



Full wwPDB EM Validation Report ⓘ

Nov 11, 2024 – 06:45 AM EST

PDB ID : 9E1C
EMDB ID : EMD-47389
Title : Structure of RyR1 in the primed state in the presence of IBMX
Authors : Miotto, M.C.; Marks, A.R.
Deposited on : 2024-10-21
Resolution : 2.63 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

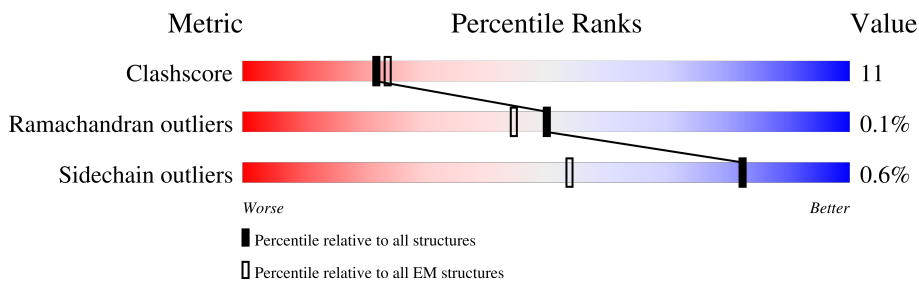
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 2.63 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 6 unique types of molecules in this entry. The entry contains 144120 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	C	N	O	S	9	0
			35150	22365	6063	6485	237		
1	B	4404	Total	C	N	O	S	9	0
			35150	22365	6063	6485	237		
1	D	4404	Total	C	N	O	S	9	0
			35150	22365	6063	6485	237		
1	C	4404	Total	C	N	O	S	9	0
			35150	22365	6063	6485	237		

- 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	C	N	O	S	0	0
			831	527	146	154	4		
2	H	107	Total	C	N	O	S	0	0
			831	527	146	154	4		
2	G	107	Total	C	N	O	S	0	0
			831	527	146	154	4		
2	F	107	Total	C	N	O	S	0	0
			831	527	146	154	4		

- 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
			Total	C	N	O	P	
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
			Total	Ca	
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).

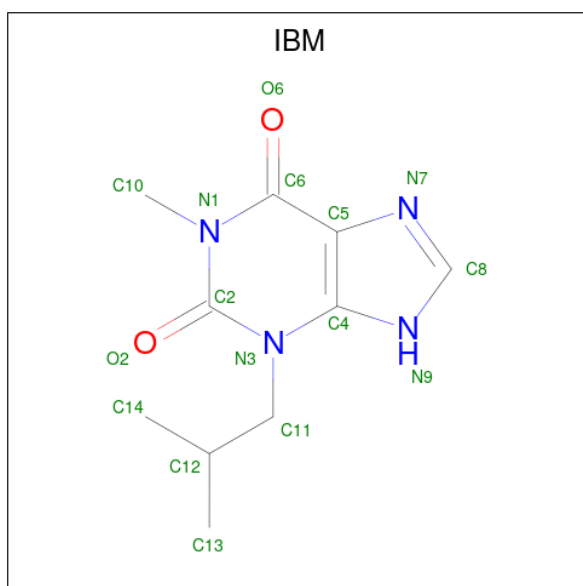
Mol	Chain	Residues	Atoms		AltConf
			Total	Zn	
5	A	1	Total	Zn	0
			1	1	

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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-ISOBUTYL-1-METHYLNANTHINE (three-letter code: IBM) (formula: $C_{10}H_{14}N_4O_2$) (labeled as "Ligand of Interest" by depositor).

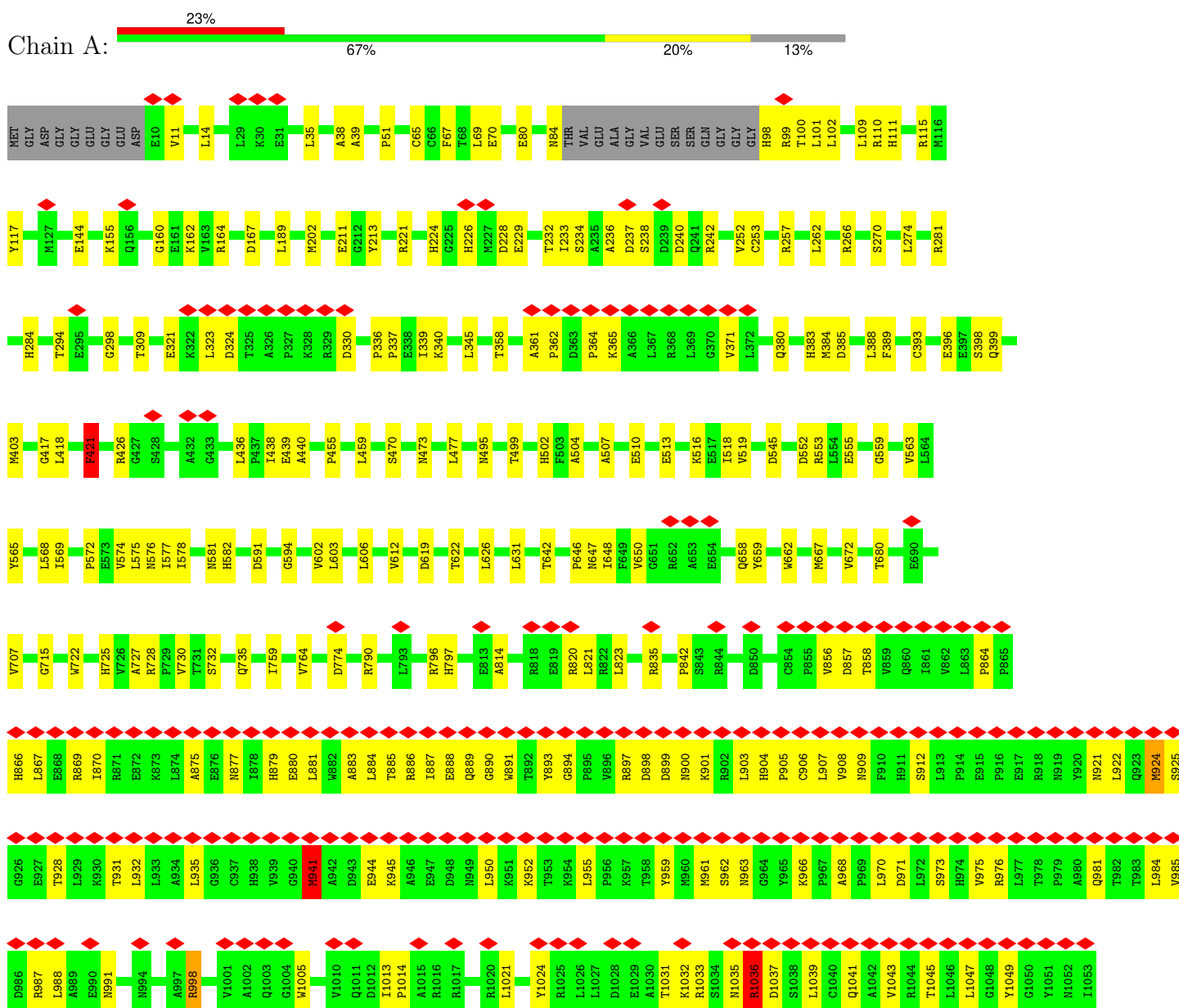


Mol	Chain	Residues	Atoms				AltConf
6	A	1	Total	C	N	O	0
			16	10	4	2	
6	B	1	Total	C	N	O	0
			16	10	4	2	
6	D	1	Total	C	N	O	0
			16	10	4	2	
6	C	1	Total	C	N	O	0
			16	10	4	2	

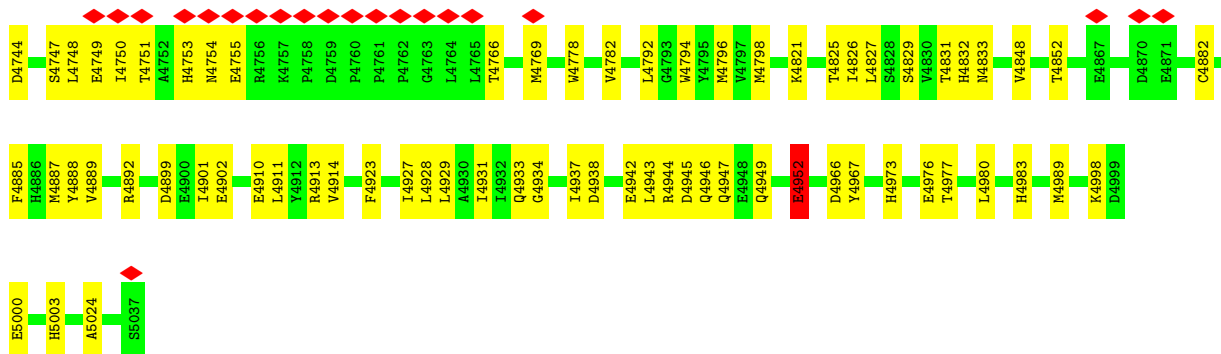
3 Residue-property plots

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.

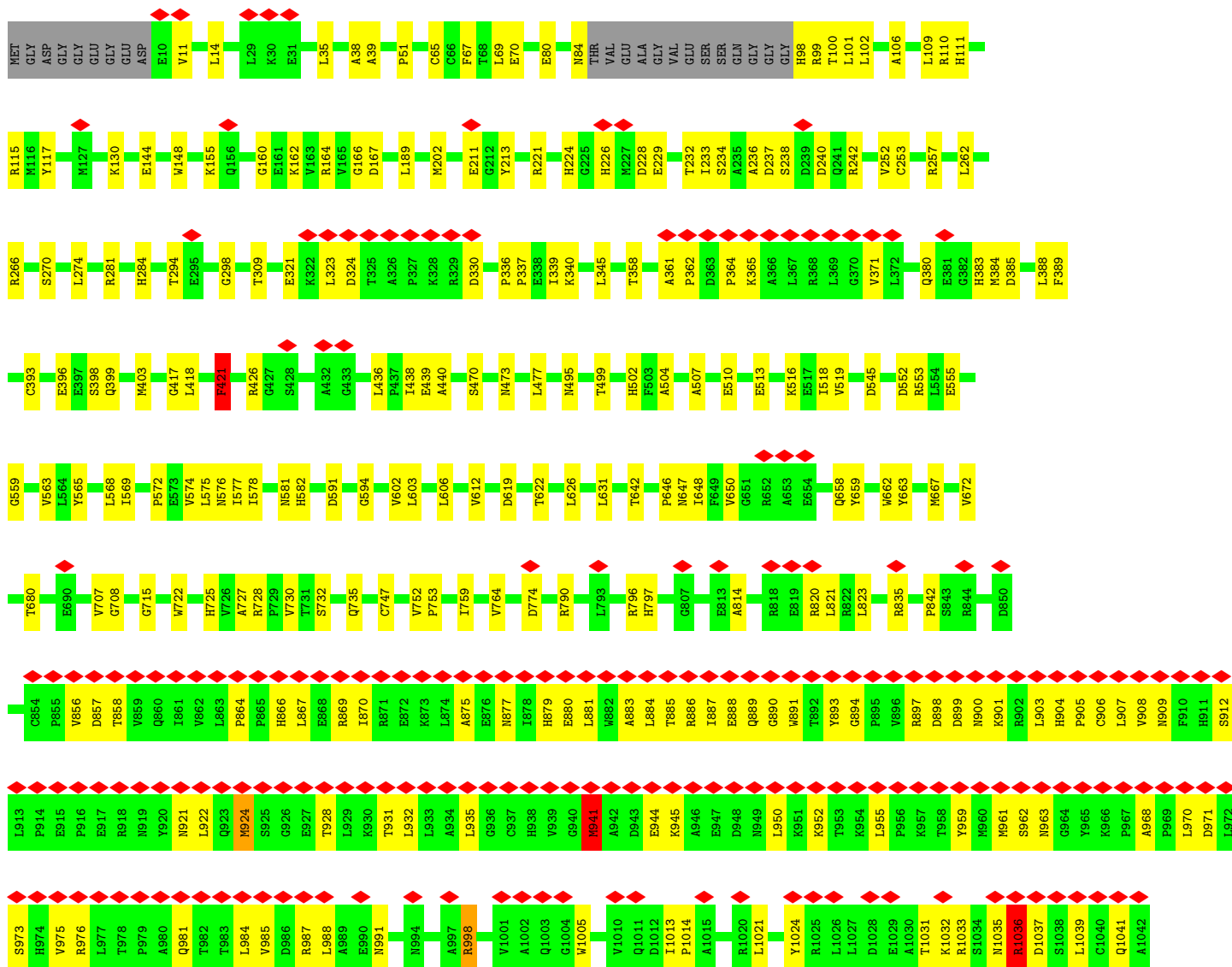
• Molecule 1: Ryanodine receptor 1

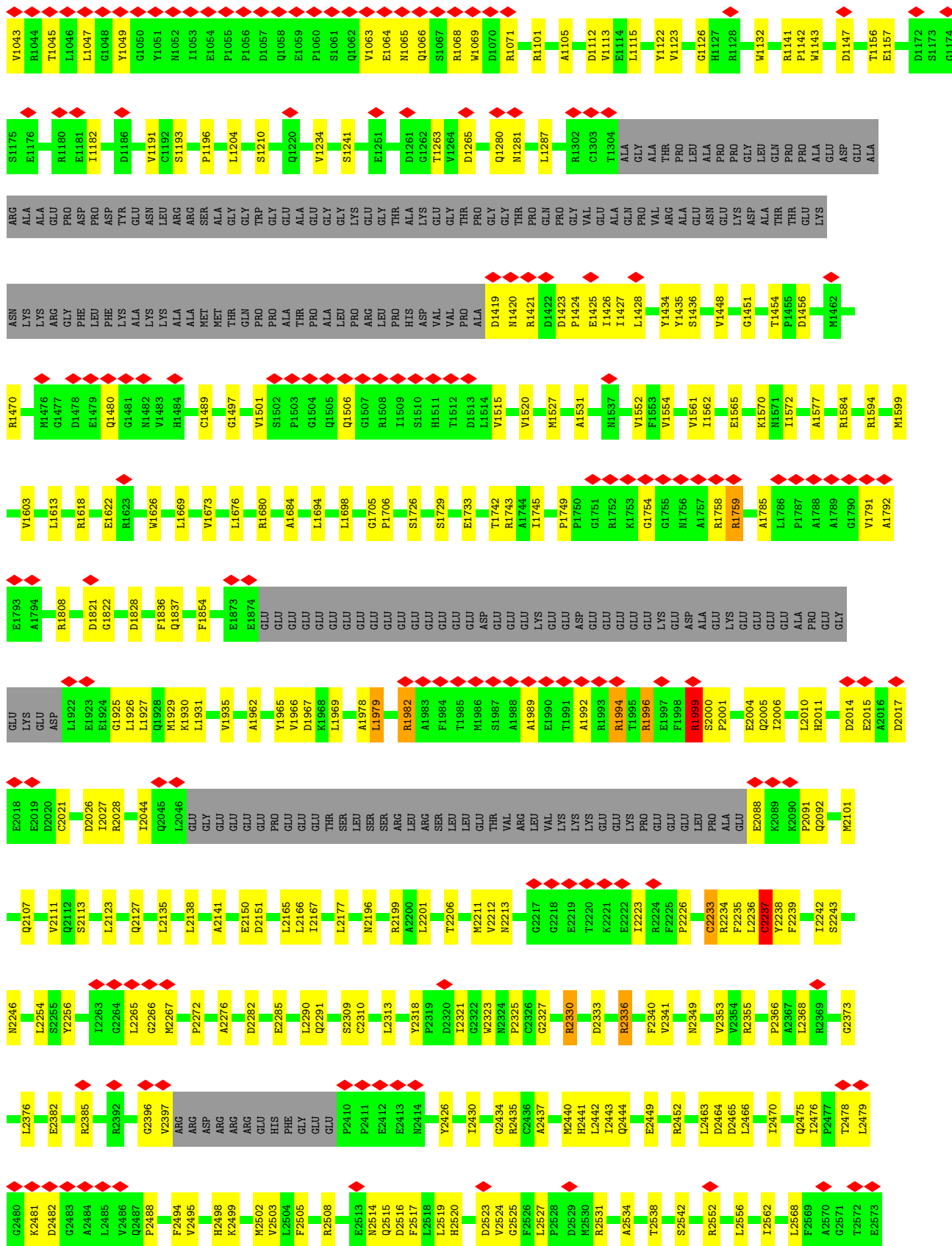


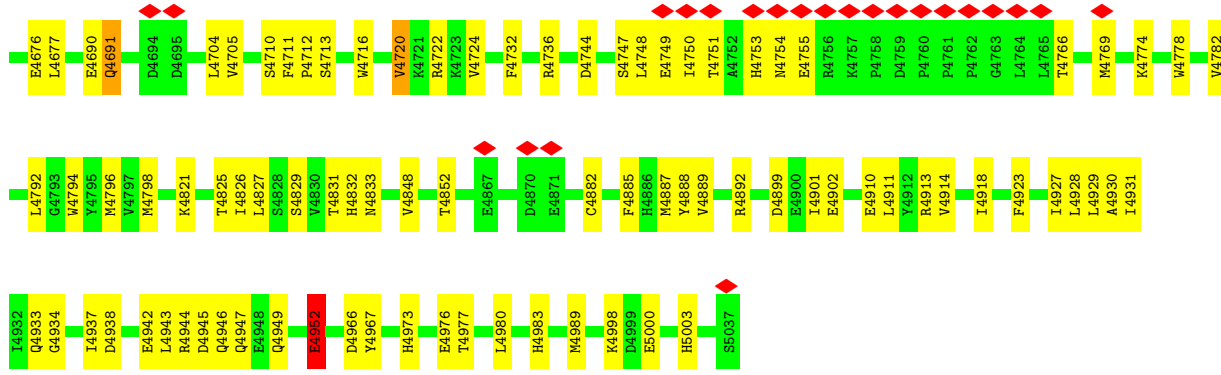
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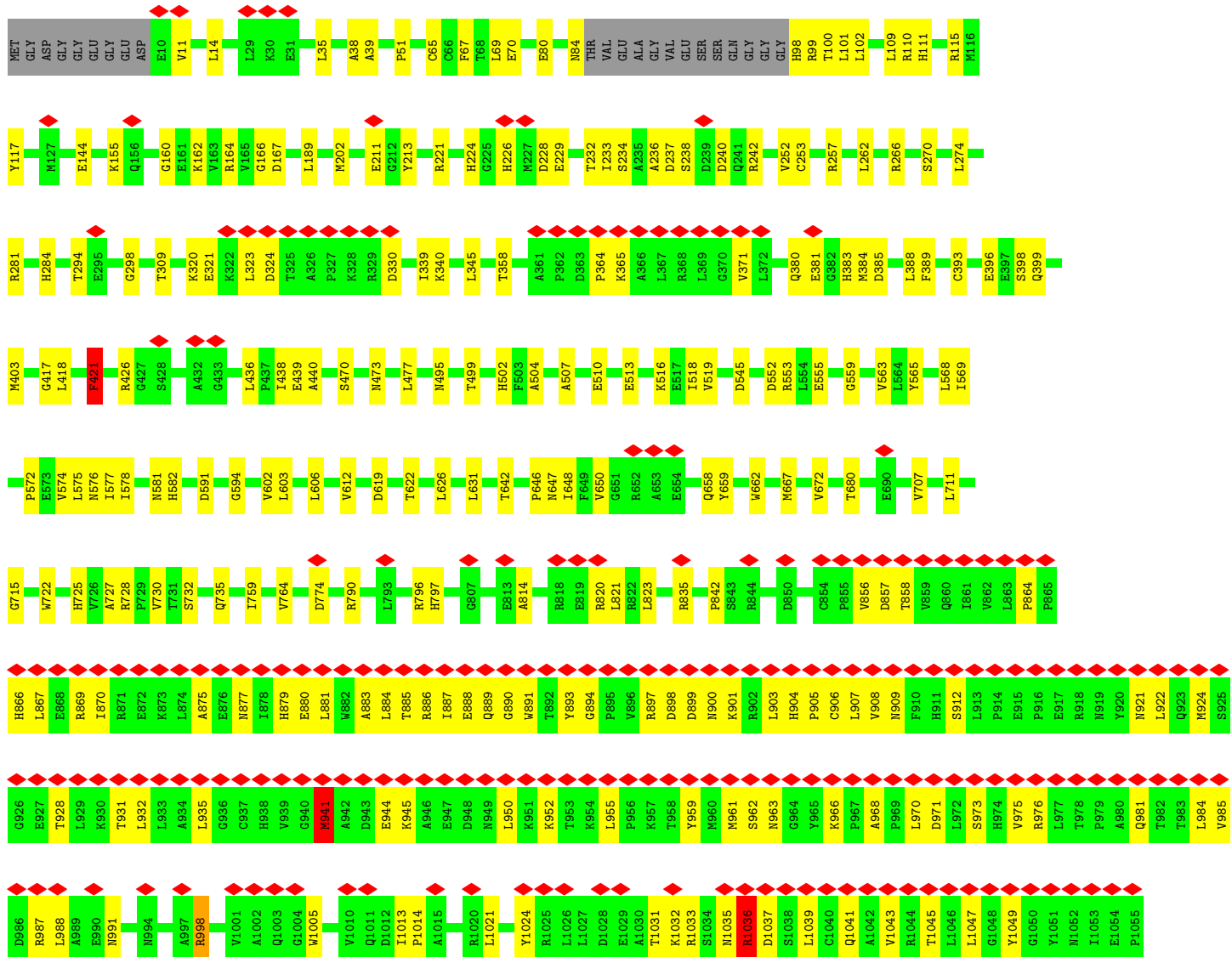
● Molecule 1: Ryanodine receptor 1

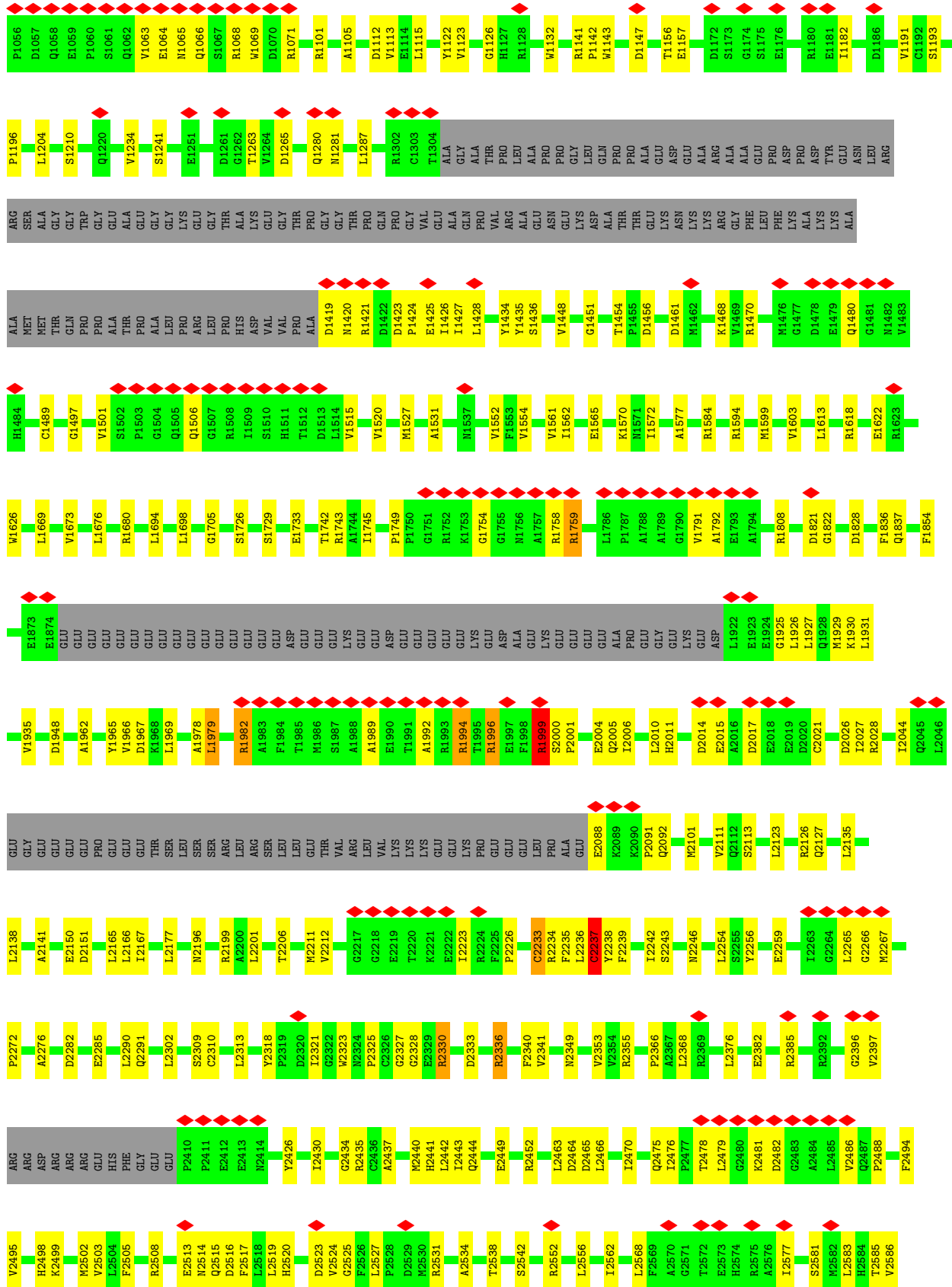




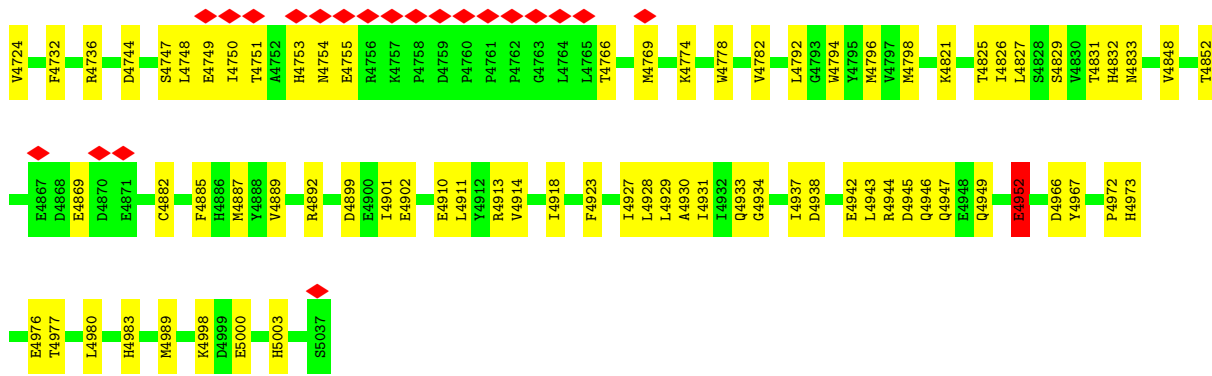


• Molecule 1: Ryanodine receptor 1

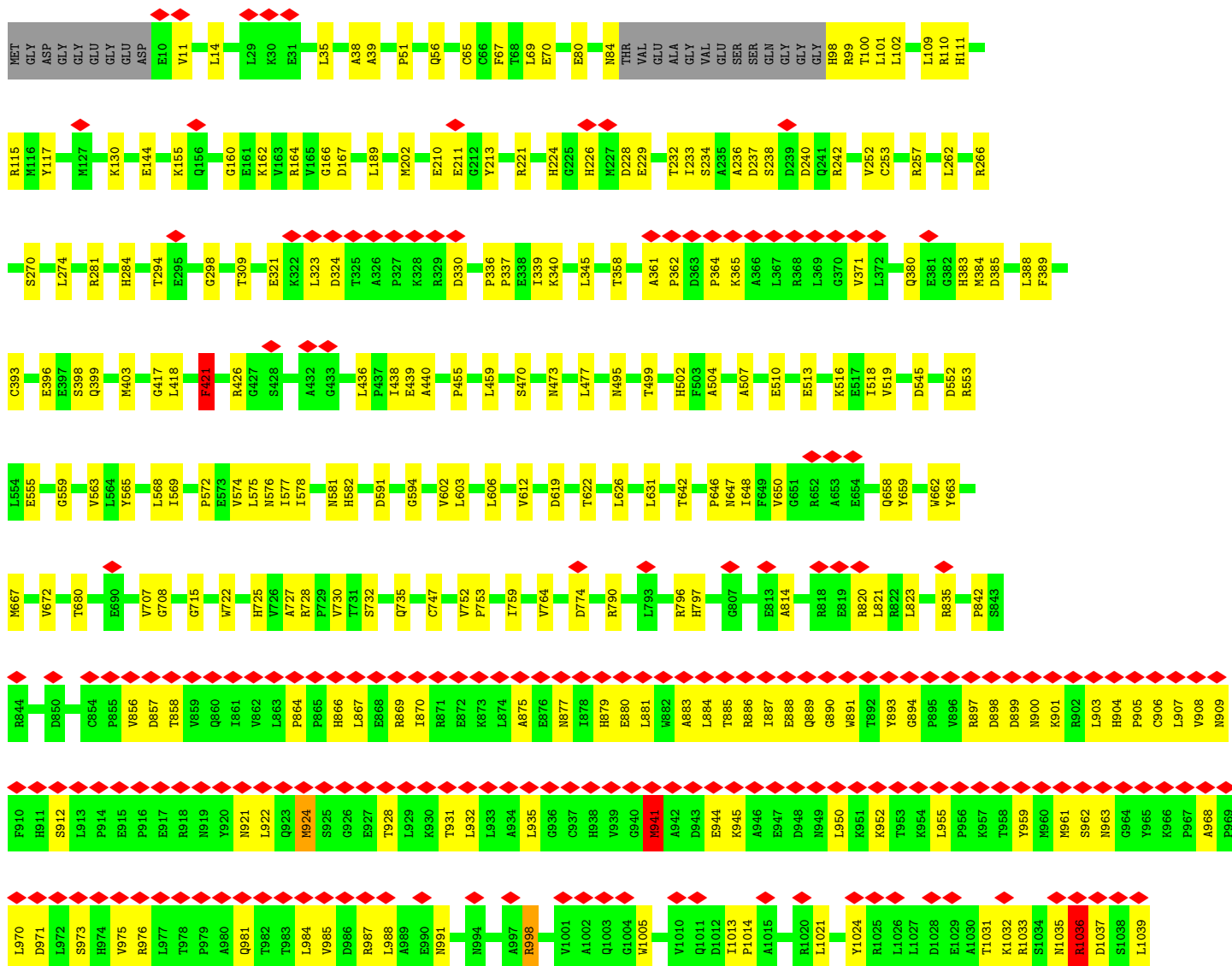


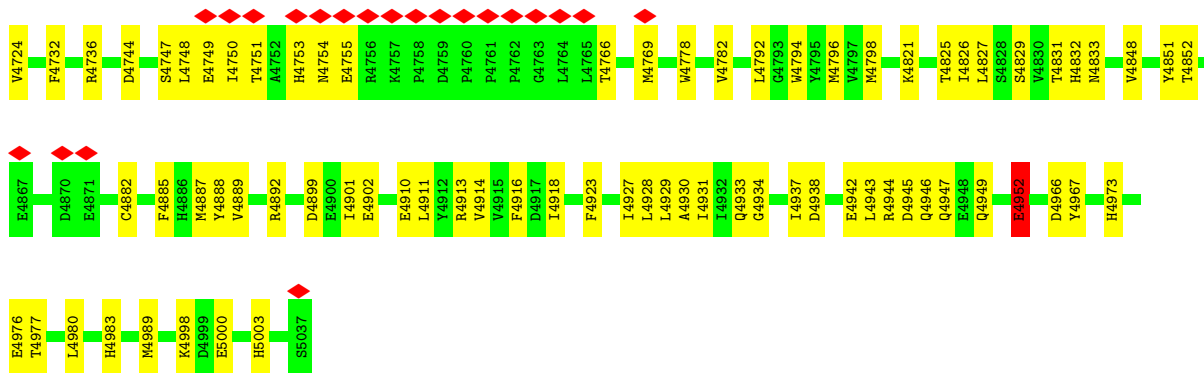


S3347	R3348	A3349	R3350	L3353	L3354	F3357	F3358	I3359	P3360	T3361	L3362	G3363	R3364	L3365	R3366	R3367	R3368	A3369	G3370	K3371	V3372	V3373	A3374	E3375	E3376	E3377	Q3378	L3379	R3380	L3381	E3382	A3383	K3384	A3385	E3386	A3387	E3388	E3389	G3390	E3391	L3392	L3393	V3394	R3395	D3396	L3401	R3402	R3403	D3404	L3405	V3406	A3407	V3409	F3410	L3411																																																														
E3286	R3287	G3288	P3289	E3290	A3291	P3292	P3293	P3294	A3295	L3296	P3297	A3298	G3299	A3300	P3301	P3302	P3303	C3304	T3305	A3306	V3307	T3308	S3309	D3310	H3311	L3312	M3313	L3316	G3317	A3318	L3319	G3320	E3321	D3322	L3323	A3324	E3325	S3326	L3327	A3328	L3329	A3330	P3331	L3332	V3333	M3334	N3335	N3336	L3337	G3338	L3339	E3340	F3341	A3342	Q3343	P3344	V3345																																																												
P3224	R3225	E3226	R3227	A3228	L3229	G3231	L3232	L3233	P3233	N3234	S3235	V3236	E3237	E3238	M3239	C3240	P3241	D3242	I3243	P3244	V3245	L3246	D3247	R3248	L3249	M3250	G3251	D3252	I3253	A3257	E3258	S3259	G3260	A3261	R3262	Y3263	T3264	E3265	M3266	F3267	H3268	V3269	E3270	I3271	V3272	T3273	L3274	P3275	M3276	L3277	C3278	S3279	L3281	P3282	R3283	P3284	W3285																																																												
E3286	R3287	G3288	P3289	E3290	A3291	P3292	P3293	P3294	A3295	L3296	P3297	A3298	G3299	A3300	P3301	P3302	P3303	C3304	T3305	A3306	V3307	T3308	S3309	D3310	H3311	L3312	M3313	L3316	G3317	A3318	L3319	G3320	E3321	D3322	L3323	A3324	E3325	S3326	L3327	A3328	L3329	A3330	P3331	L3332	V3333	M3334	N3335	N3336	L3337	G3338	L3339	E3340	F3341	A3342	Q3343	P3344	V3345																																																												
H2673	L2674	T2675	R2676	K2677	L2678	I2682	F2683	D2684	S2685	L2686	A2687	H2688	K2689	K2690	Y2691	D2692	Q2693	E2694	L2695	R2696	R2697	M2698	M2700	P2701	C2702	L2703	L2703	C2704	A2705	I2706	A2707	G2708	L2710	P2711	P2712	D2713	Y2714	V2715	D2716	R2717	C2718	V2719	S2720	S2721	L2722	P2723	L2724	G2725	L2726	T2727	E2728	S2729	L2730	T2731	R2732	A2733	E2734	K2735	L2736	L2737	L2738	L2739	L2740	E2741	T2742	L2743	N2744	V2745	L2746	L2747	P2748	E2749	K2750	L2751	D2752	S2753	F2754	L2755	N2756	K2757	P2758	A2759	E2760	Y2761	T2762	E2763	H2764	K2765	W2766	A2767	L2768	D2769	K2770	I2771	D2772	N2773	M2774	S2775	K2776	Y2777	G2778	E2779	M2780	A2781	L2782	Y2783	Y2784	L2785	K2786	L2787	H2788	P2789	M2790	L2791	R2792	P2793	Y2794
K2795	T2796	F2797	S2798	E2799	K2800	D2801	K2802	E2803	L2804	Y2805	R2806	W2807	P2808	L2809	K2810	S2811	E2812	L2813	K2814	A2815	R2816	L2817	A2818	W2819	E2820	W2821	L2822	L2823	E2824	K2825	A2826	R2827	E2828	G2829	E2830	GLU	GLU	ARG	THR	GLU	LYS	LYS	THR	ARG	LYS	ARG	LYS	ILE	LYS	SER	GLN	THR	ALA	GLN	THR	TVR	ASP	PRO	ARG	GLU	GLY	N2834																																																							
Y2855	N2856	P2857	Q2858	P2859	P2860	D2861	L2862	S2863	Q2864	V2865	T2866	L2867	S2868	R2869	E2870	L2871	Q2872	A2873	M2874	E2875	Q2877	L2878	A2879	E2880	N2881	Y2882	H2883	N2884	T2885	W2886	Q2887	R2888	K2889	K2890	K2891	Q2892	E2893	L2894	E2895	A2896	K2897	G2898	G2899	Q2900	T2901	H2902	P2903	L2904	L2905	V2906	P2907	Y2908	D2909	T2910	L2911	T2912	A2913	K2914																																																											
E2915	K2916	A2917	R2918	D2919	R2920	E2921	K2922	G2923	Q2924	E2925	L2926	K2927	K2928	F2929	L2930	Q2931	M2932	N2933	G2934	Y2935	A2936	V2937	T2938	R2939	GLY	LEU	LYS	ASP	MET	GLU	L2946	D2947	T2948	S2949	S2950	L2951	K2953	R2954	F2955	F2957	G2958	F2959	L2960	L2963	L2964	R2965	W2966	K2967	D2968	L2969	G3028	G3029	H3030	A3031	S3032	L2973	T2974	A2975																																																											
H2976	L2977	E2978	A2979	V2980	V2981	S2982	S2983	G2984	R2985	V2986	E2987	K2988	S2989	P2990	H2991	E2992	Q2993	E2994	I2995	K2996	F2997	F2998	A2999	K3000	I3001	L3002	L3003	P3004	L3005	I3006	N3007	Q3008	Y3009	T3010	N3011	N3012	C3014	L3015	F2955	A2956	G2958	F2959	L2960	L3018	S3019	T3020	P3021	A3022	K3023	V3024	L3025	G3026	S3027	G3028	G3029	H3030	A3031	S3032	L2973	T2974	A2975																																																								
K3036	E3037	H3038	I3039	L3042	F3043	L3046	A3047	A3048	L3049	H3052	R3053	V3054	S3055	L3056	F3057	G3058	T3059	D3060	A3061	F3062	A3063	V3064	Y3065	M3066	C3067	L3068	H3069	I3070	L3071	A3072	R3073	S3074	L3075	D3076	A3077	R3078	T3079	V3080	M3081	K3082	S3083	G3084	F3085	E3086	I3087	V3088	K3089	A3090	G3091	L3092	R3093	S3094	F3095	F3096	E3097	S3098	E3095																																																												
A3099	S3100	E3101	D3102	T3103	E3104	K3105	M3106	V3107	E3108	N3109	L3110	R3111	L3112	G3113	K3114	V3115	S3116	GLN	ALA	ARG	THR	GLN	VAL	K3123	G3124	V3125	G3126	Q3127	N3128	L3129	T3130	Y3131	T3132	T3133	V3134	A3135	L3136	L3137	P3138	V3139	L3140	L3143	F3144	Q3145	Q3149	H3150	Q3151	G3152	G3153	D3154	D3155	V3156	I3157	L3158	D3159	D3160	V3161																																																												
C3165	Y3166	R3167	T3168	L3169	C3170	S3171	I3172	Y3173	S3174	L3175	G3176	T3177	T3178	K3179	N3180	T3181	Y3182	V3183	E3184	L3186	R3187	A3188	L3189	G3191	E3192	C3193	L3194	A3195	R3196	L3197	A3198	A3199	A3200	M3201	P3202	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	G3231	L3232	L3233	P3233	N3234	S3235	V3236	E3237	E3238	M3239	C3240	P3241	D3242	I3243	P3244	V3245	L3246	D3247	R3248	L3249	M3250	G3251	D3252	I3253	A3257	E3258	S3259	G3260	A3261	R3262	Y3263	T3264	E3265	M3266	F3267	H3268	V3269	E3270	I3271	V3272	T3273	L3274	P3275	M3276	L3277	C3278	S3279	L3281	P3282	R3283	P3284	W3285																																																												
E3286	R3287	G3288	P3289	E3290	A3291	P3292	P3293	P3294	A3295	L3296	P3297	A3298	G3299	A3300	P3301	P3302	P3303	C3304	T3305	A3306	V3307	T3308	S3309	D3310	H3311	L3312	M3313	L3316	G3317	A3318	L3319	G3320	E3321	D3322	L3323	A3324	E3325	S3326	L3327	A3328	L3329	A3330	P3331	L3332	V3333	M3334	N3335	N3336	L3337	G3338	L3339	E3340	F3341	A3342	Q3343	P3344	V3345																																																												
H3347	L3348	E3349	F3350	L3353	L3354	F3357	F3358	I3359	P3360	T3361	L3362	G3363	R3364	L3365	R3366	R3367	R3368	A3369	G3370	K3371	V3372	V3373	A3374	E3375	E3376	E3377	Q3378	L3379	R3380	L3381	E3382	A3383	K3384	A3385	E3386	A3387	E3388	E3389	G3390	E3391	L3392	L3393	V3394	R3395	D3396	L3401	R3402	R3403	D3404	L3405	V3406	A3407	V3409	F3410	L3411																																																														

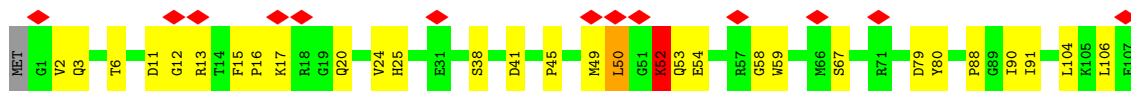


• Molecule 1: Ryanodine receptor 1

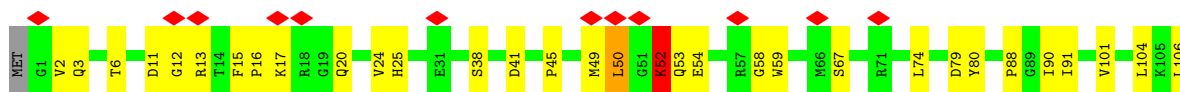




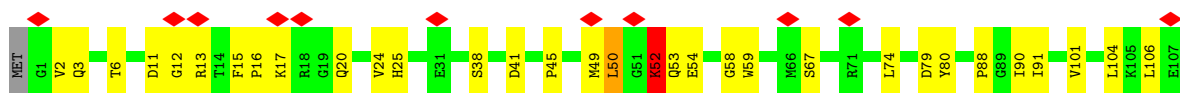
• Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1A



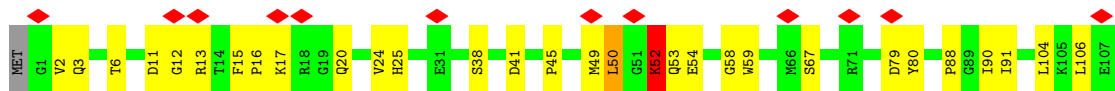
• Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1A



• Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1A



• Molecule 2: Peptidyl-prolyl cis-trans isomerase FKBP1A



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	128983	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	0.485	Depositor
Minimum map value	-0.205	Depositor
Average map value	0.001	Depositor
Map value standard deviation	0.019	Depositor
Recommended contour level	0.1	Depositor
Map size (Å)	427.52, 427.52, 427.52	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.835, 0.835, 0.835	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: IBM, CA, ATP, ZN

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	0.30	6/35977 (0.0%)	0.55	19/48726 (0.0%)
1	B	0.30	6/35977 (0.0%)	0.55	19/48726 (0.0%)
1	C	0.30	6/35977 (0.0%)	0.55	19/48726 (0.0%)
1	D	0.30	6/35977 (0.0%)	0.55	19/48726 (0.0%)
2	E	0.44	0/850	0.84	3/1146 (0.3%)
2	F	0.44	0/850	0.84	3/1146 (0.3%)
2	G	0.44	0/850	0.84	3/1146 (0.3%)
2	H	0.44	0/850	0.84	3/1146 (0.3%)
All	All	0.30	24/147308 (0.0%)	0.56	88/199488 (0.0%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	7
1	B	0	7
1	C	0	7
1	D	0	7
All	All	0	28

All (24) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	A	2237	CYS	CB-SG	6.82	1.93	1.82
1	B	2237	CYS	CB-SG	6.82	1.93	1.82
1	C	2237	CYS	CB-SG	6.82	1.93	1.82
1	D	2237	CYS	CB-SG	6.81	1.93	1.82
1	D	1999	ARG	CB-CG	6.01	1.68	1.52
1	A	1999	ARG	CB-CG	5.98	1.68	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	C	1999	ARG	CB-CG	5.98	1.68	1.52
1	B	1999	ARG	CB-CG	5.96	1.68	1.52
1	D	2233	CYS	CB-SG	-5.96	1.72	1.81
1	B	2233	CYS	CB-SG	-5.95	1.72	1.81
1	A	2233	CYS	CB-SG	-5.95	1.72	1.81
1	C	2233	CYS	CB-SG	-5.95	1.72	1.81
1	A	3899	PHE	CD1-CE1	-5.55	1.28	1.39
1	B	3899	PHE	CD1-CE1	-5.54	1.28	1.39
1	C	3899	PHE	CD1-CE1	-5.53	1.28	1.39
1	D	3899	PHE	CD1-CE1	-5.52	1.28	1.39
1	A	1036	ARG	CG-CD	-5.32	1.38	1.51
1	D	1036	ARG	CG-CD	-5.31	1.38	1.51
1	C	1036	ARG	CG-CD	-5.31	1.38	1.51
1	B	1036	ARG	CG-CD	-5.29	1.38	1.51
1	C	1982	ARG	CG-CD	-5.17	1.39	1.51
1	A	1982	ARG	CG-CD	-5.14	1.39	1.51
1	B	1982	ARG	CG-CD	-5.12	1.39	1.51
1	D	1982	ARG	CG-CD	-5.12	1.39	1.51

All (88) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	H	52	LYS	CD-CE-NZ	14.60	145.27	111.70
2	G	52	LYS	CD-CE-NZ	14.57	145.21	111.70
2	F	52	LYS	CD-CE-NZ	14.56	145.19	111.70
2	E	52	LYS	CD-CE-NZ	14.56	145.18	111.70
1	D	998	ARG	CG-CD-NE	13.13	139.38	111.80
1	A	998	ARG	CG-CD-NE	13.12	139.36	111.80
1	B	998	ARG	CG-CD-NE	13.12	139.35	111.80
1	C	998	ARG	CG-CD-NE	13.11	139.32	111.80
1	D	2237	CYS	CA-CB-SG	11.54	134.78	114.00
1	A	2237	CYS	CA-CB-SG	11.51	134.72	114.00
1	B	2237	CYS	CA-CB-SG	11.51	134.72	114.00
1	C	2237	CYS	CA-CB-SG	11.50	134.71	114.00
1	B	998	ARG	CB-CG-CD	10.45	138.77	111.60
1	A	998	ARG	CB-CG-CD	10.45	138.76	111.60
1	C	998	ARG	CB-CG-CD	10.45	138.76	111.60
1	D	998	ARG	CB-CG-CD	10.43	138.71	111.60
2	H	49	MET	CB-CG-SD	9.26	140.18	112.40
2	G	49	MET	CB-CG-SD	9.25	140.15	112.40
2	E	49	MET	CB-CG-SD	9.24	140.13	112.40
2	F	49	MET	CB-CG-SD	9.24	140.12	112.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	3111	ARG	CG-CD-NE	-8.73	93.46	111.80
1	B	3111	ARG	CG-CD-NE	-8.73	93.46	111.80
1	C	3111	ARG	CG-CD-NE	-8.73	93.46	111.80
1	D	3111	ARG	CG-CD-NE	-8.72	93.48	111.80
1	D	3250	MET	CB-CG-SD	8.22	137.06	112.40
1	A	3250	MET	CB-CG-SD	8.22	137.05	112.40
1	B	3250	MET	CB-CG-SD	8.20	137.01	112.40
1	C	3250	MET	CB-CG-SD	8.19	136.98	112.40
1	C	3756	LYS	CD-CE-NZ	-7.73	93.91	111.70
1	A	3756	LYS	CD-CE-NZ	-7.73	93.93	111.70
1	D	3756	LYS	CD-CE-NZ	-7.72	93.94	111.70
1	B	3756	LYS	CD-CE-NZ	-7.71	93.97	111.70
1	A	924	MET	CB-CG-SD	7.54	135.03	112.40
1	B	924	MET	CB-CG-SD	7.54	135.02	112.40
1	D	924	MET	CB-CG-SD	7.54	135.02	112.40
1	C	924	MET	CB-CG-SD	7.54	135.03	112.40
1	D	3111	ARG	CB-CG-CD	7.51	131.13	111.60
1	C	3111	ARG	CB-CG-CD	7.51	131.13	111.60
1	A	3111	ARG	CB-CG-CD	7.51	131.12	111.60
1	B	3111	ARG	CB-CG-CD	7.49	131.07	111.60
1	B	941	MET	CB-CG-SD	7.30	134.31	112.40
1	D	941	MET	CB-CG-SD	7.29	134.27	112.40
1	A	941	MET	CB-CG-SD	7.29	134.27	112.40
1	C	941	MET	CB-CG-SD	7.28	134.24	112.40
2	F	50	LEU	CB-CG-CD2	6.90	122.73	111.00
2	E	50	LEU	CB-CG-CD2	6.89	122.72	111.00
2	H	50	LEU	CB-CG-CD2	6.88	122.69	111.00
2	G	50	LEU	CB-CG-CD2	6.87	122.68	111.00
1	D	1994	ARG	CG-CD-NE	6.80	126.08	111.80
1	B	1994	ARG	CG-CD-NE	6.79	126.05	111.80
1	A	1994	ARG	CG-CD-NE	6.78	126.04	111.80
1	C	1994	ARG	CG-CD-NE	6.76	126.01	111.80
1	D	2233	CYS	CA-CB-SG	6.50	125.70	114.00
1	B	2233	CYS	CA-CB-SG	6.50	125.69	114.00
1	A	2233	CYS	CA-CB-SG	6.49	125.68	114.00
1	C	2233	CYS	CA-CB-SG	6.45	125.62	114.00
1	B	4952	GLU	CA-CB-CG	6.38	127.42	113.40
1	D	4952	GLU	CA-CB-CG	6.38	127.43	113.40
1	A	4952	GLU	CA-CB-CG	6.37	127.42	113.40
1	C	4952	GLU	CA-CB-CG	6.37	127.42	113.40
1	B	3756	LYS	CB-CG-CD	6.10	127.47	111.60
1	C	3756	LYS	CB-CG-CD	6.09	127.45	111.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	3756	LYS	CB-CG-CD	6.09	127.44	111.60
1	D	3756	LYS	CB-CG-CD	6.08	127.41	111.60
1	C	2330	ARG	NE-CZ-NH1	6.04	123.32	120.30
1	D	2330	ARG	NE-CZ-NH1	5.99	123.29	120.30
1	A	2330	ARG	NE-CZ-NH1	5.96	123.28	120.30
1	B	2330	ARG	NE-CZ-NH1	5.86	123.23	120.30
1	B	2233	CYS	N-CA-CB	5.74	120.93	110.60
1	C	2233	CYS	N-CA-CB	5.73	120.91	110.60
1	A	2233	CYS	N-CA-CB	5.73	120.91	110.60
1	D	2233	CYS	N-CA-CB	5.72	120.89	110.60
1	A	1979	LEU	CB-CG-CD2	5.69	120.68	111.00
1	B	1979	LEU	CB-CG-CD2	5.68	120.66	111.00
1	D	1979	LEU	CB-CG-CD2	5.68	120.66	111.00
1	C	1979	LEU	CB-CG-CD2	5.67	120.63	111.00
1	B	3111	ARG	CA-CB-CG	-5.41	101.50	113.40
1	A	3111	ARG	CA-CB-CG	-5.40	101.52	113.40
1	D	3111	ARG	CA-CB-CG	-5.39	101.53	113.40
1	C	3111	ARG	CA-CB-CG	-5.38	101.57	113.40
1	B	1036	ARG	CD-NE-CZ	-5.31	116.16	123.60
1	D	1036	ARG	CD-NE-CZ	-5.31	116.17	123.60
1	A	1036	ARG	CD-NE-CZ	-5.28	116.21	123.60
1	C	1036	ARG	CD-NE-CZ	-5.28	116.21	123.60
1	B	1999	ARG	CD-NE-CZ	-5.23	116.27	123.60
1	D	1999	ARG	CD-NE-CZ	-5.20	116.32	123.60
1	A	1999	ARG	CD-NE-CZ	-5.20	116.32	123.60
1	C	1999	ARG	CD-NE-CZ	-5.16	116.37	123.60

There are no chirality outliers.

