

QUARK RESULT

PDB ID	Length	TMscore	RMSD
1B4B_A	72	0.411	8.165
1BG8_A	76	0.327	10.584
1BGF_A	124	0.372	11.116
1C5E_A	95	0.193	12.464
1CF7_B	82	0.506	6.358
1CQA_A	123	0.352	8.021
1DJ7_A	109	0.381	10.659
1DP7_P	76	0.706	3.862
1E2A_A	102	0.766	2.487
1G8Q_A	90	0.358	10.652
1H4L_D	147	0.417	10.889
1HL6_D	143	0.345	12.058
1I27_A	73	0.604	4.425
1IJY_A	122	0.302	13.825
1IM3_D	95	0.327	10.794
1IPI_A	114	0.545	4.846
1IS7_K	85	0.355	9.857
1J1V_A	94	0.808	1.963
1J3W_A	134	0.365	9.302
1JC7_A	129	0.238	12.843
1JO0_A	97	0.344	11.153
1JR8_A	105	0.431	8.37
1JSG_A	111	0.204	16.158
1K3B_A	119	0.346	14.613
1KA8_A	100	0.282	12.566
1KP6_A	79	0.251	10.988
1KPT_A	105	0.281	11.171
1KVD_B	77	0.266	12.363
1LDD_A	71	0.724	2.356
1LMI_A	131	0.327	12.053
1LWB_A	122	0.301	10.678
1MC2_A	122	0.253	13.898
1MHD_A	123	0.252	13.723
1MN8_A	95	0.383	11.171
1MWP_A	96	0.335	14.878
1N13_B	112	0.35	11.381
1N8V_A	101	0.512	6.038
1NLQ_A	105	0.373	8.201
1NQJ_B	114	0.237	15.195
1NZE_A	112	0.824	2.644
1O82_A	70	0.387	9.876
1OCZ_E	109	0.656	3.61
1OK0_A	74	0.217	10.356
1PZW_A	80	0.371	7.512
1QC7_A	101	0.731	4.097
1QYN_D	136	0.367	16.828
1R75_A	110	0.301	12.117
1R77_B	103	0.291	10.744
1R7L_B	109	0.28	13.528
1RKI_A	101	0.438	9.736
1SAU_A	114	0.526	6.232
1SMP_I	100	0.329	11.682

PDB ID	Length	TMscore	RMSD
1T92_A	108	0.536	7.705
1TSF_A	81	0.244	15.06
1U84_A	81	0.724	2.659
1UFY_A	121	0.355	10.289
1UGI_A	83	0.44	8.231
1UZ3_A	102	0.426	13.934
1V14_A	132	0.374	11.09
1V1J_A	150	0.429	9.077
1V54_H	79	0.535	5.855
1VAS_A	137	0.27	13.792
1VCC_A	77	0.342	10.252
1VMH_A	128	0.254	15.295
1VR7_A	120	0.501	10.291
1VYI_A	111	0.296	15.433
1W53_A	84	0.373	12.073
1WCK_A	136	0.179	15.943
1WN2_A	118	0.295	13.721
1WPU_A	147	0.329	15.636
1WXC_B	72	0.335	9.928
1XAW_A	107	0.853	1.929
1XEQ_A	89	0.341	9.028
1XPX_A	144	0.311	10.728
1XZQ_B	117	0.341	12.731
1Y71_B	112	0.395	11.358
1Y96_B	85	0.326	14.379
1YKU_A	132	0.626	4.253
1YO3_A	85	0.559	5.405
1ZKE_A	81	0.413	7.312
2AIB_A	98	0.389	12.279
2AXW_A	134	0.223	19.596
2AYD_A	76	0.298	8.702
2B91_A	129	0.275	14.68
2B97_A	70	0.326	9.102
2BHM_A	134	0.396	10.376
2BL7_A	79	0.603	3.373
2BSE_A	107	0.26	13.296
2BT9_A	90	0.412	6.824
2CHH_A	113	0.288	14.328
2CIU_A	123	0.386	16.137
2CZS_A	70	0.335	9.418
2CZV_D	119	0.451	12.661
2D5R_B	116	0.403	13.404
2D68_B	77	0.295	12.16
2E12_A	93	0.314	10.314
2E79_A	108	0.309	12.309
2F5G_A	130	0.231	19.524
2FCW_A	106	0.834	2.034
2FIO_B	113	0.391	11.268
2FYG_A	128	0.213	16.477
2G38_C	76	0.512	6.564
2GJV_A	136	0.379	9.471
2GWF_B	121	0.293	14.243

