

**Table S4. The performance of DockRank on superimposed models.** The local identity between proteins A and A' is the percentage of identical residues in the local alignment (Num\_identical\_residues/local\_alignment\_length\*100%). The global identity between proteins A and A' is the mean of the percentage of identical residues in each protein ( $[\text{Num\_identical\_residues}/\text{length}(A) + \text{Num\_identical\_residues}/\text{length}(A')] / 2 * 100\%$ ). Out of all 51 dimer cases, there are: **section A**) 37% (19/51) cases for which all superimposed models are near-native (L-RMSD  $\leq 10$  Å); **section B**) 37% (19/51) cases for which at least one superimposed model is near-native and at least one model is incorrect; and **section C**) 25% (13/51) cases for which none of the superimposed models are near-native. The five bold cases in section C) are cases with reliable interface predictions from PS-HomPPI.

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
<b>A. Cases with at least one near-native model (L-RMSD <math>\leq 10</math> angstroms) and one incorrect model (L-RMSD <math>&gt; 10</math> Å)</b>								
1KAC	2j12	A:B	39.15	100	39.07	97.27	0.206	3.00
1KAC	2wbw	A:B	39.15	100	39.07	96.46	0.059	46.76
1AK4	1m9x	F:G	100	96.32	99.7	92.67	0.162	7.34
1AK4	1m9x	B:C	100	96.32	99.7	92.67	0.162	7.45
1AK4	1m9x	A:D	100	96.32	99.7	92.67	0.149	3.39
1AK4	4dga	B:D	98.78	93.33	98.48	89.13	0.146	5.72
1AK4	1m9x	E:H	100	96.32	99.7	92.67	0.146	3.46
1AK4	4dga	A:C	98.78	93.33	98.48	89.13	0.143	3.31
1AK4	1m9d	A:D	100	94.12	99.7	90.55	0.143	3.93
1AK4	4dge	A:C	98.17	93.33	97.87	89.13	0.141	11.60
1AK4	1m9d	B:C	100	94.12	99.7	90.55	0.136	6.65
1AK4	1m9x	A:G	100	96.32	99.7	92.67	0.070	64.97
1AVX	3i29	A:B	81.17	46.86	81.17	46.19	0.160	5.46
1AVX	2qyi	A:B	81.17	46.29	81.17	46.1	0.157	6.40
1AVX	2qyi	C:D	81.17	46.29	81.17	46.1	0.151	6.12
1AVX	2qyi	A:D	81.17	46.29	81.17	46.1	0.076	44.06
1B6C	3mdy	B:A	100	67.17	99.08	66.27	0.183	4.94
1B6C	3mdy	D:C	100	67.17	99.08	66.27	0.178	5.09
1B6C	3h9r	B:A	100	66.36	99.08	64.55	0.172	2.51
1B6C	3mdy	D:A	100	67.17	99.08	66.27	0.086	69.38
1B6C	3mdy	B:C	100	67.17	99.08	66.27	0.086	69.59
1EAW	4dg4	D:F	41.18	100	42.21	100	0.229	2.07
1EAW	4dg4	B:C	41.18	100	42.21	100	0.229	2.25
