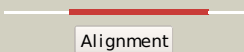

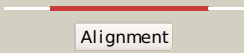







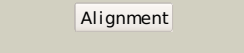

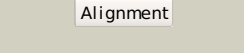

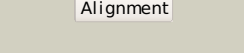



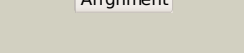

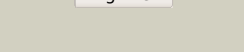








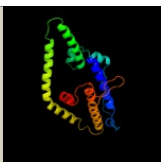
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d3dhwa1	 Alignment		99.9	15	Fold: MetI-like Superfamily: MetI-like Family: MetI-like
2	c2onkC_	 Alignment		99.8	12	PDB header: membrane protein Chain: C: PDB Molecule: molybdate/tungstate abc transporter, permease PDBTitle: abc transporter modbc in complex with its binding protein2 moda
3	d2onkc1	 Alignment		99.8	12	Fold: MetI-like Superfamily: MetI-like Family: MetI-like
4	d3d31c1	 Alignment		99.8	16	Fold: MetI-like Superfamily: MetI-like Family: MetI-like
5	c3d31D_	 Alignment		99.8	16	PDB header: transport protein Chain: D: PDB Molecule: sulfate/molybdate abc transporter, permease PDBTitle: modbc from methanosarcina acetivorans
6	c3fh6F_	 Alignment		99.7	12	PDB header: transport protein Chain: F: PDB Molecule: maltose transport system permease protein malf; PDBTitle: crystal structure of the resting state maltose transporter from e.2 coli
7	d2r6gf2	 Alignment		99.7	10	Fold: MetI-like Superfamily: MetI-like Family: MetI-like
8	c2r6gF_	 Alignment		99.7	9	PDB header: hydrolase/transport protein Chain: F: PDB Molecule: maltose transport system permease protein malf; PDBTitle: the crystal structure of the e. coli maltose transporter
9	d2r6gg1	 Alignment		99.4	16	Fold: MetI-like Superfamily: MetI-like Family: MetI-like
10	c1ciiA_	 Alignment		17.1	7	PDB header: transmembrane protein Chain: A: PDB Molecule: colicin ia; PDBTitle: colicin ia
11	c1wz4A_	 Alignment		13.8	67	PDB header: gene regulation Chain: A: PDB Molecule: major surface antigen; PDBTitle: solution conformation of adr subtype hbv pre-s2 epitope

12	c2cw1A_	Alignment		10.3	28	PDB header: de novo protein Chain: A: PDB Molecule: sn4m; PDBTitle: solution structure of the de novo-designed lambda cro fold2 protein
13	c2jwaA_	Alignment		6.8	20	PDB header: transferase Chain: A: PDB Molecule: receptor tyrosine-protein kinase erbb-2; PDBTitle: erbb2 transmembrane segment dimer spatial structure
14	d1tdpa_	Alignment		6.5	19	Fold: Bromodomain-like Superfamily: Bacteriocin immunity protein-like Family: Carnobacteriocin B2 immunity protein
15	d3ci0k2	Alignment		5.5	20	Fold: SAM domain-like Superfamily: GspK insert domain-like Family: GspK insert domain-like
16	c2b9sB_	Alignment		5.3	31	PDB header: isomerase/dna Chain: B: PDB Molecule: dna topoisomerase i-like protein; PDBTitle: crystal structure of heterodimeric l. donovani2 topoisomerase i-vanadate-dna complex

17 [d2nwwa1](#)

Alignment



5.3

11

Fold: Proton glutamate symport protein
Superfamily: Proton glutamate symport protein
Family: Proton glutamate symport protein