

Target		Template		Most accurate cluster										
				Size (%)	Min RMSD (Å)	Best model by S			Best model by S <sub>SAXS</sub>			Best model by S <sub>DOPE</sub>		
PDB	# Rigid bodies	PDB	Seq Id%			RMSD (Å)	$\Delta r$ (Å)	$\Delta\alpha$ (°)	RMSD (Å)	$\Delta r$ (Å)	$\Delta\alpha$ (°)	RMSD (Å)	$\Delta r$ (Å)	$\Delta\alpha$ (°)
1ha0	2	2viuA	100	52	2.2	2.9	0.5	0.8	2.7	0.7	1.2	2.9	0.5	0.8
1gv2	2	1h88C	100	37	1.2	2.8	5.0	24.6	2.6	4.3	25.0	5.0	3.8	45.2
1mdtA	2	1ddtA	100	17	1.4	1.5	1.5	6.7	1.9	2.0	6.8	1.5	1.5	6.7
1a62	2	1pv0A	100	53	1.3	1.6	3.2	19.2	1.5	2.7	19.2	1.4	1.2	12.2
1iknA	2	1svcP	46	46	8.3	12.6	14.1	166.	12.7	15.0	160.	12.6	14.1	171.8
1ko9A	3	1fn7A	100	27	1.8	2.1	1.7	10.8	2.4	2.6	15.3	1.8	1.0	8.7
1o0vA	3	1a6tB	33	17	27.2	50.6	58.2	172.	46.9	55.2	164.	50.6	58.2	172.5
1exmA	3	1f60A	35	24	25.4	29.8	37.8	157.	29.6	37.6	149.	25.4	31.8	117.8
1cb6	4	1h76A	63	27	4.5	4.5	3.6	11.5	23.9	27.8	98.3	8.7	5.4	18.8