1EAW	4dg4	A:E	41.18	100	42.21	100	0.229	2.12
1EAW	4dg4	G:H	41.18	100	42.21	100	0.229	2.23
1EAW	3fp6	E:I	39.92	100	41.01	100	0.229	2.10
1EAW	3fp8	E:I	39.5	100	40.58	100	0.229	2.15
1EAW	4dg4	G:F	41.18	100	42.21	100	0.060	30.11
1EAW	4dg4	D:H	41.18	100	42.21	100	0.060	30.20
1EAW	4dg4	A:C	41.18	100	42.21	100	0.060	30.24
1EAW	4dg4	B:E	41.18	100	42.21	100	0.060	29.94
1D6R	1tab	E:I	98.65	81.03	99.32	69.17	0.250	2.16
1D6R	1sbw	A:I	98.65	53.57	99.32	34.36	0.241	2.67
1D6R	3myw	B:I	82.06	67.24	82.62	60.7	0.237	1.86
1D6R	1smf	E:I	98.65	59.09	99.32	40.75	0.237	2.23
1D6R	2g81	E:I	98.65	81.03	99.32	68.83	0.237	2.57
1D6R	3ru4	T:B	98.65	81.03	99.32	79.04	0.237	2.35
1D6R	1tx6	A:I	81.17	41.3	81.72	23.98	0.228	2.36
1D6R	1tx6	D:J	81.17	41.3	81.72	23.98	0.207	6.92

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
1D6R	1g9i	E:I	98.65	63.64	99.32	43.89	0.137	21.41
1D6R	3myw	A:I	82.06	67.24	82.62	60.7	0.116	21.35
1FQJ	1fqk	C:D	91.46	95.65	90.19	94.52	0.161	4.40
1FQJ	1fqk	A:B	91.46	95.65	90.19	94.52	0.161	4.38
1FQJ	2ihb	A:B	67.2	41.38	66.05	33.73	0.160	4.59
1FQJ	2gtp	B:C	69.75	42.06	68.56	38.2	0.159	4.64
1FQJ	2gtp	A:D	69.75	42.06	68.56	38.2	0.159	4.60
1FQJ	2ik8	C:D	69.84	38.66	68.76	33.73	0.155	4.49
1FQJ	2ik8	A:B	69.84	38.66	68.76	33.73	0.152	4.62
1FQJ	2ik8	C:B	69.84	38.66	68.76	33.73	0.082	56.64
1FQJ	2ik8	A:D	69.84	38.66	68.76	33.73	0.081	56.47
1FQJ	1fqk	A:D	91.46	95.65	90.19	94.52	0.069	61.66
1GPW	2wjz	A:B	99.6	99	99.6	98.75	0.170	2.17
1GPW	2wjz	C:D	99.6	99	99.6	98.75	0.167	1.92
1GPW	2wjz	E:F	99.6	99	99.6	98.75	0.167	2.34
1GPW	2wjz	C:B	99.6	99	99.6	98.75	0.066	58.33
1IBR	2bku	A:B	98.84	40.82	94.22	30.91	0.175	4.10
1IBR	2bku	C:D	98.84	40.82	94.22	30.91	0.175	4.12
1IBR	3ea5	A:B	98.91	40.82	91.58	30.91	0.172	4.35
1IBR	3ea5	C:D	98.91	40.82	91.58	30.91	0.167	4.59
1IBR	1wa5	A:B	99.42	24.73	94.5	9.36	0.065	53.47
1SBB	1l0x	C:D	99.58	50	99.79	51.38	0.119	2.75
1SBB	1l0x	A:B	99.58	50	99.79	51.38	0.119	2.46
1SBB	1l0y	C:D	99.58	50	99.37	51.38	0.119	2.66
1SBB	1l0y	A:B	99.58	50	99.37	51.38	0.119	2.65
1SBB	1jck	C:D	99.58	65.42	99.79	65.69	0.118	4.48
1SBB	1jck	A:B	99.58	65.42	99.79	65.69	0.118	4.48
1SBB	1l0y	A:D	99.58	50	99.37	51.38	0.073	81.40
1SBB	1l0x	A:D	99.58	50	99.79	51.38	0.072	81.35
1SBB	1jck	C:B	99.58	65.42	99.79	65.69	0.071	81.52