All (28) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	1994	ARG	Sidechain
1	A	1996	ARG	Sidechain
1	A	1999	ARG	Sidechain
1	A	2237	CYS	Peptide
1	A	2947	ASP	Peptide
1	A	3111	ARG	Sidechain
1	A	421	PHE	Sidechain
1	B	1994	ARG	Sidechain
1	B	1996	ARG	Sidechain
1	B	1999	ARG	Sidechain
1	B	2237	CYS	Peptide

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Mol	Chain	Res	Type	Group
1	B	2947	ASP	Peptide
1	B	3111	ARG	Sidechain
1	B	421	PHE	Sidechain
1	C	1994	ARG	Sidechain
1	C	1996	ARG	Sidechain
1	C	1999	ARG	Sidechain
1	C	2237	CYS	Peptide
1	C	2947	ASP	Peptide
1	C	3111	ARG	Sidechain
1	C	421	PHE	Sidechain
1	D	1994	ARG	Sidechain
1	D	1996	ARG	Sidechain
1	D	1999	ARG	Sidechain
1	D	2237	CYS	Peptide
1	D	2947	ASP	Peptide
1	D	3111	ARG	Sidechain
1	D	421	PHE	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	34793	805	0
1	B	35150	0	34793	818	0
1	C	35150	0	34793	802	0
1	D	35150	0	34793	798	0
2	E	831	0	831	18	0
2	F	831	0	831	21	0
2	G	831	0	831	20	0
2	H	831	0	831	19	0
3	A	31	0	12	1	0
3	B	31	0	12	1	0
3	C	31	0	12	1	0
3	D	31	0	12	1	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

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
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	16	0	14	2	0
6	B	16	0	14	2	0
6	C	16	0	14	2	0
6	D	16	0	14	2	0
All	All	144120	0	142600	3250	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 11.