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PDB ID	Length	TMscore	RMSD
1STM_A	141	0.204	17.413
1T1J_B	125	0.362	12.084
1T6U_A	117	0.471	5.688
2HDI_B	103	0.29	14.637
2HEW_F	128	0.235	16.362
2HL7_A	82	0.726	3.24
2IGP_A	114	0.62	3.573
2IP6_A	87	0.523	4.213
2J4B_B	133	0.373	13.955
2J6B_A	109	0.296	12.04
2J6Z_A	86	0.373	12.047
2J97_A	93	0.242	11.659
2J9W_A	101	0.608	4.093
2JEK_A	140	0.317	11.453
2NML_A	100	0.288	11.411
2O26_A	144	0.276	13.17
2O42_A	138	0.229	21.093
2O4T_A	90	0.628	12.032
2OFC_A	141	0.312	9.128
2OHW_A	128	0.359	11.62
2OQ8_A	150	0.424	12.528
2OQB_B	106	0.346	14.416
2PT7_G	145	0.392	12.135
2Q3T_A	121	0.303	16.773
2QM2_B	116	0.451	7.08
2QSB_A	85	0.675	3.126
2QT7_A	89	0.511	5.062
2QVT_A	113	0.195	14.395
2QYA_A	115	0.505	8.159
2R41_D	104	0.437	11.731
2RCF_B	81	0.306	8.604
2V33_A	91	0.218	13.514
2V94_B	94	0.643	4.2
2VB1_A	129	0.235	13.85
2YX8_A	81	0.576	3.945
3BJO_A	103	0.319	13.199
3BS1_A	103	0.672	3.71
3CHB_D	103	0.301	12.319
1b72A	49	0.912	0.717
1bkrA	108	0.337	10.327
1bq9A	53	0.64	3.239
1c8cA	64	0.262	9.586
1cg5B	141	0.749	2.76
1cy5A	92	0.267	13.904
1dcjA	73	0.311	11.19
1dtjA	74	0.278	11.134
1elwA	117	0.869	1.679
1fadA	92	0.273	14.051
1fo5A	85	0.362	8.909

PDB ID	Length	TMscore	RMSD
2GYB_T	83	0.786	1.844
2H5N_D	123	0.398	12.148
2H8E_A	120	0.341	10.34
1gnuA	117	0.227	13.282
1gyvA	117	0.242	13.164
1hbkA	89	0.253	14.052
1hz6A	64	0.588	3.076
1itpA	68	0.292	10.898
1kpeA	113	0.596	7.28
1kviA	68	0.313	10.429
1no5A	93	0.289	10.581
1of9A	77	0.405	9.24
1orgA	118	0.325	11.843
256bA	106	0.747	3.334
2cr7A	60	0.324	9.84
2f3nA	65	0.433	8.351
1abv	103	0.244	13.149
1acf	125	0.7	3.302
1af7	72	0.427	10.482
1ah9	63	0.417	10.163
1ail	70	0.34	9.555
1ALY	146	0.265	14.125
1aoy	65	0.359	11.259
1csp	67	0.46	7.596
1ctf	68	0.441	5.397
1GB1	56	0.509	3.118
1gpt	47	0.367	7.258
1GYZ	60	0.661	2.401
1I6C	39	0.418	5.169
1kjs	74	0.412	6.747
1lis	131	0.412	13.816
1mla	70	0.345	10.982
1opd	85	0.549	3.632
1pgx	59	0.349	9.436
1r69	61	0.61	7.186
1rnb	109	0.466	8.037
1SAP	66	0.288	12.536
1sro	71	0.324	9.492
1ten	87	0.315	8.495
1tfi	47	0.447	4.106
1tif	59	0.338	9.238
1TIG	88	0.615	3.34
1wap	69	0.284	8.372
2a0b	118	0.23	13.865
2reb	60	0.435	6.689
3GWL	106	0.861	2.477
4icb	76	0.71	2.625
4uex	81	0.807	2.122