

Table S2. ClusPro2-BM3_31 set. This set contains 31 docking cases, for which i) ClusPro 2.0 returned at least one near-native conformation (a docked conformation with $L\text{-}RMSD \leq 10$ Å), and ii) PS-HomPPI is able to generate interface predictions for the unbound component proteins.

Case ID	Bound Complex PDB ID	Case ID	Bound Complex PDB ID	Case ID	Bound Complex PDB ID	Case ID	Bound Complex PDB ID
1	1MAH	11	1F51	21	2SNI	31	1ACB
2	1B6C	12	1UDI	22	2B42		
3	1HE1	13	1DFJ	23	1PPE		
4	1AVX	14	2HRK	24	1R0R		
5	1XD3	15	1IQD	25	1RLB		
6	1CGI	16	1ML0	26	1KXQ		
7	1GPW	17	1Z5Y	27	1HIA		
8	2PCC	18	1MLC	28	1NW9		
9	1YVB	19	2NZ8	29	1EAW		
10	1EWY	20	1E6J	30	1EZU		