



Full wwPDB EM Validation Report ⓘ

Nov 5, 2024 – 01:25 AM EST

PDB ID : 9E18
EMDB ID : EMD-47385
Title : Structure of RyR1 in the primed state in the presence of pentoxifylline
Authors : Miotto, M.C.; Marks, A.R.
Deposited on : 2024-10-21
Resolution : 2.68 Å (reported)
Based on initial model : 7TZC

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

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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.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) : 2.39

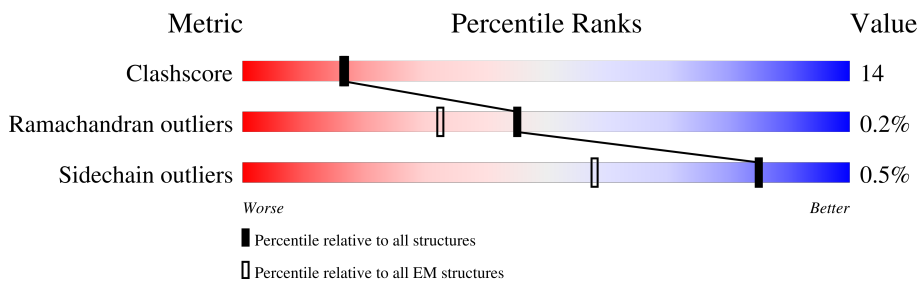
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 2.68 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	5037	
1	B	5037	
1	C	5037	
1	D	5037	
2	E	108	
2	F	108	
2	G	108	
2	H	108	

2 Entry composition

There are 7 unique types of molecules in this entry. The entry contains 144144 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 1.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
1	A	4404	Total 35150	C 22365	N 6063	O 6485	S 237	9	0
1	B	4404	Total 35150	C 22365	N 6063	O 6485	S 237	9	0
1	D	4404	Total 35150	C 22365	N 6063	O 6485	S 237	9	0
1	C	4404	Total 35150	C 22365	N 6063	O 6485	S 237	9	0

- Molecule 2 is a protein called Peptidyl-prolyl cis-trans isomerase FKBP1A.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
2	E	107	Total 831	C 527	N 146	O 154	S 4	0	0
2	H	107	Total 831	C 527	N 146	O 154	S 4	0	0
2	G	107	Total 831	C 527	N 146	O 154	S 4	0	0
2	F	107	Total 831	C 527	N 146	O 154	S 4	0	0

- 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	C	N	O	P	0
			31	10	5	13	3	
3	B	1	Total	C	N	O	P	0
			31	10	5	13	3	
3	D	1	Total	C	N	O	P	0
			31	10	5	13	3	
3	C	1	Total	C	N	O	P	0
			31	10	5	13	3	

- Molecule 4 is CALCIUM ION (three-letter code: CA) (formula: Ca).

Mol	Chain	Residues	Atoms		AltConf
4	A	1	Total	Ca	0
			1	1	
4	B	1	Total	Ca	0
			1	1	
4	D	1	Total	Ca	0
			1	1	
4	C	1	Total	Ca	0
			1	1	

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

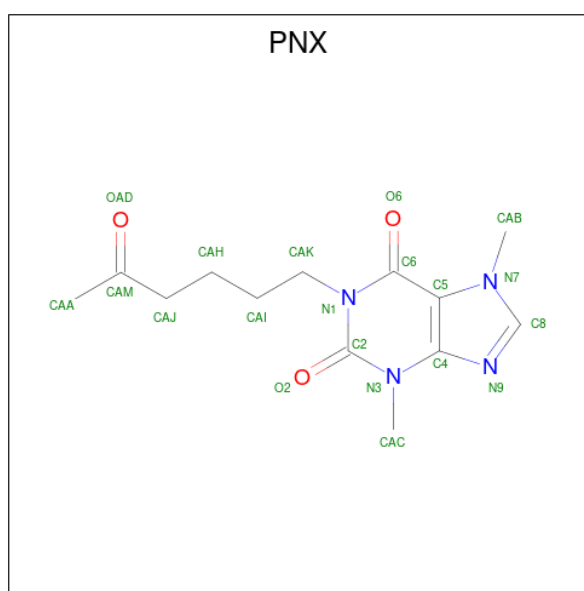
Mol	Chain	Residues	Atoms		AltConf
5	A	1	Total	Zn	0
			1	1	

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms		AltConf
5	B	1	Total	Zn	0
			1	1	
5	D	1	Total	Zn	0
			1	1	
5	C	1	Total	Zn	0
			1	1	

- Molecule 6 is 3,7-DIMETHYL-1-(5-OXOHEXYL)-3,7-DIHYDRO-1H-PURINE-2,6-DI ONE (three-letter code: PNX) (formula: C₁₃H₁₈N₄O₃) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms				AltConf
6	A	1	Total	C	N	O	0
			20	13	4	3	
6	B	1	Total	C	N	O	0
			20	13	4	3	
6	D	1	Total	C	N	O	0
			20	13	4	3	
6	C	1	Total	C	N	O	0
			20	13	4	3	

- Molecule 7 is water.

Mol	Chain	Residues	Atoms		AltConf
7	A	2	Total	O	0
			2	2	

Continued on next page...

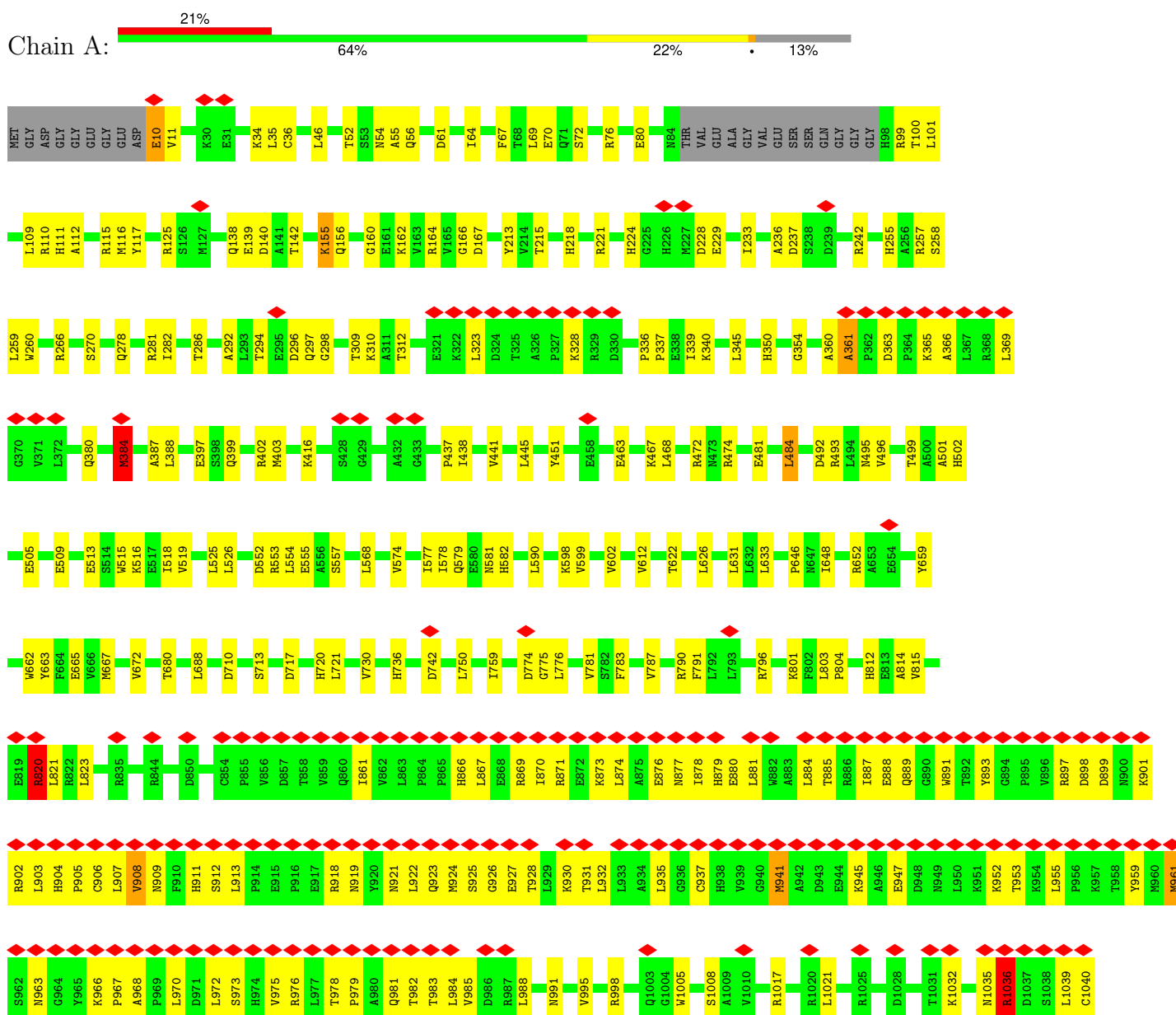
Continued from previous page...

Mol	Chain	Residues	Atoms	AltConf
7	B	2	Total O 2 2	0
7	D	2	Total O 2 2	0
7	C	2	Total O 2 2	0

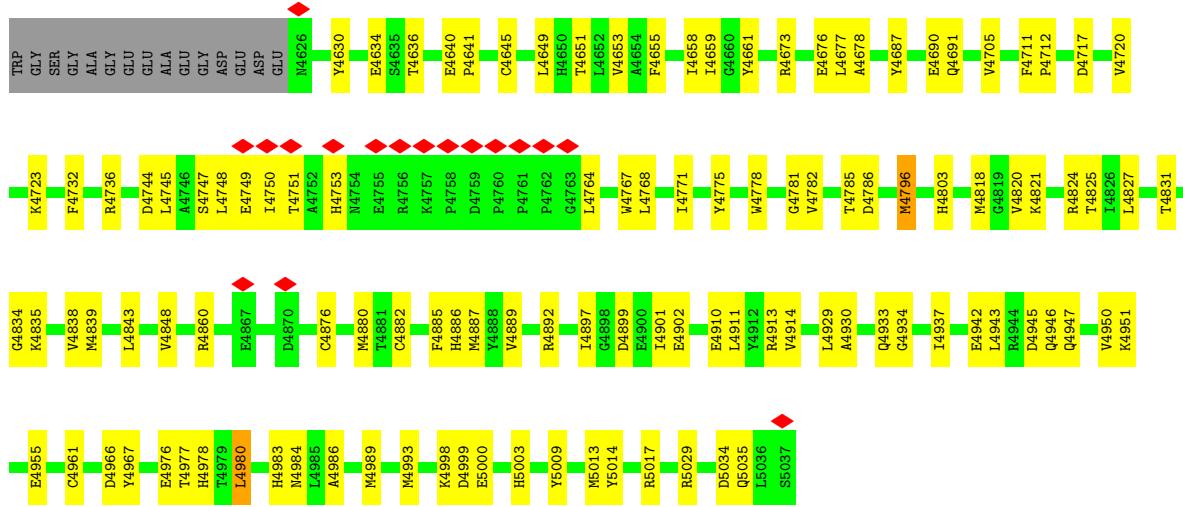
3 Residue-property plots

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.

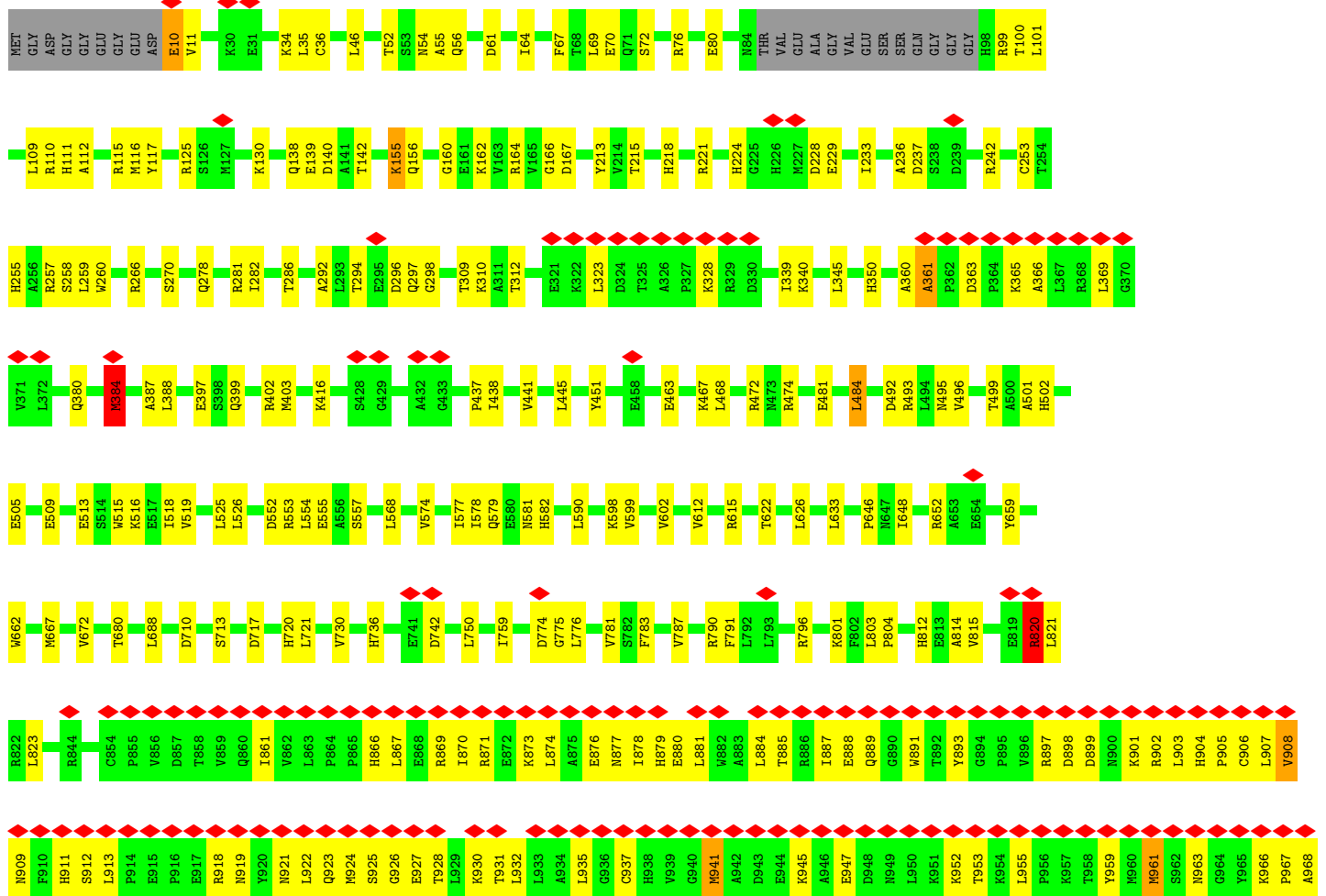
- Molecule 1: Ryanodine receptor 1

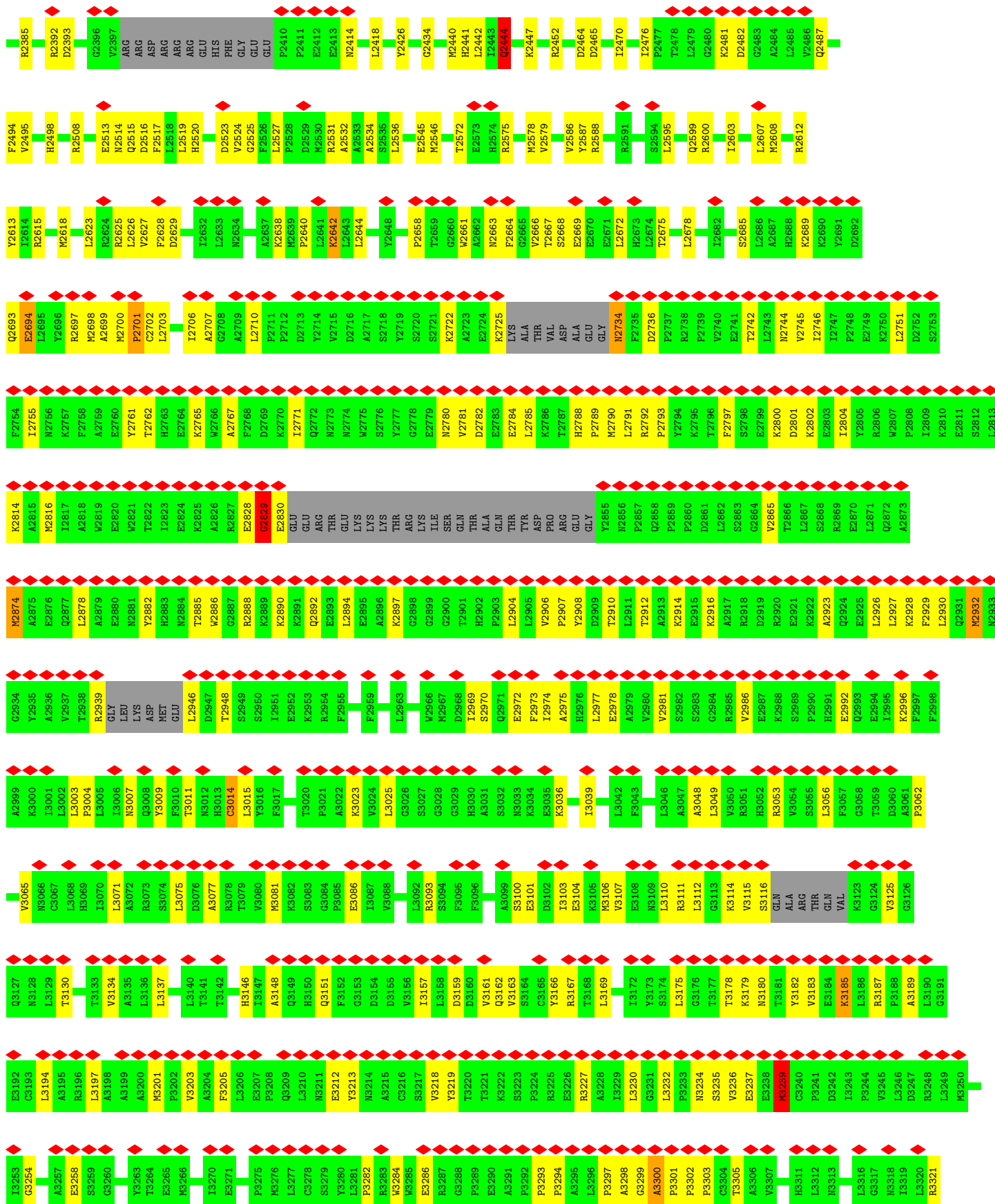


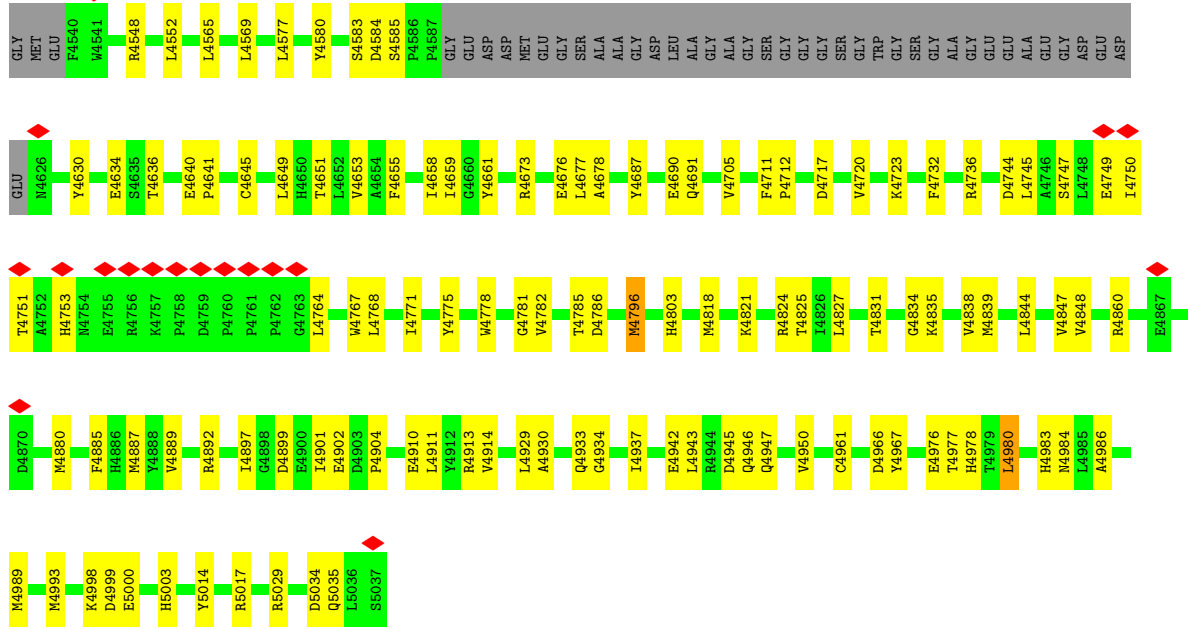
L2479	G2480	K2481	D2482	G2483	A2484	L2485	V2486	Q2487	F2494	V2495	H2498	R2508	E2513	N2514	Q2515	D2516	F2517	L2518	L2519	H2520	D2523	V2524	G2525	F2526	L2527	P2528	D2529	N2530	K2531	A2532	A2533	A2534	S2535	L2536	E2545	M2546	T2572	E2573	H2574	R2575	M2578	V2579	V2586	V2587	R2588	R2591	S2594	L2595													
Q2599	R2600	I2603	E2604	L2607	M2608	R2612	R2615	M2618	L2623	R2624	R2625	L2626	F2627	F2628	D2629	V2630	P2631	I2632	L2633	N2634	A2637	K2638	M2639	P2640	L2641	K2642	L2643	L2644	Y2648	P2658	T2659	G2660	V2661	A2662	N2663	F2664	G2665	V2666	T2667	S2668	E2669	E2670	E2671	L2672	H2673	L2674	T2675	L2678													
F2683	D2684	S2685	L2686	A2687	H2688	K2689	K2690	L2691	D2692	Q2693	E2694	L2695	Y2696	R2697	M2698	A2699	M2700	P2701	C2702	L2703	L2706	A2707	G2708	L2710	P2711	D2712	D2713	V2714	V2715	D2716	A2717	S2718	Y2719	S2720	S2721	K2722	A2723	E2724	K2725	LYS	ALA	THR	VAL	ASP	ALA	GLU	GLY	R2734	F2735	D2736	R2737	R2738	P2739	V2740	E2741	T2742					
L2743	N2744	V2745	I2746	L2747	P2748	E2749	K2750	L2751	D2752	S2753	F2754	L2755	N2756	K2757	F2758	A2759	E2760	Y2761	T2762	H2763	E2764	K2765	N2766	A2767	F2768	D2769	K2770	I2771	Q2772	Q2773	N2774	S2775	M2776	Y2777	G2778	E2779	N2780	V2781	D2782	E2783	E2784	L2785	K2786	H2787	T2788	P2789	M2790	L2791	R2792	F2793	Y2794	K2795	L2796	F2797	S2798	E2799	K2800	D2801	K2802		
E2803	L2804	Y2805	R2806	V2807	R2808	I2809	K2810	S2811	S2812	L2813	K2814	A2815	M2816	I2817	A2818	V2819	W2820	W2821	T2822	I2823	E2824	K2825	A2826	R2827	E2828	G2829	GLU	GLU	ARG	THR	ARG	LYS	LYS	THR	ARG	LYS	ILE	SER	GLN	THR	ALA	GLN	THR	TYR	ASP	PRO	ARG	GLU	GLY	Y2855	N2856	P2857	Q2858	P2859	P2860	K2861	L2862				
S2863	G2864	V2865	T2866	L2867	S2868	R2869	E2870	L2871	Q2872	A2873	M2874	A2875	E2876	Q2877	L2878	A2879	E2880	M2881	Y2882	H2883	N2884	T2885	M2886	G2887	K2888	K2889	K2890	K2891	Q2892	Q2893	E2894	F2895	E2896	A2897	A2898	K2899	G2898	G2899	G2900	T2901	H2902	F2903	L2904	L2905	V2906	P2907	Y2908	D2909	T2910	L2911	T2912	A2913	K2914	E2915	K2916	A2917	R2918	D2919	R2920	E2921	K2922
A2923	Q2924	E2925	L2926	L2927	K2928	F2929	L2930	Q2931	M2932	N2933	G2934	Y2935	A2936	V2937	T2938	R2939	LEU	LYS	ASP	MET	L2946	D2947	T2948	S2949	S2950	I2951	K2952	K2953	R2954	F2955	F2956	F2959	L2963	W2966	M2967	D2968	L2969	Q2971	E2972	F2973	I2974	A2975	Y2976	L2977	L2978	E2979	E2980	V2981	S2982	S2983	G2984	R2985	V2986	E2987							
K2988	S2989	P2990	H2991	E2992	Q2993	E2994	I2995	K2996	F2997	F2998	A2999	K3000	L3003	P3004	L3005	I3006	N3007	R3008	Y3009	T3011	N3012	H3013	C3014	L3015	Y3016	F3017	T3020	P3021	A3022	K3023	V3024	L3025	G3026	S3027	G3028	G3029	H3030	A3031	S3032	N3033	K3034	E3035	K3036	I3039	L3042	F3043	L3046	A3047	L3049	V3050	R3051	H3052									
R3053	V3054	S3055	L3056	F3057	G3058	T3059	D3060	A3061	P3062	V3065	H3066	C3067	L3068	H3069	I3070	A3071	R3072	K3073	S3074	L3075	D3076	T3077	T3079	V3080	M3081	K3082	S3083	G3084	P3085	E3086	I3087	V3088	L3092	R3093	S3094	F3095	F3096	A3099	S3100	E3101	E3102	I3103	E3104	K3105	M3106	V3107	E3108	N3109	L3110	R3111	L3112	G3113	V3114	V3115	S3116						
GLN	ALA	ARG	THR	GLN	VAL	K3123	G3124	V3125	G3126	Q3127	N3128	L3129	T3130	T3133	V3134	A3135	L3136	L3137	L3140	T3141	T3142	H3146	I3147	A3148	Q3149	H3150	Q3151	G3152	G3153	D3154	D3155	V3156	I3157	L3158	D3159	D3160	V3161	Q3162	V3163	S3164	C3165	V3166	R3167	T3168	L3169	I3172	V3173	S3174	L3175	G3176	T3177	T3178	K3179	N3180	T3181						
Y3182	V3183	E3184	K3185	L3186	R3187	P3188	A3189	L3190	G3191	E3192	G3193	L3194	A3195	R3196	L3197	A3199	A3200	M3201	P3202	V3203	A3204	F3205	L3206	E3207	P3208	Q3209	L3210	N3211	E3212	Y3213	N3214	A3215	C3216	S3217	V3218	Y3219	T3220	T3221	K3222	S3223	P3224	R3225	E3226	R3227	A3228	L3229	L3230	G3231	L3232	P3233	N3234	S3235	V3236	E3237	E3238	K3239	C3240	P3241			
D3242	I3243	P3244	V3245	L3246	D3247	R3248	L3249	M3250	I3253	G3254	G3255	L3256	A3257	E3258	S3259	Y3263	T3264	E3265	M3266	I3270	E3271	I3272	P3275	M3276	L3277	C3278	S3279	Y3280	L3281	P3282	R3283	W3284	W3285	E3286	R3287	G3288	P3289	E3290	A3291	P3292	P3293	P3294	A3295	L3296	P3297	A3298	G3299	A3300	P3301	P3302	P3303	C3304	T3305	A3306	V3307						
H3311	L3312	M3313	L3316	G3317	N3318	L3319	L3320	R3321	L3322	L3323	V3324	N3325	N3326	L3327	G3328	L3329	D3330	F3331	A3332	L3333	M3334	K3336	R3337	L3338	A3339	V3340	F3341	G3342	G3343	F3344	L3345	V3346	S3347	R3348	H3357	F3358	T3361	I3362	G3363	R3364	L3365	R3366	F3367	R3368	V3372	V3373	A3374	E3375	E3377	O3378	L3379	R3380									



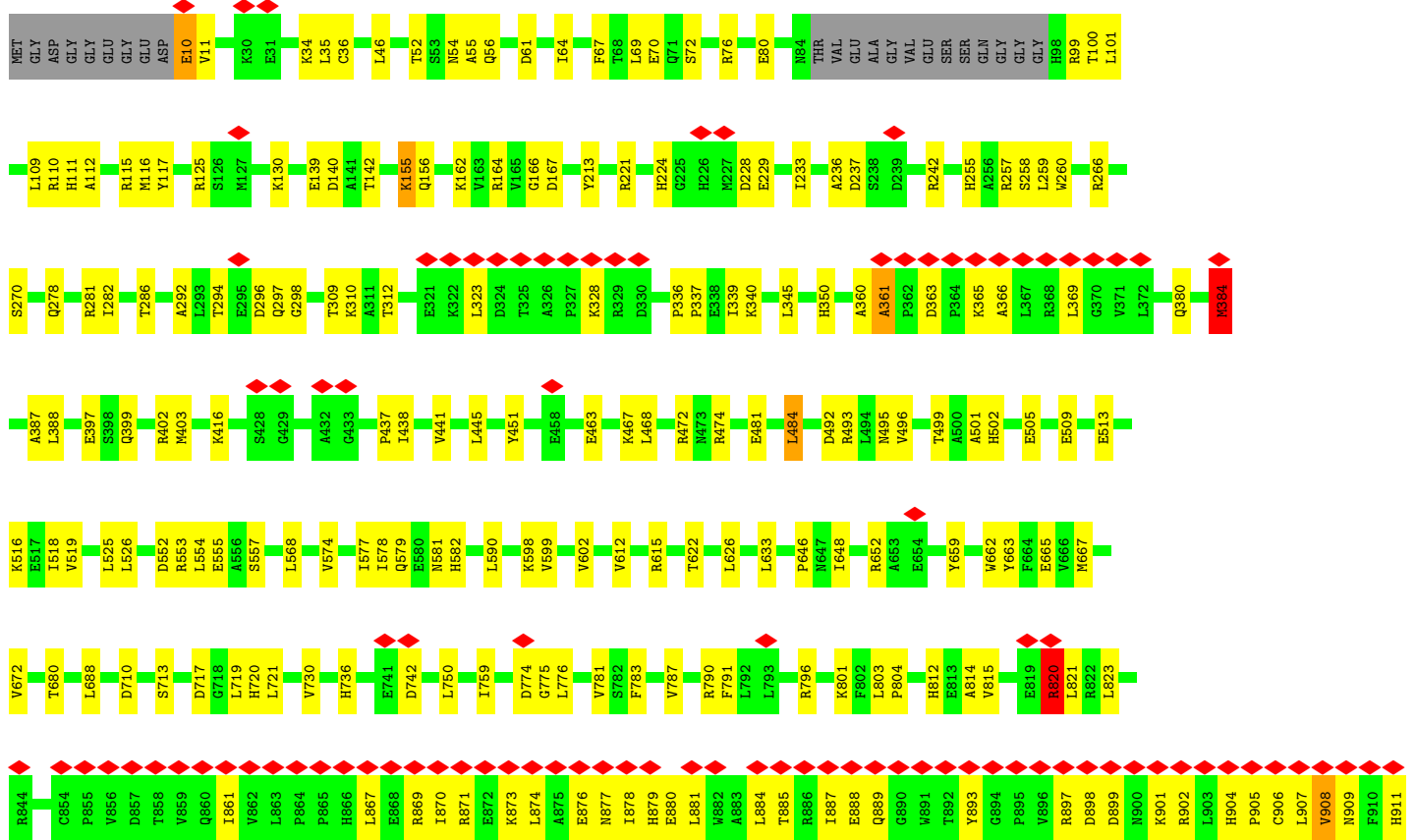
• Molecule 1: Ryanodine receptor 1

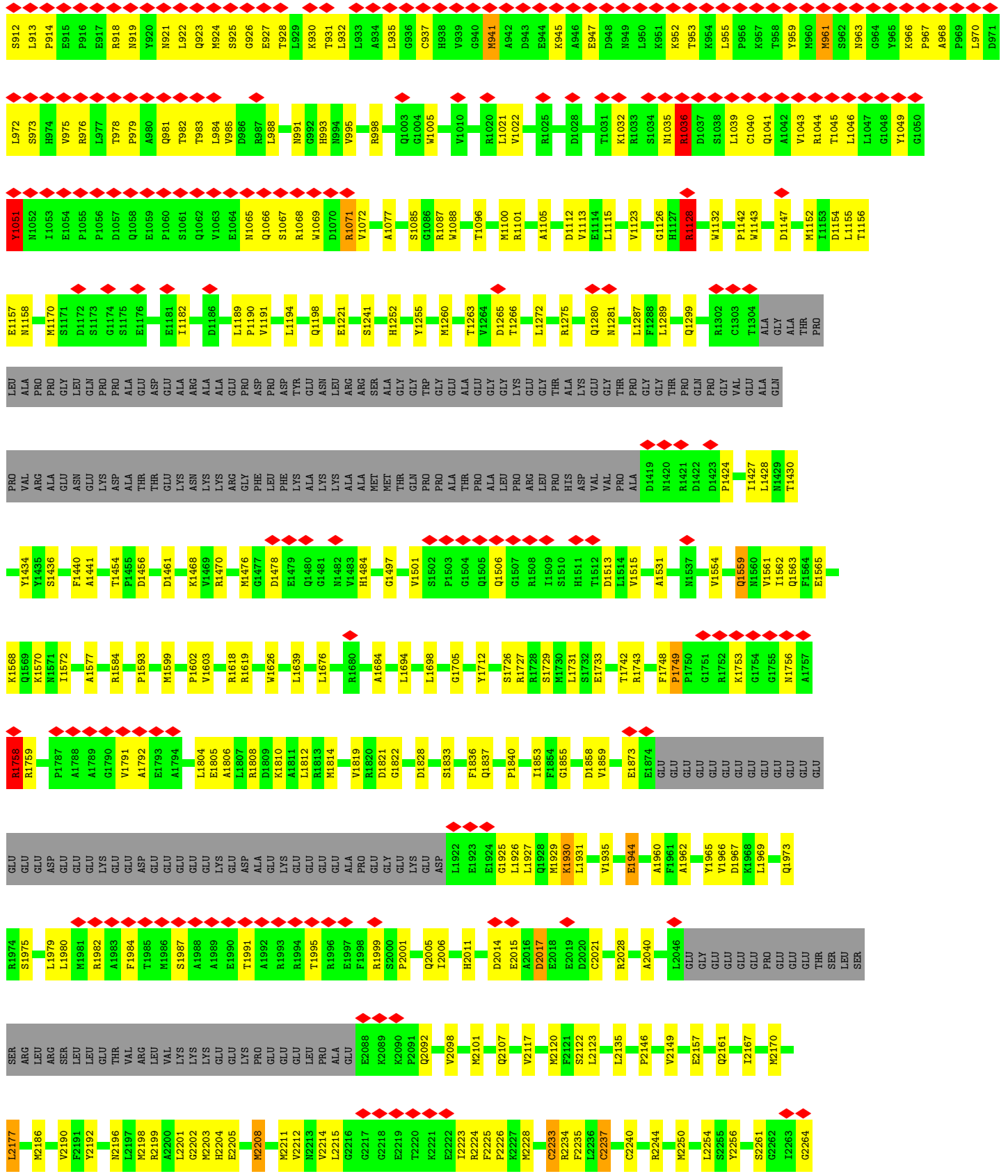




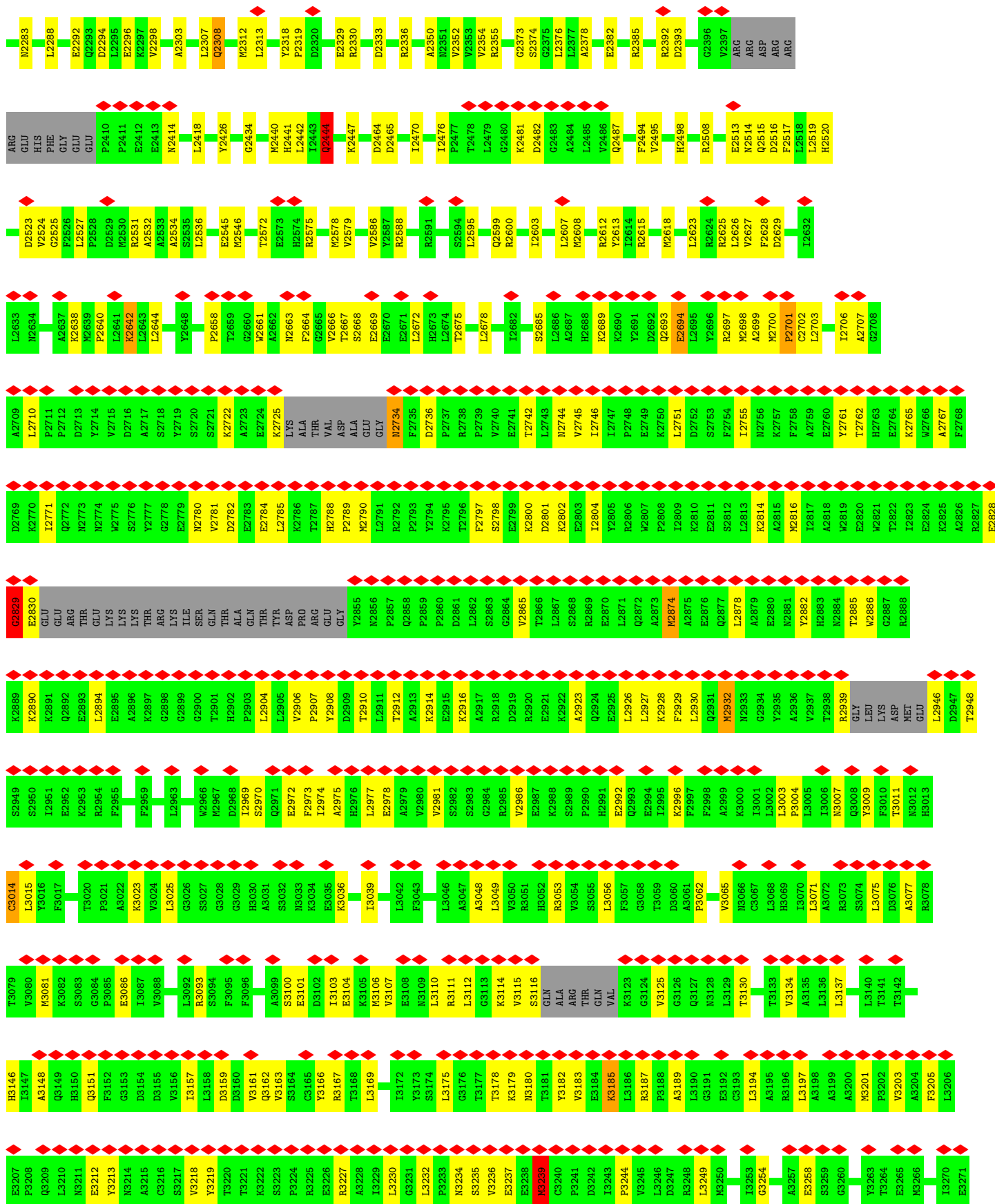


● Molecule 1: Ryanodine receptor 1

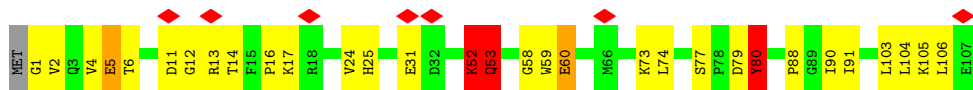




Y2192	SRR	E2088	L1980	V1791	S1436	ALA	M1170	Y1051	P969	N909
M2196	LEU	K2089	M1981	A1792	F1440	GLU	S1171	N1052	L970	F910
L2197	LEU	K2090	R1982	E1793	A1441	ASN	D1172	I1053	L971	H911
M2198	GLU	F2091	A1983	E1794	A1454	GLU	S1173	E1054	D971	S912
R2199	THR	Q2092	F1984	A1794	T1454	LYS	G1174	P1055	L972	S913
A2200	VAL	V2098	F1985	L1804	P1455	ASP	S1175	P1056	S973	L914
L2201	ARG	M2101	T1985	E1805	D1456	ALA	E1176	D1057	H974	P914
L2202	LEU	V2117	M1986	E1806	D1461	THR	E1181	Q1058	V975	E915
G2203	VAL	V2122	S1987	L1807	D1468	ASP	I1182	R1059	R976	P916
M2203	GLU	V2127	G1987	R1808	K1468	LYS	E1186	E1060	L977	E917
H2204	LYS	V2132	A1988	R1809	V1469	ASN	D1186	P1060	T978	R918
E2205	GLU	V2137	G1989	K1810	R1470	LYS	L1189	S1061	P979	N919
M2208	GLU	V2142	E1990	L1811	R1476	ALA	L1189	Q1062	A980	Y920
M2211	LYS	V2147	T1991	L1812	M1476	ARG	P1189	Q1063	Q981	N921
M2212	PRO	V2152	A1992	R1813	G1477	GLY	L1190	E1064	T982	L922
V2213	GLU	V2157	R1993	L1814	D1478	PHE	V1191	N1065	T983	Q923
V2214	GLU	V2162	A1994	M1814	E1479	LEU	L1194	Q1066	L984	M924
L2215	GLU	V2167	G1995	V1819	Q1480	PHE	Q1198	S1067	V985	S925
L2216	LEU	V2172	T1995	D1820	G1481	ALA	E1221	R1068	D986	G926
G2217	GLU	V2177	R1996	G1821	G1482	LYS	E1221	V1069	R987	E927
G2218	ALA	V2182	F1997	G1822	N1482	ALA	S1241	W1070	L988	T928
E2219	GLU	V2187	F1998	D1828	G1487	ALA	H1252	R1071	M991	L929
E2220	ALA	V2192	A1999	S1833	V1501	NET	H1252	I1072	V995	K930
K2221	PRO	V2197	S2000	F1836	S1502	NET	A1077	O1077	R998	T931
K2222	GLU	V2202	G2001	Q1837	P1503	THR	Y1255	S1085	L932	L933
E2223	GLU	V2207	Q2005	F1836	G1504	GLN	M1260	G1086	L933	L934
F2224	LYS	V2212	L2006	Q1837	Q1506	ALA	T1263	G1087	A934	A935
F2225	GLU	V2217	L2006	Q1837	Q1507	PRO	V1264	W1088	G936	G937
P2226	GLU	V2222	L2006	Q1837	R1508	ALA	V1266	W1005	L938	L939
K2227	ASP	V2227	H2011	P1840	G1509	LEU	L1272	V1010	H938	H939
K2228	GLU	V2232	D2014	I1853	G1507	THR	L1272	V1010	V939	V940
I2223	GLU	V2237	E2015	F1854	R1728	ALA	R1275	R1020	G940	G941
R2224	GLU	V2242	A2016	G1855	R1729	PRO	Q1280	L1021	M941	M942
F2225	LYS	V2247	E2017	D1856	M1730	ALA	N1281	L1113	A942	A943
F2235	GLU	V2252	E2018	E1857	L1731	VAL	M1281	L1114	D943	D944
L2236	GLU	V2257	E2019	D1858	L1732	VAL	L1287	L1115	E944	E945
C2237	GLU	V2262	E2020	K1860	E1733	VAL	L1287	L1123	A946	A947
R2244	GLU	V2267	D2020	K1860	T1742	VAL	L1287	V1126	D948	D949
L2254	GLU	V2272	C2021	P1868	R1743	VAL	L1287	G1126	N949	N950
L2255	GLU	V2277	R2028	T1872	F1749	ALA	L1287	H1127	L950	L951
S2255	GLU	V2282	A2040	E1873	P1750	ALA	L1288	R1128	K951	K952
Y2256	GLU	V2287	L2046	E1874	G1751	ALA	L1289	R1128	K953	K954
S2261	GLU	V2292	L2046	GLU	R1752	ALA	Q1299	R1128	L955	L956
G2262	GLU	V2297	L2046	GLU	R1753	ALA	R1302	R1128	L957	L958
I2263	GLU	V2302	L2046	GLU	K1753	ALA	C1303	R1128	L959	L960
G2264	GLU	V2307	L2046	GLU	G1754	ALA	T1304	R1128	M961	M962
L2265	GLU	V2312	L2046	GLU	G1755	ALA	ALA	R1128	N963	N964
L2266	GLU	V2317	L2046	GLU	N1756	ALA	GLY	R1128	G964	G965
G2266	GLU	V2322	L2046	GLU	N1756	ALA	GLY	R1128	K966	K967
M2267	PRO	V2327	L2046	GLU	A1757	ALA	ALA	R1128	A968	A969
M2267	GLU	V2332	L2046	GLU	R1758	ALA	ALA	R1128		
T2271	GLU	V2337	L2046	GLU	R1759	ALA	ALA	R1128		
T2271	THR	V2342	L2046	GLU	P1787	ALA	GLY	R1128		
V2275	SER	V2347	L2046	GLU	A1788	ALA	THR	R1128		
A2276	SER	V2352	L2046	GLU	A1789	ALA	THR	R1128		
V2280	SER	V2357	L2046	GLU	G1790	ALA	THR	R1128		



PRO	GLY	L3408	GLN	R3550	SS615	A3796	M3858	L9880	V4081	T4241	ARG	ALA
GLY	LEU	Y3409	SER	E3551	K3616	D3727	V3859	L3980	F4093	M4245	SER	LYS
ALA	VAL	P3410	GLY	F3552	K3617	M3728	M3860	L3985	K4094	M4254	ARG	LYS
THR	THR	L3277	GLY	L3553	A3618	M3729	E3861	L3988	K4095	G4254	ARG	VAL
VAL	VAL	C3278	ASP	Q3554	V3619	K3731	D3862	V3989	A4096	GLU	ARG	THR
VAL	VAL	S3279	ASP	M3555	V3620	S3732	G3863	V3990	M4097	GLY	ARG	THR
ALA	ALA	Y3280	GLN	N3556	H3621	C3733	T3864	G3991	M4098	PRU	ARG	LEU
ASP	THR	P3281	ARG	L3557	K3622	H3734	V3865	V3995	S4099	ALA	ARG	LEU
GLY	THR	L3282	THR	L3558	L3623	L3735	V3866	F3996	Q4100	ALA	ARG	ALA
THR	LYS	R3283	LYS	H3558	L3624	E3736	I3867	A3997	I4108	GLU	ARG	GLY
ASP	LYS	H3284	LYS	L3559	SS625	GLU	R3868	M4000	D4118	ASP	THR	PRO
GLY	LYS	H3285	LYS	Q3560	K3626	GLY	Q3869	L4003	D4119	GLY	ALA	ASP
GLY	LYS	E3286	LYS	R3498	Q3627	GLY	Q3870	L4006	M4122	GLY	ALA	PRO
GLY	LYS	R3287	R3499	G3561	Q3628	GLY	M3871	D4006	I4139	GLY	ALA	PRO
GLY	LYS	G3288	G3500	K3562	R3628	ASN	V3874	S4007	M4142	GLY	ALA	VAL
GLY	LYS	P3289	D3501	V3563	R3629	GLU	D3877	S4008	T4148	GLY	ALA	VAL
GLY	LYS	E3290	R3502	V3564	R3630	ALA	D3878	Q4009	E4161	GLY	ALA	VAL
GLY	LYS	A3291	Y3503	G3565	R3631	GLU	E3879	L4016	E4172	GLY	ALA	VAL
GLY	LYS	P3292	S3504	S3566	V3632	GLU	F3880	L4017	R4175	GLY	ALA	VAL
GLY	LYS	P3293	Q3505	S3567	V3633	GLU	V3883	E4032	R4176	GLY	ALA	VAL
GLY	LYS	P3294	Q3506	S3568	A3634	GLU	D3888	M4039	P4176	GLY	ALA	VAL
GLY	LYS	A3295	T3507	R3570	A3635	GLU	L3891	E4041	G4179	GLY	ALA	VAL
GLY	LYS	R3296	S3508	M3573	F3636	GLU	L3892	R4042	R4180	GLY	ALA	VAL
GLY	LYS	P3297	L3509	A3574	R3637	GLU	C3897	V4045	E4181	GLY	ALA	VAL
GLY	LYS	A3298	I3510	G3577	V3638	GLU	M3897	V4049	E4182	GLY	ALA	VAL
GLY	LYS	G3299	V3511	G3578	P3640	GLU	D3898	M4057	E4183	GLY	ALA	VAL
GLY	LYS	A3300	L3514	C3579	L3641	GLU	F3899	A4041	E4187	GLY	ALA	VAL
GLY	LYS	P3301	M3517	P3580	Y3642	GLU	Q3900	R4042	E4188	GLY	ALA	VAL
GLY	LYS	P3302	G3521	G3581	N3651	GLU	L3901	V4042	E4189	GLY	ALA	VAL
GLY	LYS	P3303	L3522	R3582	E3655	GLU	L3903	V4045	E4192	GLY	ALA	VAL
GLY	LYS	C3304	N3523	E3583	K3658	GLU	R3904	V4049	E4197	GLY	ALA	VAL
GLY	LYS	T3305	K3524	E3584	A3659	GLU	T3910	M4057	E4205	GLY	ALA	VAL
GLY	LYS	A3306	R3524	D3585	A3660	GLU	T3911	M4057	E4209	GLY	ALA	VAL
GLY	LYS	V3307	L3526	A3586	V3662	GLU	I3916	M4057	E4222	GLY	ALA	VAL
GLY	LYS	H3311	P3527	D3587	L3663	GLU	D3941	K4060	G4225	GLY	ALA	VAL
GLY	LYS	L3312	T3528	D3588	L3663	GLU	V3942	L4068	G4226	GLY	ALA	VAL
GLY	LYS	H3313	T3529	P3589	D3671	GLU	L3943	D4063	E4227	GLY	ALA	VAL
GLY	LYS	L3316	D3531	K3591	K3672	GLU	Q3946	M4064	M4231	GLY	ALA	VAL
GLY	LYS	G3317	L3532	I3592	E3682	GLU	G3947	K4067	E4232	GLY	ALA	VAL
GLY	LYS	H3318	I3533	V3593	Q3683	GLU	E3947	L4068	A4075	GLY	ALA	VAL
GLY	LYS	L3319	K3534	R3594	K3684	GLU	K3948	L4069	F4077	GLY	ALA	VAL
GLY	LYS	L3320	L3535	R3594	E3684	GLU	R3949	D4070	G4235	GLY	ALA	VAL
GLY	LYS	R3321	K3536	E3598	E3685	GLU	V3957	M4072		GLY	ALA	VAL
GLY	LYS	I3322	T3538	V3600	E3686	GLU	V3961	V4075		GLY	ALA	VAL
GLY	LYS	I3323	T3538	L3603	E3687	GLU	V3966	A4076		GLY	ALA	VAL
GLY	LYS	V3324	R3539	Y3604	E3688	GLU	T3966	F4077		GLY	ALA	VAL
GLY	LYS	N3325	R3395	L3605	E3689	GLU	I3969			GLY	ALA	VAL
GLY	LYS	H3326	D3396	H3605	V3690	GLU	G3971			GLY	ALA	VAL
GLY	LYS	L3327	E3397	E3607	E3691	GLU	P3972			GLY	ALA	VAL
GLY	LYS	G3328	F3398	T3609	K3693	GLU				GLY	ALA	VAL
GLY	LYS	D3329	L3401	T3545	K3694	GLU				GLY	ALA	VAL
GLY	LYS	D3330	C3402	T3546		GLU				GLY	ALA	VAL
GLY	LYS	E3331	R3403	E3546		GLU				GLY	ALA	VAL
GLY	LYS	A3332	R3403	E3547		GLU				GLY	ALA	VAL
GLY	LYS	T3333	D3404	E3548		GLU				GLY	ALA	VAL
GLY	LYS	W3334	L3405	V3549		GLU				GLY	ALA	VAL
GLY	LYS	H3335	A3407			GLU				GLY	ALA	VAL



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C4	Depositor
Number of particles used	64353	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	58	Depositor
Minimum defocus (nm)	500	Depositor
Maximum defocus (nm)	1500	Depositor
Magnification	Not provided	
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	0.555	Depositor
Minimum map value	-0.247	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.022	Depositor
Recommended contour level	0.1	Depositor
Map size (Å)	428.544, 428.544, 428.544	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.837, 0.837, 0.837	Depositor

5 Model quality i

5.1 Standard geometry i

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

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	0.54	50/35977 (0.1%)	0.67	65/48726 (0.1%)
1	B	0.54	50/35977 (0.1%)	0.67	65/48726 (0.1%)
1	C	0.54	49/35977 (0.1%)	0.67	65/48726 (0.1%)
1	D	0.54	50/35977 (0.1%)	0.67	65/48726 (0.1%)
2	E	2.56	15/850 (1.8%)	3.24	24/1146 (2.1%)
2	F	2.56	15/850 (1.8%)	3.25	24/1146 (2.1%)
2	G	2.56	15/850 (1.8%)	3.24	24/1146 (2.1%)
2	H	2.56	15/850 (1.8%)	3.24	24/1146 (2.1%)
All	All	0.66	259/147308 (0.2%)	0.82	356/199488 (0.2%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	14
1	B	0	14
1	C	0	14
1	D	0	14
2	E	0	5
2	F	0	5
2	G	0	5
2	H	0	5
All	All	0	76

All (259) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	H	80	TYR	CD1-CE1	47.33	2.10	1.39
2	F	80	TYR	CD1-CE1	47.31	2.10	1.39

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	E	80	TYR	CD1-CE1	47.30	2.10	1.39
2	G	80	TYR	CD1-CE1	47.26	2.10	1.39
2	G	80	TYR	CD2-CE2	36.38	1.94	1.39
2	H	80	TYR	CD2-CE2	36.34	1.93	1.39
2	F	80	TYR	CD2-CE2	36.33	1.93	1.39
2	E	80	TYR	CD2-CE2	36.32	1.93	1.39
1	A	2237	CYS	CB-SG	31.27	2.35	1.82
1	D	2237	CYS	CB-SG	31.27	2.35	1.82
1	B	2237	CYS	CB-SG	31.25	2.35	1.82
1	C	2237	CYS	CB-SG	31.25	2.35	1.82
1	A	2233	CYS	CB-SG	-30.56	1.30	1.82
1	D	2233	CYS	CB-SG	-30.56	1.30	1.82
1	C	2233	CYS	CB-SG	-30.56	1.30	1.82
1	B	2233	CYS	CB-SG	-30.53	1.30	1.82
2	F	52	LYS	CE-NZ	-23.43	0.90	1.49
2	G	52	LYS	CE-NZ	-23.40	0.90	1.49
2	H	52	LYS	CE-NZ	-23.39	0.90	1.49
2	E	52	LYS	CE-NZ	-23.38	0.90	1.49
1	D	1036	ARG	CZ-NH1	-22.60	1.03	1.33
1	A	1036	ARG	CZ-NH1	-22.58	1.03	1.33
1	B	1036	ARG	CZ-NH1	-22.58	1.03	1.33
1	C	1036	ARG	CZ-NH1	-22.58	1.03	1.33
1	B	2444	GLN	CG-CD	-19.03	1.07	1.51
1	D	2444	GLN	CG-CD	-19.03	1.07	1.51
1	A	2444	GLN	CG-CD	-19.01	1.07	1.51
1	C	2444	GLN	CG-CD	-19.01	1.07	1.51
1	A	3239	MET	CB-CG	18.46	2.10	1.51
1	D	3239	MET	CB-CG	18.46	2.10	1.51
1	B	3239	MET	CB-CG	18.45	2.10	1.51
1	C	3239	MET	CB-CG	18.45	2.10	1.51
2	H	80	TYR	CE1-CZ	-16.76	1.16	1.38
2	E	80	TYR	CE1-CZ	-16.74	1.16	1.38
2	G	80	TYR	CE1-CZ	-16.70	1.16	1.38
2	F	80	TYR	CE1-CZ	-16.69	1.16	1.38
1	B	1036	ARG	CB-CG	-16.04	1.09	1.52
1	D	1036	ARG	CB-CG	-16.02	1.09	1.52
1	A	1036	ARG	CB-CG	-16.01	1.09	1.52
1	C	1036	ARG	CB-CG	-16.01	1.09	1.52
1	B	2642	LYS	CG-CD	15.61	2.05	1.52
1	C	2642	LYS	CG-CD	15.60	2.05	1.52
1	A	2642	LYS	CG-CD	15.60	2.05	1.52
1	D	2642	LYS	CG-CD	15.58	2.05	1.52

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	H	80	TYR	CB-CG	-12.74	1.32	1.51
2	E	80	TYR	CB-CG	-12.71	1.32	1.51
2	F	80	TYR	CB-CG	-12.70	1.32	1.51
2	G	80	TYR	CB-CG	-12.70	1.32	1.51
1	A	1036	ARG	CG-CD	-12.48	1.20	1.51
1	B	1036	ARG	CG-CD	-12.48	1.20	1.51
1	C	1036	ARG	CG-CD	-12.48	1.20	1.51
1	D	1036	ARG	CG-CD	-12.47	1.20	1.51
1	C	416	LYS	CD-CE	-12.34	1.20	1.51
1	A	416	LYS	CD-CE	-12.32	1.20	1.51
1	B	416	LYS	CD-CE	-12.32	1.20	1.51
1	D	416	LYS	CD-CE	-12.29	1.20	1.51
1	D	820	ARG	CZ-NH1	12.21	1.49	1.33
1	A	820	ARG	CZ-NH1	12.19	1.48	1.33
1	B	820	ARG	CZ-NH1	12.19	1.48	1.33
1	C	820	ARG	CZ-NH1	12.19	1.48	1.33
1	A	3014	CYS	CB-SG	-11.73	1.62	1.82
1	D	3014	CYS	CB-SG	-11.73	1.62	1.82
1	B	3014	CYS	CB-SG	-11.70	1.62	1.82
1	C	3014	CYS	CB-SG	-11.70	1.62	1.82
1	C	3239	MET	CG-SD	11.04	2.09	1.81
1	A	3239	MET	CG-SD	11.01	2.09	1.81
1	B	3239	MET	CG-SD	10.99	2.09	1.81
1	D	3239	MET	CG-SD	10.99	2.09	1.81
1	D	2308	GLN	CB-CG	-10.38	1.24	1.52
1	C	2308	GLN	CB-CG	-10.37	1.24	1.52
1	A	2308	GLN	CB-CG	-10.37	1.24	1.52
1	B	2308	GLN	CB-CG	-10.34	1.24	1.52
2	E	5	GLU	CD-OE1	-10.05	1.14	1.25
2	H	5	GLU	CD-OE1	-10.05	1.14	1.25
2	G	5	GLU	CD-OE1	-10.02	1.14	1.25
2	F	5	GLU	CD-OE1	-10.00	1.14	1.25
2	G	53	GLN	CG-CD	10.00	1.74	1.51
2	H	53	GLN	CG-CD	9.97	1.74	1.51
2	F	53	GLN	CG-CD	9.97	1.74	1.51
2	E	53	GLN	CG-CD	9.94	1.74	1.51
1	B	3557	LEU	CG-CD1	-9.73	1.15	1.51
1	A	3557	LEU	CG-CD1	-9.71	1.16	1.51
1	C	3557	LEU	CG-CD1	-9.71	1.16	1.51
1	D	3557	LEU	CG-CD1	-9.69	1.16	1.51
1	A	1036	ARG	NE-CZ	-9.63	1.20	1.33
1	C	1036	ARG	NE-CZ	-9.63	1.20	1.33

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	B	1036	ARG	NE-CZ	-9.59	1.20	1.33
1	D	1036	ARG	NE-CZ	-9.59	1.20	1.33
1	D	1036	ARG	CD-NE	-9.30	1.30	1.46
1	B	1036	ARG	CD-NE	-9.30	1.30	1.46
1	A	1036	ARG	CD-NE	-9.29	1.30	1.46
1	C	1036	ARG	CD-NE	-9.29	1.30	1.46
1	D	2308	GLN	CD-OE1	-9.26	1.03	1.24
1	A	2308	GLN	CD-OE1	-9.23	1.03	1.24
1	B	2308	GLN	CD-OE1	-9.23	1.03	1.24
1	C	2308	GLN	CD-OE1	-9.23	1.03	1.24
2	F	53	GLN	CB-CG	-9.15	1.27	1.52
2	H	53	GLN	CB-CG	-9.14	1.27	1.52
2	G	53	GLN	CB-CG	-9.14	1.27	1.52
2	E	53	GLN	CB-CG	-9.12	1.27	1.52
1	B	941	MET	CG-SD	-8.89	1.58	1.81
1	A	941	MET	CG-SD	-8.88	1.58	1.81
1	D	941	MET	CG-SD	-8.88	1.58	1.81
1	C	941	MET	CG-SD	-8.88	1.58	1.81
1	A	1559	GLN	CD-NE2	-8.86	1.10	1.32
1	B	1559	GLN	CD-NE2	-8.86	1.10	1.32
1	D	1559	GLN	CD-NE2	-8.86	1.10	1.32
1	C	1559	GLN	CD-NE2	-8.86	1.10	1.32
2	H	80	TYR	CE2-CZ	-8.82	1.27	1.38
2	E	80	TYR	CE2-CZ	-8.82	1.27	1.38
2	G	80	TYR	CE2-CZ	-8.81	1.27	1.38
2	F	80	TYR	CE2-CZ	-8.80	1.27	1.38
2	H	105	LYS	CE-NZ	-8.62	1.27	1.49
1	B	2308	GLN	CD-NE2	-8.60	1.11	1.32
2	G	105	LYS	CE-NZ	-8.59	1.27	1.49
2	E	105	LYS	CE-NZ	-8.59	1.27	1.49
1	A	2308	GLN	CD-NE2	-8.58	1.11	1.32
2	F	105	LYS	CE-NZ	-8.58	1.27	1.49
1	D	2308	GLN	CD-NE2	-8.57	1.11	1.32
1	C	2308	GLN	CD-NE2	-8.57	1.11	1.32
1	C	2444	GLN	CB-CG	-8.06	1.30	1.52
1	A	2444	GLN	CB-CG	-8.05	1.30	1.52
1	B	2444	GLN	CB-CG	-8.04	1.30	1.52
1	D	2444	GLN	CB-CG	-8.04	1.30	1.52
1	C	1051	TYR	CD1-CE1	8.01	1.51	1.39
1	A	1051	TYR	CD1-CE1	7.96	1.51	1.39
1	D	1051	TYR	CD1-CE1	7.96	1.51	1.39
2	H	80	TYR	CZ-OH	-7.93	1.24	1.37

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	E	80	TYR	CZ-OH	-7.93	1.24	1.37
2	G	80	TYR	CZ-OH	-7.91	1.24	1.37
1	B	1051	TYR	CD1-CE1	7.91	1.51	1.39
2	F	80	TYR	CZ-OH	-7.90	1.24	1.37
1	B	2177	LEU	CB-CG	-7.81	1.29	1.52
1	D	2177	LEU	CB-CG	-7.81	1.29	1.52
1	A	2177	LEU	CB-CG	-7.80	1.29	1.52
1	C	2177	LEU	CB-CG	-7.79	1.29	1.52
2	H	80	TYR	CA-CB	-7.67	1.37	1.53
2	G	80	TYR	CA-CB	-7.64	1.37	1.53
2	F	80	TYR	CA-CB	-7.63	1.37	1.53
2	E	80	TYR	CA-CB	-7.63	1.37	1.53
1	B	1559	GLN	CB-CG	-7.55	1.32	1.52
1	D	1559	GLN	CB-CG	-7.55	1.32	1.52
1	C	1559	GLN	CB-CG	-7.55	1.32	1.52
1	A	1559	GLN	CB-CG	-7.55	1.32	1.52
1	C	3478	MET	CB-CG	-7.42	1.27	1.51
1	A	3478	MET	CB-CG	-7.39	1.27	1.51
1	D	3478	MET	CB-CG	-7.39	1.27	1.51
1	B	3478	MET	CB-CG	-7.38	1.27	1.51
1	C	1944	GLU	CD-OE2	7.35	1.33	1.25
1	B	2734	ASN	CB-CG	-7.32	1.34	1.51
1	A	2734	ASN	CB-CG	-7.30	1.34	1.51
1	C	2734	ASN	CB-CG	-7.30	1.34	1.51
1	A	1944	GLU	CD-OE2	7.29	1.33	1.25
1	D	1944	GLU	CD-OE2	7.26	1.33	1.25
1	D	2734	ASN	CB-CG	-7.25	1.34	1.51
1	B	1944	GLU	CD-OE2	7.21	1.33	1.25
1	D	2722	LYS	CE-NZ	-6.99	1.31	1.49
1	A	2722	LYS	CE-NZ	-6.97	1.31	1.49
1	B	2722	LYS	CE-NZ	-6.97	1.31	1.49
1	A	1260	MET	CG-SD	-6.96	1.63	1.81
1	B	1260	MET	CG-SD	-6.96	1.63	1.81
1	D	1260	MET	CG-SD	-6.96	1.63	1.81
1	C	1260	MET	CG-SD	-6.96	1.63	1.81
1	C	2722	LYS	CE-NZ	-6.95	1.31	1.49
1	B	3467	MET	CB-CG	6.72	1.72	1.51
1	D	3467	MET	CB-CG	6.72	1.72	1.51
1	C	3467	MET	CB-CG	6.72	1.72	1.51
1	A	3467	MET	CB-CG	6.72	1.72	1.51
1	A	2874	MET	CB-CG	-6.70	1.29	1.51
1	B	2874	MET	CB-CG	-6.70	1.29	1.51

