
1:C:641:ASP:O	1:C:642:LEU:HB3	2.21	0.41
1:C:828:PRO:HG2	1:C:1033:VAL:HG21	2.02	0.41
1:C:1305:SER:N	1:C:1588:HIS:O	2.54	0.41
1:C:2132:ARG:HG2	1:C:2133:MET:H	1.85	0.41
1:C:2224:SER:HB2	1:C:2239:LEU:HB2	2.03	0.41
1:C:2392:ALA:HB2	1:C:2463:HIS:CD2	2.56	0.41
1:C:2549:HIS:H	1:C:2549:HIS:HD1	1.68	0.41
1:C:2918:LYS:HA	1:C:2999:GLU:OE2	2.21	0.41
1:E:251:GLU:HG3	1:E:252:HIS:ND1	2.36	0.41
1:E:1686:LEU:HD13	1:E:1707:LEU:HD13	2.02	0.41
1:E:2775:LYS:HE3	1:E:2775:LYS:HB3	1.89	0.41
1:E:2974:PHE:HD1	1:E:2979:LEU:HD12	1.86	0.41
1:F:14:LEU:HD11	1:F:214:VAL:HG21	2.01	0.41
1:F:2132:ARG:HG2	1:F:2133:MET:H	1.85	0.41
1:F:2918:LYS:HA	1:F:2999:GLU:OE2	2.21	0.41
1:F:3124:ASP:OD1	1:F:3124:ASP:N	2.54	0.41
1:F:3369:TYR:HA	1:F:3372:LEU:HB3	2.03	0.41
1:F:4579:THR:HG1	1:F:4732:HIS:CD2	2.36	0.41
2:I:48:VAL:HG12	2:I:49:ALA:H	1.86	0.41
2:I:104:TYR:HD1	2:I:106:PRO:HD3	1.84	0.41
1:A:173:GLU:HA	1:E:3938:ARG:NH1	2.36	0.40
1:A:541:ILE:HD11	1:A:574:VAL:HG13	2.04	0.40
1:A:601:LEU:HD13	1:A:610:VAL:HB	2.03	0.40
1:A:1420:LEU:O	1:A:1423:THR:HG22	2.21	0.40
1:A:1749:LEU:HD23	1:A:1844:LEU:HD12	2.03	0.40
1:A:1838:GLU:H	1:A:1838:GLU:CD	2.25	0.40
1:A:2099:ARG:O	1:A:2103:LYS:NZ	2.34	0.40
1:A:2621:CYS:HA	1:A:2676:PRO:HG3	2.02	0.40
1:A:3029:VAL:HG12	1:A:3033:HIS:NE2	2.36	0.40
1:A:3386:GLU:HB3	1:A:3535:ASN:ND2	2.36	0.40
1:C:626:ARG:HH21	1:C:2131:VAL:HG11	1.84	0.40
1:C:908:ARG:NE	2:D:104:TYR:HB3	2.35	0.40
1:C:2680:MET:HE3	1:C:2680:MET:HA	2.02	0.40
1:C:2922:TYR:CD1	1:C:3002:MET:HE1	2.56	0.40
1:C:3046:LYS:HD2	1:C:3046:LYS:O	2.21	0.40
1:E:514:PHE:HD2	1:E:523:GLY:HA2	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:921:PHE:HA	1:E:924:LEU:HB2	2.02	0.40
1:E:1978:PHE:CZ	1:E:1995:LEU:HD23	2.56	0.40
1:E:2481:ASP:OD1	1:E:2482:PHE:N	2.54	0.40
1:E:3105:SER:HB3	1:E:3156:GLU:HG2	2.04	0.40
1:E:3273:ASN:ND2	1:E:3313:LEU:HG	2.37	0.40
1:E:4020:LEU:HD12	1:E:4124:VAL:HG12	2.03	0.40
1:F:612:ASP:OD1	1:F:1657:HIS:ND1	2.50	0.40
1:F:921:PHE:HA	1:F:924:LEU:HB2	2.02	0.40
1:F:1177:LEU:HB2	1:F:1182:LEU:HD21	2.02	0.40
1:F:1838:GLU:CD	1:F:1838:GLU:H	2.25	0.40
1:F:1967:PRO:O	1:F:1971:GLN:HG3	2.21	0.40
1:F:1999:HIS:CG	1:F:3627:TRP:HD1	2.39	0.40
1:F:2455:MET:HE3	1:F:2456:SER:HB2	2.03	0.40
1:F:2554:LEU:HG	1:F:2568:ILE:HD13	2.03	0.40
1:F:2577:GLY:O	1:F:2615:ARG:HD2	2.21	0.40
1:F:3007:PHE:HA	1:F:3035:LEU:HD13	2.03	0.40
1:F:3276:LEU:HD22	1:F:3309:VAL:HG11	2.04	0.40
1:F:3348:SER:HA	1:F:3351:GLU:HB2	2.04	0.40
1:F:4756:LEU:O	1:F:4759:VAL:HG12	2.21	0.40
2:G:52:THR:HA	2:G:71:ARG:HH11	1.85	0.40
2:I:64:LYS:HE2	2:I:64:LYS:HB3	1.94	0.40
1:A:314:LEU:O	1:A:315:LEU:HD23	2.22	0.40
1:A:514:PHE:HD2	1:A:523:GLY:HA2	1.86	0.40
1:A:555:LEU:HD23	1:A:555:LEU:HA	1.93	0.40
1:A:1257:GLN:HB2	1:A:1596:LEU:HD21	2.03	0.40
1:A:1967:PRO:O	1:A:1971:GLN:HG3	2.21	0.40
1:A:1999:HIS:CG	1:A:3627:TRP:HD1	2.39	0.40
1:A:2202:PHE:O	1:A:2205:ILE:HG12	2.21	0.40
1:A:2481:ASP:OD1	1:A:2482:PHE:N	2.54	0.40
1:A:3348:SER:HA	1:A:3351:GLU:HB2	2.04	0.40
1:A:3369:TYR:HA	1:A:3372:LEU:HB3	2.03	0.40
1:A:3488:ILE:HA	1:A:3550:VAL:HG13	2.03	0.40
1:A:3998:MET:O	1:A:4002:LEU:HG	2.21	0.40
1:A:4479:TRP:O	1:A:4483:ILE:HG12	2.22	0.40
1:C:314:LEU:O	1:C:315:LEU:HD23	2.21	0.40
1:C:430:ILE:HD11	1:C:501:CYS:HB3	2.02	0.40
1:C:513:HIS:O	1:C:517:VAL:HG23	2.22	0.40
1:C:1257:GLN:HG2	1:C:1451:HIS:CE1	2.56	0.40
1:C:2581:PRO:HD2	1:C:2629:PHE:HD2	1.85	0.40
1:C:2621:CYS:HA	1:C:2676:PRO:HG3	2.02	0.40
1:C:3502:GLU:OE1	1:C:3502:GLU:N	2.40	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:525:SER:O	1:E:529:ILE:HG13	2.21	0.40
1:E:1273:ILE:HB	1:E:1287:GLN:OE1	2.21	0.40
1:E:1704:TYR:CG	1:E:1821:PRO:HB2	2.57	0.40
1:E:2426:ILE:O	1:E:2475:TYR:OH	2.37	0.40
1:E:2859:LEU:HD13	1:E:2866:ASN:HA	2.02	0.40
1:E:3007:PHE:HA	1:E:3035:LEU:HD13	2.03	0.40
1:E:3029:VAL:HG12	1:E:3033:HIS:NE2	2.36	0.40
1:E:3046:LYS:O	1:E:3046:LYS:HD2	2.21	0.40
1:E:3616:VAL:O	1:E:3620:LEU:HG	2.21	0.40
1:E:4505:LEU:HD13	1:E:4744:ASP:HB3	2.02	0.40
1:F:1310:CYS:HA	1:F:1514:ALA:HB1	2.03	0.40
1:F:1978:PHE:CZ	1:F:1995:LEU:HD23	2.56	0.40
1:F:2481:ASP:OD1	1:F:2482:PHE:N	2.54	0.40
1:F:2579:LEU:N	1:F:2615:ARG:HH12	2.18	0.40
1:F:2667:CYS:O	1:F:2671:VAL:HG23	2.21	0.40
1:F:3214:MET:SD	1:F:3271:HIS:NE2	2.94	0.40
1:F:3943:VAL:HG23	1:F:3977:MET:SD	2.61	0.40
1:A:430:ILE:HD11	1:A:501:CYS:HB3	2.02	0.40
1:A:705:PRO:HG3	1:A:857:LEU:HD12	2.03	0.40
1:A:921:PHE:HB2	1:A:929:ARG:NH1	2.36	0.40
1:A:1166:VAL:HG22	1:A:1173:MET:HB2	2.02	0.40
1:A:1273:ILE:HB	1:A:1287:GLN:OE1	2.21	0.40
1:A:1978:PHE:CZ	1:A:1995:LEU:HD23	2.56	0.40
1:A:2348:GLU:O	1:A:2352:ILE:HG12	2.21	0.40
1:A:2549:HIS:HD1	1:A:2549:HIS:H	1.68	0.40
1:A:2999:GLU:O	1:A:3003:VAL:HG23	2.22	0.40
1:A:3273:ASN:ND2	1:A:3313:LEU:HG	2.37	0.40
1:A:3276:LEU:HD22	1:A:3309:VAL:HG11	2.04	0.40
1:A:3383:TRP:HH2	1:A:3394:LEU:HD23	1.85	0.40
1:A:4188:PHE:CE1	1:A:4914:LEU:HD22	2.56	0.40
1:A:4521:SER:O	1:A:4556:VAL:N	2.54	0.40
1:C:541:ILE:HD11	1:C:574:VAL:HG13	2.04	0.40
1:C:2459:PHE:CE1	1:C:2464:LYS:HG3	2.56	0.40
1:C:2481:ASP:OD1	1:C:2482:PHE:N	2.54	0.40
1:C:2535:ALA:HB1	1:C:2578:GLN:O	2.21	0.40
1:C:2554:LEU:HG	1:C:2568:ILE:HD13	2.03	0.40
1:C:3007:PHE:HA	1:C:3035:LEU:HD13	2.04	0.40
1:E:541:ILE:HD11	1:E:574:VAL:HG13	2.03	0.40
1:E:744:PRO:HD3	1:E:776:GLN:HE21	1.86	0.40
1:E:1305:SER:N	1:E:1588:HIS:O	2.54	0.40
1:E:2202:PHE:O	1:E:2205:ILE:HG12	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:2224:SER:HB2	1:E:2239:LEU:HB2	2.03	0.40
1:E:2261:LEU:HD12	1:E:2261:LEU:HA	1.94	0.40
1:E:2392:ALA:HB2	1:E:2463:HIS:CD2	2.56	0.40
1:E:2535:ALA:HB1	1:E:2578:GLN:O	2.22	0.40
1:E:2667:CYS:O	1:E:2671:VAL:HG23	2.21	0.40
1:E:3014:VAL:HG12	1:E:3095:TYR:CE1	2.57	0.40
1:E:3030:ASN:HA	1:E:3033:HIS:CD2	2.56	0.40
1:F:64:ILE:HA	1:F:123:HIS:CE1	2.55	0.40
1:F:541:ILE:HD11	1:F:574:VAL:HG13	2.04	0.40
1:F:900:LEU:HB2	1:F:902:TRP:HD1	1.86	0.40
1:F:2224:SER:HB2	1:F:2239:LEU:HB2	2.03	0.40
1:F:2996:SER:O	1:F:2999:GLU:HB2	2.21	0.40
1:F:2999:GLU:O	1:F:3003:VAL:HG23	2.22	0.40
1:F:3426:ASN:O	1:F:3430:LEU:HG	2.21	0.40
1:F:3612:ARG:HH11	1:F:3612:ARG:HA	1.87	0.40
1:F:4113:ARG:H	1:F:4113:ARG:HG2	1.60	0.40
2:I:69:ILE:HB	2:I:80:LEU:HD13	2.04	0.40
1:A:76:ARG:O	1:A:80:GLU:HG2	2.22	0.40
1:A:1704:TYR:CG	1:A:1821:PRO:HB2	2.57	0.40
1:A:2154:PHE:CE1	1:A:2205:ILE:HD13	2.56	0.40
1:A:2580:ARG:HG2	1:A:2583:MET:HE3	2.01	0.40
1:A:2633:SER:OG	1:A:2636:GLU:HG3	2.22	0.40
1:A:3014:VAL:HG12	1:A:3095:TYR:CE1	2.57	0.40
1:A:3105:SER:HB3	1:A:3156:GLU:HG2	2.04	0.40
1:A:3612:ARG:O	1:A:3616:VAL:HG12	2.22	0.40
1:A:4618:GLU:OE1	1:A:4618:GLU:N	2.54	0.40
1:C:1303:ARG:HD3	1:C:1446:ILE:HD11	2.03	0.40
1:C:1588:HIS:CE1	1:C:1590:GLN:HE21	2.39	0.40
1:C:2428:LEU:O	1:C:2432:VAL:HG23	2.22	0.40
1:C:3754:VAL:HA	1:C:3757:THR:HG22	2.04	0.40
1:E:970:TYR:HB2	1:E:971:GLN:H	1.70	0.40
1:E:1030:PRO:HB2	1:E:1031:ARG:NH1	2.36	0.40
1:E:2633:SER:OG	1:E:2636:GLU:HG3	2.22	0.40
1:E:2765:LYS:O	1:E:2769:ILE:HG22	2.22	0.40
1:E:3870:ILE:O	1:E:3874:VAL:HG23	2.20	0.40
1:E:3900:GLN:HA	1:E:3903:ARG:HH11	1.87	0.40
1:E:3943:VAL:HG23	1:E:3977:MET:SD	2.61	0.40
1:E:4618:GLU:OE1	1:E:4618:GLU:N	2.54	0.40
1:F:641:ASP:O	1:F:642:LEU:HB3	2.21	0.40
1:F:1257:GLN:HB2	1:F:1596:LEU:HD21	2.03	0.40
1:F:1843:ILE:HD13	1:F:1843:ILE:HA	1.92	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:4030:THR:HG23	1:F:4054:HIS:CE1	2.57	0.40
1:F:4751:THR:O	1:F:4755:ILE:HG13	2.21	0.40
2:G:48:VAL:HG12	2:G:49:ALA:H	1.86	0.40
1:A:660:PHE:HB3	1:A:787:LEU:HD22	2.04	0.40
1:A:897:LYS:HE2	1:A:915:HIS:NE2	2.36	0.40
1:A:908:ARG:NE	2:B:104:TYR:HB3	2.35	0.40
1:A:1124:PRO:HG3	1:A:1597:TRP:CE2	2.57	0.40
1:A:1588:HIS:CE1	1:A:1590:GLN:HE21	2.39	0.40
1:A:1686:LEU:HD13	1:A:1707:LEU:HD13	2.02	0.40
1:A:1802:GLU:HA	1:A:1805:LEU:HG	2.02	0.40
1:A:2535:ALA:HB1	1:A:2578:GLN:O	2.21	0.40
1:A:2676:PRO:HA	1:A:2677:PRO:HD3	1.99	0.40
1:A:3612:ARG:HA	1:A:3612:ARG:HH11	1.87	0.40
1:A:3754:VAL:HA	1:A:3757:THR:HG22	2.04	0.40
1:A:3811:ASN:O	1:A:3814:GLU:HG2	2.21	0.40
2:B:69:ILE:HB	2:B:80:LEU:HD13	2.04	0.40
1:C:143:LEU:HB3	1:C:190:ARG:HH21	1.87	0.40
1:C:227:TYR:HA	1:C:355:LYS:HA	2.03	0.40
1:C:921:PHE:HA	1:C:924:LEU:HB2	2.02	0.40
1:C:1838:GLU:H	1:C:1838:GLU:CD	2.25	0.40
1:C:1978:PHE:CZ	1:C:1995:LEU:HD23	2.56	0.40
1:C:1999:HIS:CG	1:C:3627:TRP:HD1	2.39	0.40
1:C:2455:MET:HE3	1:C:2456:SER:HB2	2.04	0.40
1:C:2588:LEU:HD21	1:C:2612:HIS:NE2	2.37	0.40
1:C:3214:MET:SD	1:C:3271:HIS:NE2	2.94	0.40
1:C:3386:GLU:HB3	1:C:3535:ASN:ND2	2.36	0.40
1:C:4044:LYS:HB2	1:C:4075:GLU:OE2	2.21	0.40
1:C:4168:LYS:HB3	1:C:4168:LYS:HE2	1.86	0.40
2:D:48:VAL:HG12	2:D:49:ALA:H	1.86	0.40
1:E:897:LYS:HE2	1:E:915:HIS:NE2	2.36	0.40
1:E:1420:LEU:O	1:E:1423:THR:HG22	2.21	0.40
1:E:1967:PRO:O	1:E:1971:GLN:HG3	2.21	0.40
1:E:2276:CYS:HB2	1:E:2290:ASN:ND2	2.36	0.40
1:E:2549:HIS:HD1	1:E:2549:HIS:H	1.68	0.40
1:E:2554:LEU:HG	1:E:2568:ILE:HD13	2.03	0.40
1:E:3644:ALA:HB2	1:E:3663:LEU:HD13	2.04	0.40
1:E:3811:ASN:O	1:E:3814:GLU:HG2	2.21	0.40
1:F:525:SER:O	1:F:529:ILE:HG13	2.21	0.40
1:F:705:PRO:HG3	1:F:857:LEU:HD12	2.03	0.40
1:F:1124:PRO:HG3	1:F:1597:TRP:CE2	2.57	0.40
1:F:2154:PHE:CE1	1:F:2205:ILE:HD13	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:2221:LEU:HD12	1:F:2260:ASP:HB2	2.03	0.40
1:F:2564:GLN:O	1:F:2568:ILE:HG13	2.22	0.40
1:F:2850:ILE:HD12	1:F:2850:ILE:HA	1.96	0.40
1:F:3169:PHE:HE1	1:F:3244:TYR:OH	2.05	0.40
1:F:3369:TYR:CE2	1:F:3465:ILE:HA	2.57	0.40
1:F:3374:ARG:NH2	1:F:3431:ILE:O	2.55	0.40
1:F:3616:VAL:O	1:F:3620:LEU:HG	2.21	0.40
1:F:3754:VAL:HA	1:F:3757:THR:HG22	2.04	0.40
1:F:3902:GLN:HE21	1:F:3963:GLN:HG2	1.87	0.40
1:F:4020:LEU:HD12	1:F:4124:VAL:HG12	2.03	0.40
1:F:4199:MET:HB2	1:F:4199:MET:HE2	1.76	0.40