All (3250) 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:1032:LYS:HB3	1:D:1036:ARG:HH12	1.14	1.13
1:A:935:LEU:HD13	1:A:987:ARG:HE	1.16	1.10
1:C:1032:LYS:HB3	1:C:1036:ARG:HH12	1.14	1.09
1:B:935:LEU:HD13	1:B:987:ARG:HE	1.16	1.08
1:A:1032:LYS:HB3	1:A:1036:ARG:HH12	1.14	1.08
1:B:1032:LYS:HB3	1:B:1036:ARG:HH12	1.14	1.08
1:D:935:LEU:HD13	1:D:987:ARG:HE	1.16	1.05
1:C:935:LEU:HD13	1:C:987:ARG:HE	1.16	1.05
1:A:2868:SER:OG	1:A:2870[B]:GLU:OE2	1.83	0.96
1:B:2868:SER:OG	1:B:2870[B]:GLU:OE2	1.83	0.96
1:C:2868:SER:OG	1:C:2870[B]:GLU:OE2	1.83	0.96
1:D:2868:SER:OG	1:D:2870[B]:GLU:OE2	1.83	0.94
1:D:1032:LYS:CB	1:D:1036:ARG:HH12	1.85	0.90
1:C:1032:LYS:CB	1:C:1036:ARG:HH12	1.85	0.90
1:B:1032:LYS:CB	1:B:1036:ARG:HH12	1.85	0.90
1:D:1032:LYS:HB3	1:D:1036:ARG:NH1	1.87	0.90
1:C:1032:LYS:HB3	1:C:1036:ARG:NH1	1.87	0.90
1:B:1032:LYS:HB3	1:B:1036:ARG:NH1	1.87	0.90
1:A:1032:LYS:CB	1:A:1036:ARG:HH12	1.85	0.90
1:B:935:LEU:HD13	1:B:987:ARG:NE	1.88	0.90
1:A:1032:LYS:HB3	1:A:1036:ARG:NH1	1.87	0.89
1:A:2452:ARG:NH1	1:B:144:GLU:OE1	2.05	0.89
1:C:935:LEU:HD13	1:C:987:ARG:NE	1.88	0.89
1:A:1561:VAL:HG12	1:A:1562:ILE:HG23	1.56	0.88
1:D:935:LEU:HD13	1:D:987:ARG:NE	1.88	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3817:LEU:HB2	1:B:3899:PHE:HE1	1.39	0.88
1:C:3817:LEU:HB2	1:C:3899:PHE:HE1	1.38	0.87
1:D:144:GLU:OE1	1:C:2452:ARG:NH1	2.06	0.87
1:A:935:LEU:HD13	1:A:987:ARG:NE	1.88	0.87
1:B:2452:ARG:NH1	1:C:144:GLU:OE1	2.07	0.86
1:D:1561:VAL:HG12	1:D:1562:ILE:HG23	1.56	0.86
1:C:1561:VAL:HG12	1:C:1562:ILE:HG23	1.56	0.86
1:A:3817:LEU:HB2	1:A:3899:PHE:HE1	1.38	0.85
1:A:3300:ALA:HB3	1:A:3301:PRO:HD3	1.57	0.85
1:B:1561:VAL:HG12	1:B:1562:ILE:HG23	1.56	0.85
1:D:3300:ALA:HB3	1:D:3301:PRO:HD3	1.57	0.85
1:C:3733:CYS:O	1:C:3766:GLN:NE2	2.09	0.85
1:B:4902:GLU:O	1:B:4913:ARG:NH1	2.10	0.85
1:C:4902:GLU:O	1:C:4913:ARG:NH1	2.10	0.85
1:B:3733:CYS:O	1:B:3766:GLN:NE2	2.09	0.85
1:D:4902:GLU:O	1:D:4913:ARG:NH1	2.10	0.85
1:A:3733:CYS:O	1:A:3766:GLN:NE2	2.09	0.84
1:B:3300:ALA:HB3	1:B:3301:PRO:HD3	1.57	0.84
1:D:3733:CYS:O	1:D:3766:GLN:NE2	2.09	0.84
1:A:4902:GLU:O	1:A:4913:ARG:NH1	2.10	0.84
1:C:3391:GLU:OE2	1:C:3450:ASN:ND2	2.11	0.84
1:D:3391:GLU:OE2	1:D:3450:ASN:ND2	2.11	0.84
1:D:3817:LEU:HB2	1:D:3899:PHE:HE1	1.38	0.84
1:C:3300:ALA:HB3	1:C:3301:PRO:HD3	1.57	0.84
1:A:3391:GLU:OE2	1:A:3450:ASN:ND2	2.11	0.84
1:C:667:MET:SD	1:C:790:ARG:NH2	2.51	0.84
1:B:884:LEU:HD22	1:B:955:LEU:HD11	1.60	0.83
1:B:667:MET:SD	1:B:790:ARG:NH2	2.51	0.83
1:D:1156:THR:OG1	1:D:1157:GLU:OE1	1.97	0.83
1:B:3391:GLU:OE2	1:B:3450:ASN:ND2	2.11	0.83
1:A:667:MET:SD	1:A:790:ARG:NH2	2.51	0.82
1:B:2534:ALA:HB1	1:B:2588:ARG:HE	1.44	0.82
1:C:884:LEU:HD22	1:C:955:LEU:HD11	1.60	0.82
1:A:144:GLU:OE1	1:D:2452:ARG:NH1	2.13	0.82
1:A:884:LEU:HD22	1:A:955:LEU:HD11	1.60	0.82
1:D:667:MET:SD	1:D:790:ARG:NH2	2.51	0.82
1:B:1156:THR:OG1	1:B:1157:GLU:OE1	1.97	0.82
1:D:884:LEU:HD22	1:D:955:LEU:HD11	1.60	0.82
1:A:2534:ALA:HB1	1:A:2588:ARG:HE	1.44	0.82
1:C:2534:ALA:HB1	1:C:2588:ARG:HE	1.44	0.82
1:A:1156:THR:OG1	1:A:1157:GLU:OE1	1.97	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:110:ARG:NH1	1:D:117:TYR:OH	2.14	0.81
1:D:2534:ALA:HB1	1:D:2588:ARG:HE	1.44	0.81
1:D:3713:LYS:NZ	1:D:3715:LYS:O	2.14	0.81
1:C:3713:LYS:NZ	1:C:3715:LYS:O	2.14	0.81
1:B:110:ARG:NH1	1:B:117:TYR:OH	2.14	0.81
1:C:110:ARG:NH1	1:C:117:TYR:OH	2.14	0.80
1:C:1156:THR:OG1	1:C:1157:GLU:OE1	1.97	0.80
1:B:1037:ASP:O	1:B:1041:GLN:NE2	2.15	0.80
1:C:1037:ASP:O	1:C:1041:GLN:NE2	2.15	0.80
1:D:1037:ASP:O	1:D:1041:GLN:NE2	2.15	0.80
1:A:110:ARG:NH1	1:A:117:TYR:OH	2.14	0.79
1:A:4899:ASP:OD1	1:D:4892:ARG:NH1	2.15	0.79
1:A:3713:LYS:NZ	1:A:3715:LYS:O	2.14	0.79
1:B:396:GLU:OE2	1:B:470:SER:OG	2.00	0.79
1:A:1037:ASP:O	1:A:1041:GLN:NE2	2.15	0.79
1:A:4744:ASP:OD2	1:A:4747:SER:OG	2.00	0.79
1:C:4744:ASP:OD2	1:C:4747:SER:OG	2.00	0.79
1:A:2694:GLU:OE1	1:A:2697:ARG:NH2	2.16	0.79
1:D:4744:ASP:OD2	1:D:4747:SER:OG	2.00	0.79
1:C:396:GLU:OE2	1:C:470:SER:OG	2.00	0.79
1:A:396:GLU:OE2	1:A:470:SER:OG	2.00	0.78
1:B:2382:GLU:OE1	1:B:2385:ARG:NH1	2.17	0.78
1:A:1263:THR:OG1	1:A:1265:ASP:OD1	2.00	0.78
1:D:396:GLU:OE2	1:D:470:SER:OG	2.00	0.78
1:C:2382:GLU:OE1	1:C:2385:ARG:NH1	2.17	0.78
1:C:1978:ALA:HB1	1:C:1982:ARG:HH12	1.47	0.78
2:F:11:ASP:OD2	2:F:67:SER:OG	2.02	0.78
1:D:2382:GLU:OE1	1:D:2385:ARG:NH1	2.17	0.78
1:D:2694:GLU:OE1	1:D:2697:ARG:NH2	2.16	0.78
1:B:1263:THR:OG1	1:B:1265:ASP:OD1	2.00	0.78
1:B:2694:GLU:OE1	1:B:2697:ARG:NH2	2.16	0.78
1:C:2694:GLU:OE1	1:C:2697:ARG:NH2	2.16	0.78
1:B:3713:LYS:NZ	1:B:3715:LYS:O	2.14	0.78
1:C:1263:THR:OG1	1:C:1265:ASP:OD1	2.00	0.78
1:D:935:LEU:CD1	1:D:987:ARG:HE	1.95	0.77
1:A:935:LEU:CD1	1:A:987:ARG:HE	1.95	0.77
1:A:1978:ALA:HB1	1:A:1982:ARG:HH12	1.48	0.77
2:E:11:ASP:OD2	2:E:67:SER:OG	2.02	0.77
1:D:1263:THR:OG1	1:D:1265:ASP:OD1	2.00	0.77
1:A:2382:GLU:OE1	1:A:2385:ARG:NH1	2.17	0.77
1:B:4744:ASP:OD2	1:B:4747:SER:OG	2.00	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:11:ASP:OD2	2:G:67:SER:OG	2.02	0.77
1:B:883:ALA:HB1	1:B:907:LEU:HD12	1.67	0.77
1:B:1978:ALA:HB1	1:B:1982:ARG:HH12	1.47	0.77
1:B:4892:ARG:NH1	1:C:4899:ASP:OD1	2.17	0.77
1:A:883:ALA:HB1	1:A:907:LEU:HD12	1.67	0.76
1:A:1821:ASP:OD1	1:A:1822:GLY:N	2.18	0.76
2:H:11:ASP:OD2	2:H:67:SER:OG	2.02	0.76
1:D:1978:ALA:HB1	1:D:1982:ARG:HH12	1.47	0.76
1:C:4885:PHE:O	1:C:4889:VAL:HG22	1.86	0.76
1:D:4899:ASP:OD1	1:C:4892:ARG:NH1	2.17	0.76
1:D:1821:ASP:OD1	1:D:1822:GLY:N	2.18	0.76
1:A:1996:ARG:CZ	1:A:1999:ARG:HH12	1.99	0.76
1:B:3107:VAL:HB	1:B:3111:ARG:HH12	1.51	0.76
1:B:4933:GLN:OE1	1:B:4933:GLN:N	2.19	0.76
1:D:4885:PHE:O	1:D:4889:VAL:HG22	1.86	0.76
1:C:4933:GLN:OE1	1:C:4933:GLN:N	2.19	0.76
1:D:1996:ARG:CZ	1:D:1999:ARG:HH12	1.99	0.75
1:B:1821:ASP:OD1	1:B:1822:GLY:N	2.18	0.75
1:A:4885:PHE:O	1:A:4889:VAL:HG22	1.86	0.75
1:D:266:ARG:O	1:D:270:SER:OG	2.04	0.75
1:D:883:ALA:HB1	1:D:907:LEU:HD12	1.67	0.75
1:C:1821:ASP:OD1	1:C:1822:GLY:N	2.18	0.75
1:B:2777:TYR:HD1	1:B:2791:LEU:HD12	1.52	0.75
1:C:421:PHE:CE2	1:C:507:ALA:HB2	2.22	0.75
1:B:421:PHE:CE2	1:B:507:ALA:HB2	2.22	0.75
1:C:266:ARG:O	1:C:270:SER:OG	2.04	0.75
1:C:935:LEU:CD1	1:C:987:ARG:HE	1.95	0.75
1:C:2777:TYR:HD1	1:C:2791:LEU:HD12	1.52	0.75
1:B:4885:PHE:O	1:B:4889:VAL:HG22	1.86	0.75
1:C:1996:ARG:CZ	1:C:1999:ARG:HH12	1.99	0.75
1:D:3107:VAL:HB	1:D:3111:ARG:HH12	1.51	0.75
1:B:935:LEU:CD1	1:B:987:ARG:HE	1.95	0.75
1:B:3172:ILE:HA	1:B:3175:LEU:HD23	1.68	0.75
1:B:2785:LEU:HB3	1:B:2787:THR:HG23	1.69	0.74
1:D:421:PHE:CE2	1:D:507:ALA:HB2	2.22	0.74
1:C:3817:LEU:HB2	1:C:3899:PHE:CE1	2.22	0.74
1:A:421:PHE:CE2	1:A:507:ALA:HB2	2.22	0.74
1:A:3107:VAL:HB	1:A:3111:ARG:NH1	2.02	0.74
1:B:2946:LEU:O	1:B:2946:LEU:HD23	1.87	0.74
1:D:4933:GLN:N	1:D:4933:GLN:OE1	2.19	0.74
1:B:266:ARG:O	1:B:270:SER:OG	2.04	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2946:LEU:O	1:D:2946:LEU:HD23	1.87	0.74
1:D:3172:ILE:HA	1:D:3175:LEU:HD23	1.69	0.74
1:C:2946:LEU:HD23	1:C:2946:LEU:O	1.87	0.74
1:A:266:ARG:O	1:A:270:SER:OG	2.04	0.74
1:A:2777:TYR:HD1	1:A:2791:LEU:HD12	1.52	0.74
1:B:3107:VAL:HB	1:B:3111:ARG:NH1	2.02	0.74
1:B:3817:LEU:HB2	1:B:3899:PHE:CE1	2.22	0.74
1:B:950:LEU:HD13	1:B:970:LEU:HG	1.69	0.74
1:D:3817:LEU:HB2	1:D:3899:PHE:CE1	2.22	0.74
1:A:2785:LEU:HB3	1:A:2787:THR:HG23	1.69	0.74
1:C:3107:VAL:HB	1:C:3111:ARG:HH12	1.51	0.74
1:A:950:LEU:HD13	1:A:970:LEU:HG	1.69	0.74
1:C:883:ALA:HB1	1:C:907:LEU:HD12	1.67	0.74
1:C:950:LEU:HD13	1:C:970:LEU:HG	1.69	0.74
1:C:3172:ILE:HA	1:C:3175:LEU:HD23	1.69	0.74
1:A:3817:LEU:HB2	1:A:3899:PHE:CE1	2.22	0.73
1:A:4933:GLN:OE1	1:A:4933:GLN:N	2.19	0.73
1:D:2777:TYR:HD1	1:D:2791:LEU:HD12	1.52	0.73
1:C:1927:LEU:HD22	1:C:2101:MET:SD	2.28	0.73
1:A:3107:VAL:HB	1:A:3111:ARG:HH12	1.51	0.73
1:B:1996:ARG:CZ	1:B:1999:ARG:HH12	1.99	0.73
1:D:2233:CYS:O	1:D:2234:ARG:C	2.27	0.73
1:A:2946:LEU:O	1:A:2946:LEU:HD23	1.87	0.73
1:D:3107:VAL:HB	1:D:3111:ARG:NH1	2.02	0.73
1:C:2233:CYS:O	1:C:2234:ARG:C	2.26	0.73
1:A:2011:HIS:NE2	1:A:2017:ASP:OD2	2.21	0.73
1:B:1927:LEU:HD22	1:B:2101:MET:SD	2.28	0.73
1:D:950:LEU:HD13	1:D:970:LEU:HG	1.69	0.73
1:D:1032:LYS:C	1:D:1036:ARG:NH1	2.42	0.73
1:D:2011:HIS:NE2	1:D:2017:ASP:OD2	2.21	0.73
1:C:2785:LEU:HB3	1:C:2787:THR:HG23	1.69	0.73
1:A:1927:LEU:HD22	1:A:2101:MET:SD	2.28	0.73
1:C:3107:VAL:HB	1:C:3111:ARG:NH1	2.02	0.73
1:A:3172:ILE:HA	1:A:3175:LEU:HD23	1.68	0.73
1:D:1927:LEU:HD22	1:D:2101:MET:SD	2.28	0.73
1:C:2644:LEU:HD13	1:C:2678:LEU:HD21	1.71	0.73
1:A:1032:LYS:C	1:A:1036:ARG:NH1	2.42	0.73
1:B:1032:LYS:C	1:B:1036:ARG:NH1	2.42	0.72
1:B:1743[A]:ARG:NH2	1:B:1967:ASP:OD2	2.22	0.72
1:B:2644:LEU:HD13	1:B:2678:LEU:HD21	1.71	0.72
1:A:2226:PRO:O	1:A:2267:MET:HE1	1.89	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2226:PRO:O	1:C:2267:MET:HE1	1.89	0.72
1:D:4069:LYS:NZ	1:D:4130:ASN:OD1	2.23	0.72
1:A:3627:GLN:O	1:A:3630:ARG:NH1	2.23	0.72
1:B:2011:HIS:NE2	1:B:2017:ASP:OD2	2.21	0.72
1:D:3627:GLN:O	1:D:3630:ARG:NH1	2.23	0.72
1:C:1032:LYS:C	1:C:1036:ARG:NH1	2.42	0.72
1:C:2011:HIS:NE2	1:C:2017:ASP:OD2	2.21	0.72
1:A:2233:CYS:O	1:A:2234:ARG:C	2.26	0.72
1:B:2233:CYS:O	1:B:2234:ARG:C	2.27	0.72
1:D:2226:PRO:O	1:D:2267:MET:HE1	1.90	0.72
1:D:2644:LEU:HD13	1:D:2678:LEU:HD21	1.71	0.72
1:D:2785:LEU:HB3	1:D:2787:THR:HG23	1.69	0.72
1:A:4069:LYS:NZ	1:A:4130:ASN:OD1	2.23	0.72
1:B:3627:GLN:O	1:B:3630:ARG:NH1	2.23	0.72
1:D:1743[A]:ARG:NH2	1:D:1967:ASP:OD2	2.22	0.72
1:D:2265:LEU:HB2	1:D:2330:ARG:HH21	1.55	0.72
1:C:1743[A]:ARG:NH2	1:C:1967:ASP:OD2	2.22	0.72
1:A:2644:LEU:HD13	1:A:2678:LEU:HD21	1.71	0.71
1:B:835:ARG:NH2	1:B:1210:SER:O	2.23	0.71
1:B:2226:PRO:O	1:B:2267:MET:HE1	1.89	0.71
1:C:221:ARG:NH1	1:C:253:CYS:O	2.24	0.71
1:D:1996:ARG:NH2	1:D:1999:ARG:HH12	1.89	0.71
1:A:3514:LEU:HD21	1:A:3602:VAL:CG1	2.21	0.71
1:B:1996:ARG:NH2	1:B:1999:ARG:HH12	1.89	0.71
1:B:3514:LEU:HD21	1:B:3602:VAL:CG1	2.21	0.71
1:D:221:ARG:NH1	1:D:253:CYS:O	2.24	0.71
1:C:1996:ARG:NH2	1:C:1999:ARG:HH12	1.89	0.71
1:D:1112:ASP:OD1	1:D:1113:VAL:N	2.24	0.71
1:C:1112:ASP:OD1	1:C:1113:VAL:N	2.24	0.71
1:C:3514:LEU:HD21	1:C:3602:VAL:CG1	2.21	0.71
1:A:1112:ASP:OD1	1:A:1113:VAL:N	2.24	0.71
1:A:1743[A]:ARG:NH2	1:A:1967:ASP:OD2	2.22	0.71
1:B:1749:PRO:HD2	1:B:1758:ARG:NH1	2.06	0.71
1:A:221:ARG:NH1	1:A:253:CYS:O	2.24	0.71
1:D:835:ARG:NH2	1:D:1210:SER:O	2.23	0.71
1:D:1749:PRO:HD2	1:D:1758:ARG:NH1	2.06	0.71
1:A:4063:ASP:OD1	1:A:4064:MET:N	2.24	0.71
1:C:835:ARG:NH2	1:C:1210:SER:O	2.23	0.71
1:B:4069:LYS:NZ	1:B:4130:ASN:OD1	2.23	0.70
1:A:835:ARG:NH2	1:A:1210:SER:O	2.23	0.70
1:A:1749:PRO:HD2	1:A:1758:ARG:NH1	2.06	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1996:ARG:NH2	1:A:1999:ARG:HH12	1.89	0.70
1:A:2265:LEU:HB2	1:A:2330:ARG:HH21	1.55	0.70
1:C:1749:PRO:HD2	1:C:1758:ARG:NH1	2.06	0.70
1:D:3514:LEU:HD21	1:D:3602:VAL:CG1	2.21	0.70
1:C:3627:GLN:O	1:C:3630:ARG:NH1	2.23	0.70
1:B:221:ARG:NH1	1:B:253:CYS:O	2.24	0.70
1:B:4063:ASP:OD1	1:B:4064:MET:N	2.24	0.70
1:C:2265:LEU:HB2	1:C:2330:ARG:HH21	1.55	0.70
1:A:3699:HIS:CE1	1:A:3703:LEU:HD11	2.27	0.70
1:C:4063:ASP:OD1	1:C:4064:MET:N	2.24	0.70
1:A:2980:VAL:HG21	1:A:2986:VAL:HG12	1.74	0.70
1:D:4063:ASP:OD1	1:D:4064:MET:N	2.24	0.70
1:D:2980:VAL:HG21	1:D:2986:VAL:HG12	1.74	0.70
1:C:2327:GLY:HA2	1:C:2330:ARG:CZ	2.22	0.70
1:C:4722:ARG:HE	1:C:4748:LEU:HD11	1.55	0.70
1:A:4892:ARG:NH1	1:B:4899:ASP:OD1	2.24	0.70
1:B:1112:ASP:OD1	1:B:1113:VAL:N	2.24	0.70
1:D:4722:ARG:HE	1:D:4748:LEU:HD11	1.55	0.70
1:B:2265:LEU:HB2	1:B:2330:ARG:HH21	1.55	0.69
1:D:3699:HIS:CE1	1:D:3703:LEU:HD11	2.27	0.69
1:A:393:CYS:SG	1:A:398:SER:OG	2.50	0.69
1:A:4722:ARG:HE	1:A:4748:LEU:HD11	1.55	0.69
1:B:393:CYS:SG	1:B:398:SER:OG	2.50	0.69
1:B:2980:VAL:HG21	1:B:2986:VAL:HG12	1.74	0.69
1:C:1999:ARG:O	1:C:3638:MET:CE	2.40	0.69
1:A:1999:ARG:O	1:A:3638:MET:CE	2.40	0.69
1:D:2514:ASN:ND2	1:D:2516:ASP:OD1	2.26	0.69
1:C:3699:HIS:CE1	1:C:3703:LEU:HD11	2.27	0.69
1:C:4158:PRO:HA	1:C:4161:ARG:HE	1.58	0.69
1:B:1999:ARG:O	1:B:3638:MET:CE	2.40	0.69
1:D:2233:CYS:O	1:D:2235:PHE:N	2.26	0.69
1:C:393:CYS:SG	1:C:398:SER:OG	2.50	0.69
1:B:4158:PRO:HA	1:B:4161:ARG:HE	1.58	0.69
1:C:2166:LEU:HD11	1:C:2206:THR:HG23	1.75	0.69
1:A:2514:ASN:ND2	1:A:2516:ASP:OD1	2.26	0.69
1:A:2869:ARG:HD3	1:A:2947:ASP:OD2	1.93	0.69
1:B:2327:GLY:HA2	1:B:2330:ARG:CZ	2.22	0.69
1:B:4722:ARG:HE	1:B:4748:LEU:HD11	1.56	0.69
1:D:393:CYS:SG	1:D:398:SER:OG	2.50	0.69
1:D:1999:ARG:O	1:D:3638:MET:CE	2.40	0.69
1:D:2327:GLY:HA2	1:D:2330:ARG:CZ	2.22	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2514:ASN:ND2	1:C:2516:ASP:OD1	2.25	0.69
1:B:2514:ASN:ND2	1:B:2516:ASP:OD1	2.26	0.69
1:B:3699:HIS:CE1	1:B:3703:LEU:HD11	2.27	0.69
1:C:2233:CYS:O	1:C:2235:PHE:N	2.26	0.69
1:A:2327:GLY:HA2	1:A:2330:ARG:CZ	2.22	0.68
1:C:975:VAL:HG13	1:C:1047:LEU:HD23	1.76	0.68
1:C:4069:LYS:NZ	1:C:4130:ASN:OD1	2.23	0.68
1:B:2869:ARG:HD3	1:B:2947:ASP:OD2	1.93	0.68
1:D:887:ILE:HD11	1:D:907:LEU:HD11	1.75	0.68
1:A:887:ILE:HD11	1:A:907:LEU:HD11	1.75	0.68
1:A:2233:CYS:O	1:A:2235:PHE:N	2.26	0.68
1:B:975:VAL:HG13	1:B:1047:LEU:HD23	1.76	0.68
1:A:975:VAL:HG13	1:A:1047:LEU:HD23	1.76	0.68
1:B:2233:CYS:O	1:B:2235:PHE:N	2.26	0.68
1:D:975:VAL:HG13	1:D:1047:LEU:HD23	1.76	0.68
1:D:2869:ARG:HD3	1:D:2947:ASP:OD2	1.93	0.68
1:C:2980:VAL:HG21	1:C:2986:VAL:HG12	1.74	0.68
1:A:1577:ALA:O	1:A:1584:ARG:NH1	2.27	0.68
1:B:2135:LEU:HD12	1:B:3658:LYS:HZ3	1.57	0.68
1:D:2166:LEU:HD11	1:D:2206:THR:HG23	1.75	0.68
1:B:2166:LEU:HD11	1:B:2206:THR:HG23	1.75	0.67
1:C:2675:THR:HB	1:C:2710:LEU:HD21	1.76	0.67
1:C:2869:ARG:HD3	1:C:2947:ASP:OD2	1.93	0.67
1:C:4998:LYS:HB3	1:C:5003:HIS:HE1	1.60	0.67
1:A:4158:PRO:HA	1:A:4161:ARG:HE	1.58	0.67
1:D:4655:PHE:CA	1:D:4796:MET:HE1	2.25	0.67
1:C:1101:ARG:NH1	1:C:1115:LEU:O	2.28	0.67
1:C:4827:LEU:O	1:C:4831:THR:HG23	1.94	0.67
1:A:4929:LEU:O	1:A:4933:GLN:OE1	2.13	0.67
1:A:4966:ASP:OD1	1:A:4967:TYR:N	2.28	0.67
1:C:2747:ILE:HD11	1:C:2751:LEU:HD22	1.77	0.67
1:A:2675:THR:HB	1:A:2710:LEU:HD21	1.76	0.67
1:B:4966:ASP:OD1	1:B:4967:TYR:N	2.28	0.67
1:B:4998:LYS:HB3	1:B:5003:HIS:HE1	1.60	0.67
1:D:4827:LEU:O	1:D:4831:THR:HG23	1.94	0.67
1:A:2166:LEU:HD11	1:A:2206:THR:HG23	1.75	0.67
1:B:4827:LEU:O	1:B:4831:THR:HG23	1.94	0.67
1:B:1577:ALA:O	1:B:1584:ARG:NH1	2.27	0.67
1:A:3531:ASP:OD1	1:A:3532:LEU:N	2.28	0.66
1:A:4827:LEU:O	1:A:4831:THR:HG23	1.94	0.66
1:B:887:ILE:HD11	1:B:907:LEU:HD11	1.75	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1101:ARG:NH1	1:B:1115:LEU:O	2.28	0.66
1:D:4158:PRO:HA	1:D:4161:ARG:HE	1.58	0.66
1:C:3531:ASP:OD1	1:C:3532:LEU:N	2.29	0.66
1:B:2747:ILE:HD11	1:B:2751:LEU:HD22	1.77	0.66
1:A:3346:VAL:HG11	1:A:3414:ARG:HB2	1.77	0.66
1:C:887:ILE:HD11	1:C:907:LEU:HD11	1.75	0.66
1:C:2330:ARG:HA	1:C:2333:ASP:OD2	1.96	0.66
1:D:4722:ARG:HE	1:D:4748:LEU:CD1	2.08	0.66
1:D:4966:ASP:OD1	1:D:4967:TYR:N	2.28	0.66
1:D:4998:LYS:HB3	1:D:5003:HIS:HE1	1.60	0.66
1:C:228:ASP:OD1	1:C:229:GLU:N	2.29	0.66
1:C:3346:VAL:HG11	1:C:3414:ARG:HB2	1.77	0.66
1:C:4929:LEU:O	1:C:4933:GLN:OE1	2.13	0.66
1:A:2970:SER:HA	1:A:2973:PHE:CE1	2.30	0.66
1:A:3534:MET:O	1:A:3538:THR:HG23	1.96	0.66
1:B:2330:ARG:HA	1:B:2333:ASP:OD2	1.96	0.66
1:B:4929:LEU:O	1:B:4933:GLN:OE1	2.13	0.66
1:D:100:THR:HG21	1:D:162:LYS:HE3	1.78	0.66
1:C:100:THR:HG21	1:C:162:LYS:HE3	1.78	0.66
1:B:3834:ALA:O	1:B:3838:THR:HG23	1.96	0.66
1:D:1101:ARG:NH1	1:D:1115:LEU:O	2.28	0.66
1:D:2970:SER:HA	1:D:2973:PHE:CE1	2.30	0.66
1:D:4929:LEU:O	1:D:4933:GLN:OE1	2.13	0.66
1:C:1577:ALA:O	1:C:1584:ARG:NH1	2.27	0.66
1:B:3346:VAL:HG11	1:B:3414:ARG:HB2	1.77	0.66
1:D:2330:ARG:HA	1:D:2333:ASP:OD2	1.96	0.66
1:D:3346:VAL:HG11	1:D:3414:ARG:HB2	1.77	0.66
1:A:2747:ILE:HD11	1:A:2751:LEU:HD22	1.77	0.66
2:H:11:ASP:OD2	2:H:12:GLY:N	2.29	0.66
1:B:2970:SER:HA	1:B:2973:PHE:CE1	2.30	0.66
1:D:1577:ALA:O	1:D:1584:ARG:NH1	2.27	0.66
1:D:1999:ARG:O	1:D:1999:ARG:HG2	1.96	0.66
1:A:572:PRO:O	1:A:576:ASN:ND2	2.29	0.65
1:A:1101:ARG:NH1	1:A:1115:LEU:O	2.28	0.65
1:B:3111:ARG:NH2	1:B:3174:SER:OG	2.30	0.65
1:D:2675:THR:HB	1:D:2710:LEU:HD21	1.76	0.65
1:C:3534:MET:O	1:C:3538:THR:HG23	1.96	0.65
1:C:4722:ARG:HE	1:C:4748:LEU:CD1	2.08	0.65
1:C:4910:GLU:O	1:C:4914:VAL:HG13	1.96	0.65
1:B:663:TYR:HD1	1:B:747:CYS:HG	1.43	0.65
1:B:4722:ARG:HE	1:B:4748:LEU:CD1	2.08	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:228:ASP:OD1	1:D:229:GLU:N	2.29	0.65
1:C:663:TYR:HD1	1:C:747:CYS:HG	1.43	0.65
1:A:4910:GLU:O	1:A:4914:VAL:HG13	1.96	0.65
1:B:1999:ARG:O	1:B:1999:ARG:HG2	1.96	0.65
1:B:4711:PHE:HB3	1:B:4712:PRO:HD3	1.78	0.65
1:B:572:PRO:O	1:B:576:ASN:ND2	2.29	0.65
1:B:3103:ILE:O	1:B:3107:VAL:HG13	1.97	0.65
1:D:572:PRO:O	1:D:576:ASN:ND2	2.29	0.65
1:D:2747:ILE:HD11	1:D:2751:LEU:HD22	1.77	0.65
1:D:4711:PHE:HB3	1:D:4712:PRO:HD3	1.78	0.65
1:C:4966:ASP:OD1	1:C:4967:TYR:N	2.28	0.65
1:A:3103:ILE:O	1:A:3107:VAL:HG13	1.97	0.65
1:A:4998:LYS:HB3	1:A:5003:HIS:HE1	1.60	0.65
1:B:228:ASP:OD1	1:B:229:GLU:N	2.29	0.65
1:B:2675:THR:HB	1:B:2710:LEU:HD21	1.76	0.65
1:C:2520:HIS:O	1:C:2524:VAL:HG22	1.97	0.65
1:C:2970:SER:HA	1:C:2973:PHE:CE1	2.30	0.65
1:C:4655:PHE:CA	1:C:4796:MET:HE1	2.26	0.65
1:A:3834:ALA:O	1:A:3838:THR:HG23	1.96	0.65
2:G:11:ASP:OD2	2:G:12:GLY:N	2.29	0.65
1:D:3111:ARG:NH2	1:D:3174:SER:OG	2.30	0.65
1:D:3534:MET:O	1:D:3538:THR:HG23	1.96	0.65
1:D:4910:GLU:O	1:D:4914:VAL:HG13	1.96	0.65
1:A:228:ASP:OD1	1:A:229:GLU:N	2.29	0.65
1:A:1999:ARG:O	1:A:1999:ARG:HG2	1.96	0.65
1:A:2330:ARG:HA	1:A:2333:ASP:OD2	1.96	0.65
2:F:11:ASP:OD2	2:F:12:GLY:N	2.29	0.65
1:B:4655:PHE:CA	1:B:4796:MET:HE1	2.27	0.65
1:D:3531:ASP:OD1	1:D:3532:LEU:N	2.28	0.65
1:C:1999:ARG:O	1:C:1999:ARG:HG2	1.96	0.65
1:C:3111:ARG:NH2	1:C:3174:SER:OG	2.30	0.65
1:B:3531:ASP:OD1	1:B:3532:LEU:N	2.28	0.65
1:B:4910:GLU:O	1:B:4914:VAL:HG13	1.96	0.65
1:D:1996:ARG:HE	1:D:1999:ARG:NH1	1.95	0.65
1:B:100:THR:HG21	1:B:162:LYS:HE3	1.78	0.65
1:B:672:VAL:O	1:B:680:THR:OG1	2.08	0.65
1:C:1996:ARG:HE	1:C:1999:ARG:NH1	1.95	0.65
1:A:100:THR:HG21	1:A:162:LYS:HE3	1.78	0.64
1:B:581:ASN:OD1	1:B:582:HIS:N	2.30	0.64
1:C:3834:ALA:O	1:C:3838:THR:HG23	1.96	0.64
1:C:4711:PHE:HB3	1:C:4712:PRO:HD3	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3111:ARG:NH2	1:A:3174:SER:OG	2.30	0.64
1:A:4722:ARG:HE	1:A:4748:LEU:CD1	2.08	0.64
2:E:11:ASP:OD2	2:E:12:GLY:N	2.29	0.64
1:B:3110:LEU:HD23	1:B:3183:VAL:HG22	1.79	0.64
1:A:1996:ARG:HE	1:A:1999:ARG:NH1	1.95	0.64
1:A:581:ASN:OD1	1:A:582:HIS:N	2.30	0.64
1:A:418:LEU:HA	1:A:421:PHE:CE1	2.33	0.64
1:A:971:ASP:OD2	1:A:973:SER:OG	2.14	0.64
2:G:52:LYS:HB2	2:G:54:GLU:OE1	1.97	0.64
1:B:418:LEU:HA	1:B:421:PHE:CE1	2.33	0.64
1:B:2520:HIS:O	1:B:2524:VAL:HG22	1.97	0.64
1:D:3103:ILE:O	1:D:3107:VAL:HG13	1.97	0.64
1:D:3834:ALA:O	1:D:3838:THR:HG23	1.96	0.64
1:C:572:PRO:O	1:C:576:ASN:ND2	2.29	0.64
1:C:3103:ILE:O	1:C:3107:VAL:HG13	1.97	0.64
1:A:1726:SER:O	1:A:1729:SER:OG	2.16	0.64
1:A:3110:LEU:HD23	1:A:3183:VAL:HG22	1.79	0.64
1:B:1996:ARG:HE	1:B:1999:ARG:NH1	1.95	0.64
1:C:1726:SER:O	1:C:1729:SER:OG	2.16	0.64
1:D:418:LEU:HA	1:D:421:PHE:CE1	2.33	0.64
1:C:581:ASN:OD1	1:C:582:HIS:N	2.30	0.64
1:A:1454:THR:OG1	1:A:1456:ASP:OD1	2.12	0.64
2:G:25:HIS:CD2	2:G:104:LEU:HD11	2.33	0.64
1:B:4091:LYS:NZ	1:B:4121:GLU:OE1	2.29	0.64
1:D:581:ASN:OD1	1:D:582:HIS:N	2.30	0.64
1:D:1999:ARG:O	1:D:3638:MET:HE3	1.98	0.64
1:A:2894:LEU:O	1:A:2897:LYS:N	2.32	0.63
2:F:52:LYS:HB2	2:F:54:GLU:OE1	1.97	0.63
1:C:2894:LEU:O	1:C:2897:LYS:N	2.32	0.63
1:A:1065:ASN:OD1	1:A:1066:GLN:N	2.32	0.63
1:A:4711:PHE:HB3	1:A:4712:PRO:HD3	1.78	0.63
1:D:2135:LEU:HD12	1:D:3658:LYS:HZ3	1.63	0.63
1:B:1726:SER:O	1:B:1729:SER:OG	2.16	0.63
1:B:2212:VAL:HG21	1:B:2256:TYR:OH	1.99	0.63
1:B:3534:MET:O	1:B:3538:THR:HG23	1.96	0.63
1:D:2520:HIS:O	1:D:2524:VAL:HG22	1.97	0.63
1:C:3284:TRP:O	1:C:3305:THR:HG21	1.99	0.63
1:C:3310:ASP:OD1	1:C:3311:HIS:N	2.31	0.63
1:A:1931:LEU:HD13	1:A:1935:VAL:HG11	1.81	0.63
1:A:1996:ARG:NE	1:A:1999:ARG:HH12	1.97	0.63
1:A:3310:ASP:OD1	1:A:3311:HIS:N	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:52:LYS:HB2	2:E:54:GLU:OE1	1.97	0.63
1:B:3284:TRP:O	1:B:3305:THR:HG21	1.99	0.63
1:D:3284:TRP:O	1:D:3305:THR:HG21	1.99	0.63
1:C:418:LEU:HA	1:C:421:PHE:CE1	2.33	0.63
1:C:1931:LEU:HD13	1:C:1935:VAL:HG11	1.80	0.63
1:C:1999:ARG:O	1:C:3638:MET:HE3	1.97	0.63
1:A:672:VAL:O	1:A:680:THR:OG1	2.08	0.63
2:H:52:LYS:HB2	2:H:54:GLU:OE1	1.97	0.63
1:B:2894:LEU:O	1:B:2897:LYS:N	2.32	0.63
1:D:1996:ARG:NE	1:D:1999:ARG:HH12	1.96	0.63
1:C:3110:LEU:HD23	1:C:3183:VAL:HG22	1.79	0.63
1:A:1033:ARG:HA	1:A:1036:ARG:CZ	2.29	0.63
1:A:2212:VAL:HG21	1:A:2256:TYR:OH	1.99	0.63
1:A:2520:HIS:O	1:A:2524:VAL:HG22	1.97	0.63
2:H:25:HIS:CD2	2:H:104:LEU:HD11	2.33	0.63
1:B:971:ASP:OD2	1:B:973:SER:OG	2.14	0.63
1:D:1065:ASN:OD1	1:D:1066:GLN:N	2.32	0.63
1:D:2894:LEU:O	1:D:2897:LYS:N	2.31	0.63
1:C:1033:ARG:HA	1:C:1036:ARG:CZ	2.29	0.63
1:A:1999:ARG:O	1:A:3638:MET:HE1	1.99	0.63
1:A:4655:PHE:CA	1:A:4796:MET:HE1	2.29	0.63
1:D:1726:SER:O	1:D:1729:SER:OG	2.16	0.63
2:E:25:HIS:CD2	2:E:104:LEU:HD11	2.33	0.62
1:B:3310:ASP:OD1	1:B:3311:HIS:N	2.31	0.62
1:D:1033:ARG:HA	1:D:1036:ARG:CZ	2.29	0.62
1:A:3442:PHE:HE1	1:A:3511:VAL:HG12	1.64	0.62
1:B:1033:ARG:HA	1:B:1036:ARG:CZ	2.29	0.62
1:B:1996:ARG:NE	1:B:1999:ARG:HH12	1.97	0.62
1:D:3110:LEU:HD23	1:D:3183:VAL:HG22	1.79	0.62
1:C:864:PRO:HD2	1:C:867:LEU:HD12	1.81	0.62
1:C:1996:ARG:NE	1:C:1999:ARG:HH12	1.97	0.62
1:A:3086:GLU:OE2	1:A:3093:ARG:NH1	2.33	0.62
2:F:25:HIS:CD2	2:F:104:LEU:HD11	2.33	0.62
1:B:1066:GLN:O	1:B:1071:ARG:NH2	2.33	0.62
1:B:2265:LEU:HD12	1:B:2266:GLY:N	2.15	0.62
1:D:3442:PHE:HE1	1:D:3511:VAL:HG12	1.64	0.62
1:A:2265:LEU:HD12	1:A:2266:GLY:N	2.15	0.62
1:B:864:PRO:HD2	1:B:867:LEU:HD12	1.81	0.62
1:B:928:THR:O	1:B:932:LEU:HD23	2.00	0.62
1:C:2265:LEU:HB2	1:C:2330:ARG:NH2	2.14	0.62
1:C:3442:PHE:HE1	1:C:3511:VAL:HG12	1.64	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2265:LEU:HB2	1:A:2330:ARG:NH2	2.14	0.62
1:A:3284:TRP:O	1:A:3305:THR:HG21	1.99	0.62
1:D:864:PRO:HD2	1:D:867:LEU:HD12	1.81	0.62
1:D:2212:VAL:HG21	1:D:2256:TYR:OH	1.99	0.62
1:D:3310:ASP:OD1	1:D:3311:HIS:N	2.31	0.62
1:C:2212:VAL:HG21	1:C:2256:TYR:OH	1.99	0.62
1:C:2265:LEU:HD12	1:C:2266:GLY:N	2.15	0.62
1:C:3635:CYS:HA	1:C:3638:MET:HE2	1.82	0.62
1:A:885:THR:O	1:A:889:GLN:NE2	2.33	0.62
1:D:1454:THR:OG1	1:D:1456:ASP:OD1	2.12	0.62
1:C:1065:ASN:OD1	1:C:1066:GLN:N	2.32	0.62
1:C:4091:LYS:NZ	1:C:4121:GLU:OE1	2.29	0.62
1:A:4942:GLU:O	1:A:4945:ASP:OD1	2.18	0.62
1:B:3635:CYS:HA	1:B:3638:MET:HE2	1.81	0.62
1:B:4942:GLU:O	1:B:4945:ASP:OD1	2.18	0.62
1:D:885:THR:O	1:D:889:GLN:NE2	2.33	0.62
1:D:2744:ASN:OD1	1:D:2745:VAL:HG23	2.00	0.62
1:A:928:THR:O	1:A:932:LEU:HD23	2.00	0.61
1:B:1931:LEU:HD13	1:B:1935:VAL:HG11	1.80	0.61
1:B:2923:ALA:O	1:B:2927:LEU:HD23	2.00	0.61
1:D:1066:GLN:O	1:D:1071:ARG:NH2	2.33	0.61
1:C:928:THR:O	1:C:932:LEU:HD23	2.00	0.61
1:C:1066:GLN:O	1:C:1071:ARG:NH2	2.33	0.61
1:C:2923:ALA:O	1:C:2927:LEU:HD23	2.00	0.61
1:A:1066:GLN:O	1:A:1071:ARG:NH2	2.33	0.61
1:A:3984:ARG:NH2	1:B:160:GLY:O	2.30	0.61
1:B:1065:ASN:OD1	1:B:1066:GLN:N	2.32	0.61
1:D:928:THR:O	1:D:932:LEU:HD23	2.00	0.61
1:D:1931:LEU:HD13	1:D:1935:VAL:HG11	1.81	0.61
1:C:885:THR:O	1:C:889:GLN:NE2	2.33	0.61
1:A:4091:LYS:NZ	1:A:4121:GLU:OE1	2.29	0.61
1:B:973:SER:O	1:B:976:ARG:NH1	2.34	0.61
1:D:2265:LEU:HD12	1:D:2266:GLY:N	2.15	0.61
1:C:2744:ASN:OD1	1:C:2745:VAL:HG23	2.00	0.61
1:A:2923:ALA:O	1:A:2927:LEU:HD23	2.00	0.61
2:H:25:HIS:HD2	2:H:104:LEU:HD21	1.65	0.61
1:B:2265:LEU:HB2	1:B:2330:ARG:NH2	2.14	0.61
1:B:885:THR:O	1:B:889:GLN:NE2	2.33	0.61
1:B:1999:ARG:O	1:B:3638:MET:HE1	2.00	0.61
1:B:3086:GLU:OE2	1:B:3093:ARG:NH1	2.33	0.61
1:C:4942:GLU:O	1:C:4945:ASP:OD1	2.18	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4091:LYS:NZ	1:D:4121:GLU:OE1	2.29	0.61
1:A:2871:LEU:HD13	1:A:2874:MET:HE1	1.81	0.61
1:A:3635:CYS:HA	1:A:3638:MET:HE2	1.81	0.61
1:D:973:SER:O	1:D:976:ARG:NH1	2.33	0.61
1:D:2923:ALA:O	1:D:2927:LEU:HD23	2.00	0.61
1:A:864:PRO:HD2	1:A:867:LEU:HD12	1.81	0.61
1:A:2744:ASN:OD1	1:A:2745:VAL:HG23	2.00	0.61
1:D:4942:GLU:O	1:D:4945:ASP:OD1	2.18	0.61
1:C:884:LEU:CD2	1:C:955:LEU:HD11	2.31	0.61
2:F:25:HIS:HD2	2:F:104:LEU:HD21	1.65	0.61
1:D:2516:ASP:OD1	1:D:2517:PHE:N	2.33	0.61
1:D:3086:GLU:OE2	1:D:3093:ARG:NH1	2.33	0.61
1:C:973:SER:O	1:C:976:ARG:NH1	2.34	0.61
1:D:887:ILE:O	1:D:890:GLY:N	2.34	0.61
1:D:971:ASP:OD2	1:D:973:SER:OG	2.14	0.61
1:D:2265:LEU:HB2	1:D:2330:ARG:NH2	2.14	0.61
1:B:2516:ASP:OD1	1:B:2517:PHE:N	2.33	0.60
1:B:2744:ASN:OD1	1:B:2745:VAL:HG23	2.00	0.60
1:B:3442:PHE:HE1	1:B:3511:VAL:HG12	1.64	0.60
1:A:887:ILE:HD11	1:A:961:MET:HE3	1.83	0.60
1:A:2475:GLN:NE2	1:A:2476:ILE:O	2.35	0.60
1:B:887:ILE:O	1:B:890:GLY:N	2.34	0.60
1:D:2376:LEU:HD21	1:D:2502:MET:HE3	1.83	0.60
1:A:2516:ASP:OD1	1:A:2517:PHE:N	2.33	0.60
1:B:884:LEU:CD2	1:B:955:LEU:HD11	2.31	0.60
1:B:1979:LEU:HD22	1:B:1982:ARG:NH2	2.17	0.60
1:A:973:SER:O	1:A:976:ARG:NH1	2.34	0.60
2:G:25:HIS:HD2	2:G:104:LEU:HD21	1.65	0.60
1:D:3635:CYS:HA	1:D:3638:MET:HE2	1.82	0.60
1:C:887:ILE:O	1:C:890:GLY:N	2.34	0.60
1:D:2475:GLN:NE2	1:D:2476:ILE:O	2.35	0.60
1:D:3097:GLU:O	1:D:3100:SER:OG	2.19	0.60
1:A:887:ILE:O	1:A:890:GLY:N	2.34	0.60
2:E:25:HIS:HD2	2:E:104:LEU:HD21	1.65	0.60
1:B:2475:GLN:NE2	1:B:2476:ILE:O	2.35	0.60
1:D:3786:CYS:SG	1:D:3831:SER:OG	2.60	0.60
1:C:238:SER:OG	1:C:240:ASP:OD1	2.20	0.60
1:C:2376:LEU:HD21	1:C:2502:MET:HE3	1.83	0.60
1:A:3357:HIS:O	1:A:3361:THR:HG23	2.01	0.60
1:D:1520:VAL:HG12	1:D:1527:MET:HG2	1.83	0.60
1:D:2464:ASP:OD1	1:D:2465:ASP:N	2.35	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3529:ASP:OD2	1:C:3595:ARG:NH2	2.35	0.60
1:C:3786:CYS:SG	1:C:3831:SER:OG	2.60	0.60
1:D:988:LEU:HB2	1:D:1039:LEU:HD21	1.83	0.60
1:D:1979:LEU:HD22	1:D:1982:ARG:NH2	2.17	0.60
1:D:3357:HIS:O	1:D:3361:THR:HG23	2.01	0.60
1:A:1520:VAL:HG12	1:A:1527:MET:HG2	1.83	0.60
1:B:2464:ASP:OD1	1:B:2465:ASP:N	2.35	0.60
1:B:2523:ASP:OD1	1:B:2524:VAL:HG13	2.02	0.59
1:B:3786:CYS:SG	1:B:3831:SER:OG	2.60	0.59
1:B:887:ILE:HD11	1:B:961:MET:HE3	1.84	0.59
1:C:3086:GLU:OE2	1:C:3093:ARG:NH1	2.33	0.59
1:A:2523:ASP:OD1	1:A:2524:VAL:HG13	2.02	0.59
1:B:568:LEU:HD12	1:B:602:VAL:HG13	1.84	0.59
1:B:988:LEU:HB2	1:B:1039:LEU:HD21	1.83	0.59
1:B:4749:GLU:OE1	1:B:4753:HIS:CD2	2.56	0.59
1:D:2523:ASP:OD1	1:D:2524:VAL:HG13	2.02	0.59
1:C:568:LEU:HD12	1:C:602:VAL:HG13	1.84	0.59
1:C:1979:LEU:HD22	1:C:1982:ARG:NH2	2.17	0.59
1:B:1454:THR:OG1	1:B:1456:ASP:OD1	2.12	0.59
1:C:2475:GLN:NE2	1:C:2476:ILE:O	2.35	0.59
1:A:238:SER:OG	1:A:240:ASP:OD1	2.20	0.59
1:A:884:LEU:CD2	1:A:955:LEU:HD11	2.31	0.59
1:C:1520:VAL:HG12	1:C:1527:MET:HG2	1.83	0.59
1:C:2464:ASP:OD1	1:C:2465:ASP:N	2.35	0.59
1:A:4217:PHE:O	1:A:4221:VAL:HG22	2.03	0.59
1:D:4749:GLU:OE1	1:D:4753:HIS:CD2	2.56	0.59
1:C:2516:ASP:OD1	1:C:2517:PHE:N	2.33	0.59
1:C:3357:HIS:O	1:C:3361:THR:HG23	2.02	0.59
1:A:553:ARG:NE	1:A:555:GLU:OE2	2.36	0.59
1:A:1979:LEU:HD22	1:A:1982:ARG:NH2	2.17	0.59
1:B:553:ARG:NE	1:B:555:GLU:OE2	2.36	0.59
1:B:797:HIS:ND1	1:B:821:LEU:HD23	2.18	0.59
1:B:3357:HIS:O	1:B:3361:THR:HG23	2.01	0.59
1:D:3308:THR:OG1	1:D:3310:ASP:OD1	2.21	0.59
1:D:4705:VAL:HG22	1:D:4711:PHE:HD1	1.68	0.59
1:A:3308:THR:OG1	1:A:3310:ASP:OD1	2.21	0.59
1:A:4749:GLU:OE1	1:A:4753:HIS:CD2	2.56	0.59
1:B:1520:VAL:HG12	1:B:1527:MET:HG2	1.83	0.59
1:D:553:ARG:NE	1:D:555:GLU:OE2	2.36	0.59
1:D:1024:TYR:OH	1:D:1036:ARG:NH1	2.36	0.59
1:C:797:HIS:ND1	1:C:821:LEU:HD23	2.18	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2534:ALA:HB3	1:B:2588:ARG:HH21	1.68	0.59
1:B:3825:GLU:OE1	1:B:3825:GLU:N	2.36	0.59
1:D:4754:ASN:OD1	1:D:4755:GLU:N	2.36	0.59
1:A:3825:GLU:OE1	1:A:3825:GLU:N	2.36	0.59
1:B:984:LEU:HG	1:B:987:ARG:NH2	2.18	0.59
1:C:2534:ALA:HB3	1:C:2588:ARG:HH21	1.68	0.59
1:C:4217:PHE:O	1:C:4221:VAL:HG22	2.03	0.59
1:A:797:HIS:ND1	1:A:821:LEU:HD23	2.18	0.58
1:A:2534:ALA:HB3	1:A:2588:ARG:HH21	1.68	0.58
1:A:3786:CYS:SG	1:A:3831:SER:OG	2.60	0.58
1:B:224:HIS:HA	1:B:388:LEU:HD23	1.85	0.58
1:B:4754:ASN:OD1	1:B:4755:GLU:N	2.36	0.58
1:C:2515:GLN:NE2	1:C:2568:LEU:O	2.36	0.58
1:A:224:HIS:HA	1:A:388:LEU:HD23	1.85	0.58
1:A:1570:LYS:O	1:A:1572:ILE:HD12	2.03	0.58
1:B:238:SER:OG	1:B:240:ASP:OD1	2.20	0.58
1:B:1552:VAL:HG11	1:B:1562:ILE:HD13	1.84	0.58
1:B:1570:LYS:O	1:B:1572:ILE:HD12	2.03	0.58
1:D:2515:GLN:NE2	1:D:2568:LEU:O	2.36	0.58
1:C:2523:ASP:OD1	1:C:2524:VAL:HG13	2.02	0.58
1:A:2376:LEU:HD21	1:A:2502:MET:HE3	1.84	0.58
1:D:1552:VAL:HG11	1:D:1562:ILE:HD13	1.84	0.58
1:C:984:LEU:HG	1:C:987:ARG:NH2	2.18	0.58
1:C:988:LEU:HB2	1:C:1039:LEU:HD21	1.83	0.58
1:A:167:ASP:HA	1:D:384:MET:HE1	1.85	0.58
1:A:1024:TYR:OH	1:A:1036:ARG:NH1	2.36	0.58
1:D:2534:ALA:HB3	1:D:2588:ARG:HH21	1.68	0.58
1:C:1570:LYS:O	1:C:1572:ILE:HD12	2.03	0.58
1:A:988:LEU:HB2	1:A:1039:LEU:HD21	1.83	0.58
1:A:4754:ASN:OD1	1:A:4755:GLU:N	2.36	0.58
1:D:224:HIS:HA	1:D:388:LEU:HD23	1.86	0.58
1:C:3097:GLU:O	1:C:3100:SER:OG	2.19	0.58
1:C:3825:GLU:N	1:C:3825:GLU:OE1	2.36	0.58
1:C:4754:ASN:OD1	1:C:4755:GLU:N	2.36	0.58
1:A:568:LEU:HD12	1:A:602:VAL:HG13	1.84	0.58
1:A:2464:ASP:OD1	1:A:2465:ASP:N	2.35	0.58
1:B:2466:LEU:HD23	1:B:2502:MET:SD	2.44	0.58
1:B:3097:GLU:O	1:B:3100:SER:OG	2.19	0.58
1:B:4705:VAL:HG22	1:B:4711:PHE:HD1	1.68	0.58
1:D:473:ASN:O	1:D:477:LEU:HD13	2.03	0.58
1:D:984:LEU:HG	1:D:987:ARG:NH2	2.18	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:887:ILE:CD1	1:C:907:LEU:HD11	2.34	0.58
1:C:3052:HIS:CD2	1:C:3127:GLN:HE22	2.22	0.58
1:A:4722:ARG:HH21	1:A:4748:LEU:HD12	1.69	0.58
1:D:167:ASP:HA	1:C:384:MET:HE1	1.86	0.58
1:D:4722:ARG:HH21	1:D:4748:LEU:HD12	1.69	0.58
1:C:4749:GLU:OE1	1:C:4753:HIS:CD2	2.56	0.58
1:A:984:LEU:HG	1:A:987:ARG:NH2	2.18	0.58
1:A:1552:VAL:HG11	1:A:1562:ILE:HD13	1.84	0.58
1:A:2515:GLN:NE2	1:A:2568:LEU:O	2.36	0.58
1:A:3377:GLU:OE1	1:A:3380:ARG:NH2	2.37	0.58
1:A:4070:ASP:OD2	1:A:4071:ILE:N	2.37	0.58
1:A:4705:VAL:HG22	1:A:4711:PHE:HD1	1.68	0.58
1:B:3111:ARG:NH2	1:B:3175:LEU:HA	2.19	0.58
1:B:4217:PHE:O	1:B:4221:VAL:HG22	2.03	0.58
1:D:4217:PHE:O	1:D:4221:VAL:HG22	2.03	0.58
1:C:224:HIS:HA	1:C:388:LEU:HD23	1.86	0.58
1:C:2285:GLU:HG2	1:C:3858:MET:HE1	1.85	0.58
1:A:887:ILE:CD1	1:A:907:LEU:HD11	2.34	0.58
1:B:473:ASN:O	1:B:477:LEU:HD13	2.03	0.58
1:B:4070:ASP:OD2	1:B:4071:ILE:N	2.37	0.58
1:B:4821:LYS:O	1:B:4825:THR:HG23	2.04	0.58
1:D:887:ILE:HD11	1:D:961:MET:HE3	1.85	0.58
1:D:894:GLY:HA3	1:D:903:LEU:HB3	1.86	0.58
1:C:2466:LEU:HD23	1:C:2502:MET:SD	2.44	0.58
1:A:2466:LEU:HD23	1:A:2502:MET:SD	2.44	0.58
1:B:1024:TYR:OH	1:B:1036:ARG:NH1	2.36	0.58
1:B:2515:GLN:NE2	1:B:2568:LEU:O	2.36	0.58
1:B:3052:HIS:CD2	1:B:3127:GLN:HE22	2.22	0.58
1:D:568:LEU:HD12	1:D:602:VAL:HG13	1.84	0.58
1:D:887:ILE:CD1	1:D:907:LEU:HD11	2.34	0.58
1:D:2466:LEU:HD23	1:D:2502:MET:SD	2.44	0.58
1:C:3709:ALA:HB2	1:C:3782:MET:SD	2.44	0.58
1:D:238:SER:OG	1:D:240:ASP:OD1	2.20	0.57
1:D:1821:ASP:OD2	1:D:1837:GLN:NE2	2.37	0.57
1:D:3671:ASP:OD1	1:D:3672:ARG:N	2.38	0.57
1:C:473:ASN:O	1:C:477:LEU:HD13	2.03	0.57
1:A:1996:ARG:HE	1:A:1999:ARG:HH12	1.50	0.57
2:G:20:GLN:NE2	2:G:106:LEU:HD13	2.19	0.57
1:B:887:ILE:CD1	1:B:907:LEU:HD11	2.34	0.57
1:A:3111:ARG:NH2	1:A:3175:LEU:HA	2.19	0.57
1:B:2869:ARG:NH1	1:B:2947:ASP:HB2	2.19	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:797:HIS:ND1	1:D:821:LEU:HD23	2.18	0.57
1:D:3111:ARG:NH2	1:D:3175:LEU:HA	2.19	0.57
1:D:4070:ASP:OD2	1:D:4071:ILE:N	2.37	0.57
1:C:1068:ARG:HA	1:C:1071:ARG:HH11	1.69	0.57
1:A:473:ASN:O	1:A:477:LEU:HD13	2.03	0.57
1:A:894:GLY:HA3	1:A:903:LEU:HB3	1.86	0.57
1:A:2869:ARG:NH1	1:A:2947:ASP:HB2	2.19	0.57
2:H:20:GLN:NE2	2:H:106:LEU:HD13	2.19	0.57
1:B:1821:ASP:OD2	1:B:1837:GLN:NE2	2.37	0.57
1:B:3573:MET:HE2	1:B:3577:ARG:HH12	1.66	0.57
1:D:3343:GLN:O	1:D:3346:VAL:HG12	2.05	0.57
1:D:4821:LYS:O	1:D:4825:THR:HG23	2.04	0.57
1:C:553:ARG:NE	1:C:555:GLU:OE2	2.36	0.57
1:C:1552:VAL:HG11	1:C:1562:ILE:HD13	1.84	0.57
1:B:1068:ARG:HA	1:B:1071:ARG:HH11	1.69	0.57
1:B:3709:ALA:HB2	1:B:3782:MET:SD	2.44	0.57
1:D:1570:LYS:O	1:D:1572:ILE:HD12	2.03	0.57
1:D:3709:ALA:HB2	1:D:3782:MET:SD	2.44	0.57
1:C:4705:VAL:HG22	1:C:4711:PHE:HD1	1.68	0.57
1:A:3343:GLN:O	1:A:3346:VAL:HG12	2.05	0.57
2:E:20:GLN:NE2	2:E:106:LEU:HD13	2.19	0.57
1:C:3343:GLN:O	1:C:3346:VAL:HG12	2.05	0.57
1:C:3377:GLU:OE1	1:C:3380:ARG:NH2	2.37	0.57
1:D:2437:ALA:O	1:D:2508:ARG:NH2	2.38	0.57
1:C:1031:THR:O	1:C:1035:ASN:OD1	2.22	0.57
1:C:4722:ARG:NE	1:C:4748:LEU:HD11	2.20	0.57
1:B:1434:TYR:HB2	1:B:1572:ILE:HG21	1.87	0.57
1:B:3477:LYS:HZ2	1:B:3479:ALA:HB2	1.68	0.57
1:C:971:ASP:OD2	1:C:973:SER:OG	2.14	0.57
1:C:2437:ALA:O	1:C:2508:ARG:NH2	2.38	0.57
1:C:2751:LEU:HD11	1:C:2817:ILE:HD12	1.87	0.57
1:C:3111:ARG:NH2	1:C:3175:LEU:HA	2.19	0.57
1:C:3671:ASP:OD1	1:C:3672:ARG:N	2.37	0.57
1:C:4821:LYS:O	1:C:4825:THR:HG23	2.04	0.57
1:A:1821:ASP:OD2	1:A:1837:GLN:NE2	2.37	0.57
1:A:3573:MET:HE2	1:A:3577:ARG:HH12	1.70	0.57
1:B:3671:ASP:OD1	1:B:3672:ARG:N	2.37	0.57
1:D:160:GLY:O	1:C:3984:ARG:NH2	2.32	0.57
1:D:1996:ARG:HE	1:D:1999:ARG:HH12	1.50	0.57
1:D:3052:HIS:CD2	1:D:3127:GLN:HE22	2.22	0.57
1:D:4722:ARG:NE	1:D:4748:LEU:HD11	2.20	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:35:LEU:HD23	1:C:51:PRO:HA	1.86	0.57
1:A:3671:ASP:OD1	1:A:3672:ARG:N	2.37	0.57
2:E:15:PHE:O	2:E:17:LYS:NZ	2.38	0.57
2:F:15:PHE:O	2:F:17:LYS:NZ	2.38	0.57
1:B:3343:GLN:O	1:B:3346:VAL:HG12	2.05	0.57
1:D:35:LEU:HD23	1:D:51:PRO:HA	1.87	0.57
1:D:1434:TYR:HB2	1:D:1572:ILE:HG21	1.87	0.57
1:C:887:ILE:HD11	1:C:961:MET:HE3	1.85	0.57
1:C:1821:ASP:OD2	1:C:1837:GLN:NE2	2.37	0.57
1:A:1031:THR:O	1:A:1035:ASN:OD1	2.22	0.56
1:A:3097:GLU:O	1:A:3100:SER:OG	2.19	0.56
1:A:3529:ASP:OD2	1:A:3595:ARG:NH2	2.35	0.56
1:A:3709:ALA:HB2	1:A:3782:MET:SD	2.44	0.56
1:A:3941:ASP:OD1	1:A:3942:VAL:N	2.38	0.56
1:B:4722:ARG:HH21	1:B:4748:LEU:HD12	1.69	0.56
1:D:2871:LEU:HD13	1:D:2874:MET:HE1	1.86	0.56
1:D:3103:ILE:HD13	1:D:3168:THR:HG23	1.86	0.56
1:D:3365:LEU:HD11	1:D:3405:LEU:HD12	1.87	0.56
1:C:35:LEU:HD11	1:C:189:LEU:HD13	1.87	0.56
1:C:2869:ARG:NH1	1:C:2947:ASP:HB2	2.19	0.56
1:A:2437:ALA:O	1:A:2508:ARG:NH2	2.38	0.56
1:A:3695:PRO:HB3	1:A:3699:HIS:ND1	2.20	0.56
1:B:35:LEU:HD11	1:B:189:LEU:HD13	1.87	0.56
1:B:3377:GLU:OE1	1:B:3380:ARG:NH2	2.37	0.56
1:C:4722:ARG:HH21	1:C:4748:LEU:HD12	1.69	0.56
1:A:4821:LYS:O	1:A:4825:THR:HG23	2.04	0.56
1:B:1031:THR:O	1:B:1035:ASN:OD1	2.22	0.56
1:B:1999:ARG:O	1:B:3638:MET:HE3	2.05	0.56
1:B:2376:LEU:HD21	1:B:2502:MET:HE3	1.87	0.56
1:B:3401:LEU:HD23	1:B:3451:PHE:HE1	1.70	0.56
1:B:4722:ARG:NE	1:B:4748:LEU:HD11	2.20	0.56
1:D:2751:LEU:HD11	1:D:2817:ILE:HD12	1.87	0.56
1:D:2869:ARG:NH1	1:D:2947:ASP:HB2	2.19	0.56
1:D:3377:GLU:OE1	1:D:3380:ARG:NH2	2.37	0.56
1:D:3401:LEU:HD23	1:D:3451:PHE:HE1	1.70	0.56
1:D:3695:PRO:HB3	1:D:3699:HIS:ND1	2.20	0.56
1:C:1434:TYR:HB2	1:C:1572:ILE:HG21	1.87	0.56
1:C:1996:ARG:HE	1:C:1999:ARG:HH12	1.50	0.56
1:C:2927:LEU:O	1:C:2931:GLN:OE1	2.24	0.56
1:C:4070:ASP:OD2	1:C:4071:ILE:N	2.37	0.56
1:A:2534:ALA:HB1	1:A:2588:ARG:NE	2.19	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3052:HIS:CD2	1:A:3127:GLN:HE22	2.22	0.56
2:H:15:PHE:O	2:H:17:LYS:NZ	2.38	0.56
2:F:20:GLN:NE2	2:F:106:LEU:HD13	2.19	0.56
1:B:35:LEU:HD23	1:B:51:PRO:HA	1.87	0.56
1:B:384:MET:HE1	1:C:167:ASP:HA	1.87	0.56
1:B:877:ASN:ND2	1:B:1045:THR:HG21	2.21	0.56
1:A:1434:TYR:HB2	1:A:1572:ILE:HG21	1.87	0.56
1:D:3941:ASP:OD1	1:D:3942:VAL:N	2.38	0.56
1:C:3695:PRO:HB3	1:C:3699:HIS:ND1	2.20	0.56
1:A:2782:ASP:O	1:A:2786:LYS:N	2.39	0.56
1:A:3103:ILE:HD13	1:A:3168:THR:HG23	1.86	0.56
1:A:3813:GLN:HG2	1:A:3899:PHE:HE2	1.71	0.56
1:A:4722:ARG:NE	1:A:4748:LEU:HD11	2.20	0.56
1:B:321:GLU:OE1	1:B:321:GLU:N	2.39	0.56
1:B:4848:VAL:HG11	1:B:4887:MET:HG2	1.88	0.56
1:D:884:LEU:CD2	1:D:955:LEU:HD11	2.31	0.56
1:D:1031:THR:O	1:D:1035:ASN:OD1	2.22	0.56
1:D:1068:ARG:HA	1:D:1071:ARG:HH11	1.69	0.56
1:D:3248:ARG:NH2	1:D:3252:ASP:OD1	2.37	0.56
1:A:1045:THR:HG22	1:A:1049:TYR:CZ	2.41	0.56
1:B:3813:GLN:HG2	1:B:3899:PHE:HE2	1.71	0.56
1:B:4655:PHE:HA	1:B:4796:MET:CE	2.36	0.56
1:D:4655:PHE:HA	1:D:4796:MET:CE	2.36	0.56
1:A:2977:LEU:HD22	1:A:3056:LEU:HD23	1.88	0.56
1:B:894:GLY:HA3	1:B:903:LEU:HB3	1.86	0.56
1:B:1996:ARG:HE	1:B:1999:ARG:HH12	1.50	0.56
1:B:2927:LEU:O	1:B:2931:GLN:OE1	2.24	0.56
1:B:3365:LEU:HD11	1:B:3405:LEU:HD12	1.87	0.56
1:D:35:LEU:HD11	1:D:189:LEU:HD13	1.87	0.56
1:C:3103:ILE:HD13	1:C:3168:THR:HG23	1.86	0.56
1:C:3813:GLN:HG2	1:C:3899:PHE:HE2	1.71	0.56
1:A:35:LEU:HD23	1:A:51:PRO:HA	1.86	0.56
1:A:1068:ARG:HA	1:A:1071:ARG:HH11	1.69	0.56
1:A:2196:ASN:OD1	1:A:2199:ARG:NH2	2.35	0.56
1:A:3365:LEU:HD11	1:A:3405:LEU:HD12	1.86	0.56
1:B:3016:TYR:O	1:B:3031:ALA:N	2.39	0.56
1:B:3695:PRO:HB3	1:B:3699:HIS:ND1	2.20	0.56
1:D:2782:ASP:O	1:D:2786:LYS:N	2.39	0.56
1:D:2977:LEU:HD22	1:D:3056:LEU:HD23	1.88	0.56
1:C:321:GLU:OE1	1:C:321:GLU:N	2.39	0.56
1:C:3308:THR:OG1	1:C:3310:ASP:OD1	2.21	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3365:LEU:HD11	1:C:3405:LEU:HD12	1.87	0.56
1:A:2751:LEU:HD11	1:A:2817:ILE:HD12	1.87	0.56