1SBB	1jck	A:D	99.58	65.42	99.79	65.69	0.071	81.52
1R0R	3qtl	A:D	99.27	35.71	99.27	16.47	0.271	2.07
1R0R	1yu6	A:C	100	31.67	99.82	23.77	0.253	3.68
1R0R	1yu6	B:D	100	31.67	99.82	23.77	0.235	1.89
1R0R	3qtl	C:D	99.27	35.71	99.27	16.47	0.061	27.90
1UDI	2j8x	C:D	41.92	100	41.83	99.4	0.258	2.35
1UDI	2j8x	A:B	41.92	100	41.83	99.4	0.258	2.18
1UDI	1eui	B:D	46.73	100	43.86	99.4	0.253	2.03
1UDI	1lqm	A:B	46.73	100	43.76	99.4	0.253	1.98
1UDI	1lqg	B:D	46.73	100	43.76	99.4	0.253	2.06
1UDI	1eui	A:C	46.73	100	43.86	99.4	0.253	2.03
1UDI	1ugh	E:I	42.08	100	41.25	99.4	0.253	2.05
1UDI	1lqg	A:C	46.73	100	43.76	99.4	0.253	2.03
1UDI	2j8x	C:B	41.92	100	41.83	99.4	0.086	53.65
1UDI	1lqm	C:H	46.73	100	43.76	99.4	0.065	39.33
2HLE	2wo2	A:B	44.44	98.58	43.24	94.71	0.204	9.50
2HLE	1kgy	A:E	44.44	99.28	43.72	98.22	0.203	9.34
2HLE	1kgy	C:G	44.44	99.28	43.72	98.22	0.200	9.42

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
2HLE	1kgy	D:H	44.44	99.28	43.72	98.22	0.197	9.43
2HLE	1kgy	B:F	44.44	99.28	43.72	98.22	0.197	9.49
2HLE	3gxu	A:B	44.75	98.58	45.03	97.89	0.190	10.27
2HLE	1kgy	B:G	44.44	99.28	43.72	98.22	0.070	54.25
2HLE	1kgy	A:H	44.44	99.28	43.72	98.22	0.068	54.03
2HLE	1kgy	D:E	44.44	99.28	43.72	98.22	0.067	54.43
2HLE	1kgy	C:F	44.44	99.28	43.72	98.22	0.067	54.22
2NZ8	1foe	C:D	24.73	99.43	26.99	97.25	0.151	12.62
2NZ8	1foe	E:F	24.73	99.43	26.99	97.25	0.150	11.89
2NZ8	1foe	G:H	24.73	99.43	26.99	97.25	0.148	12.58
2NZ8	1foe	A:B	24.73	99.43	26.99	97.25	0.147	11.92
2NZ8	1kzg	C:D	39.56	71.51	38.2	69.02	0.146	3.95
2NZ8	1kz7	C:D	38.61	71.51	37.28	69.02	0.146	4.00
2NZ8	1kzg	A:B	39.56	71.51	38.2	69.02	0.146	4.17
2NZ8	1kz7	A:B	38.61	71.51	37.28	69.02	0.146	4.21
2NZ8	1foe	A:D	24.73	99.43	26.99	97.25	0.068	67.20
2NZ8	1foe	E:H	24.73	99.43	26.99	97.25	0.068	67.78
7CEI	3u43	A:B	66.67	74.62	64.19	73.22	0.229	4.71
7CEI	2wpt	A:B	67.05	70	68.21	68.69	0.227	7.45
7CEI	3gjn	A:B	60.92	99.23	61.27	94.98	0.218	3.44
7CEI	3gkl	D:B	62.07	99.23	62.43	94.98	0.208	4.10
7CEI	3gkl	C:A	62.07	99.23	62.43	94.98	0.208	4.02
7CEI	3gjn	D:C	60.92	99.23	61.27	94.98	0.207	3.32
7CEI	3gkl	C:B	62.07	99.23	62.43	94.98	0.089	39.12
7CEI	3gjn	D:B	60.92	99.23	61.27	94.98	0.089	38.96
7CEI	3gkl	D:A	62.07	99.23	62.43	94.98	0.083	38.89
