Supplementary Table 1:

Model	1	2	3	4	5	6	7	8	9	10
1	0									
2	1,86	0								
3	3,45	2,24	0							
4*	32,65	36,78	29,36	0						
5	6,75	7,98	5,34	6,32	0					
6	4,52	5,67	4,87	5,73	5,31	0				
7	7,87	6,98	5,76	8,65	4,32	2,32	0			
8	3,68	3,28	4,56	2,98	6,79	3,85	4,23	0		
9	4,56	4,37	5,98	3,67	6,54	4,73	4,79	2,98	0	
10*	52,12	48,56	51,62	69,94	56,98	58,86	56,78	52,45	48,31	0

* Models with higher divergence

Supplementary Table 1. Similarity of Top 10 predicted complex models measured by RMSD of amino acids main chain.