1:A:3477:LYS:HZ3	1:B:1141:ARG:HB2	1.71	0.56
1:D:1045:THR:HG22	1:D:1049:TYR:CZ	2.41	0.56
1:C:2782:ASP:O	1:C:2786:LYS:N	2.39	0.56
1:C:4747:SER:O	1:C:4751:THR:HG23	2.06	0.56
1:A:877:ASN:ND2	1:A:1045:THR:HG21	2.21	0.55
1:A:4848:VAL:HG11	1:A:4887:MET:HG2	1.88	0.55
1:B:2775:TRP:CZ3	1:B:2785:LEU:HD11	2.42	0.55
1:D:4747:SER:O	1:D:4751:THR:HG23	2.06	0.55
1:C:4848:VAL:HG11	1:C:4887:MET:HG2	1.88	0.55
1:A:1930:LYS:O	1:A:1930:LYS:HD3	2.07	0.55
1:A:2927:LEU:O	1:A:2931:GLN:OE1	2.24	0.55
1:A:4655:PHE:HA	1:A:4796:MET:CE	2.36	0.55
1:B:3529:ASP:OD2	1:B:3595:ARG:NH2	2.35	0.55
1:D:877:ASN:ND2	1:D:1045:THR:HG21	2.21	0.55
1:A:3248:ARG:NH2	1:A:3252:ASP:OD1	2.37	0.55
1:B:1930:LYS:O	1:B:1930:LYS:HD3	2.06	0.55
1:B:2239:PHE:HA	1:B:2242:ILE:HD12	1.89	0.55
1:B:3514:LEU:HD21	1:B:3602:VAL:HG11	1.88	0.55
1:D:2872:GLN:O	1:D:2876:GLU:OE1	2.25	0.55
1:D:4848:VAL:HG11	1:D:4887:MET:HG2	1.88	0.55
1:C:1024:TYR:OH	1:C:1036:ARG:NH1	2.36	0.55
1:C:2872:GLN:O	1:C:2876:GLU:OE1	2.25	0.55
1:C:3016:TYR:O	1:C:3031:ALA:N	2.39	0.55
1:A:35:LEU:HD11	1:A:189:LEU:HD13	1.87	0.55
1:A:384:MET:HE1	1:B:166:GLY:O	2.06	0.55
1:A:3401:LEU:HD23	1:A:3451:PHE:HE1	1.70	0.55
1:B:1045:THR:HG22	1:B:1049:TYR:CZ	2.41	0.55
1:B:4747:SER:O	1:B:4751:THR:HG23	2.06	0.55
1:D:3813:GLN:HG2	1:D:3899:PHE:HE2	1.71	0.55
1:A:2239:PHE:HA	1:A:2242:ILE:HD12	1.89	0.55
1:A:3940:LYS:O	1:A:4002:LYS:NZ	2.29	0.55
1:B:2782:ASP:O	1:B:2786:LYS:N	2.39	0.55
1:B:3103:ILE:HD13	1:B:3168:THR:HG23	1.86	0.55
1:B:3324:VAL:HG11	1:B:3361:THR:HG22	1.88	0.55
1:D:981:GLN:O	1:D:985:VAL:HG23	2.07	0.55
1:C:894:GLY:HA3	1:C:903:LEU:HB3	1.86	0.55
1:C:3324:VAL:HG11	1:C:3361:THR:HG22	1.88	0.55
1:C:3477:LYS:HZ2	1:C:3479:ALA:HB2	1.71	0.55
1:A:2135:LEU:HD12	1:A:3658:LYS:NZ	2.22	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:15:PHE:O	2:G:17:LYS:NZ	2.38	0.55
1:D:2927:LEU:O	1:D:2931:GLN:OE1	2.24	0.55
1:C:877:ASN:ND2	1:C:1045:THR:HG21	2.21	0.55
1:C:2135:LEU:HD12	1:C:3658:LYS:NZ	2.21	0.55
1:A:1999:ARG:O	1:A:3638:MET:HE3	2.06	0.55
1:B:2212:VAL:HG11	1:B:2256:TYR:CE2	2.42	0.55
1:B:2437:ALA:O	1:B:2508:ARG:NH2	2.38	0.55
1:C:2977:LEU:HD22	1:C:3056:LEU:HD23	1.88	0.55
1:C:3401:LEU:HD23	1:C:3451:PHE:HE1	1.70	0.55
1:C:3941:ASP:OD1	1:C:3942:VAL:N	2.38	0.55
1:A:2212:VAL:HG11	1:A:2256:TYR:CE2	2.42	0.55
1:B:1005:TRP:HB3	1:B:1021:LEU:HD22	1.89	0.55
1:D:3107:VAL:O	1:D:3111:ARG:HG3	2.07	0.55
1:C:2212:VAL:HG11	1:C:2256:TYR:CE2	2.42	0.55
1:C:3573:MET:HE2	1:C:3577:ARG:HH12	1.70	0.55
1:C:4655:PHE:HA	1:C:4796:MET:CE	2.36	0.55
1:C:4705:VAL:HG13	1:C:4711:PHE:CE1	2.42	0.55
1:A:321:GLU:OE1	1:A:321:GLU:N	2.39	0.55
1:A:1005:TRP:HB3	1:A:1021:LEU:HD22	1.89	0.55
1:A:4098:ASP:OD1	1:A:4099:SER:N	2.40	0.55
1:B:2977:LEU:HD22	1:B:3056:LEU:HD23	1.88	0.55
1:D:1999:ARG:O	1:D:3638:MET:HE1	2.06	0.55
1:D:2196:ASN:OD1	1:D:2199:ARG:NH2	2.35	0.55
1:D:3825:GLU:OE1	1:D:3825:GLU:N	2.36	0.55
1:C:981:GLN:O	1:C:985:VAL:HG23	2.07	0.55
1:A:2858:GLN:O	1:A:2860:PRO:HD3	2.07	0.55
1:A:3324:VAL:HG11	1:A:3361:THR:HG22	1.88	0.55
1:B:2135:LEU:HD12	1:B:3658:LYS:NZ	2.21	0.55
1:B:2626:LEU:HD22	1:B:2640:PRO:HB3	1.89	0.55
1:B:2872:GLN:O	1:B:2876:GLU:OE1	2.25	0.55
1:B:3084:GLY:O	1:B:3089:LYS:NZ	2.37	0.55
1:B:3941:ASP:OD1	1:B:3942:VAL:N	2.38	0.55
1:D:321:GLU:OE1	1:D:321:GLU:N	2.39	0.55
1:D:1930:LYS:O	1:D:1930:LYS:HD3	2.06	0.55
1:D:2135:LEU:HD12	1:D:3658:LYS:NZ	2.22	0.55
1:A:3061:ALA:O	1:A:3065:VAL:HG23	2.07	0.54
1:A:3084:GLY:O	1:A:3089:LYS:NZ	2.37	0.54
1:C:2858:GLN:O	1:C:2860:PRO:HD3	2.07	0.54
1:C:2948:THR:HB	1:C:2952:GLU:HB2	1.90	0.54
1:C:3061:ALA:O	1:C:3065:VAL:HG23	2.07	0.54
1:C:3107:VAL:O	1:C:3111:ARG:HG3	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4176:PRO:O	1:C:4202:ARG:NH2	2.40	0.54
1:A:897:ARG:NH2	1:A:906:CYS:SG	2.73	0.54
1:A:2660:GLY:HA3	1:A:2666:VAL:HG12	1.89	0.54
1:A:3107:VAL:O	1:A:3111:ARG:HG3	2.07	0.54
1:A:4176:PRO:O	1:A:4202:ARG:NH2	2.40	0.54
1:B:3061:ALA:O	1:B:3065:VAL:HG23	2.07	0.54
1:B:4098:ASP:OD1	1:B:4099:SER:N	2.40	0.54
1:C:4098:ASP:OD1	1:C:4099:SER:N	2.40	0.54
1:A:2872:GLN:O	1:A:2876:GLU:OE1	2.25	0.54
1:A:3514:LEU:HD21	1:A:3602:VAL:HG11	1.88	0.54
1:A:4747:SER:O	1:A:4751:THR:HG23	2.06	0.54
1:B:2026:ASP:OD1	1:B:2027:ILE:N	2.40	0.54
1:D:3061:ALA:O	1:D:3065:VAL:HG23	2.07	0.54
1:C:2239:PHE:HA	1:C:2242:ILE:HD12	1.89	0.54
1:A:2775:TRP:CZ3	1:A:2785:LEU:HD11	2.42	0.54
1:A:3891:LEU:O	1:A:3899:PHE:HD2	1.90	0.54
1:B:2948:THR:HB	1:B:2952:GLU:HB2	1.90	0.54
1:B:3308:THR:OG1	1:B:3310:ASP:OD1	2.21	0.54
1:B:4176:PRO:O	1:B:4202:ARG:NH2	2.40	0.54
1:D:1005:TRP:HB3	1:D:1021:LEU:HD22	1.89	0.54
1:D:2858:GLN:O	1:D:2860:PRO:HD3	2.07	0.54
1:C:1045:THR:HG22	1:C:1049:TYR:CZ	2.41	0.54
1:C:2775:TRP:CZ3	1:C:2785:LEU:HD11	2.42	0.54
1:C:3442:PHE:CE1	1:C:3511:VAL:HG12	2.42	0.54
1:A:2026:ASP:OD1	1:A:2027:ILE:N	2.40	0.54
1:A:2751:LEU:CD2	1:A:2755:ILE:HD11	2.38	0.54
1:B:2751:LEU:HD11	1:B:2817:ILE:HD12	1.87	0.54
1:D:2775:TRP:CZ3	1:D:2785:LEU:HD11	2.42	0.54
1:D:3442:PHE:CE1	1:D:3511:VAL:HG12	2.42	0.54
1:D:4098:ASP:OD1	1:D:4099:SER:N	2.40	0.54
1:D:4176:PRO:O	1:D:4202:ARG:NH2	2.40	0.54
1:C:1930:LYS:O	1:C:1930:LYS:HD3	2.07	0.54
1:C:2626:LEU:HD22	1:C:2640:PRO:HB3	1.89	0.54
1:C:2869:ARG:CZ	1:C:2947:ASP:HB2	2.38	0.54
1:A:2948:THR:HB	1:A:2952:GLU:HB2	1.90	0.54
1:B:213:TYR:CE1	1:B:340:LYS:HD3	2.43	0.54
1:B:2751:LEU:CD2	1:B:2755:ILE:HD11	2.38	0.54
1:D:2026:ASP:OD1	1:D:2027:ILE:N	2.40	0.54
1:D:2239:PHE:HA	1:D:2242:ILE:HD12	1.89	0.54
1:D:2626:LEU:HD22	1:D:2640:PRO:HB3	1.89	0.54
1:D:2751:LEU:CD2	1:D:2755:ILE:HD11	2.38	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3324:VAL:HG11	1:D:3361:THR:HG22	1.88	0.54
1:C:897:ARG:NH2	1:C:906:CYS:SG	2.73	0.54
1:A:981:GLN:O	1:A:985:VAL:HG23	2.07	0.54
1:A:4705:VAL:HG13	1:A:4711:PHE:CE1	2.42	0.54
1:B:3107:VAL:O	1:B:3111:ARG:HG3	2.07	0.54
1:B:3442:PHE:CE1	1:B:3511:VAL:HG12	2.42	0.54
1:D:1448:VAL:HG22	1:D:1554:VAL:HG23	1.90	0.54
1:D:2212:VAL:HG11	1:D:2256:TYR:CE2	2.42	0.54
1:A:728:ARG:NH2	1:A:1489:CYS:SG	2.81	0.54
1:A:1427:ILE:HG23	1:A:1428:LEU:HD22	1.90	0.54
1:A:2233:CYS:O	1:A:2236:LEU:N	2.41	0.54
1:B:3169:LEU:HD12	1:B:3194:LEU:HD11	1.90	0.54
1:B:3689:GLU:OE2	1:B:3694:LYS:HB2	2.08	0.54
1:D:213:TYR:CE1	1:D:340:LYS:HD3	2.43	0.54
1:D:728:ARG:NH2	1:D:1489:CYS:SG	2.81	0.54
1:D:2948:THR:HB	1:D:2952:GLU:HB2	1.90	0.54
1:D:3529:ASP:OD2	1:D:3595:ARG:NH2	2.35	0.54
1:C:3514:LEU:HD21	1:C:3602:VAL:HG11	1.88	0.54
1:A:3016:TYR:O	1:A:3031:ALA:N	2.39	0.54
1:B:281:ARG:NH2	1:B:309:THR:OG1	2.41	0.54
1:B:908:VAL:HG23	1:B:963:ASN:ND2	2.23	0.54
1:B:3891:LEU:O	1:B:3899:PHE:HD2	1.90	0.54
1:B:4705:VAL:HG13	1:B:4711:PHE:CE1	2.42	0.54
1:D:3514:LEU:HD21	1:D:3602:VAL:HG11	1.88	0.54
1:C:213:TYR:CE1	1:C:340:LYS:HD3	2.43	0.54
1:C:2751:LEU:CD2	1:C:2755:ILE:HD11	2.38	0.54
1:A:2771:ILE:HG22	1:B:1506:GLN:OE1	2.08	0.54
1:B:648:ILE:HG23	1:B:814:ALA:HB3	1.90	0.54
1:B:2233:CYS:O	1:B:2236:LEU:N	2.41	0.54
1:B:2660:GLY:HA3	1:B:2666:VAL:HG12	1.89	0.54
1:D:281:ARG:NH2	1:D:309:THR:OG1	2.41	0.54
1:D:4241:THR:HB	1:D:4989:MET:HE3	1.90	0.54
1:C:3330:ASP:OD1	1:C:3331:GLU:N	2.41	0.54
1:A:2869:ARG:CZ	1:A:2947:ASP:HB2	2.38	0.53
1:A:3169:LEU:HD12	1:A:3194:LEU:HD11	1.90	0.53
1:A:3689:GLU:OE2	1:A:3694:LYS:HB2	2.08	0.53
1:B:2858:GLN:O	1:B:2860:PRO:HD3	2.07	0.53
1:B:2871:LEU:O	1:B:2874:MET:HG2	2.08	0.53
1:D:3330:ASP:OD1	1:D:3331:GLU:N	2.41	0.53
1:A:3365:LEU:HD21	1:A:3405:LEU:HD12	1.91	0.53
1:A:3442:PHE:CE1	1:A:3511:VAL:HG12	2.42	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:728:ARG:NH2	1:B:1489:CYS:SG	2.81	0.53
1:B:981:GLN:O	1:B:985:VAL:HG23	2.07	0.53
1:D:1427:ILE:HG23	1:D:1428:LEU:HD22	1.90	0.53
1:D:3169:LEU:HD12	1:D:3194:LEU:HD11	1.90	0.53
1:D:3365:LEU:HD21	1:D:3405:LEU:HD12	1.91	0.53
1:C:908:VAL:HG13	1:C:912:SER:HB2	1.91	0.53
1:C:1999:ARG:O	1:C:3638:MET:HE1	2.07	0.53
1:C:2233:CYS:O	1:C:2236:LEU:N	2.41	0.53
1:C:3248:ARG:NH2	1:C:3252:ASP:OD1	2.37	0.53
1:D:908:VAL:HG13	1:D:912:SER:HB2	1.91	0.53
1:D:2871:LEU:O	1:D:2874:MET:HG2	2.08	0.53
1:D:3689:GLU:OE2	1:D:3694:LYS:HB2	2.08	0.53
1:D:3891:LEU:O	1:D:3899:PHE:HD2	1.90	0.53
1:C:1005:TRP:HB3	1:C:1021:LEU:HD22	1.89	0.53
1:C:4000:MET:HE1	1:C:4061:PHE:CB	2.38	0.53
1:A:545:ASP:OD1	1:A:582:HIS:NE2	2.40	0.53
1:B:2534:ALA:HB1	1:B:2588:ARG:NE	2.19	0.53
1:D:2463:LEU:HA	1:D:2466:LEU:HD12	1.91	0.53
1:D:2869:ARG:CZ	1:D:2947:ASP:HB2	2.38	0.53
1:D:4705:VAL:HG13	1:D:4711:PHE:CE1	2.42	0.53
1:C:1448:VAL:HG22	1:C:1554:VAL:HG23	1.90	0.53
1:C:2948:THR:HB	1:C:2952:GLU:CB	2.39	0.53
1:C:3689:GLU:OE2	1:C:3694:LYS:HB2	2.08	0.53
1:A:384:MET:HE1	1:B:167:ASP:HA	1.89	0.53
1:A:908:VAL:HG23	1:A:963:ASN:ND2	2.23	0.53
1:A:2626:LEU:HD22	1:A:2640:PRO:HB3	1.89	0.53
1:B:2869:ARG:CZ	1:B:2947:ASP:HB2	2.38	0.53
1:D:1141:ARG:HB2	1:C:3477:LYS:HZ3	1.73	0.53
1:C:2026:ASP:OD1	1:C:2027:ILE:N	2.40	0.53
1:C:3891:LEU:O	1:C:3899:PHE:HD2	1.90	0.53
1:A:3330:ASP:OD1	1:A:3331:GLU:N	2.41	0.53
1:D:69:LEU:HD13	1:D:101:LEU:HD11	1.91	0.53
1:D:908:VAL:HG23	1:D:963:ASN:ND2	2.23	0.53
1:D:4655:PHE:HA	1:D:4796:MET:HE1	1.91	0.53
1:C:69:LEU:HD13	1:C:101:LEU:HD11	1.91	0.53
1:C:648:ILE:HG23	1:C:814:ALA:HB3	1.91	0.53
1:C:728:ARG:NH2	1:C:1489:CYS:SG	2.81	0.53
1:C:2265:LEU:HD13	1:C:2330:ARG:NH2	2.24	0.53
1:A:281:ARG:NH2	1:A:309:THR:OG1	2.41	0.53
1:A:648:ILE:HG23	1:A:814:ALA:HB3	1.91	0.53
1:A:1978:ALA:C	1:A:1982:ARG:CZ	2.77	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1427:ILE:HG23	1:B:1428:LEU:HD22	1.90	0.53
1:B:1554:VAL:HG21	1:B:1561:VAL:CG1	2.39	0.53
1:D:2658:PRO:O	1:D:2666:VAL:HG11	2.09	0.53
1:D:2948:THR:HB	1:D:2952:GLU:CB	2.39	0.53
1:C:2658:PRO:O	1:C:2666:VAL:HG11	2.09	0.53
1:A:213:TYR:CE1	1:A:340:LYS:HD3	2.43	0.53
1:B:887:ILE:HG23	1:B:959:TYR:OH	2.09	0.53
1:B:3330:ASP:OD1	1:B:3331:GLU:N	2.41	0.53
1:D:2233:CYS:O	1:D:2236:LEU:N	2.41	0.53
1:D:2265:LEU:HD13	1:D:2330:ARG:NH2	2.24	0.53
1:D:2660:GLY:HA3	1:D:2666:VAL:HG12	1.89	0.53
1:C:281:ARG:NH2	1:C:309:THR:OG1	2.41	0.53
1:C:3365:LEU:HD21	1:C:3405:LEU:HD12	1.91	0.53
1:C:4241:THR:HB	1:C:4989:MET:HE3	1.90	0.53
1:A:2001:PRO:HG3	1:A:3638:MET:SD	2.49	0.53
1:A:2254:LEU:HD11	1:A:2276:ALA:HB1	1.91	0.53
1:A:2463:LEU:HA	1:A:2466:LEU:HD12	1.91	0.53
1:A:2871:LEU:O	1:A:2874:MET:HG2	2.08	0.53
1:B:2001:PRO:HG3	1:B:3638:MET:SD	2.49	0.53
1:B:2254:LEU:HD11	1:B:2276:ALA:HB1	1.91	0.53
1:B:2948:THR:HB	1:B:2952:GLU:CB	2.39	0.53
1:B:3365:LEU:HD21	1:B:3405:LEU:HD12	1.90	0.53
1:D:648:ILE:HG23	1:D:814:ALA:HB3	1.91	0.53
1:C:1926:LEU:HD23	1:C:1929:MET:SD	2.49	0.53
1:C:3762:ARG:NH1	1:C:4755:GLU:OE1	2.42	0.53
1:A:1448:VAL:HG22	1:A:1554:VAL:HG23	1.90	0.53
1:A:2265:LEU:HD13	1:A:2330:ARG:NH2	2.24	0.53
1:A:3177:THR:O	1:A:3179:LYS:NZ	2.36	0.53
1:A:4241:THR:HB	1:A:4989:MET:HE3	1.89	0.53
1:D:3166:TYR:O	1:D:3170:CYS:SG	2.67	0.53
1:C:887:ILE:HG23	1:C:959:TYR:OH	2.09	0.53
1:C:2001:PRO:HG3	1:C:3638:MET:SD	2.49	0.53
1:C:2660:GLY:HA3	1:C:2666:VAL:HG12	1.89	0.53
1:A:69:LEU:HD13	1:A:101:LEU:HD11	1.91	0.52
1:A:887:ILE:HG23	1:A:959:TYR:OH	2.09	0.52
1:A:2948:THR:HB	1:A:2952:GLU:CB	2.39	0.52
1:B:1926:LEU:HD23	1:B:1929:MET:SD	2.49	0.52
1:B:2265:LEU:HD13	1:B:2330:ARG:NH2	2.24	0.52
1:B:2658:PRO:O	1:B:2666:VAL:HG11	2.09	0.52
1:B:4241:THR:HB	1:B:4989:MET:HE3	1.90	0.52
1:D:672:VAL:O	1:D:680:THR:OG1	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:887:ILE:HG23	1:D:959:TYR:OH	2.09	0.52
1:C:233:ILE:HD12	1:C:242:ARG:HB3	1.92	0.52
1:A:3762:ARG:NH1	1:A:4755:GLU:OE1	2.42	0.52
1:A:4632:LEU:HB3	1:A:4634:GLU:OE1	2.10	0.52
1:B:69:LEU:HD13	1:B:101:LEU:HD11	1.91	0.52
1:B:1978:ALA:C	1:B:1982:ARG:CZ	2.77	0.52
1:B:2196:ASN:OD1	1:B:2199:ARG:NH2	2.35	0.52
1:B:3477:LYS:HZ3	1:C:1141:ARG:HB2	1.75	0.52
1:D:233:ILE:HD12	1:D:242:ARG:HB3	1.92	0.52
1:D:2001:PRO:HG3	1:D:3638:MET:SD	2.49	0.52
1:D:2138:LEU:HB2	1:D:3658:LYS:HZ1	1.74	0.52
1:D:3762:ARG:NH1	1:D:4755:GLU:OE1	2.42	0.52
1:C:908:VAL:HG23	1:C:963:ASN:ND2	2.23	0.52
1:C:2534:ALA:HB1	1:C:2588:ARG:NE	2.19	0.52
1:A:3166:TYR:O	1:A:3170:CYS:SG	2.67	0.52
1:A:4928:LEU:HA	1:A:4931:ILE:HD12	1.91	0.52
1:D:1978:ALA:C	1:D:1982:ARG:CZ	2.78	0.52
1:C:1427:ILE:HG23	1:C:1428:LEU:HD22	1.90	0.52
1:C:2668:SER:O	1:C:2669:GLU:CB	2.57	0.52
1:C:3169:LEU:HD12	1:C:3194:LEU:HD11	1.90	0.52
1:A:233:ILE:HD12	1:A:242:ARG:HB3	1.92	0.52
1:A:2871:LEU:HD13	1:A:2874:MET:CE	2.39	0.52
1:B:2908:TYR:CZ	1:B:2916:LYS:HG3	2.45	0.52
1:B:3104:GLU:O	1:B:3107:VAL:HG22	2.10	0.52
1:B:3166:TYR:O	1:B:3170:CYS:SG	2.67	0.52
1:B:3762:ARG:NH1	1:B:4755:GLU:OE1	2.42	0.52
1:D:2668:SER:O	1:D:2669:GLU:CB	2.57	0.52
1:D:2747:ILE:HD12	1:D:2814:LYS:HD3	1.91	0.52
1:D:3104:GLU:O	1:D:3107:VAL:HG22	2.10	0.52
1:C:2577:ILE:H	1:C:2577:ILE:HD12	1.75	0.52
1:C:4928:LEU:HA	1:C:4931:ILE:HD12	1.91	0.52
1:A:908:VAL:HG13	1:A:912:SER:HB2	1.91	0.52
1:A:1554:VAL:HG21	1:A:1561:VAL:CG1	2.39	0.52
1:A:3477:LYS:HZ2	1:A:3479:ALA:HB2	1.75	0.52
1:B:2871:LEU:HD13	1:B:2874:MET:CE	2.39	0.52
1:D:545:ASP:OD1	1:D:582:HIS:NE2	2.40	0.52
1:D:1554:VAL:HG21	1:D:1561:VAL:CG1	2.39	0.52
1:D:1926:LEU:HD23	1:D:1929:MET:SD	2.50	0.52
1:D:2254:LEU:HD11	1:D:2276:ALA:HB1	1.91	0.52
1:D:3573:MET:HE2	1:D:3577:ARG:HH12	1.75	0.52
1:C:545:ASP:OD1	1:C:582:HIS:NE2	2.40	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2463:LEU:HA	1:C:2466:LEU:HD12	1.91	0.52
1:C:3052:HIS:HA	1:C:3127:GLN:NE2	2.25	0.52
1:C:4720:VAL:O	1:C:4724:VAL:HG23	2.10	0.52
1:B:908:VAL:HG13	1:B:912:SER:HB2	1.91	0.52
1:B:1448:VAL:HG22	1:B:1554:VAL:HG23	1.90	0.52
1:B:3052:HIS:HA	1:B:3127:GLN:NE2	2.25	0.52
1:D:575:LEU:HD22	1:D:606:LEU:HA	1.92	0.52
1:C:2138:LEU:HB2	1:C:3658:LYS:HZ1	1.75	0.52
1:C:2254:LEU:HD11	1:C:2276:ALA:HB1	1.91	0.52
1:C:2871:LEU:O	1:C:2874:MET:HG2	2.08	0.52
1:C:2894:LEU:HD12	1:C:2897:LYS:HE2	1.92	0.52
1:A:1926:LEU:HD23	1:A:1929:MET:SD	2.49	0.52
1:B:233:ILE:HD12	1:B:242:ARG:HB3	1.92	0.52
1:B:3395:ARG:HD3	1:B:3453:ARG:NH1	2.25	0.52
1:C:672:VAL:O	1:C:680:THR:OG1	2.08	0.52
1:C:3166:TYR:O	1:C:3170:CYS:SG	2.67	0.52
1:A:575:LEU:HD22	1:A:606:LEU:HA	1.92	0.52
1:A:1039:LEU:O	1:A:1043:VAL:HG23	2.09	0.52
1:A:2658:PRO:O	1:A:2666:VAL:HG11	2.09	0.52
1:A:3052:HIS:HA	1:A:3127:GLN:NE2	2.25	0.52
2:F:6:THR:HG23	2:F:6:THR:O	2.10	0.52
1:B:4928:LEU:HA	1:B:4931:ILE:HD12	1.91	0.52
1:D:2744:ASN:OD1	1:D:2745:VAL:N	2.43	0.52
1:D:3690:VAL:O	1:D:3693:LYS:HE3	2.10	0.52
1:D:4998:LYS:HB3	1:D:5003:HIS:CE1	2.45	0.52
1:C:575:LEU:HD22	1:C:606:LEU:HA	1.92	0.52
1:C:1039:LEU:O	1:C:1043:VAL:HG23	2.09	0.52
1:C:2744:ASN:OD1	1:C:2745:VAL:N	2.43	0.52
1:A:358:THR:OG1	1:A:383:HIS:ND1	2.42	0.52
1:A:2577:ILE:H	1:A:2577:ILE:HD12	1.75	0.52
1:B:897:ARG:NH2	1:B:906:CYS:SG	2.73	0.52
1:B:1828:ASP:N	1:B:1828:ASP:OD1	2.43	0.52
1:B:2285:GLU:HG2	1:B:3858:MET:HE1	1.92	0.52
1:B:2577:ILE:H	1:B:2577:ILE:HD12	1.75	0.52
1:B:3716:LEU:HD21	1:B:3782:MET:HE3	1.92	0.52
1:B:5000:GLU:HA	1:B:5003:HIS:CD2	2.45	0.52
1:D:3395:ARG:HD3	1:D:3453:ARG:NH1	2.25	0.52
1:D:3997:ALA:HB1	1:D:4057:MET:SD	2.50	0.52
1:C:1978:ALA:C	1:C:1982:ARG:CZ	2.77	0.52
1:C:2871:LEU:HD13	1:C:2874:MET:CE	2.39	0.52
1:A:2775:TRP:CE3	1:A:2785:LEU:HD21	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1978:ALA:CB	1:B:1982:ARG:HH12	2.18	0.52
1:B:3997:ALA:HB1	1:B:4057:MET:SD	2.50	0.52
1:B:4632:LEU:HB3	1:B:4634:GLU:OE1	2.10	0.52
1:D:3230:LEU:HD23	1:D:3230:LEU:H	1.75	0.52
1:D:4720:VAL:O	1:D:4724:VAL:HG23	2.10	0.52
1:C:1424:PRO:O	1:C:1428:LEU:HD23	2.10	0.52
1:C:2196:ASN:OD1	1:C:2199:ARG:NH2	2.35	0.52
1:C:2908:TYR:CZ	1:C:2916:LYS:HG3	2.45	0.52
1:C:3690:VAL:O	1:C:3693:LYS:HE3	2.10	0.52
1:A:98:HIS:C	1:A:99:ARG:HD2	2.31	0.51
1:A:2668:SER:O	1:A:2669:GLU:CB	2.57	0.51
1:A:4778:TRP:O	1:A:4782:VAL:HG23	2.10	0.51
1:B:575:LEU:HD22	1:B:606:LEU:HA	1.92	0.51
1:B:1039:LEU:O	1:B:1043:VAL:HG23	2.09	0.51
1:B:2088:GLU:N	1:B:2088:GLU:OE1	2.43	0.51
1:B:2463:LEU:HA	1:B:2466:LEU:HD12	1.91	0.51
1:D:1039:LEU:O	1:D:1043:VAL:HG23	2.09	0.51
1:D:1749:PRO:HD2	1:D:1758:ARG:HH12	1.75	0.51
1:D:2871:LEU:HD13	1:D:2874:MET:CE	2.39	0.51
1:D:3016:TYR:O	1:D:3031:ALA:N	2.39	0.51
1:D:4158:PRO:HA	1:D:4161:ARG:NE	2.26	0.51
1:A:2744:ASN:OD1	1:A:2745:VAL:N	2.43	0.51
1:A:2894:LEU:HD12	1:A:2897:LYS:HE2	1.92	0.51
1:A:3104:GLU:O	1:A:3107:VAL:HG22	2.10	0.51
1:A:4576:ILE:HG23	1:A:4639:MET:SD	2.51	0.51
1:B:2744:ASN:OD1	1:B:2745:VAL:N	2.43	0.51
1:B:2747:ILE:HD12	1:B:2814:LYS:HD3	1.91	0.51
1:B:4720:VAL:O	1:B:4724:VAL:HG23	2.10	0.51
1:D:98:HIS:C	1:D:99:ARG:HD2	2.31	0.51
1:C:1554:VAL:HG21	1:C:1561:VAL:CG1	2.39	0.51
1:C:2775:TRP:CE3	1:C:2785:LEU:HD21	2.45	0.51
1:C:3104:GLU:O	1:C:3107:VAL:HG22	2.10	0.51
1:C:3395:ARG:HD3	1:C:3453:ARG:NH1	2.25	0.51
1:C:5000:GLU:HA	1:C:5003:HIS:CD2	2.45	0.51
1:A:2000:SER:O	1:A:2005:GLN:NE2	2.41	0.51
1:A:3690:VAL:O	1:A:3693:LYS:HE3	2.10	0.51
1:A:3997:ALA:HB1	1:A:4057:MET:SD	2.50	0.51
1:B:3984:ARG:NH2	1:C:160:GLY:O	2.32	0.51
1:D:1424:PRO:O	1:D:1428:LEU:HD23	2.10	0.51
1:D:2534:ALA:HB1	1:D:2588:ARG:NE	2.19	0.51
1:D:2908:TYR:CZ	1:D:2916:LYS:HG3	2.45	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3084:GLY:O	1:D:3089:LYS:NZ	2.37	0.51
1:D:4576:ILE:HG23	1:D:4639:MET:SD	2.51	0.51
1:C:399:GLN:O	1:C:403:MET:HG3	2.11	0.51
1:C:2021:CYS:O	1:C:2028:ARG:NH2	2.44	0.51
1:C:3997:ALA:HB1	1:C:4057:MET:SD	2.50	0.51
1:C:4584:ASP:OD2	1:C:4585:SER:N	2.44	0.51
1:A:1749:PRO:HD2	1:A:1758:ARG:HH12	1.75	0.51
1:A:2021:CYS:O	1:A:2028:ARG:NH2	2.44	0.51
1:A:2138:LEU:HB2	1:A:3658:LYS:HZ1	1.75	0.51
1:A:5000:GLU:HA	1:A:5003:HIS:CD2	2.45	0.51
1:B:1424:PRO:O	1:B:1428:LEU:HD23	2.10	0.51
1:B:2894:LEU:HD12	1:B:2897:LYS:HE2	1.92	0.51
1:D:2088:GLU:OE1	1:D:2088:GLU:N	2.43	0.51
1:D:2775:TRP:CE3	1:D:2785:LEU:HD21	2.45	0.51
1:C:3940:LYS:O	1:C:4002:LYS:NZ	2.29	0.51
1:C:4632:LEU:HB3	1:C:4634:GLU:OE1	2.10	0.51
1:A:1424:PRO:O	1:A:1428:LEU:HD23	2.10	0.51
1:A:2908:TYR:CZ	1:A:2916:LYS:HG3	2.45	0.51
1:A:4158:PRO:HA	1:A:4161:ARG:NE	2.25	0.51
1:B:774:ASP:OD2	1:B:1470:ARG:NH2	2.43	0.51
1:B:2138:LEU:HB2	1:B:3658:LYS:HZ1	1.75	0.51
1:B:2327:GLY:CA	1:B:2330:ARG:NH2	2.74	0.51
1:B:3346:VAL:HG11	1:B:3414:ARG:CB	2.41	0.51
1:B:4778:TRP:O	1:B:4782:VAL:HG23	2.10	0.51
1:D:2285:GLU:HG2	1:D:3858:MET:HE1	1.92	0.51
1:D:2894:LEU:HD12	1:D:2897:LYS:HE2	1.92	0.51
1:D:3951:PHE:CE1	1:D:3999:MET:HE1	2.46	0.51
1:C:98:HIS:C	1:C:99:ARG:HD2	2.31	0.51
1:C:4778:TRP:O	1:C:4782:VAL:HG23	2.10	0.51
1:A:2088:GLU:OE1	1:A:2088:GLU:N	2.43	0.51
1:B:399:GLN:O	1:B:403:MET:HG3	2.11	0.51
1:B:2775:TRP:CE3	1:B:2785:LEU:HD21	2.45	0.51
1:B:4158:PRO:HA	1:B:4161:ARG:NE	2.25	0.51
1:B:4576:ILE:HG23	1:B:4639:MET:SD	2.51	0.51
1:D:2021:CYS:O	1:D:2028:ARG:NH2	2.44	0.51
1:D:2623:LEU:O	1:D:2627:VAL:HG23	2.11	0.51
1:D:3104:GLU:HA	1:D:3107:VAL:HG22	1.93	0.51
1:D:4632:LEU:HB3	1:D:4634:GLU:OE1	2.10	0.51
1:D:4778:TRP:O	1:D:4782:VAL:HG23	2.10	0.51
1:C:1978:ALA:CB	1:C:1982:ARG:HH12	2.18	0.51
1:C:2088:GLU:OE1	1:C:2088:GLU:N	2.43	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:2623:LEU:O	1:C:2627:VAL:HG23	2.11	0.51
1:C:2747:ILE:HD12	1:C:2814:LYS:HD3	1.91	0.51
1:A:2327:GLY:CA	1:A:2330:ARG:NH2	2.74	0.51
1:A:2747:ILE:HD12	1:A:2814:LYS:HD3	1.91	0.51
1:A:3104:GLU:HA	1:A:3107:VAL:HG22	1.93	0.51
1:A:3346:VAL:HG11	1:A:3414:ARG:CB	2.41	0.51
1:A:4584:ASP:OD2	1:A:4585:SER:N	2.44	0.51
1:A:4720:VAL:O	1:A:4724:VAL:HG23	2.10	0.51
1:A:4998:LYS:HB3	1:A:5003:HIS:CE1	2.45	0.51
1:D:4928:LEU:HA	1:D:4931:ILE:HD12	1.91	0.51
1:D:5000:GLU:HA	1:D:5003:HIS:CD2	2.45	0.51
1:C:1421:ARG:O	1:C:1570:LYS:NZ	2.33	0.51
1:A:774:ASP:OD2	1:A:1470:ARG:NH2	2.43	0.51
2:E:52:LYS:O	2:E:53:GLN:HB2	2.11	0.51
2:G:52:LYS:O	2:G:53:GLN:HB2	2.11	0.51
1:B:232:THR:HG21	1:B:252:VAL:HG21	1.92	0.51
1:B:3752:SER:O	1:B:3756:LYS:HG3	2.11	0.51
1:B:4000:MET:HE1	1:B:4061:PHE:CB	2.41	0.51
1:D:232:THR:HG21	1:D:252:VAL:HG21	1.93	0.51
1:D:399:GLN:O	1:D:403:MET:HG3	2.11	0.51
1:D:3036:LYS:O	1:D:3039:ILE:HG22	2.11	0.51
1:C:358:THR:HG1	1:C:383:HIS:HD1	1.57	0.51
1:C:2434:GLY:O	1:C:2508:ARG:NE	2.36	0.51
1:C:2974:ILE:O	1:C:2978:GLU:OE1	2.29	0.51
1:A:1828:ASP:N	1:A:1828:ASP:OD1	2.43	0.51
1:A:3036:LYS:O	1:A:3039:ILE:HG22	2.11	0.51
1:A:3230:LEU:HD23	1:A:3230:LEU:H	1.75	0.51
2:H:16:PRO:HB2	2:H:50:LEU:HD11	1.92	0.51
2:F:16:PRO:HB2	2:F:50:LEU:HD11	1.92	0.51
1:B:2668:SER:O	1:B:2669:GLU:CB	2.57	0.51
1:B:3169:LEU:CD1	1:B:3194:LEU:HD11	2.41	0.51
1:B:3699:HIS:ND1	1:B:3703:LEU:HD11	2.26	0.51
1:D:2199:ARG:NE	1:D:2246:ASN:OD1	2.37	0.51
1:D:3316:LEU:HD21	1:D:3346:VAL:HG23	1.93	0.51
1:D:3401:LEU:O	1:D:3405:LEU:HD13	2.11	0.51
1:C:2871:LEU:HD13	1:C:2874:MET:HE1	1.92	0.51
1:C:3316:LEU:HD21	1:C:3346:VAL:HG23	1.93	0.51
1:C:3633:VAL:O	1:C:3637:ARG:HG2	2.11	0.51
1:B:80:GLU:O	1:B:84:ASN:ND2	2.44	0.51
1:B:2021:CYS:O	1:B:2028:ARG:NH2	2.44	0.51
1:B:3690:VAL:O	1:B:3693:LYS:HE3	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4655:PHE:HB2	1:B:4796:MET:HE1	1.93	0.51
1:D:2577:ILE:HD12	1:D:2577:ILE:H	1.75	0.51
1:D:3052:HIS:HA	1:D:3127:GLN:NE2	2.25	0.51
1:D:3633:VAL:O	1:D:3637:ARG:HG2	2.11	0.51
1:C:1828:ASP:OD1	1:C:1828:ASP:N	2.43	0.51
1:C:3084:GLY:O	1:C:3089:LYS:NZ	2.37	0.51
1:A:3395:ARG:HD3	1:A:3453:ARG:NH1	2.25	0.50
1:A:4655:PHE:HB2	1:A:4796:MET:HE1	1.93	0.50
2:G:6:THR:HG23	2:G:6:THR:O	2.10	0.50
1:D:2327:GLY:CA	1:D:2330:ARG:NH2	2.74	0.50
1:D:3169:LEU:CD1	1:D:3194:LEU:HD11	2.41	0.50
1:C:2327:GLY:CA	1:C:2330:ARG:NH2	2.74	0.50
1:C:3230:LEU:HD23	1:C:3230:LEU:H	1.75	0.50
1:C:3455:GLU:O	1:C:3459:VAL:HG23	2.11	0.50
1:A:80:GLU:O	1:A:84:ASN:ND2	2.44	0.50
2:E:6:THR:HG23	2:E:6:THR:O	2.10	0.50
2:G:16:PRO:HB2	2:G:50:LEU:HD11	1.93	0.50
1:B:2974:ILE:O	1:B:2978:GLU:OE1	2.29	0.50
1:B:4584:ASP:OD2	1:B:4585:SER:N	2.44	0.50
1:D:2974:ILE:O	1:D:2978:GLU:OE1	2.29	0.50
1:D:4716:TRP:CD1	6:D:5304:IBM:H12	2.47	0.50
1:C:80:GLU:O	1:C:84:ASN:ND2	2.44	0.50
1:C:3104:GLU:HA	1:C:3107:VAL:HG22	1.93	0.50
1:C:3752:SER:O	1:C:3756:LYS:HG3	2.11	0.50
1:C:4576:ILE:HG23	1:C:4639:MET:SD	2.51	0.50
1:A:160:GLY:O	1:D:3984:ARG:NH2	2.38	0.50
1:A:3169:LEU:CD1	1:A:3194:LEU:HD11	2.41	0.50
1:A:4655:PHE:HA	1:A:4796:MET:HE1	1.94	0.50
1:A:4938:ASP:OD1	1:D:4944:ARG:NE	2.44	0.50
2:E:16:PRO:HB2	2:E:50:LEU:HD11	1.92	0.50
1:B:3104:GLU:HA	1:B:3107:VAL:HG22	1.93	0.50
1:B:3248:ARG:NH2	1:B:3252:ASP:OD1	2.37	0.50
1:B:4182:GLU:OE1	1:B:4983:HIS:NE2	2.44	0.50
1:D:4584:ASP:OD2	1:D:4585:SER:N	2.44	0.50
1:C:774:ASP:OD2	1:C:1470:ARG:NH2	2.43	0.50
1:A:2552:ARG:O	1:A:2556:LEU:HD23	2.12	0.50
1:A:2623:LEU:O	1:A:2627:VAL:HG23	2.11	0.50
1:A:3633:VAL:O	1:A:3637:ARG:HG2	2.11	0.50
1:B:3036:LYS:O	1:B:3039:ILE:HG22	2.11	0.50
1:B:3455:GLU:O	1:B:3459:VAL:HG23	2.11	0.50
1:D:1828:ASP:N	1:D:1828:ASP:OD1	2.43	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3036:LYS:O	1:C:3039:ILE:HG22	2.11	0.50
1:C:3690:VAL:O	1:C:3693:LYS:CE	2.60	0.50
1:A:236:ALA:O	1:A:237:ASP:OD1	2.30	0.50
1:A:988:LEU:CB	1:A:1039:LEU:HD21	2.42	0.50
1:A:1978:ALA:CB	1:A:1982:ARG:HH12	2.18	0.50
1:A:2111:VAL:HG12	1:A:2113:SER:H	1.77	0.50
1:A:3226:GLU:O	1:A:3227:ARG:HB2	2.12	0.50
1:A:4021:LYS:O	1:A:4025:VAL:HG23	2.12	0.50
1:B:236:ALA:O	1:B:237:ASP:OD1	2.30	0.50
1:B:3690:VAL:O	1:B:3693:LYS:CE	2.60	0.50
1:D:774:ASP:OD2	1:D:1470:ARG:NH2	2.43	0.50
1:D:3445:TRP:O	1:D:3452:LYS:NZ	2.43	0.50
1:C:3699:HIS:ND1	1:C:3703:LEU:HD11	2.26	0.50
1:A:232:THR:HG21	1:A:252:VAL:HG21	1.92	0.50
1:A:3401:LEU:O	1:A:3405:LEU:HD13	2.11	0.50
1:A:3957:VAL:O	1:A:3961:VAL:HG23	2.12	0.50
2:F:52:LYS:O	2:F:53:GLN:HB2	2.11	0.50
1:B:98:HIS:C	1:B:99:ARG:HD2	2.31	0.50
1:D:80:GLU:O	1:D:84:ASN:ND2	2.44	0.50
1:D:1978:ALA:CB	1:D:1982:ARG:HH12	2.18	0.50
1:D:3455:GLU:O	1:D:3459:VAL:HG23	2.11	0.50
1:C:2685:SER:O	1:C:2689:LYS:HG2	2.12	0.50
1:A:1141:ARG:HB2	1:D:3477:LYS:HZ3	1.77	0.50
1:A:2165:LEU:HD21	1:A:2177:LEU:HD23	1.94	0.50
1:A:3813:GLN:O	1:A:3899:PHE:HZ	1.95	0.50
1:B:2165:LEU:HD21	1:B:2177:LEU:HD23	1.94	0.50
1:B:2973:PHE:CE2	1:B:2995:ILE:HG12	2.47	0.50
1:B:3633:VAL:O	1:B:3637:ARG:HG2	2.11	0.50
1:D:1978:ALA:C	1:D:1982:ARG:NH1	2.65	0.50
1:D:2165:LEU:HD21	1:D:2177:LEU:HD23	1.94	0.50
1:D:2552:ARG:O	1:D:2556:LEU:HD23	2.12	0.50
1:D:3266:MET:SD	1:D:3270:ILE:HG13	2.52	0.50
1:D:3957:VAL:O	1:D:3961:VAL:HG23	2.12	0.50
1:D:4655:PHE:CB	1:D:4796:MET:HE1	2.42	0.50
1:C:2165:LEU:HD21	1:C:2177:LEU:HD23	1.94	0.50
1:C:3957:VAL:O	1:C:3961:VAL:HG23	2.12	0.50
1:C:4583:SER:OG	1:C:4585:SER:O	2.29	0.50
1:A:4716:TRP:CD1	6:A:5304:IBM:H12	2.47	0.50
1:B:988:LEU:CB	1:B:1039:LEU:HD21	2.42	0.50
1:B:2430:ILE:HG13	1:B:2502:MET:HE1	1.94	0.50
1:B:2685:SER:O	1:B:2689:LYS:HG2	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3230:LEU:HD23	1:B:3230:LEU:H	1.75	0.50
1:B:4021:LYS:O	1:B:4025:VAL:HG23	2.12	0.50
1:D:897:ARG:NH2	1:D:906:CYS:SG	2.73	0.50
1:D:2494:PHE:HD1	1:D:2498:HIS:HD1	1.59	0.50
1:D:3226:GLU:O	1:D:3227:ARG:HB2	2.12	0.50
1:C:236:ALA:O	1:C:237:ASP:OD1	2.30	0.50
1:C:3401:LEU:O	1:C:3405:LEU:HD13	2.11	0.50
1:A:3448:SER:O	1:A:3452:LYS:NZ	2.37	0.50
1:B:2552:ARG:O	1:B:2556:LEU:HD23	2.12	0.50
1:B:4655:PHE:CB	1:B:4796:MET:HE1	2.42	0.50
1:C:232:THR:HG21	1:C:252:VAL:HG21	1.93	0.50
1:C:887:ILE:CD1	1:C:961:MET:HE3	2.42	0.50
1:C:1978:ALA:C	1:C:1982:ARG:NH1	2.65	0.50
1:A:2974:ILE:O	1:A:2978:GLU:OE1	2.29	0.49
1:A:3752:SER:O	1:A:3756:LYS:HG3	2.11	0.49
1:A:4182:GLU:OE1	1:A:4983:HIS:NE2	2.44	0.49
2:E:25:HIS:CE1	2:E:45:PRO:HG3	2.47	0.49
2:G:25:HIS:CE1	2:G:45:PRO:HG3	2.47	0.49
1:B:1749:PRO:HD2	1:B:1758:ARG:HH12	1.75	0.49
1:D:236:ALA:O	1:D:237:ASP:OD1	2.30	0.49
1:D:2440:MET:O	1:D:2443:ILE:N	2.45	0.49
1:D:2524:VAL:HG23	1:D:2525:GLY:N	2.27	0.49
1:C:257:ARG:O	1:C:284:HIS:NE2	2.43	0.49
1:C:642:THR:HG23	1:C:1613:LEU:HD12	1.94	0.49
1:C:988:LEU:CB	1:C:1039:LEU:HD21	2.42	0.49
1:C:2581:SER:O	1:C:2585:THR:HG23	2.12	0.49
1:C:3169:LEU:CD1	1:C:3194:LEU:HD11	2.41	0.49
1:A:925:SER:O	1:A:928:THR:OG1	2.23	0.49
1:A:1978:ALA:C	1:A:1982:ARG:NH1	2.65	0.49
1:A:3716:LEU:HD21	1:A:3782:MET:HE3	1.94	0.49
2:H:6:THR:HG23	2:H:6:THR:O	2.10	0.49
2:H:52:LYS:O	2:H:53:GLN:HB2	2.11	0.49
1:B:642:THR:HG23	1:B:1613:LEU:HD12	1.94	0.49
1:B:2494:PHE:HD1	1:B:2498:HIS:HD1	1.59	0.49
1:B:2623:LEU:O	1:B:2627:VAL:HG23	2.11	0.49
1:B:3401:LEU:O	1:B:3405:LEU:HD13	2.11	0.49
1:B:4998:LYS:HB3	1:B:5003:HIS:CE1	2.44	0.49
1:A:3455:GLU:O	1:A:3459:VAL:HG23	2.11	0.49
1:A:4766:THR:HA	1:A:4769:MET:HE2	1.94	0.49
1:B:1978:ALA:C	1:B:1982:ARG:NH1	2.65	0.49
1:B:1978:ALA:O	1:B:1982:ARG:NH1	2.45	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2524:VAL:HG23	1:B:2525:GLY:N	2.27	0.49
1:B:3365:LEU:HD21	1:B:3405:LEU:CD1	2.43	0.49
1:D:900:ASN:OD1	1:D:901:LYS:N	2.46	0.49
1:D:3752:SER:O	1:D:3756:LYS:HG3	2.11	0.49
1:D:4655:PHE:HB2	1:D:4796:MET:HE1	1.93	0.49
1:C:4716:TRP:CD1	6:C:5304:IBM:H12	2.47	0.49
1:A:2494:PHE:HD1	1:A:2498:HIS:HD1	1.59	0.49
1:A:2973:PHE:CE2	1:A:2995:ILE:HG12	2.47	0.49
1:B:900:ASN:OD1	1:B:901:LYS:N	2.46	0.49
1:B:2538:THR:O	1:B:2542:SER:N	2.46	0.49
1:B:2581:SER:O	1:B:2585:THR:HG23	2.12	0.49
1:B:3316:LEU:HD21	1:B:3346:VAL:HG23	1.93	0.49
1:B:4716:TRP:CD1	6:B:5304:IBM:H12	2.47	0.49
1:D:887:ILE:CD1	1:D:961:MET:HE3	2.42	0.49
1:D:2111:VAL:HG12	1:D:2113:SER:H	1.77	0.49
1:D:2285:GLU:HG2	1:D:3858:MET:CE	2.43	0.49
1:D:3690:VAL:O	1:D:3693:LYS:CE	2.60	0.49
1:D:4021:LYS:O	1:D:4025:VAL:HG23	2.12	0.49
1:C:2440:MET:O	1:C:2443:ILE:N	2.45	0.49
1:C:2524:VAL:HG23	1:C:2525:GLY:N	2.27	0.49
1:C:3226:GLU:O	1:C:3227:ARG:HB2	2.12	0.49
1:A:399:GLN:O	1:A:403:MET:HG3	2.11	0.49
1:A:900:ASN:OD1	1:A:901:LYS:N	2.46	0.49
1:A:2524:VAL:HG23	1:A:2525:GLY:N	2.27	0.49
1:A:2660:GLY:CA	1:A:2666:VAL:HG12	2.43	0.49
1:A:3690:VAL:O	1:A:3693:LYS:CE	2.60	0.49
1:B:70:GLU:OE2	1:B:110:ARG:HD2	2.13	0.49
1:B:575:LEU:HA	1:B:578:ILE:HG12	1.95	0.49
1:D:642:THR:HG23	1:D:1613:LEU:HD12	1.94	0.49
1:D:988:LEU:CB	1:D:1039:LEU:HD21	2.42	0.49
1:D:2973:PHE:CE2	1:D:2995:ILE:HG12	2.47	0.49
1:D:3365:LEU:HD21	1:D:3405:LEU:CD1	2.43	0.49
1:D:3699:HIS:ND1	1:D:3703:LEU:HD11	2.26	0.49
1:D:3813:GLN:O	1:D:3899:PHE:HZ	1.95	0.49
1:C:2494:PHE:HD1	1:C:2498:HIS:HD1	1.59	0.49
1:C:2660:GLY:CA	1:C:2666:VAL:HG12	2.43	0.49
1:C:2973:PHE:CE2	1:C:2995:ILE:HG12	2.47	0.49
1:C:2974:ILE:HD12	1:C:3049:LEU:HD11	1.95	0.49
1:C:3365:LEU:HD21	1:C:3405:LEU:CD1	2.43	0.49
1:C:4655:PHE:HB2	1:C:4796:MET:HE1	1.93	0.49
1:A:70:GLU:OE2	1:A:110:ARG:HD2	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1999:ARG:CB	1:A:1999:ARG:HH11	2.26	0.49
1:A:3266:MET:SD	1:A:3270:ILE:HG13	2.52	0.49
1:B:3226:GLU:O	1:B:3227:ARG:HB2	2.12	0.49
1:D:3477:LYS:NZ	1:D:3479:ALA:HB2	2.28	0.49
1:C:70:GLU:OE2	1:C:110:ARG:HD2	2.13	0.49
1:C:2552:ARG:O	1:C:2556:LEU:HD23	2.12	0.49
1:C:3266:MET:SD	1:C:3270:ILE:HG13	2.52	0.49
1:C:4655:PHE:HA	1:C:4796:MET:HE1	1.92	0.49
1:A:2992:GLU:HA	1:A:2995:ILE:HD12	1.95	0.49
1:A:3316:LEU:HD21	1:A:3346:VAL:HG23	1.93	0.49
1:A:4655:PHE:CB	1:A:4796:MET:HE1	2.42	0.49
1:B:418:LEU:HD23	1:B:421:PHE:CZ	2.48	0.49
1:B:603:LEU:HD23	1:B:606:LEU:HD12	1.94	0.49
1:D:418:LEU:HD23	1:D:421:PHE:CZ	2.48	0.49
1:D:1112:ASP:OD1	1:D:1113:VAL:HG23	2.13	0.49
1:D:4000:MET:HE1	1:D:4058:ILE:HA	1.94	0.49
1:C:4655:PHE:CB	1:C:4796:MET:HE1	2.42	0.49
2:E:88:PRO:O	2:E:90:ILE:HD12	2.13	0.49
1:B:3299:GLY:O	1:B:3300:ALA:HB2	2.13	0.49
1:B:3477:LYS:NZ	1:B:3479:ALA:HB2	2.28	0.49
1:B:3539:ARG:NH2	1:B:3548:GLU:OE2	2.38	0.49
1:B:3957:VAL:O	1:B:3961:VAL:HG23	2.12	0.49
1:B:4937:ILE:HG12	1:C:4934:GLY:HA2	1.95	0.49
1:D:3177:THR:O	1:D:3179:LYS:NZ	2.36	0.49
1:D:4583:SER:OG	1:D:4585:SER:O	2.29	0.49
1:C:1999:ARG:CB	1:C:1999:ARG:HH11	2.26	0.49
1:C:4021:LYS:O	1:C:4025:VAL:HG23	2.12	0.49
1:A:2801:ASP:OD1	1:A:2802:LYS:N	2.46	0.49
1:A:2806:ARG:HE	1:A:2810:LYS:NZ	2.11	0.49
1:A:3365:LEU:HD21	1:A:3405:LEU:CD1	2.43	0.49
2:H:25:HIS:CE1	2:H:45:PRO:HG3	2.47	0.49
1:B:545:ASP:OD1	1:B:582:HIS:NE2	2.40	0.49
1:B:1676:LEU:HD22	1:B:2167:ILE:HD12	1.95	0.49
1:B:2285:GLU:HG2	1:B:3858:MET:CE	2.43	0.49
1:B:2440:MET:O	1:B:2443:ILE:N	2.45	0.49
1:C:2819:TRP:O	1:C:2821:TRP:CD1	2.66	0.49
1:A:732:SER:O	1:A:735:GLN:HG3	2.13	0.49
1:B:2660:GLY:CA	1:B:2666:VAL:HG12	2.43	0.49
1:B:2974:ILE:HD12	1:B:3049:LEU:HD11	1.94	0.49
1:B:4902:GLU:O	1:B:4913:ARG:CZ	2.61	0.49
1:D:1978:ALA:O	1:D:1982:ARG:NH1	2.45	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4000:MET:HG2	1:D:4061:PHE:CE2	2.48	0.49
1:C:900:ASN:OD1	1:C:901:LYS:N	2.46	0.49
1:C:1978:ALA:O	1:C:1982:ARG:NH1	2.45	0.49
1:C:3346:VAL:HG11	1:C:3414:ARG:CB	2.41	0.49
1:C:3354:LEU:HD11	1:C:3434:LEU:HD22	1.95	0.49
1:A:418:LEU:HD23	1:A:421:PHE:CZ	2.48	0.48
1:A:1069:TRP:CZ3	1:A:1112:ASP:HB2	2.48	0.48
1:A:1962:ALA:O	1:A:1966:VAL:HG23	2.13	0.48
1:A:1978:ALA:O	1:A:1982:ARG:NH1	2.45	0.48
1:A:2440:MET:O	1:A:2443:ILE:N	2.45	0.48
1:A:2685:SER:O	1:A:2689:LYS:HG2	2.12	0.48
1:A:3199:ALA:O	1:A:3283:ARG:NE	2.40	0.48
1:A:4000:MET:HG2	1:A:4061:PHE:CE2	2.48	0.48
2:H:88:PRO:O	2:H:90:ILE:HD12	2.13	0.48
2:G:79:ASP:OD2	2:G:80:TYR:HD1	1.96	0.48
2:F:79:ASP:OD2	2:F:80:TYR:HD1	1.96	0.48
1:B:2871:LEU:HD13	1:B:2874:MET:HE1	1.93	0.48
1:D:732:SER:O	1:D:735:GLN:HG3	2.13	0.48
1:D:1962:ALA:O	1:D:1966:VAL:HG23	2.13	0.48
1:D:1999:ARG:CB	1:D:1999:ARG:HH11	2.26	0.48
1:D:4934:GLY:HA2	1:C:4937:ILE:HG12	1.95	0.48
1:C:2285:GLU:HG2	1:C:3858:MET:CE	2.43	0.48
1:C:3448:SER:O	1:C:3452:LYS:NZ	2.37	0.48
1:C:4766:THR:HA	1:C:4769:MET:HE2	1.95	0.48
1:A:1733:GLU:HG2	1:A:2201:LEU:HD23	1.95	0.48
1:A:2107:GLN:O	1:A:3694:LYS:NZ	2.34	0.48
1:A:3108:GLU:O	1:A:3111:ARG:HB2	2.14	0.48
1:A:3354:LEU:HD11	1:A:3434:LEU:HD22	1.95	0.48
1:A:3699:HIS:ND1	1:A:3703:LEU:HD11	2.26	0.48
1:A:4902:GLU:O	1:A:4913:ARG:CZ	2.61	0.48
2:E:79:ASP:OD2	2:E:80:TYR:HD1	1.96	0.48
2:H:79:ASP:OD2	2:H:80:TYR:HD1	1.96	0.48
2:F:25:HIS:CE1	2:F:45:PRO:HG3	2.47	0.48
1:B:732:SER:O	1:B:735:GLN:HG3	2.13	0.48
1:B:3108:GLU:O	1:B:3111:ARG:HB2	2.14	0.48
1:B:3266:MET:SD	1:B:3270:ILE:HG13	2.52	0.48
1:B:4000:MET:HG2	1:B:4061:PHE:CE2	2.48	0.48
1:B:4565:LEU:O	1:B:4569:LEU:HD13	2.13	0.48
1:D:70:GLU:OE2	1:D:110:ARG:HD2	2.13	0.48
1:D:879:HIS:CE1	1:D:921:ASN:HD22	2.31	0.48
1:D:1676:LEU:HD22	1:D:2167:ILE:HD12	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2660:GLY:CA	1:D:2666:VAL:HG12	2.43	0.48
1:D:4016:LEU:O	1:D:4020:GLN:HG3	2.13	0.48
1:C:603:LEU:HD23	1:C:606:LEU:HD12	1.94	0.48
1:C:1112:ASP:OD1	1:C:1113:VAL:HG23	2.13	0.48
1:C:4016:LEU:O	1:C:4020:GLN:HG3	2.13	0.48
1:C:4690:GLU:O	1:C:4691:GLN:HB2	2.13	0.48
1:A:3299:GLY:O	1:A:3300:ALA:HB2	2.13	0.48
2:E:2:VAL:CG2	2:E:58:GLY:HA2	2.44	0.48
2:H:2:VAL:CG2	2:H:58:GLY:HA2	2.44	0.48
1:B:266:ARG:NE	1:B:330:ASP:OD2	2.46	0.48
1:B:1999:ARG:CB	1:B:1999:ARG:HH11	2.26	0.48
1:B:2111:VAL:HG12	1:B:2113:SER:H	1.77	0.48
1:B:2199:ARG:NE	1:B:2246:ASN:OD1	2.37	0.48
1:B:2801:ASP:OD1	1:B:2802:LYS:N	2.46	0.48
1:B:2806:ARG:HE	1:B:2810:LYS:NZ	2.11	0.48
1:B:2819:TRP:O	1:B:2821:TRP:CD1	2.66	0.48
1:D:2806:ARG:HE	1:D:2810:LYS:NZ	2.11	0.48
1:D:3448:SER:O	1:D:3452:LYS:NZ	2.37	0.48
1:C:877:ASN:O	1:C:881:LEU:HD23	2.13	0.48
1:C:1676:LEU:HD22	1:C:2167:ILE:HD12	1.95	0.48
1:C:3412:LEU:HD11	1:C:3434:LEU:HD21	1.96	0.48
1:C:4902:GLU:O	1:C:4913:ARG:CZ	2.61	0.48
1:A:3477:LYS:NZ	1:A:3479:ALA:HB2	2.28	0.48
1:A:4000:MET:HE1	1:A:4058:ILE:HA	1.94	0.48
1:A:4690:GLU:O	1:A:4691:GLN:HB2	2.14	0.48
2:G:88:PRO:O	2:G:90:ILE:HD12	2.13	0.48
1:B:3813:GLN:O	1:B:3899:PHE:HZ	1.95	0.48
1:B:4583:SER:OG	1:B:4585:SER:O	2.29	0.48
1:D:102:LEU:HD23	1:D:162:LYS:HA	1.96	0.48
1:D:1069:TRP:CZ3	1:D:1112:ASP:HB2	2.48	0.48
1:D:1733:GLU:HG2	1:D:2201:LEU:HD23	1.95	0.48
1:D:2819:TRP:O	1:D:2821:TRP:CD1	2.66	0.48
1:D:3299:GLY:O	1:D:3300:ALA:HB2	2.13	0.48
1:D:3477:LYS:HZ2	1:D:3479:ALA:HB2	1.79	0.48
1:D:4170:ILE:H	1:D:4170:ILE:HD12	1.79	0.48
1:C:266:ARG:NE	1:C:330:ASP:OD2	2.46	0.48
1:C:3003:LEU:HB2	1:C:3004:PRO:HD3	1.95	0.48
1:C:3299:GLY:O	1:C:3300:ALA:HB2	2.13	0.48
1:A:102:LEU:HD23	1:A:162:LYS:HA	1.96	0.48
1:A:877:ASN:O	1:A:881:LEU:HD23	2.13	0.48
1:A:1979:LEU:N	1:A:1982:ARG:NH2	2.61	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2285:GLU:HG2	1:A:3858:MET:CE	2.43	0.48
1:A:2581:SER:O	1:A:2585:THR:HG23	2.12	0.48
1:A:3412:LEU:HD11	1:A:3434:LEU:HD21	1.96	0.48
1:A:4016:LEU:O	1:A:4020:GLN:HG3	2.13	0.48
2:F:88:PRO:O	2:F:90:ILE:HD12	2.13	0.48
1:B:722:TRP:CZ2	1:B:727:ALA:HB2	2.49	0.48
1:B:887:ILE:CD1	1:B:961:MET:HE3	2.43	0.48
1:B:4655:PHE:HA	1:B:4796:MET:HE1	1.92	0.48
1:B:4766:THR:HA	1:B:4769:MET:HE2	1.96	0.48
1:D:3940:LYS:O	1:D:4002:LYS:NZ	2.29	0.48
1:C:722:TRP:CZ2	1:C:727:ALA:HB2	2.49	0.48
1:C:879:HIS:CE1	1:C:921:ASN:HD22	2.31	0.48
1:C:2806:ARG:HE	1:C:2810:LYS:NZ	2.11	0.48
1:C:4565:LEU:O	1:C:4569:LEU:HD13	2.13	0.48
1:A:603:LEU:HD23	1:A:606:LEU:HD12	1.94	0.48
1:A:619:ASP:OD2	1:A:1680:ARG:NH2	2.47	0.48
1:A:879:HIS:CE1	1:A:921:ASN:HD22	2.31	0.48
1:A:1676:LEU:HD22	1:A:2167:ILE:HD12	1.95	0.48
1:A:2974:ILE:HD12	1:A:3049:LEU:HD11	1.94	0.48
1:B:1069:TRP:CZ3	1:B:1112:ASP:HB2	2.48	0.48
1:B:1791:VAL:O	1:B:1792:ALA:HB3	2.14	0.48
1:D:2581:SER:O	1:D:2585:THR:HG23	2.12	0.48
1:D:2685:SER:O	1:D:2689:LYS:HG2	2.12	0.48
1:D:2801:ASP:OD1	1:D:2802:LYS:N	2.46	0.48
1:D:3003:LEU:HB2	1:D:3004:PRO:HD3	1.95	0.48
1:D:3412:LEU:HD11	1:D:3434:LEU:HD21	1.96	0.48
1:D:3445:TRP:NE1	1:D:3455:GLU:OE1	2.45	0.48
1:C:418:LEU:HD23	1:C:421:PHE:CZ	2.48	0.48
1:C:575:LEU:HA	1:C:578:ILE:HG12	1.95	0.48
1:C:1069:TRP:CZ3	1:C:1112:ASP:HB2	2.48	0.48
1:C:2538:THR:O	1:C:2542:SER:N	2.46	0.48
1:C:2801:ASP:OD1	1:C:2802:LYS:N	2.46	0.48
1:C:3813:GLN:O	1:C:3899:PHE:HZ	1.95	0.48
1:C:4000:MET:HG2	1:C:4061:PHE:CE2	2.48	0.48
1:A:2434:GLY:O	1:A:2508:ARG:NE	2.36	0.48
1:A:4565:LEU:O	1:A:4569:LEU:HD13	2.13	0.48
1:B:1733:GLU:HG2	1:B:2201:LEU:HD23	1.95	0.48
1:D:266:ARG:NE	1:D:330:ASP:OD2	2.46	0.48
1:D:796:ARG:O	1:D:1622:GLU:O	2.32	0.48
1:D:2000:SER:O	1:D:2005:GLN:NE2	2.41	0.48
1:D:2992:GLU:HA	1:D:2995:ILE:HD12	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1999:ARG:O	1:C:1999:ARG:CG	2.62	0.48
1:C:3108:GLU:O	1:C:3111:ARG:HB2	2.14	0.48
1:C:3477:LYS:NZ	1:C:3479:ALA:HB2	2.28	0.48
1:C:4170:ILE:H	1:C:4170:ILE:HD12	1.79	0.48
1:A:266:ARG:NE	1:A:330:ASP:OD2	2.46	0.48
1:A:642:THR:HG23	1:A:1613:LEU:HD12	1.94	0.48
1:A:1112:ASP:OD1	1:A:1113:VAL:HG23	2.13	0.48
1:A:1999:ARG:O	1:A:1999:ARG:CG	2.62	0.48
1:A:2475:GLN:OE1	1:A:2488:PRO:HB3	2.14	0.48
1:B:358:THR:OG1	1:B:383:HIS:ND1	2.42	0.48
1:B:796:ARG:O	1:B:1622:GLU:O	2.32	0.48
1:B:1979:LEU:N	1:B:1982:ARG:NH2	2.61	0.48
1:B:3412:LEU:HD11	1:B:3434:LEU:HD21	1.95	0.48
1:B:4690:GLU:O	1:B:4691:GLN:HB2	2.13	0.48
1:D:856:VAL:HG12	1:D:991:ASN:ND2	2.29	0.48
1:D:2974:ILE:HD12	1:D:3049:LEU:HD11	1.94	0.48
1:D:4766:THR:HA	1:D:4769:MET:HE2	1.94	0.48
1:D:4938:ASP:OD1	1:C:4944:ARG:NE	2.47	0.48
1:C:1962:ALA:O	1:C:1966:VAL:HG23	2.13	0.48
1:C:2111:VAL:HG12	1:C:2113:SER:H	1.77	0.48
1:C:4204:GLN:HB3	1:C:4245:MET:HG2	1.95	0.48
1:A:39:ALA:O	1:A:111:HIS:NE2	2.47	0.48
1:A:2871:LEU:HD13	1:A:2874:MET:SD	2.54	0.48
1:A:3003:LEU:HB2	1:A:3004:PRO:HD3	1.95	0.48
1:A:4170:ILE:HD12	1:A:4170:ILE:H	1.79	0.48
1:B:856:VAL:HG12	1:B:991:ASN:ND2	2.29	0.48
1:B:3615:SER:O	1:B:3616:LYS:HG2	2.14	0.48
1:B:4170:ILE:H	1:B:4170:ILE:HD12	1.79	0.48
1:D:39:ALA:O	1:D:111:HIS:NE2	2.47	0.48
1:D:575:LEU:HA	1:D:578:ILE:HG12	1.95	0.48
1:D:1676:LEU:HB3	1:D:2167:ILE:HD12	1.96	0.48
1:D:3354:LEU:HD11	1:D:3434:LEU:HD22	1.94	0.48
1:D:4182:GLU:OE1	1:D:4983:HIS:NE2	2.44	0.48
1:D:4673:ARG:O	1:D:4676:GLU:HG3	2.14	0.48