7CEI	3gjn	A:C	60.92	99.23	61.27	94.98	0.083	38.82
1Z5Y	1vrs	D:A	30	99.17	17.01	91.54	0.197	1.99
1Z5Y	1vrs	E:B	30	99.17	17.01	91.54	0.197	1.42
1Z5Y	1vrs	F:C	30	99.17	17.01	91.54	0.183	2.18
1Z5Y	1vrs	D:C	30	99.17	17.01	91.54	0.089	58.17
2B42	1t6g	B:D	95.01	46.71	97.5	42.28	0.183	1.58
2B42	3hd8	C:D	84.92	99.46	85.6	99.46	0.183	0.84
2B42	1t6g	A:C	95.01	46.71	97.5	42.28	0.183	1.56
2B42	3hd8	A:B	84.92	99.46	85.6	99.46	0.183	0.78
2B42	3hd8	A:D	84.92	99.46	85.6	99.46	0.070	53.70
2B42	3hd8	C:B	84.92	99.46	85.6	99.46	0.070	53.63
2I25	2i26	N:L	88.29	100	83.86	100	0.178	2.65
2I25	2i26	P:Q	88.29	100	83.86	100	0.169	3.29
2I25	2i26	O:M	88.29	100	83.86	100	0.166	2.98
2I25	1sq2	N:L	65.18	100	64.6	100	0.158	15.23
2I25	1t6v	O:M	65.18	100	64.6	100	0.153	15.27
2I25	1t6v	N:L	65.18	100	64.6	100	0.150	16.76
2I25	1t6v	N:M	65.18	100	64.6	100	0.068	41.05
2PCC	2bcn	A:B	100	99.07	99.49	99.07	0.155	1.82
2PCC	1u75	A:B	100	63.37	99.49	60.4	0.116	6.89
2PCC	2pcb	A:B	100	63.37	99.49	60.4	0.110	6.85
2PCC	2bcn	C:B	100	99.07	99.49	99.07	0.077	23.01

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
<b>B. Cases with superimposed models that are all near-native (L-RMSD <math>\leq</math> 10 Å)</b>								
1CGI	1z7k	A:B	43.97	51.22	43.69	35.69	0.261	2.61
1CGI	1hja	C:I	100	38.1	69.8	29.98	0.259	3.68
1CGI	1cho	F:I	100	38.1	76.73	28.57	0.257	2.74
1CGI	1hja	B:I	100	38.1	76.73	29.98	0.255	2.95
1CGI	1an1	E:I	43.97	50	43.69	21.78	0.252	2.71
1CGI	1ldt	T:L	43.97	50	43.69	21.78	0.252	2.50
1CGI	1tgs	Z:I	42.06	69.09	41.4	67.86	0.250	2.65
1CGI	1cho	G:I	100	38.1	69.8	28.57	0.247	3.27
1AY7	1x1u	A:D	33.75	98.88	26.34	98.88	0.235	2.65
1AY7	1x1u	B:E	33.75	98.88	26.34	98.88	0.235	2.43
1AY7	1bgs	C:G	33.75	98.88	26.34	98.88	0.231	2.54
1AY7	1x1u	C:F	33.75	98.88	26.34	98.88	0.230	2.85
1AY7	1bgs	B:F	33.75	98.88	26.34	98.88	0.230	2.53
1AY7	1brs	A:D	33.75	98.88	26.34	98.88	0.230	2.39
1AY7	1brs	B:E	33.75	98.88	26.34	98.88	0.229	2.54
1AY7	1brs	C:F	33.75	98.88	26.34	98.88	0.229	2.81
1AY7	1bgs	A:E	33.75	98.88	26.34	98.88	0.229	2.24
1AY7	1b27	B:E	33.75	98.88	26.34	98.33	0.228	2.49
1DFJ	1z7x	X:W	70.16	77.36	68.8	76.77	0.202	3.81
1DFJ	2q4g	X:W	70.16	77.36	68.8	76.77	0.197	3.72
1DFJ	2q4g	Z:Y	70.16	77.36	68.8	76.77	0.196	4.15
1DFJ	1z7x	Z:Y	70.16	77.36	68.8	76.77	0.190	4.20