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	D	2874	MET	CB-CG	-6.69	1.29	1.51
1	C	2874	MET	CB-CG	-6.69	1.29	1.51
1	B	1128	ARG	CB-CG	-6.67	1.34	1.52
1	A	1128	ARG	CB-CG	-6.67	1.34	1.52
1	D	1128	ARG	CB-CG	-6.63	1.34	1.52
1	C	1128	ARG	CB-CG	-6.63	1.34	1.52
1	B	416	LYS	CE-NZ	6.49	1.65	1.49
1	D	416	LYS	CE-NZ	6.49	1.65	1.49
1	B	820	ARG	CA-CB	6.49	1.68	1.53
1	C	416	LYS	CE-NZ	6.48	1.65	1.49
1	A	416	LYS	CE-NZ	6.48	1.65	1.49
1	B	2312	MET	CB-CG	-6.48	1.30	1.51
1	D	820	ARG	CA-CB	6.48	1.68	1.53
1	C	820	ARG	CA-CB	6.48	1.68	1.53
1	A	820	ARG	CA-CB	6.47	1.68	1.53
1	A	2312	MET	CB-CG	-6.46	1.30	1.51
1	D	2312	MET	CB-CG	-6.45	1.30	1.51
1	C	2312	MET	CB-CG	-6.45	1.30	1.51
1	D	820	ARG	CG-CD	-6.35	1.36	1.51
1	A	820	ARG	CG-CD	-6.31	1.36	1.51
1	B	820	ARG	CG-CD	-6.29	1.36	1.51
1	B	820	ARG	CD-NE	6.29	1.57	1.46
1	C	820	ARG	CG-CD	-6.29	1.36	1.51
1	A	820	ARG	CD-NE	6.26	1.57	1.46
1	D	820	ARG	CD-NE	6.26	1.57	1.46
1	C	820	ARG	CD-NE	6.24	1.57	1.46
1	A	1559	GLN	CD-OE1	-6.23	1.10	1.24
1	B	1559	GLN	CD-OE1	-6.23	1.10	1.24
1	D	1559	GLN	CD-OE1	-6.23	1.10	1.24
1	C	1559	GLN	CD-OE1	-6.23	1.10	1.24
1	C	2444	GLN	CD-OE1	-5.93	1.10	1.24
1	A	2444	GLN	CD-OE1	-5.92	1.10	1.24
1	B	2444	GLN	CD-OE1	-5.92	1.10	1.24
1	D	2444	GLN	CD-OE1	-5.91	1.10	1.24
1	D	961	MET	CG-SD	-5.84	1.66	1.81
1	A	961	MET	CG-SD	-5.83	1.66	1.81
1	B	961	MET	CG-SD	-5.83	1.66	1.81
1	C	961	MET	CG-SD	-5.83	1.66	1.81
2	G	73	LYS	CE-NZ	-5.72	1.34	1.49
2	F	73	LYS	CE-NZ	-5.72	1.34	1.49
2	E	73	LYS	CE-NZ	-5.71	1.34	1.49
1	B	4161	ARG	CZ-NH1	-5.70	1.25	1.33

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	D	4161	ARG	CZ-NH1	-5.70	1.25	1.33
2	H	73	LYS	CE-NZ	-5.69	1.34	1.49
1	C	4161	ARG	CZ-NH1	-5.68	1.25	1.33
1	D	2267	MET	CA-CB	-5.68	1.41	1.53
1	C	2267	MET	CA-CB	-5.68	1.41	1.53
1	A	2267	MET	CA-CB	-5.67	1.41	1.53
1	B	2267	MET	CA-CB	-5.67	1.41	1.53
1	A	4161	ARG	CZ-NH1	-5.67	1.25	1.33
1	D	820	ARG	CB-CG	5.66	1.67	1.52
1	D	4161	ARG	NE-CZ	-5.66	1.25	1.33
1	A	820	ARG	CB-CG	5.65	1.67	1.52
1	A	4161	ARG	NE-CZ	-5.65	1.25	1.33
1	B	820	ARG	CB-CG	5.65	1.67	1.52
1	C	820	ARG	CB-CG	5.65	1.67	1.52
1	B	4161	ARG	NE-CZ	-5.61	1.25	1.33
1	C	4161	ARG	NE-CZ	-5.61	1.25	1.33
1	A	4796	MET	CG-SD	-5.60	1.66	1.81
1	B	4796	MET	CG-SD	-5.59	1.66	1.81
1	D	4796	MET	CG-SD	-5.59	1.66	1.81
1	C	4796	MET	CG-SD	-5.58	1.66	1.81
2	G	80	TYR	CG-CD2	-5.57	1.31	1.39
2	F	80	TYR	CG-CD2	-5.56	1.31	1.39
2	E	80	TYR	CG-CD2	-5.54	1.31	1.39
1	A	2267	MET	CG-SD	-5.52	1.66	1.81
1	D	2267	MET	CG-SD	-5.52	1.66	1.81
1	B	2267	MET	CG-SD	-5.51	1.66	1.81
1	C	2267	MET	CG-SD	-5.51	1.66	1.81
2	H	80	TYR	CG-CD2	-5.50	1.31	1.39
1	B	1036	ARG	C-N	-5.18	1.22	1.34
1	D	1036	ARG	C-N	-5.18	1.22	1.34
1	C	1036	ARG	C-N	-5.18	1.22	1.34
1	A	1036	ARG	C-N	-5.17	1.22	1.34
2	G	52	LYS	CD-CE	-5.14	1.38	1.51
1	D	1944	GLU	CB-CG	5.14	1.61	1.52
2	E	52	LYS	CD-CE	-5.14	1.38	1.51
1	B	1944	GLU	CB-CG	5.12	1.61	1.52
2	H	52	LYS	CD-CE	-5.11	1.38	1.51
2	F	52	LYS	CD-CE	-5.11	1.38	1.51
1	A	1944	GLU	CB-CG	5.10	1.61	1.52
1	B	2694	GLU	CG-CD	-5.09	1.44	1.51
1	C	1944	GLU	CB-CG	5.08	1.61	1.52
1	A	2694	GLU	CG-CD	-5.06	1.44	1.51

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	D	2694	GLU	CG-CD	-5.04	1.44	1.51
1	C	2694	GLU	CG-CD	-5.04	1.44	1.51
1	A	2722	LYS	CD-CE	5.01	1.63	1.51
1	B	2722	LYS	CD-CE	5.01	1.63	1.51
1	D	2722	LYS	CD-CE	5.01	1.63	1.51

All (356) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	H	80	TYR	CE1-CZ-CE2	-55.59	30.86	119.80
2	F	80	TYR	CE1-CZ-CE2	-55.57	30.88	119.80
2	G	80	TYR	CE1-CZ-CE2	-55.57	30.89	119.80
2	E	80	TYR	CE1-CZ-CE2	-55.56	30.90	119.80
2	F	80	TYR	CB-CG-CD2	45.56	148.33	121.00
2	G	80	TYR	CB-CG-CD2	45.46	148.28	121.00
2	H	80	TYR	CB-CG-CD2	45.45	148.27	121.00
2	E	80	TYR	CB-CG-CD2	45.40	148.24	121.00
2	H	80	TYR	CD1-CG-CD2	-44.58	68.86	117.90
2	E	80	TYR	CD1-CG-CD2	-44.55	68.89	117.90
2	G	80	TYR	CD1-CG-CD2	-44.55	68.89	117.90
2	F	80	TYR	CD1-CG-CD2	-44.54	68.91	117.90
2	F	80	TYR	CG-CD1-CE1	-39.52	89.69	121.30
2	G	80	TYR	CG-CD1-CE1	-39.50	89.70	121.30
2	E	80	TYR	CG-CD1-CE1	-39.49	89.71	121.30
2	H	80	TYR	CG-CD1-CE1	-39.46	89.73	121.30
2	E	80	TYR	CB-CG-CD1	-24.89	106.07	121.00
2	F	80	TYR	CB-CG-CD1	-24.84	106.09	121.00
2	G	80	TYR	CB-CG-CD1	-24.84	106.10	121.00
2	H	80	TYR	CB-CG-CD1	-24.83	106.10	121.00
1	B	2642	LYS	CD-CE-NZ	-23.04	58.72	111.70
1	D	2642	LYS	CD-CE-NZ	-23.03	58.73	111.70
1	A	2642	LYS	CD-CE-NZ	-23.02	58.75	111.70
1	C	2642	LYS	CD-CE-NZ	-23.02	58.76	111.70
1	A	3239	MET	CB-CG-SD	22.53	179.99	112.40
1	B	3239	MET	CB-CG-SD	22.53	179.99	112.40
1	C	3239	MET	CB-CG-SD	22.52	179.97	112.40
1	D	3239	MET	CB-CG-SD	22.52	179.97	112.40
2	G	80	TYR	CZ-CE2-CD2	-21.84	100.14	119.80
2	F	80	TYR	CZ-CE2-CD2	-21.82	100.17	119.80
2	E	80	TYR	CZ-CE2-CD2	-21.81	100.17	119.80
2	H	80	TYR	CZ-CE2-CD2	-21.79	100.19	119.80
1	B	384	MET	CG-SD-CE	-18.39	70.77	100.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	384	MET	CG-SD-CE	-18.38	70.79	100.20
1	C	384	MET	CG-SD-CE	-18.38	70.79	100.20
1	D	384	MET	CG-SD-CE	-18.37	70.81	100.20
1	D	384	MET	CB-CG-SD	18.25	167.14	112.40
1	A	384	MET	CB-CG-SD	18.23	167.10	112.40
1	C	384	MET	CB-CG-SD	18.22	167.07	112.40
1	B	384	MET	CB-CG-SD	18.22	167.06	112.40
1	C	2312	MET	CG-SD-CE	17.91	128.85	100.20
1	A	2312	MET	CG-SD-CE	17.89	128.83	100.20
1	B	2312	MET	CG-SD-CE	17.89	128.83	100.20
1	D	2312	MET	CG-SD-CE	17.89	128.82	100.20
1	D	3239	MET	CA-CB-CG	-17.62	83.35	113.30
1	A	3239	MET	CA-CB-CG	-17.61	83.37	113.30
1	B	3239	MET	CA-CB-CG	-17.59	83.39	113.30
1	C	3239	MET	CA-CB-CG	-17.59	83.39	113.30
1	B	384	MET	CA-CB-CG	17.09	142.36	113.30
1	D	384	MET	CA-CB-CG	17.08	142.34	113.30
1	A	384	MET	CA-CB-CG	17.07	142.33	113.30
1	C	384	MET	CA-CB-CG	17.05	142.29	113.30
1	C	1036	ARG	CG-CD-NE	16.97	147.43	111.80
1	A	1036	ARG	CG-CD-NE	16.96	147.41	111.80
1	B	1036	ARG	CG-CD-NE	16.95	147.39	111.80
1	D	1036	ARG	CG-CD-NE	16.93	147.35	111.80
2	E	80	TYR	CD1-CE1-CZ	15.81	134.03	119.80
2	G	80	TYR	CD1-CE1-CZ	15.80	134.02	119.80
2	F	80	TYR	CD1-CE1-CZ	15.79	134.01	119.80
2	H	80	TYR	CD1-CE1-CZ	15.76	133.99	119.80
2	F	5	GLU	OE1-CD-OE2	-15.06	105.23	123.30
2	G	5	GLU	OE1-CD-OE2	-15.06	105.23	123.30
2	H	5	GLU	OE1-CD-OE2	-15.05	105.24	123.30
2	E	5	GLU	OE1-CD-OE2	-15.04	105.25	123.30
1	D	1036	ARG	CD-NE-CZ	-14.54	103.24	123.60
1	A	1036	ARG	CD-NE-CZ	-14.53	103.26	123.60
1	B	1036	ARG	CD-NE-CZ	-14.51	103.28	123.60
1	C	1036	ARG	CD-NE-CZ	-14.50	103.30	123.60
2	G	53	GLN	CG-CD-OE1	14.42	150.45	121.60
2	H	53	GLN	CG-CD-OE1	14.42	150.43	121.60
2	F	53	GLN	CG-CD-OE1	14.41	150.43	121.60
2	E	53	GLN	CG-CD-OE1	14.39	150.38	121.60
1	B	1128	ARG	NE-CZ-NH1	14.37	127.48	120.30
1	C	1128	ARG	NE-CZ-NH1	14.34	127.47	120.30
1	A	1128	ARG	NE-CZ-NH1	14.33	127.47	120.30

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	D	1128	ARG	NE-CZ-NH1	14.33	127.47	120.30
2	E	52	LYS	CD-CE-NZ	14.26	144.51	111.70
2	H	52	LYS	CD-CE-NZ	14.25	144.48	111.70
2	G	52	LYS	CD-CE-NZ	14.24	144.46	111.70
2	F	52	LYS	CD-CE-NZ	14.24	144.45	111.70
1	B	820	ARG	CD-NE-CZ	-13.52	104.67	123.60
1	A	820	ARG	CD-NE-CZ	-13.50	104.70	123.60
1	D	820	ARG	CD-NE-CZ	-13.49	104.71	123.60
1	C	820	ARG	CD-NE-CZ	-13.49	104.71	123.60
1	C	1128	ARG	NE-CZ-NH2	-12.52	114.04	120.30
1	A	1128	ARG	NE-CZ-NH2	-12.40	114.10	120.30
2	H	53	GLN	OE1-CD-NE2	-12.40	93.38	121.90
1	B	1128	ARG	NE-CZ-NH2	-12.40	114.10	120.30
2	E	53	GLN	OE1-CD-NE2	-12.39	93.40	121.90
2	G	53	GLN	OE1-CD-NE2	-12.38	93.42	121.90
2	F	53	GLN	OE1-CD-NE2	-12.38	93.42	121.90
1	D	1128	ARG	NE-CZ-NH2	-12.35	114.12	120.30
1	C	1036	ARG	NE-CZ-NH2	-11.96	114.32	120.30
1	B	1036	ARG	NE-CZ-NH2	-11.94	114.33	120.30
1	D	1036	ARG	NE-CZ-NH2	-11.94	114.33	120.30
1	A	1036	ARG	NE-CZ-NH2	-11.91	114.34	120.30
1	B	820	ARG	CG-CD-NE	-11.80	87.02	111.80
1	A	820	ARG	CG-CD-NE	-11.79	87.03	111.80
1	D	820	ARG	CG-CD-NE	-11.79	87.04	111.80
1	C	820	ARG	CG-CD-NE	-11.78	87.06	111.80
1	A	2267	MET	CG-SD-CE	-11.47	81.85	100.20
1	B	2267	MET	CG-SD-CE	-11.47	81.85	100.20
1	D	2267	MET	CG-SD-CE	-11.47	81.85	100.20
1	C	2267	MET	CG-SD-CE	-11.47	81.86	100.20
1	B	820	ARG	NE-CZ-NH1	10.88	125.74	120.30
1	C	820	ARG	NE-CZ-NH1	10.88	125.74	120.30
1	A	820	ARG	NE-CZ-NH1	10.79	125.69	120.30
1	D	820	ARG	NE-CZ-NH1	10.75	125.67	120.30
1	C	820	ARG	CA-CB-CG	-10.74	89.78	113.40
1	B	820	ARG	CA-CB-CG	-10.73	89.79	113.40
1	D	820	ARG	CA-CB-CG	-10.73	89.78	113.40
1	A	820	ARG	CA-CB-CG	-10.73	89.80	113.40
1	C	3557	LEU	CB-CG-CD1	10.58	128.98	111.00
1	B	3557	LEU	CB-CG-CD1	10.56	128.96	111.00
1	D	3557	LEU	CB-CG-CD1	10.56	128.96	111.00
1	A	3557	LEU	CB-CG-CD1	10.56	128.96	111.00
1	B	1559	GLN	CA-CB-CG	10.31	136.08	113.40

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	C	1559	GLN	CA-CB-CG	10.31	136.08	113.40
1	D	1559	GLN	CA-CB-CG	10.31	136.08	113.40
1	A	1559	GLN	CA-CB-CG	10.31	136.07	113.40
1	D	3478	MET	CB-CG-SD	9.77	141.70	112.40
1	C	3478	MET	CB-CG-SD	9.76	141.67	112.40
1	A	3478	MET	CB-CG-SD	9.75	141.66	112.40
1	B	3478	MET	CB-CG-SD	9.74	141.62	112.40
2	H	105	LYS	CD-CE-NZ	9.57	133.72	111.70
2	F	105	LYS	CD-CE-NZ	9.57	133.71	111.70
2	E	105	LYS	CD-CE-NZ	9.55	133.67	111.70
2	G	105	LYS	CD-CE-NZ	9.55	133.66	111.70
2	H	53	GLN	CG-CD-NE2	-9.47	93.98	116.70
2	G	53	GLN	CG-CD-NE2	-9.46	93.99	116.70
2	F	53	GLN	CG-CD-NE2	-9.46	94.00	116.70
2	E	53	GLN	CG-CD-NE2	-9.46	94.00	116.70
1	B	2017	ASP	CB-CG-OD2	9.45	126.81	118.30
1	A	2017	ASP	CB-CG-OD2	9.42	126.78	118.30
1	D	2017	ASP	CB-CG-OD2	9.39	126.75	118.30
1	D	1036	ARG	NE-CZ-NH1	-9.37	115.62	120.30
1	C	2017	ASP	CB-CG-OD2	9.37	126.73	118.30
1	B	1036	ARG	NE-CZ-NH1	-9.31	115.64	120.30
1	A	1036	ARG	NE-CZ-NH1	-9.31	115.64	120.30
1	C	1036	ARG	NE-CZ-NH1	-9.31	115.64	120.30
1	B	4796	MET	CG-SD-CE	-9.28	85.36	100.20
1	D	4796	MET	CG-SD-CE	-9.28	85.36	100.20
1	C	4796	MET	CG-SD-CE	-9.27	85.36	100.20
1	A	4796	MET	CG-SD-CE	-9.27	85.37	100.20
1	D	3014	CYS	CA-CB-SG	9.17	130.51	114.00
1	A	3014	CYS	CA-CB-SG	9.16	130.49	114.00
1	B	3014	CYS	CA-CB-SG	9.16	130.48	114.00
1	C	3014	CYS	CA-CB-SG	9.14	130.45	114.00
1	D	3731	LYS	CD-CE-NZ	8.83	132.01	111.70
1	A	3731	LYS	CD-CE-NZ	8.82	131.99	111.70
1	D	4161	ARG	NE-CZ-NH1	-8.82	115.89	120.30
1	C	3731	LYS	CD-CE-NZ	8.82	131.98	111.70
1	B	3731	LYS	CD-CE-NZ	8.81	131.97	111.70
1	A	4161	ARG	NE-CZ-NH1	-8.77	115.92	120.30
1	B	4161	ARG	NE-CZ-NH1	-8.76	115.92	120.30
1	C	4161	ARG	NE-CZ-NH1	-8.73	115.93	120.30
1	D	3858	MET	CB-CG-SD	8.58	138.15	112.40
1	C	3858	MET	CB-CG-SD	8.58	138.13	112.40
1	A	3858	MET	CB-CG-SD	8.58	138.13	112.40

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	B	3858	MET	CB-CG-SD	8.57	138.12	112.40
1	D	2312	MET	CB-CG-SD	8.33	137.38	112.40
1	B	2312	MET	CB-CG-SD	8.32	137.37	112.40
1	A	2312	MET	CB-CG-SD	8.32	137.35	112.40
1	C	2312	MET	CB-CG-SD	8.31	137.32	112.40
1	C	2308	GLN	CA-CB-CG	8.26	131.56	113.40
1	D	2308	GLN	CA-CB-CG	8.26	131.56	113.40
1	A	2308	GLN	CA-CB-CG	8.25	131.55	113.40
1	B	2308	GLN	CA-CB-CG	8.23	131.51	113.40
1	A	2444	GLN	CA-CB-CG	8.01	131.02	113.40
1	C	2444	GLN	CA-CB-CG	8.01	131.01	113.40
1	B	2444	GLN	CA-CB-CG	8.00	131.01	113.40
1	D	2444	GLN	CA-CB-CG	8.00	131.01	113.40
1	D	3478	MET	CG-SD-CE	7.94	112.91	100.20
1	A	3478	MET	CG-SD-CE	7.92	112.88	100.20
1	B	3478	MET	CG-SD-CE	7.92	112.88	100.20
1	C	3478	MET	CG-SD-CE	7.92	112.88	100.20
2	G	80	TYR	N-CA-CB	-7.86	96.45	110.60
2	F	80	TYR	N-CA-CB	-7.86	96.45	110.60
2	H	80	TYR	N-CA-CB	-7.85	96.47	110.60
2	E	80	TYR	N-CA-CB	-7.85	96.47	110.60
1	D	3239	MET	CB-CA-C	-7.82	94.75	110.40
1	A	3239	MET	CB-CA-C	-7.82	94.77	110.40
1	C	2444	GLN	CG-CD-OE1	-7.82	105.97	121.60
1	B	3239	MET	CB-CA-C	-7.81	94.78	110.40
1	C	3239	MET	CB-CA-C	-7.81	94.78	110.40
2	F	53	GLN	CA-CB-CG	7.81	130.58	113.40
1	B	1260	MET	CG-SD-CE	-7.81	87.71	100.20
1	B	2444	GLN	CG-CD-OE1	-7.81	105.98	121.60
2	H	53	GLN	CA-CB-CG	7.80	130.57	113.40
1	A	1260	MET	CG-SD-CE	-7.80	87.72	100.20
1	D	1260	MET	CG-SD-CE	-7.79	87.73	100.20
1	C	1260	MET	CG-SD-CE	-7.79	87.73	100.20
1	A	2444	GLN	CG-CD-OE1	-7.79	106.01	121.60
2	E	53	GLN	CA-CB-CG	7.79	130.53	113.40
1	D	2444	GLN	CG-CD-OE1	-7.79	106.03	121.60
2	G	80	TYR	CG-CD2-CE2	-7.78	115.07	121.30
1	D	2208	MET	CG-SD-CE	-7.78	87.75	100.20
1	B	2208	MET	CG-SD-CE	-7.78	87.76	100.20
2	G	53	GLN	CA-CB-CG	7.77	130.50	113.40
1	A	2208	MET	CG-SD-CE	-7.77	87.77	100.20
1	C	2208	MET	CG-SD-CE	-7.75	87.80	100.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	E	80	TYR	CG-CD2-CE2	-7.75	115.10	121.30
2	H	80	TYR	CG-CD2-CE2	-7.75	115.10	121.30
1	A	2177	LEU	CB-CG-CD2	-7.68	97.94	111.00
1	C	2177	LEU	CB-CG-CD2	-7.68	97.95	111.00
2	F	80	TYR	CG-CD2-CE2	-7.67	115.16	121.30
1	B	2177	LEU	CB-CG-CD2	-7.67	97.96	111.00
1	D	2177	LEU	CB-CG-CD2	-7.66	97.98	111.00
1	C	3239	MET	CA-C-O	7.60	136.06	120.10
1	A	3239	MET	CA-C-O	7.60	136.05	120.10
1	B	3239	MET	CA-C-O	7.59	136.04	120.10
1	D	3239	MET	CA-C-O	7.58	136.03	120.10
1	D	3467	MET	CB-CG-SD	7.55	135.04	112.40
1	B	3467	MET	CB-CG-SD	7.54	135.04	112.40
1	A	3467	MET	CB-CG-SD	7.54	135.03	112.40
1	C	1944	GLU	OE1-CD-OE2	-7.54	114.25	123.30
1	A	1944	GLU	OE1-CD-OE2	-7.53	114.26	123.30
1	C	3467	MET	CB-CG-SD	7.53	134.99	112.40
1	B	1944	GLU	OE1-CD-OE2	-7.52	114.27	123.30
1	D	1944	GLU	OE1-CD-OE2	-7.51	114.28	123.30
1	D	3478	MET	CA-CB-CG	-7.42	100.69	113.30
1	A	3478	MET	CA-CB-CG	-7.41	100.70	113.30
1	B	3478	MET	CA-CB-CG	-7.39	100.73	113.30
1	C	3478	MET	CA-CB-CG	-7.38	100.75	113.30
1	A	4880	MET	CA-CB-CG	-7.37	100.77	113.30
1	B	4880	MET	CA-CB-CG	-7.36	100.78	113.30
1	C	4880	MET	CA-CB-CG	-7.36	100.78	113.30
1	D	4880	MET	CA-CB-CG	-7.35	100.80	113.30
1	C	2444	GLN	CB-CG-CD	7.35	130.70	111.60
1	B	2444	GLN	CB-CG-CD	7.34	130.69	111.60
1	D	2444	GLN	CB-CG-CD	7.34	130.69	111.60
1	A	2444	GLN	CB-CG-CD	7.33	130.67	111.60
1	D	1071	ARG	CG-CD-NE	7.11	126.72	111.80
1	C	1071	ARG	CG-CD-NE	7.10	126.71	111.80
1	A	1071	ARG	CG-CD-NE	7.10	126.71	111.80
1	B	1071	ARG	CG-CD-NE	7.09	126.69	111.80
1	D	4880	MET	CB-CG-SD	7.06	133.58	112.40
2	H	80	TYR	O-C-N	7.05	133.98	122.70
1	A	4880	MET	CB-CG-SD	7.05	133.54	112.40
1	B	4880	MET	CB-CG-SD	7.05	133.54	112.40
1	C	4880	MET	CB-CG-SD	7.04	133.53	112.40
2	G	80	TYR	O-C-N	7.02	133.93	122.70
2	E	80	TYR	O-C-N	7.02	133.93	122.70

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	F	80	TYR	O-C-N	7.02	133.93	122.70
1	C	4880	MET	CG-SD-CE	7.01	111.42	100.20
1	A	4880	MET	CG-SD-CE	7.00	111.40	100.20
1	B	4880	MET	CG-SD-CE	6.98	111.36	100.20
1	D	4880	MET	CG-SD-CE	6.97	111.36	100.20
1	D	3856	LEU	CB-CG-CD1	6.97	122.85	111.00
1	A	3856	LEU	CB-CG-CD1	6.97	122.84	111.00
1	C	3856	LEU	CB-CG-CD1	6.96	122.84	111.00
1	B	3856	LEU	CB-CG-CD1	6.94	122.80	111.00
1	B	2829	GLY	C-N-CA	6.89	138.92	121.70
1	A	2829	GLY	C-N-CA	6.88	138.91	121.70
1	D	2829	GLY	C-N-CA	6.87	138.88	121.70
1	C	2829	GLY	C-N-CA	6.87	138.88	121.70
2	H	60	GLU	OE1-CD-OE2	-6.83	115.11	123.30
2	F	60	GLU	OE1-CD-OE2	-6.81	115.12	123.30
2	E	60	GLU	OE1-CD-OE2	-6.81	115.13	123.30
2	G	60	GLU	OE1-CD-OE2	-6.79	115.15	123.30
2	G	80	TYR	CB-CA-C	6.77	123.94	110.40
2	E	80	TYR	CB-CA-C	6.77	123.93	110.40
2	H	80	TYR	CB-CA-C	6.76	123.93	110.40
2	F	80	TYR	CB-CA-C	6.75	123.91	110.40
1	D	2734	ASN	CB-CG-OD1	-6.73	108.14	121.60
1	A	2734	ASN	CB-CG-OD1	-6.71	108.17	121.60
1	B	2734	ASN	CB-CG-OD1	-6.71	108.18	121.60
1	C	2734	ASN	CB-CG-OD1	-6.71	108.19	121.60
1	D	1036	ARG	CB-CG-CD	6.59	128.74	111.60
1	A	1036	ARG	CB-CG-CD	6.58	128.72	111.60
1	B	1036	ARG	CB-CG-CD	6.57	128.69	111.60
1	C	1036	ARG	CB-CG-CD	6.57	128.67	111.60
1	B	1128	ARG	NH1-CZ-NH2	-6.23	112.55	119.40
1	A	1128	ARG	NH1-CZ-NH2	-6.22	112.56	119.40
1	D	1128	ARG	NH1-CZ-NH2	-6.21	112.56	119.40
1	C	1128	ARG	NH1-CZ-NH2	-6.19	112.60	119.40
2	F	80	TYR	OH-CZ-CE2	-6.07	103.71	120.10
2	G	80	TYR	OH-CZ-CE2	-6.05	103.77	120.10
2	E	80	TYR	OH-CZ-CE2	-6.04	103.78	120.10
2	H	80	TYR	OH-CZ-CE2	-6.03	103.81	120.10
1	C	2722	LYS	CD-CE-NZ	6.00	125.51	111.70
1	D	2722	LYS	CD-CE-NZ	6.00	125.49	111.70
1	A	2722	LYS	CD-CE-NZ	5.99	125.48	111.70
1	B	2722	LYS	CD-CE-NZ	5.98	125.45	111.70
1	D	484	LEU	CB-CG-CD2	5.91	121.05	111.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	484	LEU	CB-CG-CD2	5.91	121.04	111.00
1	B	484	LEU	CB-CG-CD2	5.91	121.04	111.00
1	C	484	LEU	CB-CG-CD2	5.88	121.00	111.00
1	D	4980	LEU	CB-CG-CD2	5.78	120.83	111.00
1	A	1559	GLN	CG-CD-OE1	5.77	133.14	121.60
1	B	1559	GLN	CG-CD-OE1	5.77	133.14	121.60
1	C	1559	GLN	CG-CD-OE1	5.77	133.14	121.60
1	D	1559	GLN	CG-CD-OE1	5.76	133.12	121.60
1	A	4980	LEU	CB-CG-CD2	5.76	120.79	111.00
1	C	4980	LEU	CB-CG-CD2	5.74	120.76	111.00
1	B	4980	LEU	CB-CG-CD2	5.74	120.75	111.00
1	D	3239	MET	CA-C-N	-5.62	104.83	117.20
1	A	3239	MET	CA-C-N	-5.61	104.85	117.20
1	B	3239	MET	CA-C-N	-5.60	104.88	117.20
1	C	3239	MET	CA-C-N	-5.60	104.88	117.20
1	B	2722	LYS	CG-CD-CE	-5.49	95.42	111.90
1	A	2722	LYS	CG-CD-CE	-5.48	95.46	111.90
1	C	2722	LYS	CG-CD-CE	-5.48	95.47	111.90
1	D	2722	LYS	CG-CD-CE	-5.47	95.48	111.90
1	A	10	GLU	N-CA-C	5.46	125.74	111.00
1	B	10	GLU	N-CA-C	5.46	125.74	111.00
1	A	2237	CYS	CB-CA-C	5.45	121.31	110.40
1	C	10	GLU	N-CA-C	5.45	125.72	111.00
1	B	2237	CYS	CB-CA-C	5.45	121.31	110.40
1	C	2237	CYS	CB-CA-C	5.45	121.30	110.40
1	D	10	GLU	N-CA-C	5.45	125.71	111.00
1	D	2237	CYS	CB-CA-C	5.45	121.29	110.40
2	H	52	LYS	CG-CD-CE	-5.41	95.68	111.90
2	G	52	LYS	CG-CD-CE	-5.40	95.70	111.90
2	F	52	LYS	CG-CD-CE	-5.39	95.72	111.90
2	E	52	LYS	CG-CD-CE	-5.39	95.74	111.90
1	B	2208	MET	CB-CG-SD	5.36	128.47	112.40
1	D	2208	MET	CB-CG-SD	5.35	128.46	112.40
1	A	1758	ARG	NE-CZ-NH2	-5.35	117.63	120.30
1	C	2208	MET	CB-CG-SD	5.35	128.44	112.40
1	A	2208	MET	CB-CG-SD	5.34	128.43	112.40
1	C	1758	ARG	NE-CZ-NH2	-5.34	117.63	120.30
1	D	1758	ARG	NE-CZ-NH2	-5.29	117.65	120.30
1	B	2932	MET	CG-SD-CE	5.29	108.66	100.20
1	D	2932	MET	CG-SD-CE	5.29	108.66	100.20
1	A	2932	MET	CG-SD-CE	5.29	108.66	100.20
1	B	1260	MET	CB-CG-SD	5.28	128.25	112.40

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	C	2932	MET	CG-SD-CE	5.28	108.64	100.20
1	D	1260	MET	CB-CG-SD	5.27	128.22	112.40
1	C	1260	MET	CB-CG-SD	5.27	128.22	112.40
1	A	1260	MET	CB-CG-SD	5.27	128.22	112.40
1	B	1758	ARG	NE-CZ-NH2	-5.26	117.67	120.30
1	B	2017	ASP	OD1-CG-OD2	-5.25	113.33	123.30
1	B	820	ARG	NH1-CZ-NH2	-5.24	113.64	119.40
2	E	5	GLU	CG-CD-OE1	5.22	128.73	118.30
1	A	2017	ASP	OD1-CG-OD2	-5.21	113.40	123.30
2	F	5	GLU	CG-CD-OE1	5.21	128.72	118.30
2	H	5	GLU	CG-CD-OE1	5.21	128.72	118.30
1	D	2017	ASP	OD1-CG-OD2	-5.20	113.42	123.30
1	A	820	ARG	NH1-CZ-NH2	-5.20	113.68	119.40
1	C	820	ARG	NH1-CZ-NH2	-5.20	113.68	119.40
1	D	820	ARG	NH1-CZ-NH2	-5.19	113.69	119.40
1	C	2017	ASP	OD1-CG-OD2	-5.18	113.45	123.30
2	G	5	GLU	CG-CD-OE1	5.18	128.66	118.30
2	H	80	TYR	CA-C-O	-5.16	109.26	120.10
2	E	80	TYR	CA-C-O	-5.16	109.27	120.10
2	G	80	TYR	CA-C-O	-5.16	109.27	120.10
2	F	80	TYR	CA-C-O	-5.15	109.28	120.10
1	B	3868	ARG	CG-CD-NE	-5.11	101.08	111.80
1	C	3868	ARG	CG-CD-NE	-5.10	101.08	111.80
1	A	3868	ARG	CG-CD-NE	-5.10	101.09	111.80
1	D	3868	ARG	CG-CD-NE	-5.09	101.11	111.80
2	E	13	ARG	NE-CZ-NH2	-5.05	117.77	120.30
2	G	13	ARG	NE-CZ-NH2	-5.04	117.78	120.30
2	H	13	ARG	NE-CZ-NH2	-5.02	117.79	120.30
2	F	13	ARG	NE-CZ-NH2	-5.01	117.80	120.30

There are no chirality outliers.

All (76) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	1035	ASN	Peptide
1	A	1036	ARG	Mainchain,Sidechain
1	A	1051	TYR	Sidechain
1	A	1128	ARG	Sidechain
1	A	1749	PRO	Mainchain
1	A	1758	ARG	Sidechain
1	A	1944	GLU	Sidechain
1	A	1982	ARG	Sidechain

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
1	A	2444	GLN	Sidechain
1	A	2701	PRO	Peptide
1	A	3453	ARG	Sidechain
1	A	4161	ARG	Sidechain
1	A	820	ARG	Sidechain
1	B	1035	ASN	Peptide
1	B	1036	ARG	Mainchain,Sidechain
1	B	1051	TYR	Sidechain
1	B	1128	ARG	Sidechain
1	B	1749	PRO	Mainchain
1	B	1758	ARG	Sidechain
1	B	1944	GLU	Sidechain
1	B	1982	ARG	Sidechain
1	B	2444	GLN	Sidechain
1	B	2701	PRO	Peptide
1	B	3453	ARG	Sidechain
1	B	4161	ARG	Sidechain
1	B	820	ARG	Sidechain
1	C	1035	ASN	Peptide
1	C	1036	ARG	Mainchain,Sidechain
1	C	1051	TYR	Sidechain
1	C	1128	ARG	Sidechain
1	C	1749	PRO	Mainchain
1	C	1758	ARG	Sidechain
1	C	1944	GLU	Sidechain
1	C	1982	ARG	Sidechain
1	C	2444	GLN	Sidechain
1	C	2701	PRO	Peptide
1	C	3453	ARG	Sidechain
1	C	4161	ARG	Sidechain
1	C	820	ARG	Sidechain
1	D	1035	ASN	Peptide
1	D	1036	ARG	Mainchain,Sidechain
1	D	1051	TYR	Sidechain
1	D	1128	ARG	Sidechain
1	D	1749	PRO	Mainchain
1	D	1758	ARG	Sidechain
1	D	1944	GLU	Sidechain
1	D	1982	ARG	Sidechain
1	D	2444	GLN	Sidechain
1	D	2701	PRO	Peptide
1	D	3453	ARG	Sidechain

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Group
1	D	4161	ARG	Sidechain
1	D	820	ARG	Sidechain
2	E	5	GLU	Sidechain
2	E	52	LYS	Mainchain
2	E	53	GLN	Sidechain
2	E	60	GLU	Sidechain
2	E	80	TYR	Sidechain
2	F	5	GLU	Sidechain
2	F	52	LYS	Mainchain
2	F	53	GLN	Sidechain
2	F	60	GLU	Sidechain
2	F	80	TYR	Sidechain
2	G	5	GLU	Sidechain
2	G	52	LYS	Mainchain
2	G	53	GLN	Sidechain
2	G	60	GLU	Sidechain
2	G	80	TYR	Sidechain
2	H	5	GLU	Sidechain
2	H	52	LYS	Mainchain
2	H	53	GLN	Sidechain
2	H	60	GLU	Sidechain
2	H	80	TYR	Sidechain

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	35150	0	34792	993	0
1	B	35150	0	34792	985	0
1	C	35150	0	34792	973	0
1	D	35150	0	34792	984	0
2	E	831	0	831	23	0
2	F	831	0	831	23	0
2	G	831	0	831	23	0
2	H	831	0	831	23	0
3	A	31	0	12	0	0
3	B	31	0	12	0	0
3	C	31	0	12	0	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
3	D	31	0	12	0	0
4	A	1	0	0	0	0
4	B	1	0	0	0	0
4	C	1	0	0	0	0
4	D	1	0	0	0	0
5	A	1	0	0	0	0
5	B	1	0	0	0	0
5	C	1	0	0	0	0
5	D	1	0	0	0	0
6	A	20	0	18	3	0
6	B	20	0	18	3	0
6	C	20	0	18	3	0
6	D	20	0	18	3	0
7	A	2	0	0	0	0
7	B	2	0	0	0	0
7	C	2	0	0	0	0
7	D	2	0	0	0	0
All	All	144144	0	142612	3958	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 14.

All (3958) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3239:MET:CG	1:D:3239:MET:SD	2.09	1.41
1:A:3239:MET:CG	1:A:3239:MET:SD	2.09	1.41
1:B:3239:MET:SD	1:B:3239:MET:CG	2.09	1.39
1:C:3239:MET:SD	1:C:3239:MET:CG	2.09	1.39
1:A:2642:LYS:CD	1:A:2642:LYS:CG	2.05	1.34
1:B:2642:LYS:CG	1:B:2642:LYS:CD	2.05	1.34
1:D:2642:LYS:CG	1:D:2642:LYS:CD	2.05	1.33
1:A:1749:PRO:O	1:A:1758:ARG:NH2	1.62	1.32
1:C:2642:LYS:CG	1:C:2642:LYS:CD	2.05	1.32
1:C:1749:PRO:O	1:C:1758:ARG:NH2	1.62	1.31
1:D:3239:MET:CG	1:D:3239:MET:CB	2.10	1.30
1:A:3239:MET:CG	1:A:3239:MET:CB	2.10	1.30
1:B:1749:PRO:O	1:B:1758:ARG:NH2	1.62	1.30
1:D:1749:PRO:O	1:D:1758:ARG:NH2	1.62	1.28
1:B:3239:MET:CG	1:B:3239:MET:CB	2.10	1.28
1:C:3239:MET:CG	1:C:3239:MET:CB	2.10	1.28

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:961:MET:CE	1:C:963:ASN:HB3	1.66	1.26
1:A:961:MET:CE	1:A:963:ASN:HB3	1.66	1.25
1:D:961:MET:CE	1:D:963:ASN:HB3	1.66	1.24
1:B:961:MET:CE	1:B:963:ASN:HB3	1.66	1.23
1:B:972:LEU:HG	1:B:1041:GLN:NE2	1.59	1.18
1:C:972:LEU:HG	1:C:1041:GLN:NE2	1.59	1.18
1:A:941:MET:CE	1:A:1051:TYR:CD1	2.27	1.17
1:C:941:MET:CE	1:C:1051:TYR:CD1	2.27	1.17
1:B:941:MET:CE	1:B:1051:TYR:CD1	2.27	1.17
1:C:961:MET:HE3	1:C:963:ASN:HB3	1.24	1.17
1:D:941:MET:CE	1:D:1051:TYR:CD1	2.27	1.16
1:D:972:LEU:HG	1:D:1041:GLN:NE2	1.59	1.16
1:A:972:LEU:HG	1:A:1041:GLN:NE2	1.59	1.16
1:B:2237:CYS:SG	1:B:2237:CYS:CB	2.35	1.15
1:A:2237:CYS:CB	1:A:2237:CYS:SG	2.35	1.15
1:C:941:MET:HE1	1:C:1051:TYR:CD1	1.81	1.14
1:C:2237:CYS:SG	1:C:2237:CYS:CB	2.35	1.14
1:D:2237:CYS:SG	1:D:2275:VAL:HG22	1.88	1.13
1:D:2237:CYS:SG	1:D:2237:CYS:CB	2.35	1.13
1:C:2237:CYS:SG	1:C:2275:VAL:HG22	1.88	1.13
1:A:2237:CYS:SG	1:A:2275:VAL:HG22	1.88	1.12
1:B:2237:CYS:SG	1:B:2275:VAL:HG22	1.88	1.12
1:B:3450:ASN:OD1	1:B:3453:ARG:NH2	1.82	1.12
1:D:3450:ASN:OD1	1:D:3453:ARG:NH2	1.82	1.12
1:A:3450:ASN:OD1	1:A:3453:ARG:NH2	1.82	1.12
1:C:3450:ASN:OD1	1:C:3453:ARG:NH2	1.82	1.12
1:B:961:MET:HE3	1:B:963:ASN:HB3	1.29	1.11
1:B:941:MET:HE1	1:B:1051:TYR:CD1	1.86	1.09
1:D:961:MET:HE3	1:D:963:ASN:HB3	1.30	1.07
1:A:941:MET:HE1	1:A:1051:TYR:CD1	1.89	1.07
1:B:2233:CYS:CB	1:B:2237:CYS:SG	2.43	1.07
1:A:2233:CYS:CB	1:A:2237:CYS:SG	2.43	1.06
1:C:2233:CYS:CB	1:C:2237:CYS:SG	2.43	1.06
1:D:2233:CYS:CB	1:D:2237:CYS:SG	2.43	1.05
1:D:2642:LYS:CG	1:D:2642:LYS:NZ	2.21	1.04
1:A:961:MET:HE3	1:A:963:ASN:HB3	1.36	1.03
1:A:2642:LYS:CG	1:A:2642:LYS:NZ	2.21	1.03
1:D:941:MET:HE1	1:D:1051:TYR:CD1	1.90	1.02
1:A:2874:MET:HE2	1:A:2939:ARG:HB2	1.38	1.02
1:D:941:MET:SD	1:D:1051:TYR:CE1	2.52	1.02
1:C:972:LEU:HG	1:C:1041:GLN:HE22	0.87	1.02

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:941:MET:SD	1:A:1051:TYR:CE1	2.52	1.02
1:B:941:MET:SD	1:B:1051:TYR:CE1	2.52	1.02
1:C:941:MET:SD	1:C:1051:TYR:CE1	2.52	1.02
1:C:2642:LYS:CG	1:C:2642:LYS:NZ	2.21	1.02
1:D:2929:PHE:HA	1:D:2932:MET:HE1	1.36	1.02
1:A:2929:PHE:HA	1:A:2932:MET:HE1	1.39	1.01
1:B:2642:LYS:CG	1:B:2642:LYS:NZ	2.21	1.01
1:D:972:LEU:HG	1:D:1041:GLN:HE22	0.87	1.01
1:B:2874:MET:HE2	1:B:2939:ARG:HB2	1.42	1.00
1:A:972:LEU:HG	1:A:1041:GLN:HE22	0.87	1.00
1:B:972:LEU:HG	1:B:1041:GLN:HE22	0.87	1.00
1:C:2929:PHE:HA	1:C:2932:MET:HE1	1.44	0.99
1:B:3219:TYR:HE2	1:B:3239:MET:SD	1.86	0.99
1:C:3219:TYR:HE2	1:C:3239:MET:SD	1.86	0.99
1:D:972:LEU:CG	1:D:1041:GLN:HE22	1.77	0.98
1:A:972:LEU:CG	1:A:1041:GLN:HE22	1.77	0.98
1:A:3219:TYR:HE2	1:A:3239:MET:SD	1.86	0.98
1:B:1808:ARG:NH2	1:B:1858:ASP:OD2	1.97	0.97
1:D:3219:TYR:HE2	1:D:3239:MET:SD	1.86	0.97
1:B:2929:PHE:HA	1:B:2932:MET:HE1	1.42	0.97
1:C:972:LEU:CG	1:C:1041:GLN:HE22	1.77	0.97
1:C:2929:PHE:HA	1:C:2932:MET:CE	1.95	0.97
1:D:1808:ARG:NH2	1:D:1858:ASP:OD2	1.97	0.96
1:B:2929:PHE:HA	1:B:2932:MET:CE	1.95	0.96
1:D:2929:PHE:HA	1:D:2932:MET:CE	1.95	0.96
1:A:1808:ARG:NH2	1:A:1858:ASP:OD2	1.97	0.96
1:A:2929:PHE:HA	1:A:2932:MET:CE	1.95	0.96
1:B:2233:CYS:HB3	1:B:2237:CYS:SG	2.05	0.96
1:C:1749:PRO:C	1:C:1758:ARG:HH22	1.69	0.96
1:D:2233:CYS:HB3	1:D:2237:CYS:SG	2.04	0.96
1:C:1808:ARG:NH2	1:C:1858:ASP:OD2	1.97	0.96
1:C:2233:CYS:HB3	1:C:2237:CYS:SG	2.04	0.96
1:A:2771:ILE:HG22	1:B:1506:GLN:OE1	1.66	0.96
1:D:1749:PRO:C	1:D:1758:ARG:HH22	1.69	0.96
1:B:1749:PRO:C	1:B:1758:ARG:HH22	1.69	0.95
1:A:2233:CYS:HB3	1:A:2237:CYS:SG	2.04	0.95
1:B:972:LEU:CG	1:B:1041:GLN:HE22	1.77	0.95
1:B:3521:GLY:HA2	1:B:3524:MET:CE	1.97	0.95
1:D:3521:GLY:HA2	1:D:3524:MET:CE	1.96	0.94
1:A:3521:GLY:HA2	1:A:3524:MET:CE	1.96	0.94
1:C:3521:GLY:HA2	1:C:3524:MET:CE	1.96	0.94

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3521:GLY:HA2	1:C:3524:MET:HE3	1.49	0.94
1:D:3521:GLY:HA2	1:D:3524:MET:HE3	1.49	0.94
1:A:1749:PRO:C	1:A:1758:ARG:HH22	1.69	0.93
1:D:2874:MET:CE	1:D:2939:ARG:HB2	1.99	0.93
1:B:2874:MET:CE	1:B:2939:ARG:HB2	1.99	0.93
1:C:2874:MET:CE	1:C:2939:ARG:HB2	1.99	0.93
1:D:2874:MET:HE2	1:D:2939:ARG:HB2	1.48	0.93
1:B:2523:ASP:HB3	1:B:2578:MET:HE1	1.51	0.92
1:B:3521:GLY:HA2	1:B:3524:MET:HE3	1.49	0.92
1:C:2874:MET:HE2	1:C:2939:ARG:HB2	1.49	0.92
1:A:221:ARG:NH2	1:A:397:GLU:OE2	2.03	0.92
1:A:2874:MET:CE	1:A:2939:ARG:HB2	1.99	0.92
2:E:52:LYS:O	2:E:53:GLN:HB2	1.68	0.92
1:B:2771:ILE:HG22	1:C:1506:GLN:OE1	1.69	0.92
1:D:1506:GLN:OE1	1:C:2771:ILE:HG22	1.70	0.92
1:D:221:ARG:NH2	1:D:397:GLU:OE2	2.03	0.91
1:B:2572:THR:HG21	1:B:2579:VAL:CG2	2.00	0.91
1:C:2572:THR:HG21	1:C:2579:VAL:CG2	2.00	0.91
1:A:221:ARG:NH1	1:A:258:SER:OG	2.04	0.91
2:F:52:LYS:O	2:F:53:GLN:HB2	1.68	0.91
1:B:221:ARG:NH1	1:B:258:SER:OG	2.04	0.91
1:C:221:ARG:NH1	1:C:258:SER:OG	2.04	0.91
1:A:495:ASN:OD1	1:A:553:ARG:NH1	2.03	0.91
1:B:221:ARG:NH2	1:B:397:GLU:OE2	2.03	0.91
1:B:495:ASN:OD1	1:B:553:ARG:NH1	2.03	0.91
1:D:2011:HIS:HE1	1:D:2017:ASP:OD2	1.54	0.91
1:C:495:ASN:OD1	1:C:553:ARG:NH1	2.03	0.91
1:D:221:ARG:NH1	1:D:258:SER:OG	2.04	0.90
1:A:2011:HIS:HE1	1:A:2017:ASP:OD2	1.54	0.90
2:G:52:LYS:O	2:G:53:GLN:HB2	1.68	0.90
1:D:2628:PHE:CZ	1:D:2734:ASN:OD1	2.25	0.90
1:C:221:ARG:NH2	1:C:397:GLU:OE2	2.03	0.90
1:A:2572:THR:HG21	1:A:2579:VAL:CG2	2.00	0.90
1:D:2572:THR:HG21	1:D:2579:VAL:CG2	2.00	0.90
1:A:2628:PHE:CZ	1:A:2734:ASN:OD1	2.25	0.90
1:B:2628:PHE:CZ	1:B:2734:ASN:OD1	2.24	0.90
2:H:52:LYS:O	2:H:53:GLN:HB2	1.68	0.90
1:B:2011:HIS:HE1	1:B:2017:ASP:OD2	1.54	0.90
1:A:2226:PRO:HB3	1:A:2267:MET:HE1	1.54	0.89
1:D:495:ASN:OD1	1:D:553:ARG:NH1	2.03	0.89
1:C:2628:PHE:CZ	1:C:2734:ASN:OD1	2.25	0.89

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2011:HIS:HE1	1:C:2017:ASP:OD2	1.54	0.89
1:D:4039:MET:SD	1:D:4042:ARG:NH2	2.46	0.89
1:B:3239:MET:CG	1:B:3239:MET:CA	2.51	0.89
1:B:1454:THR:OG1	1:B:1456:ASP:OD1	1.91	0.88
1:B:2292:GLU:OE1	1:B:2352:VAL:HG11	1.74	0.88
1:C:1454:THR:OG1	1:C:1456:ASP:OD1	1.91	0.88
1:C:3239:MET:CG	1:C:3239:MET:CA	2.51	0.88
1:A:952:LYS:NZ	1:A:968:ALA:O	2.06	0.88
1:A:2292:GLU:OE1	1:A:2352:VAL:HG11	1.73	0.88
1:D:3239:MET:CG	1:D:3239:MET:CA	2.51	0.88
1:A:961:MET:SD	1:A:963:ASN:HB3	2.14	0.88
1:A:2599:GLN:O	1:A:2603:ILE:HD12	1.74	0.88
1:A:4039:MET:SD	1:A:4042:ARG:NH2	2.46	0.88
1:B:952:LYS:NZ	1:B:968:ALA:O	2.06	0.88
1:B:4039:MET:SD	1:B:4042:ARG:NH2	2.46	0.88
1:A:3239:MET:CG	1:A:3239:MET:CA	2.51	0.87
1:D:4006:ASP:OD2	1:D:4008:SER:OG	1.91	0.87
1:D:952:LYS:NZ	1:D:968:ALA:O	2.06	0.87
1:C:952:LYS:NZ	1:C:968:ALA:O	2.06	0.87
1:C:2292:GLU:OE1	1:C:2352:VAL:HG11	1.74	0.87
1:C:961:MET:SD	1:C:963:ASN:HB3	2.14	0.87
1:D:961:MET:SD	1:D:963:ASN:HB3	2.14	0.87
1:C:4006:ASP:OD2	1:C:4008:SER:OG	1.91	0.87
1:C:4039:MET:SD	1:C:4042:ARG:NH2	2.46	0.87
1:A:1506:GLN:OE1	1:D:2771:ILE:HG22	1.73	0.87
1:D:3466:ASN:ND2	1:D:3507:THR:O	2.08	0.87
1:A:1454:THR:OG1	1:A:1456:ASP:OD1	1.91	0.87
1:A:2196:ASN:OD1	1:A:2199:ARG:NH2	2.08	0.87
1:B:961:MET:SD	1:B:963:ASN:HB3	2.14	0.87
1:D:2292:GLU:OE1	1:D:2352:VAL:HG11	1.74	0.87
1:B:2196:ASN:OD1	1:B:2199:ARG:NH2	2.08	0.86
1:C:2271:THR:OG1	1:C:2330:ARG:NH2	2.08	0.86
1:C:2599:GLN:O	1:C:2603:ILE:HD12	1.74	0.86
1:B:2599:GLN:O	1:B:2603:ILE:HD12	1.74	0.86
1:D:1454:THR:OG1	1:D:1456:ASP:OD1	1.91	0.86
1:C:3466:ASN:ND2	1:C:3507:THR:O	2.08	0.86
1:D:3573:MET:HB3	1:D:3577:ARG:NH1	1.91	0.86
1:C:2196:ASN:OD1	1:C:2199:ARG:NH2	2.08	0.86
1:D:2226:PRO:HB3	1:D:2267:MET:HE1	1.57	0.86
1:A:3573:MET:HB3	1:A:3577:ARG:NH1	1.91	0.86
1:A:4006:ASP:OD2	1:A:4008:SER:OG	1.91	0.86

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2271:THR:OG1	1:B:2330:ARG:NH2	2.08	0.86
1:B:4006:ASP:OD2	1:B:4008:SER:OG	1.91	0.86
1:D:2599:GLN:O	1:D:2603:ILE:HD12	1.74	0.86
1:B:3466:ASN:ND2	1:B:3507:THR:O	2.08	0.86
1:C:3573:MET:HB3	1:C:3577:ARG:NH1	1.91	0.86
1:B:3573:MET:HB3	1:B:3577:ARG:NH1	1.91	0.86
1:B:3943:ILE:O	1:B:3948:LYS:NZ	2.09	0.86
1:A:972:LEU:CG	1:A:1041:GLN:NE2	2.38	0.85
1:A:3466:ASN:ND2	1:A:3507:THR:O	2.08	0.85
1:B:941:MET:CE	1:B:1051:TYR:CE1	2.59	0.85
1:D:3943:ILE:O	1:D:3948:LYS:NZ	2.09	0.85
1:A:3521:GLY:HA2	1:A:3524:MET:HE2	1.58	0.85
1:C:3943:ILE:O	1:C:3948:LYS:NZ	2.09	0.85
1:A:941:MET:CE	1:A:1051:TYR:CE1	2.60	0.85
1:D:2196:ASN:OD1	1:D:2199:ARG:NH2	2.08	0.85
1:D:2271:THR:OG1	1:D:2330:ARG:NH2	2.08	0.85
1:A:2271:THR:OG1	1:A:2330:ARG:NH2	2.08	0.85
1:A:3943:ILE:O	1:A:3948:LYS:NZ	2.09	0.85
1:C:3573:MET:HB3	1:C:3577:ARG:HH12	1.42	0.85
1:B:70:GLU:OE2	1:B:110:ARG:NE	2.09	0.85
1:A:70:GLU:OE2	1:A:110:ARG:NE	2.09	0.84
1:D:70:GLU:OE2	1:D:110:ARG:NE	2.09	0.84
1:C:3219:TYR:CE2	1:C:3239:MET:SD	2.70	0.84
1:C:70:GLU:OE2	1:C:110:ARG:NE	2.09	0.84
1:D:3219:TYR:CE2	1:D:3239:MET:SD	2.70	0.84
1:D:3239:MET:CA	1:D:3239:MET:HG2	2.08	0.84
1:D:3573:MET:HB3	1:D:3577:ARG:HH12	1.42	0.84
1:A:3219:TYR:CE2	1:A:3239:MET:SD	2.70	0.84
1:A:2929:PHE:HA	1:A:2932:MET:SD	2.18	0.84
1:B:2929:PHE:HA	1:B:2932:MET:SD	2.18	0.84
1:D:941:MET:CE	1:D:1051:TYR:CE1	2.60	0.84
1:B:3219:TYR:CE2	1:B:3239:MET:SD	2.70	0.83
1:C:941:MET:CE	1:C:1051:TYR:CE1	2.60	0.83
1:A:3573:MET:HB3	1:A:3577:ARG:HH12	1.42	0.83
1:C:2929:PHE:HA	1:C:2932:MET:SD	2.18	0.83
1:B:1999:ARG:HH12	1:B:3636:PHE:HA	1.44	0.83
1:B:3573:MET:HB3	1:B:3577:ARG:HH12	1.42	0.83
1:D:1999:ARG:HH12	1:D:3636:PHE:HA	1.44	0.83
1:D:2929:PHE:HA	1:D:2932:MET:SD	2.18	0.83
1:C:1156:THR:OG1	1:C:1157:GLU:OE1	1.97	0.82
1:A:1999:ARG:HH12	1:A:3636:PHE:HA	1.44	0.82

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4749:GLU:O	1:A:4753:HIS:ND1	2.13	0.82
1:C:3282:PRO:O	1:C:3286:GLU:OE1	1.97	0.82
1:B:871:ARG:HD2	1:B:926:GLY:CA	2.09	0.82
1:C:1999:ARG:HH12	1:C:3636:PHE:HA	1.44	0.82
1:A:871:ARG:HD2	1:A:926:GLY:CA	2.09	0.82
1:C:871:ARG:HD2	1:C:926:GLY:CA	2.09	0.82
1:D:871:ARG:HD2	1:D:926:GLY:CA	2.09	0.82
1:D:2237:CYS:SG	1:D:2275:VAL:CG2	2.68	0.82
1:C:4749:GLU:O	1:C:4753:HIS:ND1	2.12	0.82
1:D:1252:HIS:O	1:D:1275:ARG:NH1	2.13	0.81
1:A:3282:PRO:O	1:A:3286:GLU:OE1	1.97	0.81
1:B:3175:LEU:HD21	1:B:3183:VAL:HG13	1.63	0.81
1:D:871:ARG:HD2	1:D:926:GLY:HA2	1.62	0.81
1:D:3282:PRO:O	1:D:3286:GLU:OE1	1.97	0.81
1:D:4749:GLU:O	1:D:4753:HIS:ND1	2.13	0.81
1:C:2237:CYS:SG	1:C:2275:VAL:CG2	2.68	0.81
1:B:4961:CYS:SG	1:B:4983:HIS:ND1	2.54	0.81
1:D:2523:ASP:HB3	1:D:2578:MET:HE1	1.63	0.81
1:C:972:LEU:CG	1:C:1041:GLN:NE2	2.38	0.81
1:C:2523:ASP:HB3	1:C:2578:MET:HE1	1.61	0.81
1:B:2237:CYS:SG	1:B:2275:VAL:CG2	2.68	0.81
1:B:4749:GLU:O	1:B:4753:HIS:ND1	2.13	0.81
1:C:1252:HIS:O	1:C:1275:ARG:NH1	2.13	0.81
1:A:2237:CYS:SG	1:A:2275:VAL:CG2	2.68	0.81
1:C:4961:CYS:SG	1:C:4983:HIS:ND1	2.54	0.81
1:A:4961:CYS:SG	1:A:4983:HIS:ND1	2.54	0.81
1:B:1156:THR:OG1	1:B:1157:GLU:OE1	1.97	0.81
1:B:1252:HIS:O	1:B:1275:ARG:NH1	2.13	0.81
1:D:3175:LEU:HD21	1:D:3183:VAL:HG13	1.62	0.81
1:A:1156:THR:OG1	1:A:1157:GLU:OE1	1.97	0.81
1:D:1156:THR:OG1	1:D:1157:GLU:OE1	1.97	0.81
1:A:2523:ASP:HB3	1:A:2578:MET:CE	2.11	0.81
1:D:2523:ASP:HB3	1:D:2578:MET:CE	2.11	0.81
1:A:1252:HIS:O	1:A:1275:ARG:NH1	2.13	0.80
1:B:3465:ASN:OD1	1:B:3467:MET:HG2	1.81	0.80
1:C:885:THR:O	1:C:889:GLN:NE2	2.14	0.80
1:B:871:ARG:HD2	1:B:926:GLY:HA2	1.62	0.80
1:B:2523:ASP:HB3	1:B:2578:MET:CE	2.11	0.80
1:D:885:THR:O	1:D:889:GLN:NE2	2.14	0.80
1:B:3282:PRO:O	1:B:3286:GLU:OE1	1.97	0.80
1:C:3175:LEU:HD21	1:C:3183:VAL:HG13	1.62	0.80

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:871:ARG:HD2	1:A:926:GLY:HA2	1.62	0.80
1:A:885:THR:O	1:A:889:GLN:NE2	2.14	0.80
1:B:885:THR:O	1:B:889:GLN:NE2	2.14	0.80
1:A:3175:LEU:HD21	1:A:3183:VAL:HG13	1.62	0.80
1:D:4961:CYS:SG	1:D:4983:HIS:ND1	2.53	0.80
1:B:972:LEU:CG	1:B:1041:GLN:NE2	2.38	0.80
1:B:2226:PRO:HB3	1:B:2267:MET:HE1	1.64	0.80
1:B:3946:GLN:OE1	1:B:3949:ARG:NH1	2.15	0.80
1:A:3521:GLY:HA2	1:A:3524:MET:HE3	1.64	0.80
1:A:4075:GLU:OE1	1:B:4736:ARG:NH1	2.14	0.80
1:A:4744:ASP:OD2	1:A:4747:SER:OG	1.99	0.79
1:D:3465:ASN:OD1	1:D:3467:MET:HG2	1.82	0.79
1:C:2523:ASP:HB3	1:C:2578:MET:CE	2.11	0.79
1:C:3465:ASN:OD1	1:C:3467:MET:HG2	1.81	0.79
1:A:3465:ASN:OD1	1:A:3467:MET:HG2	1.82	0.79
1:C:4744:ASP:OD2	1:C:4747:SER:OG	1.99	0.79
1:A:3946:GLN:OE1	1:A:3949:ARG:NH1	2.15	0.79
1:B:981:GLN:O	1:B:985:VAL:HG23	1.83	0.79
1:B:1263:THR:OG1	1:B:1265:ASP:OD1	2.01	0.79
1:B:4744:ASP:OD2	1:B:4747:SER:OG	1.99	0.79
1:A:3458:PHE:HE2	1:A:3464:ILE:CD1	1.96	0.79
1:D:941:MET:HE3	1:D:1051:TYR:CD1	2.17	0.79
1:A:384:MET:SD	1:B:166:GLY:C	2.61	0.79
1:D:3946:GLN:OE1	1:D:3949:ARG:NH1	2.15	0.79
1:C:871:ARG:HD2	1:C:926:GLY:HA2	1.62	0.79
1:C:3946:GLN:OE1	1:C:3949:ARG:NH1	2.15	0.79
1:C:3458:PHE:HE2	1:C:3464:ILE:CD1	1.96	0.79
1:B:4885:PHE:O	1:B:4889:VAL:HG22	1.84	0.78
1:C:4885:PHE:O	1:C:4889:VAL:HG22	1.83	0.78
1:A:3236:VAL:HA	1:A:3239:MET:SD	2.24	0.78
1:A:4736:ARG:NH1	1:D:4075:GLU:OE1	2.16	0.78
1:B:3236:VAL:HA	1:B:3239:MET:SD	2.24	0.78
1:D:3236:VAL:HA	1:D:3239:MET:SD	2.24	0.78
1:D:1263:THR:OG1	1:D:1265:ASP:OD1	2.01	0.78
1:A:981:GLN:O	1:A:985:VAL:HG23	1.83	0.78
1:B:3458:PHE:HE2	1:B:3464:ILE:CD1	1.95	0.78
1:B:4075:GLU:OE1	1:C:4736:ARG:NH1	2.17	0.78
1:C:981:GLN:O	1:C:985:VAL:HG23	1.83	0.78
1:A:2523:ASP:HB3	1:A:2578:MET:HE1	1.66	0.78
1:A:4885:PHE:O	1:A:4889:VAL:HG22	1.84	0.78
1:D:3458:PHE:HE2	1:D:3464:ILE:CD1	1.95	0.78