There are no symmetry-related clashes.

5.3 Torsion angles ⓘ

5.3.1 Protein backbone ⓘ

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	4094/4966 (82%)	3987 (97%)	106 (3%)	1 (0%)	100	100
1	C	4094/4966 (82%)	3986 (97%)	107 (3%)	1 (0%)	100	100
1	E	4094/4966 (82%)	3986 (97%)	107 (3%)	1 (0%)	100	100
1	F	4094/4966 (82%)	3983 (97%)	110 (3%)	1 (0%)	100	100
2	B	124/137 (90%)	118 (95%)	6 (5%)	0	100	100
2	D	124/137 (90%)	118 (95%)	6 (5%)	0	100	100
2	G	124/137 (90%)	118 (95%)	6 (5%)	0	100	100
2	I	124/137 (90%)	118 (95%)	6 (5%)	0	100	100
All	All	16872/20412 (83%)	16414 (97%)	454 (3%)	4 (0%)	100	100

All (4) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	2530	CYS
1	C	2530	CYS
1	E	2530	CYS
1	F	2530	CYS

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A	3589/4355 (82%)	3516 (98%)	73 (2%)	50	70
1	C	3589/4355 (82%)	3517 (98%)	72 (2%)	50	70
1	E	3589/4355 (82%)	3516 (98%)	73 (2%)	50	70
1	F	3589/4355 (82%)	3516 (98%)	73 (2%)	50	70
2	B	103/114 (90%)	102 (99%)	1 (1%)	73	83
2	D	103/114 (90%)	102 (99%)	1 (1%)	73	83
2	G	103/114 (90%)	101 (98%)	2 (2%)	52	71
2	I	103/114 (90%)	102 (99%)	1 (1%)	73	83
All	All	14768/17876 (83%)	14472 (98%)	296 (2%)	50	70

All (296) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	36	CYS
1	A	42	PHE
1	A	137	ARG
1	A	196	TYR
1	A	241	MET
1	A	317	MET
1	A	332	ARG
1	A	356	TYR
1	A	655	MET
1	A	678	MET
1	A	904	TYR
1	A	913	ARG

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Mol	Chain	Res	Type
1	A	917	CYS
1	A	926	GLU
1	A	988	LEU
1	A	1156	TRP
1	A	1174	MET
1	A	1290	PHE
1	A	1293	GLN
1	A	1421	MET
1	A	1487	MET
1	A	1801	LYS
1	A	1915	CYS
1	A	1962	ARG
1	A	2083	MET
1	A	2184	LYS
1	A	2302	ARG
1	A	2383	MET
1	A	2406	HIS
1	A	2491	PHE
1	A	2534	PHE
1	A	2549	HIS
1	A	2604	MET
1	A	2638	HIS
1	A	2723	TYR
1	A	2740	TRP
1	A	2742	TYR
1	A	2836	LEU
1	A	2855	LYS
1	A	2918	LYS
1	A	2924	PHE
1	A	2931	TYR
1	A	2961	PHE
1	A	2973	TYR
1	A	2974	PHE
1	A	3007	PHE
1	A	3046	LYS
1	A	3244	TYR
1	A	3281	LYS
1	A	3294	TRP
1	A	3322	MET
1	A	3383	TRP
1	A	3411	PHE
1	A	3427	MET

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Mol	Chain	Res	Type
1	A	3627	TRP
1	A	3853	PHE
1	A	3888	TYR
1	A	3955	MET
1	A	3977	MET
1	A	4011	MET
1	A	4047	PHE
1	A	4051	MET
1	A	4111	ASP
1	A	4161	LYS
1	A	4512	ASN
1	A	4518	TYR
1	A	4643	TYR
1	A	4671	MET
1	A	4736	PHE
1	A	4799	ASP
1	A	4894	ASN
1	A	4922	MET
1	A	4938	TYR
2	B	114	TYR
1	C	36	CYS
1	C	42	PHE
1	C	137	ARG
1	C	196	TYR
1	C	241	MET
1	C	317	MET
1	C	332	ARG
1	C	356	TYR
1	C	655	MET
1	C	678	MET
1	C	904	TYR
1	C	913	ARG
1	C	917	CYS
1	C	926	GLU
1	C	988	LEU
1	C	1156	TRP
1	C	1174	MET
1	C	1290	PHE
1	C	1293	GLN
1	C	1421	MET
1	C	1487	MET
1	C	1801	LYS

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Mol	Chain	Res	Type
1	C	1915	CYS
1	C	1962	ARG
1	C	2083	MET
1	C	2184	LYS
1	C	2302	ARG
1	C	2383	MET
1	C	2406	HIS
1	C	2491	PHE
1	C	2534	PHE
1	C	2549	HIS
1	C	2604	MET
1	C	2638	HIS
1	C	2723	TYR
1	C	2740	TRP
1	C	2742	TYR
1	C	2836	LEU
1	C	2855	LYS
1	C	2918	LYS
1	C	2924	PHE
1	C	2931	TYR
1	C	2961	PHE
1	C	2973	TYR
1	C	2974	PHE
1	C	3007	PHE
1	C	3046	LYS
1	C	3244	TYR
1	C	3281	LYS
1	C	3294	TRP
1	C	3322	MET
1	C	3383	TRP
1	C	3411	PHE
1	C	3627	TRP
1	C	3853	PHE
1	C	3888	TYR
1	C	3955	MET
1	C	3977	MET
1	C	4011	MET
1	C	4047	PHE
1	C	4051	MET
1	C	4111	ASP
1	C	4161	LYS
1	C	4512	ASN

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Mol	Chain	Res	Type
1	C	4518	TYR
1	C	4643	TYR
1	C	4671	MET
1	C	4736	PHE
1	C	4799	ASP
1	C	4894	ASN
1	C	4922	MET
1	C	4938	TYR
2	D	114	TYR
1	E	36	CYS
1	E	42	PHE
1	E	137	ARG
1	E	196	TYR
1	E	241	MET
1	E	317	MET
1	E	332	ARG
1	E	356	TYR
1	E	655	MET
1	E	678	MET
1	E	904	TYR
1	E	913	ARG
1	E	917	CYS
1	E	926	GLU
1	E	988	LEU
1	E	1156	TRP
1	E	1174	MET
1	E	1290	PHE
1	E	1293	GLN
1	E	1421	MET
1	E	1487	MET
1	E	1801	LYS
1	E	1915	CYS
1	E	1962	ARG
1	E	2083	MET
1	E	2184	LYS
1	E	2302	ARG
1	E	2383	MET
1	E	2406	HIS
1	E	2491	PHE
1	E	2534	PHE
1	E	2549	HIS
1	E	2604	MET