1ACB	4b2a	A:B	40.95	98.48	40.69	95.67	0.243	2.40
1ACB	4b2b	A:B	40.52	98.57	40.26	97.88	0.235	2.48
1ACB	4b2c	A:B	40.95	98.57	40.69	97.88	0.235	2.64
1ACB	4b1t	A:B	41.38	98.57	41.12	98.57	0.235	2.69
1ACB	4b2c	C:D	40.95	98.57	40.69	97.88	0.226	2.38
1ACB	4b2a	C:D	40.95	98.48	40.69	95.67	0.219	1.96
1ACB	4b2b	C:D	40.52	98.57	40.26	97.88	0.219	2.34
1ACB	4b1t	C:D	41.38	98.57	41.12	98.57	0.219	2.35
1BKD	3cf6	E:R	25.81	57.49	12.96	57.66	0.203	5.29
1EFN	1avz	B:C	76.15	100	82.78	100	0.222	4.49
1GLA	1glb	G:F	99.8	100	99.4	94.64	0.132	1.71
1GLA	1gld	G:F	99.8	100	99.4	94.64	0.132	1.76
1GLA	1glc	G:F	99.8	100	99.4	94.64	0.131	1.75
1GLA	1gle	G:F	99.8	100	99.4	94.64	0.131	1.74
1GRN	2ngr	A:B	99.47	94.3	99.21	87.04	0.141	4.97
1GRN	1tx4	B:A	54.91	94.44	51.83	96.69	0.140	4.95
1GRN	1ow3	B:A	52.69	94.82	51.18	86.22	0.138	4.90
1T6B	1tzn	C:c	92.7	100	85.52	100	0.153	2.94
1T6B	1tzn	L:l	92.7	100	85.52	100	0.153	2.94
1T6B	1tzn	B:b	92.7	100	85.52	100	0.153	2.92
1T6B	1tzn	J:j	92.7	100	85.52	100	0.153	2.92
1T6B	1tzn	E:e	92.7	100	85.52	100	0.153	2.92
1T6B	1tzn	F:f	92.7	100	85.52	100	0.153	2.94

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
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1T6B	1tzn	I:i	92.7	100	85.52	100	0.153	2.94
1T6B	1tzn	K:k	92.7	100	85.52	100	0.153	2.92
1T6B	1tzn	M:m	92.7	100	85.52	100	0.153	2.92
1T6B	1tzn	D:d	92.7	100	85.52	100	0.153	2.92
1XD3	3kw5	A:B	52.73	97.18	53.6	95.28	0.201	3.75
1XD3	3ifw	A:B	52.73	97.18	53.6	95.28	0.201	3.62
1XD3	3kvf	A:B	52.27	97.18	53.13	95.28	0.192	3.75
1YVB	3ima	A:B	36.44	33.87	36.35	21.26	0.215	3.68
1YVB	3ima	C:D	36.44	33.87	36.35	21.26	0.213	3.52
1Z0K	1z0j	A:B	40.37	57.58	37.58	36.76	0.265	2.50
2HRK	2hsm	A:B	100	100	93.48	92.21	0.185	1.98
2SNI	1y3d	E:I	99.63	92.19	96.95	91.48	0.208	1.62
2SNI	1mee	A:I	85.82	37.1	85.98	35.66	0.203	4.64
2SNI	1y4d	E:I	99.63	90.62	96.95	89.92	0.200	2.37
2SNI	1y4a	E:I	99.63	90.62	96.95	89.92	0.198	3.43
2SNI	1sib	E:I	99.64	35.48	99.82	32.63	0.198	3.91
2SNI	1cse	E:I	69.45	37.1	69.7	34.12	0.198	4.50
2SNI	1sbn	E:I	99.64	37.1	99.82	34.12	0.196	3.83
2SNI	2sec	E:I	69.45	37.1	69.7	34.12	0.190	4.52
1HE1	1g4u	S:R	26.98	100	17.41	98.09	0.155	5.57
1IRA	4dep	B:A	100	29.93	98.29	29.09	0.246	2.78
1IRA	4dep	E:D	100	29.93	98.29	29.09	0.243	2.37