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4885:PHE:O	1:D:4889:VAL:HG22	1.84	0.78
1:A:3236:VAL:HG23	1:A:3239:MET:SD	2.25	0.77
1:A:4892:ARG:NH1	1:B:4899:ASP:OD1	2.17	0.77
1:B:110:ARG:NH2	1:B:117:TYR:OH	2.17	0.77
1:D:4736:ARG:NH1	1:C:4075:GLU:OE1	2.18	0.77
1:C:3236:VAL:HA	1:C:3239:MET:SD	2.24	0.77
1:D:961:MET:SD	1:D:963:ASN:CB	2.72	0.77
1:D:4744:ASP:OD2	1:D:4747:SER:OG	1.99	0.77
1:A:110:ARG:NH2	1:A:117:TYR:OH	2.17	0.77
1:A:961:MET:SD	1:A:963:ASN:CB	2.72	0.77
1:D:2224:ARG:HG3	1:D:2225:PHE:CD2	2.20	0.77
1:C:110:ARG:NH2	1:C:117:TYR:OH	2.17	0.77
1:C:3236:VAL:HG23	1:C:3239:MET:SD	2.24	0.77
1:A:2642:LYS:HG3	1:A:2698:MET:CE	2.15	0.77
1:B:3733:CYS:O	1:B:3766:GLN:NE2	2.18	0.77
1:C:961:MET:SD	1:C:963:ASN:CB	2.72	0.77
1:C:2742:THR:HG23	1:C:2814:LYS:HE3	1.67	0.77
1:B:961:MET:SD	1:B:963:ASN:CB	2.72	0.77
1:D:2642:LYS:HG3	1:D:2698:MET:CE	2.15	0.77
1:C:4687:TYR:O	1:C:4691:GLN:NE2	2.18	0.77
1:C:3300:ALA:HB3	1:C:3301:PRO:HD3	1.67	0.77
1:A:2742:THR:HG23	1:A:2814:LYS:HE3	1.67	0.77
1:D:1753:LYS:O	1:D:1758:ARG:HD2	1.85	0.77
1:D:2742:THR:HG23	1:D:2814:LYS:HE3	1.67	0.77
1:D:3236:VAL:HG23	1:D:3239:MET:SD	2.25	0.77
1:C:2642:LYS:HG3	1:C:2698:MET:CE	2.15	0.77
1:D:972:LEU:CG	1:D:1041:GLN:NE2	2.38	0.77
1:B:4687:TYR:O	1:B:4691:GLN:NE2	2.18	0.77
1:D:2233:CYS:CB	1:D:2237:CYS:HG	1.89	0.77
1:D:4687:TYR:O	1:D:4691:GLN:NE2	2.18	0.77
1:C:3733:CYS:O	1:C:3766:GLN:NE2	2.18	0.77
1:A:4687:TYR:O	1:A:4691:GLN:NE2	2.18	0.77
1:B:2642:LYS:HG3	1:B:2698:MET:CE	2.15	0.77
1:D:110:ARG:NH2	1:D:117:TYR:OH	2.17	0.77
1:B:2628:PHE:CE2	1:B:2734:ASN:OD1	2.38	0.76
1:D:981:GLN:O	1:D:985:VAL:HG23	1.83	0.76
1:A:2224:ARG:HG3	1:A:2225:PHE:CD2	2.20	0.76
1:A:2233:CYS:CB	1:A:2237:CYS:HG	1.91	0.76
1:B:3236:VAL:HG23	1:B:3239:MET:SD	2.25	0.76
1:C:941:MET:HE1	1:C:1051:TYR:CG	2.20	0.76
1:A:3733:CYS:O	1:A:3766:GLN:NE2	2.18	0.76

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2628:PHE:CE2	1:C:2734:ASN:OD1	2.38	0.76
1:C:1753:LYS:O	1:C:1758:ARG:HD2	1.85	0.76
1:B:2742:THR:HG23	1:B:2814:LYS:HE3	1.67	0.76
1:A:2638:LYS:HE3	1:A:2694:GLU:OE1	1.86	0.76
1:B:3300:ALA:HB3	1:B:3301:PRO:HD3	1.67	0.76
1:A:710:ASP:OD1	1:A:713:SER:OG	2.03	0.76
1:B:1255:TYR:O	1:B:1275:ARG:NH1	2.19	0.76
1:D:3733:CYS:O	1:D:3766:GLN:NE2	2.18	0.76
1:B:1753:LYS:O	1:B:1758:ARG:HD2	1.85	0.76
1:C:1263:THR:OG1	1:C:1265:ASP:OD1	2.01	0.76
1:C:2224:ARG:HG3	1:C:2225:PHE:CD2	2.20	0.76
1:A:1255:TYR:O	1:A:1275:ARG:NH1	2.19	0.76
1:A:1263:THR:OG1	1:A:1265:ASP:OD1	2.01	0.76
1:C:2638:LYS:HE3	1:C:2694:GLU:OE1	1.86	0.76
1:B:1561:VAL:HG12	1:B:1562:ILE:HG23	1.67	0.75
1:D:1255:TYR:O	1:D:1275:ARG:NH1	2.19	0.75
1:D:3300:ALA:HB3	1:D:3301:PRO:HD3	1.67	0.75
1:A:1753:LYS:O	1:A:1758:ARG:HD2	1.85	0.75
1:A:2212:VAL:HG21	1:A:2256:TYR:OH	1.86	0.75
1:B:2224:ARG:HG3	1:B:2225:PHE:CD2	2.20	0.75
1:B:4902:GLU:O	1:B:4913:ARG:NH1	2.20	0.75
1:D:2628:PHE:CE2	1:D:2734:ASN:OD1	2.38	0.75
1:D:2638:LYS:HE3	1:D:2694:GLU:OE1	1.86	0.75
1:D:4902:GLU:O	1:D:4913:ARG:NH1	2.20	0.75
1:C:4902:GLU:O	1:C:4913:ARG:NH1	2.20	0.75
1:D:2212:VAL:HG21	1:D:2256:TYR:OH	1.86	0.75
1:D:1561:VAL:HG12	1:D:1562:ILE:HG23	1.67	0.75
1:A:2628:PHE:CE2	1:A:2734:ASN:OD1	2.38	0.75
1:C:1561:VAL:HG12	1:C:1562:ILE:HG23	1.67	0.75
1:A:1561:VAL:HG12	1:A:1562:ILE:HG23	1.67	0.75
1:C:1255:TYR:O	1:C:1275:ARG:NH1	2.19	0.75
1:A:870:ILE:O	1:A:874:LEU:HD12	1.87	0.74
1:A:4902:GLU:O	1:A:4913:ARG:NH1	2.20	0.74
1:B:2638:LYS:HE3	1:B:2694:GLU:OE1	1.86	0.74
1:B:2212:VAL:HG21	1:B:2256:TYR:OH	1.86	0.74
1:D:941:MET:HE3	1:D:1051:TYR:CE1	2.21	0.74
1:A:941:MET:HE3	1:A:1051:TYR:CE1	2.22	0.74
1:A:3300:ALA:HB3	1:A:3301:PRO:HD3	1.67	0.74
1:B:1999:ARG:NH1	1:B:3636:PHE:HD1	1.86	0.74
1:D:2233:CYS:HB3	1:D:2237:CYS:HG	1.47	0.74
1:C:2212:VAL:HG21	1:C:2256:TYR:OH	1.86	0.74

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3159:ASP:HA	1:A:3218:VAL:HG22	1.69	0.74
1:B:941:MET:SD	1:B:1051:TYR:CD1	2.81	0.74
1:D:941:MET:SD	1:D:1051:TYR:CD1	2.81	0.74
1:C:2226:PRO:HB3	1:C:2267:MET:HE1	1.70	0.74
1:A:2638:LYS:CE	1:A:2694:GLU:OE1	2.36	0.74
1:B:941:MET:HE3	1:B:1051:TYR:CD1	2.22	0.74
1:B:2638:LYS:CE	1:B:2694:GLU:OE1	2.36	0.74
1:D:961:MET:CE	1:D:963:ASN:CB	2.59	0.74
1:D:870:ILE:O	1:D:874:LEU:HD12	1.87	0.73
1:A:3458:PHE:HE2	1:A:3464:ILE:HD12	1.53	0.73
1:B:384:MET:SD	1:C:166:GLY:C	2.66	0.73
1:B:2974:ILE:HD13	1:B:3049:LEU:HD12	1.71	0.73
1:D:774:ASP:OD2	1:D:1470:ARG:NH2	2.21	0.73
1:A:166:GLY:C	1:D:384:MET:SD	2.67	0.73
1:A:2186:MET:O	1:A:2192:TYR:OH	2.05	0.73
1:D:3159:ASP:HA	1:D:3218:VAL:HG22	1.69	0.73
1:A:774:ASP:OD2	1:A:1470:ARG:NH2	2.21	0.73
1:A:941:MET:HE3	1:A:1051:TYR:CD1	2.19	0.73
1:B:870:ILE:O	1:B:874:LEU:HD12	1.87	0.73
1:C:2186:MET:O	1:C:2192:TYR:OH	2.05	0.73
1:A:2974:ILE:HD13	1:A:3049:LEU:HD12	1.71	0.73
1:D:911:HIS:O	1:D:918:ARG:NH2	2.22	0.73
1:D:1999:ARG:NH1	1:D:3636:PHE:HD1	1.86	0.73
1:C:870:ILE:O	1:C:874:LEU:HD12	1.87	0.73
1:C:1999:ARG:NH1	1:C:3636:PHE:HD1	1.86	0.73
1:C:2974:ILE:HD13	1:C:3049:LEU:HD12	1.71	0.73
1:C:2929:PHE:O	1:C:2932:MET:SD	2.47	0.73
1:A:911:HIS:O	1:A:918:ARG:NH2	2.22	0.72
1:B:911:HIS:O	1:B:918:ARG:NH2	2.22	0.72
1:B:961:MET:CE	1:B:963:ASN:CB	2.59	0.72
1:B:3159:ASP:HA	1:B:3218:VAL:HG22	1.69	0.72
1:C:911:HIS:O	1:C:918:ARG:NH2	2.22	0.72
1:C:2642:LYS:NZ	1:C:2642:LYS:HG3	2.04	0.72
1:A:2929:PHE:O	1:A:2932:MET:SD	2.47	0.72
1:A:2874:MET:HE2	1:A:2939:ARG:CB	2.18	0.72
2:G:77:SER:OG	2:G:79:ASP:OD1	2.08	0.72
1:B:941:MET:HE3	1:B:1051:TYR:CE1	2.24	0.72
1:B:3458:PHE:HE2	1:B:3464:ILE:HD12	1.53	0.72
1:D:2638:LYS:CE	1:D:2694:GLU:OE1	2.36	0.72
1:D:166:GLY:C	1:C:384:MET:SD	2.67	0.72
1:D:2186:MET:O	1:D:2192:TYR:OH	2.05	0.72