1:D:4902:GLU:O	1:D:4913:ARG:CZ	2.61	0.48
1:C:1791:VAL:O	1:C:1792:ALA:HB3	2.14	0.48
1:A:3996:PHE:HD1	1:A:4016:LEU:HD11	1.79	0.48
1:B:1112:ASP:OD1	1:B:1113:VAL:HG23	2.13	0.48
1:B:3003:LEU:HB2	1:B:3004:PRO:HD3	1.95	0.48
1:B:3337:ARG:O	1:B:3340:VAL:HG22	2.14	0.48
1:B:3354:LEU:HD11	1:B:3434:LEU:HD22	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3996:PHE:HD1	1:B:4016:LEU:HD11	1.79	0.48
1:D:417:GLY:O	1:D:421:PHE:CD1	2.67	0.48
1:D:603:LEU:HD23	1:D:606:LEU:HD12	1.94	0.48
1:D:877:ASN:O	1:D:881:LEU:HD23	2.13	0.48
1:D:2871:LEU:HD13	1:D:2874:MET:SD	2.54	0.48
1:C:309:THR:O	1:C:309:THR:HG22	2.14	0.48
1:C:732:SER:O	1:C:735:GLN:HG3	2.13	0.48
1:C:1733:GLU:HG2	1:C:2201:LEU:HD23	1.95	0.48
1:C:2992:GLU:HA	1:C:2995:ILE:HD12	1.95	0.48
1:A:417:GLY:O	1:A:421:PHE:CD1	2.67	0.47
1:A:884:LEU:O	1:A:888:GLU:OE1	2.32	0.47
1:A:1791:VAL:O	1:A:1792:ALA:HB3	2.14	0.47
1:A:4204:GLN:HB3	1:A:4245:MET:HG2	1.96	0.47
2:H:20:GLN:HE21	2:H:106:LEU:HB3	1.78	0.47
2:G:2:VAL:CG2	2:G:58:GLY:HA2	2.44	0.47
1:B:879:HIS:CE1	1:B:921:ASN:HD22	2.31	0.47
1:B:886:ARG:HG3	1:B:891:TRP:HB2	1.96	0.47
1:B:2000:SER:O	1:B:2005:GLN:NE2	2.41	0.47
1:D:358:THR:OG1	1:D:383:HIS:ND1	2.42	0.47
1:D:886:ARG:HG3	1:D:891:TRP:HB2	1.96	0.47
1:D:2538:THR:O	1:D:2542:SER:N	2.46	0.47
1:D:3346:VAL:HG11	1:D:3414:ARG:CB	2.41	0.47
1:D:4690:GLU:O	1:D:4691:GLN:HB2	2.13	0.47
1:C:1749:PRO:HD2	1:C:1758:ARG:HH12	1.75	0.47
1:C:2475:GLN:OE1	1:C:2488:PRO:HB3	2.14	0.47
1:C:2904:LEU:O	1:C:2906:VAL:HG22	2.14	0.47
1:A:856:VAL:HG12	1:A:991:ASN:ND2	2.29	0.47
1:B:39:ALA:O	1:B:111:HIS:NE2	2.47	0.47
1:B:4204:GLN:HB3	1:B:4245:MET:HG2	1.96	0.47
1:D:1979:LEU:N	1:D:1982:ARG:NH2	2.61	0.47
1:D:2475:GLN:OE1	1:D:2488:PRO:HB3	2.14	0.47
1:C:891:TRP:CZ2	1:C:899:ASP:HA	2.50	0.47
1:C:1979:LEU:HD22	1:C:1982:ARG:HH21	1.78	0.47
1:C:1979:LEU:N	1:C:1982:ARG:NH2	2.61	0.47
1:C:3985:LEU:HD11	1:C:4026:MET:HE1	1.96	0.47
1:A:2430:ILE:HG13	1:A:2502:MET:HE1	1.96	0.47
1:A:2515:GLN:O	1:A:2519:LEU:HG	2.15	0.47
2:E:20:GLN:HE21	2:E:106:LEU:HB3	1.78	0.47
1:B:309:THR:HG22	1:B:309:THR:O	2.14	0.47
1:B:2475:GLN:OE1	1:B:2488:PRO:HB3	2.14	0.47
1:B:2583:LEU:O	1:B:2586:VAL:HG12	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2992:GLU:HA	1:B:2995:ILE:HD12	1.95	0.47
1:B:3985:LEU:HD11	1:B:4026:MET:HE1	1.96	0.47
1:B:4732:PHE:O	1:B:4736:ARG:NH1	2.45	0.47
1:D:3337:ARG:O	1:D:3340:VAL:HG22	2.15	0.47
1:D:3615:SER:O	1:D:3616:LYS:HG2	2.14	0.47
1:D:3985:LEU:O	1:D:3989:VAL:HG23	2.14	0.47
1:C:102:LEU:HD23	1:C:162:LYS:HA	1.96	0.47
1:C:856:VAL:HG12	1:C:991:ASN:ND2	2.29	0.47
1:C:4158:PRO:HA	1:C:4161:ARG:NE	2.25	0.47
1:A:2819:TRP:O	1:A:2821:TRP:CD1	2.66	0.47
1:A:3105:LYS:O	1:A:3109:ASN:ND2	2.48	0.47
1:A:3615:SER:O	1:A:3616:LYS:HG2	2.14	0.47
1:A:3985:LEU:O	1:A:3989:VAL:HG23	2.14	0.47
1:B:619:ASP:OD2	1:B:1680:ARG:NH2	2.47	0.47
1:B:1962:ALA:O	1:B:1966:VAL:HG23	2.13	0.47
1:B:2515:GLN:O	1:B:2519:LEU:HG	2.15	0.47
1:B:3379:LEU:HD23	1:B:3382:GLU:OE2	2.15	0.47
1:C:417:GLY:O	1:C:421:PHE:CD1	2.67	0.47
1:C:886:ARG:HD3	1:C:891:TRP:CE3	2.50	0.47
1:C:2583:LEU:O	1:C:2586:VAL:HG12	2.15	0.47
1:C:3615:SER:O	1:C:3616:LYS:HG2	2.14	0.47
1:A:309:THR:O	1:A:309:THR:HG22	2.14	0.47
1:A:891:TRP:CZ2	1:A:899:ASP:HA	2.50	0.47
1:A:1979:LEU:HD22	1:A:1982:ARG:HH21	1.78	0.47
1:A:3878:ASP:N	1:A:3878:ASP:OD1	2.48	0.47
2:F:2:VAL:CG2	2:F:58:GLY:HA2	2.44	0.47
1:B:3765:TYR:CE1	1:B:4753:HIS:ND1	2.83	0.47
1:D:1506:GLN:OE1	1:C:2771:ILE:HG22	2.14	0.47
1:D:1979:LEU:HD22	1:D:1982:ARG:HH21	1.78	0.47
1:D:3108:GLU:O	1:D:3111:ARG:HB2	2.14	0.47
1:D:4565:LEU:O	1:D:4569:LEU:HD13	2.13	0.47
1:C:2355:ARG:NH2	1:C:2449:GLU:OE2	2.48	0.47
1:B:887:ILE:HD11	1:B:961:MET:CE	2.45	0.47
1:B:1979:LEU:HD22	1:B:1982:ARG:HH21	1.78	0.47
1:B:3528:THR:HG23	1:B:3573:MET:SD	2.55	0.47
1:D:886:ARG:HD3	1:D:891:TRP:CE3	2.50	0.47
1:D:891:TRP:CZ2	1:D:899:ASP:HA	2.50	0.47
1:D:1791:VAL:O	1:D:1792:ALA:HB3	2.14	0.47
1:D:3765:TYR:CE1	1:D:4753:HIS:ND1	2.83	0.47
1:D:3893:GLU:HA	1:D:3967:GLU:OE2	2.15	0.47
1:C:3379:LEU:HD23	1:C:3382:GLU:OE2	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3985:LEU:O	1:C:3989:VAL:HG23	2.14	0.47
1:C:3996:PHE:HD1	1:C:4016:LEU:HD11	1.79	0.47
1:C:4673:ARG:O	1:C:4676:GLU:HG3	2.14	0.47
1:A:575:LEU:HA	1:A:578:ILE:HG12	1.95	0.47
1:A:886:ARG:HD3	1:A:891:TRP:CE3	2.50	0.47
1:A:3505:VAL:O	1:A:3511:VAL:HG21	2.15	0.47
2:G:20:GLN:HE21	2:G:106:LEU:HB3	1.79	0.47
2:F:20:GLN:HE21	2:F:106:LEU:HB3	1.78	0.47
1:B:38:ALA:HB2	1:B:65:CYS:SG	2.55	0.47
1:B:102:LEU:HD23	1:B:162:LYS:HA	1.96	0.47
1:B:417:GLY:O	1:B:421:PHE:CD1	2.67	0.47
1:B:436:LEU:O	1:B:438:ILE:HD12	2.15	0.47
1:B:877:ASN:O	1:B:881:LEU:HD23	2.13	0.47
1:B:886:ARG:HD3	1:B:891:TRP:CE3	2.50	0.47
1:B:891:TRP:CZ2	1:B:899:ASP:HA	2.50	0.47
1:B:1999:ARG:O	1:B:1999:ARG:CG	2.62	0.47
1:B:2327:GLY:HA2	1:B:2330:ARG:NH2	2.29	0.47
1:B:2355:ARG:NH2	1:B:2449:GLU:OE2	2.48	0.47
1:B:2872:GLN:O	1:B:2875:ALA:N	2.48	0.47
1:B:2904:LEU:O	1:B:2906:VAL:HG22	2.14	0.47
1:B:3448:SER:O	1:B:3452:LYS:NZ	2.37	0.47
1:D:2583:LEU:O	1:D:2586:VAL:HG12	2.15	0.47
1:D:2904:LEU:O	1:D:2906:VAL:HG22	2.14	0.47
1:D:3505:VAL:O	1:D:3511:VAL:HG21	2.15	0.47
1:D:3996:PHE:HD1	1:D:4016:LEU:HD11	1.79	0.47
1:D:4204:GLN:HB3	1:D:4245:MET:HG2	1.96	0.47
1:D:4655:PHE:CA	1:D:4796:MET:CE	2.93	0.47
1:C:39:ALA:O	1:C:111:HIS:NE2	2.47	0.47
1:C:436:LEU:O	1:C:438:ILE:HD12	2.15	0.47
1:C:495:ASN:OD1	1:C:553:ARG:NH1	2.48	0.47
1:C:884:LEU:O	1:C:888:GLU:OE1	2.32	0.47
1:C:886:ARG:HG3	1:C:891:TRP:HB2	1.96	0.47
1:C:2872:GLN:O	1:C:2875:ALA:N	2.48	0.47
1:C:3337:ARG:O	1:C:3340:VAL:HG22	2.15	0.47
1:C:3765:TYR:CE1	1:C:4753:HIS:ND1	2.83	0.47
1:C:3878:ASP:N	1:C:3878:ASP:OD1	2.48	0.47
1:C:3893:GLU:HA	1:C:3967:GLU:OE2	2.15	0.47
1:A:796:ARG:O	1:A:1622:GLU:O	2.32	0.47
1:A:2327:GLY:HA2	1:A:2330:ARG:NH2	2.29	0.47
1:A:2913:ALA:O	1:A:2917:ALA:N	2.48	0.47
1:B:3985:LEU:O	1:B:3989:VAL:HG23	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4016:LEU:O	1:B:4020:GLN:HG3	2.13	0.47
1:D:309:THR:HG22	1:D:309:THR:O	2.14	0.47
1:D:722:TRP:CZ2	1:D:727:ALA:HB2	2.49	0.47
1:D:984:LEU:O	1:D:988:LEU:HD23	2.15	0.47
1:D:3878:ASP:N	1:D:3878:ASP:OD1	2.48	0.47
1:C:2199:ARG:NE	1:C:2246:ASN:OD1	2.37	0.47
1:C:2327:GLY:HA2	1:C:2330:ARG:NH2	2.29	0.47
1:C:2515:GLN:O	1:C:2519:LEU:HG	2.15	0.47
1:A:436:LEU:O	1:A:438:ILE:HD12	2.15	0.47
1:A:722:TRP:CZ2	1:A:727:ALA:HB2	2.49	0.47
1:A:1480:GLN:O	1:A:1480:GLN:HG2	2.15	0.47
1:A:1676:LEU:HB3	1:A:2167:ILE:HD12	1.96	0.47
1:A:3951:PHE:CE1	1:A:3999:MET:HE1	2.50	0.47
1:B:884:LEU:O	1:B:888:GLU:OE1	2.32	0.47
1:D:869:ARG:NH2	1:D:941:MET:HE1	2.29	0.47
1:D:3379:LEU:HD23	1:D:3382:GLU:OE2	2.15	0.47
1:C:796:ARG:O	1:C:1622:GLU:O	2.32	0.47
1:C:3105:LYS:O	1:C:3109:ASN:ND2	2.48	0.47
1:C:3445:TRP:O	1:C:3452:LYS:NZ	2.43	0.47
1:A:358:THR:HG1	1:A:383:HIS:HD1	1.57	0.47
1:A:880:GLU:O	1:A:884:LEU:HG	2.15	0.47
1:A:2199:ARG:NE	1:A:2246:ASN:OD1	2.37	0.47
1:A:4583:SER:OG	1:A:4585:SER:O	2.29	0.47
1:A:4673:ARG:O	1:A:4676:GLU:HG3	2.14	0.47
1:B:11:VAL:HG11	1:B:164:ARG:HD2	1.97	0.47
1:B:495:ASN:OD1	1:B:553:ARG:NH1	2.48	0.47
1:B:3105:LYS:O	1:B:3109:ASN:ND2	2.48	0.47
1:D:880:GLU:O	1:D:884:LEU:HG	2.15	0.47
1:D:1421:ARG:O	1:D:1570:LYS:NZ	2.33	0.47
1:D:1999:ARG:O	1:D:1999:ARG:CG	2.62	0.47
1:D:2515:GLN:O	1:D:2519:LEU:HG	2.15	0.47
1:D:3105:LYS:O	1:D:3109:ASN:ND2	2.48	0.47
1:D:3266:MET:SD	1:D:3266:MET:O	2.73	0.47
1:C:1032:LYS:C	1:C:1036:ARG:HH12	2.18	0.47
1:C:3445:TRP:NE1	1:C:3455:GLU:OE1	2.45	0.47
1:C:4182:GLU:OE1	1:C:4983:HIS:NE2	2.44	0.47
1:A:221:ARG:CZ	1:A:253:CYS:O	2.63	0.46
1:A:886:ARG:HG3	1:A:891:TRP:HB2	1.96	0.46
1:A:3893:GLU:HA	1:A:3967:GLU:OE2	2.15	0.46
1:B:1480:GLN:O	1:B:1480:GLN:HG2	2.15	0.46
1:B:1676:LEU:HB3	1:B:2167:ILE:HD12	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2693:GLN:OE1	1:B:2697:ARG:NH1	2.48	0.46
1:B:4677:LEU:HD23	1:B:4711:PHE:CE1	2.50	0.46
1:D:2006:ILE:O	1:D:2010:LEU:HD13	2.15	0.46
1:D:3528:THR:HG23	1:D:3573:MET:SD	2.55	0.46
1:C:3528:THR:HG23	1:C:3573:MET:SD	2.55	0.46
1:A:38:ALA:HB2	1:A:65:CYS:SG	2.55	0.46
1:A:3477:LYS:HZ3	1:B:1141:ARG:CB	2.28	0.46
1:A:3528:THR:HG23	1:A:3573:MET:SD	2.55	0.46
1:A:3573:MET:CE	1:A:3577:ARG:HH12	2.28	0.46
1:B:884:LEU:O	1:B:887:ILE:HB	2.15	0.46
1:B:2871:LEU:HD13	1:B:2874:MET:SD	2.54	0.46
1:D:884:LEU:O	1:D:888:GLU:OE1	2.32	0.46
1:D:2355:ARG:NH2	1:D:2449:GLU:OE2	2.48	0.46
1:D:2872:GLN:O	1:D:2875:ALA:N	2.48	0.46
1:C:11:VAL:HG11	1:C:164:ARG:HD2	1.97	0.46
1:C:38:ALA:HB2	1:C:65:CYS:SG	2.55	0.46
1:C:2871:LEU:HD13	1:C:2874:MET:SD	2.54	0.46
1:C:2913:ALA:O	1:C:2917:ALA:N	2.48	0.46
1:C:4677:LEU:HD23	1:C:4711:PHE:CE1	2.51	0.46
1:A:2355:ARG:NH2	1:A:2449:GLU:OE2	2.48	0.46
1:A:2583:LEU:O	1:A:2586:VAL:HG12	2.15	0.46
1:A:2693:GLN:OE1	1:A:2697:ARG:NH1	2.49	0.46
1:A:2904:LEU:O	1:A:2906:VAL:HG22	2.14	0.46
1:A:4677:LEU:HD23	1:A:4711:PHE:CE1	2.51	0.46
1:B:2861:ASP:OD1	1:B:2925:GLU:HG2	2.16	0.46
1:B:3390:GLY:O	1:B:3394:VAL:HG13	2.15	0.46
1:D:2913:ALA:O	1:D:2917:ALA:N	2.48	0.46
1:D:4749:GLU:O	1:D:4753:HIS:CD2	2.69	0.46
1:C:499:THR:HG23	1:C:502:HIS:H	1.81	0.46
1:C:3177:THR:O	1:C:3179:LYS:NZ	2.36	0.46
1:C:4655:PHE:CA	1:C:4796:MET:CE	2.93	0.46
1:A:213:TYR:HA	1:A:339:ILE:O	2.16	0.46
1:A:2006:ILE:O	1:A:2010:LEU:HD13	2.15	0.46
1:A:4934:GLY:HA2	1:D:4937:ILE:HG12	1.98	0.46
1:B:984:LEU:O	1:B:988:LEU:HD23	2.15	0.46
1:B:3445:TRP:NE1	1:B:3455:GLU:OE1	2.44	0.46
1:D:38:ALA:HB2	1:D:65:CYS:SG	2.55	0.46
1:D:221:ARG:CZ	1:D:253:CYS:O	2.63	0.46
1:D:3539:ARG:NH2	1:D:3548:GLU:OE2	2.38	0.46
1:C:1480:GLN:HG2	1:C:1480:GLN:O	2.15	0.46
1:C:3505:VAL:O	1:C:3511:VAL:HG21	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:4732:PHE:O	1:C:4736:ARG:NH1	2.45	0.46
1:A:887:ILE:CD1	1:A:961:MET:HE3	2.44	0.46
1:A:3337:ARG:O	1:A:3340:VAL:HG22	2.14	0.46
1:A:4749:GLU:O	1:A:4753:HIS:CD2	2.69	0.46
1:B:1032:LYS:C	1:B:1036:ARG:HH12	2.18	0.46
1:B:2913:ALA:O	1:B:2917:ALA:N	2.48	0.46
1:D:759:ILE:HG23	1:D:759:ILE:O	2.16	0.46
1:D:1480:GLN:O	1:D:1480:GLN:HG2	2.15	0.46
1:D:4677:LEU:HD23	1:D:4711:PHE:CE1	2.51	0.46
1:C:213:TYR:HA	1:C:339:ILE:O	2.16	0.46
1:C:323:LEU:C	1:C:324:ASP:OD1	2.54	0.46
1:C:884:LEU:O	1:C:887:ILE:HB	2.15	0.46
1:C:1676:LEU:HB3	1:C:2167:ILE:HD12	1.96	0.46
1:A:3379:LEU:HD23	1:A:3382:GLU:OE2	2.15	0.46
1:A:3765:TYR:CE1	1:A:4753:HIS:ND1	2.83	0.46
1:A:4892:ARG:HD3	1:B:4918:ILE:HD13	1.98	0.46
1:B:213:TYR:HA	1:B:339:ILE:O	2.16	0.46
1:B:1036:ARG:HH11	1:B:1036:ARG:HD3	1.48	0.46
1:B:1808:ARG:HG2	1:B:1854:PHE:CE1	2.51	0.46
1:B:2373:GLY:O	1:C:130:LYS:NZ	2.45	0.46
1:D:213:TYR:HA	1:D:339:ILE:O	2.16	0.46
1:D:2327:GLY:HA2	1:D:2330:ARG:NH2	2.29	0.46
1:D:2693:GLN:OE1	1:D:2697:ARG:NH1	2.49	0.46
1:C:619:ASP:OD2	1:C:1680:ARG:NH2	2.47	0.46
1:C:984:LEU:O	1:C:988:LEU:HD23	2.15	0.46
1:A:3573:MET:HB3	1:A:3577:ARG:NH1	2.31	0.46
1:A:4000:MET:HE1	1:A:4061:PHE:CB	2.46	0.46
1:A:4749:GLU:OE2	1:A:4750:ILE:HG23	2.16	0.46
1:A:4832:HIS:ND1	1:A:4833:ASN:OD1	2.45	0.46
1:B:221:ARG:CZ	1:B:253:CYS:O	2.63	0.46
1:B:880:GLU:O	1:B:884:LEU:HG	2.14	0.46
1:B:2313:LEU:HD12	1:B:2318:TYR:CD1	2.51	0.46
1:B:3893:GLU:HA	1:B:3967:GLU:OE2	2.15	0.46
1:B:4060:LYS:O	1:B:4063:ASP:OD1	2.34	0.46
1:D:323:LEU:C	1:D:324:ASP:OD1	2.54	0.46
1:D:887:ILE:HD11	1:D:961:MET:CE	2.45	0.46
1:D:2784:GLU:OE2	1:D:2785:LEU:HD12	2.16	0.46
1:C:221:ARG:CZ	1:C:253:CYS:O	2.63	0.46
1:C:3106:MET:O	1:C:3110:LEU:HD13	2.16	0.46
1:A:495:ASN:OD1	1:A:553:ARG:NH1	2.48	0.46
1:A:2538:THR:O	1:A:2542:SER:N	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2872:GLN:O	1:A:2875:ALA:N	2.48	0.46
1:B:421:PHE:CD2	1:B:421:PHE:O	2.69	0.46
1:B:875:ALA:CB	1:B:922:LEU:HD23	2.46	0.46
1:B:2771:ILE:HG22	1:C:1506:GLN:OE1	2.15	0.46
1:B:3505:VAL:O	1:B:3511:VAL:HG21	2.15	0.46
1:B:3573:MET:HB3	1:B:3577:ARG:NH1	2.31	0.46
1:D:421:PHE:O	1:D:421:PHE:CD2	2.69	0.46
1:D:2937:VAL:O	1:D:2937:VAL:HG13	2.16	0.46
1:D:3390:GLY:O	1:D:3394:VAL:HG13	2.15	0.46
1:C:14:LEU:HD13	1:C:202:MET:SD	2.56	0.46
1:C:421:PHE:CD2	1:C:421:PHE:O	2.69	0.46
1:C:880:GLU:O	1:C:884:LEU:HG	2.14	0.46
1:C:887:ILE:HD11	1:C:961:MET:CE	2.45	0.46
1:C:3266:MET:SD	1:C:3266:MET:O	2.74	0.46
1:C:3300:ALA:HB3	1:C:3301:PRO:CD	2.39	0.46
1:A:499:THR:HG23	1:A:502:HIS:H	1.81	0.46
1:A:875:ALA:CB	1:A:922:LEU:HD23	2.46	0.46
1:A:884:LEU:O	1:A:887:ILE:HB	2.15	0.46
1:A:3100:SER:O	1:A:3104:GLU:OE1	2.34	0.46
1:A:4242:ILE:HB	6:A:5304:IBM:H132	1.98	0.46
1:A:4640:GLU:HB3	1:A:4641:PRO:HD3	1.98	0.46
1:B:2138:LEU:O	1:B:2141:ALA:N	2.45	0.46
1:B:2949:SER:OG	1:B:2952:GLU:OE1	2.25	0.46
1:B:3100:SER:O	1:B:3104:GLU:OE1	2.34	0.46
1:B:4655:PHE:CA	1:B:4796:MET:CE	2.93	0.46
1:B:4673:ARG:O	1:B:4676:GLU:HG3	2.14	0.46
1:D:650:VAL:HA	1:D:658:GLN:NE2	2.31	0.46
1:D:891:TRP:HZ2	1:D:899:ASP:HA	1.81	0.46
1:D:3106:MET:O	1:D:3110:LEU:HD13	2.16	0.46
1:C:211:GLU:O	1:C:340:LYS:NZ	2.49	0.46
1:C:3390:GLY:O	1:C:3394:VAL:HG13	2.15	0.46
1:C:3573:MET:HB3	1:C:3577:ARG:NH1	2.31	0.46
1:C:4749:GLU:O	1:C:4753:HIS:CD2	2.69	0.46
1:A:11:VAL:HG11	1:A:164:ARG:HD2	1.97	0.46
1:A:759:ILE:O	1:A:759:ILE:HG23	2.16	0.46
1:A:984:LEU:O	1:A:988:LEU:HD23	2.15	0.46
1:A:4060:LYS:O	1:A:4063:ASP:OD1	2.34	0.46
1:A:4732:PHE:O	1:A:4736:ARG:NH1	2.45	0.46
1:B:384:MET:HG3	1:B:385:ASP:N	2.31	0.46
1:B:553:ARG:NH2	1:B:555:GLU:OE2	2.49	0.46
1:B:952:LYS:HD3	1:B:968:ALA:HB3	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3940:LYS:O	1:B:4002:LYS:NZ	2.29	0.46
1:B:4749:GLU:OE2	1:B:4750:ILE:HG23	2.16	0.46
1:D:436:LEU:O	1:D:438:ILE:HD12	2.15	0.46
1:D:884:LEU:O	1:D:887:ILE:HB	2.15	0.46
1:D:3573:MET:CE	1:D:3577:ARG:HH12	2.28	0.46
1:D:3573:MET:HB3	1:D:3577:ARG:NH1	2.31	0.46
1:D:3999:MET:HE3	1:D:3999:MET:HB3	1.80	0.46
1:C:2233:CYS:HB3	1:C:2237:CYS:HB3	1.66	0.46
1:C:2784:GLU:OE2	1:C:2785:LEU:HD12	2.16	0.46
1:C:4242:ILE:HB	6:C:5304:IBM:H132	1.98	0.46
1:C:4749:GLU:OE2	1:C:4750:ILE:HG23	2.16	0.46
1:A:650:VAL:HA	1:A:658:GLN:NE2	2.31	0.45
1:A:1141:ARG:CB	1:D:3477:LYS:HZ3	2.29	0.45
1:A:3266:MET:SD	1:A:3266:MET:O	2.74	0.45
1:A:4944:ARG:NE	1:B:4938:ASP:OD1	2.48	0.45
1:B:14:LEU:HD13	1:B:202:MET:SD	2.56	0.45
1:B:421:PHE:HE2	1:B:507:ALA:HB2	1.79	0.45
1:B:1979:LEU:HA	1:B:1982:ARG:NH2	2.31	0.45
1:B:2736:ASP:OD1	1:B:2736:ASP:O	2.35	0.45
1:B:2937:VAL:O	1:B:2937:VAL:HG13	2.16	0.45
1:B:3500:GLY:O	1:B:3504:SER:OG	2.33	0.45
1:B:4826:ILE:O	1:B:4829:SER:OG	2.25	0.45
1:D:11:VAL:HG11	1:D:164:ARG:HD2	1.97	0.45
1:D:14:LEU:HD13	1:D:202:MET:SD	2.56	0.45
1:D:619:ASP:OD2	1:D:1680:ARG:NH2	2.47	0.45
1:D:2313:LEU:HD12	1:D:2318:TYR:CD1	2.51	0.45
1:D:4242:ILE:HB	6:D:5304:IBM:H132	1.98	0.45
1:D:4794:TRP:O	1:D:4798:MET:HG2	2.16	0.45
1:C:759:ILE:HG23	1:C:759:ILE:O	2.16	0.45
1:C:891:TRP:HZ2	1:C:899:ASP:HA	1.81	0.45
1:C:2736:ASP:OD1	1:C:2736:ASP:O	2.34	0.45
1:A:211:GLU:O	1:A:340:LYS:NZ	2.49	0.45
1:A:952:LYS:HD3	1:A:968:ALA:HB3	1.98	0.45
1:A:2135:LEU:HD12	1:A:3658:LYS:HZ3	1.79	0.45
1:A:2373:GLY:O	1:B:130:LYS:NZ	2.41	0.45
1:B:1497:GLY:O	1:B:1501:VAL:HG13	2.16	0.45
1:B:2946:LEU:O	1:B:2946:LEU:CD2	2.62	0.45
1:B:3266:MET:SD	1:B:3266:MET:O	2.74	0.45
1:B:3878:ASP:OD1	1:B:3878:ASP:N	2.48	0.45
1:B:4242:ILE:HB	6:B:5304:IBM:H132	1.98	0.45
1:D:887:ILE:CD1	1:D:961:MET:CE	2.95	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:1979:LEU:HA	1:D:1982:ARG:NH2	2.31	0.45
1:D:4911:LEU:O	1:D:4914:VAL:HG22	2.16	0.45
1:C:1979:LEU:HA	1:C:1982:ARG:NH2	2.31	0.45
1:C:2313:LEU:HD12	1:C:2318:TYR:CD1	2.51	0.45
1:C:2693:GLN:OE1	1:C:2697:ARG:NH1	2.49	0.45
1:C:3500:GLY:O	1:C:3504:SER:OG	2.33	0.45
1:C:4545:GLU:OE2	1:C:4548:ARG:NH2	2.39	0.45
1:C:4794:TRP:O	1:C:4798:MET:HG2	2.16	0.45
1:C:4998:LYS:HB3	1:C:5003:HIS:CE1	2.45	0.45
1:A:323:LEU:C	1:A:324:ASP:OD1	2.54	0.45
1:A:574:VAL:HA	1:A:577:ILE:HG12	1.99	0.45
1:A:1497:GLY:O	1:A:1501:VAL:HG13	2.16	0.45
1:A:2736:ASP:O	1:A:2736:ASP:OD1	2.34	0.45
1:A:2861:ASP:OD1	1:A:2925:GLU:HG2	2.16	0.45
1:A:2937:VAL:O	1:A:2937:VAL:HG13	2.16	0.45
1:A:3207:GLU:OE1	1:A:3280:TYR:OH	2.31	0.45
1:A:4710:SER:O	1:A:4713:SER:OG	2.25	0.45
1:B:211:GLU:O	1:B:340:LYS:NZ	2.49	0.45
1:B:897:ARG:HH22	1:B:906:CYS:HG	1.63	0.45
1:B:1965:TYR:CE2	1:B:1969:LEU:HD11	2.52	0.45
1:B:3573:MET:CE	1:B:3577:ARG:HH12	2.28	0.45
1:B:4705:VAL:HG13	1:B:4711:PHE:CD1	2.51	0.45
1:B:4832:HIS:ND1	1:B:4833:ASN:OD1	2.45	0.45
1:D:495:ASN:OD1	1:D:553:ARG:NH1	2.48	0.45
1:D:2233:CYS:HB3	1:D:2237:CYS:HB3	1.67	0.45
1:D:3100:SER:O	1:D:3104:GLU:OE1	2.34	0.45
1:D:4749:GLU:OE2	1:D:4750:ILE:HG23	2.16	0.45
1:C:4095:LYS:O	1:C:4098:ASP:OD1	2.35	0.45
1:A:14:LEU:HD13	1:A:202:MET:SD	2.56	0.45
1:A:553:ARG:NH2	1:A:555:GLU:OE2	2.49	0.45
1:A:891:TRP:HZ2	1:A:899:ASP:HA	1.81	0.45
1:A:2313:LEU:HD12	1:A:2318:TYR:CD1	2.51	0.45
1:A:2340:PHE:CG	1:A:2435:ARG:HD3	2.51	0.45
1:A:3535:LEU:HD11	1:A:3559:LEU:HD22	1.99	0.45
1:A:4794:TRP:O	1:A:4798:MET:HG2	2.16	0.45
1:B:1105:ALA:HB2	1:B:1191:VAL:HG11	1.98	0.45
1:B:4794:TRP:O	1:B:4798:MET:HG2	2.16	0.45
1:D:384:MET:HG3	1:D:385:ASP:N	2.31	0.45
1:D:952:LYS:HD3	1:D:968:ALA:HB3	1.98	0.45
1:D:1497:GLY:O	1:D:1501:VAL:HG13	2.16	0.45
1:D:2340:PHE:CG	1:D:2435:ARG:HD3	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3535:LEU:HD11	1:D:3559:LEU:HD22	1.99	0.45
1:D:4658:ILE:HD13	1:D:4792:LEU:HB3	1.98	0.45
1:C:2430:ILE:HG13	1:C:2502:MET:HE1	1.97	0.45
1:C:2725:LYS:NZ	1:C:2734:ASN:O	2.49	0.45
1:A:421:PHE:CD2	1:A:421:PHE:O	2.69	0.45
1:A:2368:LEU:N	1:A:2368:LEU:HD12	2.32	0.45
1:A:4095:LYS:O	1:A:4098:ASP:OD1	2.35	0.45
1:B:336:PRO:HA	1:B:337:PRO:HD3	1.87	0.45
1:B:2014:ASP:O	1:B:2015:GLU:HB3	2.17	0.45
1:B:2233:CYS:HB3	1:B:2237:CYS:HB3	1.67	0.45
1:B:3924:LEU:O	1:B:3927:GLN:HG3	2.17	0.45
1:D:211:GLU:O	1:D:340:LYS:NZ	2.49	0.45
1:D:707:VAL:HG12	1:D:715:GLY:HA3	1.99	0.45
1:D:2725:LYS:NZ	1:D:2734:ASN:O	2.49	0.45
1:D:4060:LYS:O	1:D:4063:ASP:OD1	2.34	0.45
1:D:4705:VAL:HG13	1:D:4711:PHE:CD1	2.52	0.45
1:C:553:ARG:NH2	1:C:555:GLU:OE2	2.49	0.45
1:C:707:VAL:HG12	1:C:715:GLY:HA3	1.99	0.45
1:C:887:ILE:CD1	1:C:961:MET:CE	2.95	0.45
1:C:1280:GLN:O	1:C:1281:ASN:OD1	2.35	0.45
1:C:1965:TYR:CE2	1:C:1969:LEU:HD11	2.52	0.45
1:C:2937:VAL:HG13	1:C:2937:VAL:O	2.16	0.45
1:C:3207:GLU:OE1	1:C:3280:TYR:OH	2.31	0.45
1:C:3539:ARG:NH2	1:C:3548:GLU:OE2	2.38	0.45
1:A:565:TYR:O	1:A:569:ILE:HG12	2.17	0.45
1:A:2006:ILE:HD11	1:A:3641:LEU:HD11	1.98	0.45
1:A:2223:ILE:HG23	1:A:2223:ILE:O	2.17	0.45
1:B:323:LEU:C	1:B:324:ASP:OD1	2.54	0.45
1:B:499:THR:HG23	1:B:502:HIS:H	1.81	0.45
1:B:931:THR:O	1:B:935:LEU:HG	2.17	0.45
1:B:1996:ARG:NE	1:B:1999:ARG:NH1	2.61	0.45
1:D:730:VAL:HG21	1:D:764:VAL:HG12	1.99	0.45
1:D:931:THR:O	1:D:935:LEU:HG	2.17	0.45
1:D:1808:ARG:HG2	1:D:1854:PHE:CE1	2.51	0.45
1:D:2430:ILE:HG13	1:D:2502:MET:HE1	1.97	0.45
1:D:3924:LEU:O	1:D:3927:GLN:HG3	2.17	0.45
1:D:4000:MET:HE1	1:D:4061:PHE:CB	2.46	0.45
1:C:647:ASN:ND2	1:C:820:ARG:O	2.44	0.45
1:C:650:VAL:HA	1:C:658:GLN:NE2	2.31	0.45
1:C:857:ASP:OD2	1:C:858:THR:N	2.50	0.45
1:C:869:ARG:NH2	1:C:941:MET:HE1	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:875:ALA:CB	1:C:922:LEU:HD23	2.46	0.45
1:C:1497:GLY:O	1:C:1501:VAL:HG13	2.16	0.45
1:C:2006:ILE:O	1:C:2010:LEU:HD13	2.15	0.45
1:C:2006:ILE:HD11	1:C:3641:LEU:HD11	1.98	0.45
1:C:2340:PHE:CG	1:C:2435:ARG:HD3	2.51	0.45
1:C:3971:GLY:N	1:C:3972:PRO:HA	2.32	0.45
1:C:4640:GLU:HB3	1:C:4641:PRO:HD3	1.98	0.45
1:C:4705:VAL:HG13	1:C:4711:PHE:CD1	2.51	0.45
1:A:612:VAL:HG13	1:A:2167:ILE:O	2.17	0.45
1:A:887:ILE:CD1	1:A:961:MET:CE	2.95	0.45
1:A:941:MET:SD	1:A:944:GLU:HA	2.57	0.45
1:A:1032:LYS:O	1:A:1036:ARG:NH1	2.50	0.45
1:A:1808:ARG:HG2	1:A:1854:PHE:CE1	2.51	0.45
1:A:1979:LEU:HA	1:A:1982:ARG:NH2	2.31	0.45
1:A:3434:LEU:HD23	1:A:3517:MET:CE	2.47	0.45
1:A:3985:LEU:HD11	1:A:4026:MET:HE1	1.99	0.45
1:A:4911:LEU:O	1:A:4914:VAL:HG22	2.16	0.45
1:B:67:PHE:HB3	1:B:109:LEU:HD22	1.98	0.45
1:B:759:ILE:HG23	1:B:759:ILE:O	2.16	0.45
1:B:857:ASP:OD2	1:B:858:THR:N	2.50	0.45
1:B:891:TRP:HZ2	1:B:899:ASP:HA	1.81	0.45
1:B:1280:GLN:O	1:B:1281:ASN:OD1	2.35	0.45
1:B:3106:MET:O	1:B:3110:LEU:HD13	2.16	0.45
1:B:3229:ILE:HG13	1:B:3230:LEU:N	2.32	0.45
1:B:3300:ALA:CB	1:B:3301:PRO:HD3	2.40	0.45
1:B:3434:LEU:HD23	1:B:3517:MET:CE	2.47	0.45
1:B:4658:ILE:HD13	1:B:4792:LEU:HB3	1.98	0.45
1:D:612:VAL:HG13	1:D:2167:ILE:O	2.17	0.45
1:D:857:ASP:OD2	1:D:858:THR:N	2.50	0.45
1:D:1280:GLN:O	1:D:1281:ASN:OD1	2.35	0.45
1:D:2014:ASP:O	1:D:2015:GLU:HB3	2.17	0.45
1:D:2736:ASP:O	1:D:2736:ASP:OD1	2.34	0.45
1:D:2879:ALA:HB1	1:D:2908:TYR:OH	2.17	0.45
1:D:3229:ILE:HG13	1:D:3230:LEU:N	2.32	0.45
1:D:3300:ALA:HB3	1:D:3301:PRO:CD	2.39	0.45
1:C:421:PHE:HD1	1:C:421:PHE:H	1.64	0.45
1:C:1032:LYS:O	1:C:1036:ARG:NH1	2.50	0.45
1:C:2861:ASP:OD1	1:C:2925:GLU:HG2	2.16	0.45
1:C:3535:LEU:HD11	1:C:3559:LEU:HD22	1.99	0.45
1:C:4911:LEU:O	1:C:4914:VAL:HG22	2.16	0.45
1:A:384:MET:HG3	1:A:385:ASP:N	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1506:GLN:OE1	1:D:2771:ILE:HG22	2.17	0.45
1:A:2751:LEU:HD23	1:A:2755:ILE:CD1	2.47	0.45
1:A:3106:MET:O	1:A:3110:LEU:HD13	2.16	0.45
1:A:3390:GLY:O	1:A:3394:VAL:HG13	2.15	0.45
1:A:3445:TRP:NE1	1:A:3455:GLU:OE1	2.45	0.45
2:E:90:ILE:HG22	2:E:91:ILE:HG13	1.99	0.45
1:B:612:VAL:HG13	1:B:2167:ILE:O	2.17	0.45
1:B:3971:GLY:N	1:B:3972:PRO:HA	2.32	0.45
1:B:4095:LYS:O	1:B:4098:ASP:OD1	2.35	0.45
1:D:426:ARG:NH1	1:D:504:ALA:O	2.50	0.45
1:D:499:THR:HG23	1:D:502:HIS:H	1.81	0.45
1:D:875:ALA:CB	1:D:922:LEU:HD23	2.46	0.45
1:D:941:MET:SD	1:D:944:GLU:HA	2.57	0.45
1:D:1105:ALA:HB2	1:D:1191:VAL:HG11	1.98	0.45
1:D:1965:TYR:CE2	1:D:1969:LEU:HD11	2.52	0.45
1:D:2223:ILE:HG23	1:D:2223:ILE:O	2.17	0.45
1:D:2368:LEU:HD12	1:D:2368:LEU:N	2.32	0.45
1:D:2861:ASP:OD1	1:D:2925:GLU:HG2	2.16	0.45
1:D:3624:LEU:O	1:D:3628:ARG:HG2	2.17	0.45
1:D:3971:GLY:N	1:D:3972:PRO:HA	2.32	0.45
1:D:4095:LYS:O	1:D:4098:ASP:OD1	2.35	0.45
1:D:4826:ILE:O	1:D:4829:SER:OG	2.25	0.45
1:C:1808:ARG:HG2	1:C:1854:PHE:CE1	2.51	0.45
1:C:3573:MET:CE	1:C:3577:ARG:HH12	2.28	0.45
1:C:4060:LYS:O	1:C:4063:ASP:OD1	2.34	0.45
1:A:659:TYR:O	1:A:662:TRP:NE1	2.47	0.45
1:A:887:ILE:HD11	1:A:961:MET:CE	2.45	0.45
1:A:2226:PRO:HA	1:A:2267:MET:HE1	1.99	0.45
1:A:3791:GLY:O	1:A:3794:VAL:HG12	2.17	0.45
1:A:4655:PHE:CA	1:A:4796:MET:CE	2.93	0.45
2:F:54:GLU:O	1:B:1785:ALA:N	2.36	0.45
1:B:659:TYR:O	1:B:662:TRP:NE1	2.47	0.45
1:B:823:LEU:HD13	1:B:1626:TRP:CE3	2.52	0.45
1:B:1032:LYS:O	1:B:1036:ARG:NH1	2.50	0.45
1:B:2006:ILE:O	1:B:2010:LEU:HD13	2.15	0.45
1:B:2340:PHE:CG	1:B:2435:ARG:HD3	2.51	0.45
1:B:2879:ALA:HB1	1:B:2908:TYR:OH	2.17	0.45
1:B:4640:GLU:HB3	1:B:4641:PRO:HD3	1.98	0.45
1:B:4911:LEU:O	1:B:4914:VAL:HG22	2.16	0.45
1:D:877:ASN:HA	1:D:880:GLU:HG2	1.99	0.45
1:D:2234:ARG:O	1:D:2238:TYR:HD2	2.00	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3292:PRO:O	1:D:3294:PRO:HD3	2.17	0.45
1:C:384:MET:HG3	1:C:385:ASP:N	2.31	0.45
1:C:2000:SER:O	1:C:2005:GLN:NE2	2.41	0.45
1:C:2135:LEU:HD12	1:C:3658:LYS:HZ3	1.80	0.45
1:C:3199:ALA:O	1:C:3283:ARG:NE	2.40	0.45
1:C:3434:LEU:HD23	1:C:3517:MET:CE	2.47	0.45
1:C:3791:GLY:O	1:C:3794:VAL:HG12	2.17	0.45
1:C:4832:HIS:ND1	1:C:4833:ASN:OD1	2.45	0.45
1:A:823:LEU:HD13	1:A:1626:TRP:CE3	2.52	0.45
1:A:3229:ILE:HG13	1:A:3230:LEU:N	2.32	0.45
1:A:3300:ALA:CB	1:A:3301:PRO:HD3	2.40	0.45
1:B:650:VAL:HA	1:B:658:GLN:NE2	2.31	0.45
1:B:4749:GLU:O	1:B:4753:HIS:CD2	2.69	0.45
1:D:553:ARG:NH2	1:D:555:GLU:OE2	2.49	0.45
1:D:2309:SER:HB2	1:D:2321:ILE:O	2.17	0.45
1:D:3359:ILE:HD13	1:D:3434:LEU:HD13	1.99	0.45
1:D:3500:GLY:O	1:D:3504:SER:OG	2.33	0.45
1:D:3540:TYR:HB3	1:D:3604:TYR:CD2	2.52	0.45
1:C:725:HIS:ND1	1:C:725:HIS:O	2.50	0.45
1:A:67:PHE:HB3	1:A:109:LEU:HD22	1.98	0.44
1:A:234:SER:OG	1:A:240:ASP:OD2	2.29	0.44
1:A:421:PHE:HD1	1:A:421:PHE:H	1.64	0.44
1:A:1280:GLN:O	1:A:1281:ASN:OD1	2.35	0.44
1:A:1287:LEU:HD11	1:A:1599:MET:HE3	1.98	0.44
1:A:3289:PRO:O	1:A:3290:GLU:HB2	2.18	0.44
1:A:3359:ILE:HD13	1:A:3434:LEU:HD13	1.99	0.44
1:B:426:ARG:NH1	1:B:504:ALA:O	2.50	0.44
1:B:565:TYR:O	1:B:569:ILE:HG12	2.17	0.44
1:B:2784:GLU:OE2	1:B:2785:LEU:HD12	2.16	0.44
1:B:3292:PRO:O	1:B:3294:PRO:HD3	2.17	0.44
1:B:3535:LEU:HD11	1:B:3559:LEU:HD22	1.99	0.44
1:B:3868:ARG:HH12	1:B:3870:ASN:HB3	1.82	0.44
1:B:4944:ARG:NE	1:C:4938:ASP:OD1	2.47	0.44
1:D:823:LEU:HD13	1:D:1626:TRP:CE3	2.52	0.44
1:D:2738:ARG:HG2	1:D:2738:ARG:O	2.17	0.44
1:D:2821:TRP:CH2	1:D:2877:GLN:OE1	2.70	0.44
1:D:3632:VAL:O	1:D:3636:PHE:HD2	2.00	0.44
1:D:4732:PHE:O	1:D:4736:ARG:NH1	2.45	0.44
1:C:2014:ASP:O	1:C:2015:GLU:HB3	2.17	0.44
1:C:2309:SER:HB2	1:C:2321:ILE:O	2.17	0.44
1:C:3229:ILE:HG13	1:C:3230:LEU:N	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3359:ILE:HB	1:C:3360:PRO:HD3	2.00	0.44
1:C:3624:LEU:O	1:C:3628:ARG:HG2	2.17	0.44
1:A:257:ARG:O	1:A:284:HIS:NE2	2.43	0.44
1:A:2738:ARG:O	1:A:2738:ARG:HG2	2.17	0.44
1:A:2784:GLU:OE2	1:A:2785:LEU:HD12	2.16	0.44
1:A:4705:VAL:HG13	1:A:4711:PHE:CD1	2.52	0.44
1:B:1999:ARG:HH11	1:B:1999:ARG:HB3	1.83	0.44
1:B:2368:LEU:HD12	1:B:2368:LEU:N	2.32	0.44
1:B:3624:LEU:O	1:B:3628:ARG:HG2	2.17	0.44
1:B:3817:LEU:HD21	1:B:3898:ASP:HB3	1.99	0.44
1:C:2004:GLU:OE1	1:C:2004:GLU:N	2.51	0.44
1:C:2784:GLU:OE2	1:C:2785:LEU:CD1	2.65	0.44
1:C:2879:ALA:HB1	1:C:2908:TYR:OH	2.17	0.44
1:C:3924:LEU:O	1:C:3927:GLN:HG3	2.17	0.44
1:A:426:ARG:NH1	1:A:504:ALA:O	2.50	0.44
1:A:730:VAL:HG21	1:A:764:VAL:HG12	1.98	0.44
1:A:1105:ALA:HB2	1:A:1191:VAL:HG11	1.98	0.44
1:A:2309:SER:HB2	1:A:2321:ILE:O	2.17	0.44
1:A:2821:TRP:CH2	1:A:2877:GLN:OE1	2.70	0.44
1:A:2946:LEU:O	1:A:2946:LEU:CD2	2.61	0.44
1:A:3359:ILE:HB	1:A:3360:PRO:HD3	2.00	0.44
1:A:3868:ARG:HH12	1:A:3870:ASN:HB3	1.82	0.44
1:A:3924:LEU:O	1:A:3927:GLN:HG3	2.17	0.44
1:B:384:MET:HE1	1:C:166:GLY:O	2.17	0.44
1:B:707:VAL:HG12	1:B:715:GLY:HA3	1.99	0.44
1:B:2751:LEU:HD23	1:B:2755:ILE:CD1	2.47	0.44
1:B:2784:GLU:OE2	1:B:2785:LEU:CD1	2.65	0.44
1:B:2821:TRP:CH2	1:B:2877:GLN:OE1	2.70	0.44
1:B:3589:PRO:O	1:B:3593:VAL:HG13	2.18	0.44
1:D:725:HIS:ND1	1:D:725:HIS:O	2.50	0.44
1:D:797:HIS:CG	1:D:821:LEU:HD23	2.52	0.44
1:D:3434:LEU:HD23	1:D:3517:MET:CE	2.47	0.44
1:D:4943:LEU:O	1:D:4947:GLN:HG2	2.18	0.44
1:C:612:VAL:HG13	1:C:2167:ILE:O	2.17	0.44
1:C:1105:ALA:HB2	1:C:1191:VAL:HG11	1.98	0.44
1:C:1126:GLY:O	1:C:1142:PRO:HA	2.18	0.44
1:C:1999:ARG:HH11	1:C:1999:ARG:HB3	1.83	0.44
1:C:3100:SER:O	1:C:3104:GLU:OE1	2.34	0.44
1:C:3292:PRO:O	1:C:3294:PRO:HD3	2.17	0.44
1:C:3632:VAL:O	1:C:3636:PHE:HD2	2.00	0.44
1:A:725:HIS:ND1	1:A:725:HIS:O	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:857:ASP:OD2	1:A:858:THR:N	2.50	0.44
1:A:931:THR:O	1:A:935:LEU:HG	2.17	0.44
1:A:2004:GLU:N	1:A:2004:GLU:OE1	2.51	0.44
1:A:2785:LEU:O	1:A:2786:LYS:HG2	2.18	0.44
1:A:4658:ILE:HD13	1:A:4792:LEU:HB3	1.98	0.44
1:A:4937:ILE:HG12	1:B:4934:GLY:HA2	2.00	0.44
1:B:421:PHE:H	1:B:421:PHE:HD1	1.64	0.44
1:B:869:ARG:NH2	1:B:941:MET:HE1	2.32	0.44
1:B:887:ILE:CD1	1:B:961:MET:CE	2.95	0.44
1:B:941:MET:SD	1:B:944:GLU:HA	2.57	0.44
1:B:1126:GLY:O	1:B:1142:PRO:HA	2.18	0.44
1:B:1435:TYR:OH	1:B:1451:GLY:O	2.35	0.44
1:B:2309:SER:HB2	1:B:2321:ILE:O	2.17	0.44
1:B:3371:LYS:HE2	1:B:3371:LYS:HA	2.00	0.44
1:B:3527:PRO:HD2	1:B:3573:MET:SD	2.57	0.44
1:D:2751:LEU:CD1	1:D:2817:ILE:CD1	2.96	0.44
1:D:2785:LEU:O	1:D:2786:LYS:HG2	2.18	0.44
1:D:3359:ILE:HB	1:D:3360:PRO:HD3	2.00	0.44
1:D:3527:PRO:HD2	1:D:3573:MET:SD	2.57	0.44
1:D:3559:LEU:HD23	1:D:3559:LEU:O	2.18	0.44
1:D:3589:PRO:O	1:D:3593:VAL:HG13	2.18	0.44
1:D:3716:LEU:HD21	1:D:3782:MET:HE3	1.99	0.44
1:D:4640:GLU:HB3	1:D:4641:PRO:HD3	1.98	0.44
1:D:4983:HIS:O	3:D:5301:ATP:N6	2.51	0.44
1:C:823:LEU:HD13	1:C:1626:TRP:CE3	2.52	0.44
1:C:897:ARG:HG3	1:C:905:PRO:HD2	2.00	0.44
1:C:952:LYS:HD3	1:C:968:ALA:HB3	1.98	0.44
1:C:1287:LEU:HD11	1:C:1599:MET:HE3	1.98	0.44
1:C:2821:TRP:CH2	1:C:2877:GLN:OE1	2.70	0.44
1:C:3589:PRO:O	1:C:3593:VAL:HG13	2.18	0.44
1:A:2014:ASP:O	1:A:2015:GLU:HB3	2.17	0.44
1:A:2784:GLU:OE2	1:A:2785:LEU:CD1	2.65	0.44
1:A:3500:GLY:O	1:A:3504:SER:OG	2.33	0.44
1:A:3559:LEU:HD23	1:A:3559:LEU:O	2.18	0.44
1:A:4943:LEU:O	1:A:4947:GLN:HG2	2.18	0.44
1:B:262:LEU:HD12	1:B:274:LEU:HD11	2.00	0.44
1:B:646:PRO:HB2	1:B:648:ILE:CD1	2.48	0.44
1:B:725:HIS:O	1:B:725:HIS:ND1	2.50	0.44
1:B:1115:LEU:HD12	1:B:1193:SER:HB3	2.00	0.44
1:B:3359:ILE:HB	1:B:3360:PRO:HD3	2.00	0.44
1:B:4748:LEU:O	1:B:4751:THR:OG1	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4852:THR:HG23	1:B:4882:CYS:HB3	1.99	0.44
1:B:4943:LEU:O	1:B:4947:GLN:HG2	2.18	0.44
1:D:2703:LEU:N	1:D:2703:LEU:HD12	2.33	0.44
1:D:3199:ALA:O	1:D:3283:ARG:NE	2.40	0.44
1:C:67:PHE:HB3	1:C:109:LEU:HD22	1.98	0.44
1:C:552:ASP:OD1	1:C:553:ARG:N	2.51	0.44
1:C:931:THR:O	1:C:935:LEU:HG	2.17	0.44
1:C:941:MET:SD	1:C:944:GLU:HA	2.57	0.44
1:A:897:ARG:HG3	1:A:905:PRO:HD2	2.00	0.44
1:A:1965:TYR:CE2	1:A:1969:LEU:HD11	2.52	0.44
1:A:3589:PRO:O	1:A:3593:VAL:HG13	2.18	0.44
1:A:3971:GLY:N	1:A:3972:PRO:HA	2.32	0.44
1:B:897:ARG:HG3	1:B:905:PRO:HD2	2.00	0.44
1:B:1419:ASP:CG	1:B:1420:ASN:H	2.21	0.44
1:B:2006:ILE:HD11	1:B:3641:LEU:HD11	1.99	0.44
1:B:2527:LEU:O	1:B:2531:ARG:HG3	2.18	0.44
1:B:2785:LEU:O	1:B:2786:LYS:HG2	2.18	0.44
1:B:3289:PRO:O	1:B:3290:GLU:HB2	2.18	0.44
1:D:67:PHE:HB3	1:D:109:LEU:HD22	1.98	0.44
1:D:897:ARG:HG3	1:D:905:PRO:HD2	2.00	0.44
1:D:1435:TYR:OH	1:D:1451:GLY:O	2.35	0.44
1:D:2336:ARG:HG2	1:D:2435:ARG:HD2	2.00	0.44
1:D:3791:GLY:O	1:D:3794:VAL:HG12	2.17	0.44
1:D:3817:LEU:HD21	1:D:3898:ASP:HB3	1.99	0.44
1:D:4852:THR:HG23	1:D:4882:CYS:HB3	1.99	0.44
1:C:262:LEU:HD12	1:C:274:LEU:HD11	2.00	0.44
1:C:426:ARG:NH1	1:C:504:ALA:O	2.50	0.44
1:C:565:TYR:O	1:C:569:ILE:HG12	2.17	0.44
1:C:646:PRO:HB2	1:C:648:ILE:CD1	2.48	0.44
1:C:2234:ARG:O	1:C:2238:TYR:HD2	2.00	0.44
1:C:3127:GLN:O	1:C:3130:THR:HG22	2.18	0.44
1:C:4943:LEU:O	1:C:4947:GLN:HG2	2.18	0.44
1:A:262:LEU:HD12	1:A:274:LEU:HD11	2.00	0.44
1:A:2703:LEU:N	1:A:2703:LEU:HD12	2.32	0.44
1:A:2755:ILE:HD13	1:A:2810:LYS:HG2	2.00	0.44
1:B:552:ASP:OD1	1:B:553:ARG:N	2.51	0.44
1:B:1287:LEU:HD11	1:B:1599:MET:HE3	1.99	0.44
1:B:2234:ARG:O	1:B:2238:TYR:HD2	2.00	0.44
1:B:3199:ALA:O	1:B:3283:ARG:NE	2.40	0.44
1:D:294:THR:O	1:D:298:GLY:N	2.51	0.44
1:D:2123:LEU:O	1:D:2127:GLN:HG2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:877:ASN:HA	1:C:880:GLU:HG2	1.99	0.44
1:C:2223:ILE:O	1:C:2223:ILE:HG23	2.17	0.44
1:C:3527:PRO:HD2	1:C:3573:MET:SD	2.57	0.44
1:C:3540:TYR:HB3	1:C:3604:TYR:CD2	2.52	0.44
1:C:4658:ILE:HD13	1:C:4792:LEU:HB3	1.99	0.44
1:A:866:HIS:O	1:A:869:ARG:HG2	2.18	0.44
1:A:1126:GLY:O	1:A:1142:PRO:HA	2.18	0.44
1:A:1999:ARG:HH11	1:A:1999:ARG:HB3	1.83	0.44
1:A:2123:LEU:O	1:A:2127:GLN:HG2	2.18	0.44
1:A:2234:ARG:O	1:A:2238:TYR:HD2	2.00	0.44
1:A:2285:GLU:HG2	1:A:3858:MET:HE1	2.00	0.44
1:A:2336:ARG:HG2	1:A:2435:ARG:HD2	2.00	0.44
1:A:2751:LEU:HD11	1:A:2817:ILE:CD1	2.48	0.44
1:A:2756:ASN:HA	1:A:2806:ARG:HH12	1.83	0.44
1:B:574:VAL:HA	1:B:577:ILE:HG12	1.98	0.44
1:B:730:VAL:HG21	1:B:764:VAL:HG12	1.98	0.44
1:B:2091:PRO:O	1:B:2092:GLN:HB2	2.18	0.44
1:B:3365:LEU:HD11	1:B:3405:LEU:CD1	2.48	0.44
1:B:3540:TYR:HB3	1:B:3604:TYR:CD2	2.52	0.44
1:B:4000:MET:HE1	1:B:4058:ILE:HA	1.99	0.44
1:D:262:LEU:HD12	1:D:274:LEU:HD11	2.00	0.44
1:D:552:ASP:OD1	1:D:553:ARG:N	2.51	0.44
1:D:2006:ILE:HD11	1:D:3641:LEU:HD11	1.98	0.44
1:D:2527:LEU:O	1:D:2531:ARG:HG3	2.18	0.44
1:D:2625:ARG:NH1	1:D:2629:ASP:OD1	2.51	0.44
1:D:2755:ILE:HD13	1:D:2810:LYS:HG2	2.00	0.44
1:D:2784:GLU:OE2	1:D:2785:LEU:CD1	2.65	0.44
1:D:4710:SER:O	1:D:4713:SER:OG	2.25	0.44
1:C:421:PHE:HE2	1:C:507:ALA:HB2	1.79	0.44
1:C:559:GLY:O	1:C:563:VAL:HG23	2.18	0.44
1:C:574:VAL:HA	1:C:577:ILE:HG12	1.99	0.44
1:C:2091:PRO:O	1:C:2092:GLN:HB2	2.18	0.44
1:C:2368:LEU:N	1:C:2368:LEU:HD12	2.32	0.44
1:A:707:VAL:HG12	1:A:715:GLY:HA3	1.99	0.44
1:A:797:HIS:CG	1:A:821:LEU:HD23	2.52	0.44
1:A:2527:LEU:O	1:A:2531:ARG:HG3	2.18	0.44
1:A:3266:MET:SD	1:A:3270:ILE:HD11	2.58	0.44
1:A:3292:PRO:O	1:A:3294:PRO:HD3	2.17	0.44
1:A:3365:LEU:HD11	1:A:3405:LEU:CD1	2.48	0.44
1:A:3624:LEU:O	1:A:3628:ARG:HG2	2.17	0.44
1:A:4024:VAL:HG11	1:A:4142:ASN:HB3	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4933:GLN:HG3	1:B:4930:ALA:CB	2.48	0.44
1:B:234:SER:OG	1:B:240:ASP:OD2	2.29	0.44
1:B:797:HIS:CG	1:B:821:LEU:HD23	2.52	0.44
1:B:2434:GLY:O	1:B:2508:ARG:NE	2.36	0.44
1:B:2725:LYS:NZ	1:B:2734:ASN:O	2.49	0.44
1:B:2751:LEU:HD11	1:B:2817:ILE:CD1	2.48	0.44
1:B:3127:GLN:O	1:B:3130:THR:HG22	2.18	0.44
1:B:3266:MET:SD	1:B:3270:ILE:HD11	2.58	0.44
1:B:3791:GLY:O	1:B:3794:VAL:HG12	2.17	0.44
1:D:417:GLY:O	1:D:421:PHE:HD1	2.01	0.44
1:D:421:PHE:HD1	1:D:421:PHE:H	1.64	0.44
1:D:574:VAL:O	1:D:578:ILE:HG12	2.18	0.44
1:D:866:HIS:O	1:D:869:ARG:HG2	2.18	0.44
1:D:893:TYR:HB3	1:D:962:SER:OG	2.18	0.44
1:D:1999:ARG:HH11	1:D:1999:ARG:HB3	1.83	0.44
1:D:2751:LEU:HD23	1:D:2755:ILE:CD1	2.47	0.44
1:C:1436:SER:OG	1:C:1565:GLU:HB2	2.18	0.44
1:A:552:ASP:OD1	1:A:553:ARG:N	2.51	0.43
1:A:2879:ALA:HB1	1:A:2908:TYR:OH	2.17	0.43
1:A:3104:GLU:OE2	1:A:3167:ARG:HB3	2.18	0.43
2:G:90:ILE:HD11	1:C:1684:ALA:HA	2.00	0.43
1:B:2223:ILE:HG23	1:B:2223:ILE:O	2.17	0.43
1:B:2703:LEU:HD12	1:B:2703:LEU:N	2.32	0.43
1:B:2738:ARG:HG2	1:B:2738:ARG:O	2.17	0.43
1:B:2751:LEU:CD1	1:B:2817:ILE:CD1	2.96	0.43
1:B:3632:VAL:O	1:B:3636:PHE:HD2	2.00	0.43
1:D:574:VAL:HA	1:D:577:ILE:HG12	1.99	0.43
1:D:2004:GLU:OE1	1:D:2004:GLU:N	2.51	0.43
1:D:2091:PRO:O	1:D:2092:GLN:HB2	2.18	0.43
1:D:2756:ASN:HA	1:D:2806:ARG:HH12	1.83	0.43
1:C:3300:ALA:CB	1:C:3301:PRO:HD3	2.40	0.43
1:C:3999:MET:HE3	1:C:3999:MET:HB3	1.75	0.43
1:C:4710:SER:O	1:C:4713:SER:OG	2.24	0.43
1:A:574:VAL:O	1:A:578:ILE:HG12	2.18	0.43
1:A:1435:TYR:OH	1:A:1451:GLY:O	2.35	0.43
1:A:2212:VAL:HG11	1:A:2256:TYR:CZ	2.54	0.43
1:A:2499:LYS:O	1:A:2503:VAL:HG23	2.18	0.43
1:A:3391:GLU:HG2	1:A:3395:ARG:HH21	1.83	0.43
1:A:3540:TYR:HB3	1:A:3604:TYR:CD2	2.52	0.43
1:A:3891:LEU:O	1:A:3899:PHE:CD2	2.71	0.43
1:A:4826:ILE:O	1:A:4829:SER:OG	2.25	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4852:THR:HG23	1:A:4882:CYS:HB3	1.99	0.43
2:G:90:ILE:HG22	2:G:91:ILE:HG13	1.99	0.43
2:F:90:ILE:HG22	2:F:91:ILE:HG13	1.99	0.43
1:B:559:GLY:O	1:B:563:VAL:HG23	2.18	0.43
1:B:877:ASN:HA	1:B:880:GLU:HG2	1.99	0.43
1:B:2004:GLU:OE1	1:B:2004:GLU:N	2.51	0.43
1:B:2336:ARG:HG2	1:B:2435:ARG:HD2	2.00	0.43
1:D:1032:LYS:O	1:D:1036:ARG:NH1	2.50	0.43
1:D:3104:GLU:OE2	1:D:3167:ARG:HB3	2.18	0.43
1:D:3127:GLN:O	1:D:3130:THR:HG22	2.18	0.43
1:D:3266:MET:SD	1:D:3270:ILE:HD11	2.58	0.43
1:D:3289:PRO:O	1:D:3290:GLU:HB2	2.18	0.43
1:C:294:THR:O	1:C:298:GLY:N	2.50	0.43
1:C:797:HIS:CG	1:C:821:LEU:HD23	2.52	0.43
1:C:893:TYR:HB3	1:C:962:SER:OG	2.18	0.43
1:C:2703:LEU:HD12	1:C:2703:LEU:N	2.33	0.43
1:C:2751:LEU:HD23	1:C:2755:ILE:CD1	2.47	0.43
1:C:2751:LEU:CD1	1:C:2817:ILE:CD1	2.96	0.43
1:A:1419:ASP:CG	1:A:1420:ASN:H	2.21	0.43
1:A:1436:SER:OG	1:A:1565:GLU:HB2	2.18	0.43
1:A:3409:TYR:N	1:A:3410:PRO:HD2	2.33	0.43
1:A:3539:ARG:NH2	1:A:3548:GLU:OE2	2.38	0.43
2:E:38:SER:HB3	2:E:41:ASP:OD1	2.19	0.43
1:B:2499:LYS:O	1:B:2503:VAL:HG23	2.18	0.43
1:B:2755:ILE:HD13	1:B:2810:LYS:HG2	2.00	0.43
1:B:3359:ILE:HD13	1:B:3434:LEU:HD13	1.99	0.43
1:B:3559:LEU:O	1:B:3559:LEU:HD23	2.18	0.43
1:D:565:TYR:O	1:D:569:ILE:HG12	2.17	0.43
1:D:1287:LEU:HD11	1:D:1599:MET:HE3	2.00	0.43
1:D:2499:LYS:O	1:D:2503:VAL:HG23	2.19	0.43
1:C:345:LEU:HD21	1:C:389:PHE:CZ	2.54	0.43
1:C:417:GLY:O	1:C:421:PHE:HD1	2.01	0.43
1:C:730:VAL:HG21	1:C:764:VAL:HG12	1.98	0.43
1:C:1122:TYR:CE2	1:C:1182:ILE:HD12	2.54	0.43
1:C:1419:ASP:CG	1:C:1420:ASN:H	2.21	0.43
1:C:2738:ARG:O	1:C:2738:ARG:HG2	2.17	0.43
1:C:2785:LEU:O	1:C:2786:LYS:HG2	2.18	0.43
1:C:3266:MET:SD	1:C:3270:ILE:HD11	2.58	0.43
1:C:3391:GLU:HG2	1:C:3395:ARG:HH21	1.83	0.43
1:C:3409:TYR:N	1:C:3410:PRO:HD2	2.33	0.43
1:A:646:PRO:HB2	1:A:648:ILE:CD1	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:H:90:ILE:HG22	2:H:91:ILE:HG13	1.99	0.43
1:B:893:TYR:HB3	1:B:962:SER:OG	2.18	0.43
1:B:2604:GLU:O	1:B:2608:MET:HG2	2.19	0.43
1:B:4253:GLU:OE1	1:B:4557:ARG:NE	2.52	0.43
1:D:867:LEU:HA	1:D:870:ILE:HG22	2.01	0.43
1:D:1122:TYR:CE2	1:D:1182:ILE:HD12	2.53	0.43
1:D:2668:SER:O	1:D:2669:GLU:HB2	2.18	0.43
1:C:574:VAL:O	1:C:578:ILE:HG12	2.19	0.43
1:C:1435:TYR:OH	1:C:1451:GLY:O	2.35	0.43
1:C:1989:ALA:O	1:C:1992:ALA:HB3	2.19	0.43
1:C:2336:ARG:HG2	1:C:2435:ARG:HD2	2.00	0.43
1:C:2499:LYS:O	1:C:2503:VAL:HG23	2.19	0.43
1:C:3559:LEU:O	1:C:3559:LEU:HD23	2.18	0.43
1:A:877:ASN:HA	1:A:880:GLU:HG2	1.99	0.43
1:A:2091:PRO:O	1:A:2092:GLN:HB2	2.18	0.43
1:A:2751:LEU:CD1	1:A:2817:ILE:CD1	2.96	0.43
1:A:3632:VAL:O	1:A:3636:PHE:HD2	2.00	0.43
1:A:4253:GLU:OE1	1:A:4557:ARG:NE	2.52	0.43
1:A:4545:GLU:OE2	1:A:4548:ARG:NH2	2.39	0.43
1:B:2123:LEU:O	1:B:2127:GLN:HG2	2.18	0.43
1:B:4024:VAL:HG11	1:B:4142:ASN:HB3	2.00	0.43
1:B:4933:GLN:HG3	1:C:4930:ALA:CB	2.48	0.43
1:D:2751:LEU:HD11	1:D:2817:ILE:CD1	2.48	0.43
1:D:3434:LEU:HB3	1:D:3517:MET:HE1	2.01	0.43
1:D:4930:ALA:CB	1:C:4933:GLN:HG3	2.48	0.43
1:C:3365:LEU:HD11	1:C:3405:LEU:CD1	2.48	0.43
1:C:4901:ILE:HG13	1:C:4913:ARG:NH2	2.34	0.43
1:A:345:LEU:HD21	1:A:389:PHE:CZ	2.54	0.43
1:A:380:GLN:OE1	1:A:380:GLN:N	2.47	0.43
1:A:4901:ILE:HG13	1:A:4913:ARG:NH2	2.34	0.43
2:H:38:SER:HB3	2:H:41:ASP:OD1	2.19	0.43
2:G:38:SER:HB3	2:G:41:ASP:OD1	2.19	0.43
2:F:38:SER:HB3	2:F:41:ASP:OD1	2.19	0.43