1KTZ	3kfd	D:H	76.79	99.04	76.79	93.45	0.165	1.34
1KTZ	3kfd	B:F	76.79	99.04	76.79	93.45	0.165	1.33
1KTZ	3kfd	A:E	76.79	99.04	76.79	93.45	0.160	3.83
1KTZ	3kfd	C:G	76.79	99.04	76.79	93.45	0.160	3.86
1MAH	2x8b	A:B	87.22	100	84.58	100	0.250	2.59
1MAH	1f8u	A:B	87.22	100	84.58	100	0.250	2.60
1MAH	1b41	A:B	87.38	100	87.87	100	0.250	2.66
1PPE	2sta	E:I	64.86	96.55	65.15	96.55	0.380	1.82
1PPE	1h9i	E:I	81.17	66.67	81.72	56.04	0.365	1.80
1PPE	2stb	E:I	64.86	89.66	65.15	89.66	0.363	1.79
1PPE	1h9h	E:I	78.35	66.67	80.31	54.72	0.361	1.78
1PPE	1f2s	E:I	98.65	65.52	99.32	66.69	0.346	1.76
1PPE	1mct	A:I	80.27	57.14	80.82	56.15	0.345	1.80
1PPE	2btc	E:I	98.65	89.66	99.32	89.66	0.342	1.74

### C. Cases with superimposed models that are all incorrect (L-RMSD > 10 Å)

<b>1H1V</b>	1esv	A:S	99.73	44.55	98.94	22.96	0.097	54.41
<b>1H1V</b>	1p8z	A:G	100	44.55	99.2	21.38	0.097	54.58
<b>1H1V</b>	1rgi	A:G	100	33.53	99.2	24.08	0.097	53.66
<b>1H1V</b>	1eqy	A:S	99.73	44.55	98.94	22.96	0.096	54.40
<b>1H1V</b>	1mdu	E:D	99.73	44.55	98.94	22.96	0.096	53.82
<b>1H1V</b>	1t44	A:G	100	44.55	99.6	20.03	0.096	54.47
<b>1H1V</b>	3cjc	A:G	100	44.55	99.2	22.96	0.096	54.12

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
1H1V	3cjb	A:G	100	44.55	99.2	22.96	0.096	54.39
1H1V	1mdu	B:A	99.73	44.55	98.94	22.96	0.096	53.99
1H1V	1mdu	E:A	99.73	44.55	98.94	22.96	0.063	45.94
2OT3	2efe	C:D	37.22	39.38	26.22	36.5	0.179	17.16
2OT3	2efd	C:D	37.22	39.38	26.22	36.5	0.176	16.87
2OT3	2efd	A:B	37.22	39.38	26.22	36.5	0.173	16.94
2OT3	2efc	A:B	37.22	39.38	26.22	36.5	0.173	17.11
2OT3	2efc	C:D	37.22	39.38	26.22	36.5	0.173	17.20
2OT3	2efe	A:B	37.22	39.38	26.22	36.5	0.173	17.11
2OT3	2efh	C:D	36.67	39.38	25.83	36.5	0.164	17.30
2OT3	2efh	A:B	36.67	39.38	25.83	36.5	0.164	17.23
2OT3	2efc	C:B	37.22	39.38	26.22	36.5	0.066	47.70
2OT3	2efe	C:B	37.22	39.38	26.22	36.5	0.066	47.82
2CFH	2j3t	A:B	98.11	52.53	91.91	55.12	0.214	13.39
2CFH	2c0j	A:B	98.09	52.53	96.25	54.96	0.214	13.65
1ZHI	1zbx	A:B	93.46	88.28	94.93	89.92	0.159	11.48
2AJF	3sci	B:F	100	98.91	99.5	89.15	0.149	12.89
2AJF	3sci	A:E	100	98.91	99.5	89.15	0.144	12.27
1EWY	1krh	B:A	25	32.22	21.9	19.08	0.000	57.37
1EWY	1krh	A:B	25	32.22	21.9	19.08	0.000	60.09
1GCQ	1g83	B:A	32.76	46.15	10.26	12.59	0.071	53.71