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3458:PHE:HE2	1:D:3464:ILE:HD12	1.53	0.72
1:C:3343:GLN:OE1	1:C:3414:ARG:NE	2.23	0.72
1:A:1999:ARG:NH1	1:A:3636:PHE:HD1	1.86	0.72
1:B:774:ASP:OD2	1:B:1470:ARG:NH2	2.21	0.72
1:A:941:MET:SD	1:A:1051:TYR:CD1	2.81	0.72
1:B:213:TYR:CD1	1:B:340:LYS:HG2	2.25	0.72
1:B:3690:VAL:O	1:B:3693:LYS:HE3	1.90	0.72
1:C:774:ASP:OD2	1:C:1470:ARG:NH2	2.21	0.72
1:C:2638:LYS:CE	1:C:2694:GLU:OE1	2.36	0.72
1:B:941:MET:HE1	1:B:1051:TYR:CG	2.24	0.72
1:B:4063:ASP:OD1	1:B:4064:MET:N	2.23	0.72
1:D:2628:PHE:HZ	1:D:2734:ASN:OD1	1.73	0.72
1:D:2929:PHE:O	1:D:2932:MET:SD	2.47	0.72
1:B:710:ASP:OD1	1:B:713:SER:OG	2.03	0.72
1:D:213:TYR:CD1	1:D:340:LYS:HG2	2.25	0.72
1:D:4899:ASP:OD1	1:C:4892:ARG:NH1	2.23	0.72
1:C:3458:PHE:HE2	1:C:3464:ILE:HD12	1.53	0.72
1:A:213:TYR:CD1	1:A:340:LYS:HG2	2.25	0.72
1:B:2642:LYS:NZ	1:B:2642:LYS:HG3	2.04	0.72
1:D:2572:THR:HG21	1:D:2579:VAL:HG21	1.71	0.72
1:D:2974:ILE:HD13	1:D:3049:LEU:HD12	1.71	0.72
1:D:3690:VAL:O	1:D:3693:LYS:HE3	1.90	0.72
1:B:2628:PHE:HZ	1:B:2734:ASN:OD1	1.73	0.71
1:B:2929:PHE:O	1:B:2932:MET:SD	2.47	0.71
1:D:4063:ASP:OD1	1:D:4064:MET:N	2.23	0.71
1:C:3891:LEU:HD13	1:C:3899:PHE:CZ	2.25	0.71
1:C:4063:ASP:OD1	1:C:4064:MET:N	2.23	0.71
1:A:961:MET:CE	1:A:963:ASN:CB	2.59	0.71
1:A:4063:ASP:OD1	1:A:4064:MET:N	2.23	0.71
1:D:2642:LYS:NZ	1:D:2642:LYS:HG3	2.04	0.71
1:A:2572:THR:HG21	1:A:2579:VAL:HG21	1.71	0.71
1:A:3690:VAL:O	1:A:3693:LYS:HE3	1.90	0.71
2:F:77:SER:OG	2:F:79:ASP:OD1	2.08	0.71
1:C:213:TYR:CD1	1:C:340:LYS:HG2	2.25	0.71
1:C:2572:THR:HG21	1:C:2579:VAL:HG21	1.71	0.71
1:C:3690:VAL:O	1:C:3693:LYS:HE3	1.90	0.71
1:B:1087:ARG:NH1	1:B:1221:GLU:O	2.23	0.71
1:C:941:MET:SD	1:C:1051:TYR:CD1	2.81	0.71
1:A:384:MET:SD	1:B:166:GLY:O	2.48	0.71
1:B:3891:LEU:HD13	1:B:3899:PHE:CZ	2.25	0.71
1:D:3891:LEU:HD13	1:D:3899:PHE:CZ	2.25	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3159:ASP:HA	1:C:3218:VAL:HG22	1.69	0.71
1:A:3321:ARG:NH1	1:A:3325:ASN:OD1	2.23	0.71
1:B:3343:GLN:OE1	1:B:3414:ARG:NE	2.23	0.71
1:D:2761:TYR:HB3	1:D:2765:LYS:NZ	2.06	0.71
2:E:77:SER:OG	2:E:79:ASP:OD1	2.08	0.71
1:C:2761:TYR:HB3	1:C:2765:LYS:NZ	2.06	0.71
1:A:3891:LEU:HD13	1:A:3899:PHE:CZ	2.25	0.71
1:D:3321:ARG:NH1	1:D:3325:ASN:OD1	2.23	0.71
1:A:4899:ASP:OD1	1:D:4892:ARG:NH1	2.23	0.71
1:B:3321:ARG:NH1	1:B:3325:ASN:OD1	2.23	0.71
1:A:3534:MET:O	1:A:3538:THR:HG23	1.91	0.70
2:E:80:TYR:CE2	2:E:80:TYR:CD1	2.38	0.70
1:B:2572:THR:HG21	1:B:2579:VAL:HG21	1.71	0.70
1:B:2761:TYR:HB3	1:B:2765:LYS:NZ	2.06	0.70
1:D:4961:CYS:SG	1:D:4978:HIS:CE1	2.84	0.70
1:A:2628:PHE:HZ	1:A:2734:ASN:OD1	1.73	0.70
1:A:2642:LYS:NZ	1:A:2642:LYS:HG3	2.04	0.70
1:A:2011:HIS:CE1	1:A:2017:ASP:OD2	2.43	0.70
1:B:4961:CYS:SG	1:B:4978:HIS:CE1	2.84	0.70
1:D:1821:ASP:OD1	1:D:1822:GLY:N	2.25	0.70
1:C:3321:ARG:NH1	1:C:3325:ASN:OD1	2.23	0.70
1:A:4961:CYS:SG	1:A:4978:HIS:CE1	2.84	0.70
2:H:77:SER:OG	2:H:79:ASP:OD1	2.08	0.70
2:G:1:GLY:HA2	2:G:80:TYR:CE2	2.27	0.70
1:D:1087:ARG:NH1	1:D:1221:GLU:O	2.24	0.70
1:C:871:ARG:NH1	1:C:922:LEU:O	2.24	0.70
1:A:2226:PRO:HB3	1:A:2267:MET:CE	2.22	0.70
1:B:871:ARG:NH1	1:B:922:LEU:O	2.24	0.70
1:B:4892:ARG:NH1	1:C:4899:ASP:OD1	2.24	0.70
1:A:871:ARG:NH1	1:A:922:LEU:O	2.24	0.70
1:D:887:ILE:HD11	1:D:907:LEU:HD22	1.73	0.70
1:C:1087:ARG:NH1	1:C:1221:GLU:O	2.23	0.70
1:A:1087:ARG:NH1	1:A:1221:GLU:O	2.24	0.70
2:H:1:GLY:HA2	2:H:80:TYR:CE2	2.27	0.70
1:B:887:ILE:HD11	1:B:907:LEU:CD2	2.22	0.70
1:B:2226:PRO:HB3	1:B:2267:MET:CE	2.22	0.70
1:A:2761:TYR:HB3	1:A:2765:LYS:NZ	2.06	0.70
1:B:3521:GLY:HA2	1:B:3524:MET:HE2	1.74	0.70
1:D:887:ILE:HD11	1:D:907:LEU:CD2	2.22	0.70
1:A:1821:ASP:OD1	1:A:1822:GLY:N	2.24	0.69
1:B:3550:ARG:NH1	1:B:3597:GLN:OE1	2.25	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2226:PRO:HB3	1:D:2267:MET:CE	2.22	0.69
1:C:710:ASP:OD1	1:C:713:SER:OG	2.03	0.69
1:C:3550:ARG:NH1	1:C:3597:GLN:OE1	2.25	0.69
1:A:1733:GLU:HG2	1:A:2201:LEU:HD23	1.74	0.69
1:A:2244:ARG:NH2	1:A:3858:MET:SD	2.65	0.69
1:B:887:ILE:HD11	1:B:907:LEU:HD22	1.73	0.69
1:D:3550:ARG:NH1	1:D:3597:GLN:OE1	2.25	0.69
1:C:69:LEU:HD13	1:C:101:LEU:HD11	1.74	0.69
1:A:3343:GLN:OE1	1:A:3414:ARG:NE	2.23	0.69
1:D:871:ARG:NH1	1:D:922:LEU:O	2.24	0.69
1:D:2244:ARG:NH2	1:D:3858:MET:SD	2.65	0.69
1:A:887:ILE:HD11	1:A:907:LEU:CD2	2.22	0.69
1:A:3550:ARG:NH1	1:A:3597:GLN:OE1	2.25	0.69
1:A:4767:TRP:CE3	1:A:4768:LEU:HD23	2.28	0.69
1:B:3534:MET:O	1:B:3538:THR:HG23	1.91	0.69
1:D:1987:SER:O	1:D:1991:THR:OG1	2.08	0.69
1:C:887:ILE:HD11	1:C:907:LEU:CD2	2.22	0.69
1:A:69:LEU:HD13	1:A:101:LEU:HD11	1.75	0.69
1:D:1808:ARG:HD2	1:D:1853:ILE:HG22	1.75	0.69
1:D:3534:MET:O	1:D:3538:THR:HG23	1.91	0.69
1:C:4961:CYS:SG	1:C:4978:HIS:CE1	2.84	0.69
1:A:941:MET:HE1	1:A:1051:TYR:CG	2.27	0.69
1:A:2011:HIS:HE1	1:A:2017:ASP:CG	1.95	0.69
1:B:4767:TRP:CE3	1:B:4768:LEU:HD23	2.28	0.69
1:D:672:VAL:O	1:D:680:THR:OG1	2.08	0.69
1:D:2970:SER:HA	1:D:2973:PHE:CE1	2.28	0.69
1:C:2244:ARG:NH2	1:C:3858:MET:SD	2.66	0.69
1:C:2628:PHE:HZ	1:C:2734:ASN:OD1	1.73	0.69
2:F:1:GLY:HA2	2:F:80:TYR:CE2	2.27	0.69
1:B:2244:ARG:NH2	1:B:3858:MET:SD	2.65	0.69
1:B:3343:GLN:O	1:B:3346:VAL:HG12	1.93	0.69
1:D:4767:TRP:CE3	1:D:4768:LEU:HD23	2.27	0.69
1:C:925:SER:O	1:C:928:THR:OG1	2.10	0.69
1:C:1821:ASP:OD1	1:C:1822:GLY:N	2.25	0.69
1:C:2226:PRO:HB3	1:C:2267:MET:CE	2.22	0.69
1:C:3343:GLN:O	1:C:3346:VAL:HG12	1.93	0.69
1:A:581:ASN:OD1	1:A:582:HIS:N	2.26	0.69
1:A:918:ARG:O	1:A:922:LEU:HD23	1.93	0.69
1:A:1808:ARG:HD2	1:A:1853:ILE:HG22	1.75	0.69
1:A:2516:ASP:OD1	1:A:2517:PHE:N	2.26	0.69
2:E:1:GLY:HA2	2:E:80:TYR:CE2	2.27	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:69:LEU:HD13	1:B:101:LEU:HD11	1.75	0.69
1:B:384:MET:SD	1:C:166:GLY:O	2.51	0.69
1:B:1821:ASP:OD1	1:B:1822:GLY:N	2.25	0.69
1:D:69:LEU:HD13	1:D:101:LEU:HD11	1.75	0.69
1:D:451:TYR:O	1:D:474:ARG:NH2	2.26	0.69
1:D:581:ASN:OD1	1:D:582:HIS:N	2.26	0.69
1:D:2516:ASP:OD1	1:D:2517:PHE:N	2.26	0.69
1:D:3293:PRO:HB2	1:D:3298:ALA:HB2	1.75	0.69
1:C:887:ILE:HD11	1:C:907:LEU:HD22	1.73	0.69
1:C:4767:TRP:CE3	1:C:4768:LEU:HD23	2.28	0.69
1:A:2021:CYS:O	1:A:2028:ARG:NH2	2.26	0.69
1:B:1733:GLU:HG2	1:B:2201:LEU:HD23	1.74	0.69
1:D:1733:GLU:HG2	1:D:2201:LEU:HD23	1.74	0.69
1:D:2350:ALA:O	1:D:2354:VAL:HG23	1.93	0.69
1:D:3458:PHE:CE2	1:D:3464:ILE:HD12	2.28	0.69
1:C:2011:HIS:HE1	1:C:2017:ASP:CG	1.95	0.69
1:C:3284:TRP:O	1:C:3305:THR:HG21	1.93	0.69
1:C:3293:PRO:HB2	1:C:3298:ALA:HB2	1.75	0.69
1:C:3534:MET:O	1:C:3538:THR:HG23	1.91	0.69
2:F:80:TYR:CE2	2:F:80:TYR:CD1	2.38	0.69
1:A:2970:SER:HA	1:A:2973:PHE:CE1	2.28	0.68
1:D:2021:CYS:O	1:D:2028:ARG:NH2	2.26	0.68
1:B:2288:LEU:O	1:B:3849:ARG:NH1	2.26	0.68
1:B:5014:TYR:CD1	6:B:5304:PNX:HAA3	2.29	0.68
1:D:820:ARG:NH1	1:D:821:LEU:H	1.92	0.68
1:D:3343:GLN:OE1	1:D:3414:ARG:NE	2.23	0.68
1:C:918:ARG:O	1:C:922:LEU:HD23	1.93	0.68
1:C:1808:ARG:HD2	1:C:1853:ILE:HG22	1.74	0.68
1:C:2970:SER:HA	1:C:2973:PHE:CE1	2.28	0.68
1:A:820:ARG:NH1	1:A:821:LEU:H	1.92	0.68
1:A:3343:GLN:O	1:A:3346:VAL:HG12	1.93	0.68
1:D:2211:MET:O	1:D:2214:VAL:HG12	1.93	0.68
1:C:2021:CYS:O	1:C:2028:ARG:NH2	2.26	0.68
1:A:887:ILE:HD11	1:A:907:LEU:HD22	1.73	0.68
2:H:11:ASP:OD2	2:H:12:GLY:N	2.27	0.68
1:D:918:ARG:O	1:D:922:LEU:HD23	1.93	0.68
1:D:3284:TRP:O	1:D:3305:THR:HG21	1.93	0.68
1:C:2011:HIS:CE1	1:C:2017:ASP:OD2	2.43	0.68
1:A:3458:PHE:CE2	1:A:3464:ILE:HD12	2.28	0.68
1:D:941:MET:HE1	1:D:1051:TYR:CG	2.29	0.68
1:D:2011:HIS:HE1	1:D:2017:ASP:CG	1.96	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2929:PHE:CA	1:D:2932:MET:HE1	2.19	0.68
1:D:5014:TYR:CD1	6:D:5304:PNX:HAA3	2.29	0.68
1:C:581:ASN:OD1	1:C:582:HIS:N	2.26	0.68
1:C:941:MET:HE3	1:C:1051:TYR:CE1	2.28	0.68
1:C:2211:MET:O	1:C:2214:VAL:HG12	1.93	0.68
1:C:5014:TYR:CD1	6:C:5304:PNX:HAA3	2.28	0.68
1:A:54:ASN:O	1:A:56:GLN:N	2.27	0.68
1:B:581:ASN:OD1	1:B:582:HIS:N	2.26	0.68
1:B:3458:PHE:CE2	1:B:3464:ILE:HD12	2.28	0.68
1:D:1821:ASP:OD2	1:D:1837:GLN:NE2	2.26	0.68
1:D:3521:GLY:HA2	1:D:3524:MET:HE2	1.74	0.68
1:C:820:ARG:NH1	1:C:821:LEU:H	1.92	0.68
1:C:941:MET:HE3	1:C:1051:TYR:CD1	2.27	0.68
1:A:1980:LEU:HD11	1:A:1991:THR:HG23	1.75	0.68
1:A:2350:ALA:O	1:A:2354:VAL:HG23	1.93	0.68
2:E:11:ASP:OD2	2:E:12:GLY:N	2.27	0.68
1:B:1808:ARG:HD2	1:B:1853:ILE:HG22	1.74	0.68
1:B:2970:SER:HA	1:B:2973:PHE:CE1	2.28	0.68
1:B:3293:PRO:HB2	1:B:3298:ALA:HB2	1.75	0.68
1:D:2288:LEU:O	1:D:3849:ARG:NH1	2.26	0.68
1:D:3343:GLN:O	1:D:3346:VAL:HG12	1.93	0.68
1:D:4929:LEU:O	1:D:4933:GLN:OE1	2.12	0.68
1:C:1980:LEU:HD11	1:C:1991:THR:HG23	1.75	0.68
1:A:3293:PRO:HB2	1:A:3298:ALA:HB2	1.75	0.68
2:F:11:ASP:OD2	2:F:12:GLY:N	2.27	0.68
1:B:2011:HIS:HE1	1:B:2017:ASP:CG	1.96	0.68
1:A:4929:LEU:O	1:A:4933:GLN:OE1	2.12	0.68
1:B:451:TYR:O	1:B:474:ARG:NH2	2.26	0.68
1:B:820:ARG:NH1	1:B:821:LEU:H	1.92	0.68
1:B:2021:CYS:O	1:B:2028:ARG:NH2	2.26	0.68
1:B:2211:MET:O	1:B:2214:VAL:HG12	1.93	0.68
1:B:2350:ALA:O	1:B:2354:VAL:HG23	1.93	0.68
1:D:4749:GLU:HB3	1:D:4753:HIS:HE1	1.59	0.68
1:C:54:ASN:O	1:C:56:GLN:N	2.27	0.68
1:C:1821:ASP:OD2	1:C:1837:GLN:NE2	2.26	0.68
1:C:2350:ALA:O	1:C:2354:VAL:HG23	1.93	0.68
1:C:2516:ASP:OD1	1:C:2517:PHE:N	2.26	0.68
1:A:2211:MET:O	1:A:2214:VAL:HG12	1.93	0.67
1:B:1980:LEU:HD11	1:B:1991:THR:HG23	1.75	0.67
1:C:2288:LEU:O	1:C:3849:ARG:NH1	2.26	0.67
1:A:5014:TYR:CD1	6:A:5304:PNX:HAA3	2.28	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:11:ASP:OD2	2:G:12:GLY:N	2.27	0.67
1:B:54:ASN:O	1:B:56:GLN:N	2.27	0.67
1:B:918:ARG:O	1:B:922:LEU:HD23	1.93	0.67
1:B:2186:MET:O	1:B:2192:TYR:OH	2.05	0.67
1:A:4749:GLU:HB3	1:A:4753:HIS:HE1	1.59	0.67
1:B:1821:ASP:OD2	1:B:1837:GLN:NE2	2.26	0.67
1:B:2516:ASP:OD1	1:B:2517:PHE:N	2.26	0.67
1:A:1821:ASP:OD2	1:A:1837:GLN:NE2	2.26	0.67
1:A:2288:LEU:O	1:A:3849:ARG:NH1	2.26	0.67
1:B:3284:TRP:O	1:B:3305:THR:HG21	1.93	0.67
1:D:3324:VAL:HG11	1:D:3361:THR:HG22	1.76	0.67
1:C:961:MET:CE	1:C:963:ASN:CB	2.59	0.67
1:A:166:GLY:O	1:D:384:MET:SD	2.52	0.67
1:B:3239:MET:SD	1:B:3239:MET:HG2	2.31	0.67
1:B:3324:VAL:HG11	1:B:3361:THR:HG22	1.76	0.67
1:C:908:VAL:O	1:C:963:ASN:ND2	2.28	0.67
1:C:3458:PHE:CE2	1:C:3464:ILE:HD12	2.28	0.67
1:C:493:ARG:O	1:C:496:VAL:HG22	1.94	0.67
1:C:4749:GLU:HB3	1:C:4753:HIS:HE1	1.59	0.67
1:A:3825:GLU:OE1	1:A:3825:GLU:N	2.28	0.67
1:D:166:GLY:O	1:C:384:MET:SD	2.53	0.67
1:C:3239:MET:SD	1:C:3239:MET:HG2	2.31	0.67
1:A:493:ARG:O	1:A:496:VAL:HG22	1.94	0.67
1:A:4070:ASP:OD1	1:A:4071:ILE:N	2.28	0.67
1:B:961:MET:SD	1:B:963:ASN:N	2.68	0.67
1:B:4749:GLU:HB3	1:B:4753:HIS:HE1	1.59	0.67
1:C:1733:GLU:HG2	1:C:2201:LEU:HD23	1.74	0.67
1:D:54:ASN:O	1:D:56:GLN:N	2.27	0.67
1:D:908:VAL:O	1:D:963:ASN:ND2	2.28	0.67
1:D:3239:MET:HG2	1:D:3239:MET:HA	1.76	0.67
1:C:3521:GLY:HA2	1:C:3524:MET:HE2	1.74	0.67
1:A:2233:CYS:CA	1:A:2237:CYS:SG	2.83	0.67
1:A:3324:VAL:HG11	1:A:3361:THR:HG22	1.76	0.67
1:B:4929:LEU:O	1:B:4933:GLN:OE1	2.12	0.67
1:D:1980:LEU:HD11	1:D:1991:THR:HG23	1.75	0.67
1:C:3324:VAL:HG11	1:C:3361:THR:HG22	1.76	0.67
1:B:493:ARG:O	1:B:496:VAL:HG22	1.94	0.66
1:B:2874:MET:HE2	1:B:2939:ARG:CB	2.22	0.66
1:B:3825:GLU:OE1	1:B:3825:GLU:N	2.28	0.66
1:C:4929:LEU:O	1:C:4933:GLN:OE1	2.12	0.66
1:A:2572:THR:O	1:A:2572:THR:HG22	1.95	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:493:ARG:O	1:D:496:VAL:HG22	1.94	0.66
1:C:4070:ASP:OD1	1:C:4071:ILE:N	2.28	0.66
1:A:451:TYR:O	1:A:474:ARG:NH2	2.26	0.66
1:D:1980:LEU:HD21	1:D:1995:THR:HG22	1.78	0.66
1:C:2625:ARG:NH1	1:C:2629:ASP:OD1	2.25	0.66
1:A:908:VAL:O	1:A:963:ASN:ND2	2.28	0.66
1:A:925:SER:O	1:A:928:THR:OG1	2.10	0.66
1:A:2233:CYS:O	1:A:2234:ARG:C	2.34	0.66
1:A:3284:TRP:O	1:A:3305:THR:HG21	1.93	0.66
2:H:80:TYR:CE2	2:H:80:TYR:CD1	2.38	0.66
1:B:2011:HIS:CE1	1:B:2017:ASP:OD2	2.43	0.66
1:C:961:MET:SD	1:C:963:ASN:N	2.68	0.66
1:A:1068:ARG:O	1:A:1072:VAL:HG12	1.96	0.66
1:A:4910:GLU:O	1:A:4914:VAL:HG13	1.96	0.66
2:F:1:GLY:HA2	2:F:80:TYR:HE2	1.60	0.66
1:D:2642:LYS:CG	1:D:2642:LYS:CE	2.74	0.66
1:C:1980:LEU:HD21	1:C:1995:THR:HG22	1.78	0.66
1:B:4070:ASP:OD1	1:B:4071:ILE:N	2.28	0.66
1:D:4070:ASP:OD1	1:D:4071:ILE:N	2.28	0.66
1:B:908:VAL:O	1:B:963:ASN:ND2	2.28	0.66
1:D:1424:PRO:O	1:D:1428:LEU:HD23	1.96	0.66
1:D:61:ASP:OD2	1:D:402:ARG:NH2	2.28	0.66
1:D:2233:CYS:CA	1:D:2237:CYS:SG	2.84	0.66
1:D:4910:GLU:O	1:D:4914:VAL:HG13	1.96	0.66
1:C:2233:CYS:CB	1:C:2237:CYS:HG	2.02	0.66
1:A:61:ASP:OD2	1:A:402:ARG:NH2	2.28	0.66
1:B:2572:THR:O	1:B:2572:THR:HG22	1.95	0.66
1:D:925:SER:O	1:D:928:THR:OG1	2.10	0.66
1:D:2572:THR:HG21	1:D:2579:VAL:HG23	1.78	0.66
1:C:796:ARG:HH12	1:C:1619:ARG:HH12	1.44	0.66
1:C:3825:GLU:N	1:C:3825:GLU:OE1	2.28	0.66
1:A:2572:THR:HG21	1:A:2579:VAL:HG23	1.78	0.66
1:D:1980:LEU:HD21	1:D:1995:THR:CG2	2.26	0.66
1:D:3235:SER:OG	1:D:3237:GLU:OE1	2.14	0.66
1:C:2642:LYS:CG	1:C:2642:LYS:CE	2.74	0.66
1:A:1980:LEU:HD21	1:A:1995:THR:HG22	1.78	0.65
1:D:796:ARG:HH12	1:D:1619:ARG:HH12	1.44	0.65
1:D:4148:THR:HG21	1:D:4180:ARG:HH21	1.62	0.65
1:A:796:ARG:HH12	1:A:1619:ARG:HH12	1.44	0.65
2:E:1:GLY:HA2	2:E:80:TYR:HE2	1.60	0.65
1:B:796:ARG:HH12	1:B:1619:ARG:HH12	1.44	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1068:ARG:O	1:D:1072:VAL:HG12	1.96	0.65
1:D:3897:ASN:OD1	1:D:3901:ASN:ND2	2.29	0.65
1:C:1975:SER:O	1:C:1979:LEU:HD23	1.96	0.65
1:C:2233:CYS:CA	1:C:2237:CYS:SG	2.84	0.65
1:C:3897:ASN:OD1	1:C:3901:ASN:ND2	2.29	0.65
1:A:1424:PRO:O	1:A:1428:LEU:HD23	1.96	0.65
1:A:1980:LEU:HD21	1:A:1995:THR:CG2	2.26	0.65
1:A:2642:LYS:CG	1:A:2642:LYS:CE	2.74	0.65
1:A:3897:ASN:OD1	1:A:3901:ASN:ND2	2.29	0.65
1:B:2233:CYS:CA	1:B:2237:CYS:SG	2.83	0.65
1:B:3235:SER:OG	1:B:3237:GLU:OE1	2.14	0.65
1:B:3897:ASN:OD1	1:B:3901:ASN:ND2	2.29	0.65
1:C:1424:PRO:O	1:C:1428:LEU:HD23	1.96	0.65
1:B:2642:LYS:CG	1:B:2642:LYS:CE	2.74	0.65
1:C:2572:THR:HG21	1:C:2579:VAL:HG23	1.78	0.65
1:B:2625:ARG:NH1	1:B:2629:ASP:OD1	2.25	0.65
1:B:3239:MET:SD	1:B:3239:MET:HG3	2.31	0.65
1:D:1032:LYS:O	1:D:1036:ARG:HG3	1.97	0.65
1:D:2625:ARG:NH1	1:D:2629:ASP:OD1	2.25	0.65
1:C:451:TYR:O	1:C:474:ARG:NH2	2.26	0.65
1:C:4148:THR:HG21	1:C:4180:ARG:HH21	1.62	0.65
1:A:941:MET:CE	1:A:1051:TYR:CG	2.79	0.65
1:B:1980:LEU:HD21	1:B:1995:THR:HG22	1.78	0.65
1:C:1980:LEU:HD21	1:C:1995:THR:CG2	2.26	0.65
1:C:2440:MET:O	1:C:2444:GLN:OE1	2.15	0.65
1:A:1032:LYS:O	1:A:1036:ARG:HG3	1.97	0.65
1:B:1980:LEU:HD21	1:B:1995:THR:CG2	2.26	0.65
1:B:2224:ARG:HE	1:B:2225:PHE:H	1.44	0.65
1:B:2523:ASP:OD1	1:B:2524:VAL:HG13	1.97	0.65
1:D:2751:LEU:O	1:D:2755:ILE:HD12	1.97	0.65
1:C:61:ASP:OD2	1:C:402:ARG:NH2	2.28	0.65
2:G:1:GLY:HA2	2:G:80:TYR:HE2	1.60	0.65
1:B:2440:MET:O	1:B:2444:GLN:OE1	2.15	0.65
1:B:4148:THR:HG21	1:B:4180:ARG:HH21	1.62	0.65
1:D:941:MET:CE	1:D:1051:TYR:CG	2.80	0.65
1:D:2523:ASP:OD1	1:D:2524:VAL:HG13	1.97	0.65
1:C:2572:THR:HG22	1:C:2572:THR:O	1.95	0.65
1:C:2751:LEU:O	1:C:2755:ILE:HD12	1.97	0.65
1:C:3235:SER:OG	1:C:3237:GLU:OE1	2.14	0.65
1:A:924:MET:O	1:A:928:THR:HG23	1.97	0.65
1:A:1975:SER:O	1:A:1979:LEU:HD23	1.96	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4148:THR:HG21	1:A:4180:ARG:HH21	1.62	0.65
1:B:61:ASP:OD2	1:B:402:ARG:NH2	2.28	0.65
1:B:1424:PRO:O	1:B:1428:LEU:HD23	1.96	0.65
1:B:4910:GLU:O	1:B:4914:VAL:HG13	1.96	0.65
1:C:568:LEU:HD12	1:C:602:VAL:HG13	1.80	0.65
1:C:1068:ARG:O	1:C:1072:VAL:HG12	1.96	0.65
1:A:2440:MET:O	1:A:2444:GLN:OE1	2.14	0.64
1:A:2751:LEU:O	1:A:2755:ILE:HD12	1.97	0.64
1:A:3235:SER:OG	1:A:3237:GLU:OE1	2.14	0.64
1:B:568:LEU:HD12	1:B:602:VAL:HG13	1.79	0.64
1:D:3239:MET:SD	1:D:3239:MET:HG3	2.31	0.64
1:B:1032:LYS:O	1:B:1036:ARG:HG3	1.97	0.64
1:B:1068:ARG:O	1:B:1072:VAL:HG12	1.96	0.64
1:D:2572:THR:O	1:D:2572:THR:HG22	1.95	0.64
1:C:2874:MET:CE	1:C:2939:ARG:CB	2.75	0.64
1:D:1975:SER:O	1:D:1979:LEU:HD23	1.96	0.64
1:D:924:MET:O	1:D:928:THR:HG23	1.97	0.64
1:D:959:TYR:HD2	1:D:966:LYS:HE2	1.63	0.64
1:C:2226:PRO:CB	1:C:2267:MET:HE3	2.28	0.64
1:A:568:LEU:HD12	1:A:602:VAL:HG13	1.79	0.64
1:A:2625:ARG:NH1	1:A:2629:ASP:OD1	2.25	0.64
1:A:4818:MET:O	1:A:4824:ARG:NH2	2.29	0.64
2:H:1:GLY:HA2	2:H:80:TYR:HE2	1.60	0.64
1:C:4910:GLU:O	1:C:4914:VAL:HG13	1.96	0.64
1:A:2224:ARG:HE	1:A:2225:PHE:H	1.44	0.64
1:A:2523:ASP:OD1	1:A:2524:VAL:HG13	1.97	0.64
1:D:721:LEU:HD12	1:D:1476:MET:HE1	1.80	0.64
1:D:2233:CYS:O	1:D:2234:ARG:C	2.34	0.64
1:C:924:MET:O	1:C:928:THR:HG23	1.97	0.64
1:C:2523:ASP:OD1	1:C:2524:VAL:HG13	1.97	0.64
1:A:2829:GLY:O	1:A:2830:GLU:OE1	2.16	0.64
1:B:941:MET:CE	1:B:1051:TYR:CG	2.79	0.64
1:B:1975:SER:O	1:B:1979:LEU:HD23	1.96	0.64
1:B:2874:MET:CE	1:B:2939:ARG:CB	2.75	0.64
1:D:2440:MET:O	1:D:2444:GLN:OE1	2.15	0.64
1:D:2615:ARG:NH1	1:D:2663:ASN:O	2.31	0.64
1:D:3825:GLU:OE1	1:D:3825:GLU:N	2.28	0.64
1:C:1032:LYS:O	1:C:1036:ARG:HG3	1.97	0.64
1:C:2224:ARG:HE	1:C:2225:PHE:H	1.45	0.64
1:B:959:TYR:HD2	1:B:966:LYS:HE2	1.63	0.64
1:B:3840:SER:OG	1:B:3877:ASP:OD1	2.15	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2011:HIS:CE1	1:D:2017:ASP:OD2	2.43	0.64
1:A:2874:MET:CE	1:A:2939:ARG:CB	2.75	0.64
1:B:2829:GLY:O	1:B:2830:GLU:OE1	2.16	0.64
1:D:2224:ARG:HE	1:D:2225:PHE:H	1.44	0.64
1:A:959:TYR:HD2	1:A:966:LYS:HE2	1.63	0.64
1:D:568:LEU:HD12	1:D:602:VAL:HG13	1.80	0.64
1:D:880:GLU:OE1	1:D:967:PRO:HB2	1.98	0.64
1:C:880:GLU:OE1	1:C:967:PRO:HB2	1.98	0.64
1:C:4658:ILE:HD12	1:C:4796:MET:SD	2.38	0.64
1:C:4747:SER:O	1:C:4751:THR:HG23	1.98	0.64
1:D:961:MET:SD	1:D:963:ASN:N	2.68	0.63
1:D:3458:PHE:CE2	1:D:3464:ILE:CD1	2.80	0.63
1:C:2829:GLY:O	1:C:2830:GLU:OE1	2.16	0.63
1:A:961:MET:SD	1:A:963:ASN:N	2.68	0.63
1:B:925:SER:O	1:B:928:THR:OG1	2.10	0.63
1:B:2751:LEU:O	1:B:2755:ILE:HD12	1.97	0.63
1:D:2829:GLY:O	1:D:2830:GLU:OE1	2.16	0.63
1:C:3840:SER:OG	1:C:3877:ASP:OD1	2.15	0.63
1:C:4711:PHE:HB3	1:C:4712:PRO:HD3	1.80	0.63
1:A:3239:MET:SD	1:A:3239:MET:HG2	2.31	0.63
1:D:2874:MET:CE	1:D:2939:ARG:CB	2.75	0.63
1:B:880:GLU:OE1	1:B:967:PRO:HB2	1.98	0.63
1:B:2615:ARG:NH1	1:B:2663:ASN:O	2.31	0.63
1:C:2644:LEU:HD13	1:C:2678:LEU:HD21	1.80	0.63
1:B:924:MET:O	1:B:928:THR:HG23	1.97	0.63
1:B:3239:MET:CG	1:B:3239:MET:CE	2.77	0.63
1:C:959:TYR:HD2	1:C:966:LYS:HE2	1.63	0.63
1:C:1987:SER:O	1:C:1991:THR:OG1	2.08	0.63
1:C:2233:CYS:O	1:C:2234:ARG:C	2.34	0.63
1:A:4711:PHE:HB3	1:A:4712:PRO:HD3	1.80	0.63
1:C:870:ILE:HD11	1:C:1049:TYR:CD2	2.34	0.63
1:C:2615:ARG:NH1	1:C:2663:ASN:O	2.31	0.63
1:C:2623:LEU:O	1:C:2627:VAL:HG23	1.99	0.63
1:A:2615:ARG:NH1	1:A:2663:ASN:O	2.31	0.63
1:A:3368:ARG:O	1:A:3372:VAL:HG23	1.99	0.63
1:B:870:ILE:HD11	1:B:1049:TYR:CD2	2.34	0.63
1:D:881:LEU:O	1:D:885:THR:HG23	1.99	0.63
1:D:2623:LEU:O	1:D:2627:VAL:HG23	1.99	0.63
1:A:2157:GLU:O	1:A:2161:GLN:NE2	2.32	0.63
1:A:2644:LEU:HD13	1:A:2678:LEU:HD21	1.80	0.63
1:B:871:ARG:CD	1:B:926:GLY:HA2	2.29	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:881:LEU:O	1:B:885:THR:HG23	1.99	0.63
1:B:2233:CYS:O	1:B:2234:ARG:C	2.34	0.63
1:D:710:ASP:OD1	1:D:713:SER:OG	2.03	0.63
1:C:881:LEU:O	1:C:885:THR:HG23	1.99	0.63
1:A:3239:MET:SD	1:A:3239:MET:HG3	2.31	0.62
1:A:4658:ILE:HD12	1:A:4796:MET:SD	2.38	0.62
1:B:2233:CYS:CB	1:B:2237:CYS:HG	2.04	0.62
1:B:2572:THR:HG21	1:B:2579:VAL:HG23	1.78	0.62
1:B:2644:LEU:HD13	1:B:2678:LEU:HD21	1.80	0.62
1:B:4711:PHE:HB3	1:B:4712:PRO:HD3	1.80	0.62
1:D:4658:ILE:HD12	1:D:4796:MET:SD	2.38	0.62
1:C:3368:ARG:O	1:C:3372:VAL:HG23	1.99	0.62
1:A:880:GLU:OE1	1:A:967:PRO:HB2	1.98	0.62
1:A:881:LEU:O	1:A:885:THR:HG23	1.99	0.62
1:B:2623:LEU:O	1:B:2627:VAL:HG23	1.99	0.62
1:C:871:ARG:CD	1:C:926:GLY:HA2	2.29	0.62
1:C:3834:ALA:O	1:C:3838:THR:HG23	1.99	0.62
1:A:1987:SER:O	1:A:1991:THR:OG1	2.08	0.62
1:A:3239:MET:CG	1:A:3239:MET:CE	2.77	0.62
1:A:4630:TYR:OH	1:B:4860:ARG:NH1	2.30	0.62
1:B:2157:GLU:O	1:B:2161:GLN:NE2	2.32	0.62
1:B:3834:ALA:O	1:B:3838:THR:HG23	1.99	0.62
1:D:3368:ARG:O	1:D:3372:VAL:HG23	1.99	0.62
1:D:4747:SER:O	1:D:4751:THR:HG23	1.98	0.62
1:A:1577:ALA:O	1:A:1584:ARG:NH1	2.32	0.62
1:A:3458:PHE:CE2	1:A:3464:ILE:CD1	2.81	0.62
1:A:4747:SER:O	1:A:4751:THR:HG23	1.98	0.62
1:D:988:LEU:HB3	1:D:1039:LEU:HD11	1.81	0.62
1:D:3840:SER:OG	1:D:3877:ASP:OD1	2.15	0.62
1:A:972:LEU:CD1	1:A:1041:GLN:NE2	2.62	0.62
1:A:3834:ALA:O	1:A:3838:THR:HG23	1.99	0.62
1:A:3840:SER:OG	1:A:3877:ASP:OD1	2.15	0.62
1:B:4658:ILE:HD12	1:B:4796:MET:SD	2.38	0.62
1:C:941:MET:CE	1:C:1051:TYR:CG	2.79	0.62
1:A:2623:LEU:O	1:A:2627:VAL:HG23	1.99	0.62
1:B:4747:SER:O	1:B:4751:THR:HG23	1.98	0.62
1:D:3834:ALA:O	1:D:3838:THR:HG23	1.99	0.62
1:C:3239:MET:CG	1:C:3239:MET:CE	2.77	0.62
1:A:3969:ILE:HG21	1:A:3980:LEU:HD12	1.82	0.62
1:B:672:VAL:O	1:B:680:THR:OG1	2.08	0.62
1:B:3368:ARG:O	1:B:3372:VAL:HG23	1.99	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4818:MET:O	1:B:4824:ARG:NH2	2.29	0.62
1:D:870:ILE:HD11	1:D:1049:TYR:CD2	2.34	0.62
1:D:2782:ASP:OD2	1:D:2785:LEU:HD22	1.99	0.62
1:D:3213:TYR:CE1	1:D:3302:PRO:HG2	2.34	0.62
1:C:2157:GLU:O	1:C:2161:GLN:NE2	2.32	0.62
1:B:988:LEU:HB3	1:B:1039:LEU:HD11	1.81	0.62
1:B:3671:ASP:OD1	1:B:3672:ARG:N	2.33	0.62
1:D:972:LEU:CD1	1:D:1041:GLN:NE2	2.62	0.62
1:D:2157:GLU:O	1:D:2161:GLN:NE2	2.32	0.62
1:D:2644:LEU:HD13	1:D:2678:LEU:HD21	1.80	0.62
1:A:870:ILE:HD11	1:A:1049:TYR:CD2	2.34	0.62
1:A:1280:GLN:O	1:A:1281:ASN:OD1	2.18	0.62
1:A:4745:LEU:O	1:A:4749:GLU:OE1	2.18	0.62
1:B:972:LEU:CD1	1:B:1041:GLN:NE2	2.62	0.62
1:D:4711:PHE:HB3	1:D:4712:PRO:HD3	1.80	0.62
1:A:3213:TYR:CE1	1:A:3302:PRO:HG2	2.34	0.62
1:D:3239:MET:CG	1:D:3239:MET:CE	2.77	0.62
1:C:721:LEU:HD12	1:C:1476:MET:HE1	1.82	0.62
1:A:2744:ASN:OD1	1:A:2745:VAL:N	2.33	0.61
1:A:3671:ASP:OD1	1:A:3672:ARG:N	2.33	0.61
2:F:16:PRO:O	2:F:17:LYS:HE2	2.00	0.61
1:B:2226:PRO:CB	1:B:2267:MET:CE	2.78	0.61
1:B:3213:TYR:CE1	1:B:3302:PRO:HG2	2.34	0.61
1:C:3213:TYR:CE1	1:C:3302:PRO:HG2	2.34	0.61
1:C:3239:MET:SD	1:C:3239:MET:HG3	2.31	0.61
1:C:3573:MET:CB	1:C:3577:ARG:HH12	2.12	0.61
1:A:2782:ASP:OD2	1:A:2785:LEU:HD22	1.99	0.61
2:E:79:ASP:OD1	2:E:80:TYR:N	2.33	0.61
1:B:281:ARG:NH2	1:B:309:THR:OG1	2.33	0.61
1:B:2744:ASN:OD1	1:B:2745:VAL:N	2.33	0.61
1:D:2265:LEU:HD12	1:D:2266:GLY:N	2.15	0.61
1:D:2642:LYS:HG3	1:D:2698:MET:HE3	1.81	0.61
1:D:3969:ILE:HG21	1:D:3980:LEU:HD12	1.82	0.61
1:A:281:ARG:NH2	1:A:309:THR:OG1	2.33	0.61
2:G:16:PRO:O	2:G:17:LYS:HE2	2.00	0.61
1:B:1577:ALA:O	1:B:1584:ARG:NH1	2.32	0.61
1:B:2782:ASP:OD2	1:B:2785:LEU:HD22	1.99	0.61
1:B:3969:ILE:HG21	1:B:3980:LEU:HD12	1.82	0.61
1:B:4584:ASP:OD1	1:B:4585:SER:N	2.33	0.61
1:C:1280:GLN:O	1:C:1281:ASN:OD1	2.18	0.61
1:C:3671:ASP:OD1	1:C:3672:ARG:N	2.33	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4634:GLU:OE1	1:C:4636:THR:OG1	2.16	0.61
1:A:2226:PRO:CB	1:A:2267:MET:CE	2.78	0.61
1:B:906:CYS:HB2	1:B:913:LEU:HB2	1.83	0.61
1:B:1929:MET:O	1:B:1930:LYS:CB	2.49	0.61
1:B:3891:LEU:HB3	1:B:3899:PHE:CE2	2.35	0.61
1:C:988:LEU:HB3	1:C:1039:LEU:HD11	1.81	0.61
1:C:2226:PRO:CB	1:C:2267:MET:CE	2.78	0.61
1:A:988:LEU:HB3	1:A:1039:LEU:HD11	1.81	0.61
1:B:1965:TYR:O	1:B:1969:LEU:HD13	2.00	0.61
1:D:871:ARG:CD	1:D:926:GLY:HA2	2.29	0.61
1:D:906:CYS:HB2	1:D:913:LEU:HB2	1.83	0.61
1:C:281:ARG:NH2	1:C:309:THR:OG1	2.33	0.61
1:C:2782:ASP:OD2	1:C:2785:LEU:HD22	1.99	0.61
1:C:3458:PHE:CE2	1:C:3464:ILE:CD1	2.81	0.61
1:C:3891:LEU:HB3	1:C:3899:PHE:CE2	2.35	0.61
1:A:906:CYS:HB2	1:A:913:LEU:HB2	1.83	0.61
1:B:1280:GLN:O	1:B:1281:ASN:OD1	2.18	0.61
1:B:2265:LEU:HD12	1:B:2266:GLY:N	2.15	0.61
1:B:4745:LEU:O	1:B:4749:GLU:OE1	2.18	0.61
1:D:1929:MET:O	1:D:1930:LYS:CB	2.49	0.61
1:D:2744:ASN:OD1	1:D:2745:VAL:N	2.33	0.61
1:D:4745:LEU:O	1:D:4749:GLU:OE1	2.18	0.61
1:C:972:LEU:CD1	1:C:1041:GLN:NE2	2.62	0.61
1:C:2744:ASN:OD1	1:C:2745:VAL:N	2.33	0.61
1:C:3969:ILE:HG21	1:C:3980:LEU:HD12	1.82	0.61
1:C:4584:ASP:OD1	1:C:4585:SER:N	2.34	0.61
1:A:721:LEU:HD12	1:A:1476:MET:CE	2.31	0.61
1:A:1965:TYR:O	1:A:1969:LEU:HD13	2.01	0.61
1:A:2929:PHE:CA	1:A:2932:MET:HE1	2.23	0.61
1:A:4584:ASP:OD1	1:A:4585:SER:N	2.34	0.61
1:A:4634:GLU:OE1	1:A:4636:THR:OG1	2.16	0.61
1:A:4705:VAL:HG13	1:A:4711:PHE:CE1	2.36	0.61
2:H:16:PRO:O	2:H:17:LYS:HE2	2.00	0.61
1:B:2658:PRO:O	1:B:2666:VAL:HG11	2.01	0.61
1:D:721:LEU:HD12	1:D:1476:MET:CE	2.31	0.61
1:D:1965:TYR:O	1:D:1969:LEU:HD13	2.01	0.61
1:C:2874:MET:HE2	1:C:2939:ARG:CB	2.28	0.61
1:C:4705:VAL:HG13	1:C:4711:PHE:CE1	2.36	0.61
1:A:672:VAL:O	1:A:680:THR:OG1	2.08	0.61
1:A:3891:LEU:HB3	1:A:3899:PHE:CE2	2.35	0.61
1:B:3007:ASN:O	1:B:3011:THR:OG1	2.15	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3573:MET:CB	1:B:3577:ARG:HH12	2.12	0.61
1:B:4705:VAL:HG13	1:B:4711:PHE:CE1	2.36	0.61
1:D:140:ASP:OD2	1:D:142:THR:OG1	2.19	0.61
1:D:1280:GLN:O	1:D:1281:ASN:OD1	2.18	0.61
1:C:1577:ALA:O	1:C:1584:ARG:NH1	2.32	0.61
1:C:2265:LEU:HD12	1:C:2266:GLY:N	2.15	0.61
1:C:2658:PRO:O	1:C:2666:VAL:HG11	2.01	0.61
1:C:2977:LEU:O	1:C:2981:VAL:HG23	2.01	0.61
2:E:16:PRO:O	2:E:17:LYS:HE2	2.00	0.61
1:D:2382:GLU:OE1	1:D:2385:ARG:NH1	2.34	0.61
1:D:3634:ALA:O	1:D:3638:MET:HG3	2.01	0.61
1:D:3671:ASP:OD1	1:D:3672:ARG:N	2.33	0.61
1:C:4745:LEU:O	1:C:4749:GLU:OE1	2.18	0.61
1:A:1929:MET:O	1:A:1930:LYS:CB	2.49	0.60
1:A:2265:LEU:HD12	1:A:2266:GLY:N	2.15	0.60
1:B:2977:LEU:O	1:B:2981:VAL:HG23	2.01	0.60
1:D:904:HIS:ND1	1:D:906:CYS:SG	2.73	0.60
1:D:4818:MET:O	1:D:4824:ARG:NH2	2.29	0.60
1:A:2977:LEU:O	1:A:2981:VAL:HG23	2.01	0.60
1:B:140:ASP:OD2	1:B:142:THR:OG1	2.19	0.60
1:D:3692:GLU:OE1	1:D:3694:LYS:N	2.35	0.60
1:D:3891:LEU:HB3	1:D:3899:PHE:CE2	2.35	0.60
1:C:1929:MET:O	1:C:1930:LYS:CB	2.49	0.60
1:A:2890:LYS:O	1:A:2894:LEU:HD13	2.02	0.60
1:B:2890:LYS:O	1:B:2894:LEU:HD13	2.02	0.60
1:D:294:THR:O	1:D:298:GLY:N	2.35	0.60
1:D:941:MET:HE3	1:D:1051:TYR:CZ	2.36	0.60
1:D:2977:LEU:O	1:D:2981:VAL:HG23	2.01	0.60
1:C:906:CYS:HB2	1:C:913:LEU:HB2	1.83	0.60
1:C:970:LEU:HD11	1:C:1049:TYR:CE1	2.37	0.60
1:C:1965:TYR:O	1:C:1969:LEU:HD13	2.00	0.60
1:C:4677:LEU:HD23	1:C:4711:PHE:CE1	2.37	0.60
1:A:871:ARG:HD2	1:A:926:GLY:N	2.17	0.60
1:A:941:MET:HE3	1:A:1051:TYR:CZ	2.37	0.60
2:H:79:ASP:OD1	2:H:80:TYR:N	2.33	0.60
1:D:281:ARG:NH2	1:D:309:THR:OG1	2.33	0.60
1:D:1577:ALA:O	1:D:1584:ARG:NH1	2.32	0.60
1:D:2226:PRO:CB	1:D:2267:MET:CE	2.78	0.60
1:A:3573:MET:CB	1:A:3577:ARG:HH12	2.12	0.60
1:A:4976:GLU:O	1:A:4980:LEU:HD13	2.02	0.60
1:B:871:ARG:HD2	1:B:926:GLY:N	2.16	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4885:PHE:CE2	1:B:4897:ILE:HD11	2.36	0.60
1:D:2464:ASP:OD1	1:D:2465:ASP:N	2.35	0.60
1:D:3573:MET:CB	1:D:3577:ARG:HH12	2.12	0.60
1:D:4677:LEU:HD23	1:D:4711:PHE:CE1	2.37	0.60
1:A:2382:GLU:OE1	1:A:2385:ARG:NH1	2.34	0.60
1:A:2642:LYS:HG3	1:A:2698:MET:HE3	1.83	0.60
1:B:1441:ALA:N	1:B:1513:ASP:OD1	2.34	0.60
1:B:2382:GLU:OE1	1:B:2385:ARG:NH1	2.34	0.60
1:D:871:ARG:HD2	1:D:926:GLY:N	2.17	0.60
1:D:4584:ASP:OD1	1:D:4585:SER:N	2.33	0.60
1:B:296:ASP:CG	1:B:297:GLN:OE1	2.40	0.60
1:B:3458:PHE:CE2	1:B:3464:ILE:CD1	2.81	0.60
1:B:3634:ALA:O	1:B:3638:MET:HG3	2.01	0.60
1:D:970:LEU:HD11	1:D:1049:TYR:CE1	2.37	0.60
1:D:4885:PHE:CE2	1:D:4897:ILE:HD11	2.36	0.60
1:C:294:THR:O	1:C:298:GLY:N	2.35	0.60
1:C:984:LEU:O	1:C:988:LEU:HD23	2.01	0.60
1:C:2464:ASP:OD1	1:C:2465:ASP:N	2.35	0.60
1:C:4976:GLU:O	1:C:4980:LEU:HD13	2.02	0.60
1:A:932:LEU:HD12	1:A:937:CYS:SG	2.42	0.60
1:A:3227:ARG:HB3	1:A:3232:LEU:HD12	1.83	0.60
1:D:985:VAL:HG13	1:D:1039:LEU:CD2	2.32	0.60
1:C:721:LEU:HD12	1:C:1476:MET:CE	2.31	0.60
1:C:1441:ALA:N	1:C:1513:ASP:OD1	2.34	0.60
1:C:4966:ASP:OD1	1:C:4967:TYR:N	2.35	0.60
1:A:871:ARG:CD	1:A:926:GLY:HA2	2.29	0.60
1:A:3634:ALA:O	1:A:3638:MET:HG3	2.02	0.60
1:D:4705:VAL:HG13	1:D:4711:PHE:CE1	2.36	0.60
1:C:140:ASP:OD2	1:C:142:THR:OG1	2.19	0.60
1:C:2382:GLU:OE1	1:C:2385:ARG:NH1	2.34	0.60
1:C:4827:LEU:O	1:C:4831:THR:HG23	2.02	0.60
1:A:296:ASP:CG	1:A:297:GLN:OE1	2.40	0.60
1:A:2233:CYS:HB3	1:A:2237:CYS:HG	1.49	0.60
1:A:2658:PRO:O	1:A:2666:VAL:HG11	2.01	0.60
1:A:4677:LEU:HD23	1:A:4711:PHE:CE1	2.37	0.60
1:B:3692:GLU:OE1	1:B:3694:LYS:N	2.35	0.60
1:B:4976:GLU:O	1:B:4980:LEU:HD13	2.02	0.60
1:D:932:LEU:HD12	1:D:937:CYS:SG	2.42	0.60
1:D:984:LEU:O	1:D:988:LEU:HD23	2.01	0.60
1:D:2658:PRO:O	1:D:2666:VAL:HG11	2.01	0.60
1:D:4976:GLU:O	1:D:4980:LEU:HD13	2.02	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:985:VAL:HG13	1:A:1039:LEU:CD2	2.32	0.59
1:B:721:LEU:HD12	1:B:1476:MET:CE	2.31	0.59
1:B:884:LEU:O	1:B:888:GLU:OE1	2.20	0.59
1:B:932:LEU:HD12	1:B:937:CYS:SG	2.42	0.59
1:C:3692:GLU:OE1	1:C:3694:LYS:N	2.35	0.59
1:A:140:ASP:OD2	1:A:142:THR:OG1	2.19	0.59
1:A:721:LEU:HD12	1:A:1476:MET:HE1	1.84	0.59
1:A:984:LEU:O	1:A:988:LEU:HD23	2.01	0.59
1:B:970:LEU:HD11	1:B:1049:TYR:CE1	2.37	0.59
1:B:4677:LEU:HD23	1:B:4711:PHE:CE1	2.37	0.59
1:B:4827:LEU:O	1:B:4831:THR:HG23	2.02	0.59
1:D:4634:GLU:OE1	1:D:4636:THR:OG1	2.16	0.59
1:C:871:ARG:HD2	1:C:926:GLY:N	2.17	0.59
1:C:932:LEU:HD12	1:C:937:CYS:SG	2.42	0.59
1:C:2890:LYS:O	1:C:2894:LEU:HD13	2.02	0.59
1:C:4885:PHE:CE2	1:C:4897:ILE:HD11	2.36	0.59
1:A:11:VAL:HG11	1:A:164:ARG:HD3	1.85	0.59
1:B:985:VAL:HG13	1:B:1039:LEU:CD2	2.32	0.59
1:D:2890:LYS:O	1:D:2894:LEU:HD13	2.02	0.59
1:A:970:LEU:HD11	1:A:1049:TYR:CE1	2.37	0.59
1:A:3327:LEU:HD12	1:A:3364:ARG:NH1	2.18	0.59
1:A:3692:GLU:OE1	1:A:3694:LYS:N	2.35	0.59
1:A:4827:LEU:O	1:A:4831:THR:HG23	2.02	0.59
1:B:233:ILE:O	1:B:257:ARG:NH1	2.35	0.59
1:B:984:LEU:O	1:B:988:LEU:HD23	2.01	0.59
1:B:3327:LEU:HD12	1:B:3364:ARG:NH1	2.18	0.59
1:D:4966:ASP:OD1	1:D:4967:TYR:N	2.35	0.59
1:C:296:ASP:CG	1:C:297:GLN:OE1	2.40	0.59
1:C:3157:ILE:HG23	1:C:3161:VAL:HG12	1.85	0.59
1:C:3634:ALA:O	1:C:3638:MET:HG3	2.02	0.59
1:A:884:LEU:O	1:A:888:GLU:OE1	2.20	0.59
1:A:4885:PHE:CE2	1:A:4897:ILE:HD11	2.36	0.59
2:G:79:ASP:OD1	2:G:80:TYR:N	2.33	0.59
2:F:79:ASP:OD1	2:F:80:TYR:N	2.33	0.59
1:D:11:VAL:HG11	1:D:164:ARG:HD3	1.85	0.59
1:D:2874:MET:HE2	1:D:2939:ARG:CB	2.27	0.59
1:C:978:THR:HG1	1:C:981:GLN:CD	2.06	0.59
1:A:64:ILE:O	1:A:111:HIS:NE2	2.36	0.59
1:A:1427:ILE:HG23	1:A:1428:LEU:HD22	1.84	0.59
1:C:985:VAL:HG13	1:C:1039:LEU:CD2	2.32	0.59
1:C:2376:LEU:HD11	1:C:2426:TYR:HB3	1.85	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:294:THR:O	1:A:298:GLY:N	2.35	0.59
2:F:17:LYS:HE2	2:F:17:LYS:HA	1.85	0.59
1:B:978:THR:HG1	1:B:981:GLN:CD	2.06	0.59
1:B:2376:LEU:HD11	1:B:2426:TYR:HB3	1.85	0.59
1:B:2464:ASP:OD1	1:B:2465:ASP:N	2.35	0.59
1:D:1427:ILE:HG23	1:D:1428:LEU:HD22	1.84	0.59
1:D:3327:LEU:HD12	1:D:3364:ARG:NH1	2.18	0.59
1:C:2642:LYS:HG3	1:C:2698:MET:HE1	1.82	0.59
1:C:3236:VAL:O	1:C:3239:MET:HG2	2.02	0.59
1:C:3327:LEU:HD12	1:C:3364:ARG:NH1	2.18	0.59
1:C:3357:HIS:O	1:C:3361:THR:HG23	2.03	0.59
1:A:4930:ALA:HA	1:A:4933:GLN:OE1	2.03	0.59
1:B:3157:ILE:HG23	1:B:3161:VAL:HG12	1.85	0.59
1:D:3985:LEU:O	1:D:3989:VAL:HG23	2.03	0.59
1:A:2376:LEU:HD11	1:A:2426:TYR:HB3	1.85	0.59
1:A:3985:LEU:O	1:A:3989:VAL:HG23	2.03	0.59
2:E:17:LYS:HE2	2:E:17:LYS:HA	1.85	0.59
1:B:11:VAL:HG11	1:B:164:ARG:HD3	1.85	0.59
1:B:2761:TYR:HB3	1:B:2765:LYS:HZ3	1.68	0.59
1:B:3227:ARG:HB3	1:B:3232:LEU:HD12	1.83	0.59
1:D:228:ASP:OD1	1:D:229:GLU:N	2.36	0.59
1:C:3227:ARG:HB3	1:C:3232:LEU:HD12	1.83	0.59
1:A:3157:ILE:HG23	1:A:3161:VAL:HG12	1.85	0.58
2:G:17:LYS:HE2	2:G:17:LYS:HA	1.85	0.58
1:B:3985:LEU:O	1:B:3989:VAL:HG23	2.03	0.58
1:B:4930:ALA:HA	1:B:4933:GLN:OE1	2.03	0.58
1:D:2974:ILE:HD13	1:D:3049:LEU:CD1	2.33	0.58
1:C:11:VAL:HG11	1:C:164:ARG:HD3	1.85	0.58
1:C:2874:MET:HE1	1:C:2939:ARG:HB2	1.83	0.58
1:A:228:ASP:OD1	1:A:229:GLU:N	2.36	0.58
1:A:2464:ASP:OD1	1:A:2465:ASP:N	2.35	0.58
1:A:3324:VAL:CG1	1:A:3361:THR:HG22	2.33	0.58
1:A:4998:LYS:HB3	1:A:5003:HIS:HE1	1.68	0.58
1:B:64:ILE:O	1:B:111:HIS:NE2	2.36	0.58
1:B:3324:VAL:CG1	1:B:3361:THR:HG22	2.33	0.58
1:B:3357:HIS:O	1:B:3361:THR:HG23	2.03	0.58
1:B:4749:GLU:C	1:B:4753:HIS:HD1	2.07	0.58
1:A:4966:ASP:OD1	1:A:4967:TYR:N	2.35	0.58
1:D:978:THR:HG1	1:D:981:GLN:CD	2.06	0.58
1:D:3239:MET:SD	1:D:3239:MET:HG2	2.31	0.58
1:C:64:ILE:O	1:C:111:HIS:NE2	2.36	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4930:ALA:HA	1:C:4933:GLN:OE1	2.03	0.58
1:A:1112:ASP:OD1	1:A:1113:VAL:HG23	2.03	0.58
1:A:3559:LEU:HD23	1:A:3559:LEU:O	2.03	0.58
2:H:17:LYS:HE2	2:H:17:LYS:HA	1.85	0.58
1:B:941:MET:HE3	1:B:1051:TYR:CZ	2.38	0.58
1:D:296:ASP:CG	1:D:297:GLN:OE1	2.40	0.58
1:D:2376:LEU:HD11	1:D:2426:TYR:HB3	1.85	0.58
1:D:3559:LEU:HD23	1:D:3559:LEU:O	2.03	0.58
1:D:4827:LEU:O	1:D:4831:THR:HG23	2.02	0.58
1:C:1427:ILE:HG23	1:C:1428:LEU:HD22	1.84	0.58
1:C:1925:GLY:O	1:C:1929:MET:HG3	2.04	0.58
1:C:2974:ILE:HD13	1:C:3049:LEU:CD1	2.33	0.58
1:C:4818:MET:O	1:C:4824:ARG:NH2	2.29	0.58
1:A:3357:HIS:O	1:A:3361:THR:HG23	2.03	0.58
1:B:228:ASP:OD1	1:B:229:GLU:N	2.36	0.58
1:B:294:THR:O	1:B:298:GLY:N	2.35	0.58
1:B:2226:PRO:CB	1:B:2267:MET:HE3	2.33	0.58
1:B:2572:THR:CG2	1:B:2579:VAL:HG21	2.33	0.58
1:B:2781:VAL:HG23	1:B:2789:PRO:HG3	1.85	0.58
1:D:69:LEU:CD1	1:D:101:LEU:HD11	2.34	0.58
1:C:1112:ASP:OD1	1:C:1113:VAL:HG23	2.03	0.58
1:A:978:THR:HG1	1:A:981:GLN:CD	2.06	0.58
1:A:2226:PRO:CB	1:A:2267:MET:HE1	2.31	0.58
1:D:64:ILE:O	1:D:111:HIS:NE2	2.36	0.58
1:D:3157:ILE:HG23	1:D:3161:VAL:HG12	1.85	0.58
1:D:3324:VAL:CG1	1:D:3361:THR:HG22	2.33	0.58
1:D:4930:ALA:HA	1:D:4933:GLN:OE1	2.03	0.58
1:A:3236:VAL:HA	1:A:3239:MET:HG2	1.85	0.58
1:A:3398:PHE:CE2	1:A:3450:ASN:HB2	2.38	0.58
1:B:1112:ASP:OD1	1:B:1113:VAL:HG23	2.03	0.58
1:B:4966:ASP:OD1	1:B:4967:TYR:N	2.35	0.58
1:D:2572:THR:CG2	1:D:2579:VAL:HG21	2.33	0.58
1:C:69:LEU:CD1	1:C:101:LEU:HD11	2.34	0.58
1:C:4749:GLU:C	1:C:4753:HIS:HD1	2.07	0.58
1:A:1925:GLY:O	1:A:1929:MET:HG3	2.04	0.58
1:C:228:ASP:OD1	1:C:229:GLU:N	2.36	0.58
1:C:884:LEU:O	1:C:888:GLU:OE1	2.20	0.58
1:C:3985:LEU:O	1:C:3989:VAL:HG23	2.03	0.58
1:A:904:HIS:ND1	1:A:906:CYS:SG	2.73	0.58
1:A:2572:THR:CG2	1:A:2579:VAL:HG21	2.33	0.58
1:B:3398:PHE:CE2	1:B:3450:ASN:HB2	2.38	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3442:PHE:CG	1:B:3514:LEU:HD22	2.39	0.58
1:B:4998:LYS:HB3	1:B:5003:HIS:HE1	1.68	0.58
1:D:884:LEU:O	1:D:888:GLU:OE1	2.20	0.58
1:D:3227:ARG:HB3	1:D:3232:LEU:HD12	1.83	0.58
1:A:2974:ILE:HD13	1:A:3049:LEU:CD1	2.33	0.58
1:A:3372:VAL:CG1	1:A:3398:PHE:CE1	2.87	0.58
1:B:1925:GLY:O	1:B:1929:MET:HG3	2.04	0.58
1:B:3559:LEU:O	1:B:3559:LEU:HD23	2.03	0.58
1:C:3398:PHE:CE2	1:C:3450:ASN:HB2	2.38	0.58
1:C:3442:PHE:CG	1:C:3514:LEU:HD22	2.39	0.58
1:A:2781:VAL:HG23	1:A:2789:PRO:HG3	1.86	0.57
1:B:110:ARG:HD3	1:B:115:ARG:HE	1.70	0.57
1:D:3398:PHE:CE2	1:D:3450:ASN:HB2	2.39	0.57
1:B:1427:ILE:HG23	1:B:1428:LEU:HD22	1.84	0.57
1:D:110:ARG:HD3	1:D:115:ARG:HE	1.70	0.57
1:D:233:ILE:O	1:D:257:ARG:NH1	2.35	0.57
1:D:4749:GLU:C	1:D:4753:HIS:HD1	2.07	0.57
1:C:233:ILE:O	1:C:257:ARG:NH1	2.35	0.57
1:C:3559:LEU:O	1:C:3559:LEU:HD23	2.03	0.57
1:A:69:LEU:CD1	1:A:101:LEU:HD11	2.34	0.57
1:B:3566:SER:HB3	1:B:3569:LEU:HD12	1.86	0.57
1:D:1112:ASP:OD1	1:D:1113:VAL:HG23	2.03	0.57
1:D:3357:HIS:O	1:D:3361:THR:HG23	2.03	0.57
1:C:2586:VAL:HG13	1:C:2607:LEU:HD13	1.87	0.57
1:C:3566:SER:HB3	1:C:3569:LEU:HD12	1.86	0.57
1:A:1441:ALA:N	1:A:1513:ASP:OD1	2.34	0.57
1:B:69:LEU:CD1	1:B:101:LEU:HD11	2.34	0.57
1:B:1748:PHE:HB3	1:B:1758:ARG:NH1	2.20	0.57
1:B:4634:GLU:OE1	1:B:4636:THR:OG1	2.16	0.57
1:D:554:LEU:HD11	1:D:1593:PRO:CG	2.35	0.57
1:D:3442:PHE:CG	1:D:3514:LEU:HD22	2.39	0.57
1:C:904:HIS:ND1	1:C:906:CYS:SG	2.73	0.57
1:C:2572:THR:CG2	1:C:2579:VAL:HG21	2.33	0.57
1:A:1748:PHE:HB3	1:A:1758:ARG:NH1	2.20	0.57
1:B:3372:VAL:CG1	1:B:3398:PHE:CE1	2.87	0.57
1:B:4630:TYR:OH	1:C:4860:ARG:NH1	2.37	0.57
1:D:4998:LYS:HB3	1:D:5003:HIS:HE1	1.68	0.57
1:C:2382:GLU:OE2	1:C:2392:ARG:NH2	2.38	0.57
1:C:3372:VAL:CG1	1:C:3398:PHE:CE1	2.87	0.57
1:C:4998:LYS:HB3	1:C:5003:HIS:HE1	1.68	0.57
1:A:3254:GLY:O	1:A:3258:GLU:OE1	2.23	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1441:ALA:N	1:D:1513:ASP:OD1	2.34	0.57
1:D:3498:ARG:O	1:D:3501:ASP:N	2.38	0.57
1:C:1748:PHE:HB3	1:C:1758:ARG:NH1	2.20	0.57
1:C:3254:GLY:O	1:C:3258:GLU:OE1	2.23	0.57
1:C:3324:VAL:CG1	1:C:3361:THR:HG22	2.33	0.57
1:A:110:ARG:HD3	1:A:115:ARG:HE	1.70	0.57
1:A:554:LEU:HD11	1:A:1593:PRO:CG	2.35	0.57
1:A:2382:GLU:OE2	1:A:2392:ARG:NH2	2.38	0.57
1:A:3442:PHE:CG	1:A:3514:LEU:HD22	2.39	0.57
1:D:1748:PHE:HB3	1:D:1758:ARG:NH1	2.19	0.57
1:D:1925:GLY:O	1:D:1929:MET:HG3	2.04	0.57
1:D:3179:LYS:O	1:D:3180:ASN:ND2	2.38	0.57
1:C:1497:GLY:O	1:C:1501:VAL:HG13	2.05	0.57
1:A:3498:ARG:O	1:A:3501:ASP:N	2.38	0.57
1:B:3179:LYS:O	1:B:3180:ASN:ND2	2.38	0.57
1:B:3236:VAL:HA	1:B:3239:MET:HG2	1.86	0.57
1:D:1497:GLY:O	1:D:1501:VAL:HG13	2.05	0.57
1:B:554:LEU:HD11	1:B:1593:PRO:CG	2.35	0.57
1:B:2974:ILE:HD13	1:B:3049:LEU:CD1	2.33	0.57
1:D:3372:VAL:CG1	1:D:3398:PHE:CE1	2.87	0.57
1:C:554:LEU:HD11	1:C:1593:PRO:CG	2.35	0.57
1:C:3179:LYS:O	1:C:3180:ASN:ND2	2.38	0.57
1:A:2761:TYR:HB3	1:A:2765:LYS:HZ3	1.68	0.57
1:A:2782:ASP:N	1:A:2782:ASP:OD1	2.38	0.57
1:B:2382:GLU:OE2	1:B:2392:ARG:NH2	2.38	0.57
1:B:2586:VAL:HG13	1:B:2607:LEU:HD13	1.87	0.57
1:B:3498:ARG:O	1:B:3501:ASP:N	2.38	0.57
1:A:3179:LYS:O	1:A:3180:ASN:ND2	2.38	0.56
1:A:4749:GLU:C	1:A:4753:HIS:HD1	2.07	0.56
1:B:3137:LEU:HD23	1:B:3189:ALA:HB1	1.86	0.56
1:B:3254:GLY:O	1:B:3258:GLU:OE1	2.23	0.56
1:D:2780:ASN:OD1	1:D:2781:VAL:N	2.38	0.56
1:C:3137:LEU:HD23	1:C:3189:ALA:HB1	1.86	0.56
1:A:907:LEU:O	1:A:908:VAL:HB	2.05	0.56
1:B:812:HIS:O	1:B:815:VAL:HG12	2.05	0.56
1:D:2226:PRO:CB	1:D:2267:MET:HE1	2.34	0.56
1:D:2586:VAL:HG13	1:D:2607:LEU:HD13	1.87	0.56
1:D:3254:GLY:O	1:D:3258:GLU:OE1	2.23	0.56
1:C:110:ARG:HD3	1:C:115:ARG:HE	1.70	0.56
1:C:2780:ASN:OD1	1:C:2781:VAL:N	2.38	0.56
1:B:553:ARG:NE	1:B:555:GLU:OE2	2.38	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:904:HIS:ND1	1:B:906:CYS:SG	2.73	0.56
1:B:5034:ASP:OD1	1:B:5035:GLN:N	2.38	0.56
1:D:812:HIS:O	1:D:815:VAL:HG12	2.05	0.56
1:D:1980:LEU:CD1	1:D:1991:THR:HG23	2.36	0.56
1:D:2382:GLU:OE2	1:D:2392:ARG:NH2	2.38	0.56
1:D:2782:ASP:N	1:D:2782:ASP:OD1	2.38	0.56
1:D:4860:ARG:NH1	1:C:4630:TYR:OH	2.36	0.56
1:C:812:HIS:O	1:C:815:VAL:HG12	2.05	0.56
1:C:2781:VAL:HG23	1:C:2789:PRO:HG3	1.86	0.56
1:C:3498:ARG:O	1:C:3501:ASP:N	2.38	0.56
1:B:721:LEU:HD12	1:B:1476:MET:HE1	1.87	0.56
1:B:907:LEU:O	1:B:908:VAL:HB	2.05	0.56
1:D:1461:ASP:OD2	1:D:1468:LYS:NZ	2.37	0.56
1:A:233:ILE:O	1:A:257:ARG:NH1	2.35	0.56
1:A:384:MET:CE	1:B:167:ASP:OD1	2.54	0.56
1:A:1980:LEU:CD1	1:A:1991:THR:HG23	2.35	0.56
1:A:3137:LEU:HD23	1:A:3189:ALA:HB1	1.86	0.56
1:B:991:ASN:O	1:B:995:VAL:HG23	2.05	0.56
1:B:2780:ASN:OD1	1:B:2781:VAL:N	2.38	0.56
1:D:2572:THR:O	1:D:2618:MET:HE2	2.06	0.56
1:D:2781:VAL:HG23	1:D:2789:PRO:HG3	1.85	0.56
1:D:3137:LEU:HD23	1:D:3189:ALA:HB1	1.86	0.56
1:D:5034:ASP:OD1	1:D:5035:GLN:N	2.38	0.56
1:C:1461:ASP:OD2	1:C:1468:LYS:NZ	2.37	0.56
1:C:1980:LEU:CD1	1:C:1991:THR:HG23	2.36	0.56
1:C:3629:ARG:O	1:C:3633:VAL:HG23	2.06	0.56
1:A:167:ASP:HA	1:D:384:MET:SD	2.45	0.56
1:A:1497:GLY:O	1:A:1501:VAL:HG13	2.05	0.56
1:A:5034:ASP:OD1	1:A:5035:GLN:N	2.38	0.56
1:B:861:ILE:HD13	1:B:930:LYS:HD3	1.87	0.56
1:B:1497:GLY:O	1:B:1501:VAL:HG13	2.05	0.56
1:B:3327:LEU:HD12	1:B:3364:ARG:HH12	1.70	0.56
1:B:3398:PHE:HE2	1:B:3450:ASN:HB2	1.71	0.56
1:B:4768:LEU:HA	1:B:4771:ILE:HD12	1.87	0.56
1:D:897:ARG:HB2	1:D:905:PRO:HD3	1.88	0.56
1:D:3114:LYS:HD2	1:D:3125:VAL:HG21	1.88	0.56
1:C:907:LEU:O	1:C:908:VAL:HB	2.05	0.56
1:C:5034:ASP:OD1	1:C:5035:GLN:N	2.38	0.56
1:A:812:HIS:O	1:A:815:VAL:HG12	2.05	0.56
1:A:3566:SER:HB3	1:A:3569:LEU:HD12	1.86	0.56
1:A:3629:ARG:O	1:A:3633:VAL:HG23	2.06	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2608:MET:O	1:D:2612[B]:ARG:HG3	2.06	0.56
1:C:991:ASN:O	1:C:995:VAL:HG23	2.05	0.56
1:A:554:LEU:HD11	1:A:1593:PRO:HG3	1.88	0.56
1:D:991:ASN:O	1:D:995:VAL:HG23	2.05	0.56
1:D:2212:VAL:HG21	1:D:2256:TYR:HH	1.71	0.56
1:D:4768:LEU:HA	1:D:4771:ILE:HD12	1.87	0.56
1:C:553:ARG:NE	1:C:555:GLU:OE2	2.38	0.56
1:C:3114:LYS:HD2	1:C:3125:VAL:HG21	1.88	0.56
1:C:3398:PHE:HE2	1:C:3450:ASN:HB2	1.71	0.56
1:A:2523:ASP:HB3	1:A:2578:MET:HE2	1.88	0.56
1:A:2586:VAL:HG13	1:A:2607:LEU:HD13	1.87	0.56
1:A:3852:LYS:O	1:A:3856:LEU:HG	2.06	0.56
1:B:1157:GLU:OE1	1:B:1157:GLU:N	2.39	0.56
1:B:4061:PHE:HA	1:B:4064:MET:HE3	1.88	0.56
1:D:554:LEU:HD11	1:D:1593:PRO:HG3	1.88	0.56
1:D:3566:SER:HB3	1:D:3569:LEU:HD12	1.86	0.56
1:D:3629:ARG:O	1:D:3633:VAL:HG23	2.06	0.56
1:C:897:ARG:HB2	1:C:905:PRO:HD3	1.88	0.56
1:C:941:MET:HE3	1:C:1051:TYR:CZ	2.41	0.56
1:A:167:ASP:OD1	1:D:384:MET:CE	2.54	0.56
1:A:991:ASN:O	1:A:995:VAL:HG23	2.05	0.56
1:A:3166:TYR:OH	1:A:3203:VAL:HG11	2.06	0.56
1:A:3372:VAL:HG12	1:A:3398:PHE:HE1	1.71	0.56
1:A:3398:PHE:HE2	1:A:3450:ASN:HB2	1.71	0.56
1:A:4768:LEU:HA	1:A:4771:ILE:HD12	1.87	0.56
1:C:125:ARG:HB3	1:C:125:ARG:HH11	1.71	0.56
1:C:499:THR:HG23	1:C:502:HIS:H	1.71	0.56
1:B:499:THR:HG23	1:B:502:HIS:H	1.71	0.55
1:B:897:ARG:HB2	1:B:905:PRO:HD3	1.87	0.55
1:B:2608:MET:O	1:B:2612[B]:ARG:HG3	2.06	0.55
1:D:213:TYR:CE1	1:D:340:LYS:HG2	2.42	0.55
1:D:3327:LEU:HD12	1:D:3364:ARG:HH12	1.70	0.55
1:C:2572:THR:O	1:C:2618:MET:HE2	2.06	0.55
1:C:4651:THR:CB	1:C:4803:HIS:HE2	2.19	0.55
1:A:2780:ASN:OD1	1:A:2781:VAL:N	2.38	0.55
1:A:3327:LEU:HD12	1:A:3364:ARG:HH12	1.70	0.55
1:B:1980:LEU:CD1	1:B:1991:THR:HG23	2.36	0.55
1:B:2642:LYS:HG3	1:B:2698:MET:HE3	1.86	0.55
1:B:3852:LYS:O	1:B:3856:LEU:HG	2.06	0.55
1:D:861:ILE:HD13	1:D:930:LYS:HD3	1.87	0.55
1:D:2211:MET:O	1:D:2215:LEU:HD13	2.06	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4749:GLU:HB3	1:D:4753:HIS:CE1	2.41	0.55
1:C:2608:MET:O	1:C:2612[B]:ARG:HG3	2.06	0.55
1:B:3166:TYR:OH	1:B:3203:VAL:HG11	2.06	0.55
1:B:3239:MET:CG	1:B:3239:MET:HA	2.35	0.55
1:D:125:ARG:HH11	1:D:125:ARG:HB3	1.71	0.55
1:D:907:LEU:O	1:D:908:VAL:HB	2.05	0.55
1:D:928:THR:O	1:D:932:LEU:HD23	2.07	0.55
1:C:213:TYR:CE1	1:C:340:LYS:HG2	2.42	0.55
1:C:861:ILE:HD13	1:C:930:LYS:HD3	1.87	0.55
1:C:4061:PHE:HA	1:C:4064:MET:HE3	1.88	0.55
1:A:67:PHE:HB3	1:A:109:LEU:HD22	1.89	0.55
1:A:1157:GLU:OE1	1:A:1157:GLU:N	2.39	0.55
1:A:3114:LYS:HD2	1:A:3125:VAL:HG21	1.88	0.55
1:D:3372:VAL:HG12	1:D:3398:PHE:HE1	1.71	0.55
1:C:2211:MET:O	1:C:2215:LEU:HD13	2.06	0.55
1:C:3327:LEU:HD12	1:C:3364:ARG:HH12	1.70	0.55
1:A:1931:LEU:HD22	1:A:1935:VAL:HG11	1.88	0.55
1:A:4860:ARG:NH1	1:D:4630:TYR:OH	2.39	0.55
1:B:928:THR:O	1:B:932:LEU:HD23	2.07	0.55
1:B:3629:ARG:O	1:B:3633:VAL:HG23	2.06	0.55
1:C:928:THR:O	1:C:932:LEU:HD23	2.06	0.55
1:C:1931:LEU:HD22	1:C:1935:VAL:HG11	1.88	0.55
1:C:3635:CYS:HA	1:C:3638:MET:HG3	1.88	0.55
1:C:4749:GLU:HB3	1:C:4753:HIS:CE1	2.41	0.55
1:A:861:ILE:HD13	1:A:930:LYS:HD3	1.87	0.55
1:A:897:ARG:HB2	1:A:905:PRO:HD3	1.88	0.55
1:A:2211:MET:O	1:A:2215:LEU:HD13	2.06	0.55
1:B:2211:MET:O	1:B:2215:LEU:HD13	2.06	0.55
1:D:499:THR:HG23	1:D:502:HIS:H	1.71	0.55
1:D:4061:PHE:HA	1:D:4064:MET:HE3	1.89	0.55
1:A:553:ARG:NE	1:A:555:GLU:OE2	2.38	0.55
1:A:908:VAL:O	1:A:963:ASN:CG	2.45	0.55
1:B:67:PHE:HB3	1:B:109:LEU:HD22	1.89	0.55
1:B:1726:SER:O	1:B:1729:SER:OG	2.23	0.55
1:B:4651:THR:CB	1:B:4803:HIS:HE2	2.19	0.55
1:D:2874:MET:HE1	1:D:2939:ARG:HB2	1.84	0.55
1:D:3398:PHE:HE2	1:D:3450:ASN:HB2	1.71	0.55
1:D:4205:TRP:CE3	1:D:4989:MET:HE2	2.41	0.55
1:C:2233:CYS:HB3	1:C:2237:CYS:HG	1.60	0.55
1:A:112:ALA:O	1:A:115:ARG:NH1	2.39	0.55
1:A:499:THR:HG23	1:A:502:HIS:H	1.71	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1158:ASN:ND2	1:B:1182:ILE:O	2.40	0.55
1:B:3635:CYS:HA	1:B:3638:MET:HG3	1.88	0.55
1:D:3852:LYS:O	1:D:3856:LEU:HG	2.06	0.55
1:D:4651:THR:CB	1:D:4803:HIS:HE2	2.19	0.55
1:C:1157:GLU:OE1	1:C:1157:GLU:N	2.39	0.55
1:C:1726:SER:O	1:C:1729:SER:OG	2.23	0.55
1:C:2782:ASP:OD1	1:C:2782:ASP:N	2.38	0.55
1:A:125:ARG:HB3	1:A:125:ARG:HH11	1.71	0.55
1:A:928:THR:O	1:A:932:LEU:HD23	2.06	0.55
1:A:2608:MET:O	1:A:2612[B]:ARG:HG3	2.06	0.55
1:A:4061:PHE:HA	1:A:4064:MET:HE3	1.88	0.55
1:B:112:ALA:O	1:B:115:ARG:NH1	2.38	0.55
1:B:2904:LEU:O	1:B:2904:LEU:HD23	2.07	0.55
1:C:1158:ASN:ND2	1:C:1182:ILE:O	2.40	0.55
1:C:4768:LEU:HA	1:C:4771:ILE:HD12	1.87	0.55
1:A:213:TYR:CE1	1:A:340:LYS:HG2	2.42	0.55
1:B:908:VAL:HG13	1:B:912:SER:HB2	1.89	0.55
1:C:112:ALA:O	1:C:115:ARG:NH1	2.39	0.55
1:C:931:THR:O	1:C:935:LEU:HD23	2.07	0.55
1:A:4749:GLU:HB3	1:A:4753:HIS:CE1	2.41	0.54
1:B:554:LEU:HD11	1:B:1593:PRO:HG3	1.88	0.54
1:B:3372:VAL:HG12	1:B:3398:PHE:HE1	1.71	0.54
1:D:1931:LEU:HD22	1:D:1935:VAL:HG11	1.88	0.54
1:D:2790:MET:SD	1:D:2797:PHE:HZ	2.30	0.54
1:D:2907:PRO:HB2	1:D:2910:THR:HG23	1.89	0.54
1:D:3635:CYS:HA	1:D:3638:MET:HG3	1.88	0.54
1:C:3166:TYR:OH	1:C:3203:VAL:HG11	2.06	0.54
1:A:2790:MET:SD	1:A:2797:PHE:HZ	2.30	0.54
1:A:2907:PRO:HB2	1:A:2910:THR:HG23	1.89	0.54
1:A:3635:CYS:HA	1:A:3638:MET:HG3	1.88	0.54
1:B:985:VAL:HG13	1:B:1039:LEU:HD22	1.89	0.54
1:B:2625:ARG:NH1	1:B:2629:ASP:CG	2.61	0.54
1:B:3114:LYS:HD2	1:B:3125:VAL:HG21	1.88	0.54
1:D:908:VAL:HG13	1:D:912:SER:HB2	1.89	0.54
1:D:931:THR:O	1:D:935:LEU:HD23	2.07	0.54
1:C:554:LEU:HD11	1:C:1593:PRO:HG3	1.88	0.54
1:C:908:VAL:O	1:C:963:ASN:CG	2.45	0.54
1:C:2790:MET:SD	1:C:2797:PHE:HZ	2.30	0.54
1:C:3372:VAL:HG12	1:C:3398:PHE:HE1	1.71	0.54
1:A:384:MET:SD	1:B:167:ASP:HA	2.47	0.54
1:A:1158:ASN:ND2	1:A:1182:ILE:O	2.40	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4651:THR:CB	1:A:4803:HIS:HE2	2.19	0.54
1:B:213:TYR:CE1	1:B:340:LYS:HG2	2.42	0.54
1:B:908:VAL:O	1:B:963:ASN:CG	2.45	0.54
1:B:1931:LEU:HD22	1:B:1935:VAL:HG11	1.88	0.54
1:D:67:PHE:HB3	1:D:109:LEU:HD22	1.89	0.54
1:D:2761:TYR:HB3	1:D:2765:LYS:HZ3	1.73	0.54
1:A:877:ASN:O	1:A:880:GLU:HG2	2.07	0.54
1:A:931:THR:O	1:A:935:LEU:HD23	2.07	0.54
1:B:2642:LYS:HG3	1:B:2698:MET:HE1	1.88	0.54
1:B:2907:PRO:HB2	1:B:2910:THR:HG23	1.89	0.54
1:B:4749:GLU:HB3	1:B:4753:HIS:CE1	2.41	0.54
1:D:1045:THR:HG22	1:D:1049:TYR:CE2	2.43	0.54
1:D:1157:GLU:OE1	1:D:1157:GLU:N	2.39	0.54
1:D:3442:PHE:HE1	1:D:3511:VAL:HG12	1.73	0.54
1:C:1045:THR:HG22	1:C:1049:TYR:CE2	2.43	0.54
1:C:2625:ARG:NH1	1:C:2629:ASP:CG	2.61	0.54
1:C:2907:PRO:HB2	1:C:2910:THR:HG23	1.89	0.54
1:B:125:ARG:HB3	1:B:125:ARG:HH11	1.71	0.54
1:B:931:THR:O	1:B:935:LEU:HD23	2.07	0.54
1:B:3527:PRO:HD2	1:B:3573:MET:SD	2.48	0.54
1:D:908:VAL:O	1:D:963:ASN:CG	2.45	0.54
1:C:3442:PHE:HE1	1:C:3511:VAL:HG12	1.73	0.54
1:A:3957:VAL:O	1:A:3961:VAL:HG23	2.08	0.54
1:B:1833:SER:HG	1:B:1836:PHE:HD1	1.54	0.54
1:B:4180:ARG:HD3	1:B:4192:ARG:HD3	1.89	0.54
1:D:2904:LEU:HD23	1:D:2904:LEU:O	2.07	0.54
1:C:3162:GLN:OE1	1:C:3218:VAL:HG23	2.08	0.54
1:C:3682:GLU:O	1:C:3685:GLU:N	2.41	0.54
1:B:1749:PRO:N	1:B:1758:ARG:HH12	2.06	0.54
1:B:2790:MET:SD	1:B:2797:PHE:HZ	2.30	0.54
1:D:985:VAL:HG13	1:D:1039:LEU:HD22	1.89	0.54
1:C:3527:PRO:HD2	1:C:3573:MET:SD	2.48	0.54
1:C:3727:ASP:O	1:C:3731:LYS:HG3	2.08	0.54
1:C:3852:LYS:O	1:C:3856:LEU:HG	2.06	0.54
1:A:1045:THR:HG22	1:A:1049:TYR:CE2	2.43	0.54
1:A:3197:LEU:C	1:A:3197:LEU:HD23	2.29	0.54
2:E:90:ILE:HG22	2:E:91:ILE:HG13	1.90	0.54
1:B:1045:THR:HG22	1:B:1049:TYR:CE2	2.43	0.54
1:B:1101:ARG:NH1	1:B:1115:LEU:O	2.40	0.54
1:B:3957:VAL:O	1:B:3961:VAL:HG23	2.08	0.54
1:D:3162:GLN:OE1	1:D:3218:VAL:HG23	2.08	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4945:ASP:OD1	1:D:4946:GLN:N	2.41	0.54
1:C:1101:ARG:NH1	1:C:1115:LEU:O	2.40	0.54
1:C:2904:LEU:O	1:C:2904:LEU:HD23	2.07	0.54
1:A:3379:LEU:HD12	1:A:3382:GLU:OE2	2.08	0.54
1:B:3442:PHE:HE1	1:B:3511:VAL:HG12	1.73	0.54
1:D:590:LEU:HG	1:D:599:VAL:HG11	1.90	0.54
1:A:590:LEU:HG	1:A:599:VAL:HG11	1.90	0.54
1:A:2642:LYS:NZ	1:A:2698:MET:SD	2.81	0.54
1:A:2904:LEU:HD23	1:A:2904:LEU:O	2.07	0.54
1:A:4945:ASP:OD1	1:A:4946:GLN:N	2.41	0.54
1:B:877:ASN:O	1:B:880:GLU:HG2	2.07	0.54
1:D:445:LEU:HD23	1:D:525:LEU:HD22	1.90	0.54
1:D:1101:ARG:NH1	1:D:1115:LEU:O	2.40	0.54
1:D:3786:CYS:SG	1:D:3831:SER:OG	2.65	0.54
1:C:877:ASN:O	1:C:880:GLU:HG2	2.07	0.54
1:A:2625:ARG:NH1	1:A:2629:ASP:CG	2.61	0.53
1:A:3527:PRO:HD2	1:A:3573:MET:SD	2.48	0.53
1:B:3727:ASP:O	1:B:3731:LYS:HG3	2.08	0.53
1:D:877:ASN:O	1:D:880:GLU:HG2	2.07	0.53
1:D:3166:TYR:OH	1:D:3203:VAL:HG11	2.06	0.53
1:D:3197:LEU:C	1:D:3197:LEU:HD23	2.29	0.53
1:C:590:LEU:HG	1:C:599:VAL:HG11	1.90	0.53
1:C:2294:ASP:O	1:C:2298:VAL:HG23	2.08	0.53
1:C:2638:LYS:HE2	1:C:2694:GLU:OE1	2.08	0.53
1:A:906:CYS:HB2	1:A:913:LEU:CB	2.38	0.53
1:A:1461:ASP:OD2	1:A:1468:LYS:NZ	2.37	0.53
1:B:2627:VAL:HA	1:B:2678:LEU:HD13	1.91	0.53
1:B:2638:LYS:HE2	1:B:2694:GLU:OE1	2.08	0.53
1:B:3379:LEU:HD12	1:B:3382:GLU:OE2	2.08	0.53
1:B:3682:GLU:O	1:B:3685:GLU:N	2.41	0.53
1:D:2294:ASP:O	1:D:2298:VAL:HG23	2.08	0.53
1:D:3727:ASP:O	1:D:3731:LYS:HG3	2.08	0.53
1:C:67:PHE:HB3	1:C:109:LEU:HD22	1.89	0.53
1:C:1749:PRO:N	1:C:1758:ARG:HH12	2.06	0.53
1:C:3379:LEU:HD12	1:C:3382:GLU:OE2	2.08	0.53
1:A:1749:PRO:N	1:A:1758:ARG:HH12	2.06	0.53
1:A:2882:TYR:O	1:A:2885:THR:HG22	2.09	0.53
2:G:80:TYR:CE2	2:G:80:TYR:CD1	2.38	0.53
1:B:437:PRO:O	1:B:441:VAL:HG23	2.09	0.53
1:B:3197:LEU:HD23	1:B:3197:LEU:C	2.29	0.53
1:D:1158:ASN:ND2	1:D:1182:ILE:O	2.40	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:672:VAL:O	1:C:680:THR:OG1	2.08	0.53
1:C:1833:SER:HG	1:C:1836:PHE:HD1	1.56	0.53
1:C:4945:ASP:OD1	1:C:4946:GLN:N	2.41	0.53
1:A:908:VAL:HG13	1:A:912:SER:HB2	1.89	0.53
1:A:1999:ARG:HH12	1:A:3636:PHE:CA	2.18	0.53
1:A:2294:ASP:O	1:A:2298:VAL:HG23	2.08	0.53
1:A:3727:ASP:O	1:A:3731:LYS:HG3	2.08	0.53
2:F:4:VAL:HG22	2:F:74:LEU:CD2	2.39	0.53
1:B:906:CYS:HB2	1:B:913:LEU:CB	2.38	0.53
1:B:1066:GLN:O	1:B:1071:ARG:NE	2.42	0.53
1:B:1999:ARG:HH12	1:B:3636:PHE:CA	2.18	0.53
1:B:2642:LYS:NZ	1:B:2698:MET:SD	2.81	0.53
1:B:2782:ASP:N	1:B:2782:ASP:OD1	2.38	0.53
1:D:906:CYS:HB2	1:D:913:LEU:CB	2.38	0.53
1:D:1123:VAL:HG23	1:D:1132:TRP:HB2	1.91	0.53
1:D:2882:TYR:O	1:D:2885:THR:HG22	2.09	0.53
1:C:2627:VAL:HA	1:C:2678:LEU:HD13	1.91	0.53
1:C:2761:TYR:HB3	1:C:2765:LYS:HZ3	1.73	0.53
1:C:2882:TYR:O	1:C:2885:THR:HG22	2.09	0.53
1:C:4180:ARG:HD3	1:C:4192:ARG:HD3	1.89	0.53
1:A:1066:GLN:O	1:A:1071:ARG:NE	2.42	0.53
1:A:3442:PHE:HE1	1:A:3511:VAL:HG12	1.73	0.53
2:G:4:VAL:HG22	2:G:74:LEU:CD2	2.39	0.53
1:B:1987:SER:O	1:B:1991:THR:OG1	2.08	0.53
1:D:69:LEU:HD23	1:D:109:LEU:CD2	2.39	0.53
1:D:553:ARG:NE	1:D:555:GLU:OE2	2.38	0.53
1:D:2642:LYS:NZ	1:D:2698:MET:SD	2.81	0.53
1:D:2969:ILE:O	1:D:2973:PHE:CD1	2.62	0.53
1:C:286:THR:HG21	1:C:481:GLU:OE1	2.09	0.53
1:C:437:PRO:O	1:C:441:VAL:HG23	2.09	0.53
1:C:906:CYS:HB2	1:C:913:LEU:CB	2.38	0.53
1:A:445:LEU:HD23	1:A:525:LEU:HD22	1.91	0.53
1:A:1287:LEU:HD11	1:A:1599:MET:HE3	1.90	0.53
1:A:3682:GLU:O	1:A:3685:GLU:N	2.41	0.53
1:D:2625:ARG:NH1	1:D:2629:ASP:CG	2.61	0.53
1:C:3957:VAL:O	1:C:3961:VAL:HG23	2.08	0.53
1:C:4222:VAL:HG11	1:C:4950:VAL:HA	1.90	0.53
1:A:56:GLN:O	1:A:281:ARG:NH2	2.42	0.53
1:A:652:ARG:NH1	1:A:750:LEU:O	2.42	0.53
1:A:1833:SER:HG	1:A:1836:PHE:HD1	1.56	0.53
1:A:2974:ILE:O	1:A:2978:GLU:OE1	2.27	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3372:VAL:HG12	1:B:3398:PHE:CE1	2.44	0.53
1:B:4945:ASP:OD1	1:B:4946:GLN:N	2.41	0.53
1:D:112:ALA:O	1:D:115:ARG:NH1	2.38	0.53
1:D:437:PRO:O	1:D:441:VAL:HG23	2.09	0.53
1:D:3527:PRO:HD2	1:D:3573:MET:SD	2.48	0.53
1:D:3682:GLU:O	1:D:3685:GLU:N	2.41	0.53
1:C:652:ARG:NH1	1:C:750:LEU:O	2.42	0.53
1:C:1123:VAL:HG23	1:C:1132:TRP:HB2	1.91	0.53
1:C:2874:MET:HE1	1:C:2939:ARG:CB	2.37	0.53
1:A:2969:ILE:O	1:A:2973:PHE:CD1	2.62	0.53
2:H:4:VAL:HG22	2:H:74:LEU:CD2	2.39	0.53
2:H:90:ILE:HG22	2:H:91:ILE:HG13	1.90	0.53
2:G:90:ILE:HG22	2:G:91:ILE:HG13	1.90	0.53
2:F:90:ILE:HD11	1:B:1684:ALA:HA	1.91	0.53
1:B:2294:ASP:O	1:B:2298:VAL:HG23	2.08	0.53
1:D:652:ARG:NH1	1:D:750:LEU:O	2.42	0.53
1:C:961:MET:HE3	1:C:963:ASN:CB	2.17	0.53
1:C:3239:MET:CG	1:C:3239:MET:HA	2.35	0.53
1:C:3755:GLU:O	1:C:3759:GLU:OE1	2.27	0.53
1:C:4998:LYS:HB3	1:C:5003:HIS:CE1	2.44	0.53
1:A:437:PRO:O	1:A:441:VAL:HG23	2.09	0.53
1:A:3755:GLU:O	1:A:3759:GLU:OE1	2.27	0.53
1:B:286:THR:HG21	1:B:481:GLU:OE1	2.09	0.53
1:B:590:LEU:HG	1:B:599:VAL:HG11	1.90	0.53
1:B:1123:VAL:HG23	1:B:1132:TRP:HB2	1.91	0.53
1:B:2233:CYS:HA	1:B:2237:CYS:SG	2.49	0.53
1:B:2874:MET:HE1	1:B:2939:ARG:HB2	1.90	0.53
1:B:4222:VAL:HG11	1:B:4950:VAL:HA	1.90	0.53
1:D:1287:LEU:HD11	1:D:1599:MET:HE3	1.90	0.53
1:D:1749:PRO:N	1:D:1758:ARG:HH12	2.06	0.53
1:D:3379:LEU:HD12	1:D:3382:GLU:OE2	2.08	0.53
1:C:445:LEU:HD23	1:C:525:LEU:HD22	1.91	0.53
1:C:908:VAL:HG13	1:C:912:SER:HB2	1.89	0.53
1:C:985:VAL:HG13	1:C:1039:LEU:HD22	1.89	0.53
1:C:1287:LEU:HD11	1:C:1599:MET:HE3	1.91	0.53
1:C:2642:LYS:NZ	1:C:2698:MET:SD	2.81	0.53
1:C:3197:LEU:HD23	1:C:3197:LEU:C	2.29	0.53
1:C:4205:TRP:CE3	1:C:4989:MET:HE2	2.44	0.53
1:A:3239:MET:CG	1:A:3239:MET:HA	2.34	0.53
1:A:4222:VAL:HG11	1:A:4950:VAL:HA	1.90	0.53
1:B:213:TYR:CE1	1:B:340:LYS:CG	2.92	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1855:GLY:O	1:B:1859:VAL:HG23	2.09	0.53
1:B:3162:GLN:OE1	1:B:3218:VAL:HG23	2.08	0.53
1:D:286:THR:HG21	1:D:481:GLU:OE1	2.09	0.53
1:D:2974:ILE:O	1:D:2978:GLU:OE1	2.27	0.53
1:C:633:LEU:HD13	1:C:1639:LEU:HD21	1.91	0.53
1:C:887:ILE:CG2	1:C:959:TYR:OH	2.57	0.53
1:C:1066:GLN:O	1:C:1071:ARG:NE	2.42	0.53
1:C:1749:PRO:C	1:C:1758:ARG:NH2	2.45	0.53
1:C:2969:ILE:O	1:C:2973:PHE:CD1	2.62	0.53
1:A:803:LEU:HD12	1:A:804:PRO:HD2	1.91	0.52
1:A:3162:GLN:OE1	1:A:3218:VAL:HG23	2.08	0.52
1:A:4984:ASN:OD1	1:A:4986:ALA:HB3	2.09	0.52
1:B:633:LEU:HD13	1:B:1639:LEU:HD21	1.91	0.52
1:B:2974:ILE:O	1:B:2978:GLU:OE1	2.27	0.52
1:D:803:LEU:HD12	1:D:804:PRO:HD2	1.91	0.52
1:D:4222:VAL:HG11	1:D:4950:VAL:HA	1.90	0.52
1:C:69:LEU:HD23	1:C:109:LEU:CD2	2.39	0.52
1:C:236:ALA:O	1:C:237:ASP:OD2	2.28	0.52
1:C:1855:GLY:O	1:C:1859:VAL:HG23	2.09	0.52
1:C:1926:LEU:HD23	1:C:1929:MET:SD	2.49	0.52
1:A:213:TYR:CE1	1:A:340:LYS:CG	2.92	0.52
1:A:228:ASP:OD2	1:B:155:LYS:HE2	2.08	0.52
1:A:985:VAL:HG13	1:A:1039:LEU:HD22	1.89	0.52
1:A:4180:ARG:HD3	1:A:4192:ARG:HD3	1.89	0.52
1:B:56:GLN:O	1:B:281:ARG:NH2	2.42	0.52
1:B:445:LEU:HD23	1:B:525:LEU:HD22	1.91	0.52
1:D:236:ALA:O	1:D:237:ASP:OD2	2.28	0.52
1:D:1066:GLN:O	1:D:1071:ARG:NE	2.42	0.52
1:D:1999:ARG:HH12	1:D:3636:PHE:CA	2.18	0.52
1:D:2233:CYS:HA	1:D:2237:CYS:SG	2.50	0.52
1:D:3957:VAL:O	1:D:3961:VAL:HG23	2.08	0.52
1:D:4180:ARG:HD3	1:D:4192:ARG:HD3	1.89	0.52
1:A:236:ALA:O	1:A:237:ASP:OD2	2.27	0.52
1:A:973:SER:O	1:A:976:ARG:NH1	2.43	0.52