1:B:866:HIS:O	1:B:869:ARG:HG2	2.18	0.43
1:B:1931:LEU:HD13	1:B:1935:VAL:CG1	2.47	0.43
1:B:4205:TRP:CZ2	1:B:4214:LYS:HD3	2.54	0.43
1:D:234:SER:OG	1:D:240:ASP:OD2	2.29	0.43
1:D:1115:LEU:HD12	1:D:1193:SER:HB3	2.00	0.43
1:D:1241:SER:HA	1:D:1603:VAL:HG22	2.01	0.43
1:D:1436:SER:OG	1:D:1565:GLU:HB2	2.18	0.43
1:D:2323:TRP:CZ3	1:D:2325:PRO:HG3	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:2478:THR:HG22	1:D:2479:LEU:N	2.33	0.43
1:D:2534:ALA:CB	1:D:2588:ARG:HE	2.24	0.43
1:D:4024:VAL:HG11	1:D:4142:ASN:HB3	2.00	0.43
1:D:4973:HIS:O	1:D:4977:THR:HG23	2.19	0.43
1:C:1978:ALA:HB1	1:C:1982:ARG:NH1	2.26	0.43
1:C:2751:LEU:HD11	1:C:2817:ILE:CD1	2.48	0.43
1:C:3087:ILE:H	1:C:3087:ILE:HD12	1.84	0.43
1:C:3817:LEU:HD21	1:C:3898:ASP:HB3	1.99	0.43
1:C:3951:PHE:CE1	1:C:3999:MET:HE1	2.53	0.43
1:A:336:PRO:HA	1:A:337:PRO:HD3	1.87	0.43
1:A:647:ASN:ND2	1:A:820:ARG:O	2.44	0.43
1:A:2226:PRO:O	1:A:2267:MET:CE	2.65	0.43
1:A:2290:LEU:HD23	1:A:2291:GLN:N	2.34	0.43
1:A:2309:SER:OG	1:A:2310:CYS:N	2.52	0.43
1:A:2470:ILE:HG22	1:A:2525:GLY:HA3	2.01	0.43
1:B:2323:TRP:CZ3	1:B:2325:PRO:HG3	2.54	0.43
1:B:3087:ILE:HD12	1:B:3087:ILE:H	1.84	0.43
1:B:3233:PRO:CB	1:B:3238:GLU:HB2	2.49	0.43
1:B:4983:HIS:O	3:B:5301:ATP:N6	2.51	0.43
1:D:646:PRO:HB2	1:D:648:ILE:CD1	2.48	0.43
1:D:2902:HIS:CE1	1:D:2904:LEU:HB3	2.54	0.43
1:D:3365:LEU:HD11	1:D:3405:LEU:CD1	2.48	0.43
1:C:1115:LEU:HD12	1:C:1193:SER:HB3	2.00	0.43
1:C:1241:SER:HA	1:C:1603:VAL:HG22	2.01	0.43
1:C:2323:TRP:CZ3	1:C:2325:PRO:HG3	2.54	0.43
1:C:2946:LEU:O	1:C:2946:LEU:CD2	2.62	0.43
1:C:3233:PRO:CB	1:C:3238:GLU:HB2	2.49	0.43
1:C:3289:PRO:O	1:C:3290:GLU:HB2	2.18	0.43
1:C:4253:GLU:OE1	1:C:4557:ARG:NE	2.52	0.43
1:A:559:GLY:O	1:A:563:VAL:HG23	2.18	0.43
1:A:869:ARG:NH2	1:A:941:MET:HE1	2.33	0.43
1:A:1978:ALA:O	1:A:1982:ARG:CZ	2.67	0.43
1:A:1989:ALA:O	1:A:1992:ALA:HB3	2.19	0.43
1:A:2902:HIS:CE1	1:A:2904:LEU:HB3	2.54	0.43
1:A:3450:ASN:HA	1:A:3453:ARG:HG2	2.01	0.43
1:A:3527:PRO:HD2	1:A:3573:MET:SD	2.57	0.43
1:A:3817:LEU:HD21	1:A:3898:ASP:HB3	1.99	0.43
1:A:4983:HIS:O	3:A:5301:ATP:N6	2.51	0.43
1:B:2290:LEU:HD23	1:B:2291:GLN:N	2.34	0.43
1:B:2675:THR:CG2	1:B:2706:ILE:HG23	2.49	0.43
1:B:2902:HIS:CE1	1:B:2904:LEU:HB3	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:4901:ILE:HG13	1:B:4913:ARG:NH2	2.34	0.43
1:D:1126:GLY:O	1:D:1142:PRO:HA	2.18	0.43
1:D:2675:THR:CG2	1:D:2706:ILE:HG23	2.49	0.43
1:D:2773:ASN:OD1	1:D:2773:ASN:C	2.57	0.43
1:D:3087:ILE:H	1:D:3087:ILE:HD12	1.84	0.43
1:D:3546:ASP:O	1:D:3549:VAL:HG22	2.19	0.43
1:D:3969:ILE:O	1:D:3972:PRO:HA	2.19	0.43
1:C:1143:TRP:HB2	1:C:1147:ASP:HB2	2.01	0.43
1:C:1461:ASP:OD2	1:C:1468:LYS:NZ	2.49	0.43
1:C:2123:LEU:O	1:C:2127:GLN:HG2	2.18	0.43
1:C:2135:LEU:O	1:C:3658:LYS:NZ	2.52	0.43
1:C:2675:THR:CG2	1:C:2706:ILE:HG23	2.49	0.43
1:C:3371:LYS:HE2	1:C:3371:LYS:HA	2.00	0.43
1:C:3868:ARG:HH12	1:C:3870:ASN:HB3	1.82	0.43
1:A:1036:ARG:HH11	1:A:1036:ARG:HD3	1.48	0.43
1:A:2784:GLU:CD	1:A:2785:LEU:HD12	2.39	0.43
1:A:3233:PRO:CB	1:A:3238:GLU:HB2	2.49	0.43
1:A:3969:ILE:O	1:A:3972:PRO:HA	2.19	0.43
1:A:4973:HIS:O	1:A:4977:THR:HG23	2.19	0.43
1:B:345:LEU:HD21	1:B:389:PHE:CZ	2.54	0.43
1:B:364:PRO:O	1:B:365:LYS:HB3	2.19	0.43
1:B:886:ARG:HG2	1:B:904:HIS:CD2	2.54	0.43
1:B:1978:ALA:O	1:B:1982:ARG:CZ	2.67	0.43
1:D:648:ILE:HD11	1:D:821:LEU:HD11	2.00	0.43
1:D:1068:ARG:CA	1:D:1071:ARG:HH11	2.32	0.43
1:D:1143:TRP:HB2	1:D:1147:ASP:HB2	2.01	0.43
1:D:1743[B]:ARG:NH2	1:D:1967:ASP:OD2	2.52	0.43
1:D:2604:GLU:O	1:D:2608:MET:HG2	2.19	0.43
1:D:3233:PRO:CB	1:D:3238:GLU:HB2	2.49	0.43
1:D:3409:TYR:N	1:D:3410:PRO:HD2	2.33	0.43
1:D:3868:ARG:HH12	1:D:3870:ASN:HB3	1.82	0.43
1:D:4253:GLU:OE1	1:D:4557:ARG:NE	2.52	0.43
1:C:69:LEU:CD1	1:C:101:LEU:HD11	2.49	0.43
1:C:622:THR:HG23	1:C:626:LEU:HD12	2.00	0.43
1:C:866:HIS:O	1:C:869:ARG:HG2	2.18	0.43
1:C:867:LEU:HA	1:C:870:ILE:HG22	2.01	0.43
1:C:886:ARG:HG2	1:C:904:HIS:CD2	2.54	0.43
1:C:1925:GLY:O	1:C:1929:MET:HG3	2.19	0.43
1:C:2226:PRO:HA	1:C:2267:MET:HE1	2.01	0.43
1:C:2290:LEU:HD23	1:C:2291:GLN:N	2.34	0.43
1:C:2309:SER:OG	1:C:2310:CYS:N	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3104:GLU:OE2	1:C:3167:ARG:HB3	2.18	0.43
1:C:3546:ASP:O	1:C:3549:VAL:HG22	2.19	0.43
1:C:4973:HIS:O	1:C:4977:THR:HG23	2.19	0.43
1:A:1115:LEU:HD12	1:A:1193:SER:HB3	2.00	0.43
1:A:1122:TYR:CE2	1:A:1182:ILE:HD12	2.53	0.43
1:A:2135:LEU:O	1:A:3658:LYS:NZ	2.52	0.43
1:A:2226:PRO:CA	1:A:2267:MET:HE1	2.48	0.43
1:A:2478:THR:HG22	1:A:2479:LEU:N	2.33	0.43
1:A:3127:GLN:O	1:A:3130:THR:HG22	2.18	0.43
1:B:648:ILE:HD11	1:B:821:LEU:HD11	2.00	0.43
1:B:1989:ALA:O	1:B:1992:ALA:HB3	2.19	0.43
1:B:2478:THR:HG22	1:B:2479:LEU:N	2.33	0.43
1:B:2630:VAL:N	1:B:2631:PRO:CD	2.82	0.43
1:B:4253:GLU:OE1	1:B:4253:GLU:N	2.52	0.43
1:B:4973:HIS:O	1:B:4977:THR:HG23	2.19	0.43
1:D:69:LEU:CD1	1:D:101:LEU:HD11	2.49	0.43
1:D:166:GLY:O	1:C:384:MET:HE1	2.18	0.43
1:D:510:GLU:HA	1:D:513:GLU:OE2	2.19	0.43
1:D:2630:VAL:N	1:D:2631:PRO:CD	2.82	0.43
1:D:2821:TRP:CD1	1:D:2939:ARG:O	2.72	0.43
1:D:3371:LYS:HE2	1:D:3371:LYS:HA	2.00	0.43
1:D:4901:ILE:HG13	1:D:4913:ARG:NH2	2.34	0.43
1:D:4976:GLU:O	1:D:4980:LEU:HD23	2.19	0.43
1:C:1931:LEU:HD13	1:C:1935:VAL:CG1	2.47	0.43
1:C:2212:VAL:HG11	1:C:2256:TYR:CZ	2.54	0.43
1:C:2527:LEU:O	1:C:2531:ARG:HG3	2.18	0.43
1:C:2604:GLU:O	1:C:2608:MET:HG2	2.19	0.43
1:C:2668:SER:O	1:C:2669:GLU:HB2	2.18	0.43
1:C:4024:VAL:HG11	1:C:4142:ASN:HB3	2.00	0.43
1:A:510:GLU:HA	1:A:513:GLU:OE2	2.19	0.42
1:A:893:TYR:HB3	1:A:962:SER:OG	2.18	0.42
1:A:1421:ARG:O	1:A:1570:LYS:NZ	2.33	0.42
1:A:1925:GLY:O	1:A:1929:MET:HG3	2.19	0.42
1:A:2604:GLU:O	1:A:2608:MET:HG2	2.19	0.42
1:A:2668:SER:O	1:A:2669:GLU:HB2	2.18	0.42
1:A:2675:THR:CG2	1:A:2706:ILE:HG23	2.49	0.42
1:A:2770:LYS:O	1:A:2773:ASN:OD1	2.37	0.42
1:A:3456:GLN:HG3	1:A:3503:TYR:CD2	2.54	0.42
1:A:4003:LEU:HA	1:A:4009:GLN:OE1	2.19	0.42
1:A:4205:TRP:CZ2	1:A:4214:LYS:HD3	2.54	0.42
1:B:1436:SER:OG	1:B:1565:GLU:HB2	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:2212:VAL:HG11	1:B:2256:TYR:CZ	2.54	0.42
1:B:2650:ARG:HG3	1:B:2651:CYS:SG	2.59	0.42
1:B:3456:GLN:HG3	1:B:3503:TYR:CD2	2.54	0.42
1:B:3969:ILE:O	1:B:3972:PRO:HA	2.19	0.42
1:B:4749:GLU:OE1	1:B:4753:HIS:NE2	2.52	0.42
1:B:4888:TYR:HA	1:C:4918:ILE:HD11	2.01	0.42
1:D:1989:ALA:O	1:D:1992:ALA:HB3	2.19	0.42
1:D:2212:VAL:HG11	1:D:2256:TYR:CZ	2.54	0.42
1:D:2784:GLU:CD	1:D:2785:LEU:HD12	2.39	0.42
1:D:3531:ASP:O	1:D:3535:LEU:HG	2.19	0.42
1:D:4205:TRP:CZ2	1:D:4214:LYS:HD3	2.54	0.42
1:D:4832:HIS:ND1	1:D:4833:ASN:OD1	2.45	0.42
1:C:659:TYR:O	1:C:662:TRP:NE1	2.47	0.42
1:C:2650:ARG:HG3	1:C:2651:CYS:SG	2.59	0.42
1:C:3725:TYR:HA	1:C:3728:ILE:HD12	2.01	0.42
1:C:4748:LEU:O	1:C:4751:THR:OG1	2.32	0.42
1:A:648:ILE:HD11	1:A:821:LEU:HD11	2.00	0.42
1:B:257:ARG:O	1:B:284:HIS:NE2	2.43	0.42
1:B:1122:TYR:CE2	1:B:1182:ILE:HD12	2.54	0.42
1:B:1421:ARG:O	1:B:1570:LYS:NZ	2.33	0.42
1:B:1423:ASP:HB3	1:B:1426:ILE:HD12	2.01	0.42
1:B:2756:ASN:HA	1:B:2806:ARG:HH12	1.83	0.42
1:D:320:LYS:NZ	1:D:381:GLU:O	2.43	0.42
1:D:908:VAL:HG13	1:D:909:ASN:N	2.35	0.42
1:D:1419:ASP:CG	1:D:1420:ASN:H	2.21	0.42
1:D:1925:GLY:O	1:D:1929:MET:HG3	2.19	0.42
1:D:2949:SER:OG	1:D:2952:GLU:OE1	2.25	0.42
1:C:56:GLN:O	1:C:309:THR:OG1	2.29	0.42
1:C:648:ILE:HD11	1:C:821:LEU:HD11	2.00	0.42
1:C:908:VAL:HG13	1:C:909:ASN:N	2.35	0.42
1:C:1758:ARG:CZ	1:C:1759:ARG:NH1	2.82	0.42
1:C:3359:ILE:HD13	1:C:3434:LEU:HD13	1.99	0.42
1:C:4976:GLU:O	1:C:4980:LEU:HD23	2.19	0.42
1:A:364:PRO:O	1:A:365:LYS:HB3	2.19	0.42
1:A:901:LYS:HG3	1:A:903:LEU:HD12	2.01	0.42
1:A:1743[B]:ARG:NH2	1:A:1967:ASP:OD2	2.52	0.42
1:A:1758:ARG:CZ	1:A:1759:ARG:NH1	2.82	0.42
1:A:2323:TRP:CZ3	1:A:2325:PRO:HG3	2.54	0.42
1:A:2481:LYS:O	1:A:2482:ASP:OD2	2.38	0.42
1:A:2821:TRP:CD1	1:A:2939:ARG:O	2.72	0.42
1:A:3087:ILE:H	1:A:3087:ILE:HD12	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:3531:ASP:O	1:A:3535:LEU:HG	2.19	0.42
1:B:622:THR:HG23	1:B:626:LEU:HD12	2.00	0.42
1:B:2481:LYS:O	1:B:2482:ASP:OD2	2.38	0.42
1:B:3409:TYR:N	1:B:3410:PRO:HD2	2.33	0.42
1:D:622:THR:HG23	1:D:626:LEU:HD12	2.00	0.42
1:D:886:ARG:HG2	1:D:904:HIS:CD2	2.54	0.42
1:D:1978:ALA:O	1:D:1982:ARG:CZ	2.67	0.42
1:D:4749:GLU:OE1	1:D:4753:HIS:NE2	2.52	0.42
1:C:380:GLN:OE1	1:C:380:GLN:N	2.47	0.42
1:C:2000:SER:OG	1:C:2004:GLU:HB2	2.20	0.42
1:C:2481:LYS:O	1:C:2482:ASP:OD2	2.38	0.42
1:C:3321:ARG:O	1:C:3324:VAL:HG22	2.19	0.42
1:C:3716:LEU:HD21	1:C:3782:MET:HE3	2.02	0.42
1:C:4205:TRP:CZ2	1:C:4214:LYS:HD3	2.54	0.42
1:C:4214:LYS:O	1:C:4218:ILE:HG12	2.20	0.42
1:C:4852:THR:HG23	1:C:4882:CYS:HB3	1.99	0.42
1:A:870:ILE:HD11	1:A:1049:TYR:CD2	2.55	0.42
1:A:1758:ARG:NH2	1:A:1759:ARG:CZ	2.83	0.42
1:A:3546:ASP:O	1:A:3549:VAL:HG22	2.19	0.42
1:A:4976:GLU:O	1:A:4980:LEU:HD23	2.19	0.42
1:B:901:LYS:HG3	1:B:903:LEU:HD12	2.02	0.42
1:B:2000:SER:OG	1:B:2004:GLU:HB2	2.19	0.42
1:B:2668:SER:O	1:B:2669:GLU:HB2	2.18	0.42
1:B:2821:TRP:CD1	1:B:2939:ARG:O	2.72	0.42
1:B:3321:ARG:O	1:B:3324:VAL:HG22	2.20	0.42
1:B:3546:ASP:O	1:B:3549:VAL:HG22	2.19	0.42
1:D:870:ILE:HD11	1:D:1049:TYR:CD2	2.55	0.42
1:D:901:LYS:HG3	1:D:903:LEU:HD12	2.01	0.42
1:D:2000:SER:OG	1:D:2004:GLU:HB2	2.20	0.42
1:D:2309:SER:OG	1:D:2310:CYS:N	2.52	0.42
1:D:3296:LEU:HB3	1:D:3297:PRO:HD3	2.02	0.42
1:D:3985:LEU:HD11	1:D:4026:MET:HE1	2.02	0.42
1:C:510:GLU:HA	1:C:513:GLU:OE2	2.19	0.42
1:C:648:ILE:HD11	1:C:821:LEU:CD1	2.50	0.42
1:C:2625:ARG:NH1	1:C:2629:ASP:OD1	2.51	0.42
1:C:2630:VAL:N	1:C:2631:PRO:CD	2.82	0.42
1:C:2755:ILE:HD13	1:C:2810:LYS:HG2	2.00	0.42
1:C:2902:HIS:CE1	1:C:2904:LEU:HB3	2.54	0.42
1:A:2773:ASN:OD1	1:A:2773:ASN:C	2.57	0.42
1:A:3371:LYS:HE2	1:A:3371:LYS:HA	2.00	0.42
1:B:417:GLY:O	1:B:421:PHE:HD1	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1758:ARG:HG2	1:B:1759:ARG:N	2.35	0.42
1:B:3107:VAL:HG12	1:B:3175:LEU:HD21	2.02	0.42
1:B:3531:ASP:O	1:B:3535:LEU:HG	2.19	0.42
1:B:4573:ILE:HG23	1:B:4643:LEU:HD11	2.02	0.42
1:D:559:GLY:O	1:D:563:VAL:HG23	2.18	0.42
1:D:1141:ARG:CB	1:C:3477:LYS:HZ3	2.32	0.42
1:D:2290:LEU:HD23	1:D:2291:GLN:N	2.34	0.42
1:D:3321:ARG:O	1:D:3324:VAL:HG22	2.19	0.42
1:D:3456:GLN:HG3	1:D:3503:TYR:CD2	2.54	0.42
1:D:3985:LEU:HD11	1:D:4026:MET:CE	2.50	0.42
1:C:901:LYS:HG3	1:C:903:LEU:HD12	2.01	0.42
1:C:1068:ARG:CA	1:C:1071:ARG:HH11	2.32	0.42
1:C:1758:ARG:NH2	1:C:1759:ARG:CZ	2.83	0.42
1:C:2440:MET:O	1:C:2444:GLN:OE1	2.38	0.42
1:C:2478:THR:HG22	1:C:2479:LEU:N	2.33	0.42
1:C:4749:GLU:OE1	1:C:4753:HIS:NE2	2.52	0.42
1:C:4983:HIS:O	3:C:5301:ATP:N6	2.51	0.42
1:A:417:GLY:O	1:A:421:PHE:HD1	2.01	0.42
1:A:648:ILE:HD11	1:A:821:LEU:CD1	2.50	0.42
1:A:898:ASP:HB2	1:A:901:LYS:HB2	2.02	0.42
1:A:1032:LYS:C	1:A:1036:ARG:HH12	2.18	0.42
1:A:1068:ARG:CA	1:A:1071:ARG:HH11	2.32	0.42
1:A:1171:SER:OG	1:A:1173:SER:OG	2.33	0.42
1:A:2650:ARG:HG3	1:A:2651:CYS:SG	2.59	0.42
1:A:3294:PRO:HB2	1:A:3297:PRO:HD2	2.02	0.42
1:A:3874:VAL:O	1:A:3953:LYS:NZ	2.53	0.42
1:A:4214:LYS:O	1:A:4218:ILE:HG12	2.20	0.42
1:B:69:LEU:CD1	1:B:101:LEU:HD11	2.49	0.42
1:B:226:HIS:ND1	1:B:226:HIS:O	2.53	0.42
1:B:510:GLU:HA	1:B:513:GLU:OE2	2.19	0.42
1:B:1143:TRP:HB2	1:B:1147:ASP:HB2	2.01	0.42
1:B:1241:SER:HA	1:B:1603:VAL:HG22	2.01	0.42
1:B:2226:PRO:HA	1:B:2267:MET:HE1	2.02	0.42
1:B:2625:ARG:NH1	1:B:2629:ASP:OD1	2.51	0.42
1:B:3391:GLU:HG2	1:B:3395:ARG:HH21	1.83	0.42
1:B:4710:SER:O	1:B:4713:SER:OG	2.25	0.42
1:D:659:TYR:O	1:D:662:TRP:NE1	2.47	0.42
1:D:898:ASP:HB2	1:D:901:LYS:HB2	2.02	0.42
1:D:1758:ARG:CZ	1:D:1759:ARG:NH1	2.82	0.42
1:D:2770:LYS:O	1:D:2773:ASN:OD1	2.37	0.42
1:D:2967:MET:HG3	1:D:3042:LEU:HD12	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:870:ILE:HD11	1:C:1049:TYR:CD2	2.55	0.42
1:C:1515:VAL:O	1:C:1531:ALA:O	2.38	0.42
1:C:2770:LYS:O	1:C:2773:ASN:OD1	2.37	0.42
1:C:3456:GLN:HG3	1:C:3503:TYR:CD2	2.54	0.42
1:C:4161:ARG:HA	1:C:4164:LEU:HD12	2.01	0.42
1:A:867:LEU:HA	1:A:870:ILE:HG22	2.01	0.42
1:A:2211:MET:SD	1:A:2272:PRO:HG3	2.60	0.42
1:A:3321:ARG:O	1:A:3324:VAL:HG22	2.19	0.42
1:A:4749:GLU:OE1	1:A:4753:HIS:NE2	2.52	0.42
1:B:867:LEU:HA	1:B:870:ILE:HG22	2.01	0.42
1:B:870:ILE:HD11	1:B:1049:TYR:CD2	2.55	0.42
1:B:2211:MET:SD	1:B:2272:PRO:HG3	2.60	0.42
1:B:2495:VAL:HG22	1:B:2498:HIS:CE1	2.55	0.42
1:B:2770:LYS:O	1:B:2773:ASN:OD1	2.37	0.42
1:B:3296:LEU:HB3	1:B:3297:PRO:HD3	2.02	0.42
1:B:3450:ASN:HA	1:B:3453:ARG:HG2	2.01	0.42
1:B:3686:GLU:O	1:B:3687:GLU:HB3	2.20	0.42
1:B:3951:PHE:CE1	1:B:3999:MET:HE1	2.54	0.42
1:B:4923:PHE:HA	1:B:4927:ILE:HD12	2.01	0.42
1:D:226:HIS:ND1	1:D:226:HIS:O	2.53	0.42
1:D:1754:GLY:O	1:D:1758:ARG:HB2	2.20	0.42
1:D:2969:ILE:O	1:D:2973:PHE:CD1	2.73	0.42
1:C:1978:ALA:O	1:C:1982:ARG:CZ	2.67	0.42
1:C:2969:ILE:O	1:C:2973:PHE:CD1	2.73	0.42
1:C:3296:LEU:HB3	1:C:3297:PRO:HD3	2.02	0.42
1:C:4573:ILE:HG23	1:C:4643:LEU:HD11	2.02	0.42
1:A:110:ARG:HH21	1:A:115:ARG:HD2	1.85	0.42
1:A:1241:SER:HA	1:A:1603:VAL:HG22	2.01	0.42
1:A:2495:VAL:HG22	1:A:2498:HIS:CE1	2.55	0.42
1:A:2630:VAL:N	1:A:2631:PRO:CD	2.82	0.42
1:A:2701:PRO:C	1:A:2702:CYS:SG	2.98	0.42
1:A:2777:TYR:CD1	1:A:2791:LEU:HD12	2.43	0.42
1:A:3107:VAL:HG12	1:A:3175:LEU:HD21	2.02	0.42
1:A:3686:GLU:O	1:A:3687:GLU:HB3	2.20	0.42
1:A:3725:TYR:HA	1:A:3728:ILE:HD12	2.01	0.42
1:A:3985:LEU:HD11	1:A:4026:MET:CE	2.50	0.42
1:B:574:VAL:O	1:B:578:ILE:HG12	2.18	0.42
1:B:1515:VAL:O	1:B:1531:ALA:O	2.38	0.42
1:B:2309:SER:OG	1:B:2310:CYS:N	2.52	0.42
1:B:2784:GLU:CD	1:B:2785:LEU:HD12	2.39	0.42
1:B:2969:ILE:O	1:B:2973:PHE:CD1	2.73	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:3268:HIS:ND1	1:B:3272:ILE:HD12	2.35	0.42
1:B:3725:TYR:O	1:B:3729:MET:HG3	2.20	0.42
1:B:3999:MET:HE3	1:B:3999:MET:HB3	1.75	0.42
1:B:4704:LEU:O	1:B:4774:LYS:NZ	2.52	0.42
1:D:110:ARG:HH21	1:D:115:ARG:HD2	1.85	0.42
1:D:345:LEU:HD21	1:D:389:PHE:CZ	2.54	0.42
1:D:2481:LYS:O	1:D:2482:ASP:OD2	2.38	0.42
1:C:898:ASP:HB2	1:C:901:LYS:HB2	2.02	0.42
1:C:2773:ASN:OD1	1:C:2773:ASN:C	2.57	0.42
1:C:2784:GLU:CD	1:C:2785:LEU:HD12	2.39	0.42
1:C:2801:ASP:HA	1:C:2804:ILE:HG12	2.02	0.42
1:A:69:LEU:CD1	1:A:101:LEU:HD11	2.49	0.42
1:A:622:THR:HG23	1:A:626:LEU:HD12	2.00	0.42
1:A:1423:ASP:HB3	1:A:1426:ILE:HD12	2.01	0.42
1:A:1758:ARG:HG2	1:A:1759:ARG:N	2.35	0.42
1:A:2562:ILE:HG22	1:A:2610:LEU:HD21	2.02	0.42
1:A:3725:TYR:O	1:A:3729:MET:HG3	2.20	0.42
1:A:4573:ILE:HG23	1:A:4643:LEU:HD11	2.02	0.42
1:A:4923:PHE:HA	1:A:4927:ILE:HD12	2.01	0.42
1:B:98:HIS:N	1:B:99:ARG:HH11	2.18	0.42
1:B:380:GLN:OE1	1:B:380:GLN:N	2.47	0.42
1:B:2799:GLU:O	1:B:2803:GLU:OE1	2.38	0.42
1:B:4003:LEU:HA	1:B:4009:GLN:OE1	2.19	0.42
1:B:4545:GLU:OE2	1:B:4548:ARG:NH2	2.39	0.42
1:B:4976:GLU:O	1:B:4980:LEU:HD23	2.19	0.42
1:D:866:HIS:HA	1:D:869:ARG:HG2	2.02	0.42
1:D:1758:ARG:HG2	1:D:1759:ARG:N	2.35	0.42
1:D:2650:ARG:HG3	1:D:2651:CYS:SG	2.59	0.42
1:D:2801:ASP:HA	1:D:2804:ILE:HG12	2.02	0.42
1:D:3905:THR:HA	1:D:3912:THR:OG1	2.20	0.42
1:D:4086:GLY:O	1:D:4125:PHE:CE2	2.73	0.42
1:D:4573:ILE:HG23	1:D:4643:LEU:HD11	2.02	0.42
1:D:4949:GLN:HA	1:D:4952:GLU:HG3	2.02	0.42
1:C:438:ILE:HG23	1:C:518:ILE:HD11	2.02	0.42
1:C:887:ILE:HG13	1:C:907:LEU:CD1	2.50	0.42
1:C:1423:ASP:HB3	1:C:1426:ILE:HD12	2.01	0.42
1:C:1758:ARG:HG2	1:C:1759:ARG:N	2.35	0.42
1:C:2430:ILE:HG22	1:C:2505:PHE:HB2	2.02	0.42
1:C:2821:TRP:CD1	1:C:2939:ARG:O	2.72	0.42
1:C:3107:VAL:HG12	1:C:3175:LEU:HD21	2.02	0.42
1:C:3531:ASP:O	1:C:3535:LEU:HG	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3686:GLU:O	1:C:3687:GLU:HB3	2.20	0.42
1:C:3822:ASP:OD1	1:C:3823:LYS:N	2.53	0.42
1:A:886:ARG:HG2	1:A:904:HIS:CD2	2.54	0.42
1:A:1515:VAL:O	1:A:1531:ALA:O	2.38	0.42
1:A:2376:LEU:HD11	1:A:2426:TYR:HB3	2.02	0.42
1:A:2967:MET:HG3	1:A:3042:LEU:HD12	2.02	0.42
1:A:3296:LEU:HB3	1:A:3297:PRO:HD3	2.02	0.42
2:G:24:VAL:HG21	2:G:59:TRP:HZ3	1.85	0.42
1:B:877:ASN:HD21	1:B:970:LEU:HD13	1.85	0.42
1:B:1758:ARG:CZ	1:B:1759:ARG:NH1	2.82	0.42
1:B:1978:ALA:HB1	1:B:1982:ARG:NH1	2.26	0.42
1:B:2440:MET:O	1:B:2444:GLN:OE1	2.38	0.42
1:B:3104:GLU:OE2	1:B:3167:ARG:HB3	2.18	0.42
1:B:3207:GLU:OE1	1:B:3280:TYR:OH	2.31	0.42
1:B:3395:ARG:HB3	1:B:3454:GLU:OE1	2.20	0.42
1:B:4161:ARG:HA	1:B:4164:LEU:HD12	2.01	0.42
1:D:438:ILE:HG23	1:D:518:ILE:HD11	2.02	0.42
1:D:516:LYS:O	1:D:519:VAL:HG22	2.20	0.42
1:D:647:ASN:ND2	1:D:820:ARG:O	2.44	0.42
1:D:3725:TYR:HA	1:D:3728:ILE:HD12	2.01	0.42
1:D:4918:ILE:HD11	1:C:4888:TYR:HA	2.01	0.42
1:C:226:HIS:ND1	1:C:226:HIS:O	2.53	0.42
1:C:364:PRO:O	1:C:365:LYS:HB3	2.19	0.42
1:C:1694:LEU:O	1:C:1698:LEU:HG	2.20	0.42
1:C:2470:ILE:HG22	1:C:2525:GLY:HA3	2.01	0.42
1:C:3294:PRO:HB2	1:C:3297:PRO:HD2	2.02	0.42
1:C:3969:ILE:O	1:C:3972:PRO:HA	2.19	0.42
1:A:2799:GLU:O	1:A:2803:GLU:OE1	2.38	0.41
1:A:3268:HIS:ND1	1:A:3272:ILE:HD12	2.35	0.41
1:A:4086:GLY:O	1:A:4125:PHE:CE2	2.73	0.41
1:B:1925:GLY:O	1:B:1929:MET:HG3	2.19	0.41
1:B:3249:LEU:HD11	1:B:3273:THR:HG21	2.01	0.41
1:B:3985:LEU:HD11	1:B:4026:MET:CE	2.50	0.41
1:D:1758:ARG:NH2	1:D:1759:ARG:CZ	2.83	0.41
1:D:2044:ILE:HD12	1:D:2044:ILE:H	1.85	0.41
1:D:2434:GLY:O	1:D:2508:ARG:NE	2.36	0.41
1:D:3235:SER:OG	1:D:3237:GLU:OE1	2.38	0.41
1:D:4869:GLU:OE2	1:D:4869:GLU:N	2.53	0.41
1:C:516:LYS:O	1:C:519:VAL:HG22	2.20	0.41
1:C:2562:ILE:HG22	1:C:2610:LEU:HD21	2.02	0.41
1:C:2701:PRO:C	1:C:2702:CYS:SG	2.98	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:3235:SER:OG	1:C:3237:GLU:OE1	2.38	0.41
1:C:3395:ARG:HB3	1:C:3454:GLU:OE1	2.20	0.41
1:C:4086:GLY:O	1:C:4125:PHE:CE2	2.73	0.41
1:A:98:HIS:N	1:A:99:ARG:HH11	2.18	0.41
1:A:294:THR:O	1:A:298:GLY:N	2.50	0.41
1:A:1669:LEU:O	1:A:1673:VAL:HG22	2.20	0.41
1:A:3999:MET:HB3	1:A:3999:MET:HE3	1.77	0.41
1:B:438:ILE:HG23	1:B:518:ILE:HD11	2.02	0.41
1:B:887:ILE:HG13	1:B:907:LEU:CD1	2.50	0.41
1:B:1157:GLU:OE1	1:B:1157:GLU:N	2.53	0.41
1:B:2773:ASN:OD1	1:B:2773:ASN:C	2.57	0.41
1:B:3725:TYR:HA	1:B:3728:ILE:HD12	2.01	0.41
1:D:1694:LEU:O	1:D:1698:LEU:HG	2.20	0.41
1:D:2470:ILE:HG22	1:D:2525:GLY:HA3	2.01	0.41
1:D:3450:ASN:HA	1:D:3453:ARG:HG2	2.01	0.41
1:D:3478:MET:SD	1:D:3478:MET:O	2.78	0.41
1:D:3686:GLU:O	1:D:3687:GLU:HB3	2.20	0.41
1:C:110:ARG:HH21	1:C:115:ARG:HD2	1.85	0.41
1:C:877:ASN:HD21	1:C:970:LEU:HD13	1.85	0.41
1:C:1157:GLU:OE1	1:C:1157:GLU:N	2.53	0.41
1:C:2756:ASN:HA	1:C:2806:ARG:HH12	1.83	0.41
1:C:4666:VAL:N	1:C:4667:PRO:CD	2.83	0.41
1:A:1204:LEU:HD12	1:A:1234:VAL:HB	2.03	0.41
1:A:1694:LEU:O	1:A:1698:LEU:HG	2.20	0.41
1:A:1742:THR:HA	1:A:1745:ILE:HD12	2.02	0.41
1:A:1996:ARG:NE	1:A:1999:ARG:NH1	2.61	0.41
1:A:5024:ALA:HB3	1:D:4972:PRO:HB3	2.02	0.41
1:B:14:LEU:CB	1:B:101:LEU:HD12	2.50	0.41
1:B:648:ILE:HD11	1:B:821:LEU:CD1	2.50	0.41
1:B:1758:ARG:NH2	1:B:1759:ARG:CZ	2.83	0.41
1:B:2233:CYS:HA	1:B:2237:CYS:HB2	2.03	0.41
1:B:3294:PRO:HB2	1:B:3297:PRO:HD2	2.02	0.41
1:B:3891:LEU:O	1:B:3899:PHE:CD2	2.71	0.41
1:D:39:ALA:O	1:D:111:HIS:CE1	2.73	0.41
1:D:224:HIS:HB2	1:D:229:GLU:OE2	2.21	0.41
1:D:364:PRO:O	1:D:365:LYS:HB3	2.19	0.41
1:D:1515:VAL:O	1:D:1531:ALA:O	2.38	0.41
1:D:2138:LEU:O	1:D:2141:ALA:N	2.45	0.41
1:D:2376:LEU:HD11	1:D:2426:TYR:HB3	2.02	0.41
1:D:2440:MET:O	1:D:2444:GLN:OE1	2.38	0.41
1:D:2495:VAL:HG22	1:D:2498:HIS:CE1	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3249:LEU:HD11	1:D:3273:THR:HG21	2.01	0.41
1:D:4161:ARG:HA	1:D:4164:LEU:HD12	2.02	0.41
1:C:14:LEU:CB	1:C:101:LEU:HD12	2.51	0.41
1:C:866:HIS:HA	1:C:869:ARG:HG2	2.02	0.41
1:C:1743[B]:ARG:NH2	1:C:1967:ASP:OD2	2.52	0.41
1:C:1758:ARG:NH2	1:C:1759:ARG:NH1	2.69	0.41
1:C:2211:MET:SD	1:C:2272:PRO:HG3	2.60	0.41
1:C:2233:CYS:HA	1:C:2237:CYS:HB2	2.02	0.41
1:C:3874:VAL:O	1:C:3953:LYS:NZ	2.53	0.41
1:A:908:VAL:HG13	1:A:909:ASN:N	2.34	0.41
1:A:1143:TRP:HB2	1:A:1147:ASP:HB2	2.01	0.41
1:A:2233:CYS:HA	1:A:2237:CYS:HB2	2.03	0.41
1:A:2614:ILE:CG2	1:A:2618:MET:HB2	2.51	0.41
1:A:3822:ASP:OD1	1:A:3823:LYS:N	2.53	0.41
1:A:3905:THR:HA	1:A:3912:THR:OG1	2.20	0.41
1:A:4666:VAL:N	1:A:4667:PRO:CD	2.83	0.41
1:A:4949:GLN:HA	1:A:4952:GLU:HG3	2.02	0.41
1:B:224:HIS:HB2	1:B:229:GLU:OE2	2.21	0.41
1:B:866:HIS:HA	1:B:869:ARG:HG2	2.02	0.41
1:B:908:VAL:HG13	1:B:909:ASN:N	2.35	0.41
1:B:1068:ARG:CA	1:B:1071:ARG:HH11	2.32	0.41
1:B:1694:LEU:O	1:B:1698:LEU:HG	2.20	0.41
1:B:1754:GLY:O	1:B:1758:ARG:HB2	2.20	0.41
1:B:2282:ASP:HA	1:B:2341:VAL:HG13	2.02	0.41
1:B:3822:ASP:OD1	1:B:3823:LYS:N	2.53	0.41
1:B:3874:VAL:O	1:B:3953:LYS:NZ	2.53	0.41
1:D:1204:LEU:HD12	1:D:1234:VAL:HB	2.03	0.41
1:D:2211:MET:SD	1:D:2272:PRO:HG3	2.60	0.41
1:D:3268:HIS:ND1	1:D:3272:ILE:HD12	2.35	0.41
1:D:3391:GLU:HG2	1:D:3395:ARG:HH21	1.83	0.41
1:D:4003:LEU:HA	1:D:4009:GLN:OE1	2.20	0.41
1:D:4214:LYS:O	1:D:4218:ILE:HG12	2.20	0.41
1:D:4666:VAL:N	1:D:4667:PRO:CD	2.83	0.41
1:D:4748:LEU:O	1:D:4751:THR:OG1	2.32	0.41
1:C:98:HIS:N	1:C:99:ARG:HH11	2.18	0.41
1:C:2782:ASP:OD1	1:C:2782:ASP:N	2.54	0.41
1:C:3018:LEU:HD21	1:C:3075:LEU:O	2.20	0.41
1:C:3478:MET:O	1:C:3478:MET:SD	2.78	0.41
1:C:3504:SER:HB2	1:C:3507:THR:HG23	2.03	0.41
1:C:3985:LEU:HD11	1:C:4026:MET:CE	2.50	0.41
1:C:4000:MET:HE1	1:C:4061:PHE:HB2	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:39:ALA:O	1:A:111:HIS:CE1	2.73	0.41
1:A:438:ILE:HG23	1:A:518:ILE:HD11	2.02	0.41
1:A:2441:HIS:CE1	1:A:2442:LEU:HG	2.56	0.41
1:A:2625:ARG:NH1	1:A:2629:ASP:OD1	2.51	0.41
1:B:594:GLY:HA2	1:B:1594:ARG:HD2	2.03	0.41
1:B:1425:GLU:OE2	1:B:1425:GLU:N	2.53	0.41
1:B:1669:LEU:O	1:B:1673:VAL:HG22	2.21	0.41
1:B:2470:ILE:HG22	1:B:2525:GLY:HA3	2.01	0.41
1:B:2701:PRO:C	1:B:2702:CYS:SG	2.98	0.41
1:B:3177:THR:O	1:B:3179:LYS:NZ	2.36	0.41
1:B:3478:MET:SD	1:B:3478:MET:O	2.78	0.41
1:B:4214:LYS:O	1:B:4218:ILE:HG12	2.20	0.41
1:B:4705:VAL:HG13	1:B:4711:PHE:HE1	1.85	0.41
1:D:98:HIS:N	1:D:99:ARG:HH11	2.18	0.41
1:D:648:ILE:HD11	1:D:821:LEU:CD1	2.50	0.41
1:D:1036:ARG:HH11	1:D:1036:ARG:HD3	1.48	0.41
1:D:2614:ILE:CG2	1:D:2618:MET:HB2	2.51	0.41
1:D:2782:ASP:N	1:D:2782:ASP:OD1	2.54	0.41
1:D:3725:TYR:O	1:D:3729:MET:HG3	2.20	0.41
1:D:3822:ASP:OD1	1:D:3823:LYS:N	2.53	0.41
1:C:1036:ARG:HH11	1:C:1036:ARG:HD3	1.48	0.41
1:C:3508:SER:O	1:C:3511:VAL:HG22	2.21	0.41
1:C:3716:LEU:HD11	1:C:3782:MET:HE1	2.03	0.41
1:C:4923:PHE:HA	1:C:4927:ILE:HD12	2.01	0.41
1:A:887:ILE:HG13	1:A:907:LEU:CD1	2.50	0.41
1:A:1754:GLY:O	1:A:1758:ARG:HB2	2.20	0.41
1:A:2801:ASP:HA	1:A:2804:ILE:HG12	2.02	0.41
1:A:3445:TRP:O	1:A:3452:LYS:NZ	2.43	0.41
1:A:3478:MET:O	1:A:3478:MET:SD	2.78	0.41
2:F:24:VAL:HG21	2:F:59:TRP:HZ3	1.85	0.41
1:B:39:ALA:O	1:B:111:HIS:CE1	2.73	0.41
1:B:1204:LEU:HD12	1:B:1234:VAL:HB	2.03	0.41
1:B:2441:HIS:CE1	1:B:2442:LEU:HG	2.56	0.41
1:B:2562:ILE:HG22	1:B:2610:LEU:HD21	2.01	0.41
1:B:4000:MET:HG2	1:B:4061:PHE:CD2	2.56	0.41
1:B:4086:GLY:O	1:B:4125:PHE:CE2	2.73	0.41
1:B:4666:VAL:N	1:B:4667:PRO:CD	2.83	0.41
1:D:2135:LEU:O	1:D:3658:LYS:NZ	2.51	0.41
1:D:2946:LEU:O	1:D:2946:LEU:CD2	2.62	0.41
1:D:3107:VAL:HG12	1:D:3175:LEU:HD21	2.02	0.41
1:D:3395:ARG:HB3	1:D:3454:GLU:OE1	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:4205:TRP:CH2	1:D:4214:LYS:HD3	2.56	0.41
1:C:39:ALA:O	1:C:111:HIS:CE1	2.73	0.41
1:C:594:GLY:HA2	1:C:1594:ARG:HD2	2.03	0.41
1:C:708:GLY:HA3	1:C:722:TRP:HB3	2.02	0.41
1:C:1425:GLU:OE2	1:C:1425:GLU:N	2.53	0.41
1:C:2967:MET:HG3	1:C:3042:LEU:HD12	2.02	0.41
1:C:3450:ASN:HA	1:C:3453:ARG:HG2	2.01	0.41
1:C:4000:MET:HG2	1:C:4061:PHE:CD2	2.56	0.41
1:C:4253:GLU:OE1	1:C:4253:GLU:N	2.52	0.41
1:A:594:GLY:HA2	1:A:1594:ARG:HD2	2.03	0.41
1:A:2000:SER:OG	1:A:2004:GLU:HB2	2.19	0.41
1:A:2969:ILE:O	1:A:2973:PHE:CD1	2.73	0.41
1:A:2978:GLU:HA	1:A:2981:VAL:HG12	2.02	0.41
1:A:3249:LEU:HD11	1:A:3273:THR:HG21	2.01	0.41
1:A:3395:ARG:HB3	1:A:3454:GLU:OE1	2.20	0.41
1:A:4161:ARG:HA	1:A:4164:LEU:HD12	2.01	0.41
2:H:24:VAL:HG21	2:H:59:TRP:HZ3	1.85	0.41
1:B:266:ARG:NH2	1:B:330:ASP:OD1	2.52	0.41
1:B:1063:VAL:HG13	1:B:1064:GLU:N	2.36	0.41
1:B:1742:THR:HA	1:B:1745:ILE:HD12	2.02	0.41
1:B:1743[B]:ARG:NH2	1:B:1967:ASP:OD2	2.52	0.41
1:B:2782:ASP:OD1	1:B:2782:ASP:N	2.54	0.41
1:B:3111:ARG:NH2	1:B:3175:LEU:N	2.69	0.41
1:D:14:LEU:CB	1:D:101:LEU:HD12	2.50	0.41
1:D:1032:LYS:C	1:D:1036:ARG:HH12	2.18	0.41
1:D:1157:GLU:OE1	1:D:1157:GLU:N	2.53	0.41
1:D:1758:ARG:NH2	1:D:1759:ARG:NH1	2.68	0.41
1:D:2233:CYS:HA	1:D:2237:CYS:HB2	2.03	0.41
1:D:2562:ILE:HG22	1:D:2610:LEU:HD21	2.02	0.41
1:D:3018:LEU:HD21	1:D:3075:LEU:O	2.20	0.41
1:D:3111:ARG:NH2	1:D:3175:LEU:N	2.69	0.41
1:D:4942:GLU:O	1:D:4946:GLN:HG3	2.21	0.41
1:C:234:SER:OG	1:C:240:ASP:OD2	2.29	0.41
1:C:336:PRO:HA	1:C:337:PRO:HD3	1.87	0.41
1:C:1705:GLY:HA3	1:C:1836:PHE:CG	2.56	0.41
1:C:2226:PRO:CA	1:C:2267:MET:HE1	2.51	0.41
1:C:2441:HIS:CE1	1:C:2442:LEU:HG	2.56	0.41
1:C:2495:VAL:HG22	1:C:2498:HIS:CE1	2.55	0.41
1:C:4000:MET:HE1	1:C:4058:ILE:HA	2.02	0.41
1:C:4003:LEU:HA	1:C:4009:GLN:OE1	2.19	0.41
1:C:4705:VAL:HG13	1:C:4711:PHE:HE1	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:226:HIS:ND1	1:A:226:HIS:O	2.53	0.41
1:A:439:GLU:HG2	1:A:440:ALA:N	2.36	0.41
1:A:516:LYS:O	1:A:519:VAL:HG22	2.20	0.41
1:A:877:ASN:HD21	1:A:970:LEU:HD13	1.84	0.41
1:A:1705:GLY:HA3	1:A:1836:PHE:CG	2.56	0.41
1:A:1758:ARG:NH2	1:A:1759:ARG:NH1	2.68	0.41
1:A:2150:GLU:HG2	1:A:2151:ASP:N	2.36	0.41
1:A:3235:SER:OG	1:A:3237:GLU:OE1	2.38	0.41
1:A:4888:TYR:HA	1:B:4918:ILE:HD11	2.03	0.41
1:B:591:ASP:HA	1:B:631:LEU:HD21	2.03	0.41
1:B:898:ASP:HB2	1:B:901:LYS:HB2	2.02	0.41
1:B:924:MET:O	1:B:928:THR:HG23	2.21	0.41
1:B:1423:ASP:O	1:B:1427:ILE:HG22	2.21	0.41
1:B:2044:ILE:HD12	1:B:2044:ILE:H	1.85	0.41
1:B:2967:MET:HG3	1:B:3042:LEU:HD12	2.02	0.41
1:B:3504:SER:HB2	1:B:3507:THR:HG23	2.03	0.41
1:B:3508:SER:O	1:B:3511:VAL:HG22	2.21	0.41
1:B:3879:GLU:HG2	1:B:3880:PHE:N	2.36	0.41
1:B:4061:PHE:CE2	1:B:4065:PHE:HE2	2.39	0.41
1:D:1063:VAL:HG13	1:D:1064:GLU:N	2.36	0.41
1:D:1931:LEU:HD13	1:D:1935:VAL:CG1	2.47	0.41
1:D:2701:PRO:C	1:D:2702:CYS:SG	2.98	0.41
1:D:2760:GLU:OE2	1:D:2794:TYR:CE1	2.74	0.41
1:D:3508:SER:O	1:D:3511:VAL:HG22	2.21	0.41
1:D:3959:LYS:HG3	1:D:4022:ASP:OD2	2.21	0.41
1:D:4923:PHE:HA	1:D:4927:ILE:HD12	2.01	0.41
1:C:591:ASP:HA	1:C:631:LEU:HD21	2.03	0.41
1:C:924:MET:O	1:C:928:THR:HG23	2.21	0.41
1:C:2777:TYR:CD1	1:C:2791:LEU:HD12	2.43	0.41
1:C:3268:HIS:ND1	1:C:3272:ILE:HD12	2.35	0.41
1:A:224:HIS:HB2	1:A:229:GLU:OE2	2.21	0.41
1:A:371:VAL:HG13	1:A:371:VAL:O	2.20	0.41
1:A:866:HIS:HA	1:A:869:ARG:HG2	2.02	0.41
1:A:950:LEU:HD12	1:A:950:LEU:O	2.21	0.41
1:A:1063:VAL:HG13	1:A:1064:GLU:N	2.36	0.41
1:A:1423:ASP:O	1:A:1427:ILE:HG22	2.21	0.41
1:A:2430:ILE:HG22	1:A:2505:PHE:HB2	2.02	0.41
1:A:2440:MET:O	1:A:2444:GLN:OE1	2.38	0.41
1:A:3077:ALA:O	1:A:3081:MET:HG2	2.21	0.41
1:A:3281:LEU:HB2	1:A:3282:PRO:HD3	2.03	0.41
1:A:3879:GLU:HG2	1:A:3880:PHE:N	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:4000:MET:HG2	1:A:4061:PHE:CD2	2.56	0.41
2:E:24:VAL:HG21	2:E:59:TRP:HZ3	1.85	0.41
1:B:110:ARG:HH21	1:B:115:ARG:HD2	1.85	0.41
1:B:361:ALA:O	1:B:362:PRO:C	2.59	0.41
1:B:371:VAL:HG13	1:B:371:VAL:O	2.20	0.41
1:B:439:GLU:HG2	1:B:440:ALA:N	2.36	0.41
1:B:516:LYS:O	1:B:519:VAL:HG22	2.20	0.41
1:B:647:ASN:ND2	1:B:820:ARG:O	2.44	0.41
1:B:842:PRO:O	1:B:1196:PRO:HA	2.21	0.41
1:B:950:LEU:O	1:B:950:LEU:HD12	2.21	0.41
1:B:1758:ARG:NH2	1:B:1759:ARG:NH1	2.68	0.41
1:B:2226:PRO:O	1:B:2267:MET:CE	2.65	0.41
1:B:2777:TYR:CD1	1:B:2791:LEU:HD12	2.43	0.41
1:B:3300:ALA:HB3	1:B:3301:PRO:CD	2.39	0.41
1:B:3946:GLN:NE2	1:B:3949:ARG:HD3	2.36	0.41
1:B:3959:LYS:HG3	1:B:4022:ASP:OD2	2.21	0.41
1:B:4952:GLU:C	1:B:4952:GLU:OE1	2.60	0.41
1:D:257:ARG:O	1:D:284:HIS:NE2	2.43	0.41
1:D:591:ASP:HA	1:D:631:LEU:HD21	2.03	0.41
1:D:594:GLY:HA2	1:D:1594:ARG:HD2	2.03	0.41
1:D:877:ASN:HD21	1:D:970:LEU:HD13	1.85	0.41
1:D:887:ILE:HG13	1:D:907:LEU:CD1	2.50	0.41
1:D:1423:ASP:HB3	1:D:1426:ILE:HD12	2.01	0.41
1:D:1461:ASP:OD2	1:D:1468:LYS:NZ	2.49	0.41
1:D:1705:GLY:HA3	1:D:1836:PHE:CG	2.56	0.41
1:D:1948:ASP:OD1	1:D:2126:ARG:NH2	2.45	0.41
1:D:2256:TYR:O	1:D:2259:GLU:HG3	2.21	0.41
1:D:2282:ASP:HA	1:D:2341:VAL:HG13	2.02	0.41
1:D:2302:LEU:CD1	1:D:2328:GLY:HA2	2.51	0.41
1:D:2430:ILE:HG22	1:D:2505:PHE:HB2	2.02	0.41
1:D:2608:MET:CE	1:D:2642:LYS:NZ	2.84	0.41
1:D:2699:ALA:O	1:D:2702:CYS:HA	2.21	0.41
1:D:2978:GLU:HA	1:D:2981:VAL:HG12	2.02	0.41
1:D:3054:VAL:HG13	1:D:3055:SER:N	2.36	0.41
1:D:3294:PRO:HB2	1:D:3297:PRO:HD2	2.02	0.41
1:D:3879:GLU:HG2	1:D:3880:PHE:N	2.36	0.41
1:D:4021:LYS:N	1:D:4139:ILE:HD13	2.36	0.41
1:D:4061:PHE:CE2	1:D:4065:PHE:HE2	2.39	0.41
1:C:358:THR:OG1	1:C:383:HIS:ND1	2.42	0.41
1:C:361:ALA:O	1:C:362:PRO:C	2.59	0.41
1:C:371:VAL:O	1:C:371:VAL:HG13	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1063:VAL:HG13	1:C:1064:GLU:N	2.36	0.41
1:C:2044:ILE:H	1:C:2044:ILE:HD12	1.86	0.41
1:C:2282:ASP:HA	1:C:2341:VAL:HG13	2.02	0.41
1:C:2396:GLY:O	1:C:2397:VAL:C	2.59	0.41
1:C:2760:GLU:OE2	1:C:2794:TYR:CE1	2.74	0.41
1:C:3249:LEU:HD11	1:C:3273:THR:HG21	2.01	0.41
1:C:3769:ARG:O	1:C:3773:ARG:NH1	2.48	0.41
1:C:3946:GLN:NE2	1:C:3949:ARG:HD3	2.36	0.41
1:C:4061:PHE:CE2	1:C:4065:PHE:HE2	2.39	0.41
1:C:4942:GLU:O	1:C:4946:GLN:HG3	2.21	0.41
1:A:361:ALA:O	1:A:362:PRO:C	2.59	0.41
1:A:438:ILE:HD12	1:A:438:ILE:H	1.86	0.41
1:A:1157:GLU:OE1	1:A:1157:GLU:N	2.53	0.41
1:A:1425:GLU:OE2	1:A:1425:GLU:N	2.53	0.41
1:A:1931:LEU:HD13	1:A:1935:VAL:CG1	2.47	0.41
1:A:3946:GLN:NE2	1:A:3949:ARG:HD3	2.36	0.41
1:A:3959:LYS:HG3	1:A:4022:ASP:OD2	2.21	0.41
1:B:438:ILE:HD12	1:B:438:ILE:H	1.86	0.41
1:B:2135:LEU:O	1:B:3658:LYS:NZ	2.51	0.41
1:B:2150:GLU:HG2	1:B:2151:ASP:N	2.36	0.41
1:B:2430:ILE:HG22	1:B:2505:PHE:HB2	2.02	0.41
1:B:2760:GLU:OE2	1:B:2794:TYR:CE1	2.74	0.41
1:B:2790:MET:HA	1:B:2792:ARG:NH1	2.36	0.41
1:B:2801:ASP:HA	1:B:2804:ILE:HG12	2.02	0.41
1:B:4021:LYS:N	1:B:4139:ILE:HD13	2.36	0.41
1:D:380:GLN:OE1	1:D:380:GLN:N	2.47	0.41
1:D:1425:GLU:OE2	1:D:1425:GLU:N	2.53	0.41
1:D:2323:TRP:NE1	1:D:2366:PRO:HG3	2.36	0.41
1:D:2486:VAL:O	1:D:2486:VAL:HG13	2.21	0.41
1:D:2799:GLU:O	1:D:2803:GLU:OE1	2.38	0.41
1:D:3281:LEU:HB2	1:D:3282:PRO:HD3	2.03	0.41
1:D:4253:GLU:OE1	1:D:4253:GLU:N	2.52	0.41
1:C:1013:ILE:N	1:C:1014:PRO:CD	2.84	0.41
1:C:2978:GLU:HA	1:C:2981:VAL:HG12	2.02	0.41
1:C:3054:VAL:HG13	1:C:3055:SER:N	2.36	0.41
1:C:3111:ARG:NH2	1:C:3175:LEU:N	2.69	0.41
1:C:3905:THR:HA	1:C:3912:THR:OG1	2.20	0.41
1:C:4205:TRP:CH2	1:C:4214:LYS:HD3	2.56	0.41
1:C:4949:GLN:HA	1:C:4952:GLU:HG3	2.02	0.41
1:A:14:LEU:CB	1:A:101:LEU:HD12	2.50	0.40
1:A:924:MET:O	1:A:928:THR:HG23	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:2233:CYS:HB3	1:A:2237:CYS:HB3	1.67	0.40
1:A:2302:LEU:CD1	1:A:2328:GLY:HA2	2.51	0.40
1:A:2608:MET:CE	1:A:2642:LYS:NZ	2.84	0.40
1:A:2790:MET:HA	1:A:2792:ARG:NH1	2.36	0.40
1:A:3018:LEU:HD21	1:A:3075:LEU:O	2.20	0.40
1:A:3508:SER:O	1:A:3511:VAL:HG22	2.21	0.40
1:A:4021:LYS:N	1:A:4139:ILE:HD13	2.36	0.40
1:B:294:THR:O	1:B:298:GLY:N	2.51	0.40
1:B:2212:VAL:HG23	1:B:2213:ASN:N	2.37	0.40
1:B:2608:MET:CE	1:B:2642:LYS:NZ	2.84	0.40
1:B:2773:ASN:OD1	1:B:2774:ASN:N	2.55	0.40
1:B:3018:LEU:HD21	1:B:3075:LEU:O	2.20	0.40
1:B:3542:LEU:O	1:B:3543:LYS:HB2	2.21	0.40
1:B:3905:THR:HA	1:B:3912:THR:OG1	2.20	0.40
1:D:421:PHE:HE2	1:D:507:ALA:HB2	1.79	0.40
1:D:950:LEU:HD12	1:D:950:LEU:O	2.21	0.40
1:D:1126:GLY:HA3	1:D:1143:TRP:CE3	2.57	0.40
1:D:3680:ALA:HB1	1:D:3683:GLN:NE2	2.37	0.40
1:D:3946:GLN:NE2	1:D:3949:ARG:HD3	2.36	0.40
1:C:1033:ARG:HA	1:C:1036:ARG:NH2	2.36	0.40
1:C:1754:GLY:O	1:C:1758:ARG:HB2	2.20	0.40
1:C:3891:LEU:O	1:C:3899:PHE:CD2	2.71	0.40
1:A:455:PRO:HB2	1:A:459:LEU:HD12	2.03	0.40
1:A:869:ARG:NH2	1:A:941:MET:CE	2.85	0.40
1:A:1031:THR:O	1:A:1032:LYS:C	2.59	0.40
1:A:2699:ALA:O	1:A:2702:CYS:HA	2.21	0.40
1:A:2810:LYS:O	1:A:2814:LYS:HG3	2.22	0.40
1:A:3071:LEU:O	1:A:3075:LEU:HG	2.21	0.40
1:A:3105:LYS:O	1:A:3108:GLU:HG3	2.21	0.40
1:A:3414:ARG:NH1	1:A:3472:ALA:HB3	2.37	0.40
1:A:4061:PHE:CE2	1:A:4065:PHE:HE2	2.39	0.40
1:A:4649:LEU:O	1:A:4653:VAL:HG23	2.22	0.40
1:A:4952:GLU:C	1:A:4952:GLU:OE1	2.60	0.40
1:B:869:ARG:NH2	1:B:941:MET:CE	2.85	0.40
1:B:2376:LEU:HD11	1:B:2426:TYR:HB3	2.02	0.40
1:B:2978:GLU:HA	1:B:2981:VAL:HG12	2.02	0.40
1:B:3077:ALA:O	1:B:3081:MET:HG2	2.21	0.40
1:B:3235:SER:OG	1:B:3237:GLU:OE1	2.38	0.40
1:B:4205:TRP:CH2	1:B:4214:LYS:HD3	2.56	0.40
1:B:4649:LEU:O	1:B:4653:VAL:HG23	2.21	0.40
1:D:438:ILE:HD12	1:D:438:ILE:H	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:869:ARG:NH2	1:D:941:MET:CE	2.85	0.40
1:D:955:LEU:HD12	1:D:966:LYS:HB3	2.03	0.40
1:D:1031:THR:O	1:D:1032:LYS:C	2.59	0.40
1:D:1742:THR:HA	1:D:1745:ILE:HD12	2.02	0.40
1:D:2150:GLU:HG2	1:D:2151:ASP:N	2.36	0.40
1:D:2396:GLY:O	1:D:2397:VAL:C	2.59	0.40
1:D:3504:SER:HB2	1:D:3507:THR:HG23	2.03	0.40
1:D:3535:LEU:O	1:D:3539:ARG:HG2	2.22	0.40
1:D:3628:ARG:HB2	1:D:3631:ALA:HB3	2.03	0.40
1:D:4000:MET:HG2	1:D:4061:PHE:CD2	2.56	0.40
1:D:4704:LEU:O	1:D:4774:LYS:NZ	2.52	0.40
1:C:455:PRO:HB2	1:C:459:LEU:HD12	2.03	0.40
1:C:752:VAL:HB	1:C:753:PRO:C	2.42	0.40
1:C:1031:THR:O	1:C:1032:LYS:C	2.59	0.40
1:C:1204:LEU:HD12	1:C:1234:VAL:HB	2.03	0.40
1:C:1705:GLY:N	1:C:1706:PRO:CD	2.85	0.40
1:C:2150:GLU:HG2	1:C:2151:ASP:N	2.36	0.40
1:C:2256:TYR:O	1:C:2259:GLU:HG3	2.21	0.40
1:C:2376:LEU:HD11	1:C:2426:TYR:HB3	2.02	0.40
1:C:3879:GLU:HG2	1:C:3880:PHE:N	2.36	0.40
1:C:4826:ILE:O	1:C:4829:SER:OG	2.25	0.40
1:A:418:LEU:HD23	1:A:421:PHE:HZ	1.86	0.40
1:A:591:ASP:HA	1:A:631:LEU:HD21	2.03	0.40
1:A:955:LEU:HD12	1:A:966:LYS:HB3	2.03	0.40
1:A:2782:ASP:N	1:A:2782:ASP:OD1	2.54	0.40
1:A:3111:ARG:NH2	1:A:3175:LEU:N	2.69	0.40
1:A:3542:LEU:O	1:A:3543:LYS:HB2	2.21	0.40
1:A:4205:TRP:CH2	1:A:4214:LYS:HD3	2.56	0.40
2:H:74:LEU:HD13	2:H:101:VAL:CG2	2.51	0.40
1:B:708:GLY:HA3	1:B:722:TRP:HB3	2.02	0.40
1:B:1013:ILE:N	1:B:1014:PRO:CD	2.84	0.40
1:B:1033:ARG:HA	1:B:1036:ARG:NH2	2.36	0.40
1:B:1123:VAL:HG23	1:B:1132:TRP:HB2	2.04	0.40
1:B:1126:GLY:HA3	1:B:1143:TRP:CE3	2.57	0.40
1:B:2699:ALA:O	1:B:2702:CYS:HA	2.21	0.40
1:B:3281:LEU:HB2	1:B:3282:PRO:HD3	2.03	0.40
1:B:4949:GLN:HA	1:B:4952:GLU:HG3	2.02	0.40
1:D:711:LEU:HD23	1:D:711:LEU:HA	1.97	0.40
1:D:842:PRO:O	1:D:1196:PRO:HA	2.21	0.40
1:D:1123:VAL:HG23	1:D:1132:TRP:HB2	2.04	0.40
1:D:2349:ASN:O	1:D:2353:VAL:HG23	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:3622:LYS:HB3	1:D:3625:SER:OG	2.21	0.40
1:D:3891:LEU:O	1:D:3899:PHE:CD2	2.71	0.40
1:C:439:GLU:HG2	1:C:440:ALA:N	2.36	0.40
1:C:2799:GLU:O	1:C:2803:GLU:OE1	2.38	0.40
1:C:3725:TYR:O	1:C:3729:MET:HG3	2.20	0.40
1:A:908:VAL:HG23	1:A:963:ASN:HD22	1.86	0.40
1:A:1013:ILE:N	1:A:1014:PRO:CD	2.84	0.40
1:A:1126:GLY:HA3	1:A:1143:TRP:CE3	2.57	0.40
1:A:1979:LEU:CA	1:A:1982:ARG:NH2	2.84	0.40
1:A:2282:ASP:HA	1:A:2341:VAL:HG13	2.02	0.40
1:A:2482:ASP:OD2	1:A:2482:ASP:C	2.60	0.40
1:A:3454:GLU:HA	1:A:3454:GLU:OE2	2.22	0.40
1:A:4098:ASP:OD1	1:A:4098:ASP:C	2.60	0.40
1:A:4942:GLU:O	1:A:4946:GLN:HG3	2.21	0.40
2:G:74:LEU:HD13	2:G:101:VAL:CG2	2.51	0.40
1:B:106:ALA:HA	1:B:148:TRP:O	2.22	0.40
1:B:752:VAL:HB	1:B:753:PRO:C	2.42	0.40
1:B:1705:GLY:HA3	1:B:1836:PHE:CG	2.56	0.40
1:B:2107:GLN:O	1:B:3694:LYS:NZ	2.34	0.40
1:B:2323:TRP:NE1	1:B:2366:PRO:HG3	2.36	0.40
1:B:2349:ASN:O	1:B:2353:VAL:HG23	2.21	0.40
1:B:2396:GLY:O	1:B:2397:VAL:C	2.60	0.40
1:B:2614:ILE:CG2	1:B:2618:MET:HB2	2.51	0.40
1:B:3054:VAL:HG13	1:B:3055:SER:N	2.36	0.40
1:B:3454:GLU:HA	1:B:3454:GLU:OE2	2.22	0.40
1:B:4627:MET:HG2	1:B:4629:TYR:CZ	2.57	0.40
1:D:371:VAL:O	1:D:371:VAL:HG13	2.20	0.40
1:D:1669:LEU:O	1:D:1673:VAL:HG22	2.21	0.40
1:D:2441:HIS:CE1	1:D:2442:LEU:HG	2.56	0.40
1:C:210:GLU:HB2	1:C:213:TYR:HD2	1.86	0.40
1:C:224:HIS:HB2	1:C:229:GLU:OE2	2.21	0.40
1:C:1126:GLY:HA3	1:C:1143:TRP:CE3	2.57	0.40
1:C:2614:ILE:CG2	1:C:2618:MET:HB2	2.51	0.40
1:C:3071:LEU:O	1:C:3075:LEU:HG	2.21	0.40
1:C:3622:LYS:HB3	1:C:3625:SER:OG	2.21	0.40
1:A:842:PRO:O	1:A:1196:PRO:HA	2.21	0.40
1:A:2486:VAL:O	1:A:2486:VAL:HG13	2.21	0.40
1:A:2768:PHE:O	1:A:2772:GLN:OE1	2.40	0.40
1:A:2773:ASN:OD1	1:A:2774:ASN:N	2.55	0.40
1:A:2792:ARG:HB3	1:A:2796:THR:OG1	2.22	0.40
1:A:3680:ALA:HB1	1:A:3683:GLN:NE2	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:F:2:VAL:HG22	2:F:58:GLY:HA2	2.03	0.40
2:F:90:ILE:HD11	1:B:1684:ALA:HA	2.03	0.40
1:B:1705:GLY:N	1:B:1706:PRO:CD	2.84	0.40
1:B:1979:LEU:CA	1:B:1982:ARG:NH2	2.84	0.40
1:B:2243:SER:HB2	1:B:3861:GLU:OE2	2.22	0.40
1:B:4098:ASP:OD1	1:B:4098:ASP:C	2.60	0.40
1:B:4942:GLU:O	1:B:4946:GLN:HG3	2.21	0.40
1:D:439:GLU:HG2	1:D:440:ALA:N	2.36	0.40
1:D:1013:ILE:N	1:D:1014:PRO:CD	2.84	0.40
1:D:1978:ALA:CA	1:D:1982:ARG:HH12	2.35	0.40
1:D:2243:SER:HB2	1:D:3861:GLU:OE2	2.22	0.40
1:D:2513:GLU:N	1:D:2513:GLU:OE1	2.55	0.40
1:D:2790:MET:HA	1:D:2792:ARG:NH1	2.36	0.40
1:C:438:ILE:HD12	1:C:438:ILE:H	1.87	0.40
1:C:842:PRO:O	1:C:1196:PRO:HA	2.21	0.40
1:C:1948:ASP:OD1	1:C:2126:ARG:NH2	2.45	0.40
1:C:1978:ALA:CA	1:C:1982:ARG:HH12	2.35	0.40
1:C:2212:VAL:HG23	1:C:2213:ASN:N	2.37	0.40
1:C:2302:LEU:CD1	1:C:2328:GLY:HA2	2.51	0.40
1:C:2486:VAL:HG13	1:C:2486:VAL:O	2.21	0.40
1:C:2773:ASN:OD1	1:C:2774:ASN:N	2.55	0.40
1:C:2790:MET:HA	1:C:2792:ARG:NH1	2.36	0.40
1:C:4851:TYR:HD1	1:C:4916:PHE:CE1	2.40	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%)	4245 (97%)	136 (3%)	4 (0%)	48 65
1	B	4385/5037 (87%)	4245 (97%)	136 (3%)	4 (0%)	48 65