1GCQ	1g83	A:B	32.76	46.15	10.26	12.59	0.065	48.07
1GHQ	3oxu	A:E	100	27.74	95.72	29.35	0.116	29.92
1GHQ	3oxu	B:F	100	27.74	95.72	29.35	0.115	29.88
1GHQ	3rj3	C:D	100	27.74	95.72	29.35	0.109	30.00
1GHQ	3oxu	C:D	100	27.74	95.72	29.35	0.109	29.82
1GHQ	2xqw	A:C	100	26.67	100	27.38	0.109	30.27
1NW9	1i4o	B:D	32.08	100	32.75	10.26	0.091	50.97
1NW9	1i51	A:E	36.99	40.24	29.4	30.6	0.090	50.70
1NW9	1i4o	A:C	32.08	100	32.75	10.26	0.088	50.57
1NW9	1i51	B:E	35.16	40.24	21.85	30.6	0.087	50.37
1NW9	1i51	C:F	36.99	40.24	29.4	30.6	0.087	32.61
1NW9	1i51	D:F	35.16	40.24	21.85	30.6	0.085	32.95
1NW9	1i3o	C:F	39.6	40.24	29.05	30.13	0.082	55.41
1NW9	1i3o	A:E	39.6	40.24	29.05	30.13	0.077	55.50
2UUY	1toc	B:R	35.04	31.43	37.14	14.58	0.000	39.00
2UUY	1toc	F:T	35.04	31.43	37.14	14.58	0.000	39.39
2UUY	1toc	H:U	35.04	31.43	37.14	14.58	0.000	19.99
2UUY	1toc	D:S	35.04	31.43	37.14	14.58	0.000	20.14
2UUY	1toc	H:S	35.04	31.43	37.14	14.58	0.000	40.05
2UUY	1toc	D:T	35.04	31.43	37.14	14.58	0.000	90.68
1QA9	1a6p	A:B	44.79	26.92	34.88	17.31	0.166	26.01
1QA9	1a64	A:B	44.79	26.92	34.17	16.96	0.159	25.73
1QA9	1a6p	B:A	44.79	26.92	34.88	17.31	0.159	26.00
1QA9	1a7b	D:C	44.79	26.92	34.17	16.96	0.148	27.46
1QA9	1a7b	A:B	44.79	26.92	34.17	16.96	0.143	26.66

Case ID	Template PDB ID	Template Rec:Lig	Local Identity		Global	Global	DockRank Score	L-RMSD
			Score AA' (%)	Local Identity Score BB' (%)	Identity Score AA' (%)	Identity Score BB' (%)		
1QA9	1a7b	C:A	44.79	26.92	34.17	16.96	0.125	11.32
1QA9	1a7b	B:D	44.79	26.92	34.17	16.96	0.108	11.20
1QA9	1a7b	B:C	44.79	26.92	34.17	16.96	0.039	36.22
1QA9	1a7b	D:A	44.79	26.92	34.17	16.96	0.039	36.24
1QA9	1a7b	C:B	44.79	26.92	34.17	16.96	0.038	36.76
1GHQ	1w2s	A:B	100	98.44	97.88	92.83	0.098	49.87
1GHQ	3rj3	C:E	100	27.74	95.72	29.35	0.076	36.57
1GHQ	3oxu	B:D	100	27.74	95.72	29.35	0.076	63.94
1GHQ	2xqw	B:C	100	26.67	100	27.38	0.076	75.28
1GHQ	3oxu	A:D	100	27.74	95.72	29.35	0.076	79.90
1OPH	2d26	A:C	98.21	37.6	90.44	39.36	0.097	93.95
1KXQ	1kxt	A:B	99.4	64.84	99.4	67.26	0.173	47.03
1KXQ	1kxt	C:D	99.4	64.84	99.4	67.26	0.172	47.24
1KXQ	1kxt	E:F	99.4	64.84	99.4	67.26	0.172	47.81
1KXQ	1kxv	B:D	99.4	58.54	99.4	59.76	0.162	51.77
1KXQ	1kxv	A:C	99.4	58.54	99.4	59.76	0.157	52.49
1KXQ	1kxv	B:C	99.4	58.54	99.4	59.76	0.069	41.10