1:A:1855:GLY:O	1:A:1859:VAL:HG23	2.09	0.52
1:A:3372:VAL:HG12	1:A:3398:PHE:CE1	2.44	0.52
2:E:4:VAL:HG22	2:E:74:LEU:CD2	2.39	0.52
1:B:69:LEU:HD23	1:B:109:LEU:CD2	2.39	0.52
1:B:1287:LEU:HD11	1:B:1599:MET:HE3	1.91	0.52
1:B:2929:PHE:CA	1:B:2932:MET:HE1	2.27	0.52
1:D:973:SER:O	1:D:976:ARG:NH1	2.42	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:633:LEU:HD13	1:A:1639:LEU:HD21	1.91	0.52
1:A:1926:LEU:HD23	1:A:1929:MET:SD	2.49	0.52
1:B:3621:HIS:O	1:B:3621:HIS:ND1	2.38	0.52
1:D:1855:GLY:O	1:D:1859:VAL:HG23	2.09	0.52
1:D:4984:ASN:OD1	1:D:4986:ALA:HB3	2.09	0.52
1:C:803:LEU:HD12	1:C:804:PRO:HD2	1.91	0.52
1:A:1123:VAL:HG23	1:A:1132:TRP:HB2	1.91	0.52
1:B:688:LEU:HD12	1:B:776:LEU:O	2.10	0.52
1:D:688:LEU:HD12	1:D:776:LEU:O	2.10	0.52
1:D:1999:ARG:HH12	1:D:3636:PHE:HD1	1.58	0.52
1:D:3755:GLU:O	1:D:3759:GLU:OE1	2.27	0.52
1:A:887:ILE:CG2	1:A:959:TYR:OH	2.58	0.52
1:A:1812:LEU:HD11	1:A:1858:ASP:HB3	1.92	0.52
1:A:2233:CYS:HA	1:A:2237:CYS:SG	2.49	0.52
1:A:2626:LEU:HD22	1:A:2640:PRO:HB3	1.92	0.52
1:A:4205:TRP:CE3	1:A:4989:MET:HE2	2.44	0.52
2:F:90:ILE:HG22	2:F:91:ILE:HG13	1.90	0.52
1:B:236:ALA:O	1:B:237:ASP:OD2	2.28	0.52
1:B:3514:LEU:HD21	1:B:3602:VAL:HG13	1.92	0.52
1:D:213:TYR:CE1	1:D:340:LYS:CG	2.92	0.52
1:D:633:LEU:HD13	1:D:1639:LEU:HD21	1.91	0.52
1:D:887:ILE:CG2	1:D:959:TYR:OH	2.58	0.52
1:D:4118:ASP:OD1	1:D:4122:MET:N	2.42	0.52
1:C:973:SER:O	1:C:976:ARG:NH1	2.43	0.52
1:A:2627:VAL:HA	1:A:2678:LEU:HD13	1.90	0.52
1:A:4998:LYS:HB3	1:A:5003:HIS:CE1	2.44	0.52
1:B:652:ARG:NH1	1:B:750:LEU:O	2.42	0.52
1:B:803:LEU:HD12	1:B:804:PRO:HD2	1.91	0.52
1:B:973:SER:O	1:B:976:ARG:NH1	2.43	0.52
1:B:1926:LEU:HD23	1:B:1929:MET:SD	2.49	0.52
1:B:2969:ILE:O	1:B:2973:PHE:CD1	2.62	0.52
1:B:4677:LEU:HD23	1:B:4711:PHE:CZ	2.45	0.52
1:D:56:GLN:O	1:D:281:ARG:NH2	2.42	0.52
1:D:1812:LEU:HD11	1:D:1858:ASP:HB3	1.92	0.52
1:D:2627:VAL:HA	1:D:2678:LEU:HD13	1.90	0.52
1:D:2638:LYS:HE2	1:D:2694:GLU:OE1	2.08	0.52
1:D:3236:VAL:O	1:D:3239:MET:HG3	2.10	0.52
1:D:4241:THR:O	1:D:4245:MET:HG3	2.10	0.52
1:D:4661:TYR:OH	1:D:4786:ASP:OD2	2.26	0.52
1:C:56:GLN:O	1:C:281:ARG:NH2	2.42	0.52
1:A:69:LEU:HD23	1:A:109:LEU:CD2	2.39	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3007:ASN:O	1:A:3011:THR:OG1	2.15	0.52
1:B:3755:GLU:O	1:B:3759:GLU:OE1	2.27	0.52
1:B:4984:ASN:OD1	1:B:4986:ALA:HB3	2.09	0.52
1:D:10:GLU:OE1	1:D:10:GLU:O	2.28	0.52
1:D:3532:LEU:HA	1:D:3535:LEU:HD12	1.92	0.52
1:C:688:LEU:HD12	1:C:776:LEU:O	2.10	0.52
1:C:2874:MET:HE1	1:C:2939:ARG:H	1.74	0.52
1:C:3372:VAL:HG12	1:C:3398:PHE:CE1	2.44	0.52
1:C:3607:GLU:O	1:C:3610:GLU:HG3	2.10	0.52
1:A:663:TYR:OH	1:A:665:GLU:OE2	2.27	0.52
1:B:1461:ASP:OD2	1:B:1468:LYS:NZ	2.37	0.52
1:D:1065:ASN:OD1	1:D:1066:GLN:N	2.43	0.52
1:D:1926:LEU:HD23	1:D:1929:MET:SD	2.49	0.52
1:D:2874:MET:HE1	1:D:2939:ARG:H	1.74	0.52
1:D:4677:LEU:HD23	1:D:4711:PHE:CZ	2.45	0.52
1:C:213:TYR:CE1	1:C:340:LYS:CG	2.92	0.52
1:C:3236:VAL:HA	1:C:3239:MET:HG2	1.90	0.52
1:C:4677:LEU:HD23	1:C:4711:PHE:CZ	2.45	0.52
1:C:4902:GLU:O	1:C:4913:ARG:CZ	2.58	0.52
1:A:10:GLU:OE1	1:A:10:GLU:O	2.28	0.52
1:A:286:THR:HG21	1:A:481:GLU:OE1	2.09	0.52
1:B:887:ILE:CG2	1:B:959:TYR:OH	2.58	0.52
1:B:1065:ASN:OD1	1:B:1066:GLN:N	2.43	0.52
1:B:2233:CYS:HB3	1:B:2237:CYS:HG	1.62	0.52
1:B:4705:VAL:HG13	1:B:4711:PHE:HE1	1.75	0.52
1:D:2625:ARG:HH12	1:D:2629:ASP:CG	2.12	0.52
1:D:3621:HIS:O	1:D:3622:LYS:HG2	2.10	0.52
1:C:2974:ILE:O	1:C:2978:GLU:OE1	2.27	0.52
1:C:3532:LEU:HA	1:C:3535:LEU:HD12	1.92	0.52
1:C:4984:ASN:OD1	1:C:4986:ALA:HB3	2.09	0.52
1:A:688:LEU:HD12	1:A:776:LEU:O	2.10	0.51
1:A:3589:PRO:O	1:A:3593:VAL:HG13	2.11	0.51
1:B:2882:TYR:O	1:B:2885:THR:HG22	2.09	0.51
1:B:4998:LYS:HB3	1:B:5003:HIS:CE1	2.44	0.51
1:D:4045:VAL:O	1:D:4049:VAL:HG23	2.11	0.51
1:C:2233:CYS:HA	1:C:2237:CYS:SG	2.49	0.51
1:A:1065:ASN:OD1	1:A:1066:GLN:N	2.43	0.51
1:A:4241:THR:O	1:A:4245:MET:HG3	2.10	0.51
1:B:3607:GLU:O	1:B:3610:GLU:HG3	2.10	0.51
1:D:3372:VAL:HG12	1:D:3398:PHE:CE1	2.44	0.51
1:C:3169:LEU:HD12	1:C:3194:LEU:HD11	1.93	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3621:HIS:O	1:C:3622:LYS:HG2	2.10	0.51
1:A:3598:GLU:O	1:A:3602:VAL:HG23	2.11	0.51
1:A:4705:VAL:HG13	1:A:4711:PHE:HE1	1.75	0.51
1:C:4548:ARG:O	1:C:4552:LEU:HD13	2.11	0.51
1:A:3621:HIS:O	1:A:3622:LYS:HG2	2.10	0.51
1:D:2292:GLU:O	1:D:2296:GLU:OE1	2.29	0.51
1:D:3589:PRO:O	1:D:3593:VAL:HG13	2.11	0.51
1:D:4902:GLU:O	1:D:4913:ARG:CZ	2.58	0.51
1:C:1812:LEU:HD11	1:C:1858:ASP:HB3	1.92	0.51
1:C:2006:ILE:HD11	1:C:3641:LEU:HD11	1.93	0.51
1:C:3514:LEU:HD21	1:C:3602:VAL:HG13	1.92	0.51
1:A:2970:SER:O	1:A:2974:ILE:HG23	2.11	0.51
1:B:3598:GLU:O	1:B:3602:VAL:HG23	2.11	0.51
1:B:4548:ARG:O	1:B:4552:LEU:HD13	2.11	0.51
1:D:3183:VAL:O	1:D:3187:ARG:HG3	2.11	0.51
1:D:4548:ARG:O	1:D:4552:LEU:HD13	2.11	0.51
1:D:4998:LYS:HB3	1:D:5003:HIS:CE1	2.44	0.51
1:A:3372:VAL:CG1	1:A:3398:PHE:HE1	2.24	0.51
1:B:1812:LEU:HD11	1:B:1858:ASP:HB3	1.92	0.51
1:B:2572:THR:O	1:B:2618:MET:HE2	2.11	0.51
1:D:2006:ILE:HD11	1:D:3641:LEU:HD11	1.93	0.51
1:C:2626:LEU:HD22	1:C:2640:PRO:HB3	1.92	0.51
1:C:2970:SER:O	1:C:2974:ILE:HG23	2.11	0.51
1:C:4705:VAL:HG13	1:C:4711:PHE:HE1	1.75	0.51
1:A:3183:VAL:O	1:A:3187:ARG:HG3	2.11	0.51
2:G:6:THR:HG23	2:G:6:THR:O	2.11	0.51
1:B:646:PRO:HB2	1:B:648:ILE:CD1	2.41	0.51
1:B:2292:GLU:O	1:B:2296:GLU:OE1	2.29	0.51
1:B:3532:LEU:HA	1:B:3535:LEU:HD12	1.92	0.51
1:B:3621:HIS:O	1:B:3622:LYS:HG2	2.10	0.51
1:D:3169:LEU:HD12	1:D:3194:LEU:HD11	1.93	0.51
1:D:3607:GLU:O	1:D:3610:GLU:HG3	2.10	0.51
1:C:2798:SER:OG	1:C:2801:ASP:OD2	2.26	0.51
1:C:4241:THR:O	1:C:4245:MET:HG3	2.10	0.51
1:A:2520:HIS:O	1:A:2524:VAL:HG22	2.11	0.51
1:A:3554:GLN:OE1	1:A:3593:VAL:HG11	2.11	0.51
1:A:3607:GLU:O	1:A:3610:GLU:HG3	2.10	0.51
1:A:4548:ARG:O	1:A:4552:LEU:HD13	2.11	0.51
1:A:4677:LEU:HD23	1:A:4711:PHE:CZ	2.45	0.51
1:A:4902:GLU:O	1:A:4913:ARG:CZ	2.58	0.51
2:G:90:ILE:HD11	1:C:1684:ALA:HA	1.92	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:228:ASP:OD2	1:C:155:LYS:HE2	2.11	0.51
1:B:3169:LEU:HD12	1:B:3194:LEU:HD11	1.93	0.51
1:B:3554:GLN:OE1	1:B:3593:VAL:HG11	2.11	0.51
1:B:3625:SER:O	1:B:3629:ARG:HG2	2.11	0.51
1:B:4902:GLU:O	1:B:4913:ARG:CZ	2.58	0.51
1:D:646:PRO:HB2	1:D:648:ILE:CD1	2.41	0.51
1:D:2626:LEU:HD22	1:D:2640:PRO:HB3	1.92	0.51
1:D:3996:PHE:HD1	1:D:4016:LEU:HD11	1.76	0.51
1:C:646:PRO:HB2	1:C:648:ILE:CD1	2.41	0.51
1:C:3372:VAL:CG1	1:C:3398:PHE:HE1	2.24	0.51
1:A:1101:ARG:NH1	1:A:1115:LEU:O	2.40	0.51
1:A:3169:LEU:HD12	1:A:3194:LEU:HD11	1.93	0.51
1:A:3621:HIS:O	1:A:3621:HIS:ND1	2.38	0.51
1:B:10:GLU:OE1	1:B:10:GLU:O	2.28	0.51
1:B:4241:THR:O	1:B:4245:MET:HG3	2.10	0.51
1:D:3299:GLY:O	1:D:3300:ALA:HB2	2.11	0.51
1:D:3514:LEU:HD21	1:D:3602:VAL:HG13	1.92	0.51
1:D:4651:THR:OG1	1:D:4803:HIS:NE2	2.05	0.51
1:C:1065:ASN:OD1	1:C:1066:GLN:N	2.43	0.51
1:C:1999:ARG:HH12	1:C:3636:PHE:CA	2.18	0.51
1:C:3589:PRO:O	1:C:3593:VAL:HG13	2.11	0.51
1:C:3996:PHE:HD1	1:C:4016:LEU:HD11	1.76	0.51
1:A:1962:ALA:O	1:A:1966:VAL:HG23	2.11	0.51
1:A:2107:GLN:O	1:A:3694:LYS:NZ	2.36	0.51
1:B:2626:LEU:HD22	1:B:2640:PRO:HB3	1.92	0.51
1:B:3589:PRO:O	1:B:3593:VAL:HG13	2.11	0.51
1:C:10:GLU:OE1	1:C:10:GLU:O	2.28	0.51
1:C:3598:GLU:O	1:C:3602:VAL:HG23	2.11	0.51
1:A:3996:PHE:HD1	1:A:4016:LEU:HD11	1.76	0.50
1:A:4098:ASP:OD1	1:A:4099:SER:N	2.44	0.50
1:A:4678:ALA:HB1	1:A:4720:VAL:HG21	1.93	0.50
1:A:4961:CYS:SG	1:A:4983:HIS:CE1	3.04	0.50
1:B:4232:GLU:OE1	1:B:5017:ARG:NH2	2.44	0.50
1:B:4583:SER:OG	1:B:4585:SER:O	2.27	0.50
1:D:2874:MET:HE1	1:D:2939:ARG:CB	2.39	0.50
1:D:4209:GLN:H	1:D:4209:GLN:CD	2.14	0.50
1:C:2292:GLU:O	1:C:2296:GLU:OE1	2.29	0.50
1:C:4045:VAL:O	1:C:4049:VAL:HG23	2.11	0.50
1:A:646:PRO:HB2	1:A:648:ILE:CD1	2.41	0.50
1:A:3532:LEU:HA	1:A:3535:LEU:HD12	1.92	0.50
1:A:4209:GLN:H	1:A:4209:GLN:CD	2.14	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:6:THR:HG23	2:F:6:THR:O	2.10	0.50
1:D:4232:GLU:OE1	1:D:5017:ARG:NH2	2.45	0.50
1:C:717:ASP:OD1	1:C:720:HIS:N	2.40	0.50
1:C:1962:ALA:O	1:C:1966:VAL:HG23	2.11	0.50
1:A:2625:ARG:HH12	1:A:2629:ASP:CG	2.12	0.50
1:A:2638:LYS:HE2	1:A:2694:GLU:OE1	2.08	0.50
1:A:3372:VAL:HG11	1:A:3398:PHE:CE1	2.47	0.50
1:A:4045:VAL:O	1:A:4049:VAL:HG23	2.11	0.50
1:B:820:ARG:CZ	1:B:820:ARG:HB3	2.24	0.50
1:B:4118:ASP:OD1	1:B:4122:MET:N	2.42	0.50
1:D:4649:LEU:O	1:D:4653:VAL:HG23	2.11	0.50
1:C:2520:HIS:O	1:C:2524:VAL:HG22	2.11	0.50
1:A:3514:LEU:HD21	1:A:3602:VAL:HG13	1.92	0.50
1:B:384:MET:SD	1:C:167:ASP:HA	2.52	0.50
1:B:2006:ILE:HD11	1:B:3641:LEU:HD11	1.93	0.50
1:B:2520:HIS:O	1:B:2524:VAL:HG22	2.11	0.50
1:B:3468:SER:O	1:B:3472:ALA:HB2	2.12	0.50
1:B:4661:TYR:OH	1:B:4786:ASP:OD2	2.27	0.50
1:B:4678:ALA:HB1	1:B:4720:VAL:HG21	1.93	0.50
1:D:2675:THR:HG23	1:D:2706:ILE:HG23	1.94	0.50
1:C:3372:VAL:HG11	1:C:3398:PHE:CE1	2.47	0.50
1:C:4678:ALA:HB1	1:C:4720:VAL:HG21	1.93	0.50
1:C:4961:CYS:SG	1:C:4983:HIS:CE1	3.04	0.50
1:A:796:ARG:NH1	1:A:1619:ARG:HH22	2.10	0.50
1:A:880:GLU:HG3	1:A:881:LEU:N	2.27	0.50
1:A:2006:ILE:HD11	1:A:3641:LEU:HD11	1.93	0.50
1:A:2572:THR:CG2	1:A:2579:VAL:CG2	2.83	0.50
1:B:2929:PHE:CA	1:B:2932:MET:SD	2.98	0.50
1:B:4045:VAL:O	1:B:4049:VAL:HG23	2.11	0.50
1:B:4098:ASP:OD1	1:B:4099:SER:N	2.44	0.50
1:B:4989:MET:SD	1:B:4993:MET:SD	3.10	0.50
1:D:323:LEU:HD23	1:D:323:LEU:H	1.77	0.50
1:D:880:GLU:HG3	1:D:881:LEU:N	2.27	0.50
1:D:3468:SER:O	1:D:3472:ALA:HB2	2.12	0.50
1:D:3598:GLU:O	1:D:3602:VAL:HG23	2.11	0.50
1:C:612:VAL:HG13	1:C:2167:ILE:O	2.12	0.50
1:C:3183:VAL:O	1:C:3187:ARG:HG3	2.11	0.50
1:C:4649:LEU:O	1:C:4653:VAL:HG23	2.11	0.50
1:A:2675:THR:HG23	1:A:2706:ILE:HG23	1.94	0.50
1:A:3468:SER:O	1:A:3472:ALA:HB2	2.12	0.50
1:A:3625:SER:O	1:A:3629:ARG:HG2	2.11	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:H:6:THR:HG23	2:H:6:THR:O	2.10	0.50
1:B:796:ARG:NH1	1:B:1619:ARG:HH22	2.10	0.50
1:B:2874:MET:HE1	1:B:2939:ARG:H	1.77	0.50
1:D:167:ASP:OD1	1:C:384:MET:CE	2.60	0.50
1:D:2970:SER:O	1:D:2974:ILE:HG23	2.11	0.50
1:C:323:LEU:HD23	1:C:323:LEU:H	1.77	0.50
1:C:3299:GLY:O	1:C:3300:ALA:HB2	2.11	0.50
1:C:3554:GLN:OE1	1:C:3593:VAL:HG11	2.11	0.50
1:C:4209:GLN:H	1:C:4209:GLN:CD	2.14	0.50
1:A:72:SER:O	1:A:99:ARG:NH1	2.45	0.50
1:A:2292:GLU:O	1:A:2296:GLU:OE1	2.29	0.50
1:A:2572:THR:O	1:A:2618:MET:HE2	2.12	0.50
1:A:2969:ILE:O	1:A:2972:GLU:HG3	2.11	0.50
1:A:4232:GLU:OE1	1:A:5017:ARG:NH2	2.44	0.50
1:B:781:VAL:HG13	1:B:791:PHE:CZ	2.47	0.50
1:B:1962:ALA:O	1:B:1966:VAL:HG23	2.12	0.50
1:B:2524:VAL:HG23	1:B:2525:GLY:N	2.27	0.50
1:B:3169:LEU:HD21	1:B:3205:PHE:CD2	2.47	0.50
1:B:3372:VAL:HG11	1:B:3398:PHE:CE1	2.47	0.50
1:B:4209:GLN:CD	1:B:4209:GLN:H	2.14	0.50
1:B:4649:LEU:O	1:B:4653:VAL:HG23	2.11	0.50
1:B:4961:CYS:SG	1:B:4983:HIS:CE1	3.04	0.50
1:D:1043:VAL:HA	1:D:1046:LEU:HD12	1.94	0.50
1:D:1962:ALA:O	1:D:1966:VAL:HG23	2.11	0.50
1:D:3554:GLN:OE1	1:D:3593:VAL:HG11	2.11	0.50
1:D:4678:ALA:HB1	1:D:4720:VAL:HG21	1.93	0.50
1:D:4989:MET:SD	1:D:4993:MET:SD	3.10	0.50
1:C:2470:ILE:HG22	1:C:2525:GLY:HA3	1.93	0.50
1:C:4098:ASP:OD1	1:C:4099:SER:N	2.44	0.50
1:A:612:VAL:HG13	1:A:2167:ILE:O	2.12	0.50
1:A:781:VAL:HG13	1:A:791:PHE:CZ	2.47	0.50
1:A:898:ASP:HB2	1:A:901:LYS:HG3	1.94	0.50
1:A:2524:VAL:HG23	1:A:2525:GLY:N	2.27	0.50
1:A:3299:GLY:O	1:A:3300:ALA:HB2	2.11	0.50
1:A:4649:LEU:O	1:A:4653:VAL:HG23	2.11	0.50
1:A:4989:MET:SD	1:A:4993:MET:SD	3.10	0.50
2:E:6:THR:HG23	2:E:6:THR:O	2.10	0.50
1:B:384:MET:CE	1:C:167:ASP:OD1	2.60	0.50
1:B:612:VAL:HG13	1:B:2167:ILE:O	2.12	0.50
1:B:880:GLU:HG3	1:B:881:LEU:N	2.27	0.50
1:B:2970:SER:O	1:B:2974:ILE:HG23	2.11	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4060:LYS:O	1:B:4064:MET:HG3	2.12	0.50
1:D:781:VAL:HG13	1:D:791:PHE:CZ	2.47	0.50
1:D:2226:PRO:CB	1:D:2267:MET:HE3	2.41	0.50
1:D:3372:VAL:HG11	1:D:3398:PHE:CE1	2.46	0.50
1:C:3169:LEU:CD1	1:C:3194:LEU:HD11	2.41	0.50
1:C:3468:SER:O	1:C:3472:ALA:HB2	2.12	0.50
1:C:4989:MET:SD	1:C:4993:MET:SD	3.10	0.50
1:A:323:LEU:HD23	1:A:323:LEU:H	1.77	0.50
1:A:659:TYR:O	1:A:662:TRP:NE1	2.44	0.50
1:A:688:LEU:HD11	1:A:775:GLY:CA	2.42	0.50
1:B:323:LEU:H	1:B:323:LEU:HD23	1.77	0.50
1:B:3372:VAL:CG1	1:B:3398:PHE:HE1	2.24	0.50
1:B:3786:CYS:SG	1:B:3831:SER:OG	2.65	0.50
1:D:1748:PHE:C	1:D:1758:ARG:HH12	2.16	0.50
1:D:2135:LEU:HD21	1:D:3663:LEU:HD21	1.94	0.50
1:D:4098:ASP:OD1	1:D:4099:SER:N	2.44	0.50
1:C:781:VAL:HG13	1:C:791:PHE:CZ	2.47	0.50
1:C:1868:PRO:O	1:C:1872:THR:OG1	2.28	0.50
1:A:3835:LEU:HD22	1:A:3880:PHE:CZ	2.47	0.49
1:B:2107:GLN:O	1:B:3694:LYS:NZ	2.36	0.49
1:B:2923:ALA:O	1:B:2927:LEU:HD23	2.12	0.49
1:D:72:SER:O	1:D:99:ARG:NH1	2.45	0.49
1:D:612:VAL:HG13	1:D:2167:ILE:O	2.12	0.49
1:D:796:ARG:NH1	1:D:1619:ARG:HH22	2.10	0.49
1:D:898:ASP:HB2	1:D:901:LYS:HG3	1.94	0.49
1:D:2969:ILE:O	1:D:2972:GLU:HG3	2.11	0.49
1:D:3169:LEU:CD1	1:D:3194:LEU:HD11	2.42	0.49
1:D:3239:MET:CG	1:D:3239:MET:HA	2.34	0.49
1:D:3625:SER:O	1:D:3629:ARG:HG2	2.11	0.49
1:C:898:ASP:HB2	1:C:901:LYS:HG3	1.94	0.49
1:C:4565:LEU:O	1:C:4569:LEU:HD13	2.12	0.49
1:B:3183:VAL:O	1:B:3187:ARG:HG3	2.11	0.49
1:B:3996:PHE:HD1	1:B:4016:LEU:HD11	1.76	0.49
1:B:4835:LYS:O	1:B:4839:MET:HG2	2.12	0.49
1:D:155:LYS:HE2	1:C:228:ASP:OD2	2.11	0.49
1:D:2520:HIS:O	1:D:2524:VAL:HG22	2.11	0.49
1:D:4961:CYS:SG	1:D:4983:HIS:CE1	3.04	0.49
1:C:2675:THR:HG23	1:C:2706:ILE:HG23	1.94	0.49
1:C:2969:ILE:O	1:C:2972:GLU:HG3	2.11	0.49
1:C:3835:LEU:HD22	1:C:3880:PHE:CZ	2.47	0.49
1:C:4060:LYS:O	1:C:4064:MET:HG3	2.12	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4232:GLU:OE1	1:C:5017:ARG:NH2	2.44	0.49
1:A:554:LEU:CD1	1:A:1593:PRO:HG3	2.42	0.49
1:A:913:LEU:O	1:A:918:ARG:NH2	2.45	0.49
1:A:2135:LEU:HD21	1:A:3663:LEU:HD21	1.94	0.49
1:A:3169:LEU:CD1	1:A:3194:LEU:HD11	2.42	0.49
1:A:3435:PHE:CD2	1:A:3524:MET:HE1	2.47	0.49
1:A:4705:VAL:HG22	1:A:4711:PHE:HD1	1.78	0.49
1:B:667:MET:SD	1:B:790:ARG:NH2	2.85	0.49
1:B:898:ASP:HB2	1:B:901:LYS:HG3	1.94	0.49
1:B:2969:ILE:O	1:B:2972:GLU:HG3	2.11	0.49
1:B:3169:LEU:CD1	1:B:3194:LEU:HD11	2.42	0.49
1:B:3299:GLY:O	1:B:3300:ALA:HB2	2.11	0.49
1:B:4565:LEU:O	1:B:4569:LEU:HD13	2.12	0.49
1:B:4705:VAL:HG22	1:B:4711:PHE:HD1	1.78	0.49
1:D:4835:LYS:O	1:D:4839:MET:HG2	2.12	0.49
1:C:796:ARG:NH1	1:C:1619:ARG:HH22	2.10	0.49
1:C:1748:PHE:C	1:C:1758:ARG:HH12	2.16	0.49
1:C:2261:SER:O	1:C:2264:GLY:N	2.43	0.49
1:C:3169:LEU:HD21	1:C:3205:PHE:CD2	2.47	0.49
1:C:3625:SER:O	1:C:3629:ARG:HG2	2.11	0.49
1:A:2470:ILE:HG22	1:A:2525:GLY:HA3	1.93	0.49
1:A:2788:HIS:ND1	1:A:2790:MET:HB3	2.27	0.49
1:A:3230:LEU:HD12	1:A:3230:LEU:O	2.13	0.49
1:B:4976:GLU:HG2	1:B:4980:LEU:HD13	1.94	0.49
1:D:887:ILE:HD11	1:D:907:LEU:HD21	1.95	0.49
1:C:72:SER:O	1:C:99:ARG:NH1	2.45	0.49
1:C:2524:VAL:HG23	1:C:2525:GLY:N	2.27	0.49
1:C:2929:PHE:CA	1:C:2932:MET:HE1	2.30	0.49
1:A:1280:GLN:O	1:A:1281:ASN:CG	2.51	0.49
1:A:2751:LEU:HD23	1:A:2755:ILE:HD13	1.95	0.49
1:A:4060:LYS:O	1:A:4064:MET:HG3	2.12	0.49
1:B:3230:LEU:HD12	1:B:3230:LEU:O	2.13	0.49
1:D:688:LEU:HD11	1:D:775:GLY:CA	2.42	0.49
1:D:1726:SER:O	1:D:1729:SER:OG	2.24	0.49
1:D:1991:THR:O	1:D:1995:THR:HG23	2.13	0.49
1:D:3169:LEU:HD21	1:D:3205:PHE:CD2	2.47	0.49
1:C:880:GLU:HG3	1:C:881:LEU:N	2.27	0.49
1:C:2135:LEU:HD21	1:C:3663:LEU:HD21	1.94	0.49
1:C:4835:LYS:O	1:C:4839:MET:HG2	2.13	0.49
1:A:1991:THR:O	1:A:1995:THR:HG23	2.13	0.49
1:A:2693:GLN:HE22	1:A:2697:ARG:HH11	1.60	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2923:ALA:O	1:A:2927:LEU:HD23	2.12	0.49
1:A:2929:PHE:CA	1:A:2932:MET:SD	2.98	0.49
1:A:3786:CYS:SG	1:A:3831:SER:OG	2.65	0.49
1:B:554:LEU:CD1	1:B:1593:PRO:HG3	2.42	0.49
1:B:688:LEU:HD11	1:B:775:GLY:CA	2.42	0.49
1:B:2470:ILE:HG22	1:B:2525:GLY:HA3	1.93	0.49
1:B:2693:GLN:HE22	1:B:2697:ARG:HH1	1.60	0.49
1:B:2788:HIS:ND1	1:B:2790:MET:HB3	2.27	0.49
1:B:4732:PHE:CD2	1:B:4732:PHE:O	2.66	0.49
1:D:554:LEU:CD1	1:D:1593:PRO:HG3	2.42	0.49
1:D:667:MET:SD	1:D:790:ARG:NH2	2.86	0.49
1:D:3230:LEU:HD12	1:D:3230:LEU:O	2.12	0.49
1:D:4705:VAL:HG13	1:D:4711:PHE:HE1	1.75	0.49
1:C:554:LEU:CD1	1:C:1593:PRO:HG3	2.42	0.49
1:C:887:ILE:HD11	1:C:907:LEU:HD21	1.95	0.49
1:C:1043:VAL:HA	1:C:1046:LEU:HD12	1.94	0.49
1:C:2642:LYS:HG3	1:C:2698:MET:HE3	1.92	0.49
1:C:2751:LEU:HD23	1:C:2755:ILE:HD13	1.95	0.49
1:C:2923:ALA:O	1:C:2927:LEU:HD23	2.12	0.49
1:A:2744:ASN:OD1	1:A:2745:VAL:HG23	2.13	0.49
1:B:72:SER:O	1:B:99:ARG:NH1	2.45	0.49
1:B:296:ASP:OD1	1:B:297:GLN:OE1	2.31	0.49
1:B:1991:THR:O	1:B:1995:THR:HG23	2.13	0.49
1:B:2135:LEU:HD21	1:B:3663:LEU:HD21	1.94	0.49
1:B:4205:TRP:CE3	1:B:4989:MET:HE2	2.48	0.49
1:D:2744:ASN:OD1	1:D:2745:VAL:HG23	2.13	0.49
1:D:2923:ALA:O	1:D:2927:LEU:HD23	2.12	0.49
1:D:3835:LEU:HD22	1:D:3880:PHE:CZ	2.47	0.49
1:D:4976:GLU:HG2	1:D:4980:LEU:HD13	1.94	0.49
1:C:2929:PHE:CA	1:C:2932:MET:SD	2.98	0.49
1:C:3621:HIS:O	1:C:3621:HIS:ND1	2.38	0.49
1:A:3901:ASN:OD1	1:A:3904:ARG:NH1	2.39	0.49
1:A:4565:LEU:O	1:A:4569:LEU:HD13	2.12	0.49
1:B:979:PRO:O	1:B:983:THR:HG23	2.13	0.49
1:B:2476:ILE:HD12	1:B:2532:ALA:HB1	1.95	0.49
1:B:2751:LEU:HD23	1:B:2755:ILE:HD13	1.95	0.49
1:D:363:ASP:OD1	1:D:366:ALA:HB2	2.13	0.49
1:D:1280:GLN:O	1:D:1281:ASN:CG	2.51	0.49
1:C:913:LEU:O	1:C:918:ARG:NH2	2.45	0.49
1:C:979:PRO:O	1:C:983:THR:HG23	2.13	0.49
1:C:2476:ILE:HD12	1:C:2532:ALA:HB1	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4732:PHE:CD2	1:C:4732:PHE:O	2.66	0.49
1:A:2675:THR:HB	1:A:2710:LEU:HD21	1.95	0.49
1:B:3835:LEU:HD22	1:B:3880:PHE:CZ	2.48	0.49
1:D:4565:LEU:O	1:D:4569:LEU:HD13	2.12	0.49
1:C:1991:THR:O	1:C:1995:THR:HG23	2.12	0.49
1:C:3969:ILE:CG2	1:C:3980:LEU:HD12	2.43	0.49
1:C:4651:THR:OG1	1:C:4803:HIS:NE2	2.05	0.49
1:C:4705:VAL:HG22	1:C:4711:PHE:HD1	1.78	0.49
1:A:887:ILE:HD11	1:A:907:LEU:HD21	1.95	0.49
1:A:1743[A]:ARG:NH2	1:A:1967:ASP:OD2	2.40	0.49
1:B:360:ALA:O	1:B:361:ALA:HB3	2.13	0.49
1:B:1043:VAL:HA	1:B:1046:LEU:HD12	1.94	0.49
1:B:1810:LYS:O	1:B:1814:MET:HG3	2.13	0.49
1:B:2675:THR:HG23	1:B:2706:ILE:HG23	1.94	0.49
1:D:659:TYR:O	1:D:662:TRP:NE1	2.44	0.49
1:D:1833:SER:HG	1:D:1836:PHE:HD1	1.59	0.49
1:D:2470:ILE:HG22	1:D:2525:GLY:HA3	1.93	0.49
1:D:2524:VAL:HG23	1:D:2525:GLY:N	2.27	0.49
1:D:4705:VAL:HG22	1:D:4711:PHE:HD1	1.78	0.49
1:C:659:TYR:O	1:C:662:TRP:NE1	2.44	0.49
1:C:2974:ILE:HG13	1:C:2975:ALA:N	2.28	0.49
1:C:3590:GLU:O	1:C:3594:ARG:HG3	2.13	0.49
1:C:4976:GLU:HG2	1:C:4980:LEU:HD13	1.94	0.49
1:A:861:ILE:HD13	1:A:930:LYS:CD	2.43	0.48
1:A:1748:PHE:C	1:A:1758:ARG:HH12	2.16	0.48
1:D:2751:LEU:HD23	1:D:2755:ILE:HD13	1.95	0.48
1:D:3969:ILE:CG2	1:D:3980:LEU:HD12	2.43	0.48
1:C:667:MET:SD	1:C:790:ARG:NH2	2.85	0.48
1:C:4000:MET:SD	1:C:4020:GLN:NE2	2.79	0.48
1:A:3941:ASP:OD2	1:A:3941:ASP:N	2.46	0.48
1:A:4640:GLU:HB3	1:A:4641:PRO:HD3	1.95	0.48
1:B:861:ILE:HD13	1:B:930:LYS:CD	2.43	0.48
1:B:1999:ARG:HH12	1:B:3636:PHE:HD1	1.58	0.48
1:B:4640:GLU:HB3	1:B:4641:PRO:HD3	1.95	0.48
1:D:913:LEU:O	1:D:918:ARG:NH2	2.45	0.48
1:D:2974:ILE:HG13	1:D:2975:ALA:N	2.28	0.48
1:C:552:ASP:OD1	1:C:553:ARG:N	2.47	0.48
1:C:4834:GLY:O	1:C:4838:VAL:HG23	2.13	0.48
1:A:360:ALA:O	1:A:361:ALA:HB3	2.13	0.48
1:A:667:MET:SD	1:A:790:ARG:NH2	2.85	0.48
1:A:898:ASP:HB2	1:A:901:LYS:CG	2.43	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1043:VAL:HA	1:A:1046:LEU:HD12	1.94	0.48
1:A:1810:LYS:O	1:A:1814:MET:HG3	2.13	0.48
1:A:3442:PHE:CE1	1:A:3511:VAL:HG12	2.48	0.48
1:A:4732:PHE:O	1:A:4732:PHE:CD2	2.66	0.48
1:B:1280:GLN:O	1:B:1281:ASN:CG	2.51	0.48
1:B:3590:GLU:O	1:B:3594:ARG:HG3	2.13	0.48
1:B:4000:MET:SD	1:B:4020:GLN:NE2	2.79	0.48
1:B:5014:TYR:HD1	6:B:5304:PNX:HAA3	1.76	0.48
1:D:167:ASP:HA	1:C:384:MET:SD	2.53	0.48
1:D:663:TYR:OH	1:D:665:GLU:OE2	2.27	0.48
1:D:1810:LYS:O	1:D:1814:MET:HG3	2.13	0.48
1:D:4764:LEU:O	1:D:4768:LEU:HG	2.13	0.48
1:C:1810:LYS:O	1:C:1814:MET:HG3	2.13	0.48
1:C:4118:ASP:OD1	1:C:4122:MET:N	2.42	0.48
1:C:4717:ASP:OD2	1:C:4723:LYS:NZ	2.35	0.48
1:A:363:ASP:OD1	1:A:366:ALA:HB2	2.13	0.48
1:A:552:ASP:OD1	1:A:553:ARG:N	2.47	0.48
1:A:2642:LYS:HG3	1:A:2698:MET:HE1	1.91	0.48
1:A:2874:MET:HE1	1:A:2939:ARG:H	1.79	0.48
1:A:3169:LEU:HD21	1:A:3205:PHE:CD2	2.47	0.48
1:A:3892:CYS:HG	1:A:3899:PHE:HD2	1.61	0.48
1:A:4835:LYS:O	1:A:4839:MET:HG2	2.12	0.48
1:B:1748:PHE:C	1:B:1758:ARG:HH12	2.16	0.48
1:B:4929:LEU:C	1:B:4933:GLN:OE1	2.52	0.48
1:D:4060:LYS:O	1:D:4064:MET:HG3	2.12	0.48
1:D:4834:GLY:O	1:D:4838:VAL:HG23	2.13	0.48
1:D:5014:TYR:HD1	6:D:5304:PNX:HAA3	1.76	0.48
1:C:2742:THR:HG23	1:C:2814:LYS:CE	2.41	0.48
1:C:3442:PHE:CE1	1:C:3511:VAL:HG12	2.48	0.48
1:A:2190:VAL:HG12	1:A:2198:MET:HE3	1.96	0.48
1:A:3197:LEU:HD21	1:A:3201:MET:CE	2.44	0.48
1:A:3735:LEU:HG	1:A:3735:LEU:O	2.14	0.48
1:A:4976:GLU:HG2	1:A:4980:LEU:HD13	1.94	0.48
1:B:363:ASP:OD1	1:B:366:ALA:HB2	2.13	0.48
1:B:552:ASP:OD1	1:B:553:ARG:N	2.46	0.48
1:B:2223:ILE:HG23	1:B:2223:ILE:O	2.13	0.48
1:B:2534:ALA:HB1	1:B:2588:ARG:HD3	1.96	0.48
1:B:2974:ILE:HG13	1:B:2975:ALA:N	2.28	0.48
1:D:552:ASP:OD1	1:D:553:ARG:N	2.47	0.48
1:D:898:ASP:HB2	1:D:901:LYS:CG	2.43	0.48
1:D:2742:THR:HG23	1:D:2814:LYS:CE	2.41	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3735:LEU:HG	1:D:3735:LEU:O	2.14	0.48
1:C:360:ALA:O	1:C:361:ALA:HB3	2.13	0.48
1:C:898:ASP:HB2	1:C:901:LYS:CG	2.43	0.48
1:C:2223:ILE:O	1:C:2223:ILE:HG23	2.13	0.48
1:C:3230:LEU:O	1:C:3230:LEU:HD12	2.13	0.48
1:A:3321:ARG:O	1:A:3324:VAL:HG22	2.14	0.48
1:A:4661:TYR:OH	1:A:4786:ASP:OD2	2.27	0.48
1:B:717:ASP:OD1	1:B:720:HIS:N	2.40	0.48
1:B:2675:THR:HB	1:B:2710:LEU:HD21	1.95	0.48
1:B:3442:PHE:CE1	1:B:3511:VAL:HG12	2.48	0.48
1:D:360:ALA:O	1:D:361:ALA:HB3	2.13	0.48
1:D:2693:GLN:HE22	1:D:2697:ARG:HH11	1.60	0.48
1:C:2355:ARG:HG3	1:C:2355:ARG:HH11	1.79	0.48
1:C:2742:THR:CG2	1:C:2814:LYS:HE3	2.42	0.48
1:C:2744:ASN:OD1	1:C:2745:VAL:HG23	2.13	0.48
1:C:3197:LEU:HD21	1:C:3201:MET:CE	2.44	0.48
1:A:167:ASP:OD1	1:D:384:MET:SD	2.72	0.48
1:A:953:THR:O	1:A:968:ALA:HB1	2.14	0.48
1:A:1726:SER:O	1:A:1729:SER:OG	2.24	0.48
1:A:4929:LEU:C	1:A:4933:GLN:OE1	2.52	0.48
1:B:648:ILE:HG23	1:B:814:ALA:HB3	1.96	0.48
1:B:861:ILE:HG21	1:B:930:LYS:HD2	1.96	0.48
1:B:3471:THR:HG22	1:C:1170:MET:CE	2.44	0.48
1:B:4834:GLY:O	1:B:4838:VAL:HG23	2.13	0.48
1:D:296:ASP:OD1	1:D:297:GLN:OE1	2.31	0.48
1:D:861:ILE:HD13	1:D:930:LYS:CD	2.43	0.48
1:D:1743[A]:ARG:NH2	1:D:1967:ASP:OD2	2.39	0.48
1:D:2642:LYS:HG3	1:D:2698:MET:HE1	1.93	0.48
1:D:3941:ASP:OD2	1:D:3941:ASP:N	2.47	0.48
1:D:4911:LEU:O	1:D:4914:VAL:HG22	2.14	0.48
1:C:3007:ASN:O	1:C:3011:THR:OG1	2.15	0.48
1:C:3901:ASN:OD1	1:C:3904:ARG:NH1	2.39	0.48
1:C:5014:TYR:HD1	6:C:5304:PNX:HAA3	1.75	0.48
1:A:574:VAL:O	1:A:578:ILE:HG12	2.14	0.48
1:A:3651:ASN:O	1:A:3655:GLU:OE1	2.32	0.48
1:A:4764:LEU:O	1:A:4768:LEU:HG	2.13	0.48
1:A:4834:GLY:O	1:A:4838:VAL:HG23	2.13	0.48
1:B:913:LEU:O	1:B:918:ARG:NH2	2.45	0.48
1:B:978:THR:O	1:B:982:THR:HG23	2.14	0.48
1:B:3321:ARG:O	1:B:3324:VAL:HG22	2.14	0.48
1:B:4764:LEU:O	1:B:4768:LEU:HG	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4911:LEU:O	1:B:4914:VAL:HG22	2.14	0.48
1:D:590:LEU:HD21	1:D:599:VAL:HB	1.96	0.48
1:D:2098:VAL:HG11	1:D:2123:LEU:HG	1.96	0.48
1:D:2122:SER:HA	1:D:3721:LEU:HD11	1.96	0.48
1:D:3365:LEU:HD11	1:D:3408:LEU:HD12	1.95	0.48
1:D:3442:PHE:CE1	1:D:3511:VAL:HG12	2.48	0.48
1:D:3651:ASN:O	1:D:3655:GLU:OE1	2.32	0.48
1:C:2098:VAL:HG11	1:C:2123:LEU:HG	1.96	0.48
1:C:2204:HIS:NE2	1:C:2205:GLU:OE2	2.47	0.48
1:C:2534:ALA:HB1	1:C:2588:ARG:HD3	1.96	0.48
1:C:2788:HIS:ND1	1:C:2790:MET:HB3	2.28	0.48
1:C:3786:CYS:SG	1:C:3831:SER:OG	2.65	0.48
1:C:4929:LEU:C	1:C:4933:GLN:OE1	2.52	0.48
1:A:34:LYS:C	1:A:35:LEU:HD12	2.35	0.48
1:A:590:LEU:HD21	1:A:599:VAL:HB	1.96	0.48
1:A:861:ILE:HG21	1:A:930:LYS:HD2	1.96	0.48
1:A:2098:VAL:HG11	1:A:2123:LEU:HG	1.96	0.48
1:A:2476:ILE:HG13	1:A:2536:LEU:HD11	1.96	0.48
1:A:3627:GLN:O	1:A:3630:ARG:HG2	2.14	0.48
1:B:659:TYR:O	1:B:662:TRP:NE1	2.44	0.48
1:B:1112:ASP:OD1	1:B:1113:VAL:N	2.47	0.48
1:B:2098:VAL:HG11	1:B:2123:LEU:HG	1.96	0.48
1:B:2476:ILE:HG13	1:B:2536:LEU:HD11	1.96	0.48
1:B:3659:ALA:HA	1:B:3663:LEU:HD12	1.96	0.48
1:B:3969:ILE:CG2	1:B:3980:LEU:HD12	2.43	0.48
1:D:574:VAL:O	1:D:578:ILE:HG12	2.14	0.48
1:D:919:ASN:O	1:D:923:GLN:OE1	2.32	0.48
1:D:2534:ALA:HB1	1:D:2588:ARG:HD3	1.96	0.48
1:D:2675:THR:HB	1:D:2710:LEU:HD21	1.94	0.48
1:D:2788:HIS:ND1	1:D:2790:MET:HB3	2.27	0.48
1:D:4732:PHE:O	1:D:4732:PHE:CD2	2.66	0.48
1:C:590:LEU:HD21	1:C:599:VAL:HB	1.96	0.48
1:C:2523:ASP:HB3	1:C:2578:MET:HE2	1.93	0.48
1:C:3735:LEU:O	1:C:3735:LEU:HG	2.14	0.48
1:A:919:ASN:O	1:A:923:GLN:OE1	2.32	0.48
1:A:979:PRO:O	1:A:983:THR:HG23	2.13	0.48
1:A:2668:SER:O	1:A:2669:GLU:HB3	2.14	0.48
1:A:2974:ILE:HG13	1:A:2975:ALA:N	2.28	0.48
1:A:3659:ALA:HA	1:A:3663:LEU:HD12	1.96	0.48
1:B:2587:TYR:OH	1:B:2629:ASP:OD2	2.24	0.48
1:D:979:PRO:O	1:D:983:THR:HG23	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2107:GLN:O	1:D:3694:LYS:NZ	2.36	0.48
1:D:4640:GLU:HB3	1:D:4641:PRO:HD3	1.96	0.48
1:C:648:ILE:HG23	1:C:814:ALA:HB3	1.96	0.48
1:C:953:THR:O	1:C:968:ALA:HB1	2.14	0.48
1:C:2625:ARG:HH12	1:C:2629:ASP:CG	2.12	0.48
1:C:4911:LEU:O	1:C:4914:VAL:HG22	2.14	0.48
1:A:438:ILE:HG23	1:A:518:ILE:HD11	1.96	0.47
1:A:899:ASP:O	1:A:902:ARG:NE	2.45	0.47
1:A:4067:LYS:O	1:A:4070:ASP:OD1	2.33	0.47
1:B:3346:VAL:HG11	1:B:3414:ARG:HB2	1.96	0.47
1:D:34:LYS:C	1:D:35:LEU:HD12	2.35	0.47
1:D:1170:MET:CE	1:C:3471:THR:HG22	2.44	0.47
1:D:2329:GLU:O	1:D:2333:ASP:OD1	2.32	0.47
1:D:3627:GLN:O	1:D:3630:ARG:HG2	2.14	0.47
1:C:69:LEU:HD23	1:C:109:LEU:HD21	1.96	0.47
1:C:233:ILE:HD12	1:C:242:ARG:HB3	1.96	0.47
1:C:2392:ARG:O	1:C:2418:LEU:HD12	2.14	0.47
1:C:2693:GLN:HE22	1:C:2697:ARG:HH11	1.60	0.47
1:C:3159:ASP:O	1:C:3163:VAL:HG23	2.14	0.47
1:C:3651:ASN:O	1:C:3655:GLU:OE1	2.32	0.47
1:C:3941:ASP:OD2	1:C:3941:ASP:N	2.46	0.47
1:A:296:ASP:OD1	1:A:297:GLN:OE1	2.31	0.47
1:A:2355:ARG:HH11	1:A:2355:ARG:HG3	1.79	0.47
1:A:2373:GLY:O	1:B:130:LYS:NZ	2.44	0.47
1:A:3365:LEU:HD11	1:A:3408:LEU:HD12	1.95	0.47
1:B:221:ARG:NH2	1:B:255:HIS:O	2.48	0.47
1:B:438:ILE:HG23	1:B:518:ILE:HD11	1.96	0.47
1:B:884:LEU:HD23	1:B:955:LEU:HD11	1.97	0.47
1:B:953:THR:O	1:B:968:ALA:HB1	2.14	0.47
1:B:2204:HIS:NE2	1:B:2205:GLU:OE2	2.47	0.47
1:B:2224:ARG:HE	1:B:2225:PHE:N	2.10	0.47
1:B:2355:ARG:HH11	1:B:2355:ARG:HG3	1.79	0.47
1:B:3197:LEU:HD21	1:B:3201:MET:CE	2.44	0.47
1:B:4067:LYS:O	1:B:4070:ASP:OD1	2.33	0.47
1:D:985:VAL:HG11	1:D:1040:CYS:HB3	1.96	0.47
1:D:2929:PHE:CA	1:D:2932:MET:SD	2.98	0.47
1:D:3197:LEU:HD21	1:D:3201:MET:CE	2.44	0.47
1:D:3346:VAL:HG11	1:D:3414:ARG:HB2	1.96	0.47
1:C:363:ASP:OD1	1:C:366:ALA:HB2	2.13	0.47
1:C:688:LEU:HD11	1:C:775:GLY:CA	2.42	0.47
1:C:978:THR:O	1:C:982:THR:HG23	2.14	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1999:ARG:HH12	1:C:3636:PHE:HD1	1.58	0.47
1:C:3835:LEU:HD22	1:C:3880:PHE:HZ	1.80	0.47
1:A:46:LEU:CD2	1:A:125:ARG:HE	2.27	0.47
1:A:4118:ASP:OD1	1:A:4122:MET:N	2.42	0.47
1:B:899:ASP:O	1:B:902:ARG:NE	2.45	0.47
1:B:1749:PRO:C	1:B:1758:ARG:NH2	2.45	0.47
1:D:1112:ASP:OD1	1:D:1113:VAL:N	2.47	0.47
1:D:2204:HIS:NE2	1:D:2205:GLU:OE2	2.47	0.47
1:D:2519:LEU:HD13	1:D:2575:ARG:HG3	1.97	0.47
1:C:1280:GLN:O	1:C:1281:ASN:CG	2.51	0.47
1:C:2668:SER:O	1:C:2669:GLU:HB3	2.14	0.47
1:C:2675:THR:HB	1:C:2710:LEU:HD21	1.95	0.47
1:C:3398:PHE:CE2	1:C:3450:ASN:CB	2.97	0.47
1:A:648:ILE:HG23	1:A:814:ALA:HB3	1.96	0.47
1:A:978:THR:O	1:A:982:THR:HG23	2.14	0.47
1:A:1299:GLN:HG2	2:E:31:GLU:OE2	2.14	0.47
1:A:3159:ASP:O	1:A:3163:VAL:HG23	2.14	0.47
1:B:898:ASP:HB2	1:B:901:LYS:CG	2.43	0.47
1:B:3365:LEU:HD11	1:B:3408:LEU:HD12	1.95	0.47
1:B:3627:GLN:O	1:B:3630:ARG:HG2	2.14	0.47
1:B:4096:ALA:O	1:B:4100:GLN:HG2	2.14	0.47
1:D:380:GLN:OE1	1:D:380:GLN:N	2.45	0.47
1:D:3166:TYR:CZ	1:D:3203:VAL:HG11	2.50	0.47
1:D:3239:MET:CB	1:D:3239:MET:HG3	2.31	0.47
1:D:4929:LEU:C	1:D:4933:GLN:OE1	2.52	0.47
1:C:3627:GLN:O	1:C:3630:ARG:HG2	2.14	0.47
1:A:928:THR:O	1:A:931:THR:OG1	2.30	0.47
1:A:3346:VAL:HG11	1:A:3414:ARG:HB2	1.96	0.47
1:A:3733:CYS:HB2	1:A:3803:SER:OG	2.14	0.47
1:A:5014:TYR:HD1	6:A:5304:PNX:HAA3	1.76	0.47
2:G:2:VAL:CG2	2:G:58:GLY:HA2	2.45	0.47
1:B:3015:LEU:HD22	1:B:3025:LEU:HB3	1.96	0.47
1:B:3362:ILE:HG22	1:B:3437:MET:HB3	1.97	0.47
1:D:3362:ILE:HG22	1:D:3437:MET:HB3	1.97	0.47
1:C:1069:TRP:CE3	1:C:1602:PRO:HG2	2.50	0.47
1:C:2519:LEU:HD13	1:C:2575:ARG:HG3	1.97	0.47
1:C:4764:LEU:O	1:C:4768:LEU:HG	2.13	0.47
1:A:233:ILE:HD12	1:A:242:ARG:HB3	1.96	0.47
1:A:360:ALA:O	1:A:361:ALA:CB	2.63	0.47
1:A:384:MET:SD	1:B:167:ASP:N	2.87	0.47
1:A:2122:SER:HA	1:A:3721:LEU:HD11	1.96	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2204:HIS:NE2	1:A:2205:GLU:OE2	2.47	0.47
1:A:2476:ILE:HD12	1:A:2532:ALA:HB1	1.95	0.47
1:A:2534:ALA:HB1	1:A:2588:ARG:HD3	1.96	0.47
1:A:2742:THR:HG23	1:A:2814:LYS:CE	2.41	0.47
1:A:3362:ILE:HG22	1:A:3437:MET:HB3	1.97	0.47
1:A:3477:LYS:NZ	1:A:3479:ALA:HA	2.30	0.47
1:A:4911:LEU:O	1:A:4914:VAL:HG22	2.14	0.47
2:E:2:VAL:CG2	2:E:58:GLY:HA2	2.45	0.47
1:B:880:GLU:HB2	1:B:967:PRO:HG2	1.97	0.47
1:B:2519:LEU:HD13	1:B:2575:ARG:HG3	1.97	0.47
1:D:978:THR:O	1:D:982:THR:HG23	2.14	0.47
1:D:2223:ILE:HG23	1:D:2223:ILE:O	2.13	0.47
1:D:2392:ARG:O	1:D:2418:LEU:HD12	2.14	0.47
1:D:2742:THR:CG2	1:D:2814:LYS:HE3	2.42	0.47
1:D:3159:ASP:O	1:D:3163:VAL:HG23	2.14	0.47
1:D:3321:ARG:O	1:D:3324:VAL:HG22	2.14	0.47
1:D:3590:GLU:O	1:D:3594:ARG:HG3	2.13	0.47
1:D:3621:HIS:O	1:D:3621:HIS:ND1	2.38	0.47
1:C:574:VAL:O	1:C:578:ILE:HG12	2.14	0.47
1:C:2122:SER:HA	1:C:3721:LEU:HD11	1.96	0.47
1:C:2329:GLU:O	1:C:2333:ASP:OD1	2.32	0.47
1:C:3321:ARG:O	1:C:3324:VAL:HG22	2.14	0.47
1:C:3346:VAL:HG11	1:C:3414:ARG:HB2	1.96	0.47
1:C:3362:ILE:HG22	1:C:3437:MET:HB3	1.97	0.47
1:C:3733:CYS:HB2	1:C:3803:SER:OG	2.14	0.47
1:A:36:CYS:SG	1:A:52:THR:HG21	2.55	0.47
1:A:1112:ASP:OD1	1:A:1113:VAL:N	2.47	0.47
1:A:1427:ILE:O	1:A:1430:THR:HG22	2.15	0.47
1:A:2014:ASP:O	1:A:2015:GLU:HB3	2.15	0.47
1:A:2519:LEU:HD13	1:A:2575:ARG:HG3	1.97	0.47
1:A:2638:LYS:O	1:A:2698:MET:CE	2.63	0.47
1:A:2790:MET:SD	1:A:2797:PHE:CZ	3.08	0.47
1:A:3234:ASN:O	1:A:3234:ASN:OD1	2.33	0.47
1:A:3590:GLU:O	1:A:3594:ARG:HG3	2.13	0.47
1:A:3969:ILE:CG2	1:A:3980:LEU:HD12	2.43	0.47
1:B:233:ILE:HD12	1:B:242:ARG:HB3	1.96	0.47
1:B:360:ALA:O	1:B:361:ALA:CB	2.63	0.47
1:B:1069:TRP:CE3	1:B:1602:PRO:HG2	2.49	0.47
1:B:2011:HIS:O	1:B:2011:HIS:ND1	2.47	0.47
1:B:2261:SER:O	1:B:2264:GLY:N	2.43	0.47
1:B:2744:ASN:OD1	1:B:2745:VAL:HG23	2.13	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2790:MET:SD	1:B:2797:PHE:CZ	3.08	0.47
1:B:3166:TYR:CZ	1:B:3203:VAL:HG11	2.50	0.47
1:B:3379:LEU:HD11	1:B:3387:ALA:O	2.15	0.47
1:B:3477:LYS:NZ	1:B:3479:ALA:HA	2.30	0.47
1:D:233:ILE:HD12	1:D:242:ARG:HB3	1.96	0.47
1:D:717:ASP:OD1	1:D:720:HIS:N	2.40	0.47
1:D:953:THR:O	1:D:968:ALA:HB1	2.14	0.47
1:D:1427:ILE:O	1:D:1430:THR:HG22	2.15	0.47
1:D:2014:ASP:O	1:D:2015:GLU:HB3	2.15	0.47
1:D:2476:ILE:HD12	1:D:2532:ALA:HB1	1.95	0.47
1:D:2523:ASP:HB3	1:D:2578:MET:HE2	1.91	0.47
1:D:2668:SER:O	1:D:2669:GLU:HB3	2.14	0.47
1:D:3015:LEU:HD22	1:D:3025:LEU:HB3	1.96	0.47
1:D:3659:ALA:HA	1:D:3663:LEU:HD12	1.96	0.47
1:D:3835:LEU:HD22	1:D:3880:PHE:HZ	1.80	0.47
1:D:4583:SER:OG	1:D:4585:SER:O	2.27	0.47
1:C:221:ARG:NH2	1:C:255:HIS:O	2.48	0.47
1:C:296:ASP:OD1	1:C:297:GLN:OE1	2.31	0.47
1:C:380:GLN:OE1	1:C:380:GLN:N	2.45	0.47
1:C:861:ILE:HD13	1:C:930:LYS:CD	2.43	0.47
1:C:880:GLU:HB2	1:C:967:PRO:HG2	1.97	0.47
1:C:884:LEU:HD23	1:C:955:LEU:HD11	1.97	0.47
1:C:2014:ASP:O	1:C:2015:GLU:HB3	2.15	0.47
1:C:2224:ARG:HE	1:C:2225:PHE:N	2.10	0.47
1:C:3015:LEU:HD22	1:C:3025:LEU:HB3	1.96	0.47
1:C:3365:LEU:HD11	1:C:3408:LEU:HD12	1.95	0.47
1:C:4096:ALA:O	1:C:4100:GLN:HG2	2.14	0.47
1:C:4640:GLU:HB3	1:C:4641:PRO:HD3	1.96	0.47
1:A:2392:ARG:O	1:A:2418:LEU:HD12	2.14	0.47
1:A:3166:TYR:CZ	1:A:3203:VAL:HG11	2.50	0.47
1:A:3239:MET:CB	1:A:3239:MET:HG2	2.31	0.47
1:A:3379:LEU:HD11	1:A:3387:ALA:O	2.15	0.47
2:H:2:VAL:CG2	2:H:58:GLY:HA2	2.45	0.47
2:F:2:VAL:CG2	2:F:58:GLY:HA2	2.45	0.47
1:B:577:ILE:O	1:B:579:GLN:NE2	2.48	0.47
1:B:590:LEU:HD21	1:B:599:VAL:HB	1.96	0.47
1:B:887:ILE:HD11	1:B:907:LEU:HD21	1.95	0.47
1:B:985:VAL:HG11	1:B:1040:CYS:HB3	1.96	0.47
1:B:3234:ASN:OD1	1:B:3234:ASN:O	2.33	0.47
1:B:3835:LEU:HD22	1:B:3880:PHE:HZ	1.80	0.47
1:D:36:CYS:SG	1:D:52:THR:HG21	2.55	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:46:LEU:CD2	1:D:125:ARG:HE	2.27	0.47
1:D:648:ILE:HG23	1:D:814:ALA:HB3	1.96	0.47
1:D:3372:VAL:CG1	1:D:3398:PHE:HE1	2.24	0.47
1:C:4067:LYS:O	1:C:4070:ASP:OD1	2.33	0.47
1:A:985:VAL:HG11	1:A:1040:CYS:HB3	1.96	0.47
1:A:3398:PHE:CE2	1:A:3450:ASN:CB	2.97	0.47
1:B:69:LEU:HD23	1:B:109:LEU:HD21	1.96	0.47
1:B:919:ASN:O	1:B:923:GLN:OE1	2.32	0.47
1:B:2122:SER:HA	1:B:3721:LEU:HD11	1.96	0.47
1:B:2329:GLU:O	1:B:2333:ASP:OD1	2.32	0.47
1:B:2392:ARG:O	1:B:2418:LEU:HD12	2.14	0.47
1:B:3115:VAL:HG13	1:B:3116:SER:N	2.30	0.47
1:B:3239:MET:CB	1:B:3239:MET:HG2	2.31	0.47
1:B:3651:ASN:O	1:B:3655:GLU:OE1	2.32	0.47
1:D:69:LEU:HD23	1:D:109:LEU:HD21	1.96	0.47
1:D:438:ILE:HG23	1:D:518:ILE:HD11	1.96	0.47
1:D:2146:PRO:HA	1:D:2149:VAL:HG23	1.97	0.47
1:D:3100:SER:O	1:D:3104:GLU:OE1	2.33	0.47
1:D:3733:CYS:HB2	1:D:3803:SER:OG	2.14	0.47
1:D:4096:ALA:O	1:D:4100:GLN:HG2	2.14	0.47
1:C:34:LYS:C	1:C:35:LEU:HD12	2.35	0.47
1:C:985:VAL:HG11	1:C:1040:CYS:HB3	1.96	0.47
1:C:3166:TYR:CZ	1:C:3203:VAL:HG11	2.50	0.47
1:C:3182:TYR:O	1:C:3185:LYS:HG3	2.15	0.47
1:C:3659:ALA:HA	1:C:3663:LEU:HD12	1.96	0.47
1:A:577:ILE:O	1:A:579:GLN:NE2	2.48	0.47
1:A:2434:GLY:O	1:A:2508:ARG:NE	2.43	0.47
1:B:553:ARG:O	1:B:557:SER:OG	2.29	0.47
1:B:2190:VAL:HG12	1:B:2198:MET:HE3	1.97	0.47
1:B:2572:THR:HG23	1:B:2575:ARG:HB2	1.97	0.47
1:B:2638:LYS:O	1:B:2698:MET:CE	2.63	0.47
1:B:3159:ASP:O	1:B:3163:VAL:HG23	2.14	0.47
1:B:3733:CYS:HB2	1:B:3803:SER:OG	2.14	0.47
1:D:577:ILE:O	1:D:579:GLN:NE2	2.48	0.47
1:D:1749:PRO:C	1:D:1758:ARG:NH2	2.45	0.47
1:D:2476:ILE:HG13	1:D:2536:LEU:HD11	1.96	0.47
1:D:2572:THR:HG23	1:D:2575:ARG:HB2	1.97	0.47
1:D:2745:VAL:HG12	1:D:2746:ILE:N	2.30	0.47
1:D:2790:MET:SD	1:D:2797:PHE:CZ	3.08	0.47
1:D:4067:LYS:O	1:D:4070:ASP:OD1	2.33	0.47
1:C:919:ASN:O	1:C:923:GLN:OE1	2.32	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1570:LYS:O	1:C:1572:ILE:HD12	2.15	0.47
1:C:2146:PRO:HA	1:C:2149:VAL:HG23	1.97	0.47
1:C:2745:VAL:HG12	1:C:2746:ILE:N	2.30	0.47
1:C:3062:PRO:HA	1:C:3065:VAL:HG22	1.97	0.47
1:C:3100:SER:O	1:C:3104:GLU:OE1	2.33	0.47
1:A:1069:TRP:CE3	1:A:1602:PRO:HG2	2.50	0.46
1:A:2329:GLU:O	1:A:2333:ASP:OD1	2.32	0.46
1:A:3048:ALA:O	1:A:3053:ARG:NH2	2.49	0.46
1:A:4096:ALA:O	1:A:4100:GLN:HG2	2.14	0.46
1:B:574:VAL:O	1:B:578:ILE:HG12	2.14	0.46
1:B:2668:SER:O	1:B:2669:GLU:HB3	2.14	0.46
1:B:3157:ILE:HG22	1:B:3162:GLN:HG2	1.97	0.46
1:B:3735:LEU:HG	1:B:3735:LEU:O	2.14	0.46
1:B:4848:VAL:HG11	1:B:4887:MET:HG2	1.97	0.46
1:D:861:ILE:HG21	1:D:930:LYS:HD2	1.96	0.46
1:D:1115:LEU:HD13	1:D:1123:VAL:HG11	1.97	0.46
1:D:2190:VAL:HG12	1:D:2198:MET:HE3	1.96	0.46
1:D:2638:LYS:O	1:D:2698:MET:CE	2.63	0.46
1:D:3048:ALA:O	1:D:3053:ARG:NH2	2.49	0.46
1:D:3901:ASN:OD1	1:D:3904:ARG:NH1	2.39	0.46
1:C:1112:ASP:OD1	1:C:1113:VAL:N	2.47	0.46
1:C:2790:MET:SD	1:C:2797:PHE:CZ	3.08	0.46
1:C:3157:ILE:HG22	1:C:3162:GLN:HG2	1.97	0.46
1:A:2226:PRO:CB	1:A:2267:MET:HE3	2.44	0.46
1:A:2742:THR:CG2	1:A:2814:LYS:HE3	2.42	0.46
1:A:3015:LEU:HD22	1:A:3025:LEU:HB3	1.96	0.46
1:A:4747:SER:HA	1:A:4750:ILE:HG12	1.97	0.46
1:B:34:LYS:C	1:B:35:LEU:HD12	2.35	0.46
1:B:1115:LEU:HD13	1:B:1123:VAL:HG11	1.97	0.46
1:D:100:THR:HG21	1:D:162:LYS:HE2	1.98	0.46
1:D:884:LEU:HD23	1:D:955:LEU:HD11	1.97	0.46
1:D:1570:LYS:O	1:D:1572:ILE:HD12	2.15	0.46
1:C:46:LEU:CD2	1:C:125:ARG:HE	2.27	0.46
1:C:100:THR:HG21	1:C:162:LYS:HE2	1.98	0.46
1:C:577:ILE:O	1:C:579:GLN:NE2	2.48	0.46
1:C:3048:ALA:O	1:C:3053:ARG:NH2	2.49	0.46
1:A:961:MET:HE1	1:A:963:ASN:HB3	1.79	0.46
1:A:1115:LEU:HD13	1:A:1123:VAL:HG11	1.97	0.46
1:A:2223:ILE:HG23	1:A:2223:ILE:O	2.13	0.46
1:A:3062:PRO:HA	1:A:3065:VAL:HG22	1.97	0.46
1:A:4705:VAL:HG13	1:A:4711:PHE:CD1	2.51	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1554:VAL:HG21	1:B:1561:VAL:CG1	2.45	0.46
1:B:2014:ASP:O	1:B:2015:GLU:HB3	2.15	0.46
1:B:3100:SER:O	1:B:3104:GLU:OE1	2.33	0.46
1:D:574:VAL:HA	1:D:577:ILE:HG12	1.98	0.46
1:D:1554:VAL:HG21	1:D:1561:VAL:CG1	2.45	0.46
1:D:2355:ARG:HH11	1:D:2355:ARG:HG3	1.79	0.46
1:D:2912:THR:HG22	1:D:2914:LYS:H	1.81	0.46
1:D:3062:PRO:HA	1:D:3065:VAL:HG22	1.97	0.46
1:D:3157:ILE:HG22	1:D:3162:GLN:HG2	1.97	0.46
1:C:1115:LEU:HD13	1:C:1123:VAL:HG11	1.97	0.46
1:C:1427:ILE:O	1:C:1430:THR:HG22	2.15	0.46
1:C:2534:ALA:HB1	1:C:2588:ARG:CD	2.46	0.46
1:A:2745:VAL:HG12	1:A:2746:ILE:N	2.30	0.46
1:A:4690:GLU:O	1:A:4691:GLN:HB2	2.16	0.46
1:A:4732:PHE:HE2	1:A:4736:ARG:HG2	1.81	0.46
1:B:46:LEU:CD2	1:B:125:ARG:HE	2.27	0.46
1:B:2211:MET:O	1:B:2214:VAL:CG1	2.64	0.46
1:B:2434:GLY:O	1:B:2508:ARG:NE	2.43	0.46
1:B:2874:MET:HE1	1:B:2939:ARG:CB	2.44	0.46
1:B:3398:PHE:CE2	1:B:3450:ASN:CB	2.97	0.46
1:B:4767:TRP:CE3	1:B:4768:LEU:CD2	2.99	0.46
1:D:1069:TRP:CE3	1:D:1602:PRO:HG2	2.49	0.46
1:D:2767:ALA:O	1:D:2771:ILE:HG12	2.16	0.46
1:D:3379:LEU:HD11	1:D:3387:ALA:O	2.15	0.46
1:C:438:ILE:HG23	1:C:518:ILE:HD11	1.96	0.46
1:C:1456:ASP:OD1	1:C:1456:ASP:N	2.47	0.46
1:C:2170:MET:HG3	1:C:2225:PHE:CE2	2.50	0.46
1:C:2476:ILE:HG13	1:C:2536:LEU:HD11	1.96	0.46
1:C:3379:LEU:HD11	1:C:3387:ALA:O	2.15	0.46
1:C:4661:TYR:OH	1:C:4786:ASP:OD2	2.27	0.46
1:A:884:LEU:O	1:A:887:ILE:HB	2.16	0.46
1:A:921:ASN:HA	1:A:924:MET:SD	2.56	0.46
1:A:1973:GLN:NE2	1:A:2005:GLN:OE1	2.44	0.46
1:A:1999:ARG:HH12	1:A:3636:PHE:HD1	1.58	0.46
1:A:2011:HIS:CE1	1:A:2017:ASP:CG	2.84	0.46
1:A:3100:SER:O	1:A:3104:GLU:OE1	2.33	0.46
1:A:3182:TYR:O	1:A:3185:LYS:HG3	2.15	0.46
1:B:667:MET:SD	1:B:801:LYS:NZ	2.86	0.46
1:B:2170:MET:HG3	1:B:2225:PHE:CE2	2.50	0.46
1:B:2534:ALA:HB1	1:B:2588:ARG:CD	2.46	0.46
1:B:3048:ALA:O	1:B:3053:ARG:NH2	2.49	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3062:PRO:HA	1:B:3065:VAL:HG22	1.97	0.46
1:B:4651:THR:OG1	1:B:4803:HIS:NE2	2.05	0.46
1:D:2170:MET:HG3	1:D:2225:PHE:CE2	2.50	0.46
1:D:2233:CYS:SG	1:D:2275:VAL:CG2	3.04	0.46
1:D:2515:GLN:O	1:D:2519:LEU:HG	2.16	0.46
1:D:4848:VAL:HG11	1:D:4887:MET:HG2	1.96	0.46
1:C:574:VAL:HA	1:C:577:ILE:HG12	1.98	0.46
1:C:861:ILE:HG21	1:C:930:LYS:HD2	1.96	0.46
1:C:921:ASN:HA	1:C:924:MET:SD	2.56	0.46
1:C:2233:CYS:SG	1:C:2275:VAL:CG2	3.04	0.46
1:C:2515:GLN:O	1:C:2519:LEU:HG	2.16	0.46
1:C:2638:LYS:O	1:C:2698:MET:CE	2.63	0.46
1:C:4041:ALA:O	1:C:4045:VAL:HG23	2.16	0.46
1:C:4690:GLU:O	1:C:4691:GLN:HB2	2.16	0.46
1:A:100:THR:HG21	1:A:162:LYS:HE2	1.98	0.46
1:A:2233:CYS:SG	1:A:2275:VAL:CG2	3.04	0.46
1:A:2586:VAL:CG1	1:A:2607:LEU:HD13	2.46	0.46
1:A:2703:LEU:CD2	1:A:2706:ILE:HD12	2.46	0.46
1:A:3115:VAL:HG13	1:A:3116:SER:N	2.30	0.46
1:A:4583:SER:OG	1:A:4585:SER:O	2.27	0.46
1:A:4934:GLY:HA2	1:D:4937:ILE:HG12	1.97	0.46
1:B:884:LEU:O	1:B:887:ILE:HB	2.16	0.46
1:B:921:ASN:HA	1:B:924:MET:SD	2.56	0.46
1:B:2146:PRO:HA	1:B:2149:VAL:HG23	1.97	0.46
1:B:2742:THR:HG23	1:B:2814:LYS:CE	2.41	0.46
1:D:360:ALA:O	1:D:361:ALA:CB	2.63	0.46
1:D:2011:HIS:ND1	1:D:2011:HIS:O	2.47	0.46
1:D:3115:VAL:HG13	1:D:3116:SER:N	2.30	0.46
1:C:879:HIS:CE1	1:C:913:LEU:HD13	2.51	0.46
1:C:3234:ASN:OD1	1:C:3234:ASN:O	2.33	0.46
1:C:3339:ALA:O	1:C:3343:GLN:HG2	2.16	0.46
1:C:3477:LYS:NZ	1:C:3479:ALA:HA	2.30	0.46
1:A:880:GLU:HB2	1:A:967:PRO:HG2	1.97	0.46
1:A:884:LEU:HD23	1:A:955:LEU:HD11	1.97	0.46
1:A:3530:GLN:O	1:A:3534:MET:HG2	2.16	0.46
1:A:3835:LEU:HD22	1:A:3880:PHE:HZ	1.80	0.46
1:A:3991:GLY:O	1:A:3995:VAL:HG23	2.16	0.46
1:B:869:ARG:HD2	1:B:870:ILE:N	2.31	0.46
1:B:2233:CYS:SG	1:B:2275:VAL:CG2	3.04	0.46
1:B:2494:PHE:HD1	1:B:2498:HIS:HD1	1.64	0.46
1:B:2586:VAL:CG1	1:B:2607:LEU:HD13	2.46	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2703:LEU:CD2	1:B:2706:ILE:HD12	2.46	0.46
1:B:3901:ASN:OD1	1:B:3904:ARG:NH1	2.40	0.46
1:D:3234:ASN:OD1	1:D:3234:ASN:O	2.33	0.46
1:D:3530:GLN:O	1:D:3534:MET:HG2	2.16	0.46
1:C:884:LEU:O	1:C:887:ILE:HB	2.16	0.46
1:C:4176:PRO:O	1:C:4202:ARG:NH2	2.49	0.46
1:A:574:VAL:HA	1:A:577:ILE:HG12	1.98	0.46
1:A:2767:ALA:O	1:A:2771:ILE:HG12	2.16	0.46
1:A:3910:THR:HG23	1:A:3911:THR:HG23	1.98	0.46
1:A:4848:VAL:HG11	1:A:4887:MET:HG2	1.97	0.46
1:B:879:HIS:CE1	1:B:913:LEU:HD13	2.51	0.46
1:B:2745:VAL:HG12	1:B:2746:ILE:N	2.30	0.46
1:B:3339:ALA:O	1:B:3343:GLN:HG2	2.16	0.46
1:B:3910:THR:HG23	1:B:3911:THR:HG23	1.98	0.46
1:B:3991:GLY:O	1:B:3995:VAL:HG23	2.16	0.46
1:B:4717:ASP:OD2	1:B:4723:LYS:NZ	2.35	0.46
1:B:4732:PHE:HE2	1:B:4736:ARG:HG2	1.81	0.46
1:D:463:GLU:O	1:D:467:LYS:HG2	2.16	0.46
1:D:2198:MET:HB3	1:D:2203:MET:SD	2.56	0.46
1:D:2534:ALA:HB1	1:D:2588:ARG:CD	2.46	0.46
1:D:2661:TRP:HB3	1:D:2664:PHE:HD1	1.81	0.46
1:D:2908:TYR:OH	1:D:2916:LYS:HA	2.16	0.46
1:D:3910:THR:HG23	1:D:3911:THR:HG23	1.98	0.46
1:C:360:ALA:O	1:C:361:ALA:CB	2.63	0.46
1:C:2233:CYS:O	1:C:2235:PHE:N	2.49	0.46
1:C:2703:LEU:CD2	1:C:2706:ILE:HD12	2.46	0.46
1:C:2767:ALA:O	1:C:2771:ILE:HG12	2.16	0.46
1:A:384:MET:SD	1:B:167:ASP:OD1	2.74	0.46
1:A:1554:VAL:HG21	1:A:1561:VAL:CG1	2.45	0.46
1:A:2170:MET:HG3	1:A:2225:PHE:CE2	2.50	0.46
1:A:2224:ARG:HE	1:A:2225:PHE:N	2.10	0.46
1:A:2908:TYR:OH	1:A:2916:LYS:HA	2.16	0.46
1:B:36:CYS:SG	1:B:52:THR:HG21	2.55	0.46
1:B:2742:THR:CG2	1:B:2814:LYS:HE3	2.42	0.46
1:D:880:GLU:HB2	1:D:967:PRO:HG2	1.97	0.46
1:D:921:ASN:HA	1:D:924:MET:SD	2.56	0.46
1:D:3401:LEU:O	1:D:3405:LEU:HD13	2.16	0.46
1:D:4645:CYS:O	1:D:4649:LEU:HD13	2.16	0.46
1:D:4705:VAL:HG13	1:D:4711:PHE:CD1	2.51	0.46
1:C:2198:MET:HB3	1:C:2203:MET:SD	2.56	0.46
1:C:2586:VAL:CG1	1:C:2607:LEU:HD13	2.46	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1749:PRO:C	1:A:1758:ARG:NH2	2.45	0.46
1:A:4041:ALA:O	1:A:4045:VAL:HG23	2.16	0.46
1:B:4041:ALA:O	1:B:4045:VAL:HG23	2.16	0.46
1:B:4690:GLU:O	1:B:4691:GLN:HB2	2.16	0.46
1:D:2233:CYS:O	1:D:2235:PHE:N	2.49	0.46
1:D:2233:CYS:SG	1:D:2275:VAL:HG23	2.57	0.46
1:D:3182:TYR:O	1:D:3185:LYS:HG3	2.15	0.46
1:D:3339:ALA:O	1:D:3343:GLN:HG2	2.16	0.46
1:D:3398:PHE:CE2	1:D:3450:ASN:CB	2.98	0.46
1:C:2661:TRP:HB3	1:C:2664:PHE:HD1	1.81	0.46
1:C:2912:THR:HG22	1:C:2914:LYS:H	1.81	0.46
1:C:3401:LEU:O	1:C:3405:LEU:HD13	2.16	0.46
1:C:3910:THR:HG23	1:C:3911:THR:HG23	1.98	0.46
1:A:69:LEU:HD23	1:A:109:LEU:HD21	1.96	0.45
1:A:213:TYR:HA	1:A:339:ILE:O	2.16	0.45
1:A:2011:HIS:O	1:A:2011:HIS:ND1	2.47	0.45
1:A:2146:PRO:HA	1:A:2149:VAL:HG23	1.97	0.45
1:A:2198:MET:HB3	1:A:2203:MET:SD	2.56	0.45
1:A:2303:ALA:O	1:A:2307:LEU:HD13	2.17	0.45
1:A:2515:GLN:O	1:A:2519:LEU:HG	2.16	0.45
1:A:2782:ASP:OD2	1:A:2785:LEU:HD13	2.17	0.45
1:A:3336:LYS:O	1:A:3340:VAL:HG23	2.16	0.45
1:A:3339:ALA:O	1:A:3343:GLN:HG2	2.16	0.45
1:B:463:GLU:O	1:B:467:LYS:HG2	2.16	0.45
1:B:1427:ILE:O	1:B:1430:THR:HG22	2.15	0.45
1:B:2198:MET:HB3	1:B:2203:MET:SD	2.56	0.45
1:B:2515:GLN:O	1:B:2519:LEU:HG	2.16	0.45
1:B:2767:ALA:O	1:B:2771:ILE:HG12	2.16	0.45
1:B:2782:ASP:OD2	1:B:2785:LEU:HD13	2.17	0.45
1:B:2828:GLU:O	1:B:2829:GLY:O	2.34	0.45
1:B:2908:TYR:OH	1:B:2916:LYS:HA	2.16	0.45
1:B:2912:THR:HG22	1:B:2914:LYS:H	1.81	0.45
1:B:4176:PRO:O	1:B:4202:ARG:NH2	2.49	0.45
1:B:4705:VAL:HG13	1:B:4711:PHE:CD1	2.51	0.45
1:B:4942:GLU:O	1:B:4945:ASP:OD1	2.34	0.45
1:D:1929:MET:O	1:D:1930:LYS:HB3	2.16	0.45
1:D:2703:LEU:CD2	1:D:2706:ILE:HD12	2.46	0.45
1:D:3477:LYS:NZ	1:D:3479:ALA:HA	2.30	0.45
1:D:4041:ALA:O	1:D:4045:VAL:HG23	2.16	0.45
1:C:213:TYR:HA	1:C:339:ILE:O	2.16	0.45
1:C:869:ARG:HD2	1:C:870:ILE:N	2.31	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2494:PHE:HD1	1:C:2498:HIS:HD1	1.64	0.45
1:C:2572:THR:HG23	1:C:2575:ARG:HB2	1.97	0.45
1:C:3115:VAL:HG13	1:C:3116:SER:N	2.30	0.45
1:A:1570:LYS:O	1:A:1572:ILE:HD12	2.15	0.45
1:A:2482:ASP:C	1:A:2482:ASP:OD1	2.55	0.45
1:A:3899:PHE:CZ	1:A:3903:LEU:HD21	2.52	0.45
1:B:100:THR:HG21	1:B:162:LYS:HE2	1.98	0.45
1:B:1570:LYS:O	1:B:1572:ILE:HD12	2.15	0.45
1:B:2233:CYS:O	1:B:2235:PHE:N	2.49	0.45
1:B:2280:VAL:HG22	1:B:2280:VAL:O	2.16	0.45
1:B:4016:LEU:O	1:B:4020:GLN:HG3	2.16	0.45
1:B:4172:GLU:OE1	1:B:4175:ARG:NH1	2.50	0.45
1:D:820:ARG:HB3	1:D:820:ARG:CZ	2.24	0.45
1:D:4148:THR:HG21	1:D:4180:ARG:NH2	2.31	0.45
1:C:2011:HIS:O	1:C:2011:HIS:ND1	2.47	0.45
1:C:2928:LYS:O	1:C:2932:MET:HE3	2.17	0.45
1:C:3398:PHE:HE2	1:C:3450:ASN:CB	2.29	0.45
1:C:3530:GLN:O	1:C:3534:MET:HG2	2.16	0.45
1:C:4148:THR:HG21	1:C:4180:ARG:NH2	2.31	0.45
1:C:4172:GLU:OE1	1:C:4175:ARG:NH1	2.50	0.45
1:C:4732:PHE:HE2	1:C:4736:ARG:HG2	1.81	0.45
1:A:224:HIS:HA	1:A:388:LEU:HD23	1.98	0.45
1:A:260:TRP:CE3	1:A:282:ILE:HG22	2.51	0.45
1:A:463:GLU:O	1:A:467:LYS:HG2	2.16	0.45
1:A:1929:MET:O	1:A:1930:LYS:HB3	2.16	0.45
1:A:2280:VAL:O	1:A:2280:VAL:HG22	2.16	0.45
1:A:2534:ALA:HB1	1:A:2588:ARG:CD	2.46	0.45
1:A:3324:VAL:HA	1:A:3327:LEU:HG	1.99	0.45
1:A:4075:GLU:CD	1:B:4736:ARG:NH1	2.69	0.45
1:A:4942:GLU:O	1:A:4945:ASP:OD1	2.34	0.45
2:G:25:HIS:CD2	2:G:104:LEU:HD21	2.51	0.45
1:B:213:TYR:HA	1:B:339:ILE:O	2.16	0.45
1:B:2707:ALA:HB1	1:B:3009:TYR:CD1	2.52	0.45
1:D:867:LEU:HA	1:D:870:ILE:HG22	1.98	0.45
1:D:2707:ALA:HB1	1:D:3009:TYR:CD1	2.52	0.45
1:D:3891:LEU:HD13	1:D:3899:PHE:HZ	1.79	0.45
1:D:4577:LEU:HD23	1:D:4580:TYR:CE1	2.52	0.45
1:C:36:CYS:SG	1:C:52:THR:HG21	2.55	0.45
1:C:260:TRP:CE3	1:C:282:ILE:HG22	2.51	0.45
1:C:2233:CYS:SG	1:C:2275:VAL:HG23	2.57	0.45
1:C:4705:VAL:HG13	1:C:4711:PHE:CD1	2.51	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:221:ARG:NH2	1:A:255:HIS:O	2.48	0.45
1:A:2912:THR:HG22	1:A:2914:LYS:H	1.81	0.45
1:A:3157:ILE:HG22	1:A:3162:GLN:HG2	1.97	0.45
1:A:4172:GLU:OE1	1:A:4175:ARG:NH1	2.50	0.45
1:A:4645:CYS:O	1:A:4649:LEU:HD13	2.16	0.45
1:B:574:VAL:HA	1:B:577:ILE:HG12	1.98	0.45
1:B:927:GLU:O	1:B:931:THR:HG23	2.16	0.45
1:B:3182:TYR:O	1:B:3185:LYS:HG3	2.15	0.45
1:B:3336:LYS:O	1:B:3340:VAL:HG23	2.16	0.45
1:B:3530:GLN:O	1:B:3534:MET:HG2	2.16	0.45
1:B:4645:CYS:O	1:B:4649:LEU:HD13	2.16	0.45
1:B:4747:SER:HA	1:B:4750:ILE:HG12	1.98	0.45
1:D:1791:VAL:O	1:D:1792:ALA:HB3	2.16	0.45
1:D:2828:GLU:O	1:D:2829:GLY:O	2.34	0.45
1:D:3409:TYR:O	1:D:3412:LEU:N	2.50	0.45
1:D:4176:PRO:O	1:D:4202:ARG:NH2	2.49	0.45
1:D:4690:GLU:O	1:D:4691:GLN:HB2	2.16	0.45
1:D:4747:SER:HA	1:D:4750:ILE:HG12	1.97	0.45
1:C:867:LEU:HA	1:C:870:ILE:HG22	1.98	0.45
1:C:927:GLU:O	1:C:931:THR:HG23	2.16	0.45
1:C:985:VAL:HA	1:C:1039:LEU:HD22	1.98	0.45
1:C:1929:MET:O	1:C:1930:LYS:HB3	2.16	0.45
1:C:2280:VAL:O	1:C:2280:VAL:HG22	2.16	0.45
1:C:2782:ASP:OD2	1:C:2785:LEU:HD13	2.17	0.45
1:C:4016:LEU:O	1:C:4020:GLN:HG3	2.16	0.45
1:A:365:LYS:O	1:A:369:LEU:HG	2.17	0.45
1:A:759:ILE:O	1:A:759:ILE:HG23	2.17	0.45
1:A:869:ARG:HD2	1:A:870:ILE:N	2.31	0.45
1:A:879:HIS:CE1	1:A:913:LEU:HD13	2.51	0.45
1:A:1272:LEU:HD22	1:A:1289:LEU:HD11	1.98	0.45
1:A:2211:MET:O	1:A:2214:VAL:CG1	2.64	0.45
1:A:2233:CYS:SG	1:A:2275:VAL:HG23	2.56	0.45
1:A:2254:LEU:CD1	1:A:2276:ALA:HB1	2.47	0.45
1:A:2261:SER:O	1:A:2264:GLY:N	2.43	0.45
1:A:2572:THR:HG23	1:A:2575:ARG:HB2	1.97	0.45
1:A:3036:LYS:O	1:A:3039:ILE:HG22	2.16	0.45
1:A:3401:LEU:O	1:A:3405:LEU:HD13	2.16	0.45
1:A:4577:LEU:HD23	1:A:4580:TYR:CE1	2.52	0.45
1:A:4902:GLU:O	1:A:4913:ARG:NH2	2.50	0.45
2:F:25:HIS:HD2	2:F:104:LEU:HD21	1.82	0.45
1:B:365:LYS:O	1:B:369:LEU:HG	2.17	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:880:GLU:O	1:B:884:LEU:HD13	2.17	0.45
1:B:941:MET:SD	1:B:1051:TYR:HE1	2.32	0.45
1:B:1272:LEU:HD22	1:B:1289:LEU:HD11	1.98	0.45
1:B:2233:CYS:SG	1:B:2275:VAL:HG23	2.56	0.45
1:D:2011:HIS:CE1	1:D:2017:ASP:CG	2.84	0.45
1:D:2280:VAL:O	1:D:2280:VAL:HG22	2.16	0.45
1:D:2782:ASP:OD2	1:D:2785:LEU:HD13	2.17	0.45
1:D:4172:GLU:OE1	1:D:4175:ARG:NH1	2.50	0.45
1:C:553:ARG:NH2	1:C:555:GLU:OE2	2.50	0.45
1:C:880:GLU:O	1:C:884:LEU:HD13	2.17	0.45
1:C:1791:VAL:O	1:C:1792:ALA:HB3	2.16	0.45
1:C:2211:MET:O	1:C:2214:VAL:CG1	2.64	0.45
1:C:4645:CYS:O	1:C:4649:LEU:HD13	2.16	0.45
1:C:4848:VAL:HG11	1:C:4887:MET:HG2	1.97	0.45
1:A:2661:TRP:HB3	1:A:2664:PHE:HD1	1.81	0.45
2:H:25:HIS:CD2	2:H:104:LEU:HD21	2.51	0.45
2:F:25:HIS:CD2	2:F:104:LEU:HD21	2.51	0.45
1:B:260:TRP:CE3	1:B:282:ILE:HG22	2.51	0.45
1:B:553:ARG:NH2	1:B:555:GLU:OE2	2.50	0.45
1:B:2572:THR:CG2	1:B:2579:VAL:CG2	2.83	0.45
1:D:1272:LEU:HD22	1:D:1289:LEU:HD11	1.98	0.45
1:D:2303:ALA:O	1:D:2307:LEU:HD13	2.17	0.45
1:D:3036:LYS:O	1:D:3039:ILE:HG22	2.16	0.45
1:C:224:HIS:HA	1:C:388:LEU:HD23	1.98	0.45
1:C:759:ILE:HG23	1:C:759:ILE:O	2.17	0.45
1:C:1272:LEU:HD22	1:C:1289:LEU:HD11	1.98	0.45
1:C:1554:VAL:HG21	1:C:1561:VAL:CG1	2.45	0.45
1:C:2214:VAL:HG11	1:C:2228:MET:HE3	1.98	0.45
1:A:2494:PHE:HD1	1:A:2498:HIS:HD1	1.64	0.45
1:A:2886:TRP:O	1:A:2890:LYS:HG2	2.17	0.45
1:A:3086:GLU:OE2	1:A:3093:ARG:NH1	2.50	0.45
1:B:759:ILE:HG23	1:B:759:ILE:O	2.17	0.45
1:B:2303:ALA:O	1:B:2307:LEU:HD13	2.17	0.45
1:B:2886:TRP:O	1:B:2890:LYS:HG2	2.17	0.45
1:D:365:LYS:O	1:D:369:LEU:HG	2.17	0.45
1:D:985:VAL:HA	1:D:1039:LEU:HD22	1.98	0.45
1:D:2244:ARG:NH2	1:D:2283:ASN:OD1	2.50	0.45
1:D:3213:TYR:CD1	1:D:3302:PRO:HG2	2.52	0.45
1:D:3324:VAL:HA	1:D:3327:LEU:HG	1.99	0.45
1:D:3390:GLY:HA2	1:D:3393:LEU:HD13	1.99	0.45
1:D:3414:ARG:HH22	1:D:3472:ALA:CB	2.30	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3899:PHE:CZ	1:D:3903:LEU:HD21	2.52	0.45
1:D:3991:GLY:O	1:D:3995:VAL:HG23	2.16	0.45
1:C:463:GLU:O	1:C:467:LYS:HG2	2.16	0.45
1:C:941:MET:SD	1:C:1051:TYR:HE1	2.32	0.45
1:C:1105:ALA:O	1:C:1189:LEU:N	2.50	0.45
1:C:2254:LEU:CD1	1:C:2276:ALA:HB1	2.47	0.45
1:C:2908:TYR:OH	1:C:2916:LYS:HA	2.16	0.45
1:C:3409:TYR:O	1:C:3412:LEU:N	2.50	0.45
1:C:3899:PHE:CZ	1:C:3903:LEU:HD21	2.52	0.45
1:C:3991:GLY:O	1:C:3995:VAL:HG23	2.16	0.45
1:C:4747:SER:HA	1:C:4750:ILE:HG12	1.98	0.45
1:A:717:ASP:OD1	1:A:720:HIS:N	2.40	0.45
1:A:985:VAL:HA	1:A:1039:LEU:HD22	1.98	0.45
1:A:3106:MET:O	1:A:3110:LEU:HD23	2.17	0.45
1:A:3566:SER:CB	1:A:3569:LEU:HD12	2.47	0.45
2:E:25:HIS:HD2	2:E:104:LEU:HD21	1.82	0.45
1:B:2254:LEU:CD1	1:B:2276:ALA:HB1	2.47	0.45
1:B:3628:ARG:HD2	1:B:3632:VAL:HG13	1.99	0.45
1:D:759:ILE:HG23	1:D:759:ILE:O	2.17	0.45
1:D:1105:ALA:O	1:D:1189:LEU:N	2.50	0.45
1:D:2586:VAL:CG1	1:D:2607:LEU:HD13	2.46	0.45
1:D:3086:GLU:OE2	1:D:3093:ARG:NH1	2.50	0.45
1:C:2244:ARG:NH2	1:C:2283:ASN:OD1	2.50	0.45
1:C:2828:GLU:O	1:C:2829:GLY:O	2.34	0.45
1:C:3390:GLY:HA2	1:C:3393:LEU:HD13	1.99	0.45
1:C:3566:SER:CB	1:C:3569:LEU:HD12	2.47	0.45
1:A:2233:CYS:O	1:A:2235:PHE:N	2.49	0.45
1:A:3802:ILE:HD11	1:A:3883:ASP:O	2.17	0.45
1:A:4016:LEU:O	1:A:4020:GLN:HG3	2.16	0.45
2:G:25:HIS:HD2	2:G:104:LEU:HD21	1.81	0.45
1:B:1873:GLU:OE1	1:B:1873:GLU:N	2.50	0.45
1:B:1929:MET:O	1:B:1930:LYS:HB3	2.16	0.45
1:B:2661:TRP:HB3	1:B:2664:PHE:HD1	1.81	0.45
1:B:3398:PHE:HE2	1:B:3450:ASN:CB	2.30	0.45
1:D:260:TRP:CE3	1:D:282:ILE:HG22	2.51	0.45
1:D:2224:ARG:HE	1:D:2225:PHE:N	2.10	0.45
1:D:4732:PHE:HE2	1:D:4736:ARG:HG2	1.81	0.45
1:D:4902:GLU:O	1:D:4913:ARG:NH2	2.50	0.45
1:C:2303:ALA:O	1:C:2307:LEU:HD13	2.17	0.45
1:C:2685:SER:O	1:C:2689:LYS:HG2	2.17	0.45
1:C:3213:TYR:CD1	1:C:3302:PRO:HG2	2.52	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3324:VAL:HA	1:C:3327:LEU:HG	1.99	0.45
1:C:3414:ARG:HH22	1:C:3472:ALA:CB	2.30	0.45
1:C:3802:ILE:HD11	1:C:3883:ASP:O	2.17	0.45
1:C:4577:LEU:HD23	1:C:4580:TYR:CE1	2.52	0.45
1:C:4821:LYS:O	1:C:4825:THR:HG23	2.17	0.45
1:C:4942:GLU:O	1:C:4945:ASP:OD1	2.34	0.45
1:A:380:GLN:OE1	1:A:380:GLN:N	2.45	0.45
1:A:622:THR:HA	1:A:626:LEU:HD13	1.99	0.45
1:A:1791:VAL:O	1:A:1792:ALA:HB3	2.16	0.45
1:A:2904:LEU:O	1:A:2906:VAL:HG22	2.17	0.45
1:A:3213:TYR:CD1	1:A:3302:PRO:HG2	2.52	0.45
1:A:3414:ARG:HH22	1:A:3472:ALA:CB	2.30	0.45
1:A:4176:PRO:O	1:A:4202:ARG:NH2	2.49	0.45
1:A:4821:LYS:O	1:A:4825:THR:HG23	2.17	0.45
1:B:155:LYS:HD2	1:B:156:GLN:N	2.32	0.45
1:B:345:LEU:HD12	1:B:387:ALA:HB1	1.99	0.45
1:B:721:LEU:HD12	1:B:1476:MET:HE2	1.98	0.45
1:B:820:ARG:CZ	1:B:820:ARG:CB	2.80	0.45
1:B:1126:GLY:HA3	1:B:1143:TRP:CE3	2.52	0.45
1:B:2685:SER:O	1:B:2689:LYS:HG2	2.17	0.45
1:B:3230:LEU:HD12	1:B:3232:LEU:HG	1.99	0.45
1:B:3401:LEU:O	1:B:3405:LEU:HD13	2.16	0.45
1:B:3409:TYR:O	1:B:3412:LEU:N	2.50	0.45
1:B:3566:SER:CB	1:B:3569:LEU:HD12	2.47	0.45
1:B:3899:PHE:CZ	1:B:3903:LEU:HD21	2.52	0.45
1:D:221:ARG:NH2	1:D:255:HIS:O	2.47	0.45
1:D:879:HIS:CE1	1:D:913:LEU:HD13	2.51	0.45
1:D:2303:ALA:O	1:D:2307:LEU:CD1	2.65	0.45
1:D:3580:PRO:O	1:D:3583:GLU:OE2	2.35	0.45
1:D:3891:LEU:CB	1:D:3899:PHE:CE2	3.00	0.45
1:D:3966:THR:HG23	1:D:4029:SER:OG	2.17	0.45
1:D:4095:LYS:O	1:D:4098:ASP:OD1	2.35	0.45
1:C:365:LYS:O	1:C:369:LEU:HG	2.17	0.45
1:C:2303:ALA:O	1:C:2307:LEU:CD1	2.65	0.45
1:C:3107:VAL:O	1:C:3111:ARG:HG2	2.17	0.45
1:A:155:LYS:HD2	1:A:156:GLN:N	2.32	0.44
1:A:336:PRO:HA	1:A:337:PRO:HD3	1.91	0.44
1:A:345:LEU:HD12	1:A:387:ALA:HB1	2.00	0.44
1:A:880:GLU:O	1:A:884:LEU:HD13	2.17	0.44
1:A:927:GLU:O	1:A:931:THR:HG23	2.16	0.44
1:A:2707:ALA:HB1	1:A:3009:TYR:CD1	2.52	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3327:LEU:O	1:A:3403:ARG:NH2	2.50	0.44
1:A:3409:TYR:O	1:A:3412:LEU:N	2.50	0.44
1:B:3036:LYS:O	1:B:3039:ILE:HG22	2.16	0.44
1:B:3324:VAL:HA	1:B:3327:LEU:HG	1.99	0.44
1:B:3414:ARG:HH22	1:B:3472:ALA:CB	2.30	0.44
1:B:4148:THR:HG21	1:B:4180:ARG:NH2	2.31	0.44
1:D:155:LYS:HD2	1:D:156:GLN:N	2.32	0.44
1:D:1515:VAL:O	1:D:1531:ALA:O	2.36	0.44
1:D:1805:GLU:HG2	1:D:1806:ALA:N	2.33	0.44
1:D:2261:SER:O	1:D:2264:GLY:N	2.43	0.44
1:D:3546:ASP:O	1:D:3549:VAL:HG22	2.17	0.44
1:D:4016:LEU:O	1:D:4020:GLN:HG3	2.16	0.44
1:C:820:ARG:HB3	1:C:820:ARG:CZ	2.24	0.44
1:C:2675:THR:CG2	1:C:2706:ILE:HG23	2.48	0.44
1:A:1126:GLY:HA3	1:A:1143:TRP:CE3	2.52	0.44
1:A:1515:VAL:O	1:A:1531:ALA:O	2.36	0.44
2:H:7:ILE:HA	1:D:719:LEU:HD11	1.99	0.44
1:B:224:HIS:HA	1:B:388:LEU:HD23	1.98	0.44
1:B:1742:THR:O	1:B:1960:ALA:HB2	2.18	0.44
1:B:1805:GLU:HG2	1:B:1806:ALA:N	2.33	0.44
1:B:2373:GLY:O	1:C:130:LYS:NZ	2.49	0.44
1:B:3106:MET:O	1:B:3110:LEU:HD23	2.17	0.44
1:B:3842:LEU:HD23	1:B:3874:VAL:CG1	2.47	0.44
1:B:3941:ASP:OD2	1:B:3941:ASP:N	2.46	0.44
1:B:4095:LYS:O	1:B:4098:ASP:OD1	2.35	0.44
1:D:884:LEU:O	1:D:887:ILE:HB	2.16	0.44