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	C	4385/5037 (87%)	4245 (97%)	136 (3%)	4 (0%)	48	65
1	D	4385/5037 (87%)	4245 (97%)	136 (3%)	4 (0%)	48	65
2	E	105/108 (97%)	99 (94%)	6 (6%)	0	100	100
2	F	105/108 (97%)	99 (94%)	6 (6%)	0	100	100
2	G	105/108 (97%)	99 (94%)	6 (6%)	0	100	100
2	H	105/108 (97%)	99 (94%)	6 (6%)	0	100	100
All	All	17960/20580 (87%)	17376 (97%)	568 (3%)	16 (0%)	50	65

All (16) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	A	3300	ALA
1	B	3300	ALA
1	D	3300	ALA
1	C	3300	ALA
1	A	2669	GLU
1	B	2669	GLU
1	D	2669	GLU
1	C	2669	GLU
1	A	2948	THR
1	B	2948	THR
1	D	2948	THR
1	C	2948	THR
1	A	4691	GLN
1	B	4691	GLN
1	D	4691	GLN
1	C	4691	GLN

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%)	3815 (100%)	21 (0%)	86	93

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	B	3836/4276 (90%)	3815 (100%)	21 (0%)	86	93
1	C	3836/4276 (90%)	3815 (100%)	21 (0%)	86	93
1	D	3836/4276 (90%)	3815 (100%)	21 (0%)	86	93
2	E	89/90 (99%)	86 (97%)	3 (3%)	32	49
2	F	89/90 (99%)	86 (97%)	3 (3%)	32	49
2	G	89/90 (99%)	86 (97%)	3 (3%)	32	49
2	H	89/90 (99%)	86 (97%)	3 (3%)	32	49
All	All	15700/17464 (90%)	15604 (99%)	96 (1%)	82	92

All (96) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	155	LYS
1	A	421	PHE
1	A	941	MET
1	A	945	LYS
1	A	998	ARG
1	A	1036	ARG
1	A	1618	ARG
1	A	1759	ARG
1	A	2336	ARG
1	A	2738	ARG
1	A	2870[A]	GLU
1	A	2870[B]	GLU
1	A	3617	LYS
1	A	3630	ARG
1	A	3758	MET
1	A	3858	MET
1	A	3951	PHE
1	A	4161	ARG
1	A	4223	ASN
1	A	4720	VAL
1	A	4952	GLU
2	E	3	GLN
2	E	13	ARG
2	E	52	LYS
2	H	3	GLN
2	H	13	ARG
2	H	52	LYS
2	G	3	GLN

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Mol	Chain	Res	Type
2	G	13	ARG
2	G	52	LYS
2	F	3	GLN
2	F	13	ARG
2	F	52	LYS
1	B	155	LYS
1	B	421	PHE
1	B	941	MET
1	B	945	LYS
1	B	998	ARG
1	B	1036	ARG
1	B	1618	ARG
1	B	1759	ARG
1	B	2336	ARG
1	B	2738	ARG
1	B	2870[A]	GLU
1	B	2870[B]	GLU
1	B	3617	LYS
1	B	3630	ARG
1	B	3758	MET
1	B	3858	MET
1	B	3951	PHE
1	B	4161	ARG
1	B	4223	ASN
1	B	4720	VAL
1	B	4952	GLU
1	D	155	LYS
1	D	421	PHE
1	D	941	MET
1	D	945	LYS
1	D	998	ARG
1	D	1036	ARG
1	D	1618	ARG
1	D	1759	ARG
1	D	2336	ARG
1	D	2738	ARG
1	D	2870[A]	GLU
1	D	2870[B]	GLU
1	D	3617	LYS
1	D	3630	ARG
1	D	3758	MET
1	D	3858	MET

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Mol	Chain	Res	Type
1	D	3951	PHE
1	D	4161	ARG
1	D	4223	ASN
1	D	4720	VAL
1	D	4952	GLU
1	C	155	LYS
1	C	421	PHE
1	C	941	MET
1	C	945	LYS
1	C	998	ARG
1	C	1036	ARG
1	C	1618	ARG
1	C	1759	ARG
1	C	2336	ARG
1	C	2738	ARG
1	C	2870[A]	GLU
1	C	2870[B]	GLU
1	C	3617	LYS
1	C	3630	ARG
1	C	3758	MET
1	C	3858	MET
1	C	3951	PHE
1	C	4161	ARG
1	C	4223	ASN
1	C	4720	VAL
1	C	4952	GLU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (55) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	105	HIS
1	A	838	HIS
1	A	879	HIS
1	A	889	GLN
1	A	904	HIS
1	A	1041	GLN
1	A	2417	HIS
1	A	2902	HIS
1	A	2991	HIS
1	A	3052	HIS
1	A	3109	ASN
1	A	3127	GLN

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Mol	Chain	Res	Type
1	A	5003	HIS
2	E	25	HIS
2	H	25	HIS
2	G	25	HIS
2	F	25	HIS
1	B	105	HIS
1	B	838	HIS
1	B	879	HIS
1	B	889	GLN
1	B	904	HIS
1	B	1041	GLN
1	B	2902	HIS
1	B	2991	HIS
1	B	3052	HIS
1	B	3109	ASN
1	B	3127	GLN
1	B	5003	HIS
1	D	105	HIS
1	D	838	HIS
1	D	879	HIS
1	D	889	GLN
1	D	904	HIS
1	D	1041	GLN
1	D	2417	HIS
1	D	2902	HIS
1	D	2991	HIS
1	D	3052	HIS
1	D	3109	ASN
1	D	3127	GLN
1	D	5003	HIS
1	C	105	HIS
1	C	838	HIS
1	C	879	HIS
1	C	889	GLN
1	C	904	HIS
1	C	1041	GLN
1	C	2417	HIS
1	C	2902	HIS
1	C	2991	HIS
1	C	3052	HIS
1	C	3109	ASN
1	C	3127	GLN

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Mol	Chain	Res	Type
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	0.61	0	34,52,52	1.00	3 (8%)
6	IBM	B	5304	-	8,17,17	0.80	0	11,25,25	2.30	4 (36%)
3	ATP	A	5301	-	28,33,33	0.60	0	34,52,52	1.00	3 (8%)
6	IBM	A	5304	-	8,17,17	0.77	0	11,25,25	2.29	4 (36%)
3	ATP	C	5301	-	28,33,33	0.61	0	34,52,52	1.00	3 (8%)
6	IBM	D	5304	-	8,17,17	0.78	0	11,25,25	2.29	4 (36%)
6	IBM	C	5304	-	8,17,17	0.78	0	11,25,25	2.29	4 (36%)
3	ATP	D	5301	-	28,33,33	0.60	0	34,52,52	1.00	3 (8%)

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

There are no bond length outliers.

All (28) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	D	5304	IBM	C5-C6-N1	-4.77	113.36	118.20
6	B	5304	IBM	C5-C6-N1	-4.77	113.36	118.20
6	C	5304	IBM	C5-C6-N1	-4.75	113.37	118.20
6	A	5304	IBM	C5-C6-N1	-4.74	113.39	118.20
6	B	5304	IBM	C11-N3-C4	4.02	121.87	118.46
6	C	5304	IBM	C11-N3-C4	4.02	121.87	118.46
6	A	5304	IBM	C11-N3-C4	4.01	121.86	118.46
6	D	5304	IBM	C11-N3-C4	3.98	121.83	118.46
6	D	5304	IBM	C4-C5-C6	3.12	122.33	119.96
6	B	5304	IBM	C4-C5-C6	3.11	122.33	119.96
6	C	5304	IBM	C4-C5-C6	3.07	122.30	119.96
6	A	5304	IBM	C4-C5-C6	3.05	122.28	119.96
3	B	5301	ATP	O4'-C1'-N9	-2.45	105.49	108.75
3	D	5301	ATP	O4'-C1'-N9	-2.43	105.52	108.75
3	A	5301	ATP	O4'-C1'-N9	-2.42	105.53	108.75
6	A	5304	IBM	C5-C4-N9	-2.42	105.90	110.87
6	B	5304	IBM	C5-C4-N9	-2.41	105.92	110.87
6	C	5304	IBM	C5-C4-N9	-2.41	105.92	110.87
3	C	5301	ATP	C5-C6-N6	2.40	123.97	120.31
6	D	5304	IBM	C5-C4-N9	-2.40	105.93	110.87
3	B	5301	ATP	C5-C6-N6	2.40	123.97	120.31
3	C	5301	ATP	O4'-C1'-N9	-2.39	105.58	108.75
3	A	5301	ATP	C5-C6-N6	2.38	123.94	120.31
3	D	5301	ATP	C5-C6-N6	2.37	123.92	120.31
3	D	5301	ATP	O2'-C2'-C3'	-2.26	104.57	111.82

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	A	5301	ATP	O2'-C2'-C3'	-2.26	104.59	111.82
3	C	5301	ATP	O2'-C2'-C3'	-2.25	104.60	111.82
3	B	5301	ATP	O2'-C2'-C3'	-2.25	104.61	111.82

There are no chirality outliers.

All (36) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
6	A	5304	IBM	C12-C11-N3-C4
6	A	5304	IBM	N3-C11-C12-C13
6	A	5304	IBM	N3-C11-C12-C14
6	B	5304	IBM	C12-C11-N3-C4
6	B	5304	IBM	N3-C11-C12-C13
6	B	5304	IBM	N3-C11-C12-C14
6	D	5304	IBM	C12-C11-N3-C4
6	D	5304	IBM	N3-C11-C12-C13
6	D	5304	IBM	N3-C11-C12-C14
6	C	5304	IBM	C12-C11-N3-C4
6	C	5304	IBM	N3-C11-C12-C13
6	C	5304	IBM	N3-C11-C12-C14
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	PB-O3B-PG-O1G
3	B	5301	ATP	PB-O3B-PG-O1G
3	D	5301	ATP	PB-O3B-PG-O1G
3	C	5301	ATP	PB-O3B-PG-O1G
3	A	5301	ATP	PB-O3A-PA-O1A
3	B	5301	ATP	PB-O3A-PA-O1A
3	D	5301	ATP	PB-O3A-PA-O1A
3	C	5301	ATP	PB-O3A-PA-O1A
3	A	5301	ATP	PB-O3B-PG-O2G
3	A	5301	ATP	PB-O3B-PG-O3G
3	B	5301	ATP	PB-O3B-PG-O2G
3	B	5301	ATP	PB-O3B-PG-O3G
3	D	5301	ATP	PB-O3B-PG-O2G
3	D	5301	ATP	PB-O3B-PG-O3G
3	C	5301	ATP	PB-O3B-PG-O2G
3	C	5301	ATP	PB-O3B-PG-O3G
3	A	5301	ATP	PB-O3A-PA-O2A
3	B	5301	ATP	PB-O3A-PA-O2A

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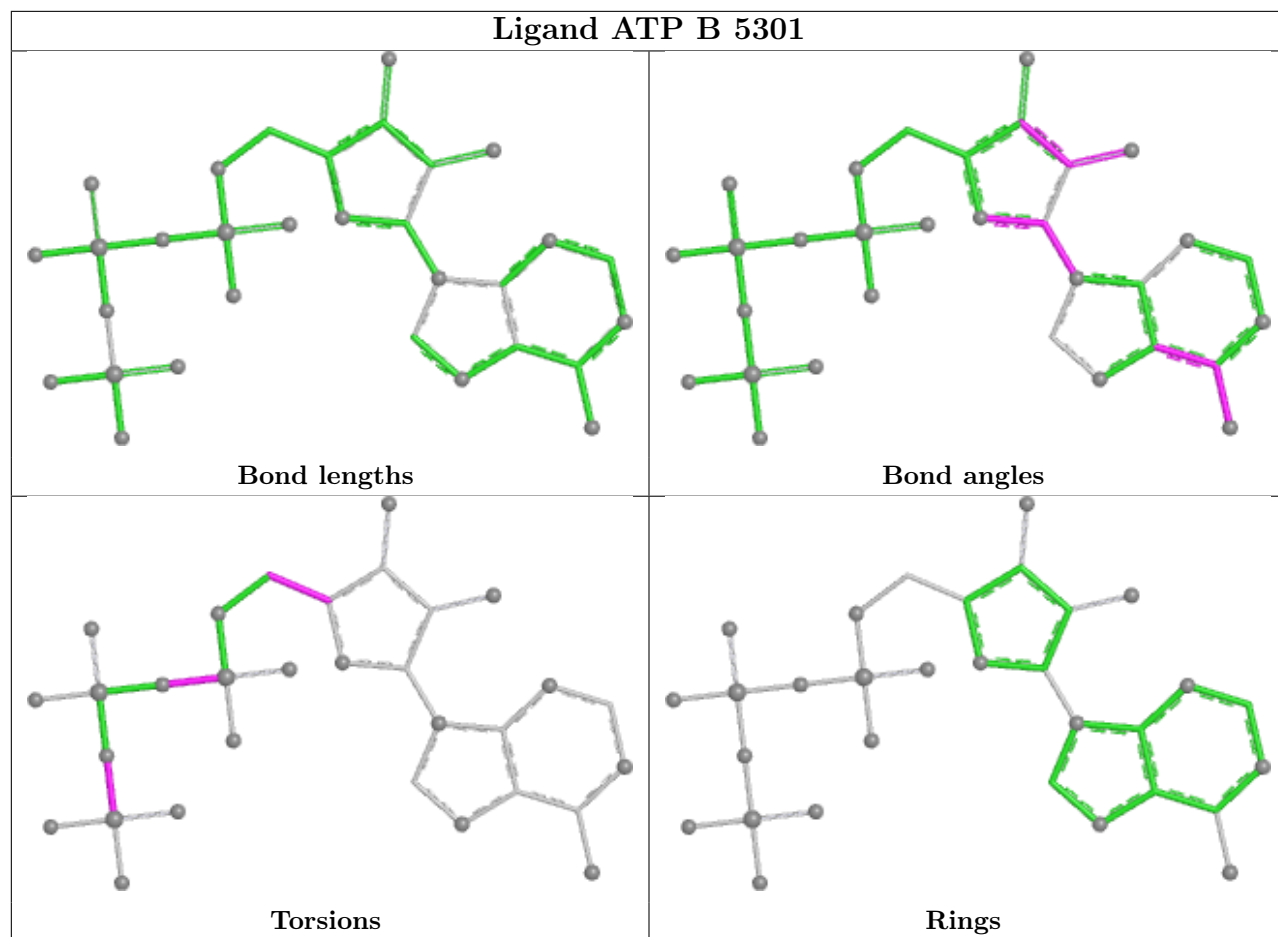
Mol	Chain	Res	Type	Atoms
3	D	5301	ATP	PB-O3A-PA-O2A
3	C	5301	ATP	PB-O3A-PA-O2A

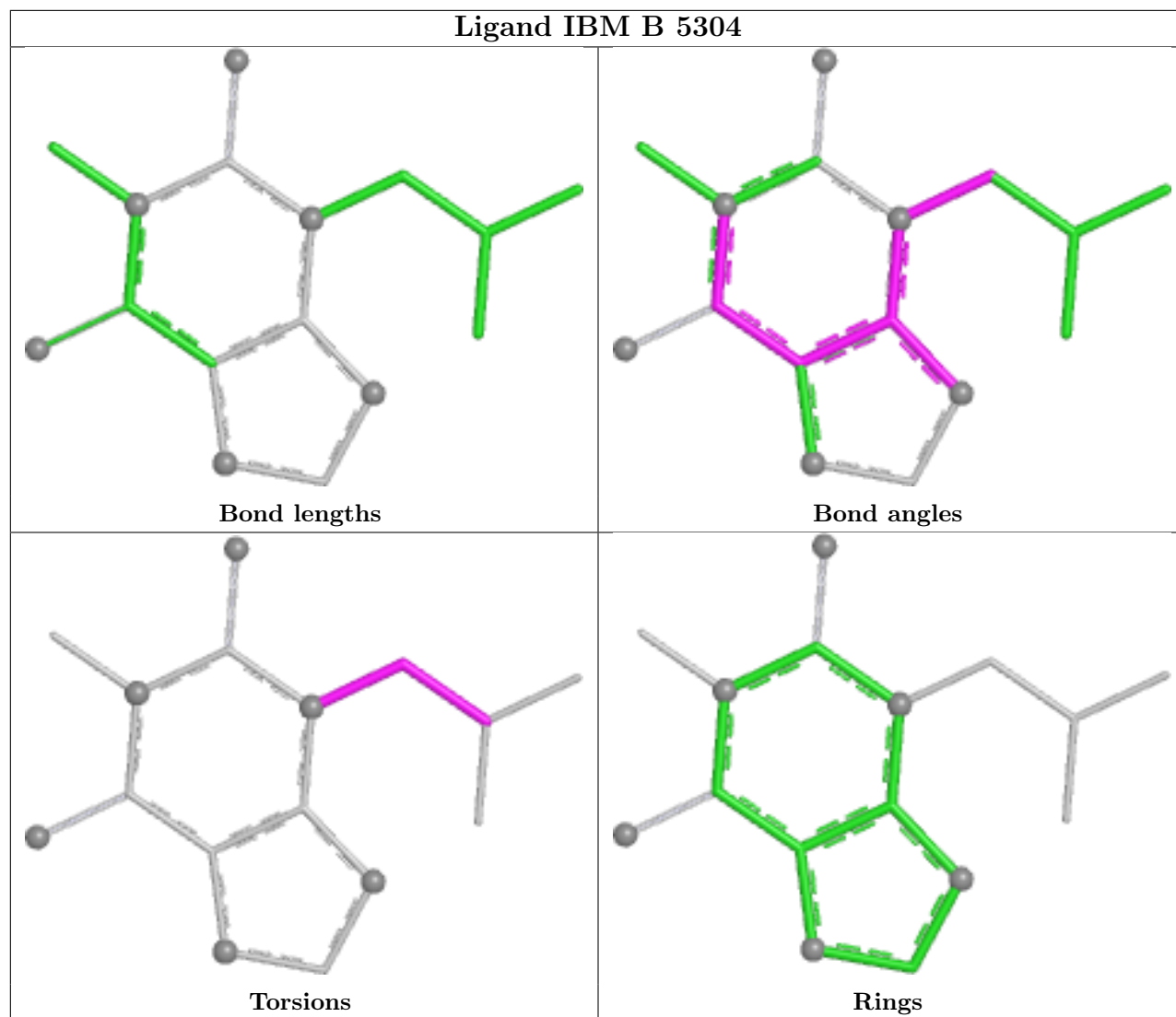
There are no ring outliers.

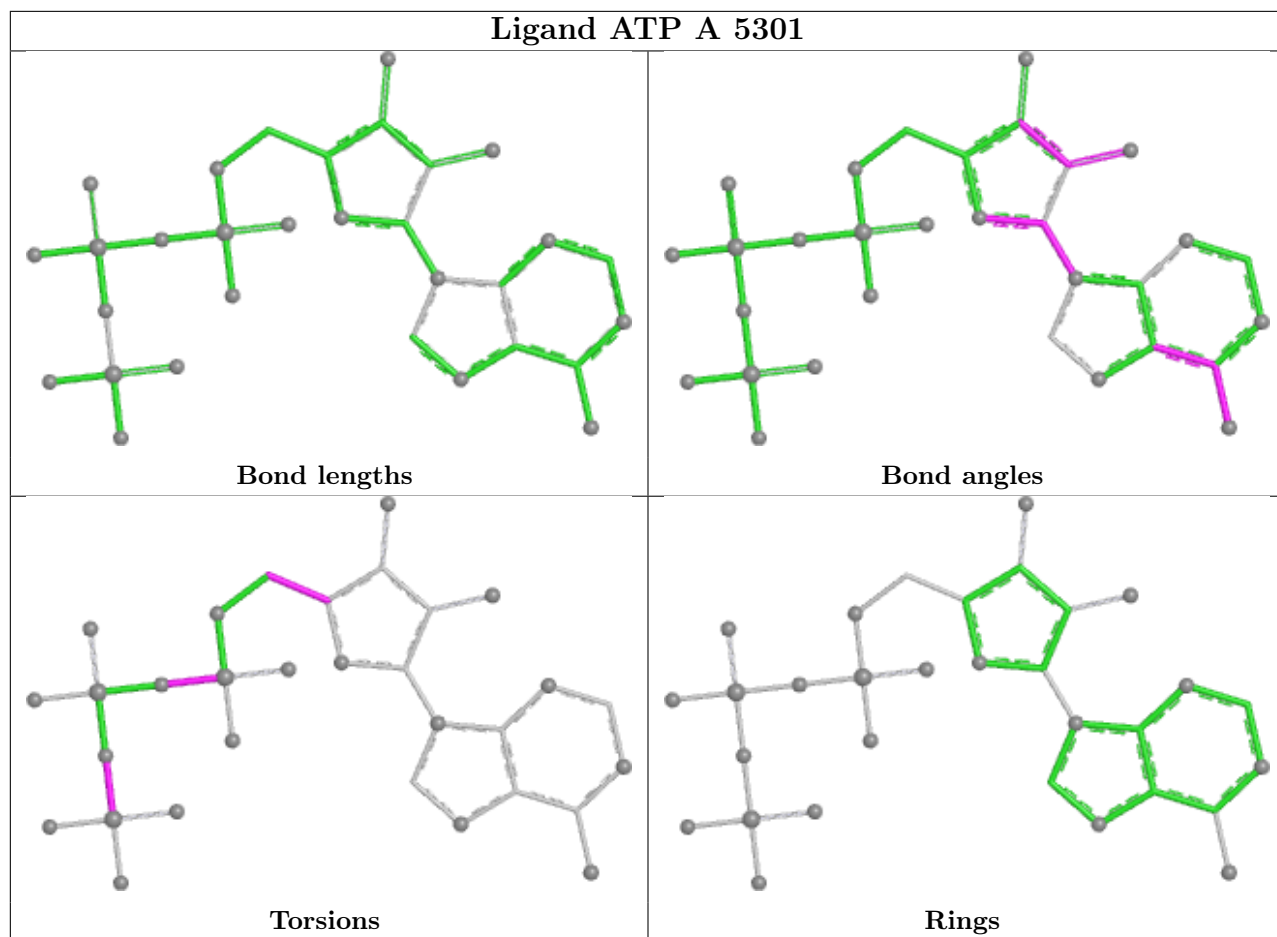
8 monomers are involved in 12 short contacts:

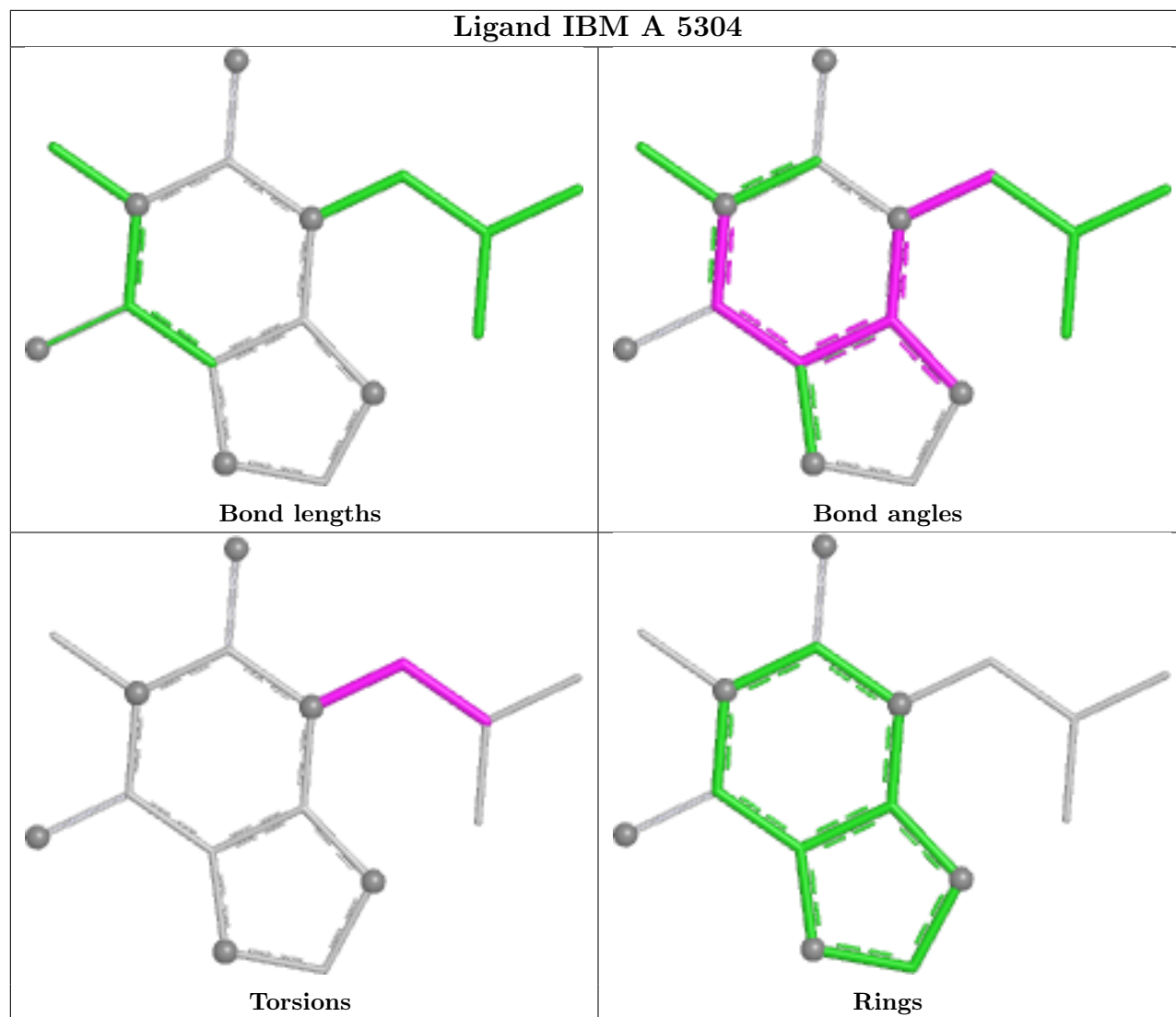
Mol	Chain	Res	Type	Clashes	Symm-Clashes
3	B	5301	ATP	1	0
6	B	5304	IBM	2	0
3	A	5301	ATP	1	0
6	A	5304	IBM	2	0
3	C	5301	ATP	1	0
6	D	5304	IBM	2	0
6	C	5304	IBM	2	0
3	D	5301	ATP	1	0

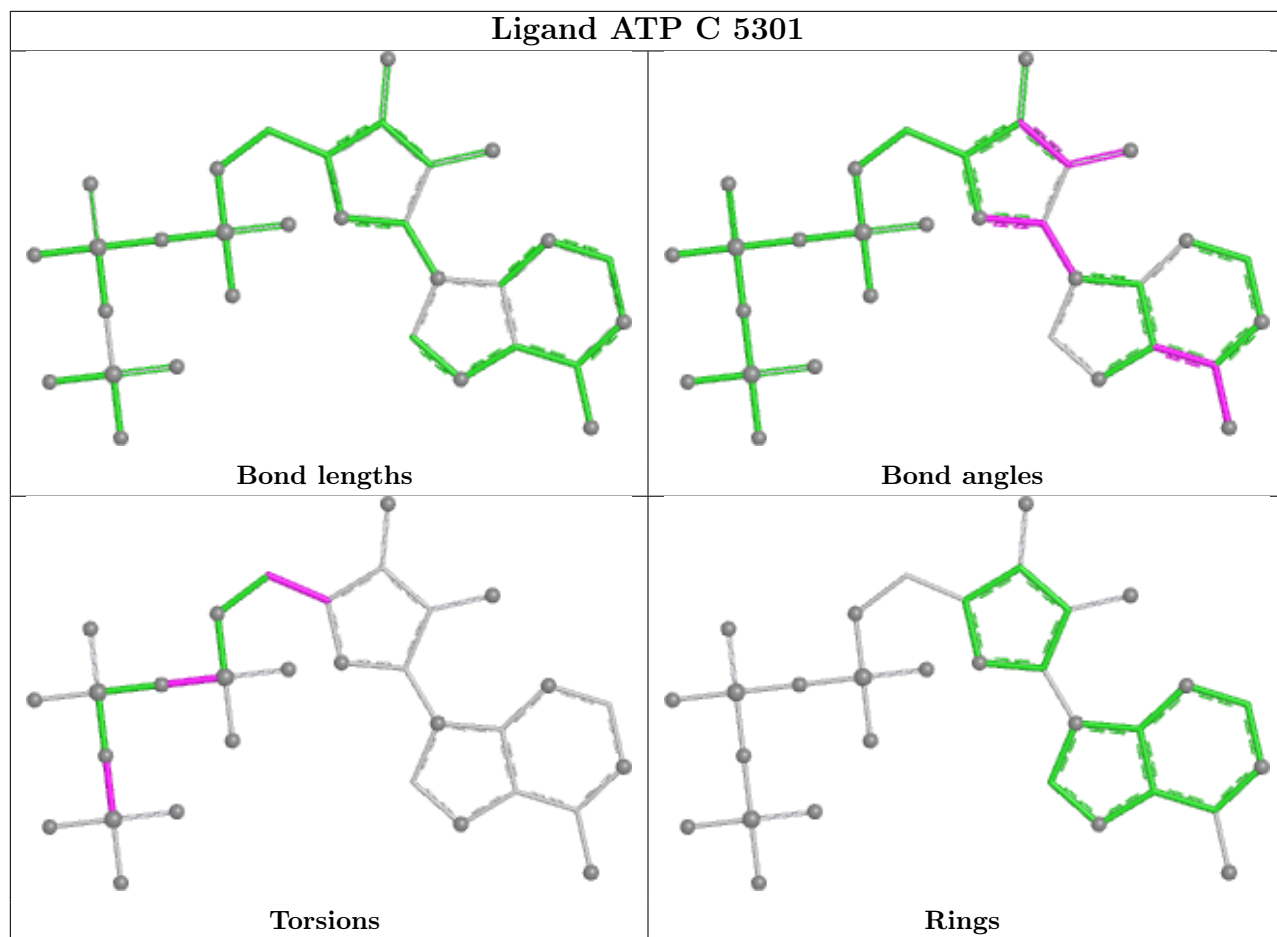
The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included.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