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Mol	Chain	Res	Type
1	E	2638	HIS
1	E	2723	TYR
1	E	2740	TRP
1	E	2742	TYR
1	E	2836	LEU
1	E	2855	LYS
1	E	2918	LYS
1	E	2924	PHE
1	E	2931	TYR
1	E	2961	PHE
1	E	2973	TYR
1	E	2974	PHE
1	E	3007	PHE
1	E	3046	LYS
1	E	3244	TYR
1	E	3281	LYS
1	E	3294	TRP
1	E	3322	MET
1	E	3383	TRP
1	E	3411	PHE
1	E	3427	MET
1	E	3627	TRP
1	E	3853	PHE
1	E	3888	TYR
1	E	3955	MET
1	E	3977	MET
1	E	4011	MET
1	E	4047	PHE
1	E	4051	MET
1	E	4111	ASP
1	E	4161	LYS
1	E	4512	ASN
1	E	4518	TYR
1	E	4643	TYR
1	E	4671	MET
1	E	4736	PHE
1	E	4799	ASP
1	E	4894	ASN
1	E	4922	MET
1	E	4938	TYR
1	F	36	CYS
1	F	42	PHE

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Mol	Chain	Res	Type
1	F	137	ARG
1	F	196	TYR
1	F	241	MET
1	F	317	MET
1	F	332	ARG
1	F	356	TYR
1	F	655	MET
1	F	678	MET
1	F	904	TYR
1	F	913	ARG
1	F	917	CYS
1	F	926	GLU
1	F	988	LEU
1	F	1156	TRP
1	F	1174	MET
1	F	1290	PHE
1	F	1293	GLN
1	F	1421	MET
1	F	1487	MET
1	F	1801	LYS
1	F	1915	CYS
1	F	1962	ARG
1	F	2083	MET
1	F	2184	LYS
1	F	2302	ARG
1	F	2383	MET
1	F	2406	HIS
1	F	2491	PHE
1	F	2534	PHE
1	F	2549	HIS
1	F	2604	MET
1	F	2638	HIS
1	F	2723	TYR
1	F	2740	TRP
1	F	2742	TYR
1	F	2836	LEU
1	F	2855	LYS
1	F	2918	LYS
1	F	2924	PHE
1	F	2931	TYR
1	F	2961	PHE
1	F	2973	TYR

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Mol	Chain	Res	Type
1	F	2974	PHE
1	F	3007	PHE
1	F	3046	LYS
1	F	3244	TYR
1	F	3281	LYS
1	F	3294	TRP
1	F	3322	MET
1	F	3383	TRP
1	F	3411	PHE
1	F	3427	MET
1	F	3627	TRP
1	F	3853	PHE
1	F	3888	TYR
1	F	3955	MET
1	F	3977	MET
1	F	4011	MET
1	F	4047	PHE
1	F	4051	MET
1	F	4111	ASP
1	F	4161	LYS
1	F	4512	ASN
1	F	4518	TYR
1	F	4643	TYR
1	F	4671	MET
1	F	4736	PHE
1	F	4799	ASP
1	F	4894	ASN
1	F	4922	MET
1	F	4938	TYR
2	G	46	GLU
2	G	114	TYR
2	I	114	TYR

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (53) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	238	HIS
1	A	375	GLN
1	A	476	GLN
1	A	1143	GLN
1	A	1296	ASN
1	A	1498	GLN

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Mol	Chain	Res	Type
1	A	1588	HIS
1	A	2480	GLN
1	A	2972	GLN
1	A	3126	GLN
1	A	3485	GLN
1	A	3954	GLN
1	A	4786	ASN
2	B	105	ASN
1	C	238	HIS
1	C	375	GLN
1	C	476	GLN
1	C	1143	GLN
1	C	1296	ASN
1	C	1498	GLN
1	C	1588	HIS
1	C	2480	GLN
1	C	2578	GLN
1	C	2972	GLN
1	C	3126	GLN
1	C	3485	GLN
1	C	4786	ASN
2	D	105	ASN
1	E	238	HIS
1	E	375	GLN
1	E	476	GLN
1	E	1143	GLN
1	E	1296	ASN
1	E	1588	HIS
1	E	1936	GLN
1	E	2480	GLN
1	E	2972	GLN
1	E	3126	GLN
1	E	3485	GLN
1	E	4786	ASN
1	F	238	HIS
1	F	375	GLN
1	F	476	GLN
1	F	1143	GLN
1	F	1296	ASN
1	F	1588	HIS
1	F	2480	GLN
1	F	2972	GLN

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Mol	Chain	Res	Type
1	F	3126	GLN
1	F	3485	GLN
1	F	4786	ASN
2	G	105	ASN
2	I	105	ASN

5.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 16 ligands modelled in this entry, 8 are monoatomic - leaving 8 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. 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 $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
3	ATP	C	5101	-	28,33,33	0.69	0	34,52,52	0.65	1 (2%)
4	CFF	A	5102	-	8,15,15	0.96	0	8,23,23	2.51	2 (25%)
3	ATP	F	5101	-	28,33,33	0.69	0	34,52,52	0.65	1 (2%)
4	CFF	E	5102	-	8,15,15	0.97	0	8,23,23	2.51	2 (25%)
3	ATP	E	5101	-	28,33,33	0.70	0	34,52,52	0.64	1 (2%)
4	CFF	C	5102	-	8,15,15	0.96	0	8,23,23	2.50	2 (25%)
3	ATP	A	5101	-	28,33,33	0.69	0	34,52,52	0.65	1 (2%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
4	CFF	F	5102	-	8,15,15	0.97	0	8,23,23	2.51	2 (25%)

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
3	ATP	C	5101	-	-	5/18/38/38	0/3/3/3
4	CFF	A	5102	-	-	-	0/2/2/2
3	ATP	F	5101	-	-	5/18/38/38	0/3/3/3
4	CFF	E	5102	-	-	-	0/2/2/2
3	ATP	E	5101	-	-	5/18/38/38	0/3/3/3
4	CFF	C	5102	-	-	-	0/2/2/2
3	ATP	A	5101	-	-	5/18/38/38	0/3/3/3
4	CFF	F	5102	-	-	-	0/2/2/2

There are no bond length outliers.

All (12) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
4	A	5102	CFF	C5-C6-N1	-6.18	111.93	118.20
4	E	5102	CFF	C5-C6-N1	-6.18	111.93	118.20
4	F	5102	CFF	C5-C6-N1	-6.16	111.94	118.20
4	C	5102	CFF	C5-C6-N1	-6.16	111.94	118.20
4	A	5102	CFF	C4-C5-C6	3.15	122.36	119.96
4	E	5102	CFF	C4-C5-C6	3.15	122.36	119.96
4	F	5102	CFF	C4-C5-C6	3.14	122.35	119.96
4	C	5102	CFF	C4-C5-C6	3.14	122.35	119.96
3	F	5101	ATP	C5-C6-N6	2.33	123.86	120.31
3	A	5101	ATP	C5-C6-N6	2.32	123.85	120.31
3	C	5101	ATP	C5-C6-N6	2.32	123.84	120.31
3	E	5101	ATP	C5-C6-N6	2.29	123.81	120.31

There are no chirality outliers.

All (20) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
3	A	5101	ATP	C5'-O5'-PA-O1A
3	A	5101	ATP	C5'-O5'-PA-O3A

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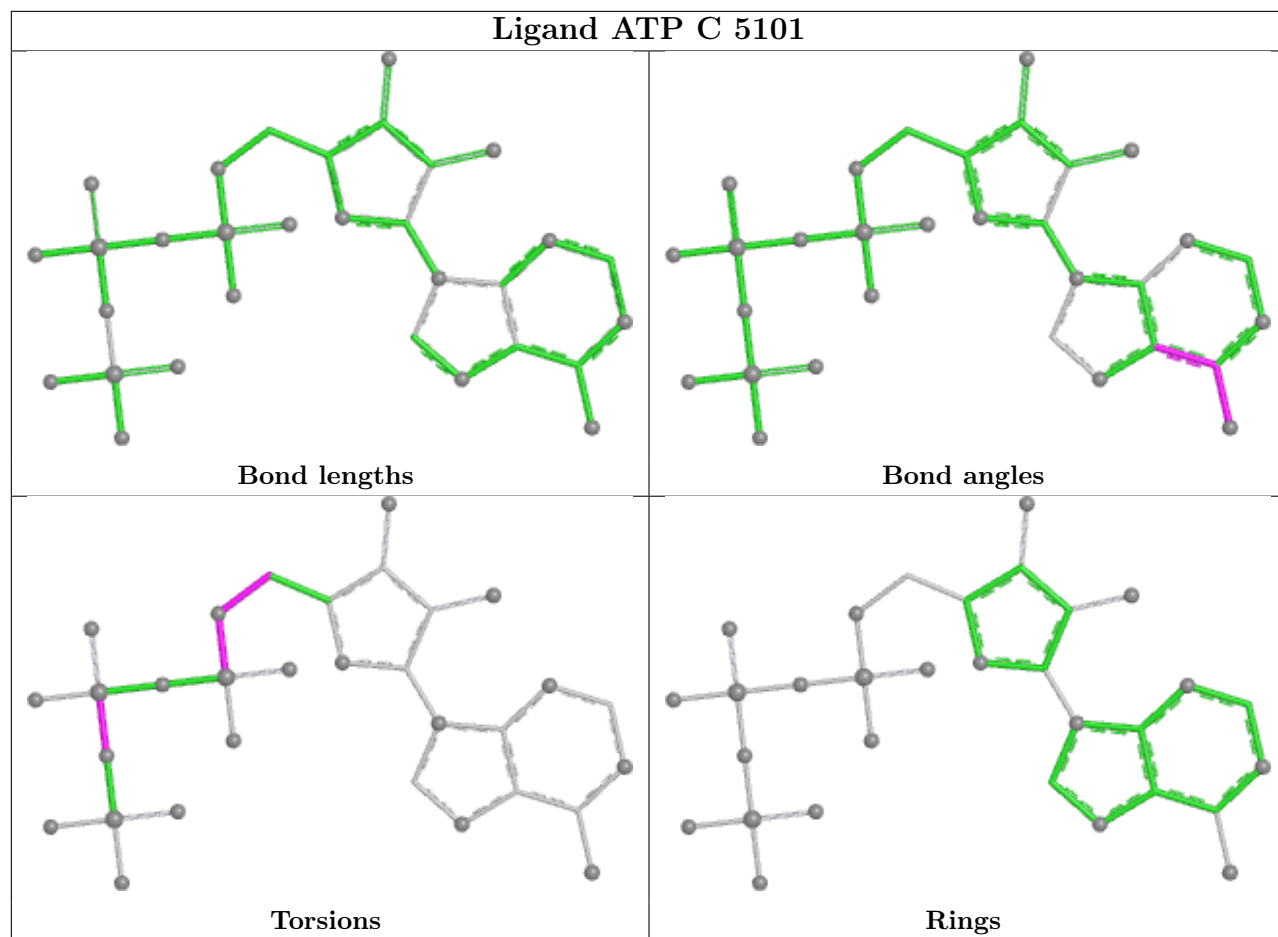
Mol	Chain	Res	Type	Atoms
3	C	5101	ATP	C5'-O5'-PA-O1A
3	C	5101	ATP	C5'-O5'-PA-O3A
3	E	5101	ATP	C5'-O5'-PA-O1A
3	E	5101	ATP	C5'-O5'-PA-O3A
3	F	5101	ATP	C5'-O5'-PA-O1A
3	F	5101	ATP	C5'-O5'-PA-O3A
3	A	5101	ATP	C4'-C5'-O5'-PA
3	C	5101	ATP	C4'-C5'-O5'-PA
3	E	5101	ATP	C4'-C5'-O5'-PA
3	F	5101	ATP	C4'-C5'-O5'-PA
3	A	5101	ATP	C5'-O5'-PA-O2A
3	C	5101	ATP	C5'-O5'-PA-O2A
3	E	5101	ATP	C5'-O5'-PA-O2A
3	F	5101	ATP	C5'-O5'-PA-O2A
3	A	5101	ATP	PG-O3B-PB-O2B
3	C	5101	ATP	PG-O3B-PB-O2B
3	E	5101	ATP	PG-O3B-PB-O2B
3	F	5101	ATP	PG-O3B-PB-O2B