1:D:2494:PHE:HD1	1:D:2498:HIS:HD1	1.64	0.44
1:D:3465:ASN:OD1	1:D:3467:MET:CG	2.60	0.44
1:D:4942:GLU:O	1:D:4945:ASP:OD1	2.34	0.44
1:C:553:ARG:O	1:C:557:SER:OG	2.29	0.44
1:C:2226:PRO:HB2	1:C:2267:MET:HE3	1.99	0.44
1:C:3036:LYS:O	1:C:3039:ILE:HG22	2.16	0.44
1:C:3580:PRO:O	1:C:3583:GLU:OE2	2.35	0.44
1:C:3966:THR:HG23	1:C:4029:SER:OG	2.17	0.44
1:C:4094:GLN:HG3	1:C:4108:ILE:HG21	1.99	0.44
1:C:4583:SER:OG	1:C:4585:SER:O	2.27	0.44
1:A:941:MET:SD	1:A:1051:TYR:HE1	2.32	0.44
1:A:2244:ARG:NH2	1:A:2283:ASN:OD1	2.50	0.44
1:A:4094:GLN:HG3	1:A:4108:ILE:HG21	2.00	0.44
1:A:4937:ILE:HG12	1:B:4934:GLY:HA2	1.99	0.44
1:B:985:VAL:HA	1:B:1039:LEU:HD22	1.98	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2214:VAL:HG11	1:B:2228:MET:CE	2.48	0.44
1:B:2303:ALA:O	1:B:2307:LEU:CD1	2.65	0.44
1:B:3107:VAL:O	1:B:3111:ARG:HG2	2.17	0.44
1:B:3802:ILE:HD11	1:B:3883:ASP:O	2.17	0.44
1:B:3966:THR:HG23	1:B:4029:SER:OG	2.17	0.44
1:B:4902:GLU:O	1:B:4913:ARG:NH2	2.50	0.44
1:D:76:ARG:O	1:D:80:GLU:HG2	2.18	0.44
1:D:213:TYR:HA	1:D:339:ILE:O	2.16	0.44
1:D:622:THR:HA	1:D:626:LEU:HD13	1.99	0.44
1:D:2482:ASP:C	1:D:2482:ASP:OD1	2.55	0.44
1:D:2791:LEU:HD13	1:D:2791:LEU:HA	1.89	0.44
1:D:3802:ILE:HD11	1:D:3883:ASP:O	2.17	0.44
1:D:4821:LYS:O	1:D:4825:THR:HG23	2.17	0.44
1:C:2482:ASP:C	1:C:2482:ASP:OD1	2.55	0.44
1:C:2572:THR:CG2	1:C:2579:VAL:CG2	2.83	0.44
1:C:3842:LEU:HD23	1:C:3874:VAL:CG1	2.47	0.44
1:C:4093:PHE:O	1:C:4097:MET:HG2	2.17	0.44
1:C:4943:LEU:O	1:C:4947:GLN:HG2	2.17	0.44
1:A:553:ARG:NH2	1:A:555:GLU:OE2	2.50	0.44
1:A:3230:LEU:HD12	1:A:3232:LEU:HG	1.99	0.44
1:A:3528:THR:HG23	1:A:3573:MET:SD	2.58	0.44
1:A:3546:ASP:O	1:A:3549:VAL:HG22	2.17	0.44
1:A:3842:LEU:HD23	1:A:3874:VAL:CG1	2.47	0.44
1:A:4820:VAL:HG11	1:B:4839:MET:CE	2.47	0.44
2:E:25:HIS:CD2	2:E:104:LEU:HD21	2.51	0.44
2:H:25:HIS:HD2	2:H:104:LEU:HD21	1.82	0.44
1:B:1791:VAL:O	1:B:1792:ALA:HB3	2.16	0.44
1:B:3327:LEU:O	1:B:3403:ARG:NH2	2.50	0.44
1:D:869:ARG:HD2	1:D:870:ILE:N	2.31	0.44
1:D:927:GLU:O	1:D:931:THR:HG23	2.16	0.44
1:D:959:TYR:CD2	1:D:966:LYS:HE2	2.50	0.44
1:D:1456:ASP:OD1	1:D:1456:ASP:N	2.47	0.44
1:D:4093:PHE:O	1:D:4097:MET:HG2	2.17	0.44
1:D:4094:GLN:HG3	1:D:4108:ILE:HG21	1.99	0.44
1:D:4943:LEU:O	1:D:4947:GLN:HG2	2.18	0.44
1:C:984:LEU:HD23	1:C:988:LEU:HD23	2.00	0.44
1:C:1515:VAL:O	1:C:1531:ALA:O	2.35	0.44
1:C:1873:GLU:N	1:C:1873:GLU:OE1	2.50	0.44
1:C:2701:PRO:C	1:C:2702:CYS:SG	2.96	0.44
1:C:2707:ALA:HB1	1:C:3009:TYR:CD1	2.52	0.44
1:C:2904:LEU:O	1:C:2906:VAL:HG22	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3336:LYS:O	1:C:3340:VAL:HG23	2.16	0.44
1:C:4029:SER:HA	1:C:4032:GLU:HG3	2.00	0.44
1:A:2675:THR:CG2	1:A:2706:ILE:HG23	2.47	0.44
1:A:2828:GLU:O	1:A:2829:GLY:O	2.34	0.44
1:A:3107:VAL:O	1:A:3111:ARG:HG2	2.17	0.44
1:A:3398:PHE:HE2	1:A:3450:ASN:CB	2.30	0.44
1:A:4736:ARG:NH1	1:D:4075:GLU:CD	2.70	0.44
1:A:4778:TRP:O	1:A:4782:VAL:HG23	2.18	0.44
1:B:380:GLN:OE1	1:B:380:GLN:N	2.45	0.44
1:B:961:MET:SD	1:B:963:ASN:HB2	2.57	0.44
1:B:2482:ASP:OD1	1:B:2482:ASP:C	2.55	0.44
1:B:2675:THR:CG2	1:B:2706:ILE:HG23	2.47	0.44
1:B:2904:LEU:O	1:B:2906:VAL:HG22	2.17	0.44
1:B:3075:LEU:O	1:B:3146:HIS:NE2	2.51	0.44
1:B:3086:GLU:OE2	1:B:3093:ARG:NH1	2.50	0.44
1:B:3604:TYR:CD1	1:B:3608:GLN:OE1	2.71	0.44
1:B:4093:PHE:O	1:B:4097:MET:HG2	2.17	0.44
1:B:4094:GLN:HG3	1:B:4108:ILE:HG21	2.00	0.44
1:B:4977:THR:OG1	1:B:5029:ARG:NH2	2.51	0.44
1:D:880:GLU:O	1:D:884:LEU:HD13	2.17	0.44
1:D:3628:ARG:HD2	1:D:3632:VAL:HG13	1.99	0.44
1:D:3842:LEU:HD23	1:D:3874:VAL:CG1	2.47	0.44
1:D:4029:SER:HA	1:D:4032:GLU:HG3	2.00	0.44
1:D:4977:THR:OG1	1:D:5029:ARG:NH2	2.51	0.44
1:C:663:TYR:OH	1:C:665:GLU:OE2	2.27	0.44
1:C:1126:GLY:HA3	1:C:1143:TRP:CE3	2.52	0.44
1:C:3086:GLU:OE2	1:C:3093:ARG:NH1	2.50	0.44
1:C:3528:THR:HG23	1:C:3573:MET:SD	2.58	0.44
1:C:3546:ASP:O	1:C:3549:VAL:HG22	2.17	0.44
1:A:1170:MET:CE	1:D:3471:THR:HG22	2.47	0.44
1:A:1805:GLU:HG2	1:A:1806:ALA:N	2.33	0.44
1:A:3580:PRO:O	1:A:3583:GLU:OE2	2.35	0.44
1:B:867:LEU:HA	1:B:870:ILE:HG22	1.98	0.44
1:B:984:LEU:HD23	1:B:988:LEU:HD23	2.00	0.44
1:B:1868:PRO:O	1:B:1872:THR:OG1	2.28	0.44
1:B:2701:PRO:C	1:B:2702:CYS:SG	2.96	0.44
1:B:4077:PHE:O	1:B:4081:VAL:HG23	2.18	0.44
1:B:4732:PHE:CE2	1:B:4736:ARG:HG2	2.53	0.44
1:D:345:LEU:HD12	1:D:387:ALA:HB1	2.00	0.44
1:D:553:ARG:NH2	1:D:555:GLU:OE2	2.50	0.44
1:D:3106:MET:O	1:D:3110:LEU:HD23	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3343:GLN:HE22	1:D:3410:PRO:HB2	1.83	0.44
1:D:3971:GLY:N	1:D:3972:PRO:HA	2.33	0.44
1:D:4778:TRP:O	1:D:4782:VAL:HG23	2.18	0.44
1:C:345:LEU:HD12	1:C:387:ALA:HB1	2.00	0.44
1:C:4902:GLU:O	1:C:4913:ARG:NH2	2.50	0.44
1:C:4977:THR:OG1	1:C:5029:ARG:NH2	2.51	0.44
1:A:155:LYS:HE2	1:D:228:ASP:OD2	2.17	0.44
1:A:345:LEU:HD12	1:A:387:ALA:CB	2.48	0.44
1:A:1873:GLU:OE1	1:A:1873:GLU:N	2.50	0.44
1:A:3984:ARG:NH1	1:B:160:GLY:O	2.48	0.44
1:A:4767:TRP:CE3	1:A:4768:LEU:CD2	2.99	0.44
1:B:1067:SER:O	1:B:1071:ARG:HG3	2.18	0.44
1:B:3537:LYS:NZ	1:B:3607:GLU:OE2	2.33	0.44
1:D:224:HIS:HA	1:D:388:LEU:HD23	1.98	0.44
1:D:598:LYS:O	1:D:602:VAL:HG23	2.18	0.44
1:D:1742:THR:O	1:D:1960:ALA:HB2	2.18	0.44
1:D:3327:LEU:O	1:D:3403:ARG:NH2	2.50	0.44
1:D:3336:LYS:O	1:D:3340:VAL:HG23	2.17	0.44
1:D:3566:SER:CB	1:D:3569:LEU:HD12	2.47	0.44
1:C:622:THR:HA	1:C:626:LEU:HD13	1.99	0.44
1:C:646:PRO:HB2	1:C:648:ILE:HD13	2.00	0.44
1:C:1805:GLU:HG2	1:C:1806:ALA:N	2.33	0.44
1:C:2886:TRP:O	1:C:2890:LYS:HG2	2.17	0.44
1:C:3455:GLU:O	1:C:3459:VAL:HG23	2.17	0.44
1:A:730:VAL:HG23	1:A:1476:MET:SD	2.58	0.44
1:A:867:LEU:HA	1:A:870:ILE:HG22	1.98	0.44
1:A:941:MET:HE3	1:A:1051:TYR:CG	2.49	0.44
1:A:1067:SER:O	1:A:1071:ARG:HG3	2.18	0.44
1:A:2303:ALA:O	1:A:2307:LEU:CD1	2.65	0.44
1:A:2701:PRO:C	1:A:2702:CYS:SG	2.96	0.44
1:A:2929:PHE:CD1	1:A:2932:MET:SD	3.11	0.44
1:A:3343:GLN:HE22	1:A:3410:PRO:HB2	1.83	0.44
1:A:3628:ARG:HD2	1:A:3632:VAL:HG13	1.99	0.44
1:A:4977:THR:OG1	1:A:5029:ARG:NH2	2.51	0.44
2:H:103:LEU:HD21	2:H:106:LEU:HD11	2.00	0.44
1:B:76:ARG:O	1:B:80:GLU:HG2	2.18	0.44
1:B:259:LEU:HD11	1:B:397:GLU:HG2	2.00	0.44
1:B:730:VAL:HG23	1:B:1476:MET:SD	2.58	0.44
1:B:3390:GLY:HA2	1:B:3393:LEU:HD13	1.99	0.44
1:B:3971:GLY:N	1:B:3972:PRO:HA	2.33	0.44
1:B:4943:LEU:O	1:B:4947:GLN:HG2	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:345:LEU:HD12	1:D:387:ALA:CB	2.48	0.44
1:D:2481:LYS:O	1:D:2482:ASP:CG	2.57	0.44
1:D:2904:LEU:O	1:D:2906:VAL:HG22	2.17	0.44
1:D:2929:PHE:CD1	1:D:2932:MET:SD	3.11	0.44
1:D:3455:GLU:O	1:D:3459:VAL:HG23	2.17	0.44
1:D:3789:GLU:OE1	1:D:3789:GLU:N	2.51	0.44
1:D:4767:TRP:CE3	1:D:4768:LEU:CD2	2.99	0.44
1:C:2214:VAL:HG11	1:C:2228:MET:CE	2.48	0.44
1:C:2330:ARG:O	1:C:2333:ASP:OD1	2.36	0.44
1:C:2527:LEU:O	1:C:2531:ARG:HG3	2.18	0.44
1:A:783:PHE:HB2	1:A:787:VAL:HG21	2.00	0.44
1:A:2330:ARG:O	1:A:2333:ASP:OD1	2.36	0.44
1:A:3390:GLY:HA2	1:A:3393:LEU:HD13	1.99	0.44
1:A:4095:LYS:O	1:A:4098:ASP:OD1	2.35	0.44
1:B:266:ARG:O	1:B:270:SER:OG	2.34	0.44
1:B:345:LEU:HD12	1:B:387:ALA:CB	2.48	0.44
1:B:1515:VAL:O	1:B:1531:ALA:O	2.36	0.44
1:B:2926:LEU:O	1:B:2930:LEU:HD13	2.18	0.44
1:B:3343:GLN:HE22	1:B:3410:PRO:HB2	1.83	0.44
1:B:4577:LEU:HD23	1:B:4580:TYR:CE1	2.52	0.44
1:D:941:MET:HE3	1:D:1051:TYR:CG	2.48	0.44
1:D:1694:LEU:O	1:D:1698:LEU:HG	2.18	0.44
1:D:2798:SER:OG	1:D:2801:ASP:OD2	2.26	0.44
1:D:5009:TYR:CZ	1:D:5013:MET:HE3	2.52	0.44
6:D:5304:PNX:HAK1	6:D:5304:PNX:HAA1	2.00	0.44
1:C:155:LYS:HD2	1:C:156:GLN:N	2.32	0.44
1:C:345:LEU:HD12	1:C:387:ALA:CB	2.48	0.44
1:C:1067:SER:O	1:C:1071:ARG:HG3	2.18	0.44
1:C:2929:PHE:CD1	1:C:2932:MET:SD	3.11	0.44
1:C:3327:LEU:O	1:C:3403:ARG:NH2	2.50	0.44
1:C:3429:ALA:HA	1:C:3432:GLU:OE1	2.18	0.44
1:C:4767:TRP:HE3	1:C:4768:LEU:HD23	1.80	0.44
1:A:1077:ALA:HB2	1:A:1190:PRO:HG2	2.00	0.43
1:A:2926:LEU:O	1:A:2930:LEU:HD13	2.18	0.43
1:A:3604:TYR:CD1	1:A:3608:GLN:OE1	2.71	0.43
1:A:4182:GLU:HB2	1:A:4983:HIS:CD2	2.53	0.43
2:F:31:GLU:OE2	1:B:1299:GLN:HG2	2.18	0.43
1:B:2481:LYS:O	1:B:2482:ASP:CG	2.57	0.43
1:B:3455:GLU:O	1:B:3459:VAL:HG23	2.17	0.43
1:B:4778:TRP:O	1:B:4782:VAL:HG23	2.18	0.43
1:D:646:PRO:HB2	1:D:648:ILE:HD13	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:667:MET:SD	1:D:801:LYS:NZ	2.85	0.43
1:D:2434:GLY:O	1:D:2508:ARG:NE	2.43	0.43
1:D:2701:PRO:C	1:D:2702:CYS:SG	2.96	0.43
1:D:3107:VAL:O	1:D:3111:ARG:HG2	2.17	0.43
1:D:3429:ALA:HA	1:D:3432:GLU:OE1	2.18	0.43
1:D:3528:THR:HG23	1:D:3573:MET:SD	2.58	0.43
1:C:76:ARG:O	1:C:80:GLU:HG2	2.18	0.43
1:C:309:THR:O	1:C:309:THR:HG22	2.18	0.43
1:C:3212:GLU:OE2	1:C:3213:TYR:CZ	2.71	0.43
1:C:3343:GLN:HE22	1:C:3410:PRO:HB2	1.83	0.43
1:C:3891:LEU:CB	1:C:3899:PHE:CE2	3.00	0.43
1:C:4095:LYS:O	1:C:4098:ASP:OD1	2.35	0.43
1:C:4182:GLU:HB2	1:C:4983:HIS:CD2	2.53	0.43
1:C:4732:PHE:CE2	1:C:4736:ARG:HG2	2.53	0.43
6:C:5304:PNX:HAA1	6:C:5304:PNX:HAK1	2.00	0.43
1:A:259:LEU:HD11	1:A:397:GLU:HG2	2.00	0.43
1:A:1694:LEU:O	1:A:1698:LEU:HG	2.18	0.43
1:A:2214:VAL:HG11	1:A:2228:MET:CE	2.48	0.43
1:A:3212:GLU:OE2	1:A:3213:TYR:CZ	2.72	0.43
1:A:4093:PHE:O	1:A:4097:MET:HG2	2.17	0.43
2:H:31:GLU:OE2	1:D:1299:GLN:HG2	2.18	0.43
1:B:598:LYS:O	1:B:602:VAL:HG23	2.18	0.43
1:B:615:ARG:NH2	1:B:1676:LEU:O	2.50	0.43
1:B:646:PRO:HB2	1:B:648:ILE:HD13	2.00	0.43
1:B:1694:LEU:O	1:B:1698:LEU:HG	2.18	0.43
1:B:2527:LEU:O	1:B:2531:ARG:HG3	2.18	0.43
1:B:2929:PHE:CD1	1:B:2932:MET:SD	3.11	0.43
1:B:3789:GLU:N	1:B:3789:GLU:OE1	2.51	0.43
1:B:3891:LEU:CB	1:B:3899:PHE:CE2	3.00	0.43
1:B:4029:SER:HA	1:B:4032:GLU:HG3	2.00	0.43
1:D:730:VAL:HG23	1:D:1476:MET:SD	2.58	0.43
1:D:1126:GLY:HA3	1:D:1143:TRP:CE3	2.52	0.43
1:D:1705:GLY:HA3	1:D:1836:PHE:CD2	2.53	0.43
1:D:1873:GLU:OE1	1:D:1873:GLU:N	2.50	0.43
1:D:2254:LEU:CD1	1:D:2276:ALA:HB1	2.47	0.43
1:D:2675:THR:CG2	1:D:2706:ILE:HG23	2.47	0.43
1:D:2685:SER:O	1:D:2689:LYS:HG2	2.17	0.43
1:D:2926:LEU:O	1:D:2930:LEU:HD13	2.18	0.43
1:D:3104:GLU:OE2	1:D:3167:ARG:HB3	2.19	0.43
1:D:3604:TYR:CD1	1:D:3608:GLN:OE1	2.71	0.43
1:C:1045:THR:HG22	1:C:1049:TYR:CZ	2.53	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1705:GLY:HA3	1:C:1836:PHE:CD2	2.53	0.43
1:C:2514:ASN:OD1	1:C:2517:PHE:HB3	2.19	0.43
1:C:3628:ARG:HD2	1:C:3632:VAL:HG13	1.99	0.43
1:C:4778:TRP:O	1:C:4782:VAL:HG23	2.18	0.43
1:A:598:LYS:O	1:A:602:VAL:HG23	2.18	0.43
1:A:646:PRO:HB2	1:A:648:ILE:HD13	2.00	0.43
1:A:1980:LEU:CD2	1:A:1995:THR:HG22	2.48	0.43
1:A:3604:TYR:CE1	1:A:3608:GLN:OE1	2.72	0.43
1:A:3789:GLU:OE1	1:A:3789:GLU:N	2.51	0.43
2:E:88:PRO:O	2:E:90:ILE:HD12	2.19	0.43
2:F:103:LEU:HD21	2:F:106:LEU:HD11	2.00	0.43
1:B:783:PHE:HB2	1:B:787:VAL:HG21	2.00	0.43
1:B:1743[A]:ARG:NH2	1:B:1967:ASP:OD2	2.40	0.43
1:B:2514:ASN:OD1	1:B:2517:PHE:HB3	2.18	0.43
1:B:2791:LEU:HD13	1:B:2791:LEU:HA	1.89	0.43
1:B:3213:TYR:CD1	1:B:3302:PRO:HG2	2.52	0.43
1:D:309:THR:O	1:D:309:THR:HG22	2.18	0.43
1:D:1077:ALA:HB2	1:D:1190:PRO:HG2	2.00	0.43
1:D:2886:TRP:O	1:D:2890:LYS:HG2	2.17	0.43
1:D:3075:LEU:O	1:D:3146:HIS:NE2	2.51	0.43
1:D:3574:ALA:HA	1:D:3577:ARG:NH2	2.33	0.43
1:D:4182:GLU:HB2	1:D:4983:HIS:CD2	2.53	0.43
1:D:4673:ARG:O	1:D:4676:GLU:HG3	2.18	0.43
1:D:4732:PHE:CE2	1:D:4736:ARG:HG2	2.53	0.43
1:C:116:MET:HG2	1:C:139:GLU:HA	2.00	0.43
1:C:336:PRO:HA	1:C:337:PRO:HD3	1.91	0.43
1:C:2092:GLN:O	1:C:2092:GLN:HG2	2.19	0.43
1:C:3604:TYR:CE1	1:C:3608:GLN:OE1	2.72	0.43
1:A:2527:LEU:O	1:A:2531:ARG:HG3	2.18	0.43
1:A:2801:ASP:OD1	1:A:2802:LYS:N	2.52	0.43
1:A:3130:THR:O	1:A:3134:VAL:HG23	2.19	0.43
1:A:3966:THR:HG23	1:A:4029:SER:OG	2.17	0.43
1:A:4029:SER:HA	1:A:4032:GLU:HG3	2.00	0.43
1:A:4651:THR:OG1	1:A:4803:HIS:NE2	2.05	0.43
1:A:4655:PHE:CE2	1:A:4659:ILE:HD11	2.54	0.43
1:A:4732:PHE:CE2	1:A:4736:ARG:HG2	2.53	0.43
1:B:116:MET:HG2	1:B:139:GLU:HA	2.00	0.43
1:B:928:THR:O	1:B:931:THR:OG1	2.30	0.43
1:B:2330:ARG:O	1:B:2333:ASP:OD1	2.36	0.43
1:B:2393:ASP:OD1	1:B:2418:LEU:N	2.52	0.43
1:B:2801:ASP:OD1	1:B:2802:LYS:N	2.52	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3162:GLN:O	1:B:3166:TYR:CD2	2.72	0.43
1:B:3580:PRO:O	1:B:3583:GLU:OE2	2.35	0.43
1:B:4060:LYS:O	1:B:4063:ASP:OD1	2.37	0.43
1:B:4821:LYS:O	1:B:4825:THR:HG23	2.17	0.43
1:D:941:MET:SD	1:D:1051:TYR:HE1	2.32	0.43
1:D:2214:VAL:HG11	1:D:2228:MET:CE	2.48	0.43
1:D:2330:ARG:O	1:D:2333:ASP:OD1	2.36	0.43
1:D:2816:MET:HG3	1:D:2878:LEU:HD13	2.01	0.43
1:D:4655:PHE:CE2	1:D:4659:ILE:HD11	2.54	0.43
1:C:959:TYR:CD2	1:C:966:LYS:HE2	2.50	0.43
1:C:3686:GLU:O	1:C:3687:GLU:HB2	2.18	0.43
1:C:3789:GLU:OE1	1:C:3789:GLU:N	2.51	0.43
1:A:76:ARG:O	1:A:80:GLU:HG2	2.18	0.43
1:A:667:MET:SD	1:A:801:LYS:NZ	2.86	0.43
1:A:3455:GLU:O	1:A:3459:VAL:HG23	2.17	0.43
1:A:3686:GLU:O	1:A:3687:GLU:HB2	2.18	0.43
1:A:3891:LEU:CB	1:A:3899:PHE:CE2	3.00	0.43
1:A:4077:PHE:O	1:A:4081:VAL:HG23	2.18	0.43
1:B:1005:TRP:HB3	1:B:1021:LEU:CD1	2.48	0.43
1:B:1077:ALA:HB2	1:B:1190:PRO:HG2	2.00	0.43
1:B:3574:ALA:HA	1:B:3577:ARG:NH2	2.33	0.43
1:B:4182:GLU:HB2	1:B:4983:HIS:CD2	2.53	0.43
6:B:5304:PNX:HAK1	6:B:5304:PNX:HAA1	2.00	0.43
1:D:615:ARG:NH2	1:D:1676:LEU:O	2.49	0.43
1:D:2393:ASP:OD1	1:D:2418:LEU:N	2.52	0.43
1:D:4717:ASP:OD2	1:D:4723:LYS:NZ	2.35	0.43
1:C:598:LYS:O	1:C:602:VAL:HG23	2.18	0.43
1:C:730:VAL:HG23	1:C:1476:MET:SD	2.58	0.43
1:C:1980:LEU:CD2	1:C:1995:THR:HG22	2.48	0.43
1:C:2393:ASP:OD1	1:C:2418:LEU:N	2.52	0.43
1:C:2481:LYS:O	1:C:2482:ASP:CG	2.56	0.43
1:C:2736:ASP:OD1	1:C:2736:ASP:O	2.36	0.43
1:C:3162:GLN:O	1:C:3166:TYR:CD2	2.72	0.43
1:A:310:LYS:O	1:A:350:HIS:NE2	2.51	0.43
1:A:985:VAL:HG13	1:A:1039:LEU:HD23	2.00	0.43
1:A:2393:ASP:OD1	1:A:2418:LEU:N	2.52	0.43
1:A:3429:ALA:HA	1:A:3432:GLU:OE1	2.18	0.43
1:A:3971:GLY:N	1:A:3972:PRO:HA	2.33	0.43
1:A:4943:LEU:O	1:A:4947:GLN:HG2	2.17	0.43
2:E:24:VAL:HG12	2:E:103:LEU:HA	2.01	0.43
2:H:11:ASP:OD1	2:H:14:THR:HG23	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:H:88:PRO:O	2:H:90:ILE:HD12	2.19	0.43
2:G:24:VAL:HG12	2:G:103:LEU:HA	2.01	0.43
2:G:88:PRO:O	2:G:90:ILE:HD12	2.19	0.43
2:G:103:LEU:HD21	2:G:106:LEU:HD11	2.00	0.43
2:F:11:ASP:OD1	2:F:14:THR:HG23	2.19	0.43
1:B:622:THR:HA	1:B:626:LEU:HD13	1.99	0.43
1:B:874:LEU:O	1:B:878:ILE:HG12	2.19	0.43
1:B:1804:LEU:HD13	1:B:1853:ILE:HD12	2.00	0.43
1:B:2092:GLN:O	1:B:2092:GLN:HG2	2.19	0.43
1:B:2736:ASP:OD1	1:B:2736:ASP:O	2.36	0.43
1:B:2992:GLU:O	1:B:2996:LYS:HG2	2.19	0.43
1:D:874:LEU:O	1:D:878:ILE:HG12	2.19	0.43
1:D:961:MET:SD	1:D:963:ASN:HB2	2.57	0.43
1:D:1126:GLY:O	1:D:1142:PRO:HA	2.19	0.43
1:D:1804:LEU:HD13	1:D:1853:ILE:HD12	2.00	0.43
1:D:2514:ASN:OD1	1:D:2517:PHE:HB3	2.19	0.43
1:D:3104:GLU:O	1:D:3107:VAL:HG22	2.19	0.43
1:D:3398:PHE:HE2	1:D:3450:ASN:CB	2.30	0.43
1:D:3623:LEU:HD12	1:D:3624:LEU:N	2.34	0.43
1:D:3686:GLU:O	1:D:3687:GLU:HB2	2.18	0.43
1:C:1694:LEU:O	1:C:1698:LEU:HG	2.18	0.43
1:C:1742:THR:O	1:C:1960:ALA:HB2	2.18	0.43
1:C:3106:MET:O	1:C:3110:LEU:HD23	2.17	0.43
1:C:3230:LEU:HD12	1:C:3232:LEU:HG	1.99	0.43
1:C:3574:ALA:HA	1:C:3577:ARG:NH2	2.33	0.43
1:C:3604:TYR:CD1	1:C:3608:GLN:OE1	2.71	0.43
1:A:1742:THR:O	1:A:1960:ALA:HB2	2.18	0.43
1:A:2514:ASN:OD1	1:A:2517:PHE:HB3	2.18	0.43
1:A:3104:GLU:O	1:A:3107:VAL:HG22	2.19	0.43
1:A:3465:ASN:OD1	1:A:3467:MET:CG	2.60	0.43
1:A:3640:PRO:HB2	1:A:3642:TYR:CE1	2.54	0.43
6:A:5304:PNX:HAK1	6:A:5304:PNX:HAA1	2.00	0.43
1:B:941:MET:HE3	1:B:1051:TYR:CG	2.52	0.43
1:B:1045:THR:HG22	1:B:1049:TYR:CZ	2.53	0.43
1:B:1705:GLY:HA3	1:B:1836:PHE:CD2	2.53	0.43
1:B:3429:ALA:HA	1:B:3432:GLU:OE1	2.18	0.43
1:B:3528:THR:HG23	1:B:3573:MET:SD	2.58	0.43
1:B:3546:ASP:O	1:B:3549:VAL:HG22	2.17	0.43
1:B:3604:TYR:CE1	1:B:3608:GLN:OE1	2.72	0.43
1:B:3693:LYS:HA	1:B:3693:LYS:HE2	2.01	0.43
1:D:984:LEU:HD23	1:D:988:LEU:HD23	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1154:ASP:OD1	1:D:1156:THR:OG1	2.37	0.43
1:D:2092:GLN:HG2	1:D:2092:GLN:O	2.19	0.43
1:C:310:LYS:O	1:C:350:HIS:NE2	2.51	0.43
1:C:1154:ASP:OD1	1:C:1156:THR:OG1	2.37	0.43
1:C:2190:VAL:HG12	1:C:2198:MET:HE3	2.00	0.43
1:C:2816:MET:HG3	1:C:2878:LEU:HD13	2.01	0.43
1:C:2986:VAL:O	1:C:2986:VAL:HG13	2.19	0.43
1:C:3640:PRO:HB2	1:C:3642:TYR:CE1	2.54	0.43
1:C:4077:PHE:O	1:C:4081:VAL:HG23	2.18	0.43
1:A:1154:ASP:OD1	1:A:1156:THR:OG1	2.37	0.43
1:A:1804:LEU:HD13	1:A:1853:ILE:HD12	2.00	0.43
1:A:3623:LEU:HD12	1:A:3624:LEU:N	2.34	0.43
1:A:3693:LYS:HE2	1:A:3693:LYS:HA	2.01	0.43
1:A:4673:ARG:O	1:A:4676:GLU:HG3	2.18	0.43
1:A:4677:LEU:HD23	1:A:4711:PHE:HE1	1.83	0.43
2:H:24:VAL:HG12	2:H:103:LEU:HA	2.01	0.43
1:B:484:LEU:HD12	1:B:526:LEU:HD12	2.01	0.43
1:B:2244:ARG:NH2	1:B:2283:ASN:OD1	2.50	0.43
1:B:3104:GLU:OE2	1:B:3167:ARG:HB3	2.19	0.43
1:D:259:LEU:HD11	1:D:397:GLU:HG2	2.00	0.43
1:D:985:VAL:HG13	1:D:1039:LEU:HD23	2.00	0.43
1:D:1005:TRP:HB3	1:D:1021:LEU:CD1	2.48	0.43
1:D:1067:SER:O	1:D:1071:ARG:HG3	2.18	0.43
1:D:2211:MET:O	1:D:2214:VAL:CG1	2.64	0.43
1:D:2801:ASP:OD1	1:D:2802:LYS:N	2.52	0.43
1:D:3197:LEU:HD21	1:D:3201:MET:HE3	2.01	0.43
1:D:4000:MET:SD	1:D:4020:GLN:NE2	2.79	0.43
1:C:259:LEU:HD11	1:C:397:GLU:HG2	2.00	0.43
1:C:2801:ASP:OD1	1:C:2802:LYS:N	2.52	0.43
1:C:2992:GLU:O	1:C:2996:LYS:HG2	2.19	0.43
1:C:3104:GLU:O	1:C:3107:VAL:HG22	2.19	0.43
1:C:3583:GLU:HB3	1:C:3585:ASP:OD1	2.19	0.43
1:C:3971:GLY:N	1:C:3972:PRO:HA	2.33	0.43
1:A:266:ARG:O	1:A:270:SER:OG	2.34	0.43
1:A:984:LEU:HD23	1:A:988:LEU:HD23	2.00	0.43
1:A:1705:GLY:HA3	1:A:1836:PHE:CD2	2.53	0.43
1:A:2800:LYS:O	1:A:2804:ILE:HG13	2.19	0.43
1:A:2816:MET:HG3	1:A:2878:LEU:HD13	2.01	0.43
1:A:3391:GLU:O	1:A:3395:ARG:HD2	2.19	0.43
1:A:3777:GLU:OE1	1:A:3777:GLU:N	2.47	0.43
1:B:309:THR:HG22	1:B:309:THR:O	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:310:LYS:O	1:B:350:HIS:NE2	2.51	0.43
1:B:1126:GLY:O	1:B:1142:PRO:HA	2.19	0.43
1:B:3212:GLU:OE2	1:B:3213:TYR:CZ	2.72	0.43
1:B:3682:GLU:O	1:B:3686:GLU:OE2	2.37	0.43
1:B:3725:TYR:O	1:B:3729:MET:HG3	2.19	0.43
1:D:1045:THR:HG22	1:D:1049:TYR:CZ	2.53	0.43
1:D:1828:ASP:N	1:D:1828:ASP:OD1	2.52	0.43
1:D:1980:LEU:CD2	1:D:1995:THR:HG22	2.48	0.43
1:D:3693:LYS:HE2	1:D:3693:LYS:HA	2.01	0.43
1:D:3777:GLU:OE1	1:D:3777:GLU:N	2.47	0.43
1:D:4946:GLN:O	1:D:4950:VAL:HG23	2.19	0.43
1:C:1077:ALA:HB2	1:C:1190:PRO:HG2	2.01	0.43
1:C:3075:LEU:O	1:C:3146:HIS:NE2	2.51	0.43
1:C:3104:GLU:OE2	1:C:3167:ARG:HB3	2.19	0.43
1:A:116:MET:HG2	1:A:139:GLU:HA	2.00	0.43
1:A:820:ARG:HB3	1:A:820:ARG:CZ	2.24	0.43
1:A:2481:LYS:O	1:A:2482:ASP:CG	2.56	0.43
1:A:2685:SER:O	1:A:2689:LYS:HG2	2.17	0.43
1:A:2736:ASP:O	1:A:2736:ASP:OD1	2.36	0.43
1:A:3104:GLU:OE2	1:A:3167:ARG:HB3	2.19	0.43
1:A:4148:THR:HG21	1:A:4180:ARG:NH2	2.31	0.43
1:A:4182:GLU:HB2	1:A:4983:HIS:NE2	2.34	0.43
2:F:24:VAL:HG12	2:F:103:LEU:HA	2.01	0.43
1:B:1440:PHE:CZ	1:B:1563:GLN:HG3	2.54	0.43
1:B:1828:ASP:OD1	1:B:1828:ASP:N	2.52	0.43
1:B:2800:LYS:O	1:B:2804:ILE:HG13	2.19	0.43
1:B:3686:GLU:O	1:B:3687:GLU:HB2	2.18	0.43
1:B:4901:ILE:HG13	1:B:4913:ARG:NH2	2.34	0.43
1:D:167:ASP:OD1	1:C:384:MET:SD	2.77	0.43
1:D:2736:ASP:O	1:D:2736:ASP:OD1	2.36	0.43
1:D:2800:LYS:O	1:D:2804:ILE:HG13	2.19	0.43
1:D:3162:GLN:O	1:D:3166:TYR:CD2	2.72	0.43
1:D:3175:LEU:O	1:D:3175:LEU:HD23	2.19	0.43
1:D:3212:GLU:OE2	1:D:3213:TYR:CZ	2.71	0.43
1:D:3230:LEU:HD12	1:D:3232:LEU:HG	1.99	0.43
1:D:4781:GLY:O	1:D:4785:THR:HG23	2.19	0.43
1:C:874:LEU:O	1:C:878:ILE:HG12	2.19	0.43
1:C:2725:LYS:HE3	1:C:2734:ASN:O	2.19	0.43
1:C:3239:MET:CB	1:C:3239:MET:HG2	2.31	0.43
1:C:3891:LEU:HD13	1:C:3899:PHE:HZ	1.79	0.43
1:C:4655:PHE:CE2	1:C:4659:ILE:HD11	2.54	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4673:ARG:O	1:C:4676:GLU:HG3	2.18	0.43
1:C:4781:GLY:O	1:C:4785:THR:HG23	2.19	0.43
1:A:908:VAL:O	1:A:909:ASN:HB2	2.19	0.42
1:A:1105:ALA:HB2	1:A:1191:VAL:HG11	2.00	0.42
1:A:1143:TRP:HB2	1:A:1147:ASP:HB2	2.01	0.42
1:A:3162:GLN:O	1:A:3166:TYR:CD2	2.72	0.42
1:A:3583:GLU:HB3	1:A:3585:ASP:OD1	2.19	0.42
1:A:4000:MET:SD	1:A:4020:GLN:NE2	2.79	0.42
1:A:4060:LYS:O	1:A:4063:ASP:OD1	2.37	0.42
1:A:4901:ILE:HG13	1:A:4913:ARG:NH2	2.34	0.42
2:E:11:ASP:OD1	2:E:14:THR:HG23	2.19	0.42
1:B:1154:ASP:OD1	1:B:1156:THR:OG1	2.37	0.42
1:B:1756:ASN:OD1	1:B:2040:ALA:HB2	2.19	0.42
1:B:3239:MET:CB	1:B:3239:MET:HG3	2.31	0.42
1:B:3380:ARG:NH1	1:B:3391:GLU:OE2	2.50	0.42
1:D:2527:LEU:O	1:D:2531:ARG:HG3	2.18	0.42
1:D:2572:THR:CG2	1:D:2579:VAL:CG2	2.83	0.42
1:D:3868:ARG:HG2	1:D:3869:GLN:N	2.34	0.42
1:D:4077:PHE:O	1:D:4081:VAL:HG23	2.18	0.42
1:C:484:LEU:HD12	1:C:526:LEU:HD12	2.01	0.42
1:C:783:PHE:HB2	1:C:787:VAL:HG21	2.00	0.42
1:C:961:MET:SD	1:C:963:ASN:HB2	2.57	0.42
1:C:1436:SER:OG	1:C:1565:GLU:HB2	2.19	0.42
1:C:2969:ILE:O	1:C:2973:PHE:HD1	2.02	0.42
1:C:3682:GLU:O	1:C:3686:GLU:OE2	2.37	0.42
1:C:3693:LYS:HE2	1:C:3693:LYS:HA	2.01	0.42
1:A:1005:TRP:HB3	1:A:1021:LEU:CD1	2.48	0.42
1:A:1045:THR:HG22	1:A:1049:TYR:CZ	2.54	0.42
1:A:1436:SER:OG	1:A:1565:GLU:HB2	2.19	0.42
1:A:1756:ASN:OD1	1:A:2040:ALA:HB2	2.19	0.42
1:A:2992:GLU:O	1:A:2996:LYS:HG2	2.19	0.42
1:A:4946:GLN:O	1:A:4950:VAL:HG23	2.19	0.42
2:G:31:GLU:OE2	1:C:1299:GLN:HG2	2.19	0.42
1:B:1105:ALA:HB2	1:B:1191:VAL:HG11	2.00	0.42
1:B:1105:ALA:O	1:B:1189:LEU:N	2.50	0.42
1:B:2595:LEU:O	1:B:2600:ARG:NH2	2.52	0.42
1:B:3573:MET:CB	1:B:3577:ARG:NH1	2.73	0.42
1:B:3623:LEU:HD12	1:B:3624:LEU:N	2.34	0.42
1:B:4673:ARG:O	1:B:4676:GLU:HG3	2.18	0.42
1:D:928:THR:O	1:D:931:THR:OG1	2.30	0.42
1:D:3604:TYR:CE1	1:D:3608:GLN:OE1	2.72	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3725:TYR:O	1:D:3729:MET:HG3	2.19	0.42
1:D:4231:MET:O	1:D:4235:VAL:HG23	2.20	0.42
1:C:1005:TRP:HB3	1:C:1021:LEU:CD1	2.48	0.42
1:C:2208:MET:O	1:C:2212:VAL:HG22	2.19	0.42
1:C:2434:GLY:O	1:C:2508:ARG:NE	2.43	0.42
1:C:3197:LEU:HD21	1:C:3201:MET:HE3	2.01	0.42
1:C:3725:TYR:O	1:C:3729:MET:HG3	2.19	0.42
1:C:4060:LYS:O	1:C:4063:ASP:OD1	2.37	0.42
1:C:4182:GLU:HB2	1:C:4983:HIS:NE2	2.34	0.42
1:A:553:ARG:O	1:A:557:SER:OG	2.29	0.42
1:A:893:TYR:HD1	1:A:907:LEU:O	2.03	0.42
1:A:921:ASN:O	1:A:924:MET:HG2	2.19	0.42
1:A:2874:MET:HE1	1:A:2939:ARG:HB2	1.95	0.42
1:A:2986:VAL:HG13	1:A:2986:VAL:O	2.19	0.42
1:A:3071:LEU:O	1:A:3075:LEU:HG	2.19	0.42
1:A:3471:THR:HG22	1:B:1170:MET:CE	2.49	0.42
1:B:921:ASN:O	1:B:924:MET:HG2	2.19	0.42
1:B:2700:MET:O	1:B:2702:CYS:SG	2.78	0.42
1:B:3130:THR:O	1:B:3134:VAL:HG23	2.19	0.42
1:D:492:ASP:O	1:D:496:VAL:HG13	2.19	0.42
1:D:2414:ASN:OD1	1:D:2414:ASN:C	2.58	0.42
1:D:4182:GLU:HB2	1:D:4983:HIS:NE2	2.34	0.42
1:C:266:ARG:O	1:C:270:SER:OG	2.34	0.42
1:C:1743[A]:ARG:NH2	1:C:1967:ASP:OD2	2.39	0.42
1:C:2865:VAL:HG11	1:C:2928:LYS:HE2	2.02	0.42
1:C:2926:LEU:O	1:C:2930:LEU:HD13	2.18	0.42
1:C:3197:LEU:HD23	1:C:3197:LEU:O	2.20	0.42
1:C:3879:GLU:HG2	1:C:3880:PHE:N	2.35	0.42
1:C:4021:LYS:N	1:C:4139:ILE:HD13	2.35	0.42
1:C:4563:ARG:NH2	1:C:4815:ASP:OD1	2.53	0.42
1:A:2414:ASN:OD1	1:A:2414:ASN:C	2.58	0.42
2:F:24:VAL:HG21	2:F:59:TRP:HZ3	1.85	0.42
1:B:2816:MET:HG3	1:B:2878:LEU:HD13	2.01	0.42
1:B:4021:LYS:N	1:B:4139:ILE:HD13	2.35	0.42
1:B:4182:GLU:HB2	1:B:4983:HIS:NE2	2.34	0.42
1:B:4231:MET:O	1:B:4235:VAL:HG23	2.20	0.42
1:D:783:PHE:HB2	1:D:787:VAL:HG21	2.00	0.42
1:D:921:ASN:O	1:D:924:MET:HG2	2.19	0.42
1:D:1194:LEU:HD23	1:D:1198:GLN:O	2.19	0.42
1:D:1434:TYR:HB2	1:D:1572:ILE:HG21	2.01	0.42
1:D:2986:VAL:HG13	1:D:2986:VAL:O	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3640:PRO:HB2	1:D:3642:TYR:CE1	2.54	0.42
1:D:3892:CYS:HG	1:D:3899:PHE:HD2	1.67	0.42
1:C:908:VAL:O	1:C:909:ASN:HB2	2.19	0.42
1:C:1194:LEU:HD23	1:C:1198:GLN:O	2.19	0.42
1:C:1828:ASP:OD1	1:C:1828:ASP:N	2.52	0.42
1:C:2612[B]:ARG:NH1	1:C:2613:TYR:OH	2.53	0.42
1:A:998:ARG:HG2	1:A:998:ARG:HH11	1.84	0.42
1:A:3682:GLU:O	1:A:3686:GLU:OE2	2.37	0.42
1:A:4781:GLY:O	1:A:4785:THR:HG23	2.19	0.42
2:E:103:LEU:HD21	2:E:106:LEU:HD11	2.00	0.42
2:G:11:ASP:OD1	2:G:14:THR:HG23	2.19	0.42
1:B:893:TYR:HD1	1:B:907:LEU:O	2.03	0.42
1:B:2703:LEU:HD22	1:B:2706:ILE:HD12	2.02	0.42
1:B:3071:LEU:O	1:B:3075:LEU:HG	2.19	0.42
1:B:3175:LEU:O	1:B:3175:LEU:HD23	2.19	0.42
1:B:3197:LEU:HD23	1:B:3197:LEU:O	2.20	0.42
1:B:3868:ARG:HG2	1:B:3869:GLN:N	2.34	0.42
1:B:3879:GLU:HG2	1:B:3880:PHE:N	2.35	0.42
1:B:4781:GLY:O	1:B:4785:THR:HG23	2.19	0.42
1:D:116:MET:HG2	1:D:139:GLU:HA	2.00	0.42
1:D:266:ARG:O	1:D:270:SER:OG	2.34	0.42
1:D:998:ARG:HG2	1:D:998:ARG:HH11	1.84	0.42
1:D:1143:TRP:HB2	1:D:1147:ASP:HB2	2.01	0.42
1:D:2595:LEU:O	1:D:2600:ARG:NH2	2.52	0.42
1:D:3879:GLU:HG2	1:D:3880:PHE:N	2.34	0.42
1:C:1105:ALA:HB2	1:C:1191:VAL:HG11	2.00	0.42
1:C:1143:TRP:HB2	1:C:1147:ASP:HB2	2.01	0.42
1:C:2595:LEU:O	1:C:2600:ARG:NH2	2.52	0.42
1:C:2667:THR:HG21	1:C:2672:LEU:HG	2.01	0.42
1:C:3294:PRO:HB2	1:C:3297:PRO:HD2	2.01	0.42
1:C:3435:PHE:CD2	1:C:3524:MET:CE	3.03	0.42
1:C:3573:MET:CB	1:C:3577:ARG:NH1	2.73	0.42
1:C:4946:GLN:O	1:C:4950:VAL:HG23	2.19	0.42
1:A:309:THR:O	1:A:309:THR:HG22	2.18	0.42
1:A:484:LEU:HD12	1:A:526:LEU:HD12	2.01	0.42
1:A:492:ASP:O	1:A:496:VAL:HG13	2.19	0.42
1:A:2700:MET:O	1:A:2702:CYS:SG	2.78	0.42
1:A:2703:LEU:HD22	1:A:2706:ILE:HD12	2.02	0.42
1:A:3725:TYR:O	1:A:3729:MET:HG3	2.19	0.42
1:A:3879:GLU:HG2	1:A:3880:PHE:N	2.35	0.42
1:A:4021:LYS:N	1:A:4139:ILE:HD13	2.35	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:908:VAL:O	1:B:909:ASN:HB2	2.19	0.42
1:B:1436:SER:OG	1:B:1565:GLU:HB2	2.20	0.42
1:B:1727:ARG:O	1:B:1731:LEU:HG	2.20	0.42
1:B:2667:THR:HG21	1:B:2672:LEU:HG	2.01	0.42
1:B:2977:LEU:HD23	1:B:3056:LEU:HD11	2.02	0.42
1:B:3435:PHE:CD2	1:B:3524:MET:CE	3.03	0.42
1:B:3640:PRO:HB2	1:B:3642:TYR:CE1	2.54	0.42
1:D:908:VAL:O	1:D:909:ASN:HB2	2.19	0.42
1:D:1927:LEU:HD22	1:D:2101:MET:SD	2.60	0.42
1:D:3610:GLU:OE1	1:D:3611:HIS:CD2	2.73	0.42
1:D:4060:LYS:O	1:D:4063:ASP:OD1	2.37	0.42
1:C:921:ASN:O	1:C:924:MET:HG2	2.20	0.42
1:C:1804:LEU:HD13	1:C:1853:ILE:HD12	2.00	0.42
1:C:3391:GLU:O	1:C:3395:ARG:HD2	2.19	0.42
1:A:1440:PHE:CZ	1:A:1563:GLN:HG3	2.54	0.42
1:A:1828:ASP:N	1:A:1828:ASP:OD1	2.52	0.42
1:A:1992:ALA:O	1:A:1995:THR:OG1	2.29	0.42
1:A:2092:GLN:HG2	1:A:2092:GLN:O	2.19	0.42
1:A:2545:GLU:HG2	1:A:2546:MET:N	2.35	0.42
1:A:2595:LEU:O	1:A:2600:ARG:NH2	2.52	0.42
1:A:3386:GLU:O	1:A:3389:GLU:O	2.38	0.42
1:A:3574:ALA:HA	1:A:3577:ARG:NH2	2.33	0.42
2:F:88:PRO:O	2:F:90:ILE:HD12	2.19	0.42
1:B:278:GLN:HG3	1:B:328:LYS:NZ	2.35	0.42
1:B:873:LYS:O	1:B:876:GLU:HG2	2.20	0.42
1:B:1194:LEU:HD23	1:B:1198:GLN:O	2.19	0.42
1:B:1276:THR:O	1:B:1279:SER:OG	2.27	0.42
1:B:1927:LEU:HD22	1:B:2101:MET:SD	2.60	0.42
1:B:2208:MET:O	1:B:2212:VAL:HG22	2.19	0.42
1:D:69:LEU:CD2	1:D:109:LEU:HD21	2.50	0.42
1:D:2208:MET:O	1:D:2212:VAL:HG22	2.19	0.42
1:D:3130:THR:O	1:D:3134:VAL:HG23	2.19	0.42
1:D:3660:ALA:HB3	1:D:3661:TRP:HD1	1.85	0.42
1:D:3899:PHE:HE2	1:D:3903:LEU:HD11	1.85	0.42
1:C:69:LEU:CD2	1:C:109:LEU:HD21	2.50	0.42
1:C:1126:GLY:O	1:C:1142:PRO:HA	2.19	0.42
1:C:1927:LEU:HD22	1:C:2101:MET:SD	2.60	0.42
1:C:3175:LEU:HD23	1:C:3175:LEU:O	2.19	0.42
1:C:4179:GLY:CA	1:C:4197:ILE:HD11	2.50	0.42
1:A:167:ASP:N	1:D:384:MET:SD	2.92	0.42
1:A:468:LEU:O	1:A:472:ARG:HG2	2.20	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:874:LEU:O	1:A:878:ILE:HG12	2.19	0.42
1:A:1194:LEU:HD23	1:A:1198:GLN:O	2.19	0.42
1:A:1927:LEU:HD22	1:A:2101:MET:SD	2.60	0.42
1:A:2798:SER:OG	1:A:2801:ASP:OD2	2.26	0.42
1:A:3478:MET:O	1:A:3478:MET:HG3	2.19	0.42
1:A:4999:ASP:OD1	1:A:5000:GLU:N	2.53	0.42
1:B:1434:TYR:HB2	1:B:1572:ILE:HG21	2.01	0.42
1:B:2414:ASN:OD1	1:B:2414:ASN:C	2.58	0.42
1:D:1756:ASN:OD1	1:D:2040:ALA:HB2	2.19	0.42
1:D:2313:LEU:HB3	1:D:2318:TYR:HB2	2.02	0.42
1:D:2587:TYR:OH	1:D:2629:ASP:OD2	2.24	0.42
1:D:2992:GLU:O	1:D:2996:LYS:HG2	2.19	0.42
1:D:3682:GLU:O	1:D:3686:GLU:OE2	2.37	0.42
1:C:292:ALA:HB2	1:C:312:THR:HG22	2.02	0.42
1:C:468:LEU:O	1:C:472:ARG:HG2	2.20	0.42
1:C:873:LYS:O	1:C:876:GLU:HG2	2.20	0.42
1:C:2001:PRO:O	1:C:2005:GLN:HG3	2.20	0.42
1:C:2545:GLU:HG2	1:C:2546:MET:N	2.35	0.42
1:C:3458:PHE:CE2	1:C:3464:ILE:HD11	2.54	0.42
1:A:869:ARG:NH2	1:A:947:GLU:OE2	2.52	0.42
1:A:1126:GLY:O	1:A:1142:PRO:HA	2.19	0.42
1:A:1434:TYR:HB2	1:A:1572:ILE:HG21	2.01	0.42
1:A:3294:PRO:HB2	1:A:3297:PRO:HD2	2.01	0.42
1:A:4843:LEU:HD21	1:D:4827:LEU:HD21	2.02	0.42
1:B:292:ALA:HB2	1:B:312:THR:HG22	2.02	0.42
1:B:309:THR:O	1:B:309:THR:CG2	2.68	0.42
1:B:468:LEU:O	1:B:472:ARG:HG2	2.20	0.42
1:B:516:LYS:O	1:B:519:VAL:HG22	2.20	0.42
1:B:3103:ILE:O	1:B:3107:VAL:HG13	2.20	0.42
1:B:3148:ALA:O	1:B:3151:GLN:NE2	2.53	0.42
1:B:3478:MET:O	1:B:3478:MET:HG3	2.19	0.42
1:B:4187:SER:O	1:B:4188:ARG:HB2	2.20	0.42
1:B:4705:VAL:HG23	1:B:4775:TYR:HD2	1.85	0.42
1:D:292:ALA:HB2	1:D:312:THR:HG22	2.02	0.42
1:D:309:THR:O	1:D:309:THR:CG2	2.68	0.42
1:D:516:LYS:O	1:D:519:VAL:HG22	2.20	0.42
1:D:887:ILE:HD13	1:D:887:ILE:HA	1.87	0.42
1:D:893:TYR:HD1	1:D:907:LEU:O	2.03	0.42
1:D:1105:ALA:HB2	1:D:1191:VAL:HG11	2.00	0.42
1:D:1265:ASP:OD1	1:D:1266:THR:N	2.53	0.42
1:D:1440:PHE:CZ	1:D:1563:GLN:HG3	2.54	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2545:GLU:HG2	1:D:2546:MET:N	2.35	0.42
1:D:3148:ALA:O	1:D:3151:GLN:NE2	2.53	0.42
1:D:3321:ARG:HA	1:D:3324:VAL:HG22	2.02	0.42
1:D:3583:GLU:HB3	1:D:3585:ASP:OD1	2.19	0.42
1:D:4021:LYS:N	1:D:4139:ILE:HD13	2.35	0.42
1:D:4179:GLY:CA	1:D:4197:ILE:HD11	2.50	0.42
1:C:1434:TYR:HB2	1:C:1572:ILE:HG21	2.01	0.42
1:C:2700:MET:O	1:C:2702:CYS:SG	2.78	0.42
1:C:2977:LEU:HD23	1:C:3056:LEU:HD11	2.02	0.42
1:C:3623:LEU:HD12	1:C:3624:LEU:N	2.34	0.42
1:A:484:LEU:HD11	1:A:526:LEU:HD11	2.02	0.42
1:A:4179:GLY:HA3	1:A:4197:ILE:HD11	2.02	0.42
2:E:24:VAL:HG21	2:E:59:TRP:HZ3	1.85	0.42
1:B:384:MET:SD	1:C:167:ASP:OD1	2.78	0.42
1:B:484:LEU:HD11	1:B:526:LEU:HD11	2.02	0.42
1:B:2865:VAL:HG11	1:B:2928:LYS:HE2	2.02	0.42
1:B:3391:GLU:O	1:B:3395:ARG:HD2	2.19	0.42
1:B:3997:ALA:HB1	1:B:4057:MET:SD	2.60	0.42
1:B:4655:PHE:CE2	1:B:4659:ILE:HD11	2.54	0.42
1:D:310:LYS:O	1:D:350:HIS:NE2	2.51	0.42
1:D:484:LEU:HD12	1:D:526:LEU:HD12	2.01	0.42
1:D:2441:HIS:CE1	1:D:2442:LEU:HD12	2.55	0.42
1:D:2612[B]:ARG:NH1	1:D:2613:TYR:OH	2.53	0.42
1:D:3197:LEU:HD23	1:D:3197:LEU:O	2.20	0.42
1:D:3478:MET:O	1:D:3478:MET:HG3	2.19	0.42
1:D:4069:LYS:O	1:D:4072:VAL:HG22	2.20	0.42
1:C:484:LEU:HD11	1:C:526:LEU:HD11	2.02	0.42
1:C:484:LEU:CD1	1:C:526:LEU:CD1	2.98	0.42
1:C:1440:PHE:CZ	1:C:1563:GLN:HG3	2.54	0.42
1:C:1756:ASN:OD1	1:C:2040:ALA:HB2	2.19	0.42
1:C:2800:LYS:O	1:C:2804:ILE:HG13	2.19	0.42
1:C:2946:LEU:N	1:C:2948:THR:HG23	2.35	0.42
1:C:3478:MET:HG3	1:C:3478:MET:O	2.19	0.42
1:C:4999:ASP:OD1	1:C:5000:GLU:N	2.53	0.42
1:A:1265:ASP:OD1	1:A:1266:THR:N	2.53	0.41
1:A:1684:ALA:HA	2:E:90:ILE:HD11	2.02	0.41
1:A:2495:VAL:HG22	1:A:2498:HIS:CE1	2.55	0.41
1:A:2667:THR:HG21	1:A:2672:LEU:HG	2.01	0.41
1:A:3111:ARG:O	1:A:3112:LEU:HB2	2.20	0.41
1:A:3148:ALA:O	1:A:3151:GLN:NE2	2.53	0.41
1:A:3175:LEU:O	1:A:3175:LEU:HD23	2.19	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3321:ARG:HA	1:A:3324:VAL:HG22	2.02	0.41
1:A:3997:ALA:HB1	1:A:4057:MET:SD	2.60	0.41
1:A:4231:MET:O	1:A:4235:VAL:HG23	2.20	0.41
1:B:484:LEU:CD1	1:B:526:LEU:CD1	2.98	0.41
1:B:998:ARG:HG2	1:B:998:ARG:HH11	1.84	0.41
1:B:2928:LYS:O	1:B:2932:MET:HE3	2.20	0.41
1:B:3077:ALA:O	1:B:3081:MET:HG2	2.20	0.41
1:B:3104:GLU:O	1:B:3107:VAL:HG22	2.19	0.41
1:B:3633:VAL:O	1:B:3637:ARG:HD3	2.20	0.41
1:B:4179:GLY:CA	1:B:4197:ILE:HD11	2.50	0.41
1:B:4179:GLY:HA3	1:B:4197:ILE:HD11	2.02	0.41
1:D:484:LEU:HD11	1:D:526:LEU:HD11	2.02	0.41
1:D:1819:VAL:HG22	1:D:1926:LEU:HD21	2.02	0.41
1:D:2495:VAL:HG22	1:D:2498:HIS:CE1	2.55	0.41
1:D:3294:PRO:HB2	1:D:3297:PRO:HD2	2.02	0.41
1:D:3458:PHE:CE2	1:D:3464:ILE:HD11	2.54	0.41
1:D:4999:ASP:OD1	1:D:5000:GLU:N	2.53	0.41
1:C:492:ASP:O	1:C:496:VAL:HG13	2.19	0.41
1:C:509:GLU:O	1:C:513:GLU:CD	2.59	0.41
1:C:926:GLY:O	1:C:930:LYS:HG3	2.20	0.41
1:C:998:ARG:HG2	1:C:998:ARG:HH11	1.84	0.41
1:C:1265:ASP:OD1	1:C:1266:THR:N	2.53	0.41
1:C:1727:ARG:O	1:C:1731:LEU:HG	2.20	0.41
1:C:2117:VAL:O	1:C:2120:MET:HG2	2.21	0.41
1:C:2135:LEU:HD12	1:C:3658:LYS:HG3	2.02	0.41
1:C:2313:LEU:HB3	1:C:2318:TYR:HB2	2.02	0.41
1:C:2487:GLN:NE2	1:C:2545:GLU:OE1	2.53	0.41
1:C:3660:ALA:HB3	1:C:3661:TRP:HD1	1.85	0.41
1:C:4231:MET:O	1:C:4235:VAL:HG23	2.20	0.41
1:A:509:GLU:O	1:A:513:GLU:CD	2.59	0.41
1:A:885:THR:HA	1:A:888:GLU:OE1	2.20	0.41
1:A:2001:PRO:O	1:A:2005:GLN:HG3	2.20	0.41
1:A:2374:SER:OG	1:A:2378:ALA:HB3	2.21	0.41
1:A:2444:GLN:OE1	1:A:2444:GLN:N	2.53	0.41
1:A:3610:GLU:OE1	1:A:3611:HIS:CD2	2.73	0.41
1:A:3899:PHE:HE2	1:A:3903:LEU:HD11	1.85	0.41
1:A:4705:VAL:HG23	1:A:4775:TYR:HD2	1.85	0.41
2:G:24:VAL:HG21	2:G:59:TRP:HZ3	1.85	0.41
1:B:384:MET:SD	1:C:167:ASP:N	2.93	0.41
1:B:492:ASP:O	1:B:496:VAL:HG13	2.19	0.41
1:B:1676:LEU:HD22	1:B:2167:ILE:HD12	2.03	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2202:GLY:HA2	1:B:2205:GLU:OE1	2.20	0.41
1:B:2487:GLN:NE2	1:B:2545:GLU:OE1	2.53	0.41
1:B:2495:VAL:HG22	1:B:2498:HIS:CE1	2.55	0.41
1:B:2725:LYS:HE3	1:B:2734:ASN:O	2.19	0.41
1:B:2986:VAL:O	1:B:2986:VAL:HG13	2.19	0.41
1:B:3003:LEU:HB2	1:B:3004:PRO:HD3	2.02	0.41
1:B:3111:ARG:HH21	1:B:3178:THR:CB	2.33	0.41
1:B:3916:ILE:HD12	1:B:3916:ILE:H	1.86	0.41
1:B:4069:LYS:O	1:B:4072:VAL:HG22	2.21	0.41
1:D:336:PRO:HA	1:D:337:PRO:HD3	1.91	0.41
1:D:873:LYS:O	1:D:876:GLU:HG2	2.20	0.41
1:D:1085:SER:N	1:D:1155:LEU:HD22	2.35	0.41
1:D:1727:ARG:O	1:D:1731:LEU:HG	2.20	0.41
1:D:3386:GLU:O	1:D:3389:GLU:O	2.38	0.41
1:D:3391:GLU:O	1:D:3395:ARG:HD2	2.19	0.41
1:D:3409:TYR:N	1:D:3410:PRO:HD2	2.35	0.41
1:D:3997:ALA:HB1	1:D:4057:MET:SD	2.60	0.41
1:D:4024:VAL:HG11	1:D:4142:ASN:HB3	2.02	0.41
1:D:4705:VAL:HG23	1:D:4775:TYR:HD2	1.85	0.41
1:C:1992:ALA:O	1:C:1995:THR:OG1	2.29	0.41
1:C:2441:HIS:CE1	1:C:2442:LEU:HD12	2.55	0.41
1:C:2642:LYS:NZ	1:C:2698:MET:HE1	2.36	0.41
1:C:3239:MET:CB	1:C:3239:MET:HG3	2.31	0.41
1:C:4024:VAL:HG11	1:C:4142:ASN:HB3	2.02	0.41
1:A:292:ALA:HB2	1:A:312:THR:HG22	2.02	0.41
1:A:399:GLN:O	1:A:403:MET:HG3	2.21	0.41
1:A:1085:SER:N	1:A:1155:LEU:HD22	2.35	0.41
1:A:1568:LYS:HB2	1:A:1572:ILE:O	2.20	0.41
1:A:1727:ARG:O	1:A:1731:LEU:HG	2.20	0.41
1:A:2208:MET:O	1:A:2212:VAL:HG22	2.19	0.41
1:A:2625:ARG:O	1:A:2628:PHE:HB3	2.21	0.41
1:A:3197:LEU:HD21	1:A:3201:MET:HE1	2.02	0.41
1:A:4069:LYS:O	1:A:4072:VAL:HG22	2.20	0.41
1:A:4187:SER:O	1:A:4188:ARG:HB2	2.20	0.41
1:A:5009:TYR:CZ	1:A:5013:MET:HE3	2.55	0.41
1:B:1999:ARG:NH1	1:B:3636:PHE:CD1	2.76	0.41
1:B:2313:LEU:HB3	1:B:2318:TYR:HB2	2.02	0.41
1:B:2374:SER:OG	1:B:2378:ALA:HB3	2.21	0.41
1:B:2444:GLN:OE1	1:B:2444:GLN:N	2.54	0.41
1:B:3583:GLU:HB3	1:B:3585:ASP:OD1	2.19	0.41
1:B:3610:GLU:OE1	1:B:3611:HIS:CD2	2.73	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3660:ALA:HB3	1:B:3661:TRP:HD1	1.85	0.41
1:B:4677:LEU:HD23	1:B:4711:PHE:HE1	1.83	0.41
1:B:4946:GLN:O	1:B:4950:VAL:HG23	2.19	0.41
1:D:2202:GLY:HA2	1:D:2205:GLU:OE1	2.20	0.41
1:D:2700:MET:O	1:D:2702:CYS:SG	2.78	0.41
1:D:3899:PHE:CE2	1:D:3903:LEU:HD11	2.55	0.41
1:D:4901:ILE:HG13	1:D:4913:ARG:NH2	2.34	0.41
1:C:1676:LEU:HD22	1:C:2167:ILE:HD12	2.03	0.41
1:C:2011:HIS:CE1	1:C:2017:ASP:CG	2.84	0.41
1:C:2703:LEU:HD22	1:C:2706:ILE:HD12	2.02	0.41
1:C:3071:LEU:O	1:C:3075:LEU:HG	2.19	0.41
1:C:3148:ALA:O	1:C:3151:GLN:NE2	2.53	0.41
1:C:3610:GLU:OE1	1:C:3611:HIS:CD2	2.73	0.41
1:C:3628:ARG:HH11	1:C:3632:VAL:HG11	1.85	0.41
1:C:3899:PHE:CE2	1:C:3903:LEU:HD11	2.55	0.41
1:C:4677:LEU:HD23	1:C:4711:PHE:HE1	1.83	0.41
1:C:5000:GLU:HA	1:C:5003:HIS:CD2	2.56	0.41
1:A:501:ALA:O	1:A:505:GLU:HG3	2.20	0.41
1:A:516:LYS:O	1:A:519:VAL:HG22	2.20	0.41
1:A:873:LYS:O	1:A:876:GLU:HG2	2.20	0.41
1:A:2487:GLN:NE2	1:A:2545:GLU:OE1	2.53	0.41
1:A:2977:LEU:HD23	1:A:3056:LEU:HD11	2.02	0.41
1:A:3003:LEU:HB2	1:A:3004:PRO:HD3	2.02	0.41
1:A:3137:LEU:HD23	1:A:3189:ALA:CB	2.50	0.41
2:F:4:VAL:HG22	2:F:74:LEU:HD22	2.03	0.41
1:B:501:ALA:O	1:B:505:GLU:HG3	2.20	0.41
1:B:926:GLY:O	1:B:930:LYS:HG3	2.20	0.41
1:B:1552:VAL:HG11	1:B:1562:ILE:HD13	2.02	0.41
1:B:1992:ALA:O	1:B:1995:THR:OG1	2.29	0.41
1:B:2452:ARG:HG3	1:C:175:SER:O	2.20	0.41
1:B:2545:GLU:HG2	1:B:2546:MET:N	2.35	0.41
1:B:4093:PHE:CE1	1:B:4097:MET:HE2	2.56	0.41
1:D:399:GLN:O	1:D:403:MET:HG3	2.21	0.41
1:D:509:GLU:O	1:D:513:GLU:CD	2.59	0.41
1:D:1088:TRP:O	1:D:1152:MET:HA	2.21	0.41
1:D:2725:LYS:HE3	1:D:2734:ASN:O	2.19	0.41
1:C:2414:ASN:C	1:C:2414:ASN:OD1	2.58	0.41
1:C:3130:THR:O	1:C:3134:VAL:HG23	2.19	0.41
1:C:3386:GLU:O	1:C:3389:GLU:O	2.38	0.41
1:C:3633:VAL:O	1:C:3637:ARG:HD3	2.20	0.41
1:C:4069:LYS:O	1:C:4072:VAL:HG22	2.20	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4187:SER:O	1:C:4188:ARG:HB2	2.20	0.41
1:C:4901:ILE:HG13	1:C:4913:ARG:NH2	2.34	0.41
1:A:1676:LEU:HD22	1:A:2167:ILE:HD12	2.03	0.41
1:A:2330:ARG:HA	1:A:2333:ASP:OD1	2.21	0.41
1:A:3075:LEU:O	1:A:3146:HIS:NE2	2.51	0.41
1:A:3114:LYS:HD3	1:A:3116:SER:H	1.85	0.41
1:A:3409:TYR:N	1:A:3410:PRO:HD2	2.35	0.41
1:A:3435:PHE:CD2	1:A:3524:MET:CE	3.03	0.41
1:A:3660:ALA:HB3	1:A:3661:TRP:HD1	1.85	0.41
1:A:3891:LEU:HD13	1:A:3899:PHE:HZ	1.79	0.41
1:A:4179:GLY:CA	1:A:4197:ILE:HD11	2.50	0.41
1:B:125:ARG:NH1	1:B:125:ARG:CB	2.84	0.41
1:B:221:ARG:NH1	1:B:255:HIS:O	2.54	0.41
1:B:869:ARG:NH2	1:B:947:GLU:OE2	2.52	0.41
1:B:1143:TRP:HB2	1:B:1147:ASP:HB2	2.01	0.41
1:B:1265:ASP:OD1	1:B:1266:THR:N	2.53	0.41
1:B:2612[B]:ARG:NH1	1:B:2613:TYR:OH	2.53	0.41
1:B:2625:ARG:O	1:B:2628:PHE:HB3	2.21	0.41
1:B:3294:PRO:HB2	1:B:3297:PRO:HD2	2.01	0.41
1:B:3458:PHE:CE2	1:B:3464:ILE:HD11	2.54	0.41
1:B:4705:VAL:HG23	1:B:4775:TYR:CD2	2.56	0.41
1:D:885:THR:HA	1:D:888:GLU:OE1	2.20	0.41
1:D:1568:LYS:HB2	1:D:1572:ILE:O	2.21	0.41
1:D:2117:VAL:O	1:D:2120:MET:HG2	2.21	0.41
1:D:2865:VAL:HG11	1:D:2928:LYS:HE2	2.02	0.41
1:D:3380:ARG:NH1	1:D:3391:GLU:OE2	2.50	0.41
1:C:125:ARG:NH1	1:C:125:ARG:CB	2.84	0.41
1:C:2330:ARG:HA	1:C:2333:ASP:OD1	2.21	0.41
1:C:3077:ALA:O	1:C:3081:MET:HG2	2.20	0.41
1:C:3103:ILE:O	1:C:3107:VAL:HG13	2.20	0.41
1:C:3380:ARG:NH2	1:C:3391:GLU:OE2	2.52	0.41
1:C:3997:ALA:HB1	1:C:4057:MET:SD	2.60	0.41
1:C:4767:TRP:CE3	1:C:4768:LEU:CD2	2.99	0.41
1:A:215:THR:HG1	1:A:218:HIS:CE1	2.37	0.41
1:A:631:LEU:HD23	1:A:631:LEU:HA	1.96	0.41
1:A:820:ARG:CZ	1:A:820:ARG:CB	2.80	0.41
1:A:973:SER:O	1:A:976:ARG:NH2	2.54	0.41
1:A:1819:VAL:HG22	1:A:1926:LEU:HD21	2.03	0.41
1:A:2476:ILE:HD11	1:A:2536:LEU:HG	2.03	0.41
1:A:2894:LEU:O	1:A:2897:LYS:HG2	2.21	0.41
1:A:2946:LEU:N	1:A:2948:THR:HG23	2.35	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2969:ILE:O	1:A:2973:PHE:HD1	2.03	0.41
1:A:3197:LEU:HD23	1:A:3197:LEU:O	2.20	0.41
2:H:90:ILE:HD11	1:D:1684:ALA:HA	2.02	0.41
1:B:399:GLN:O	1:B:403:MET:HG3	2.21	0.41
1:B:509:GLU:O	1:B:513:GLU:CD	2.59	0.41
1:B:975:VAL:HG11	1:B:1044:ARG:HA	2.03	0.41
1:B:2011:HIS:CE1	1:B:2017:ASP:CG	2.84	0.41
1:B:3434:LEU:O	1:B:3438:VAL:HG13	2.21	0.41
1:B:3729:MET:HB3	1:B:3770:LEU:HD11	2.03	0.41
1:B:3899:PHE:CE2	1:B:3903:LEU:HD11	2.55	0.41
1:B:3899:PHE:HE2	1:B:3903:LEU:HD11	1.85	0.41
1:D:1676:LEU:HD22	1:D:2167:ILE:HD12	2.03	0.41
1:D:2444:GLN:OE1	1:D:2444:GLN:N	2.54	0.41
1:D:2946:LEU:N	1:D:2948:THR:HG23	2.35	0.41
1:D:2977:LEU:HD23	1:D:3056:LEU:HD11	2.02	0.41
1:D:3628:ARG:HH11	1:D:3632:VAL:HG11	1.85	0.41
1:D:3633:VAL:O	1:D:3637:ARG:HD3	2.20	0.41
1:D:3916:ILE:H	1:D:3916:ILE:HD12	1.85	0.41
1:D:4989:MET:O	1:D:4993:MET:HG3	2.21	0.41
1:C:2202:GLY:HA2	1:C:2205:GLU:OE1	2.20	0.41
1:C:2495:VAL:HG22	1:C:2498:HIS:CE1	2.55	0.41
1:C:3409:TYR:N	1:C:3410:PRO:HD2	2.35	0.41
1:C:3435:PHE:CD2	1:C:3524:MET:HE1	2.56	0.41
1:A:484:LEU:CD1	1:A:526:LEU:CD1	2.98	0.41
1:A:1552:VAL:HG11	1:A:1562:ILE:HD13	2.03	0.41
1:A:2117:VAL:O	1:A:2120:MET:HG2	2.21	0.41
1:A:2513:GLU:OE1	1:A:2513:GLU:N	2.54	0.41
1:A:3077:ALA:O	1:A:3081:MET:HG2	2.20	0.41
1:A:3899:PHE:CE2	1:A:3903:LEU:HD11	2.55	0.41
1:B:69:LEU:CD2	1:B:109:LEU:HD21	2.50	0.41
1:B:1085:SER:N	1:B:1155:LEU:HD22	2.35	0.41
1:B:1568:LYS:HB2	1:B:1572:ILE:O	2.21	0.41
1:B:1973:GLN:NE2	1:B:2005:GLN:OE1	2.44	0.41
1:B:2135:LEU:HD12	1:B:3658:LYS:HG3	2.02	0.41
1:B:2330:ARG:HA	1:B:2333:ASP:OD1	2.21	0.41
1:B:3137:LEU:HD23	1:B:3189:ALA:CB	2.50	0.41
1:B:3433:GLU:HG3	1:B:3437:MET:HE1	2.02	0.41
1:B:3891:LEU:HD13	1:B:3899:PHE:HZ	1.79	0.41
1:B:4930:ALA:CA	1:B:4933:GLN:OE1	2.69	0.41
1:D:278:GLN:HG3	1:D:328:LYS:NZ	2.35	0.41
1:D:501:ALA:O	1:D:505:GLU:HG3	2.20	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:869:ARG:NH2	1:D:947:GLU:OE2	2.52	0.41
1:D:973:SER:O	1:D:976:ARG:NH2	2.54	0.41
1:D:2135:LEU:HD12	1:D:3658:LYS:HG3	2.02	0.41
1:D:2330:ARG:HA	1:D:2333:ASP:OD1	2.21	0.41
1:D:2667:THR:HG21	1:D:2672:LEU:HG	2.02	0.41
1:D:3071:LEU:O	1:D:3075:LEU:HG	2.19	0.41
1:D:3103:ILE:O	1:D:3107:VAL:HG13	2.20	0.41
1:D:4161:ARG:HH11	1:D:4161:ARG:HD2	1.54	0.41
1:D:4563:ARG:NH2	1:D:4815:ASP:OD1	2.53	0.41
1:C:887:ILE:HD13	1:C:887:ILE:HA	1.87	0.41
1:C:1021:LEU:HD23	1:C:1021:LEU:O	2.21	0.41
1:C:3465:ASN:OD1	1:C:3467:MET:CG	2.60	0.41
1:A:160:GLY:O	1:D:3984:ARG:NH1	2.49	0.41
1:A:278:GLN:HG3	1:A:328:LYS:NZ	2.35	0.41
1:A:1088:TRP:O	1:A:1152:MET:HA	2.21	0.41
1:A:1105:ALA:O	1:A:1189:LEU:N	2.50	0.41
1:A:1805:GLU:O	1:A:1808:ARG:HG2	2.21	0.41
1:A:2214:VAL:HG11	1:A:2228:MET:HE3	2.01	0.41
1:A:2442:LEU:HD23	1:A:2447:LYS:CE	2.51	0.41
1:A:2905:LEU:HD23	1:A:2905:LEU:HA	1.98	0.41
1:A:3103:ILE:O	1:A:3107:VAL:HG13	2.20	0.41
1:A:3434:LEU:O	1:A:3438:VAL:HG13	2.21	0.41
1:A:4717:ASP:OD2	1:A:4723:LYS:NZ	2.35	0.41
1:B:2001:PRO:O	1:B:2005:GLN:HG3	2.20	0.41
1:B:2699:ALA:O	1:B:2702:CYS:HB3	2.21	0.41
1:B:3321:ARG:HA	1:B:3324:VAL:HG22	2.02	0.41
1:B:3892:CYS:HG	1:B:3899:PHE:HD2	1.69	0.41
1:B:4999:ASP:OD1	1:B:5000:GLU:N	2.53	0.41
1:B:5000:GLU:HA	1:B:5003:HIS:CD2	2.56	0.41
1:D:468:LEU:O	1:D:472:ARG:HG2	2.20	0.41
1:D:484:LEU:CD1	1:D:526:LEU:CD1	2.98	0.41
1:D:553:ARG:O	1:D:557:SER:OG	2.29	0.41
1:D:926:GLY:O	1:D:930:LYS:HG3	2.20	0.41
1:D:1436:SER:OG	1:D:1565:GLU:HB2	2.20	0.41
1:D:1805:GLU:O	1:D:1808:ARG:HG2	2.21	0.41
1:D:1821:ASP:OD1	1:D:1821:ASP:C	2.59	0.41
1:D:1973:GLN:NE2	1:D:2005:GLN:OE1	2.44	0.41
1:D:2703:LEU:HD22	1:D:2706:ILE:HD12	2.02	0.41
1:D:3077:ALA:O	1:D:3081:MET:HG2	2.20	0.41
1:D:3114:LYS:HD3	1:D:3116:SER:H	1.85	0.41
1:D:3435:PHE:CD2	1:D:3524:MET:CE	3.03	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3604:TYR:O	1:D:3608:GLN:OE1	2.39	0.41
1:D:4179:GLY:HA3	1:D:4197:ILE:HD11	2.02	0.41
1:D:4767:TRP:HE3	1:D:4768:LEU:HD23	1.80	0.41
1:D:5000:GLU:HA	1:D:5003:HIS:CD2	2.56	0.41
1:C:874:LEU:HD21	1:C:1046:LEU:HG	2.03	0.41
1:C:885:THR:HA	1:C:888:GLU:OE1	2.20	0.41
1:C:1085:SER:N	1:C:1155:LEU:HD22	2.35	0.41
1:C:1568:LYS:HB2	1:C:1572:ILE:O	2.20	0.41
1:C:2374:SER:OG	1:C:2378:ALA:HB3	2.20	0.41
1:C:3111:ARG:HH21	1:C:3178:THR:CB	2.33	0.41
1:C:3336:LYS:HG3	1:C:3337:ARG:N	2.36	0.41
1:C:4705:VAL:HG23	1:C:4775:TYR:CD2	2.56	0.41
1:C:4989:MET:O	1:C:4993:MET:HG3	2.21	0.41
1:A:69:LEU:CD2	1:A:109:LEU:HD21	2.50	0.41
1:A:117:TYR:O	1:A:138:GLN:N	2.51	0.41
1:A:160:GLY:O	1:D:3984:ARG:NH2	2.52	0.41
1:A:721:LEU:HD12	1:A:1476:MET:HE2	2.01	0.41
1:A:823:LEU:HD13	1:A:1626:TRP:CE3	2.56	0.41
1:A:926:GLY:O	1:A:930:LYS:HG3	2.20	0.41
1:A:1021:LEU:O	1:A:1021:LEU:HD23	2.21	0.41
1:A:2202:GLY:HA2	1:A:2205:GLU:OE1	2.20	0.41
1:A:2313:LEU:HB3	1:A:2318:TYR:HB2	2.02	0.41
1:A:2725:LYS:HE3	1:A:2734:ASN:O	2.19	0.41
1:A:2865:VAL:HG11	1:A:2928:LYS:HE2	2.02	0.41
1:A:3336:LYS:HG3	1:A:3337:ARG:N	2.36	0.41
1:A:3384:LYS:H	1:A:3387:ALA:HB3	1.86	0.41
1:A:3458:PHE:HE2	1:A:3464:ILE:HD11	1.82	0.41
1:A:3573:MET:CB	1:A:3577:ARG:NH1	2.73	0.41
1:A:3604:TYR:O	1:A:3608:GLN:OE1	2.39	0.41
1:A:3628:ARG:HH11	1:A:3632:VAL:HG11	1.85	0.41
1:A:3633:VAL:O	1:A:3637:ARG:HD3	2.20	0.41
1:A:3969:ILE:HD11	1:A:4029:SER:HB2	2.02	0.41
1:A:4024:VAL:HG11	1:A:4142:ASN:HB3	2.02	0.41
1:A:4839:MET:CE	1:D:4820:VAL:HG11	2.50	0.41
1:A:5000:GLU:HA	1:A:5003:HIS:CD2	2.56	0.41
2:H:24:VAL:HG21	2:H:59:TRP:HZ3	1.85	0.41
1:B:117:TYR:O	1:B:138:GLN:N	2.51	0.41
1:B:365:LYS:HD3	1:B:369:LEU:HD21	2.03	0.41
1:B:823:LEU:HD13	1:B:1626:TRP:CE3	2.56	0.41
1:B:885:THR:HA	1:B:888:GLU:OE1	2.21	0.41
1:B:973:SER:O	1:B:976:ARG:NH2	2.54	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:985:VAL:HG13	1:B:1039:LEU:HD23	2.00	0.41
1:B:1096:THR:O	1:B:1100:MET:HE1	2.21	0.41
1:B:2441:HIS:CE1	1:B:2442:LEU:HD12	2.55	0.41
1:B:2442:LEU:HD23	1:B:2447:LYS:CE	2.51	0.41
1:B:2476:ILE:HD11	1:B:2536:LEU:HG	2.02	0.41
1:B:2946:LEU:N	1:B:2948:THR:HG23	2.35	0.41
1:B:2969:ILE:O	1:B:2973:PHE:HD1	2.02	0.41
1:B:3062:PRO:O	1:B:3065:VAL:HG22	2.21	0.41
1:B:3101:GLU:HA	1:B:3104:GLU:OE1	2.21	0.41
1:B:3114:LYS:HD3	1:B:3116:SER:H	1.85	0.41
1:B:3336:LYS:HG3	1:B:3337:ARG:N	2.36	0.41
1:B:3384:LYS:H	1:B:3387:ALA:HB3	1.86	0.41
1:B:3386:GLU:O	1:B:3389:GLU:O	2.38	0.41
1:B:3435:PHE:CD2	1:B:3524:MET:HE1	2.56	0.41
1:B:3628:ARG:HH11	1:B:3632:VAL:HG11	1.85	0.41
1:B:3969:ILE:HD11	1:B:4029:SER:HB2	2.02	0.41
1:B:4075:GLU:CD	1:C:4736:ARG:NH1	2.73	0.41
1:D:130:LYS:NZ	1:C:2373:GLY:O	2.50	0.41
1:D:874:LEU:HD21	1:D:1046:LEU:HG	2.03	0.41
1:D:877:ASN:O	1:D:880:GLU:CG	2.69	0.41
1:D:1712:TYR:CG	1:D:1840:PRO:HB3	2.56	0.41
1:D:2374:SER:OG	1:D:2378:ALA:HB3	2.21	0.41
1:D:2442:LEU:HD23	1:D:2447:LYS:CE	2.51	0.41
1:D:2487:GLN:NE2	1:D:2545:GLU:OE1	2.53	0.41
1:D:3111:ARG:HH21	1:D:3178:THR:CB	2.33	0.41
1:D:3111:ARG:O	1:D:3112:LEU:HB2	2.20	0.41
1:D:3380:ARG:NH2	1:D:3391:GLU:OE2	2.52	0.41
1:D:3434:LEU:O	1:D:3438:VAL:HG13	2.21	0.41
1:D:3545:THR:HG23	1:D:3548:GLU:H	1.86	0.41
1:D:4705:VAL:HG23	1:D:4775:TYR:CD2	2.56	0.41
1:C:215:THR:HG1	1:C:218:HIS:CE1	2.39	0.41
1:C:399:GLN:O	1:C:403:MET:HG3	2.21	0.41
1:C:985:VAL:HG13	1:C:1039:LEU:HD23	2.00	0.41
1:C:1088:TRP:O	1:C:1152:MET:HA	2.21	0.41
1:C:2212:VAL:HG21	1:C:2256:TYR:HH	1.79	0.41
1:C:2442:LEU:HD23	1:C:2447:LYS:CE	2.51	0.41
1:C:2444:GLN:OE1	1:C:2444:GLN:N	2.53	0.41
1:C:2625:ARG:O	1:C:2628:PHE:HB3	2.21	0.41
1:C:2762:THR:HA	1:C:2765:LYS:HE2	2.03	0.41
1:C:3062:PRO:O	1:C:3065:VAL:HG22	2.21	0.41
1:C:3101:GLU:HA	1:C:3104:GLU:OE1	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3111:ARG:O	1:C:3112:LEU:HB2	2.20	0.41
1:C:3434:LEU:O	1:C:3438:VAL:HG13	2.21	0.41
1:A:221:ARG:NH1	1:A:255:HIS:O	2.54	0.41
1:A:1926:LEU:HA	1:A:1929:MET:SD	2.61	0.41
1:A:2369[B]:ARG:HD3	1:A:2369[B]:ARG:HA	1.92	0.41
1:A:2441:HIS:CE1	1:A:2442:LEU:HD12	2.55	0.41
1:A:3545:THR:HG23	1:A:3548:GLU:H	1.86	0.41
1:A:4705:VAL:HG23	1:A:4775:TYR:CD2	2.56	0.41
1:B:866:HIS:CD2	1:B:867:LEU:CD2	3.04	0.41
1:B:1980:LEU:HD21	1:B:1995:THR:HG23	2.02	0.41
1:B:2762:THR:HA	1:B:2765:LYS:HE2	2.03	0.41
1:B:2792:ARG:HB3	1:B:2793:PRO:HD2	2.03	0.41
1:B:3197:LEU:HD21	1:B:3201:MET:HE1	2.02	0.41
1:D:975:VAL:HG11	1:D:1044:ARG:HA	2.03	0.41
1:D:1096:THR:O	1:D:1100:MET:HE1	2.20	0.41
1:D:1241:SER:HA	1:D:1603:VAL:HG22	2.03	0.41
1:D:1478:ASP:HB2	1:D:1484:HIS:CE1	2.56	0.41
1:D:1748:PHE:HA	1:D:1749:PRO:HD3	1.92	0.41
1:D:2625:ARG:O	1:D:2628:PHE:HB3	2.21	0.41
1:D:3336:LYS:HG3	1:D:3337:ARG:N	2.36	0.41
1:D:3666:ASP:OD1	1:D:3666:ASP:O	2.39	0.41
1:C:501:ALA:O	1:C:505:GLU:HG3	2.20	0.41
1:C:516:LYS:O	1:C:519:VAL:HG22	2.20	0.41
1:C:615:ARG:NH2	1:C:1676:LEU:O	2.50	0.41
1:C:879:HIS:HE1	1:C:913:LEU:HD13	1.86	0.41
1:C:973:SER:O	1:C:976:ARG:NH2	2.54	0.41
1:C:1241:SER:HA	1:C:1603:VAL:HG22	2.03	0.41
1:C:1712:TYR:CG	1:C:1840:PRO:HB3	2.56	0.41
1:C:3244:PRO:HD2	1:C:3249:LEU:HD21	2.03	0.41
1:C:3892:CYS:HG	1:C:3899:PHE:HD2	1.69	0.41
1:C:3899:PHE:HE2	1:C:3903:LEU:HD11	1.85	0.41
1:C:4705:VAL:HG23	1:C:4775:TYR:HD2	1.85	0.41
1:A:125:ARG:NH1	1:A:125:ARG:CB	2.84	0.40
1:A:515:TRP:O	1:A:519:VAL:HG13	2.21	0.40
1:A:1241:SER:HA	1:A:1603:VAL:HG22	2.03	0.40
1:A:3101:GLU:HA	1:A:3104:GLU:OE1	2.21	0.40
1:A:3111:ARG:HH21	1:A:3178:THR:CB	2.33	0.40
1:A:3666:ASP:O	1:A:3666:ASP:OD1	2.40	0.40
1:A:3729:MET:HB3	1:A:3770:LEU:HD11	2.03	0.40
1:A:3868:ARG:HG2	1:A:3869:GLN:N	2.34	0.40
1:A:4748:LEU:O	1:A:4751:THR:OG1	2.35	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4876:CYS:HA	1:A:4882:CYS:HB2	2.04	0.40
2:E:4:VAL:HG22	2:E:74:LEU:HD22	2.03	0.40
2:G:50:LEU:HA	2:G:50:LEU:HD12	1.89	0.40
1:B:1478:ASP:HB2	1:B:1484:HIS:CE1	2.56	0.40
1:B:1984:PHE:HA	1:B:1991:THR:HG21	2.03	0.40
1:B:2784:GLU:OE2	1:B:2785:LEU:HD12	2.21	0.40
1:B:3302:PRO:HA	1:B:3303:PRO:HD3	2.00	0.40
1:D:736:HIS:NE2	1:D:742:ASP:OD2	2.55	0.40
1:D:2319:PRO:HG2	1:D:2392:ARG:HA	2.03	0.40
1:D:3062:PRO:O	1:D:3065:VAL:HG22	2.21	0.40
1:D:3244:PRO:HD2	1:D:3249:LEU:HD21	2.03	0.40
1:D:4093:PHE:CE1	1:D:4097:MET:HE2	2.56	0.40
1:D:4187:SER:O	1:D:4188:ARG:HB2	2.20	0.40
1:C:221:ARG:NH1	1:C:255:HIS:O	2.54	0.40
1:C:626:LEU:N	1:C:627:PRO:HD2	2.37	0.40
1:C:820:ARG:CZ	1:C:820:ARG:CB	2.80	0.40
1:C:869:ARG:NH2	1:C:947:GLU:OE2	2.52	0.40
1:C:1819:VAL:HG22	1:C:1926:LEU:HD21	2.03	0.40
1:C:1856:ASP:O	1:C:1860:LYS:HG3	2.21	0.40
1:C:3321:ARG:HA	1:C:3324:VAL:HG22	2.02	0.40
1:C:3868:ARG:HG2	1:C:3869:GLN:N	2.34	0.40
1:A:1008:SER:HB3	1:A:1017:ARG:HB3	2.03	0.40
1:A:1821:ASP:OD1	1:A:1821:ASP:C	2.59	0.40
1:A:2319:PRO:HG2	1:A:2392:ARG:HA	2.03	0.40
1:A:2699:ALA:O	1:A:2702:CYS:HB3	2.21	0.40
1:A:3622:LYS:HB2	1:A:3625:SER:HB3	2.03	0.40
1:A:3916:ILE:H	1:A:3916:ILE:HD12	1.85	0.40
1:A:4989:MET:O	1:A:4993:MET:HG3	2.21	0.40
1:B:215:THR:HG1	1:B:218:HIS:CE1	2.39	0.40
1:B:891:TRP:CE3	1:B:903:LEU:HA	2.56	0.40
1:B:1021:LEU:O	1:B:1021:LEU:HD23	2.21	0.40
1:B:2319:PRO:HG2	1:B:2392:ARG:HA	2.03	0.40
1:B:2894:LEU:O	1:B:2897:LYS:HG2	2.21	0.40
1:B:4024:VAL:HG11	1:B:4142:ASN:HB3	2.03	0.40
1:D:125:ARG:NH1	1:D:125:ARG:CB	2.84	0.40
1:D:906:CYS:HA	1:D:914:PRO:HD3	2.04	0.40
1:D:1984:PHE:HA	1:D:1991:THR:HG21	2.03	0.40
1:D:2001:PRO:O	1:D:2005:GLN:HG3	2.20	0.40
1:D:2476:ILE:HD11	1:D:2536:LEU:HG	2.03	0.40
1:D:3183:VAL:HG12	1:D:3187:ARG:HE	1.87	0.40
1:C:309:THR:O	1:C:309:THR:CG2	2.68	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2513:GLU:OE1	1:C:2513:GLU:N	2.54	0.40
1:C:2699:ALA:O	1:C:2702:CYS:HB3	2.21	0.40
1:C:3114:LYS:HD3	1:C:3116:SER:H	1.86	0.40
1:C:3916:ILE:H	1:C:3916:ILE:HD12	1.85	0.40
1:A:309:THR:O	1:A:309:THR:CG2	2.68	0.40
1:A:2135:LEU:HD12	1:A:3658:LYS:HG3	2.02	0.40
1:A:2240:CYS:SG	1:A:2250:MET:HG3	2.62	0.40
1:A:2784:GLU:OE2	1:A:2785:LEU:HD12	2.21	0.40
1:A:3458:PHE:CE2	1:A:3464:ILE:HD11	2.54	0.40
1:B:221:ARG:NE	1:B:253:CYS:O	2.55	0.40
1:B:515:TRP:O	1:B:519:VAL:HG13	2.21	0.40
1:B:736:HIS:NE2	1:B:742:ASP:OD2	2.55	0.40
1:B:993:HIS:NE2	1:B:1022:VAL:O	2.53	0.40
1:B:1241:SER:HA	1:B:1603:VAL:HG22	2.03	0.40
1:B:1926:LEU:HA	1:B:1929:MET:SD	2.61	0.40
1:B:2117:VAL:O	1:B:2120:MET:HG2	2.21	0.40
1:B:2240:CYS:SG	1:B:2250:MET:HG3	2.62	0.40
1:B:2513:GLU:N	1:B:2513:GLU:OE1	2.54	0.40
1:B:3111:ARG:O	1:B:3112:LEU:HB2	2.20	0.40
1:B:3409:TYR:N	1:B:3410:PRO:HD2	2.35	0.40
1:D:870:ILE:HD12	1:D:870:ILE:HA	1.98	0.40
1:D:993:HIS:NE2	1:D:1022:VAL:O	2.53	0.40
1:D:1926:LEU:HA	1:D:1929:MET:SD	2.61	0.40
1:D:2240:CYS:SG	1:D:2250:MET:HG3	2.62	0.40
1:D:2969:ILE:O	1:D:2973:PHE:HD1	2.02	0.40
1:D:3003:LEU:HB2	1:D:3004:PRO:HD3	2.02	0.40
1:D:3532:LEU:HD23	1:D:3535:LEU:HD12	2.03	0.40
1:D:4003:LEU:HA	1:D:4009:GLN:OE1	2.22	0.40
1:C:667:MET:SD	1:C:801:LYS:NZ	2.86	0.40
1:C:891:TRP:CE3	1:C:903:LEU:HA	2.57	0.40
1:C:893:TYR:CD1	1:C:907:LEU:O	2.75	0.40
1:C:893:TYR:HD1	1:C:907:LEU:O	2.03	0.40
1:C:2476:ILE:HD11	1:C:2536:LEU:HG	2.03	0.40
1:C:2638:LYS:O	1:C:2698:MET:HE2	2.22	0.40
1:C:3254:GLY:C	1:C:3258:GLU:OE1	2.60	0.40
1:C:3532:LEU:HD23	1:C:3535:LEU:HD12	2.02	0.40
1:A:350:HIS:O	1:A:354:GLY:N	2.49	0.40
1:A:874:LEU:HD21	1:A:1046:LEU:HG	2.03	0.40
1:A:975:VAL:HG11	1:A:1044:ARG:HA	2.03	0.40
1:A:1478:ASP:HB2	1:A:1484:HIS:CE1	2.56	0.40
1:A:1712:TYR:CG	1:A:1840:PRO:HB3	2.56	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2630:VAL:N	1:A:2631:PRO:CD	2.85	0.40
1:A:3239:MET:CB	1:A:3239:MET:HG3	2.31	0.40
1:A:3254:GLY:C	1:A:3258:GLU:OE1	2.60	0.40
1:B:1821:ASP:OD1	1:B:1821:ASP:C	2.59	0.40
1:B:4844:LEU:O	1:B:4847:VAL:HG22	2.21	0.40
1:B:4989:MET:O	1:B:4993:MET:HG3	2.21	0.40
1:D:167:ASP:N	1:C:384:MET:SD	2.95	0.40
1:D:823:LEU:HD13	1:D:1626:TRP:CE3	2.56	0.40
1:D:899:ASP:O	1:D:902:ARG:NE	2.45	0.40
1:D:3969:ILE:HD11	1:D:4029:SER:HB2	2.02	0.40
1:D:4666:VAL:N	1:D:4667:PRO:CD	2.85	0.40
1:D:4844:LEU:O	1:D:4847:VAL:HG22	2.21	0.40
1:C:866:HIS:CD2	1:C:867:LEU:CD2	3.04	0.40
1:C:1805:GLU:O	1:C:1808:ARG:HG2	2.21	0.40
1:C:2319:PRO:HG2	1:C:2392:ARG:HA	2.03	0.40
1:C:2520:HIS:HA	1:C:2523:ASP:OD2	2.22	0.40
1:C:2784:GLU:OE2	1:C:2785:LEU:HD12	2.21	0.40
1:C:3003:LEU:HB2	1:C:3004:PRO:HD3	2.02	0.40
1:C:3545:THR:HG23	1:C:3548:GLU:H	1.86	0.40
1:C:3729:MET:HB3	1:C:3770:LEU:HD11	2.03	0.40
1:A:736:HIS:NE2	1:A:742:ASP:OD2	2.55	0.40
1:A:866:HIS:CD2	1:A:867:LEU:CD2	3.04	0.40
1:A:891:TRP:CE3	1:A:903:LEU:HA	2.56	0.40
1:A:973:SER:O	1:A:976:ARG:CZ	2.70	0.40
1:A:2165:LEU:HD21	1:A:2177:LEU:HD22	2.04	0.40
1:A:3239:MET:CA	1:A:3239:MET:HG2	2.46	0.40
1:A:3254:GLY:O	1:A:3255:GLY:C	2.60	0.40
1:A:4767:TRP:HE3	1:A:4768:LEU:HD23	1.80	0.40
1:A:4848:VAL:HG13	1:A:4886:HIS:HB3	2.03	0.40
1:A:4951:LYS:O	1:A:4955:GLU:OE1	2.40	0.40
1:B:1698:LEU:HD22	1:B:1712:TYR:CE2	2.57	0.40
1:B:2888:ARG:O	1:B:2892:GLN:HG2	2.22	0.40
1:B:3548:GLU:O	1:B:3551:GLU:HG3	2.22	0.40
1:B:3622:LYS:HB2	1:B:3625:SER:HB3	2.03	0.40
1:B:4118:ASP:OD1	1:B:4121:GLU:N	2.55	0.40
1:B:4904:PRO:HB3	1:B:4913:ARG:HD3	2.04	0.40
1:B:4937:ILE:HG12	1:C:4934:GLY:HA2	2.03	0.40
1:D:2442:LEU:HD23	1:D:2447:LYS:HE3	2.03	0.40
1:D:2784:GLU:OE2	1:D:2785:LEU:HD12	2.21	0.40
1:D:3101:GLU:HA	1:D:3104:GLU:OE1	2.21	0.40
1:D:3433:GLU:HG3	1:D:3437:MET:HE1	2.04	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4754:ASN:OD1	1:D:4755:GLU:N	2.55	0.40
1:D:4848:VAL:HG13	1:D:4886:HIS:HB3	2.03	0.40
1:C:278:GLN:HG3	1:C:328:LYS:NZ	2.35	0.40
1:C:365:LYS:HD3	1:C:369:LEU:HD21	2.03	0.40
1:C:4003:LEU:HA	1:C:4009:GLN:OE1	2.22	0.40
1:C:4666:VAL:N	1:C:4667:PRO:CD	2.85	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	4385/5037 (87%)	4268 (97%)	109 (2%)	8 (0%)	44	66
1	B	4385/5037 (87%)	4268 (97%)	109 (2%)	8 (0%)	44	66
1	C	4385/5037 (87%)	4268 (97%)	109 (2%)	8 (0%)	44	66
1	D	4385/5037 (87%)	4268 (97%)	109 (2%)	8 (0%)	44	66
2	E	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
2	F	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
2	G	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
2	H	105/108 (97%)	102 (97%)	3 (3%)	0	100	100
All	All	17960/20580 (87%)	17480 (97%)	448 (2%)	32 (0%)	45	66