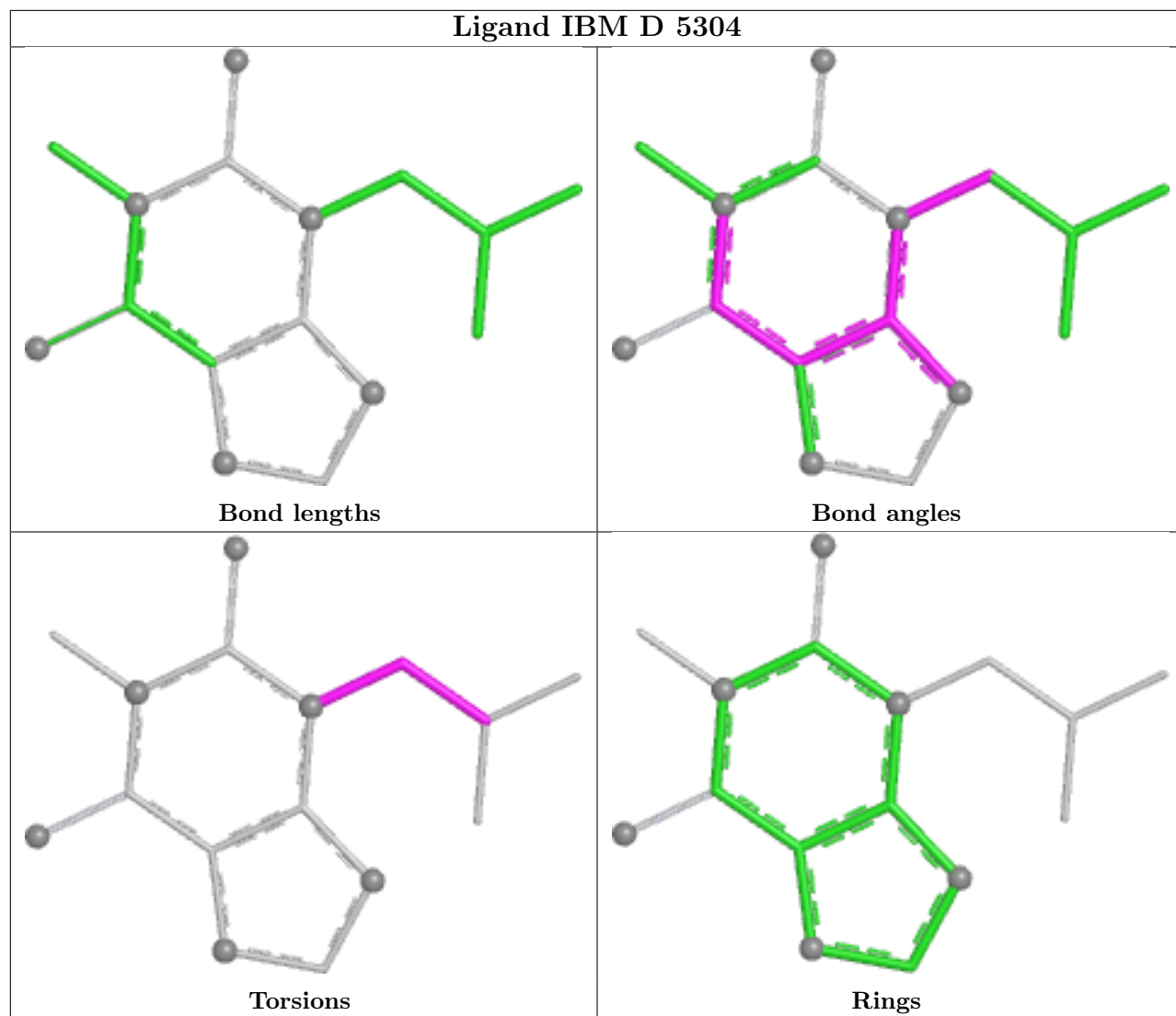


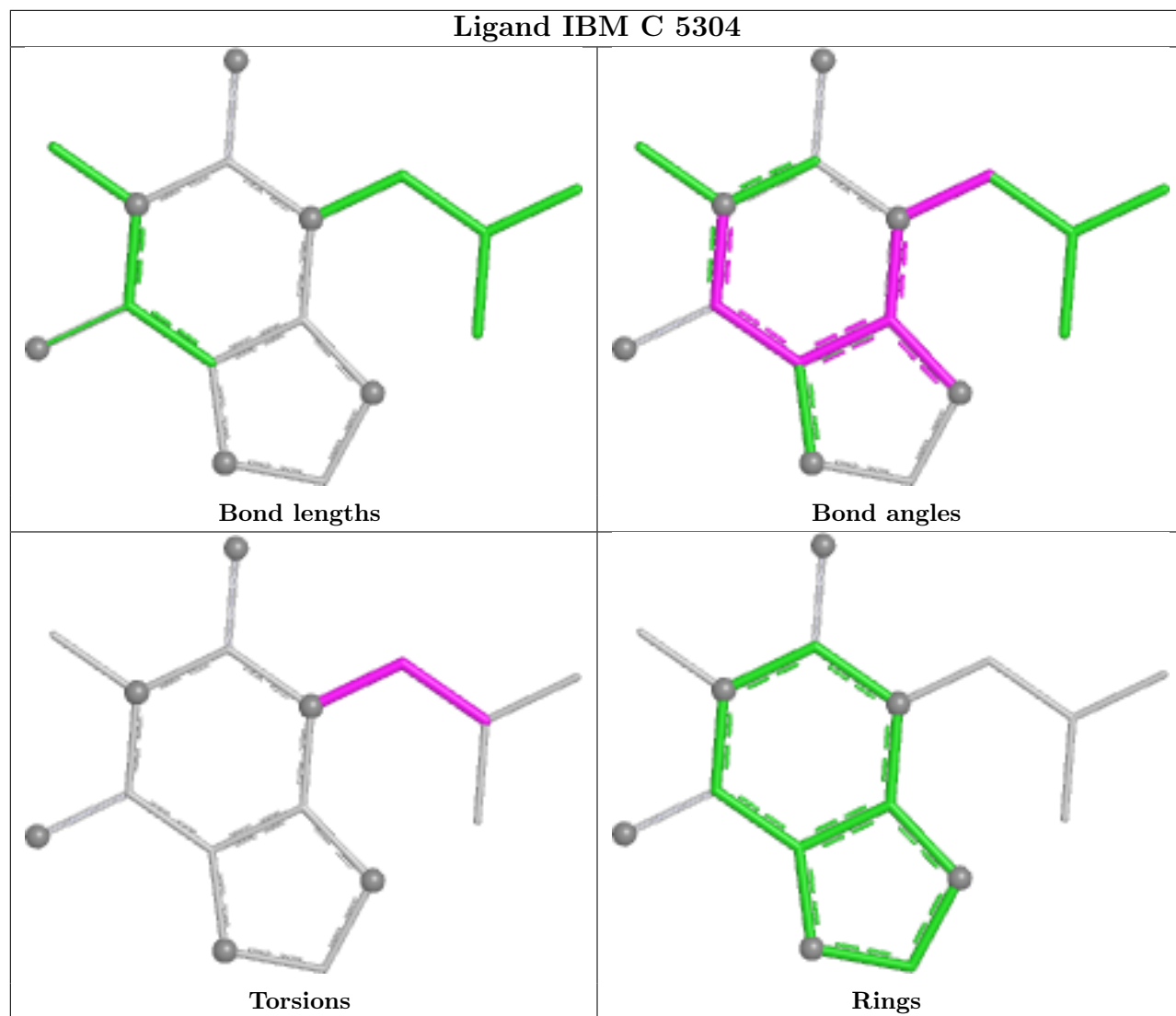


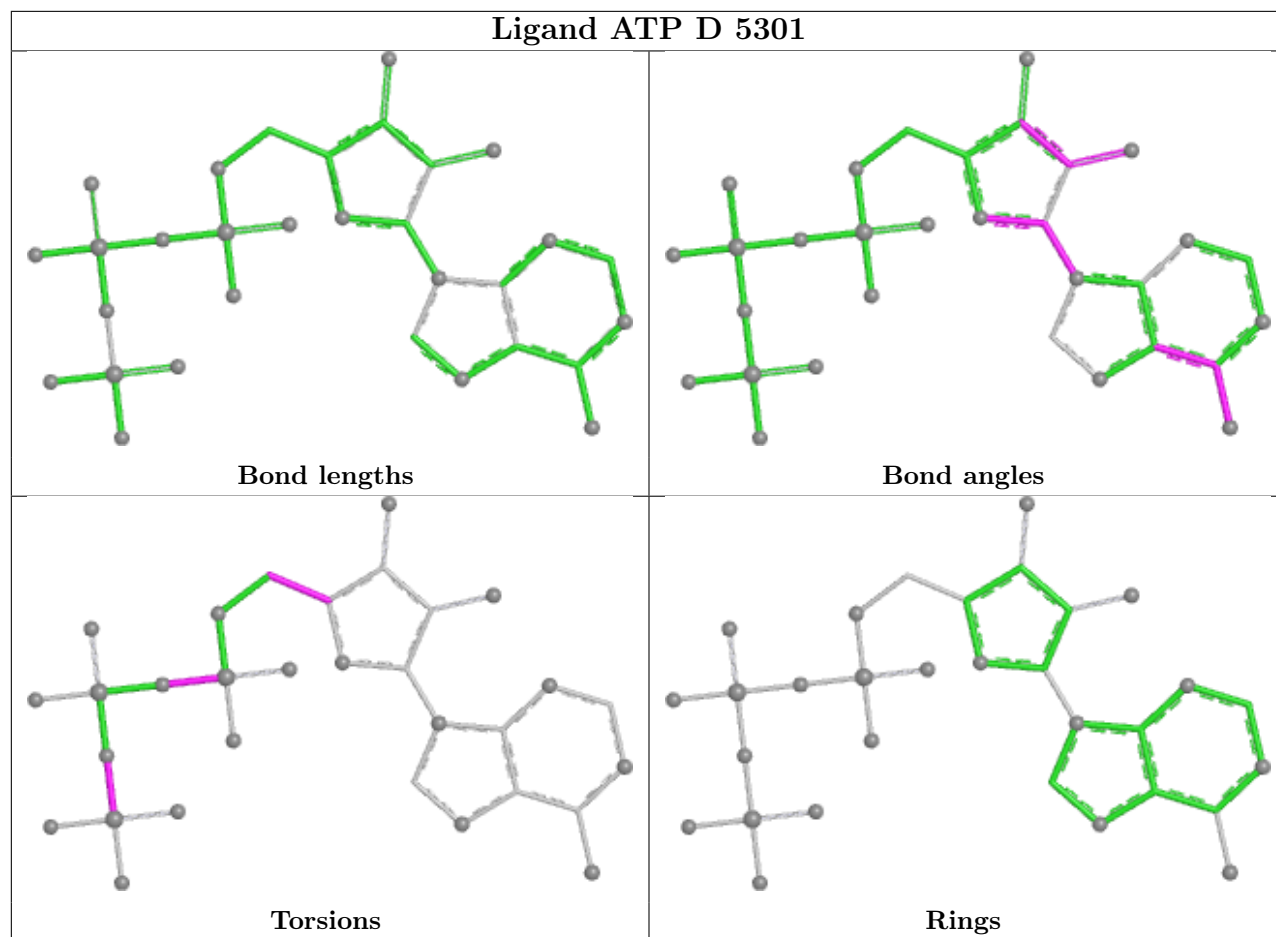












5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

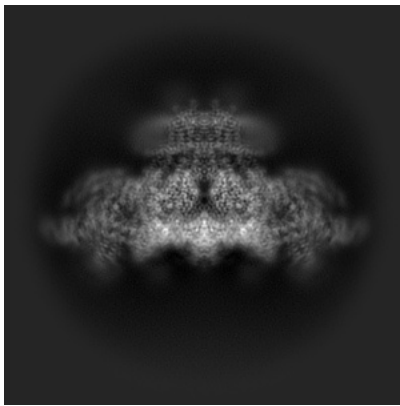
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-47389. These allow visual inspection of the internal detail of the map and identification of artifacts.

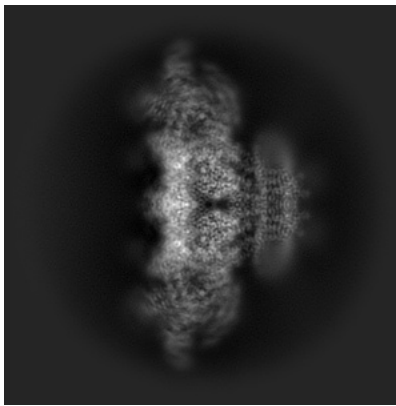
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

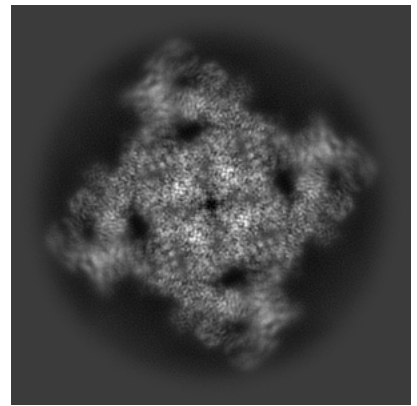
6.1.1 Primary map



X

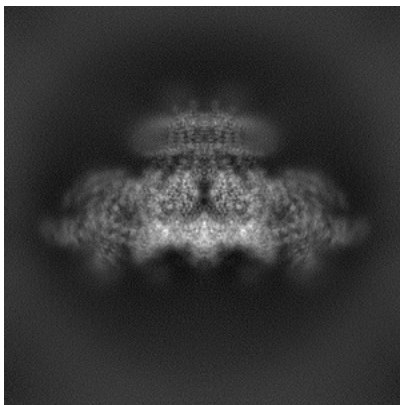


Y

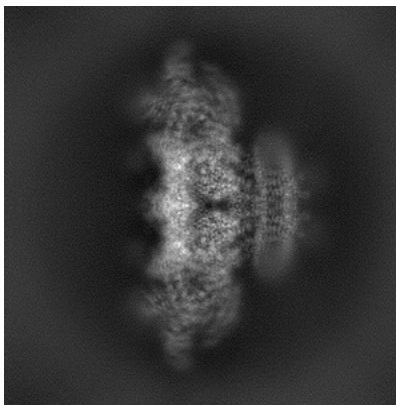


Z

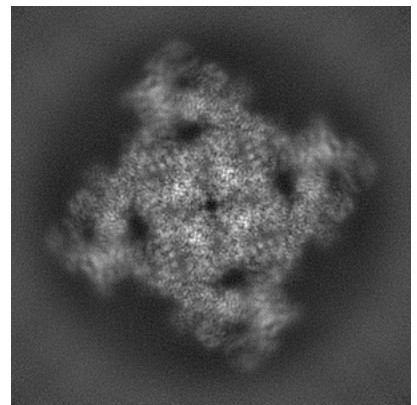
6.1.2 Raw map



X



Y

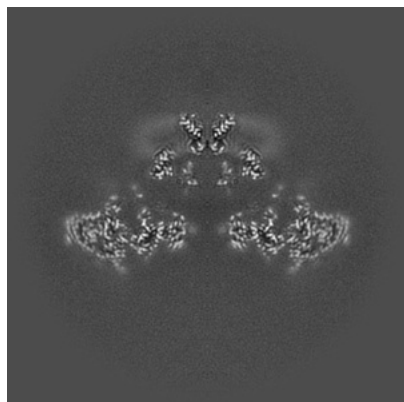


Z

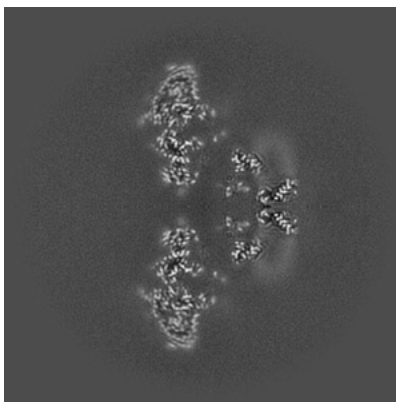
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

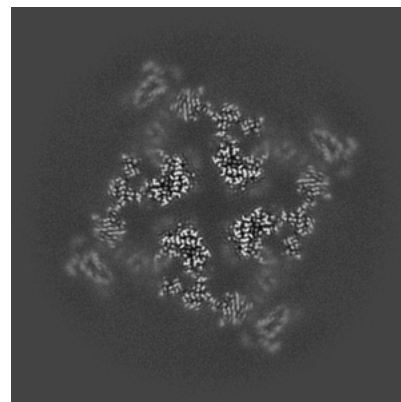
6.2.1 Primary map



X Index: 256

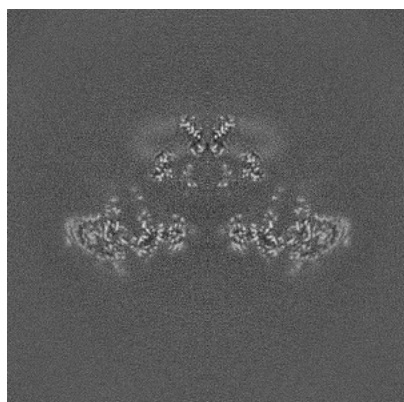


Y Index: 256

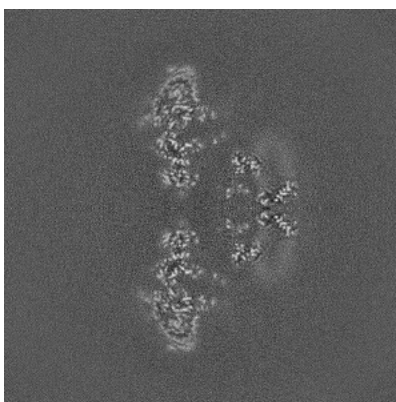


Z Index: 256

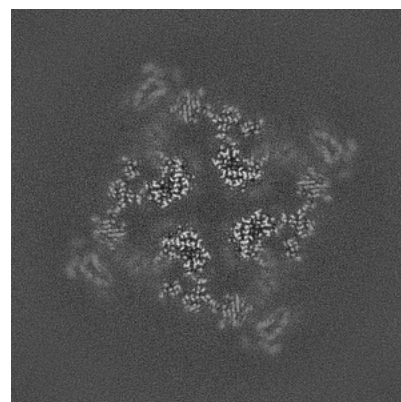
6.2.2 Raw map



X Index: 256



Y Index: 256

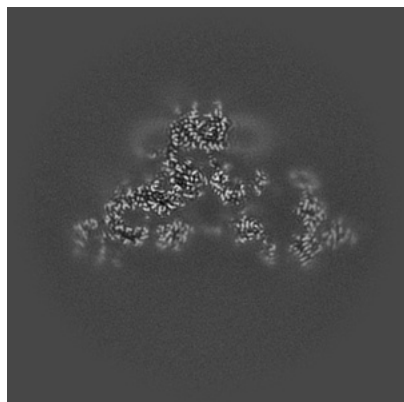


Z Index: 256

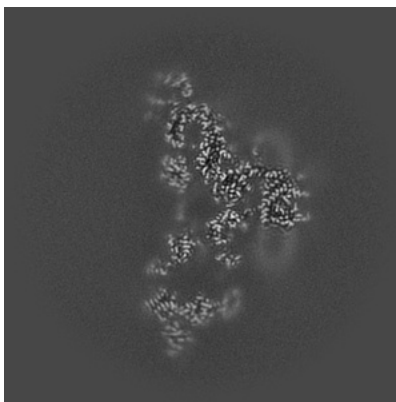
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

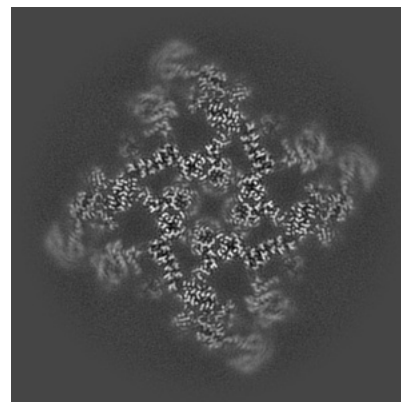
6.3.1 Primary map



X Index: 239

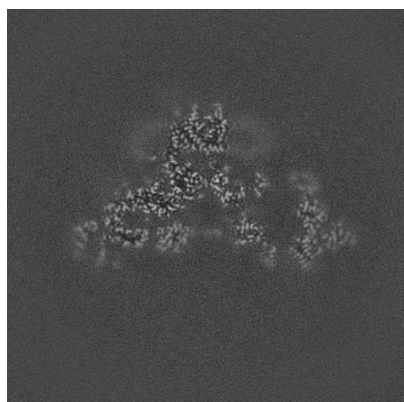


Y Index: 239

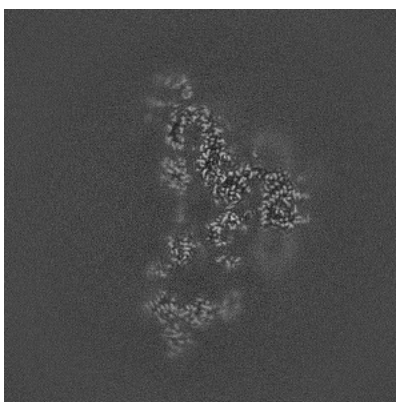


Z Index: 229

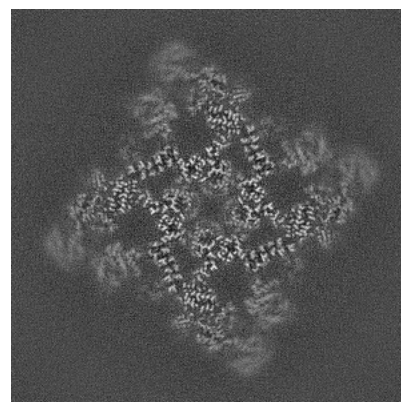
6.3.2 Raw map



X Index: 239



Y Index: 239

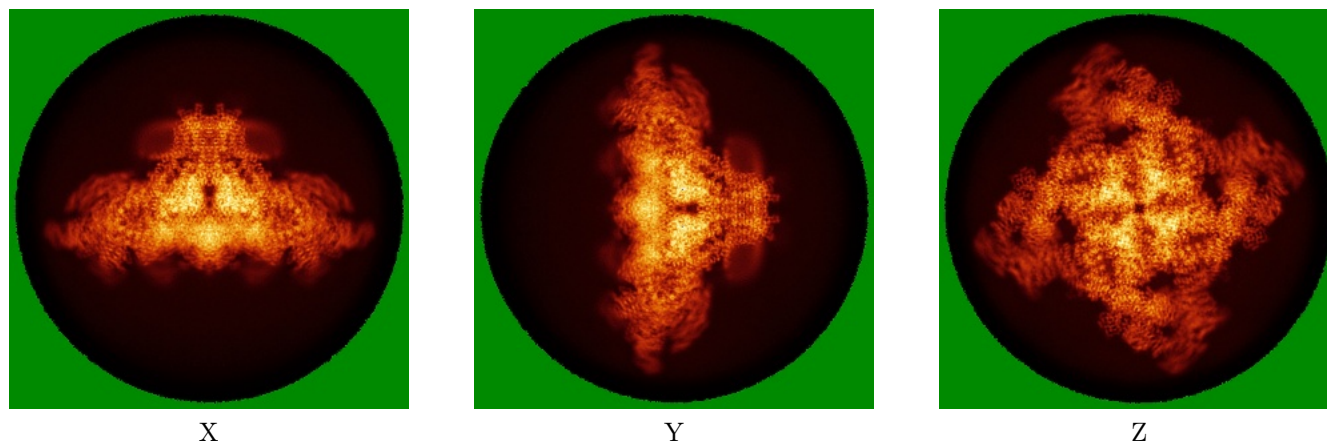


Z Index: 229

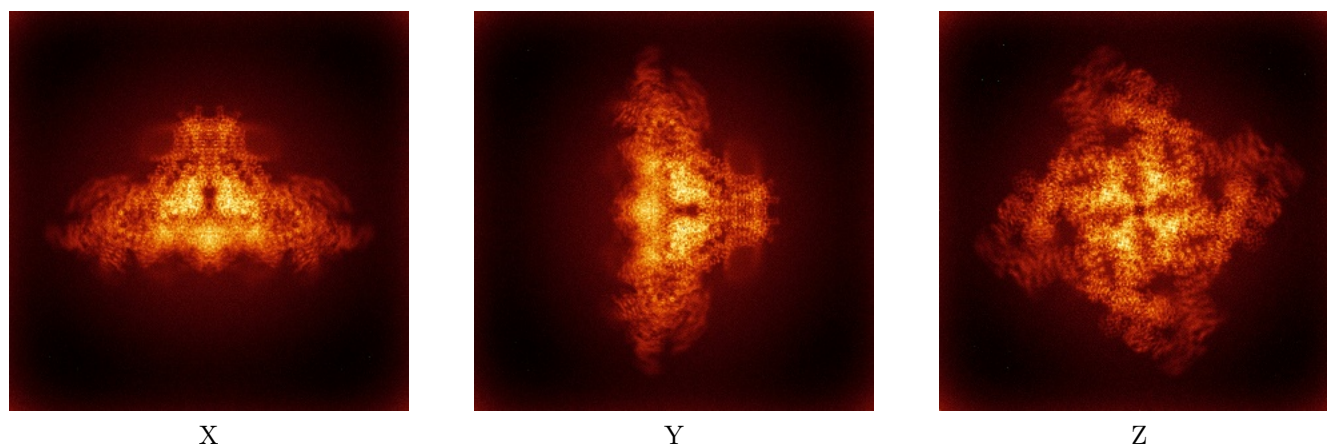
The images above show the largest variance slices of the map in three orthogonal directions.

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

6.4.1 Primary map



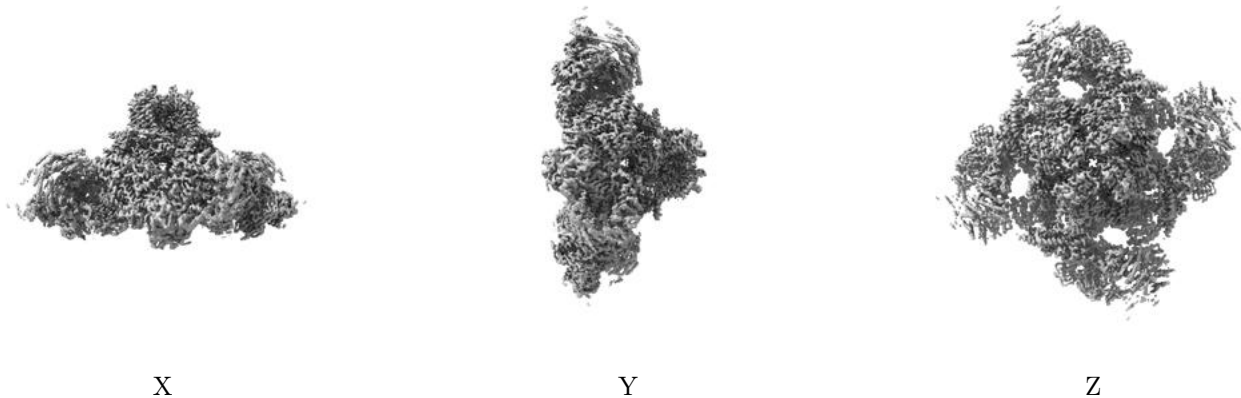
6.4.2 Raw map



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.

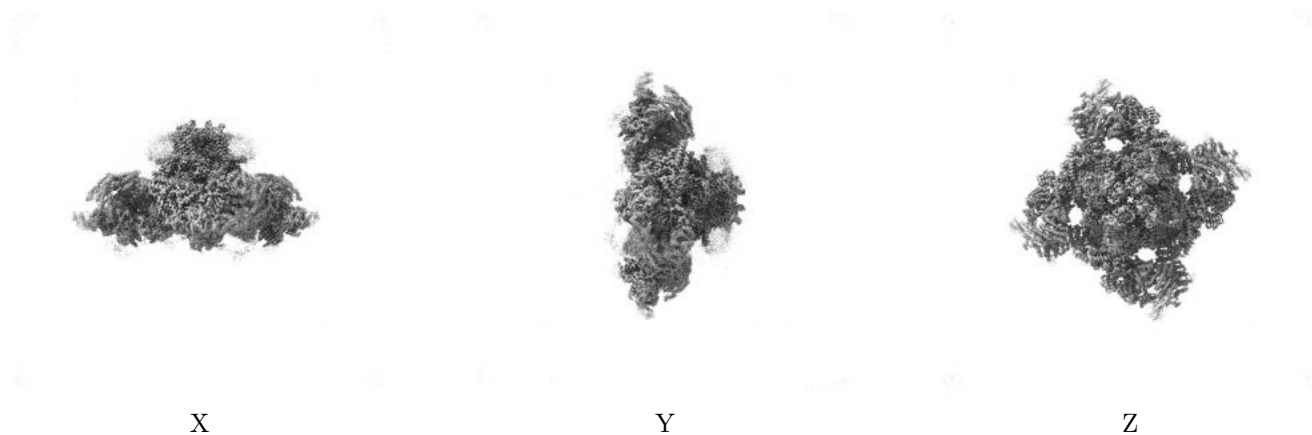
6.5 Orthogonal surface views [i](#)

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.

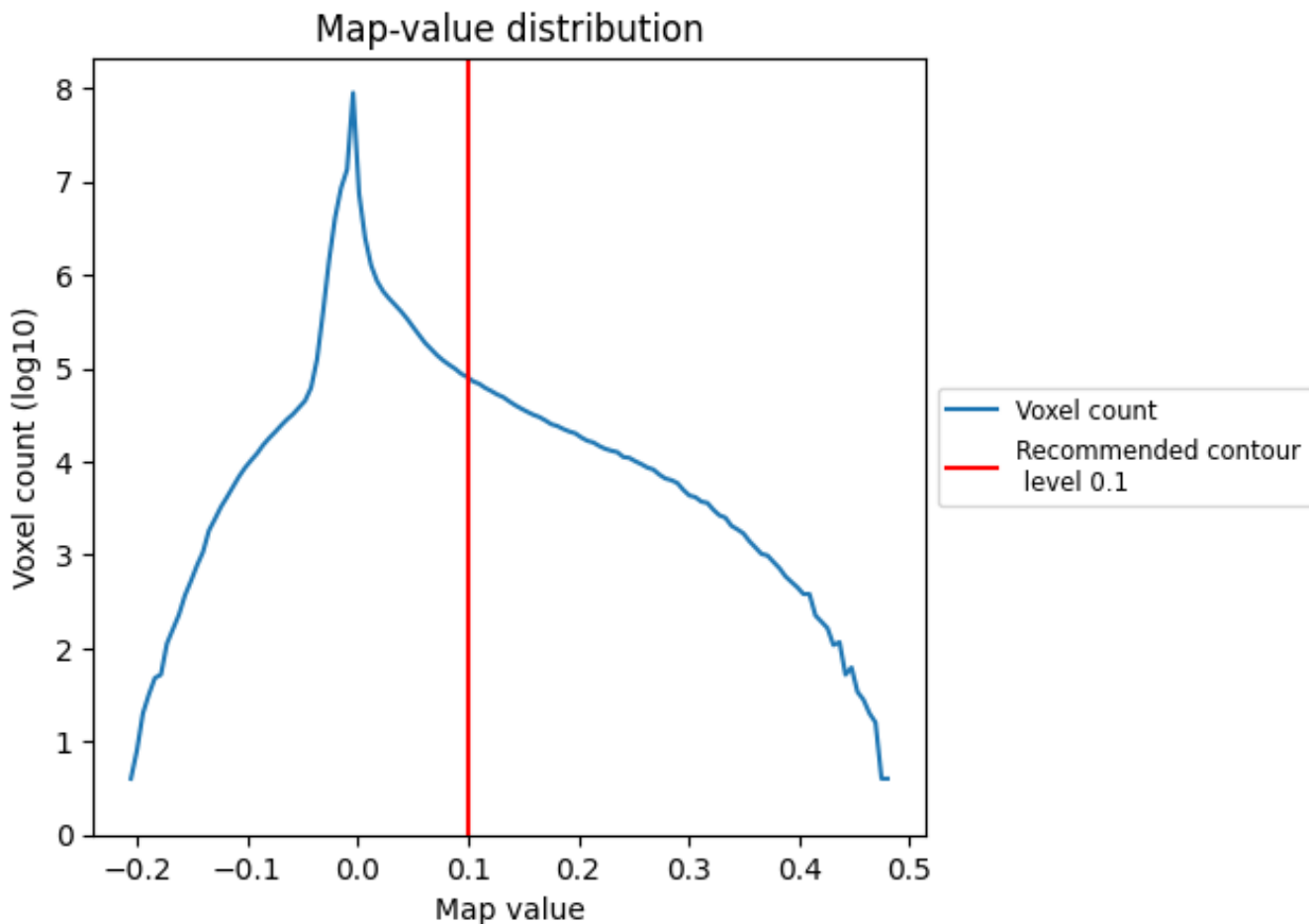
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

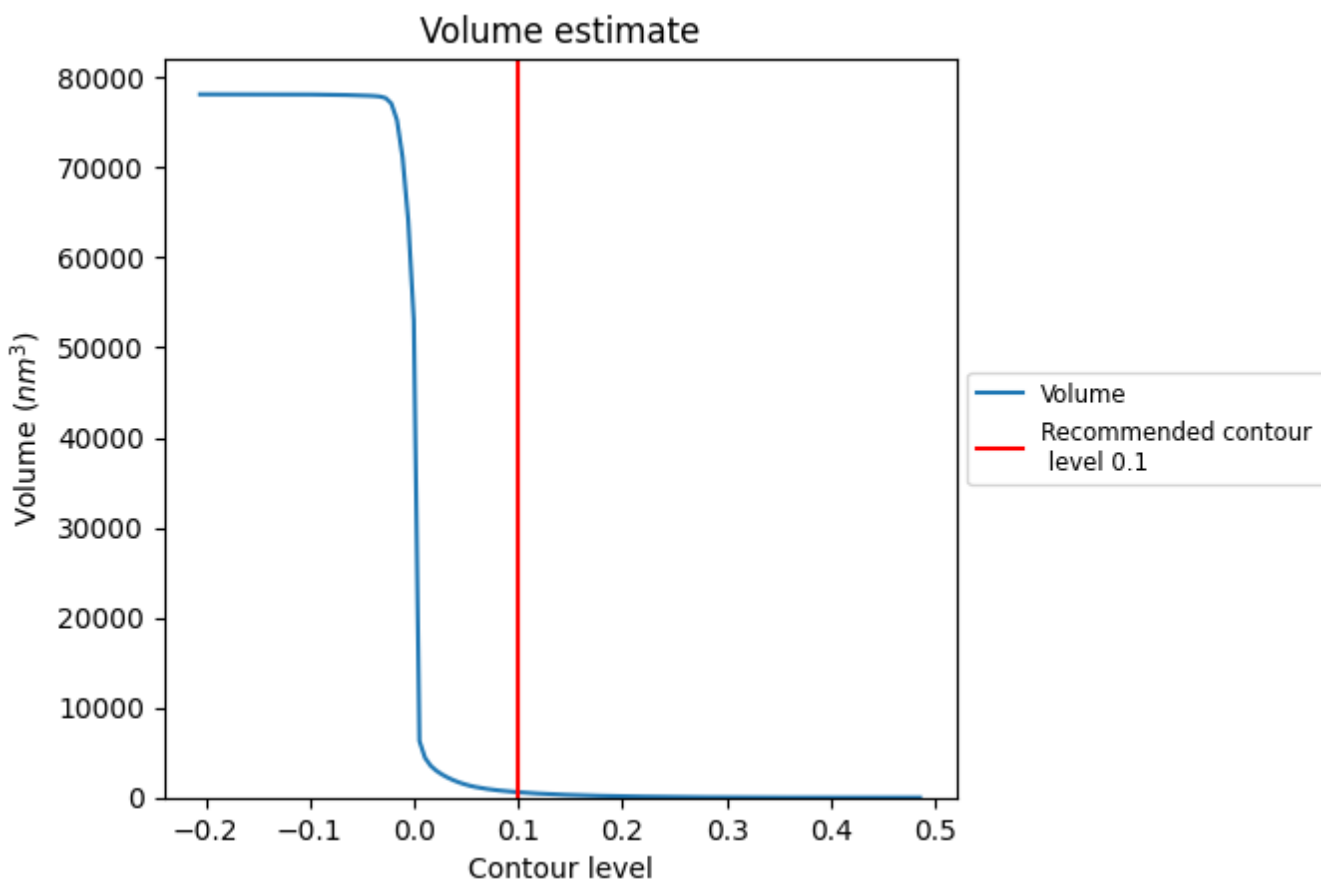
This section contains the results of statistical analysis of the map.

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.

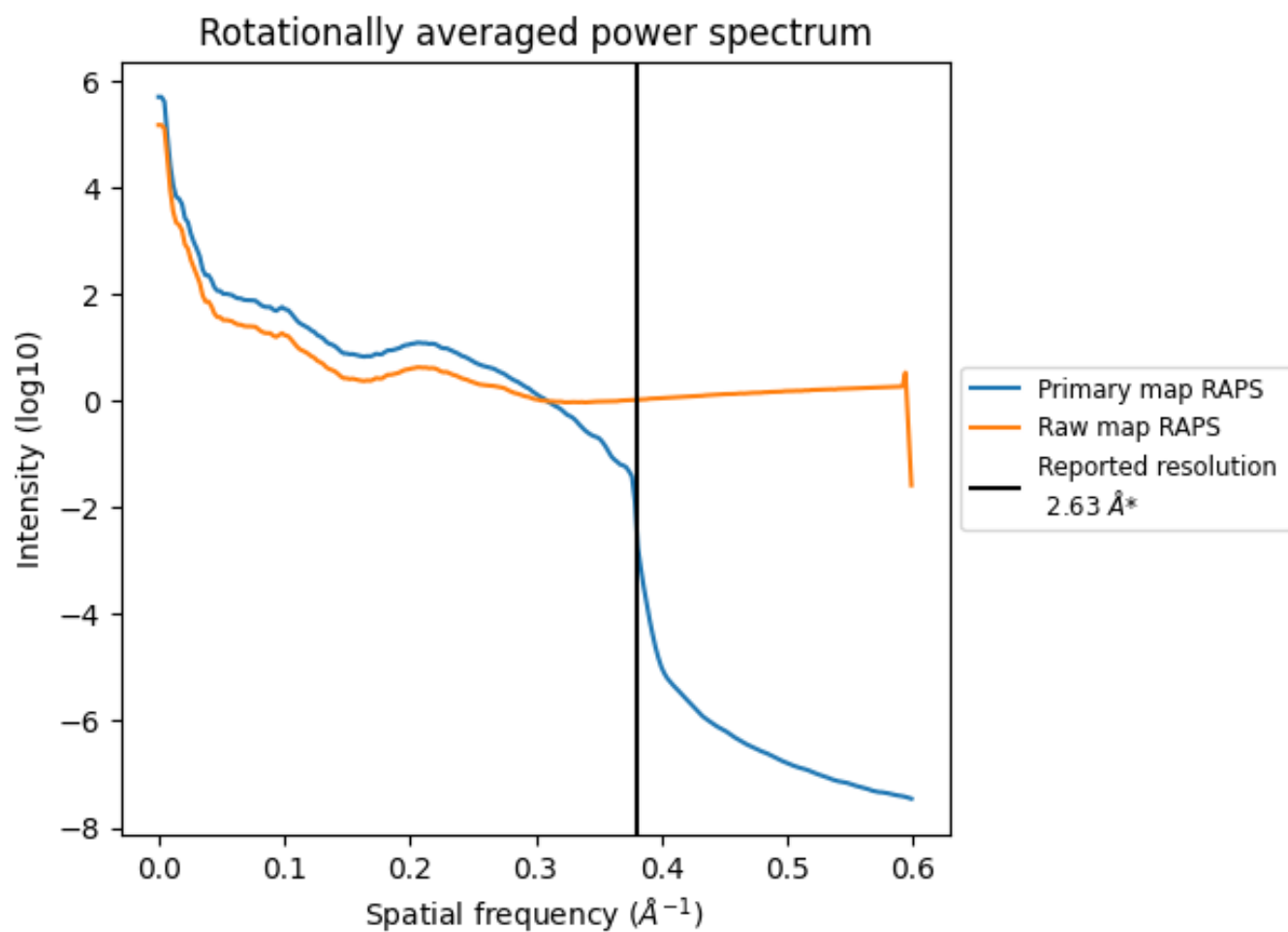
7.2 Volume estimate [\(i\)](#)



The volume at the recommended contour level is 594 nm³; this corresponds to an approximate mass of 536 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

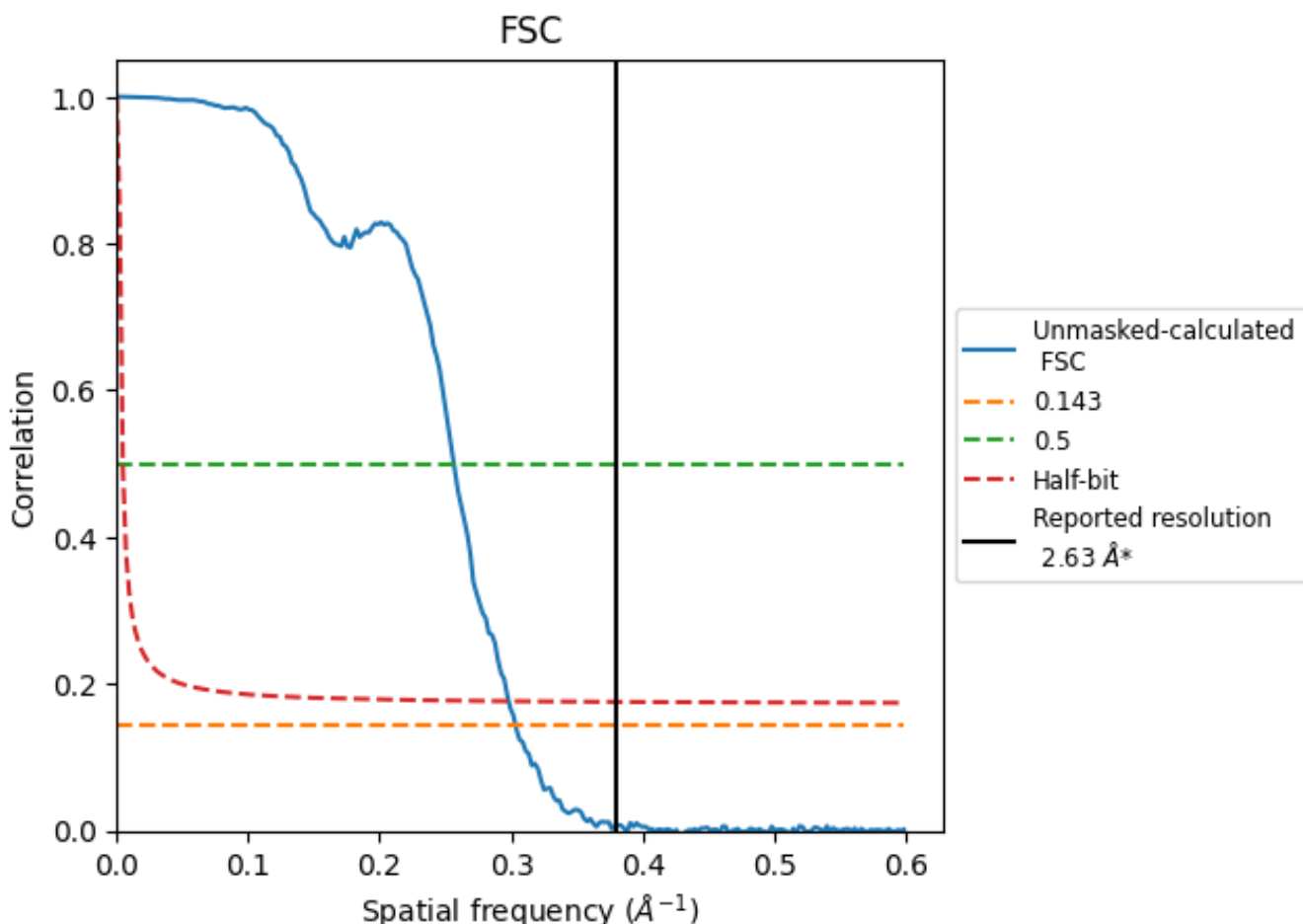


*Reported resolution corresponds to spatial frequency of 0.380 Å⁻¹

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 0.380 Å⁻¹

8.2 Resolution estimates [i](#)

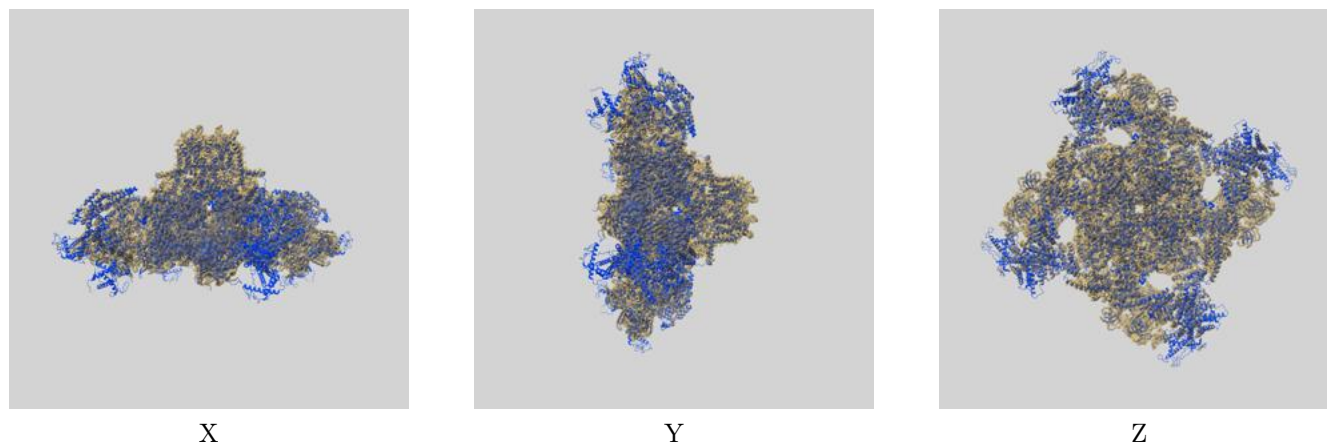
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.63	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	3.30	3.90	3.35

*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 3.30 differs from the reported value 2.63 by more than 10 %

9 Map-model fit [i](#)

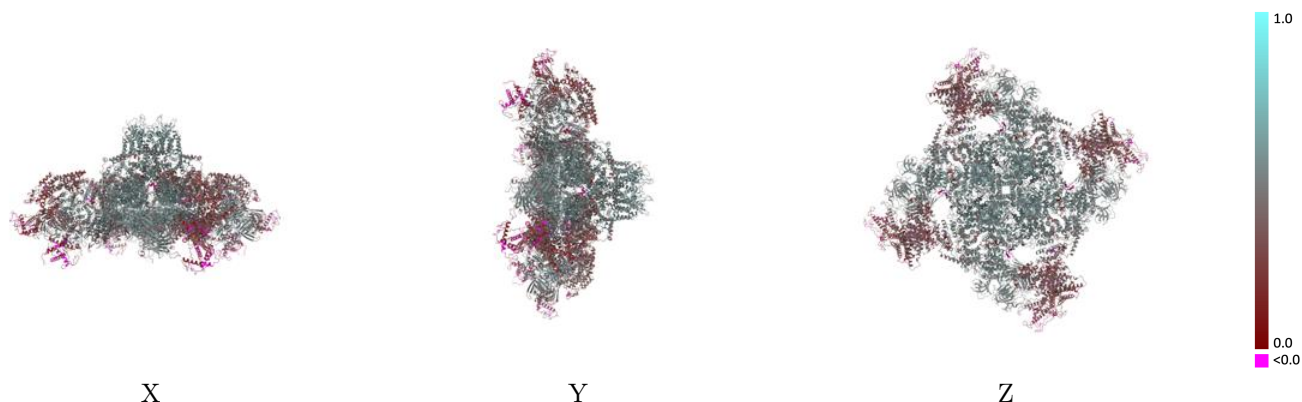
This section contains information regarding the fit between EMDB map EMD-47389 and PDB model 9E1C. Per-residue inclusion information can be found in section 3 on page 6.

9.1 Map-model overlay [i](#)



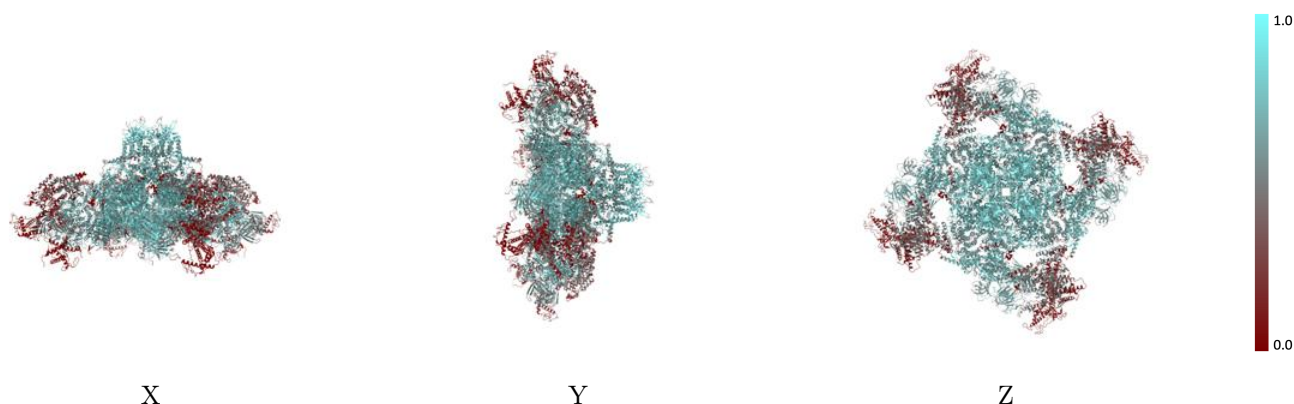
The images above show the 3D surface view of the map at the recommended contour level 0.1 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



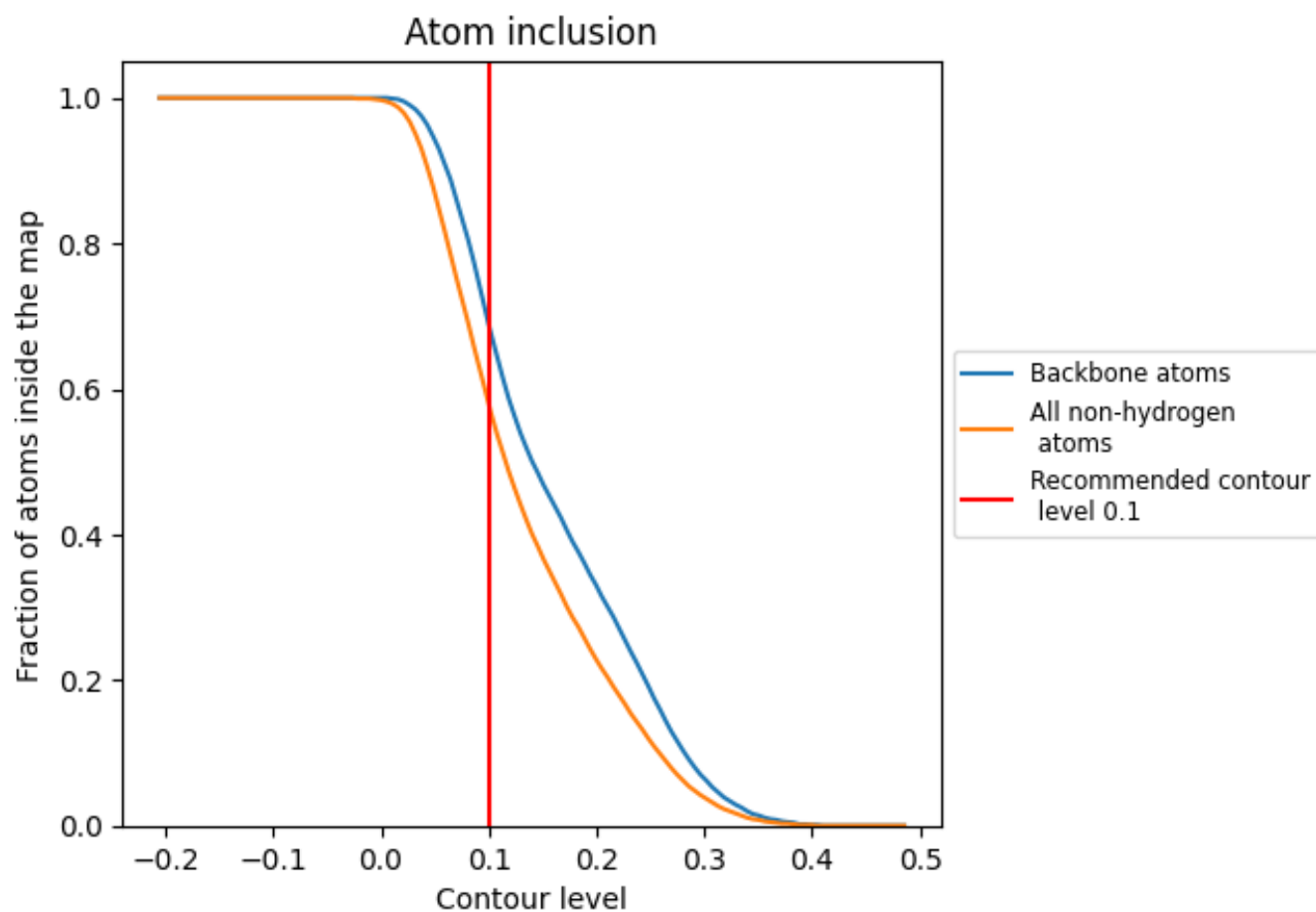
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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, 69% of all backbone atoms, 58% 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	 0.5780	 0.4410
A	 0.5780	 0.4390
B	 0.5780	 0.4400
C	 0.5780	 0.4390
D	 0.5780	 0.4390
E	 0.5880	 0.5100
F	 0.5930	 0.5090
G	 0.5930	 0.5050
H	 0.5890	 0.5050