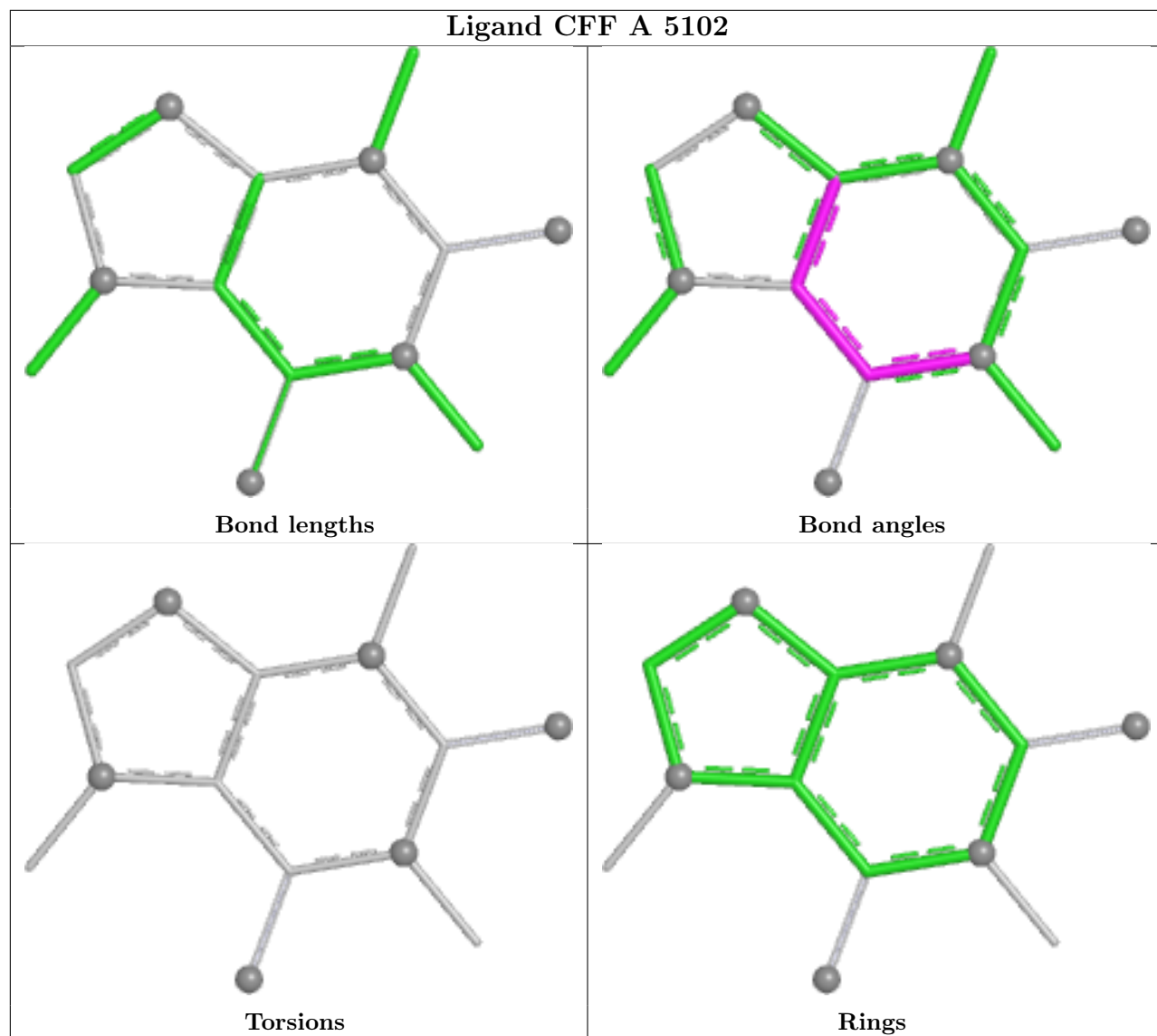
There are no ring outliers.

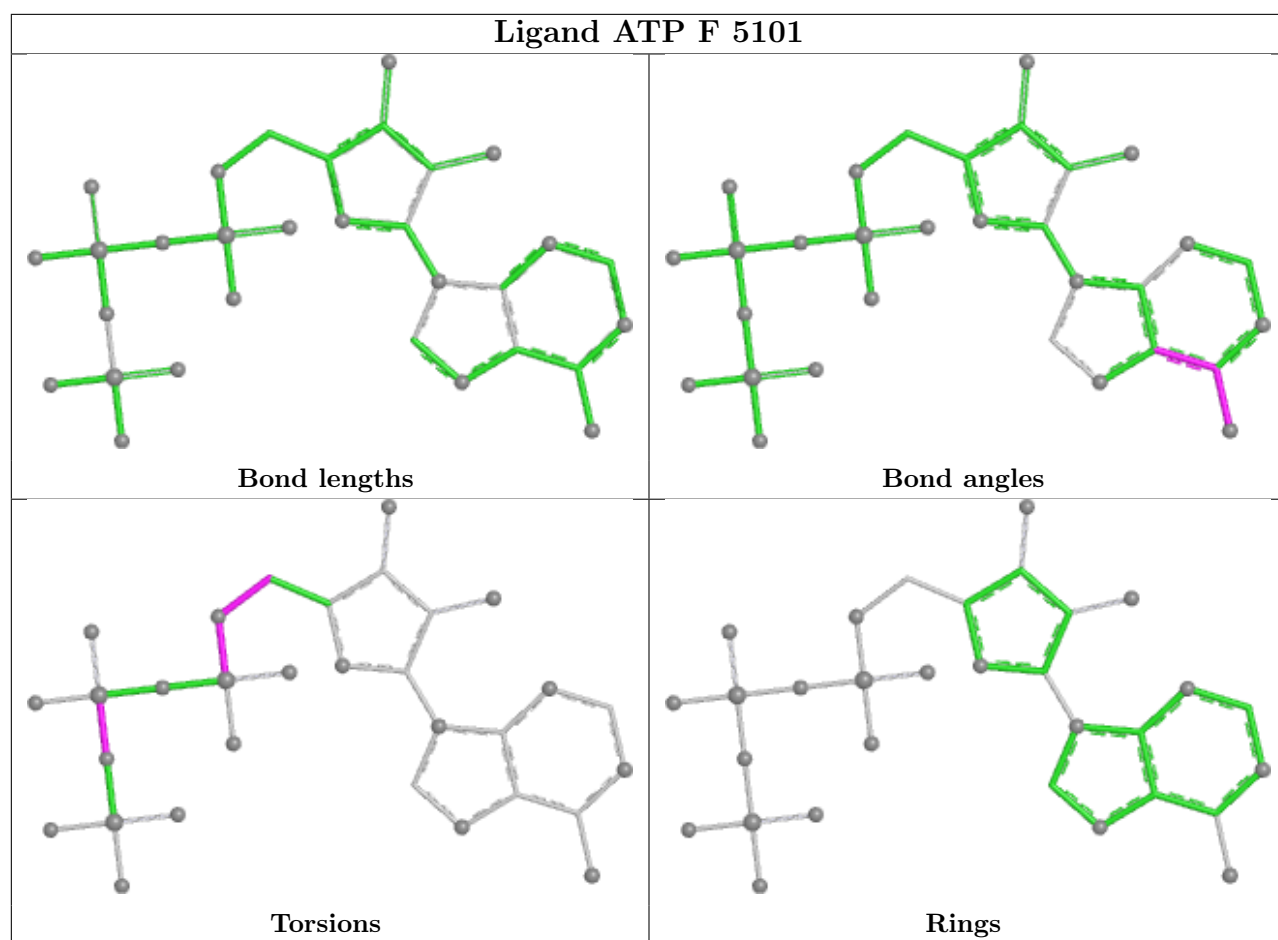
4 monomers are involved in 12 short contacts:

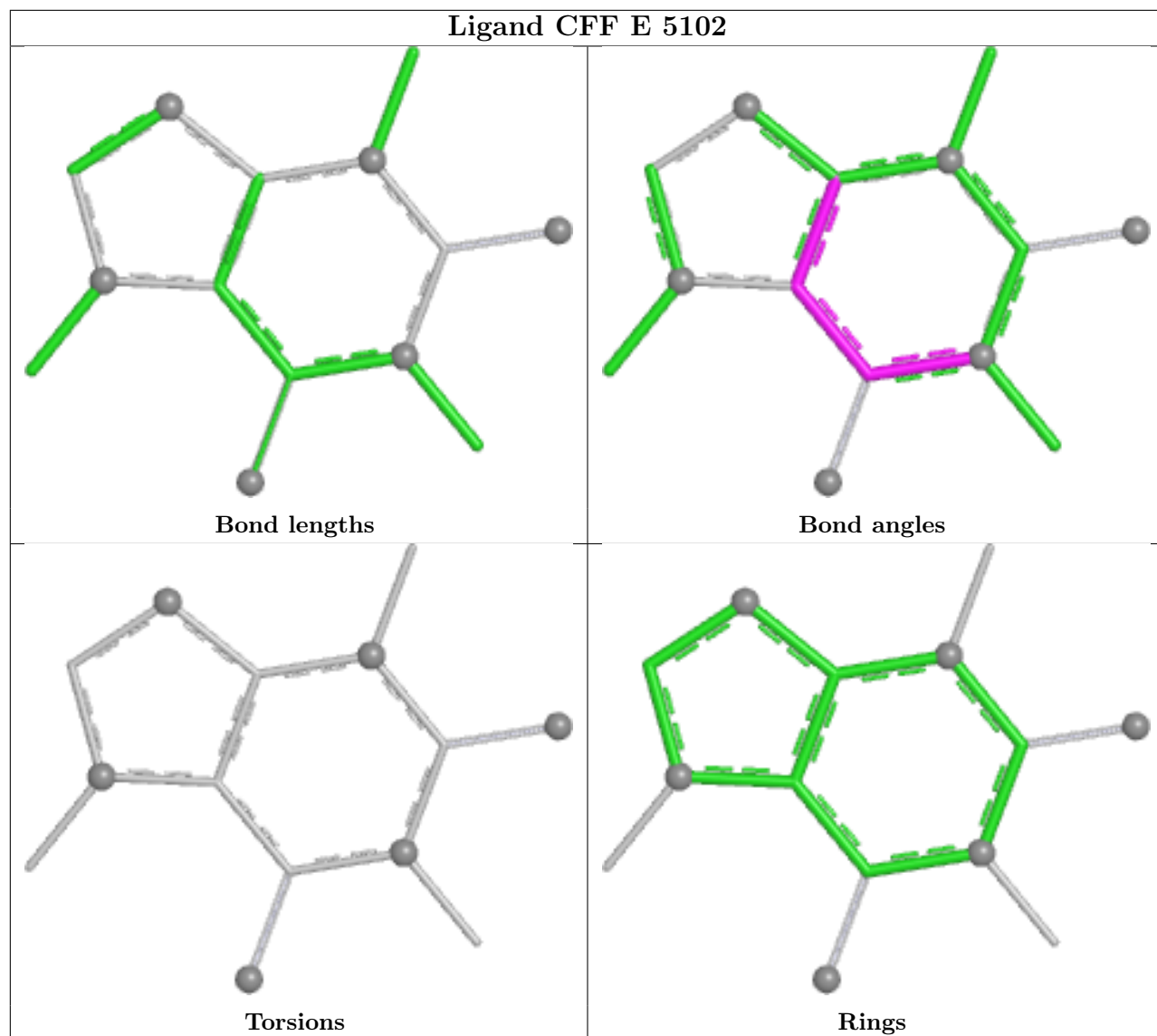
Mol	Chain	Res	Type	Clashes	Symm-Clashes
3	C	5101	ATP	3	0
3	F	5101	ATP	3	0
3	E	5101	ATP	3	0
3	A	5101	ATP	3	0