All (32) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	908	VAL
1	A	2829	GLY
1	A	3239	MET
1	A	3300	ALA

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	B	908	VAL
1	B	2829	GLY
1	B	3239	MET
1	B	3300	ALA
1	D	908	VAL
1	D	2829	GLY
1	D	3239	MET
1	D	3300	ALA
1	C	908	VAL
1	C	2829	GLY
1	C	3239	MET
1	C	3300	ALA
1	A	55	ALA
1	A	1930	LYS
1	B	55	ALA
1	B	1930	LYS
1	D	55	ALA
1	D	1930	LYS
1	C	55	ALA
1	C	1930	LYS
1	A	1036	ARG
1	B	1036	ARG
1	D	1036	ARG
1	C	1036	ARG
1	A	361	ALA
1	B	361	ALA
1	D	361	ALA
1	C	361	ALA

5.3.2 Protein sidechains [i](#)

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	3836/4276 (90%)	3816 (100%)	20 (0%)	86 95
1	B	3836/4276 (90%)	3816 (100%)	20 (0%)	86 95
1	C	3836/4276 (90%)	3816 (100%)	20 (0%)	86 95

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	D	3836/4276 (90%)	3815 (100%)	21 (0%)	86	95
2	E	89/90 (99%)	89 (100%)	0	100	100
2	F	89/90 (99%)	89 (100%)	0	100	100
2	G	89/90 (99%)	89 (100%)	0	100	100
2	H	89/90 (99%)	89 (100%)	0	100	100
All	All	15700/17464 (90%)	15619 (100%)	81 (0%)	85	95

All (81) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	155	LYS
1	A	384	MET
1	A	820	ARG
1	A	945	LYS
1	A	1036	ARG
1	A	1128	ARG
1	A	1559	GLN
1	A	1618	ARG
1	A	1759	ARG
1	A	2177	LEU
1	A	2267	MET
1	A	2308	GLN
1	A	2336	ARG
1	A	2444	GLN
1	A	3014	CYS
1	A	3023	LYS
1	A	3185	LYS
1	A	3477	LYS
1	A	3557	LEU
1	A	3617	LYS
1	B	155	LYS
1	B	384	MET
1	B	820	ARG
1	B	945	LYS
1	B	1036	ARG
1	B	1128	ARG
1	B	1559	GLN
1	B	1618	ARG
1	B	1759	ARG
1	B	2177	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	B	2267	MET
1	B	2308	GLN
1	B	2336	ARG
1	B	2444	GLN
1	B	3014	CYS
1	B	3023	LYS
1	B	3185	LYS
1	B	3477	LYS
1	B	3557	LEU
1	B	3617	LYS
1	D	155	LYS
1	D	384	MET
1	D	820	ARG
1	D	945	LYS
1	D	1036	ARG
1	D	1128	ARG
1	D	1559	GLN
1	D	1618	ARG
1	D	1759	ARG
1	D	2177	LEU
1	D	2267	MET
1	D	2308	GLN
1	D	2336	ARG
1	D	2444	GLN
1	D	3014	CYS
1	D	3023	LYS
1	D	3185	LYS
1	D	3239	MET
1	D	3477	LYS
1	D	3557	LEU
1	D	3617	LYS
1	C	155	LYS
1	C	384	MET
1	C	820	ARG
1	C	945	LYS
1	C	1036	ARG
1	C	1128	ARG
1	C	1559	GLN
1	C	1618	ARG
1	C	1759	ARG
1	C	2177	LEU
1	C	2267	MET

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	C	2308	GLN
1	C	2336	ARG
1	C	2444	GLN
1	C	3014	CYS
1	C	3023	LYS
1	C	3185	LYS
1	C	3477	LYS
1	C	3557	LEU
1	C	3617	LYS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (48) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	105	HIS
1	A	879	HIS
1	A	889	GLN
1	A	963	ASN
1	A	1041	GLN
1	A	1299	GLN
1	A	2011	HIS
1	A	2161	GLN
1	A	2734	ASN
1	A	3605	HIS
1	A	3611	HIS
1	A	5003	HIS
1	B	105	HIS
1	B	879	HIS
1	B	889	GLN
1	B	963	ASN
1	B	1041	GLN
1	B	1299	GLN
1	B	2011	HIS
1	B	2161	GLN
1	B	2734	ASN
1	B	3605	HIS
1	B	3611	HIS
1	B	5003	HIS
1	D	105	HIS
1	D	879	HIS
1	D	889	GLN
1	D	963	ASN
1	D	1041	GLN

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	D	1299	GLN
1	D	2011	HIS
1	D	2161	GLN
1	D	2734	ASN
1	D	3605	HIS
1	D	3611	HIS
1	D	5003	HIS
1	C	105	HIS
1	C	879	HIS
1	C	889	GLN
1	C	963	ASN
1	C	1041	GLN
1	C	1299	GLN
1	C	2011	HIS
1	C	2161	GLN
1	C	2734	ASN
1	C	3605	HIS
1	C	3611	HIS
1	C	5003	HIS

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	B	5301	-	28,33,33	0.63	0	34,52,52	1.04	4 (11%)
3	ATP	D	5301	-	28,33,33	0.63	0	34,52,52	1.04	4 (11%)
6	PNX	C	5304	-	13,21,21	0.79	1 (7%)	13,30,30	2.92	3 (23%)
6	PNX	A	5304	-	13,21,21	0.79	1 (7%)	13,30,30	2.92	3 (23%)
3	ATP	A	5301	-	28,33,33	0.63	0	34,52,52	1.03	4 (11%)
6	PNX	B	5304	-	13,21,21	0.79	0	13,30,30	2.91	3 (23%)
6	PNX	D	5304	-	13,21,21	0.81	1 (7%)	13,30,30	2.94	3 (23%)
3	ATP	C	5301	-	28,33,33	0.63	0	34,52,52	1.03	4 (11%)

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	B	5301	-	-	4/18/38/38	0/3/3/3
3	ATP	D	5301	-	-	4/18/38/38	0/3/3/3
6	PNX	C	5304	-	-	5/7/7/7	0/2/2/2
6	PNX	A	5304	-	-	5/7/7/7	0/2/2/2
3	ATP	A	5301	-	-	4/18/38/38	0/3/3/3
6	PNX	B	5304	-	-	5/7/7/7	0/2/2/2
6	PNX	D	5304	-	-	5/7/7/7	0/2/2/2
3	ATP	C	5301	-	-	4/18/38/38	0/3/3/3

All (3) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
6	D	5304	PNX	C6-N1	2.08	1.41	1.38
6	C	5304	PNX	C6-N1	2.04	1.41	1.38
6	A	5304	PNX	C6-N1	2.02	1.41	1.38

All (28) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	D	5304	PNX	C5-C6-N1	-9.08	111.31	120.27

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	C	5304	PNX	C5-C6-N1	-9.03	111.36	120.27
6	B	5304	PNX	C5-C6-N1	-9.03	111.37	120.27
6	A	5304	PNX	C5-C6-N1	-9.02	111.37	120.27
6	D	5304	PNX	C4-C5-C6	4.46	123.36	119.96
6	A	5304	PNX	C4-C5-C6	4.44	123.34	119.96
6	C	5304	PNX	C4-C5-C6	4.43	123.33	119.96
6	B	5304	PNX	C4-C5-C6	4.35	123.27	119.96
6	A	5304	PNX	CAC-N3-C4	2.88	121.94	118.20
6	D	5304	PNX	CAC-N3-C4	2.88	121.93	118.20
6	B	5304	PNX	CAC-N3-C4	2.85	121.90	118.20
6	C	5304	PNX	CAC-N3-C4	2.84	121.89	118.20
3	D	5301	ATP	C5-C6-N6	2.39	123.95	120.31
3	B	5301	ATP	O4'-C1'-N9	-2.36	105.61	108.75
3	B	5301	ATP	C5-C6-N6	2.36	123.91	120.31
3	A	5301	ATP	O4'-C1'-N9	-2.36	105.62	108.75
3	D	5301	ATP	O4'-C1'-N9	-2.35	105.62	108.75
3	C	5301	ATP	C5-C6-N6	2.35	123.90	120.31
3	A	5301	ATP	C5-C6-N6	2.35	123.89	120.31
3	C	5301	ATP	O4'-C1'-N9	-2.35	105.63	108.75
3	D	5301	ATP	O3'-C3'-C4'	-2.09	105.09	111.08
3	C	5301	ATP	O3'-C3'-C4'	-2.07	105.12	111.08
3	A	5301	ATP	O3'-C3'-C4'	-2.07	105.13	111.08
3	B	5301	ATP	O3'-C3'-C4'	-2.06	105.15	111.08
3	C	5301	ATP	O3'-C3'-C2'	-2.05	105.25	111.82
3	B	5301	ATP	O3'-C3'-C2'	-2.05	105.26	111.82
3	A	5301	ATP	O3'-C3'-C2'	-2.05	105.26	111.82
3	D	5301	ATP	O3'-C3'-C2'	-2.04	105.27	111.82

There are no chirality outliers.

All (36) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
3	A	5301	ATP	PB-O3B-PG-O3G
3	B	5301	ATP	PB-O3B-PG-O3G
3	D	5301	ATP	PB-O3B-PG-O3G
3	C	5301	ATP	PB-O3B-PG-O3G
6	A	5304	PNX	CAH-CAI-CAK-N1
6	B	5304	PNX	CAH-CAI-CAK-N1
6	D	5304	PNX	CAH-CAI-CAK-N1
6	C	5304	PNX	CAH-CAI-CAK-N1
6	A	5304	PNX	CAI-CAH-CAJ-CAM
6	B	5304	PNX	CAI-CAH-CAJ-CAM

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
6	D	5304	PNX	CAI-CAH-CAJ-CAM
6	C	5304	PNX	CAI-CAH-CAJ-CAM
6	A	5304	PNX	CAH-CAJ-CAM-CAA
6	A	5304	PNX	CAH-CAJ-CAM-OAD
6	B	5304	PNX	CAH-CAJ-CAM-CAA
6	B	5304	PNX	CAH-CAJ-CAM-OAD
6	D	5304	PNX	CAH-CAJ-CAM-CAA
6	D	5304	PNX	CAH-CAJ-CAM-OAD
6	C	5304	PNX	CAH-CAJ-CAM-CAA
6	C	5304	PNX	CAH-CAJ-CAM-OAD
3	A	5301	ATP	C3'-C4'-C5'-O5'
3	B	5301	ATP	C3'-C4'-C5'-O5'
3	D	5301	ATP	C3'-C4'-C5'-O5'
3	C	5301	ATP	C3'-C4'-C5'-O5'
6	D	5304	PNX	CAJ-CAH-CAI-CAK
6	B	5304	PNX	CAJ-CAH-CAI-CAK
6	C	5304	PNX	CAJ-CAH-CAI-CAK
6	A	5304	PNX	CAJ-CAH-CAI-CAK
3	A	5301	ATP	O4'-C4'-C5'-O5'
3	B	5301	ATP	O4'-C4'-C5'-O5'
3	D	5301	ATP	O4'-C4'-C5'-O5'
3	C	5301	ATP	O4'-C4'-C5'-O5'
3	A	5301	ATP	PG-O3B-PB-O1B
3	B	5301	ATP	PG-O3B-PB-O1B
3	D	5301	ATP	PG-O3B-PB-O1B
3	C	5301	ATP	PG-O3B-PB-O1B

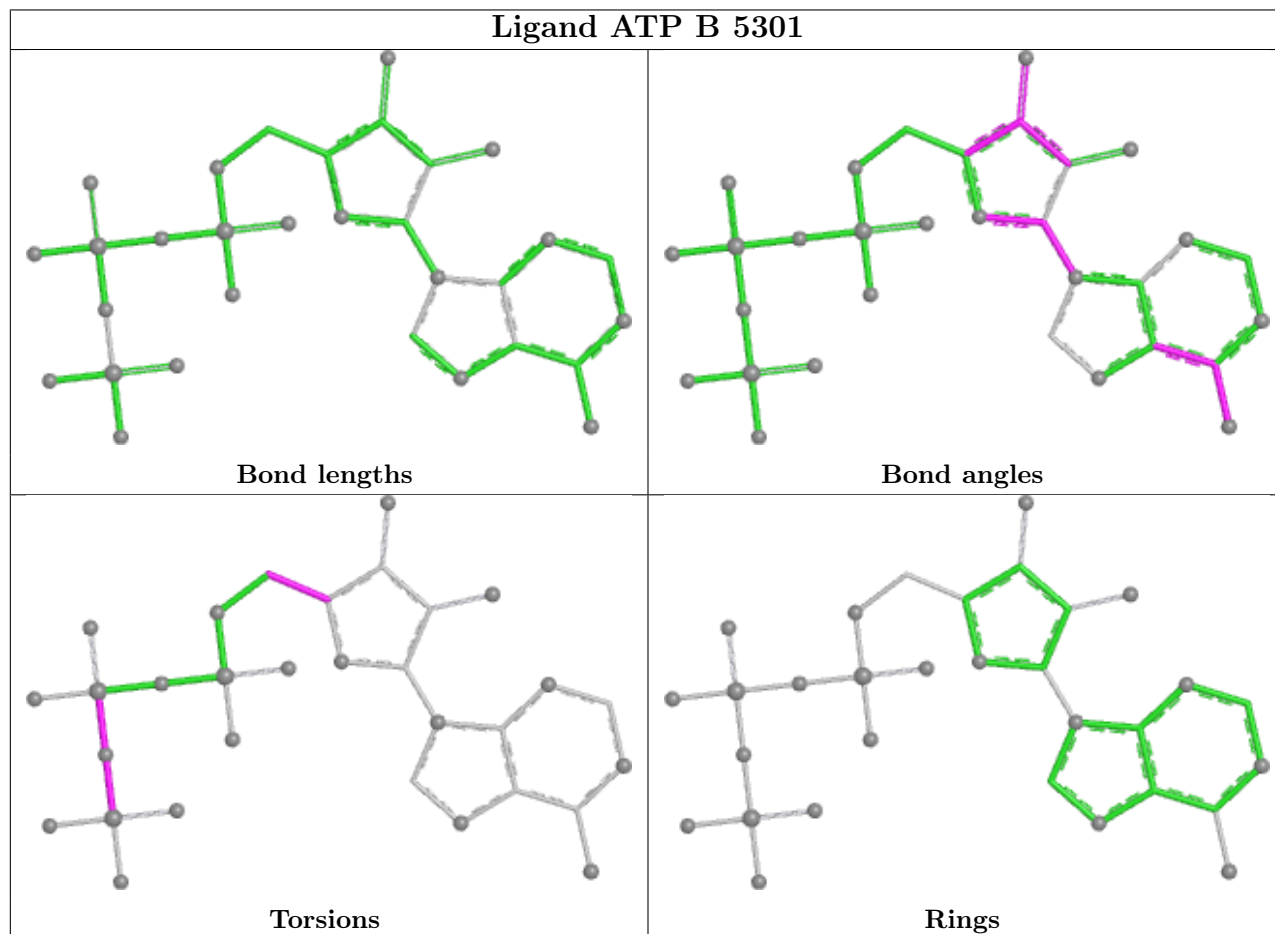
There are no ring outliers.