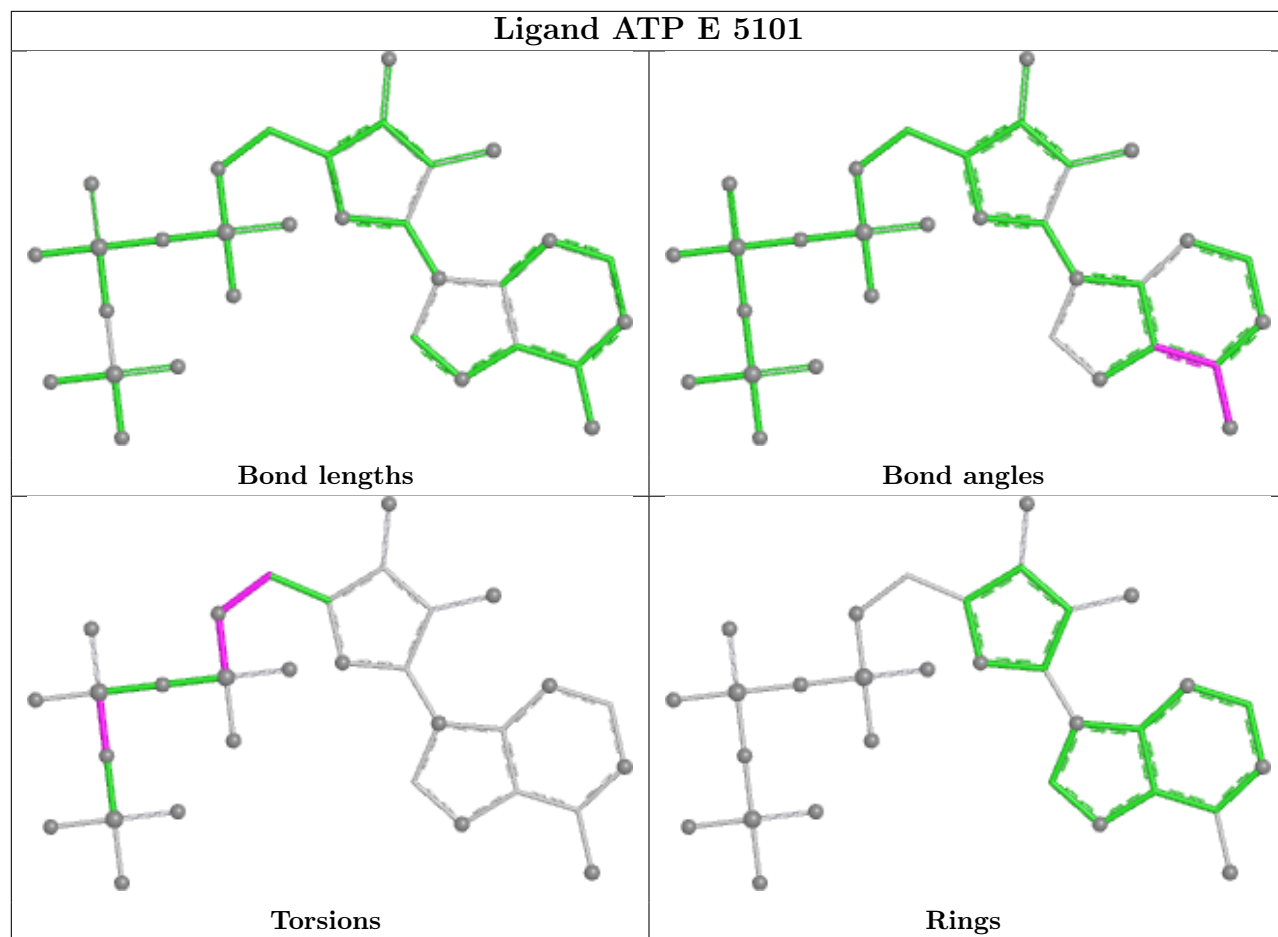
The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.

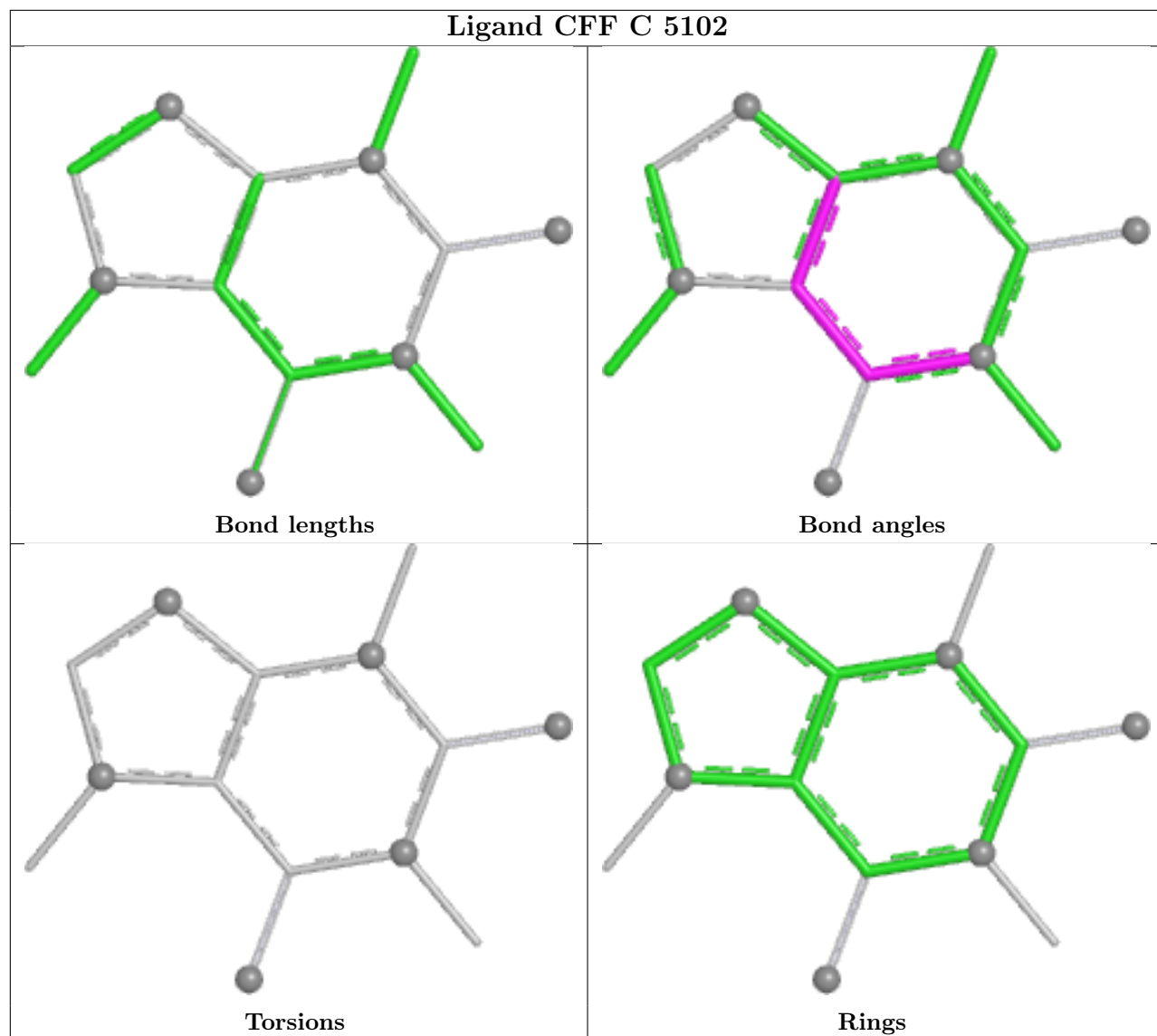


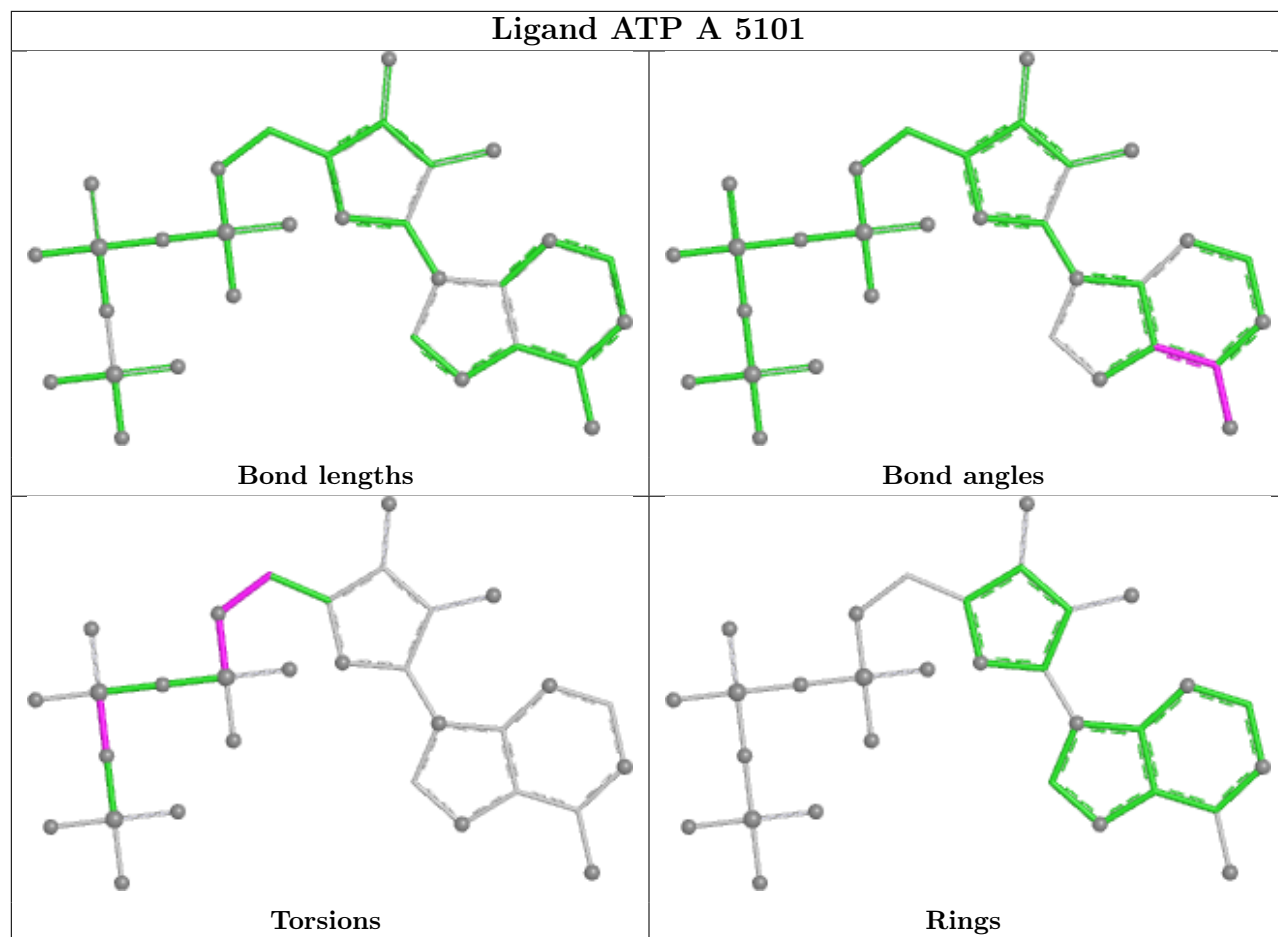


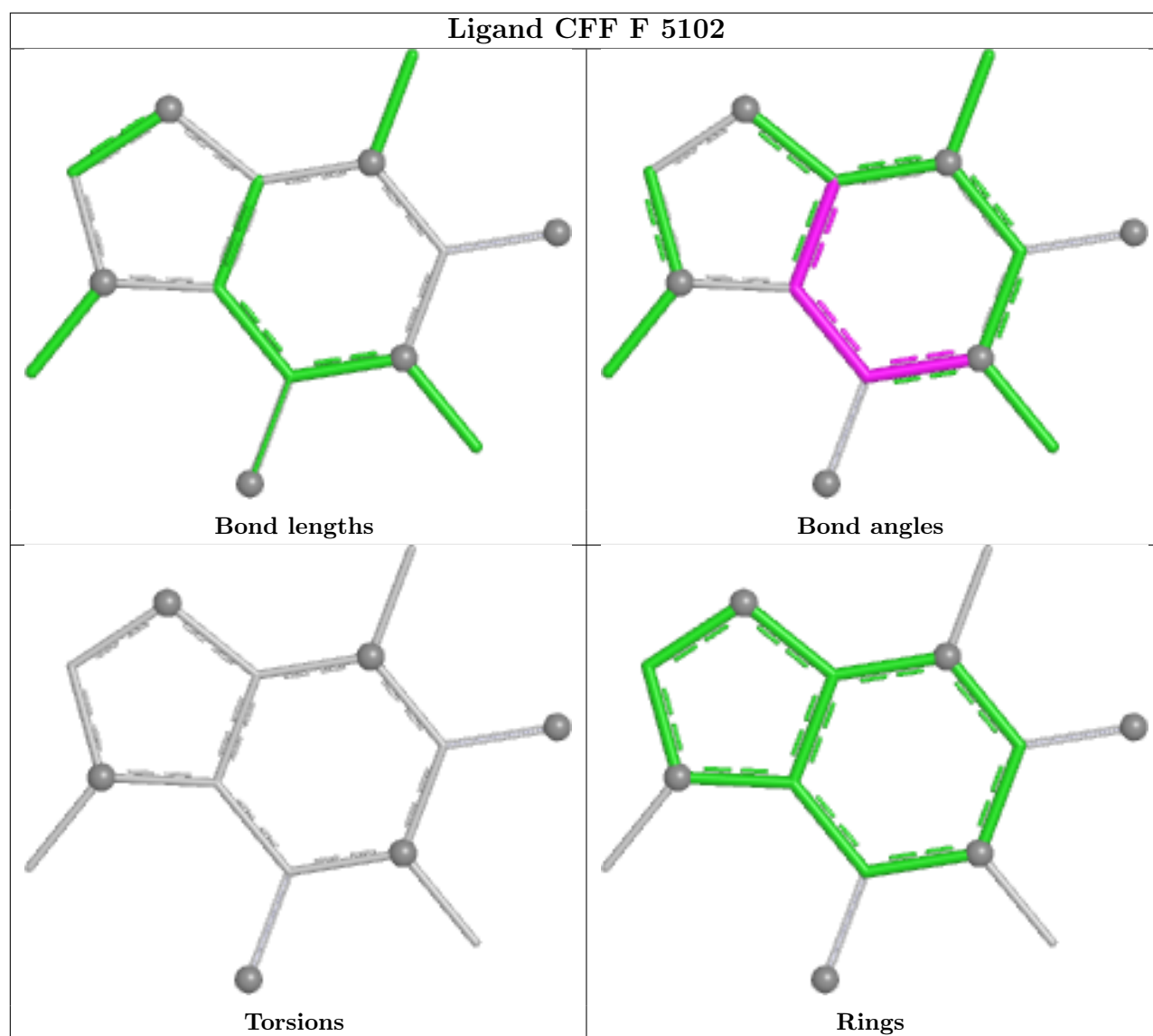












5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

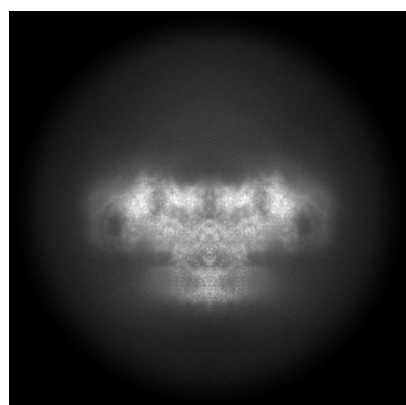
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-19463. These allow visual inspection of the internal detail of the map and identification of artifacts.

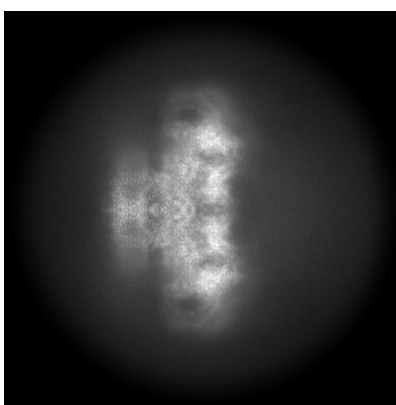
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

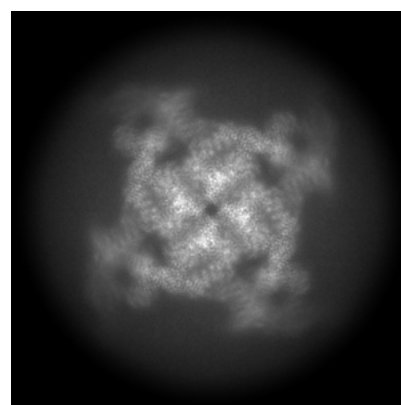
6.1.1 Primary map



X



Y

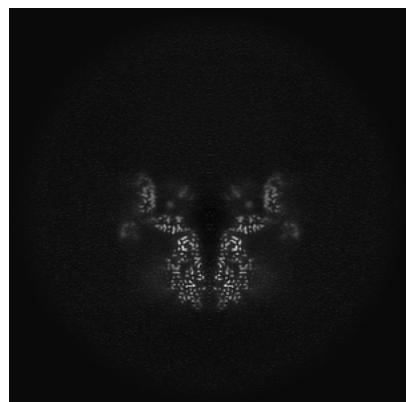


Z

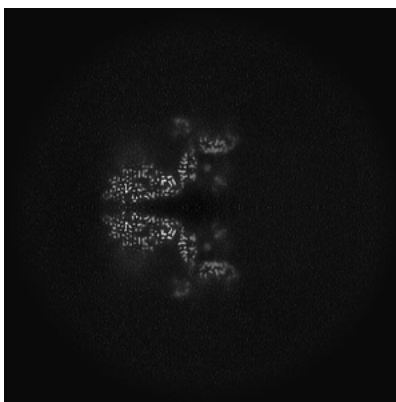
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

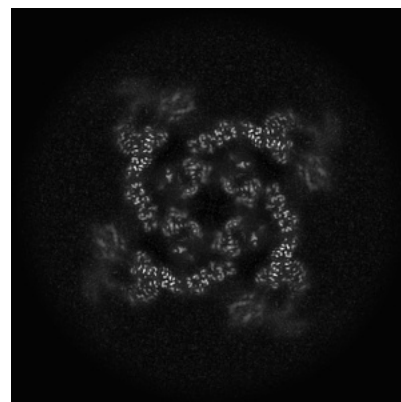
6.2.1 Primary map



X Index: 168



Y Index: 168

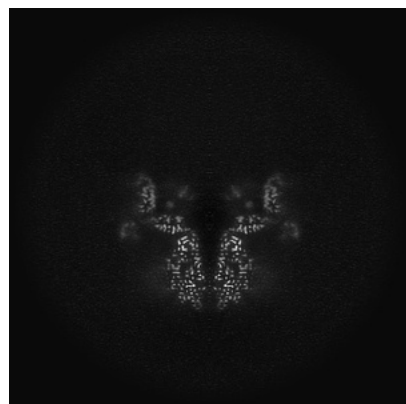


Z Index: 168

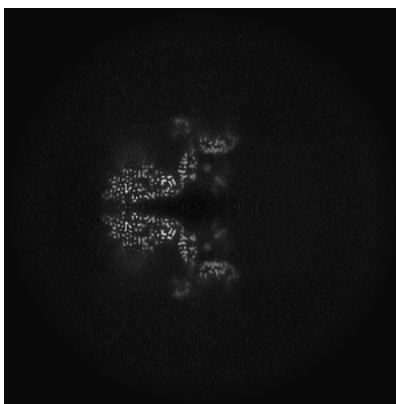
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

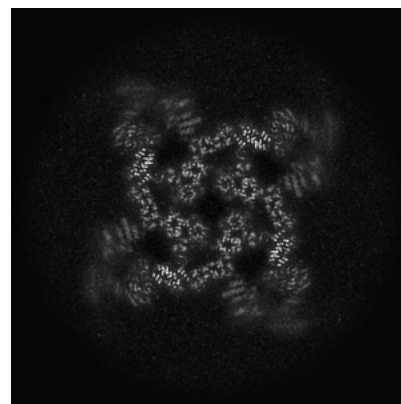
6.3.1 Primary map



X Index: 168



Y Index: 168

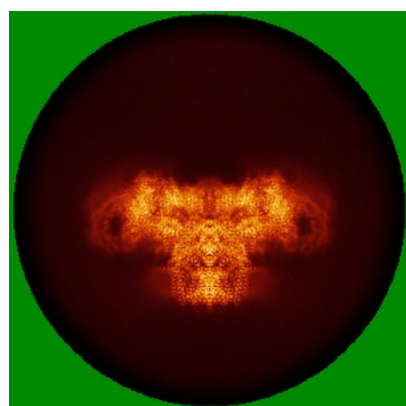


Z Index: 173

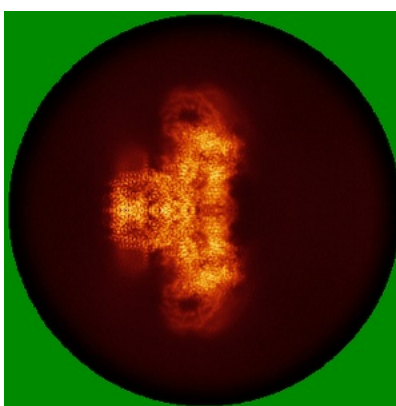
The images above show the largest variance slices of the map in three orthogonal directions.

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

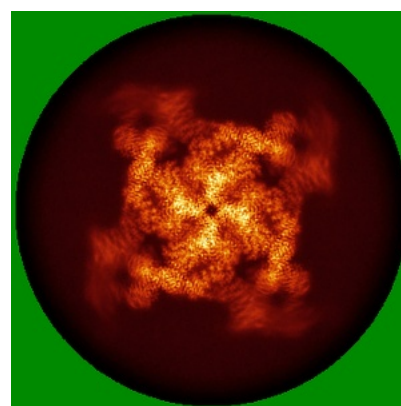
6.4.1 Primary map



X



Y

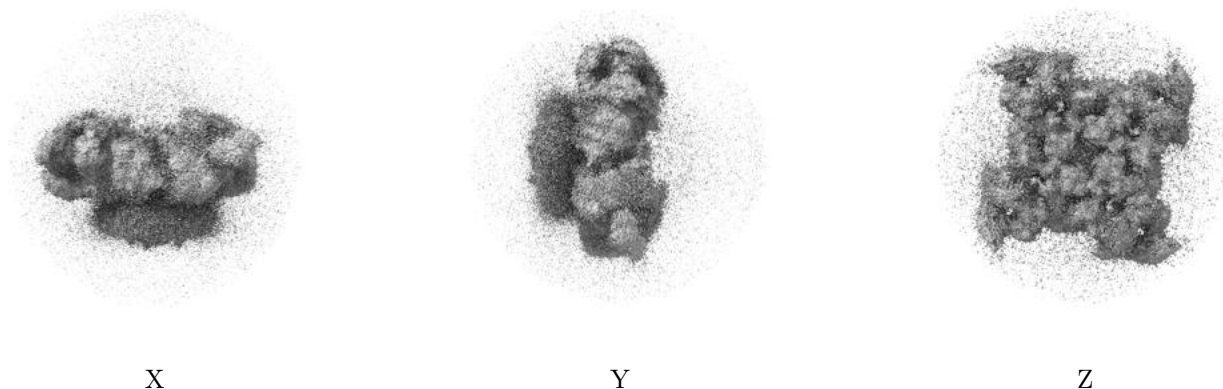


Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.35. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