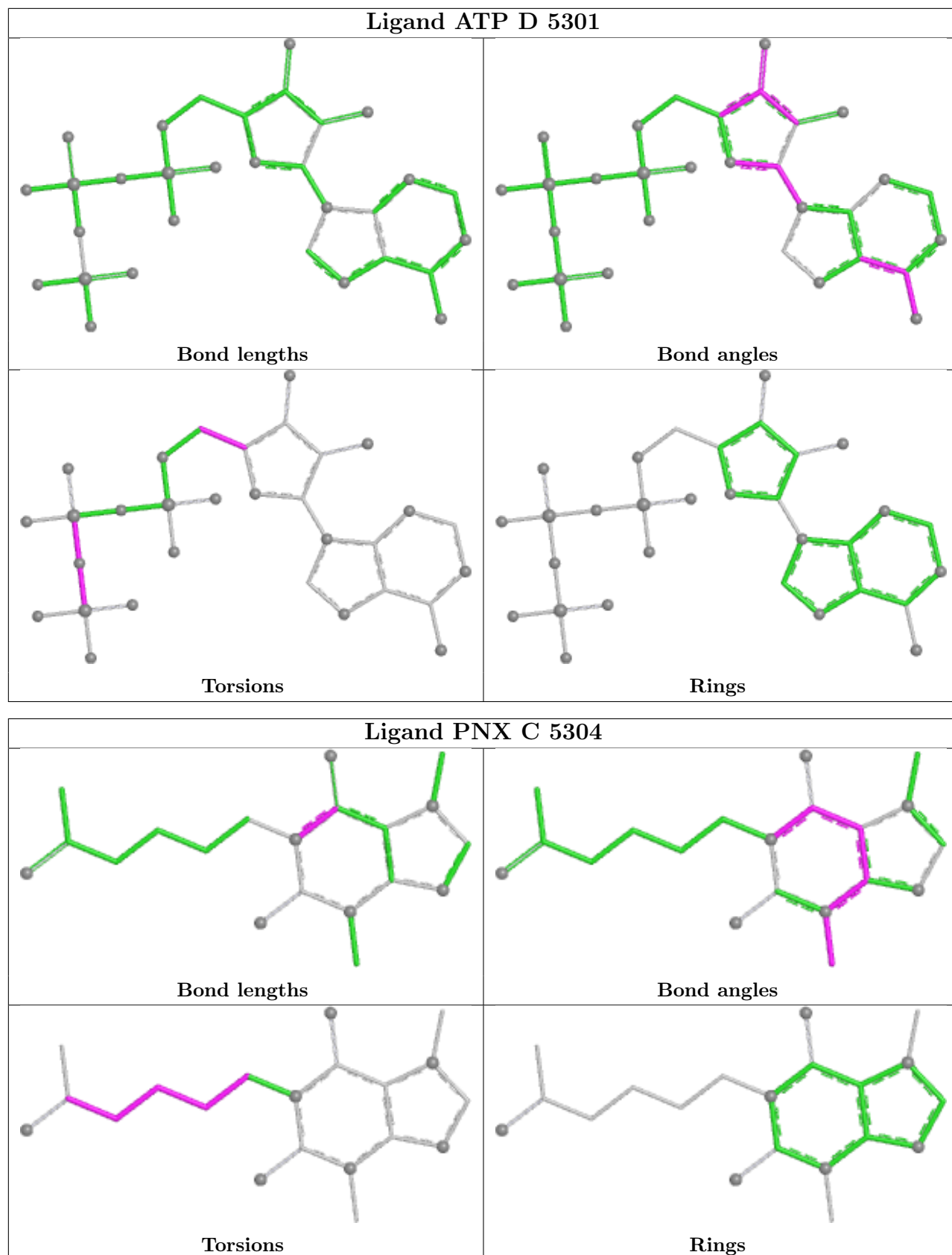
4 monomers are involved in 12 short contacts:

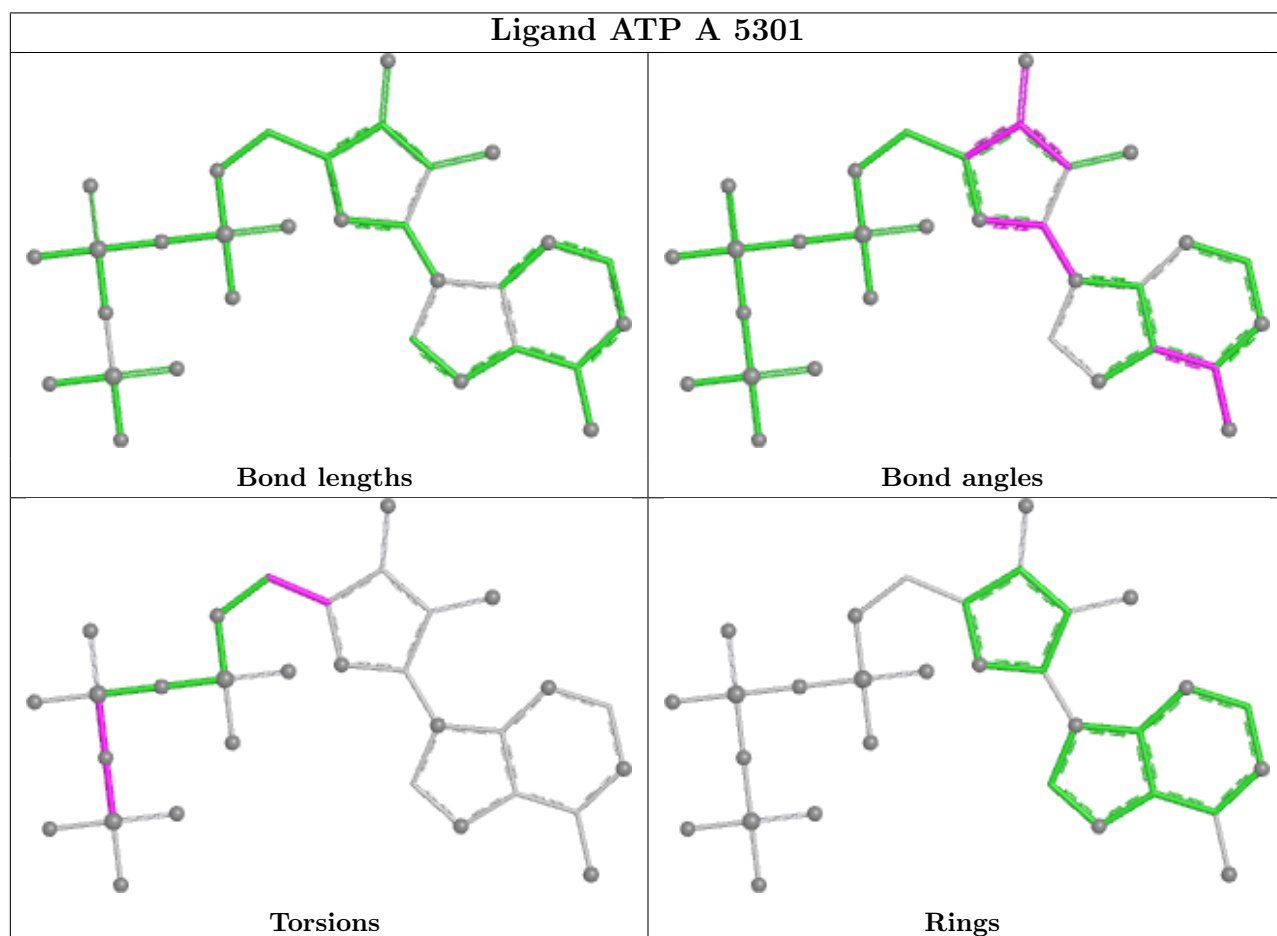
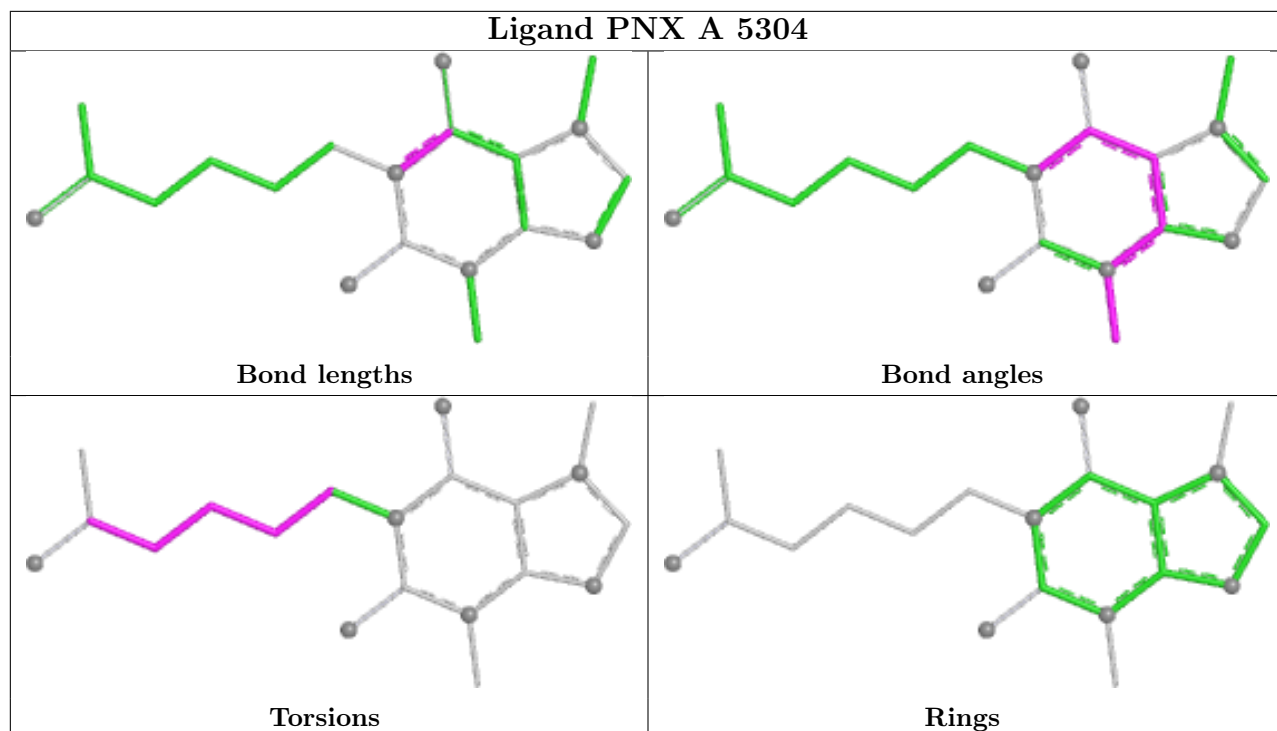
Mol	Chain	Res	Type	Clashes	Symm-Clashes
6	C	5304	PNX	3	0
6	A	5304	PNX	3	0
6	B	5304	PNX	3	0
6	D	5304	PNX	3	0

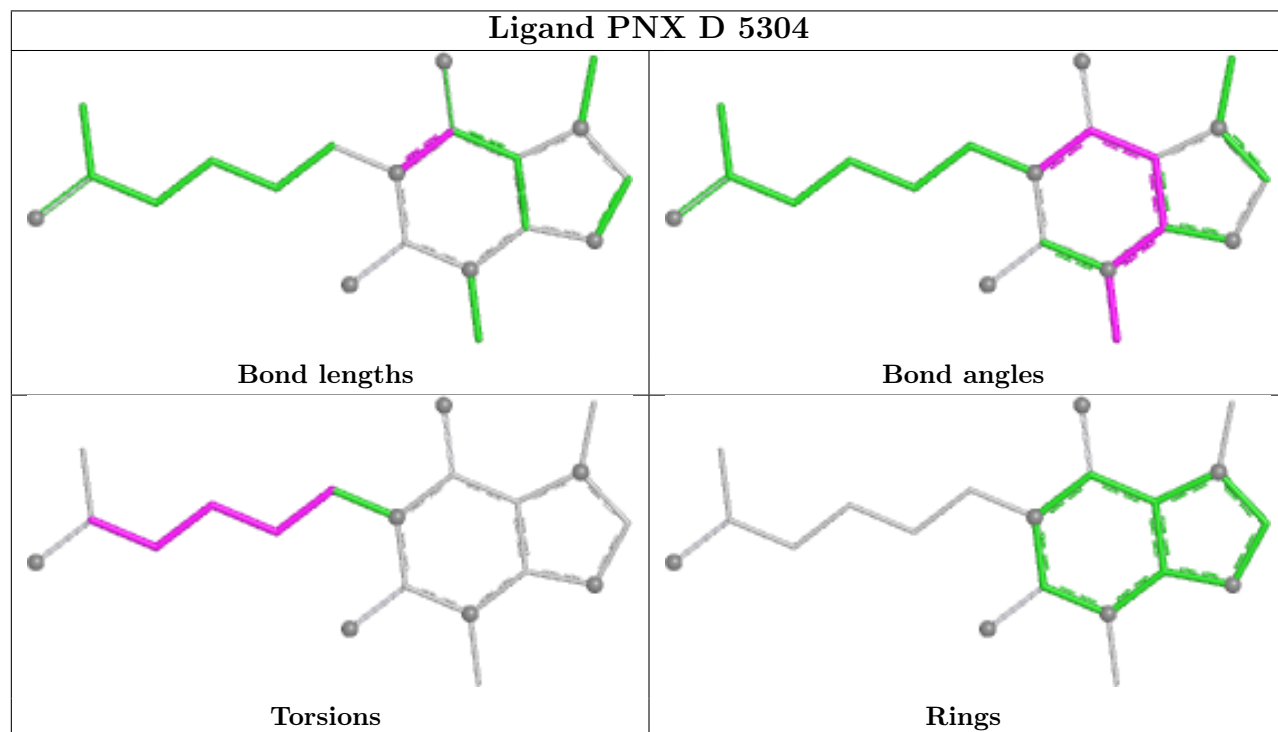
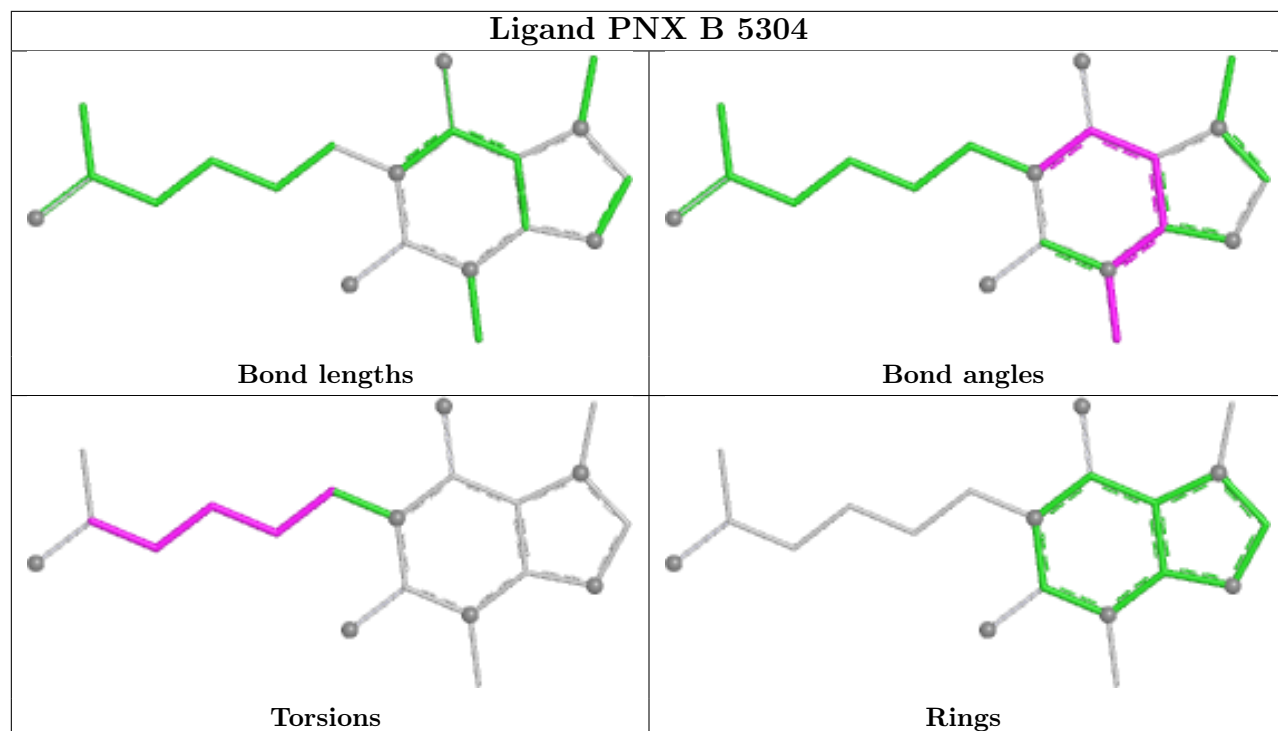
The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. 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

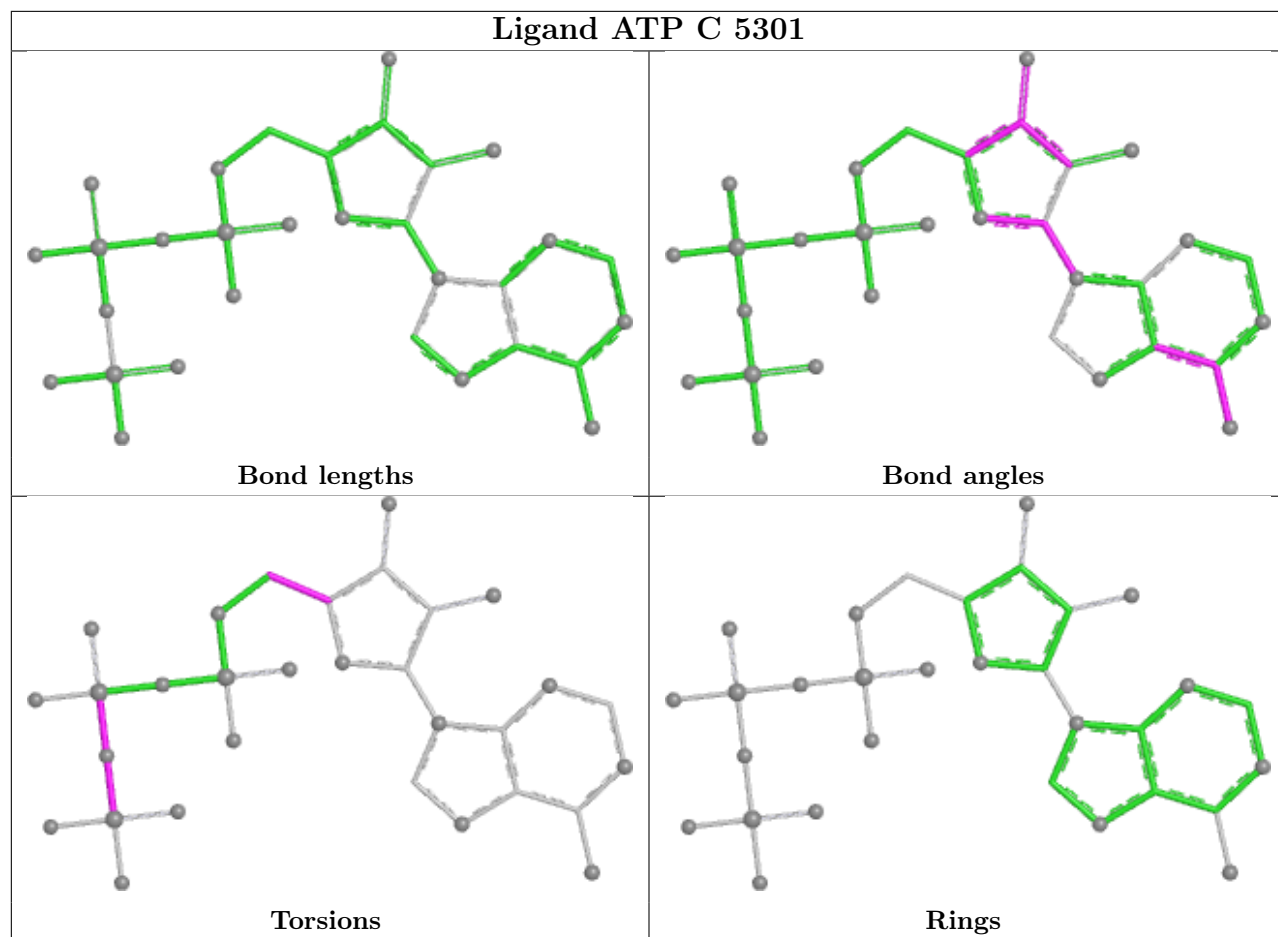
highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified 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.

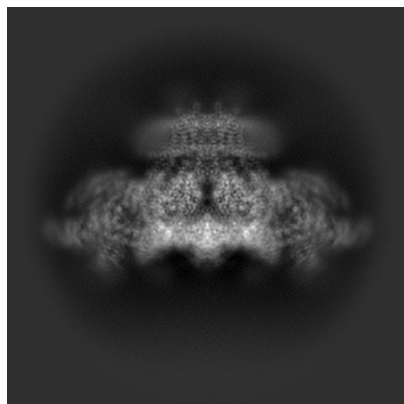
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-47385. These allow visual inspection of the internal detail of the map and identification of artifacts.

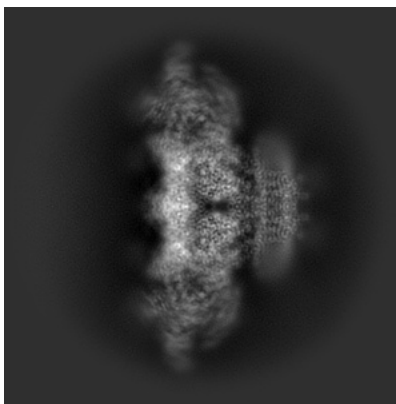
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

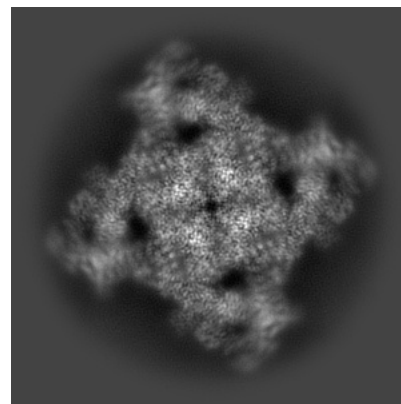
6.1.1 Primary map



X

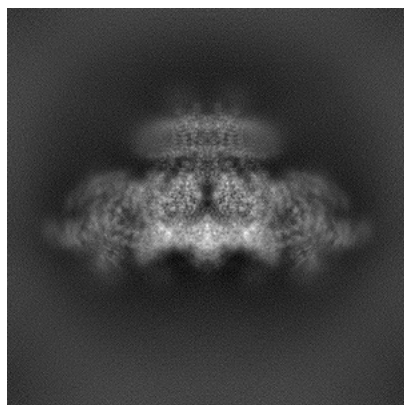


Y

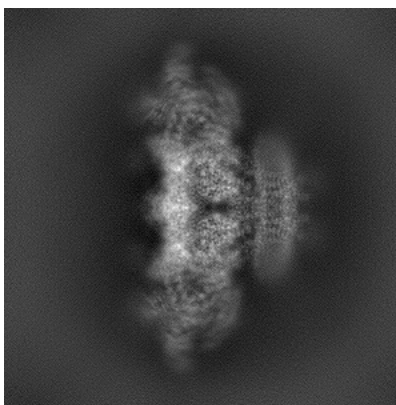


Z

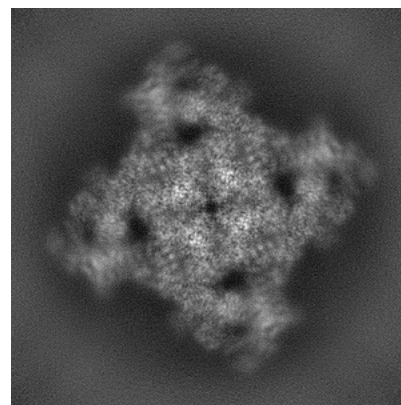
6.1.2 Raw map



X



Y

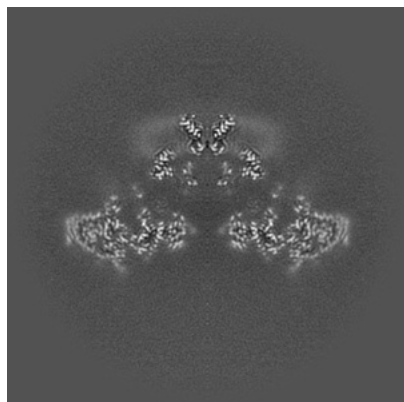


Z

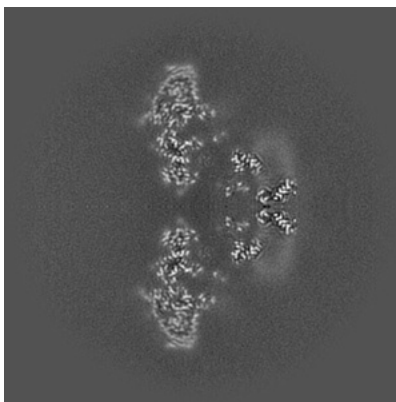
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

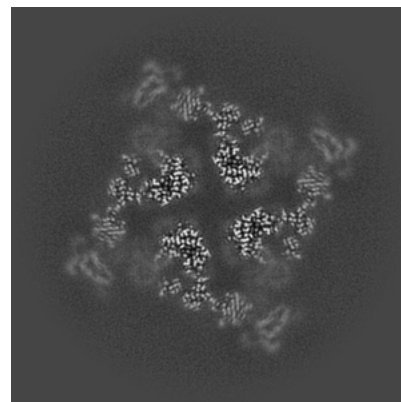
6.2.1 Primary map



X Index: 256

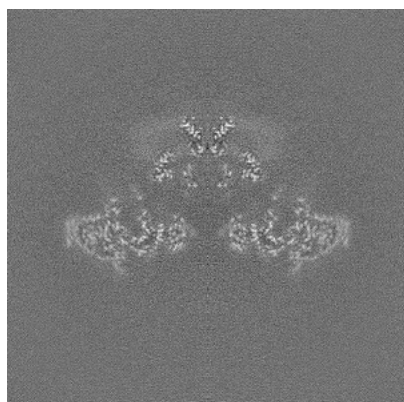


Y Index: 256

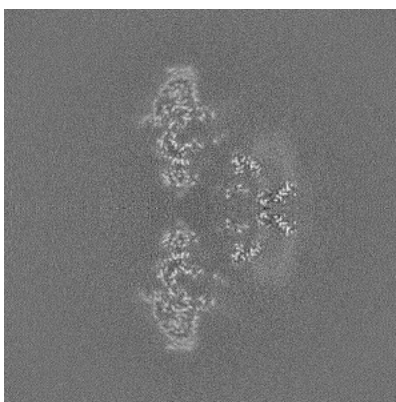


Z Index: 256

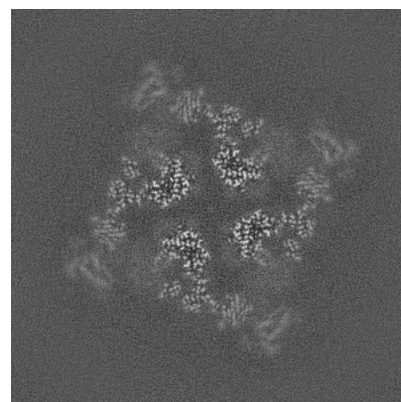
6.2.2 Raw map



X Index: 256



Y Index: 256

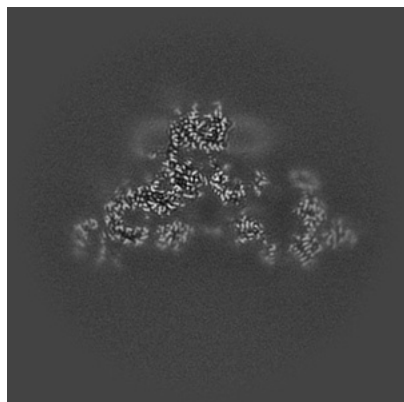


Z Index: 256

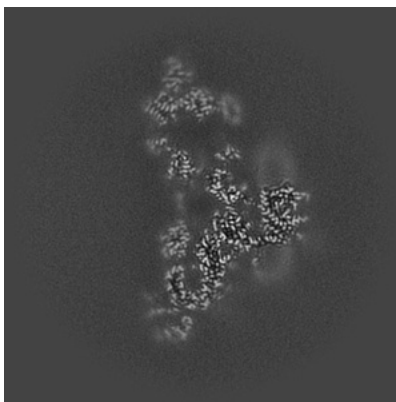
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

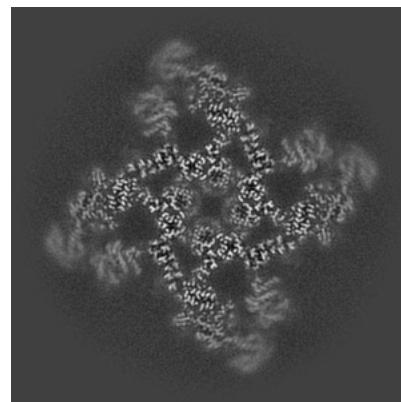
6.3.1 Primary map



X Index: 239

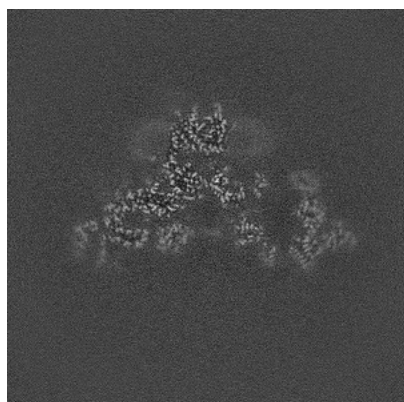


Y Index: 273



Z Index: 229

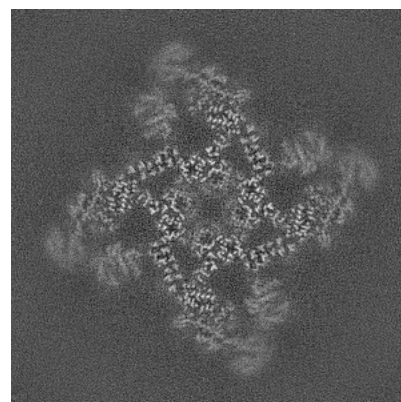
6.3.2 Raw map



X Index: 239



Y Index: 239

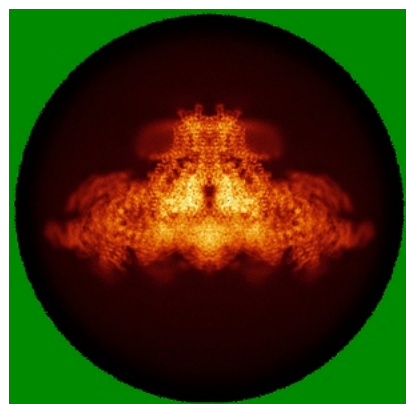


Z Index: 229

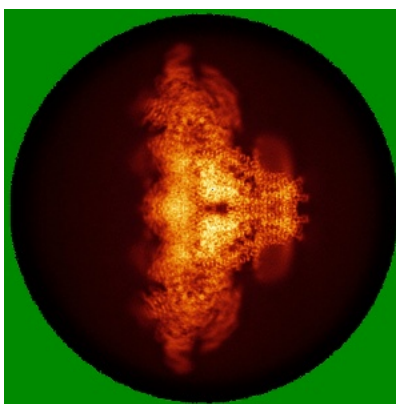
The images above show the largest variance slices of the map in three orthogonal directions.

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

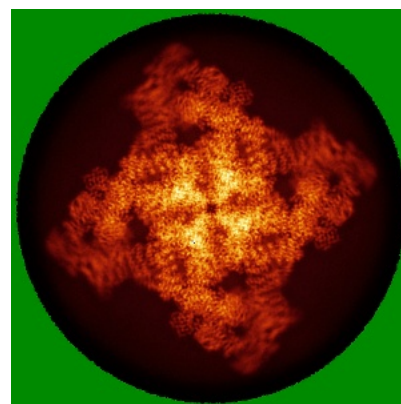
6.4.1 Primary map



X

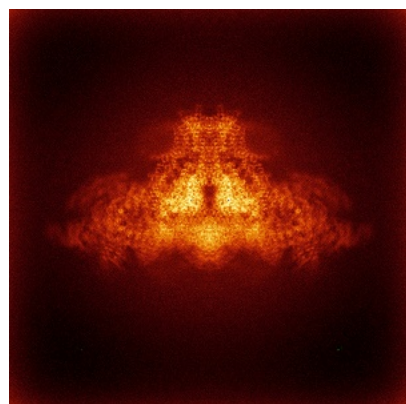


Y

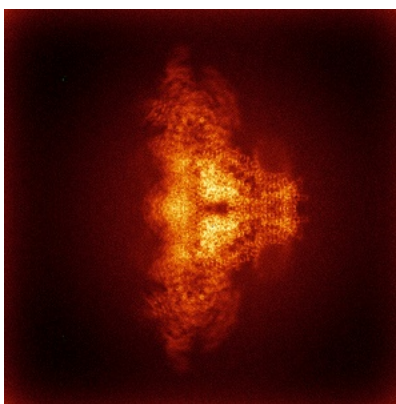


Z

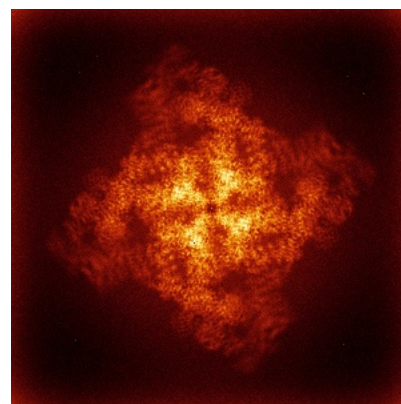
6.4.2 Raw map



X



Y

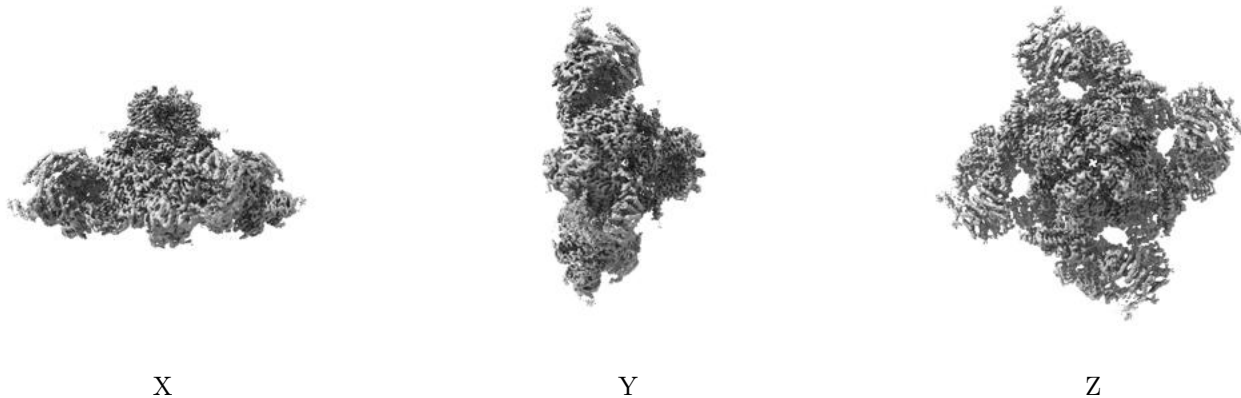


Z

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.

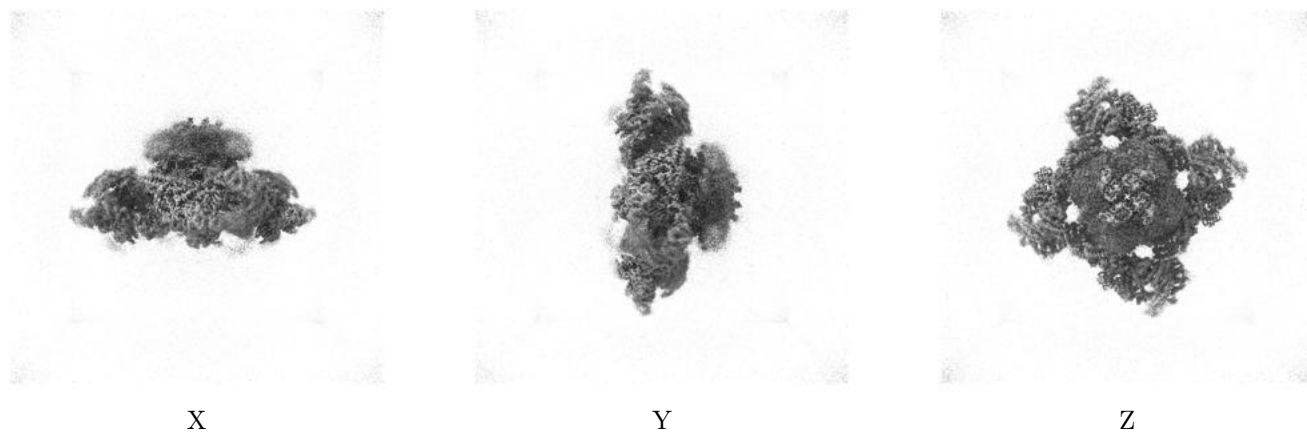
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.1. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

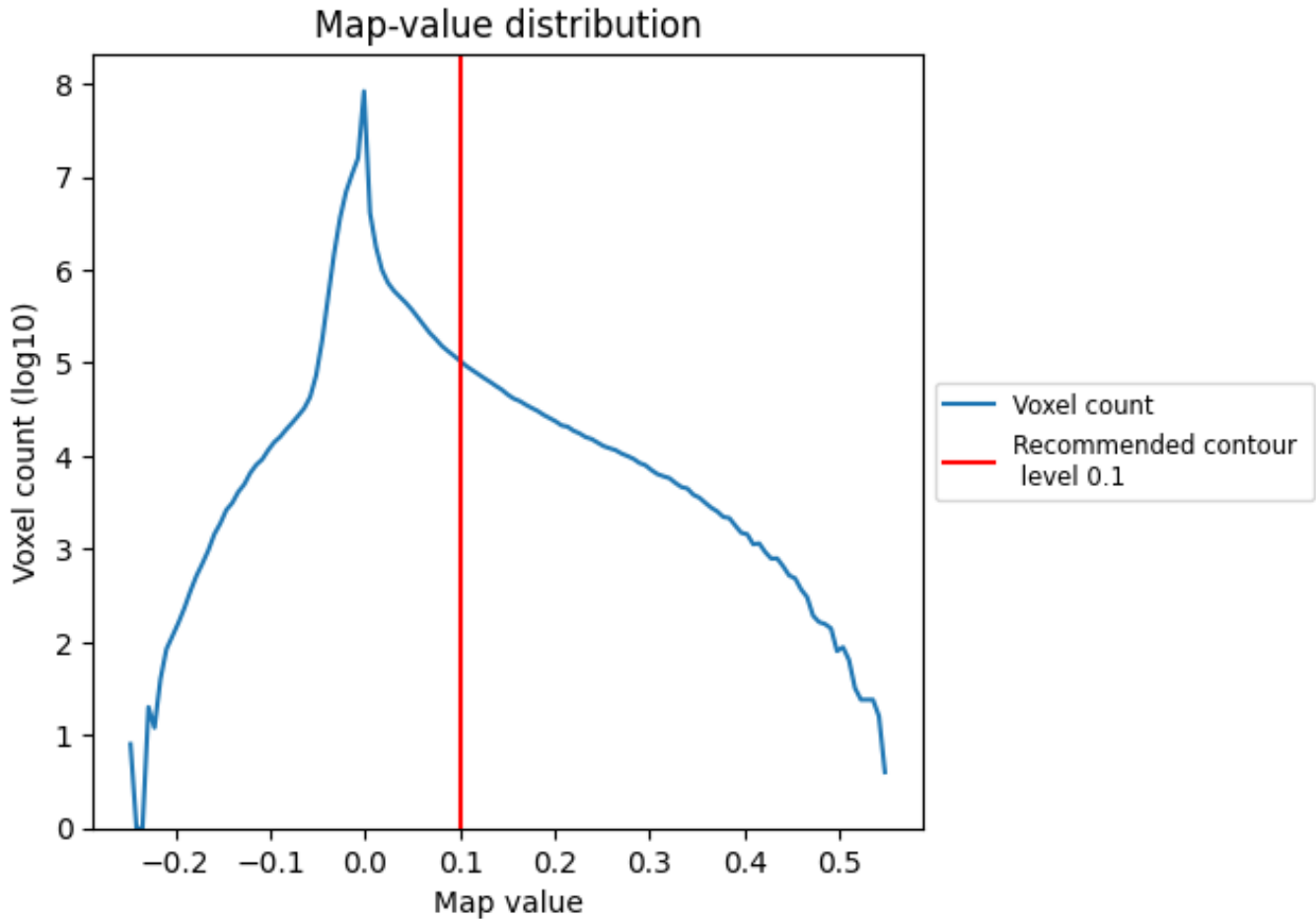
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

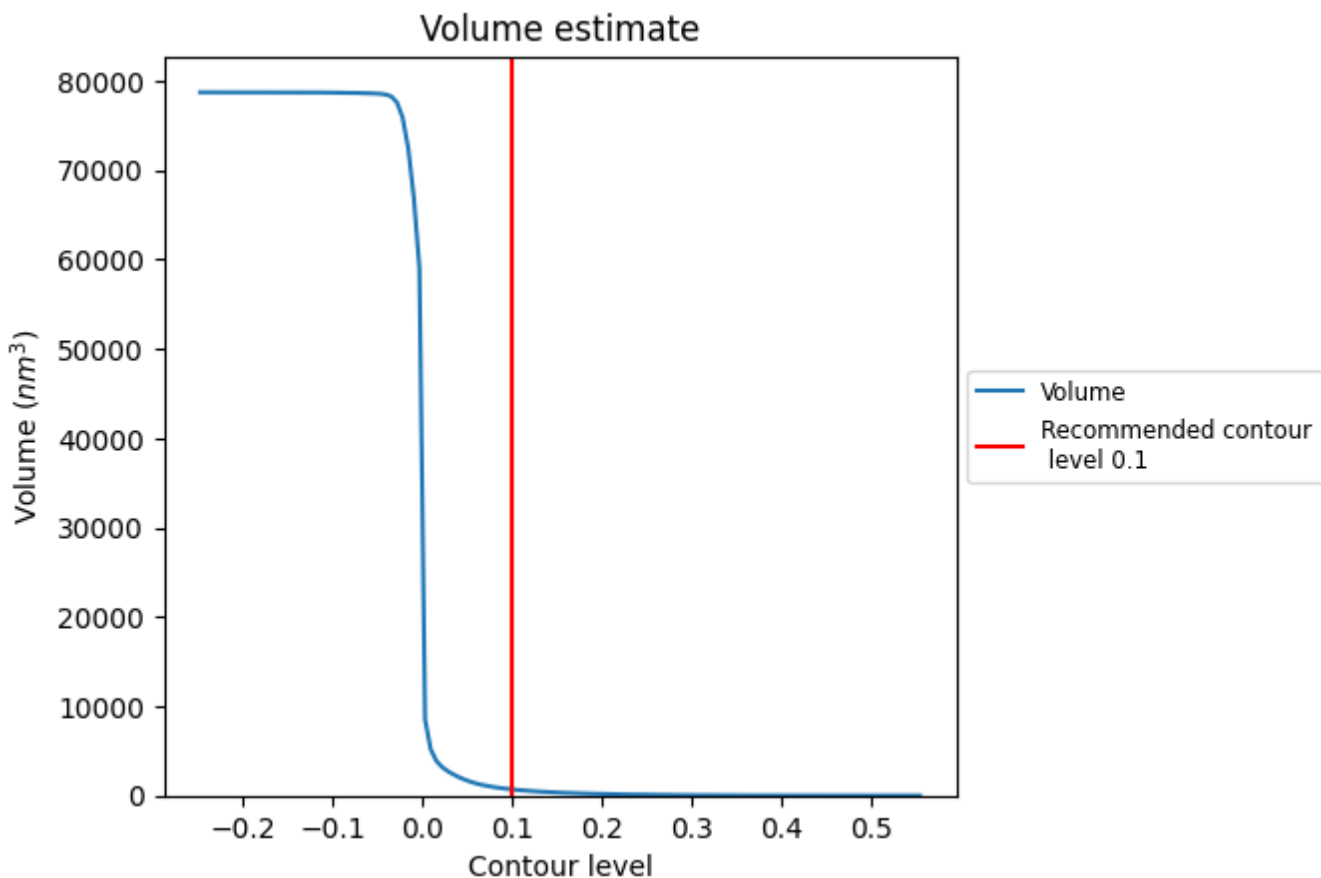
This section contains the results of statistical analysis of the map.

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.

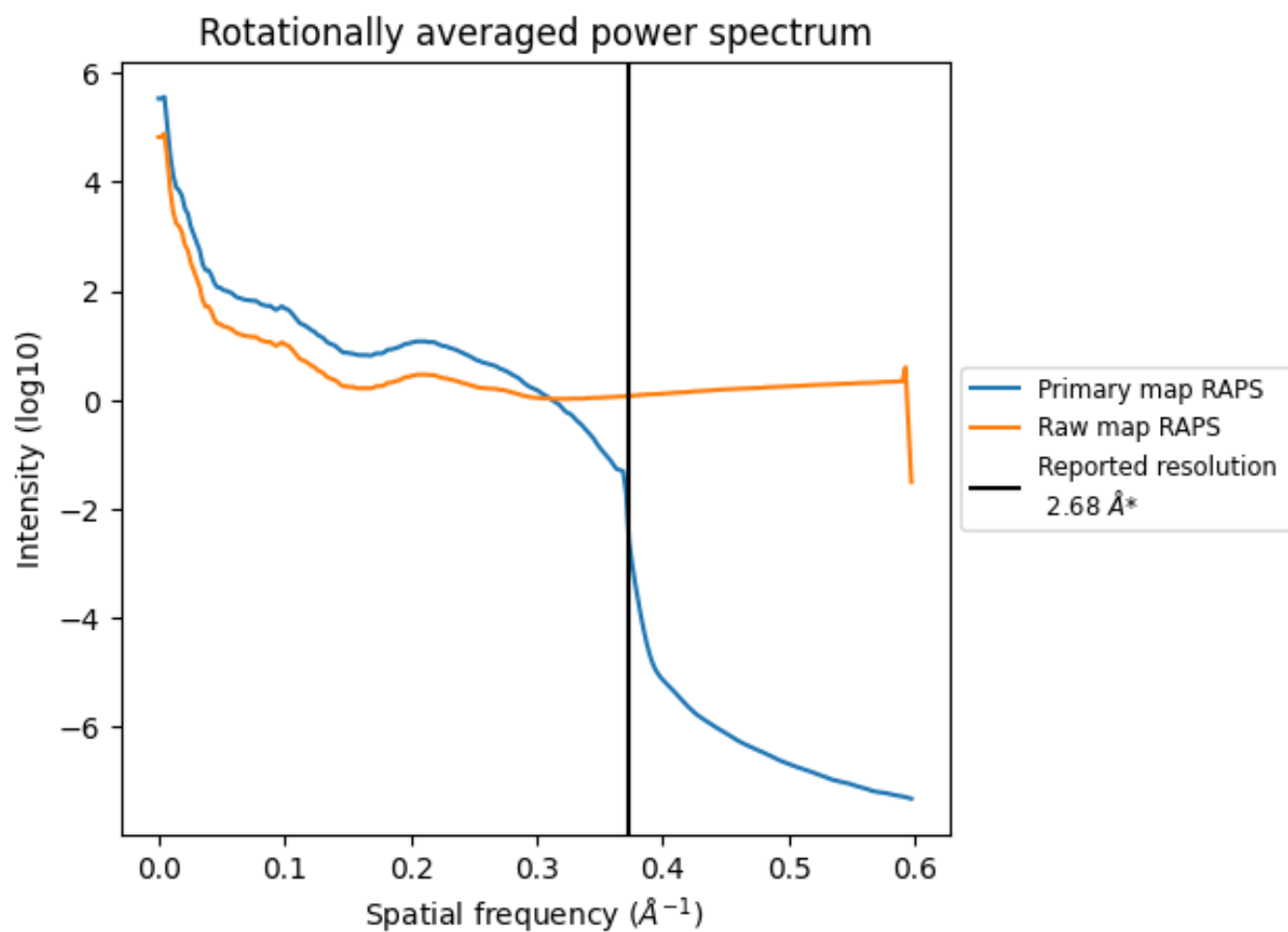
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 692 nm³; this corresponds to an approximate mass of 625 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 i

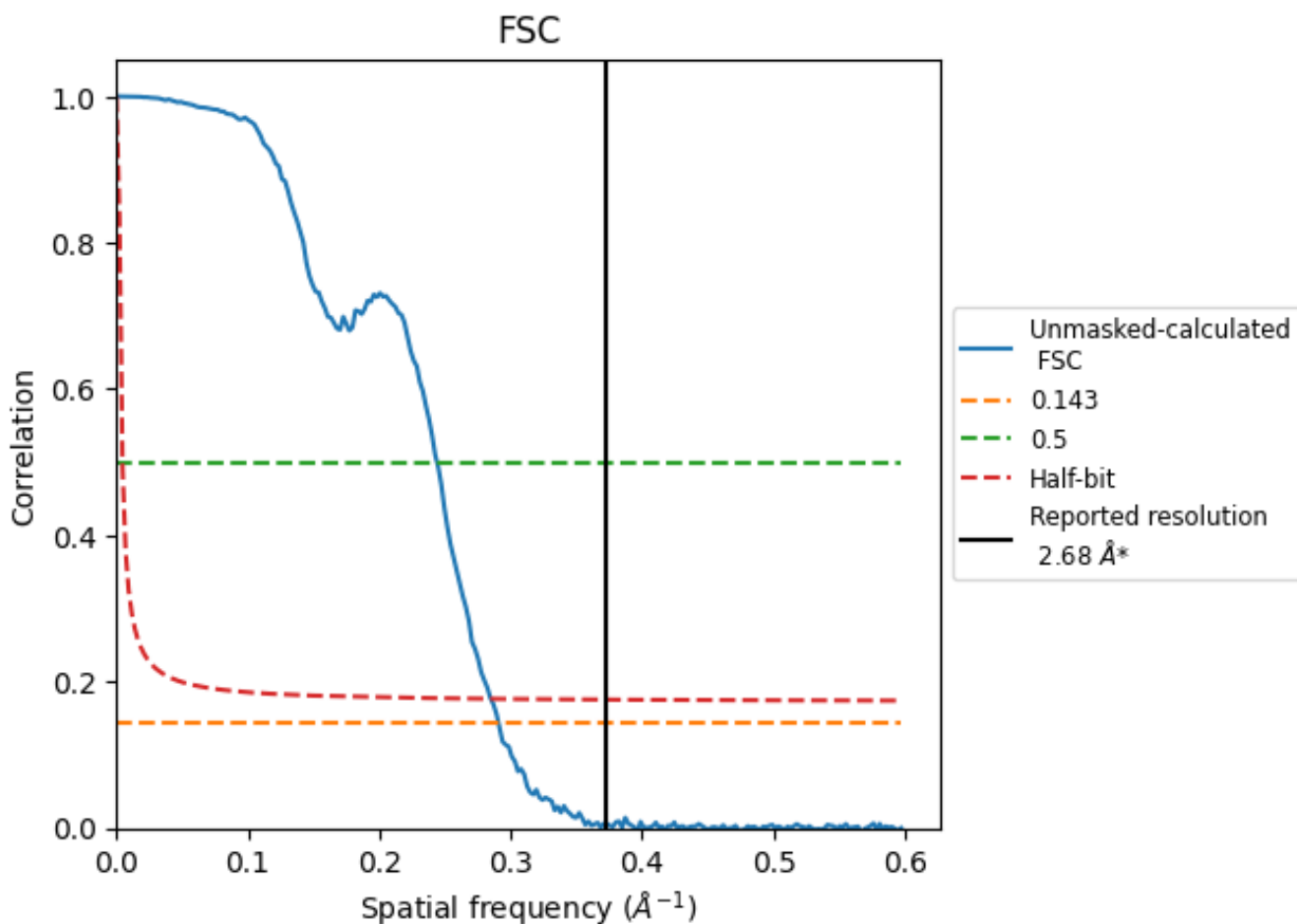


*Reported resolution corresponds to spatial frequency of 0.373 Å⁻¹

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 0.373 \AA^{-1}

8.2 Resolution estimates [i](#)

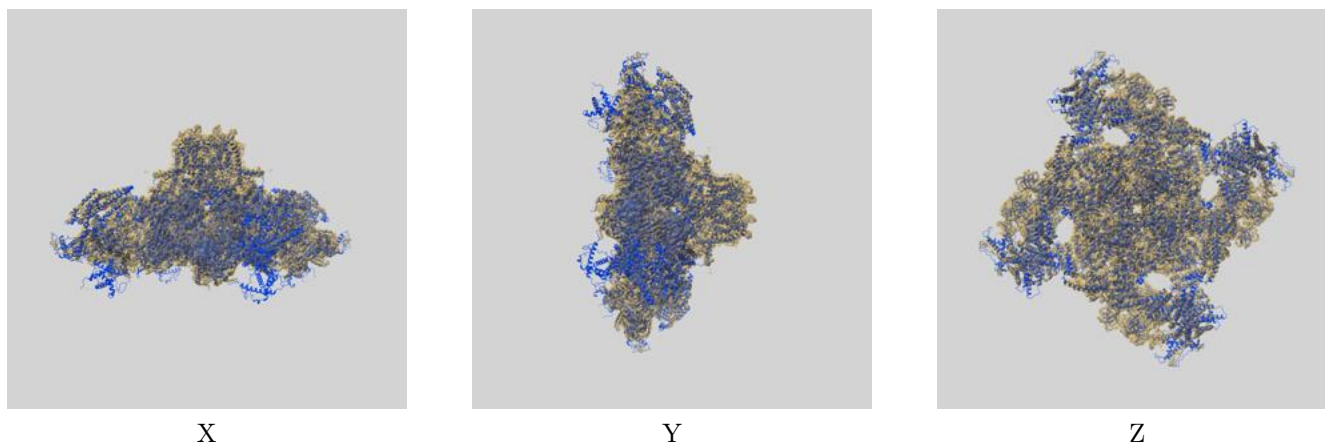
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.68	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	3.44	4.10	3.51

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.44 differs from the reported value 2.68 by more than 10 %

9 Map-model fit [i](#)

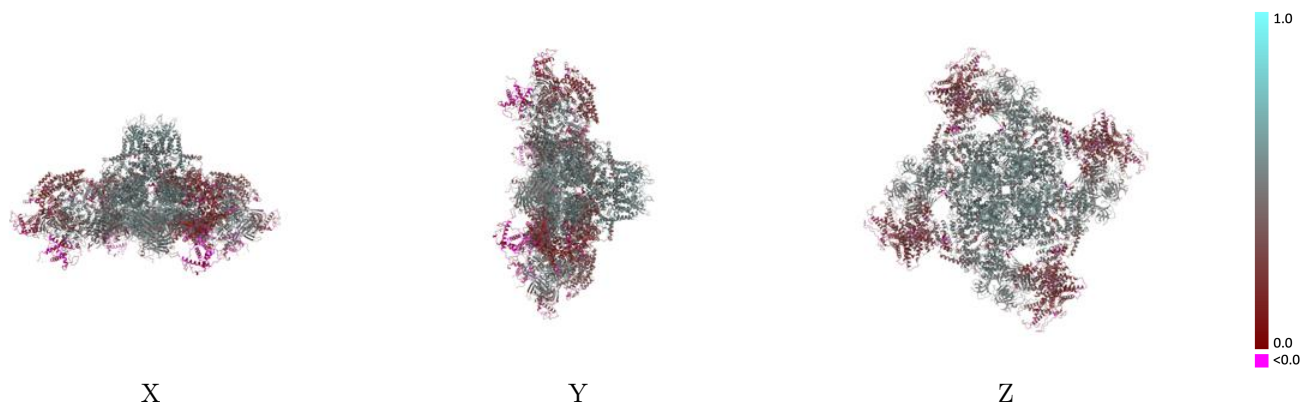
This section contains information regarding the fit between EMDB map EMD-47385 and PDB model 9E18. Per-residue inclusion information can be found in section 3 on page 7.

9.1 Map-model overlay [i](#)



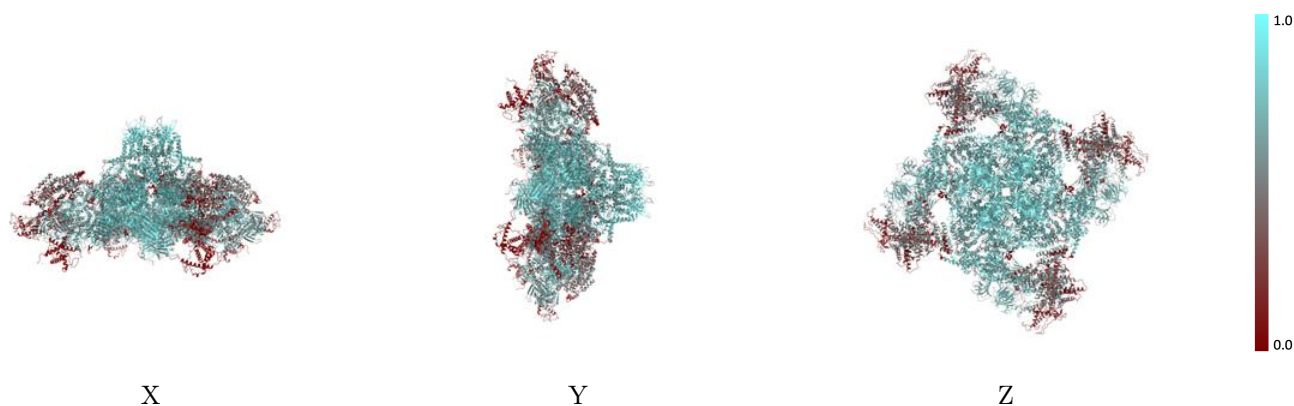
The images above show the 3D surface view of the map at the recommended contour level 0.1 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



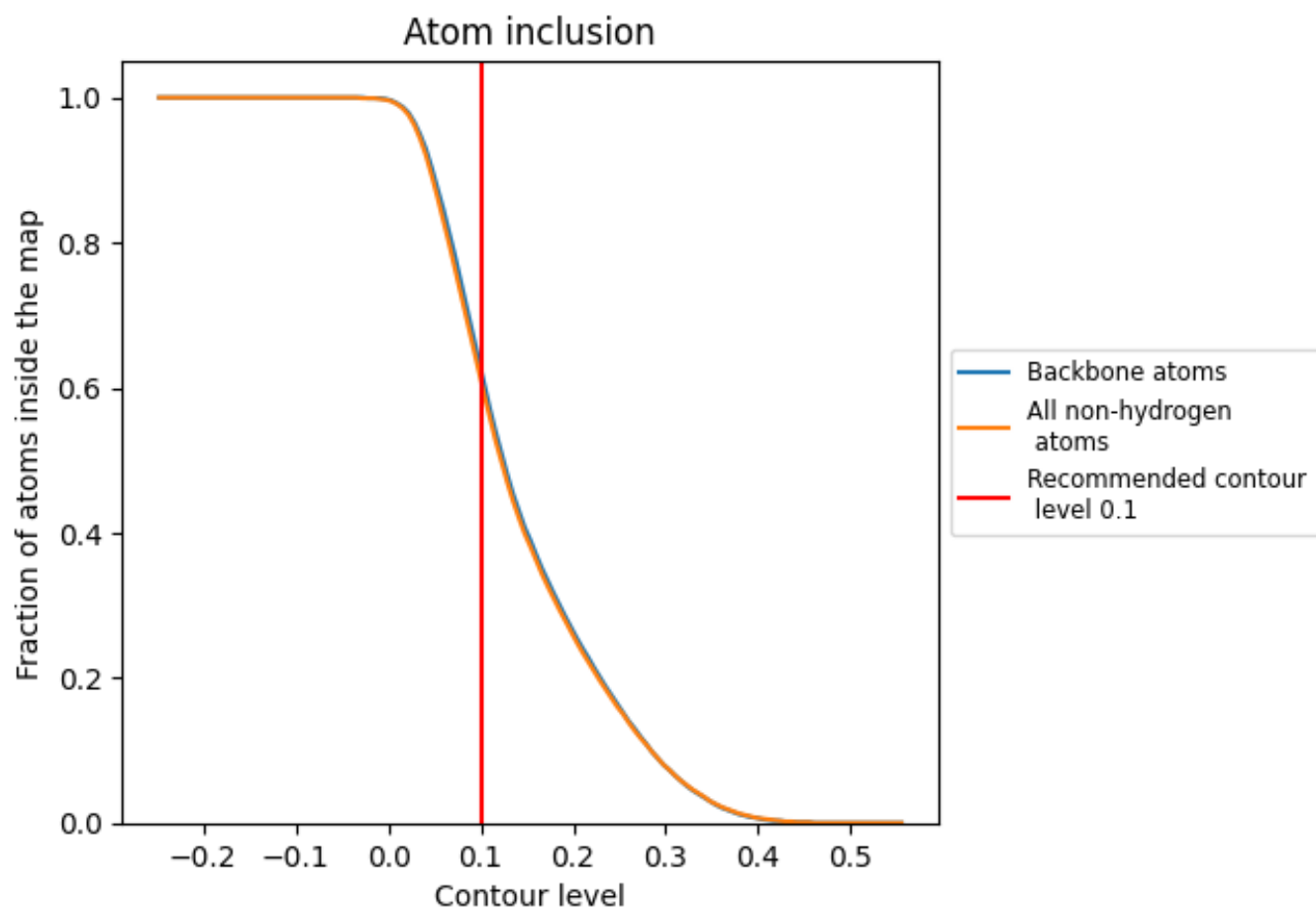
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.1).



















9.4 Atom inclusion [i](#)



At the recommended contour level, 63% of all backbone atoms, 61% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary [i](#)

The table lists the average atom inclusion at the recommended contour level (0.1) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.6110	 0.4160
A	 0.6180	 0.4140
B	 0.6180	 0.4150
C	 0.6180	 0.4140
D	 0.6180	 0.4140
E	 0.6470	 0.5000
F	 0.6500	 0.4970
G	 0.6430	 0.4940
H	 0.6470	 0.4960