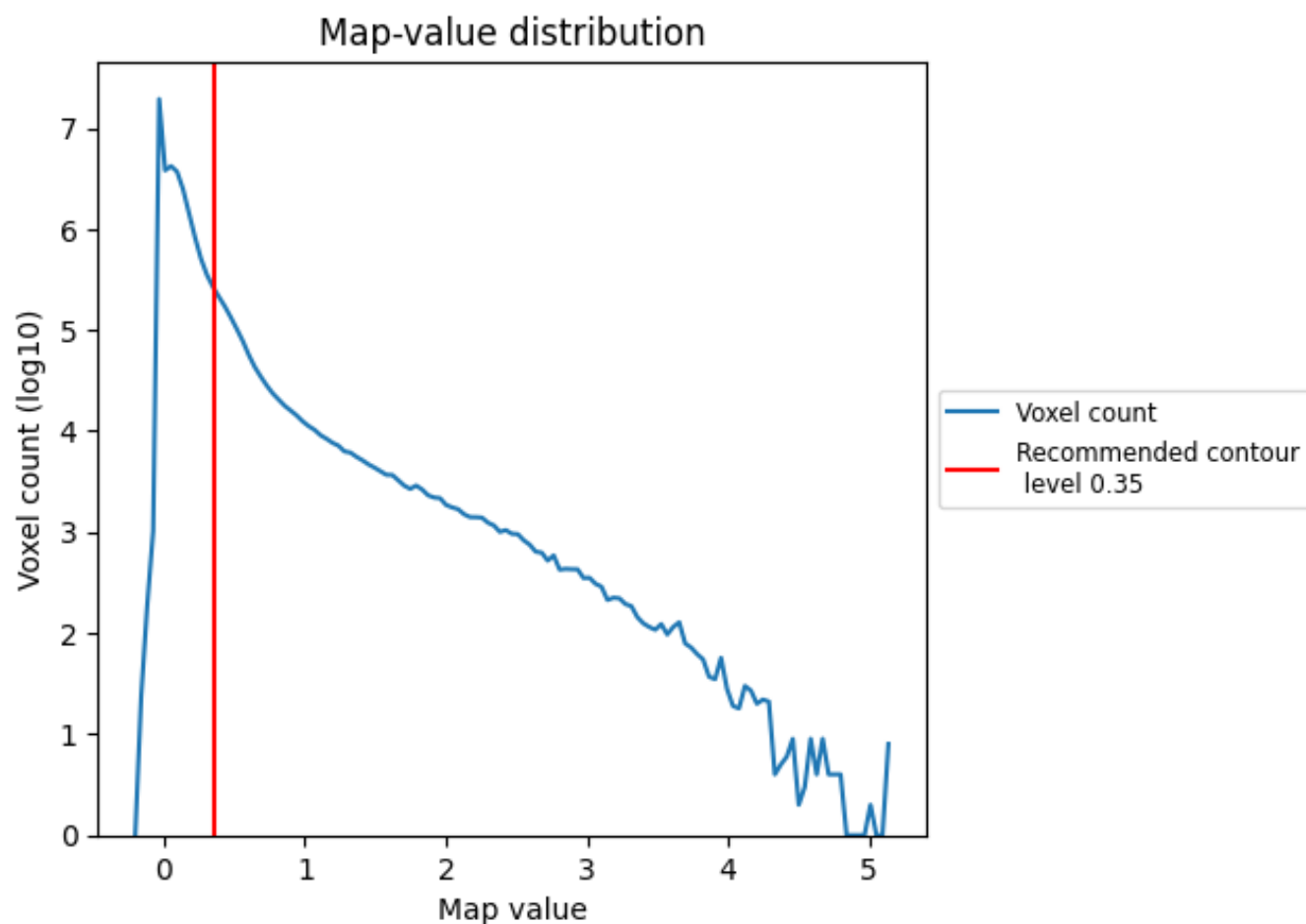
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

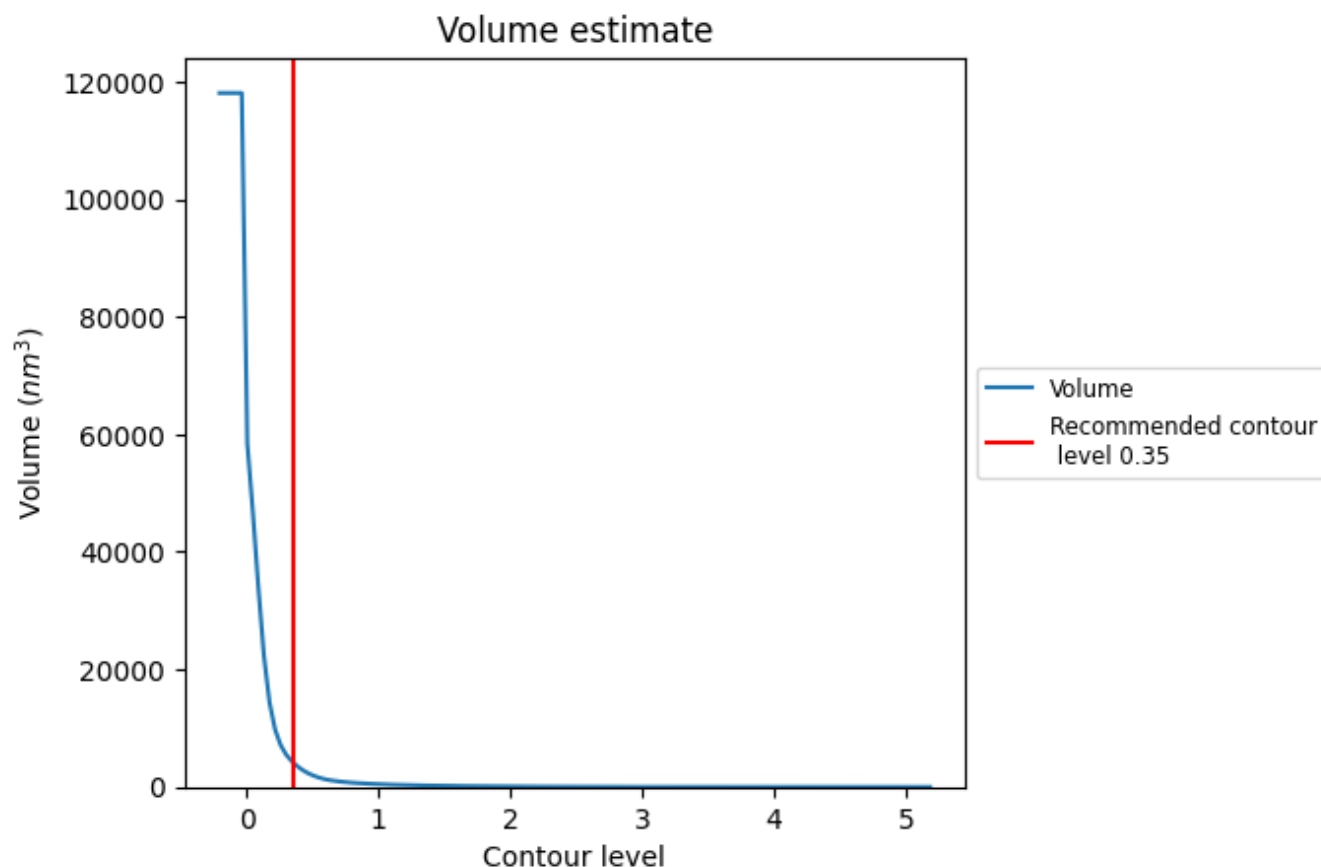
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

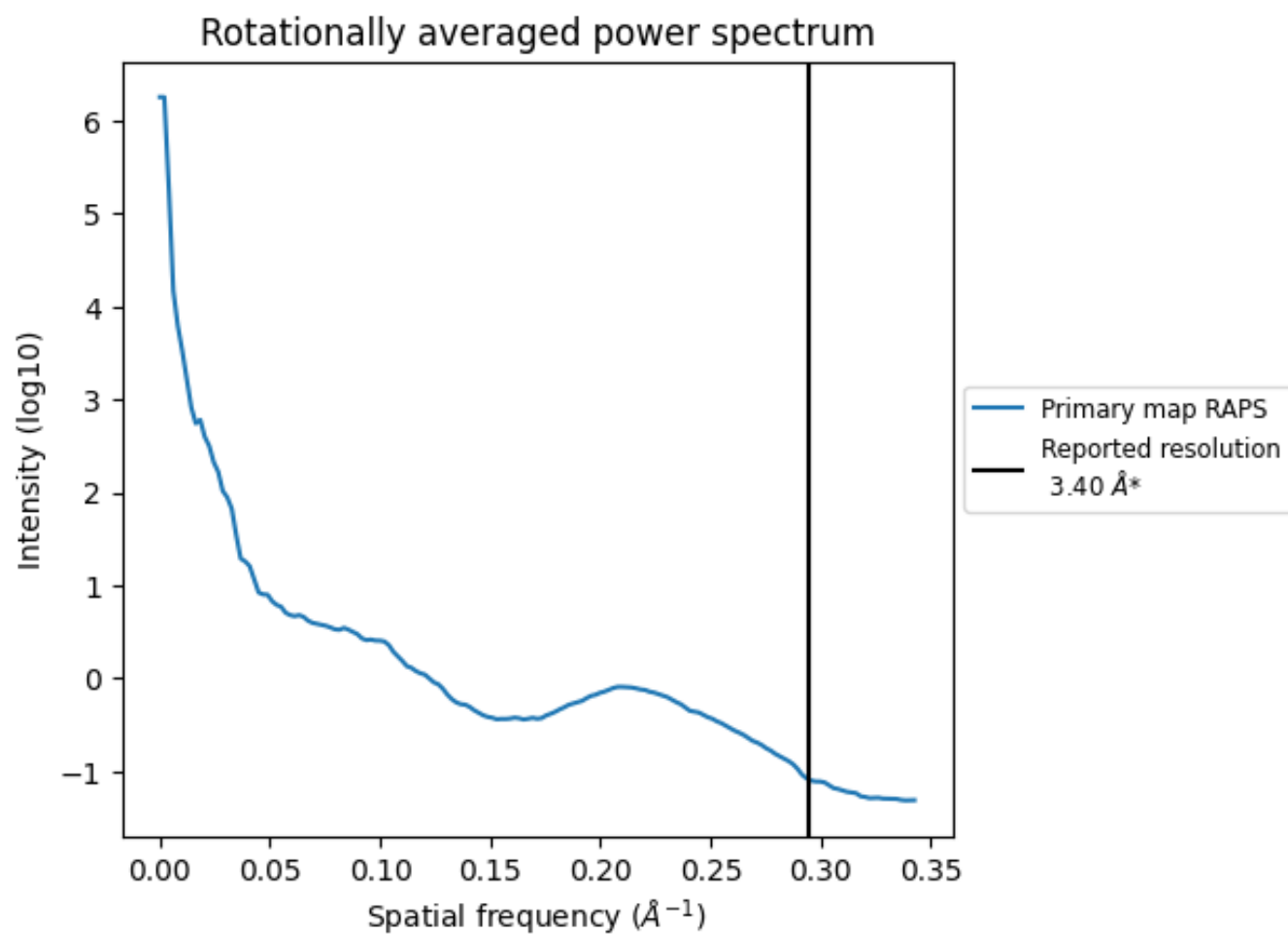
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 4230 nm³; this corresponds to an approximate mass of 3821 kDa.

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

7.3 Rotationally averaged power spectrum ⓘ

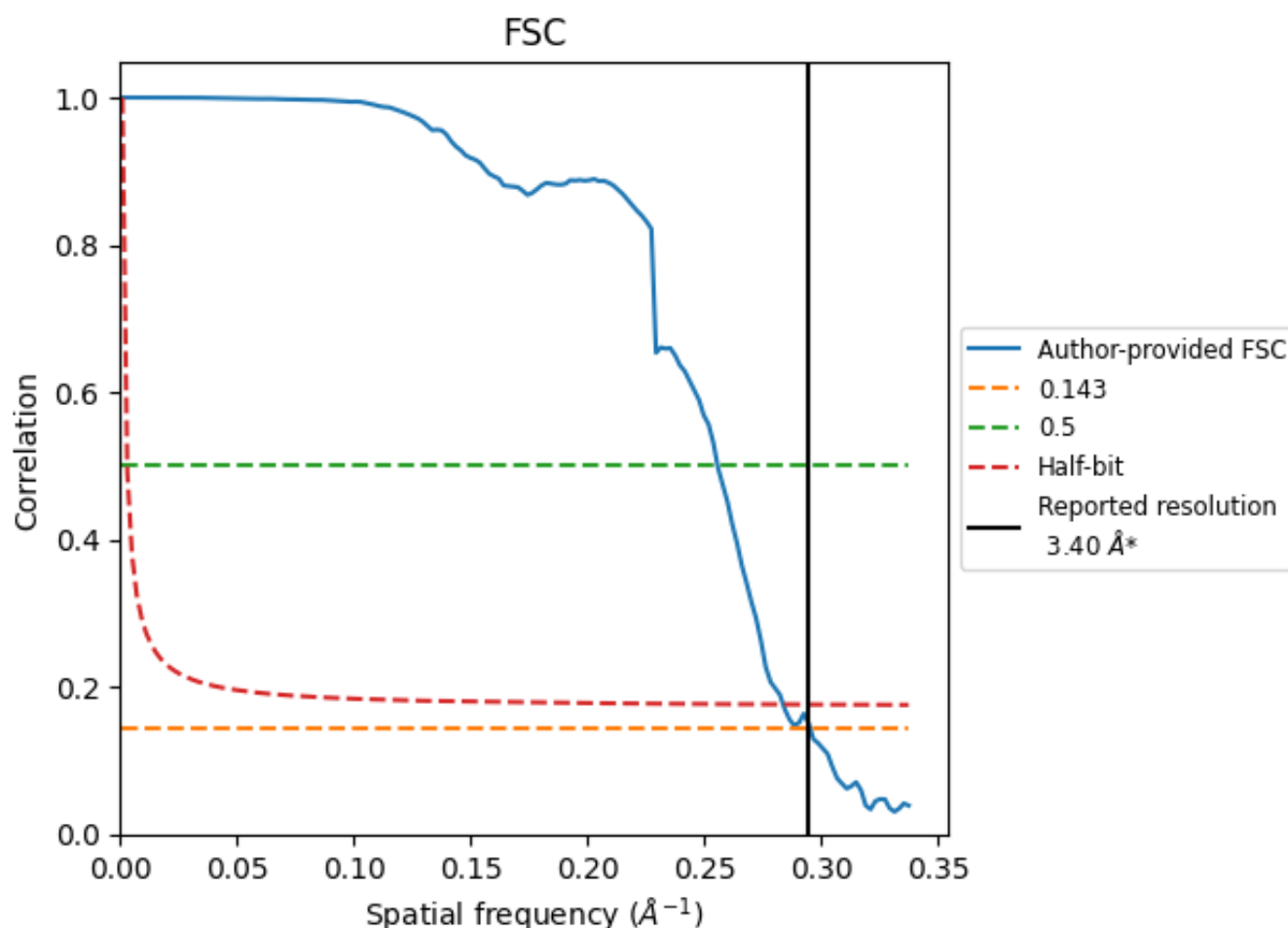


*Reported resolution corresponds to spatial frequency of 0.294 Å⁻¹

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.294 \AA^{-1}

8.2 Resolution estimates [i](#)

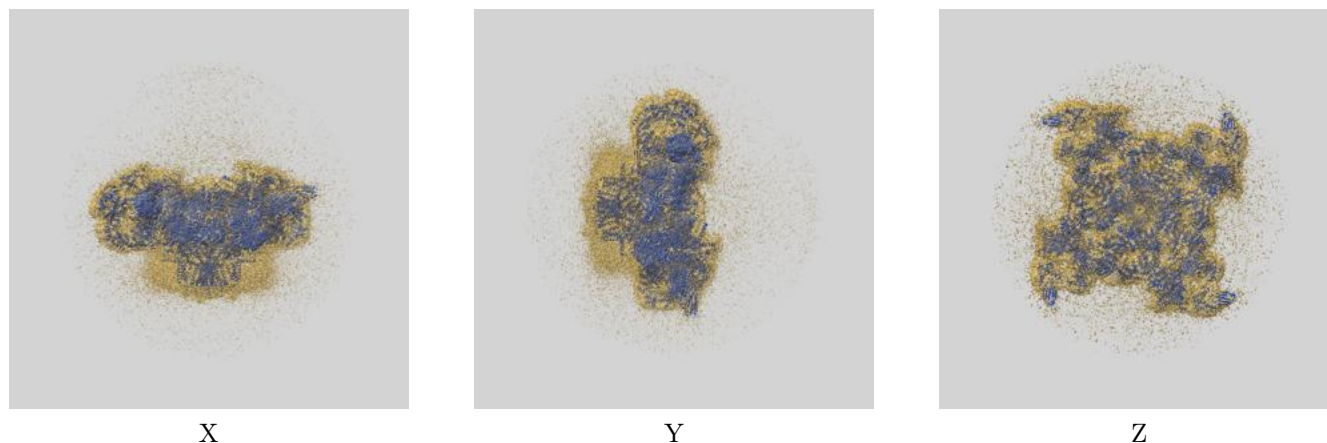
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.40	-	-
Author-provided FSC curve	3.39	3.91	3.52
Unmasked-calculated*	-	-	-

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps.

9 Map-model fit [i](#)

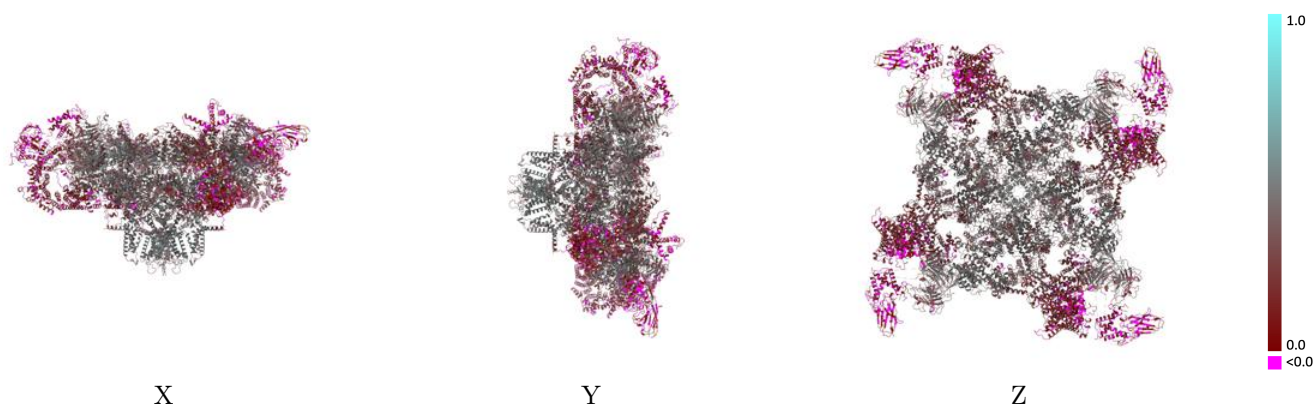
This section contains information regarding the fit between EMDB map EMD-19463 and PDB model 8RRS. Per-residue inclusion information can be found in section [3](#) on page [6](#).

9.1 Map-model overlay [i](#)



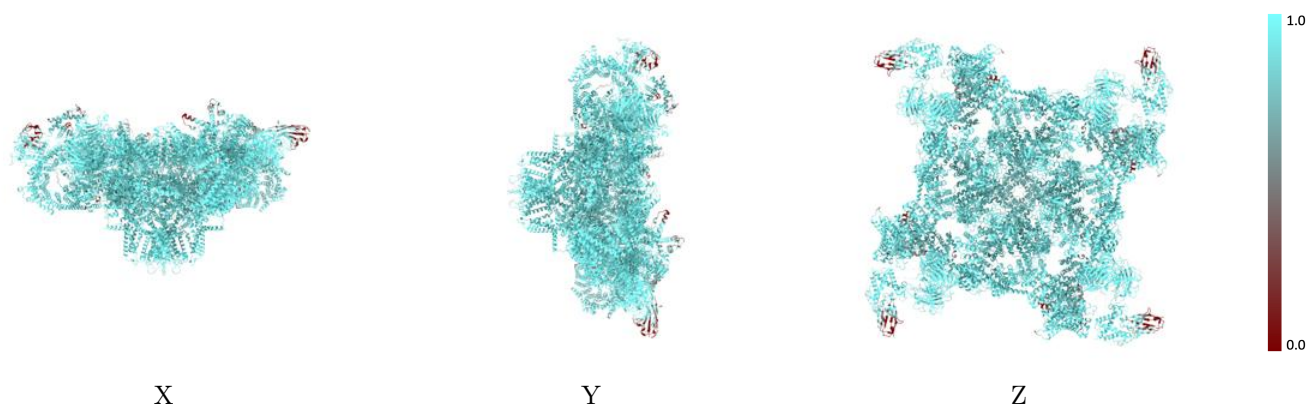
The images above show the 3D surface view of the map at the recommended contour level 0.35 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



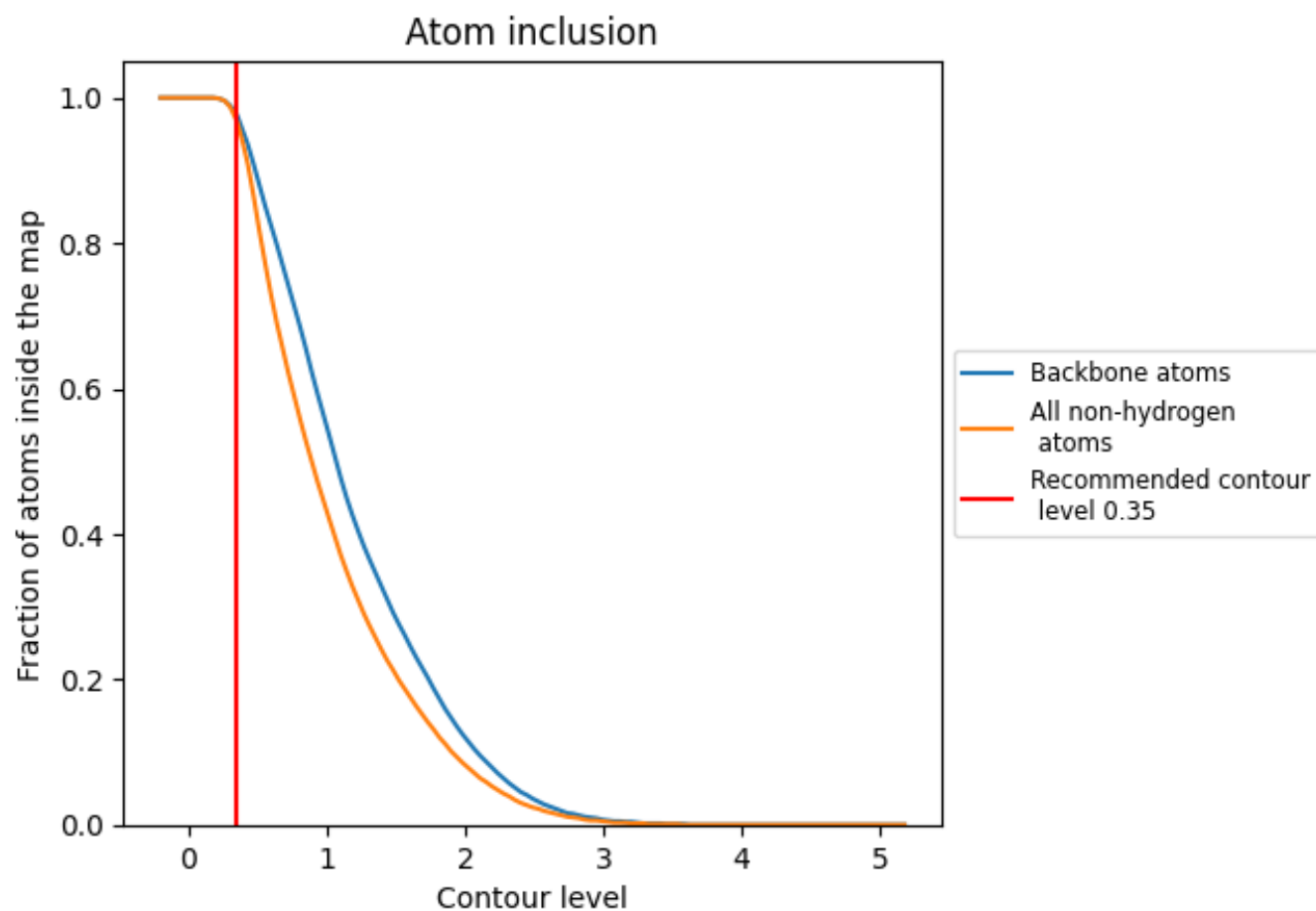
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.35).

9.4 Atom inclusion [i](#)



At the recommended contour level, 98% of all backbone atoms, 97% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.35) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	<div></div> 0.9670	<div></div> 0.3070
A	<div></div> 0.9780	<div></div> 0.3140
B	<div></div> 0.5750	<div></div> 0.0800
C	<div></div> 0.9780	<div></div> 0.3140
D	<div></div> 0.5720	<div></div> 0.0810
E	<div></div> 0.9780	<div></div> 0.3130
F	<div></div> 0.9780	<div></div> 0.3130
G	<div></div> 0.5750	<div></div> 0.0790
I	<div></div> 0.5750	<div></div> 0.0790

1.0

0.0

<0.0