
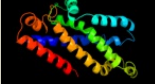


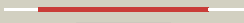



















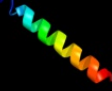





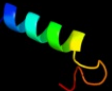


Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d3dhw1	 Alignment		99.9	20	Fold: Metl-like Superfamily: Metl-like Family: Metl-like
2	c2onkC	 Alignment		99.8	13	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	13	Fold: Metl-like Superfamily: Metl-like Family: Metl-like
4	c3d31D	 Alignment		99.8	11	PDB header: transport protein Chain: D: PDB Molecule: sulfate/molybdate abc transporter, permease PDBTitle: modbc from methanosarcina acetivorans
5	d3d31c1	 Alignment		99.8	11	Fold: Metl-like Superfamily: Metl-like Family: Metl-like
6	c3fh6F	 Alignment		99.7	10	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	12	Fold: Metl-like Superfamily: Metl-like Family: Metl-like
8	c2r6gF	 Alignment		99.6	12	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	11	Fold: Metl-like Superfamily: Metl-like Family: Metl-like
10	c1wz4A	 Alignment		22.7	22	PDB header: gene regulation Chain: A: PDB Molecule: major surface antigen; PDBTitle: solution conformation of adr subtype hbv pre-s2 epitope
11	c1ciiA	 Alignment		12.9	5	PDB header: transmembrane protein Chain: A: PDB Molecule: colicin ia; PDBTitle: colicin ia

12	d1ntca_	Alignment		12.1	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
13	d1etob_	Alignment		7.7	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
14	c2ka2B_	Alignment		7.6	12	PDB header: membrane protein Chain: B: PDB Molecule: bcl2/adenovirus e1b 19 kda protein-interacting PDBTitle: solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles with his173-ser172 intermonomer3 hydrogen bond restraints
15	c2ka1A_	Alignment		7.6	12	PDB header: membrane protein Chain: A: PDB Molecule: bcl2/adenovirus e1b 19 kda protein-interacting PDBTitle: solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles
16	c2ka1B_	Alignment		7.6	12	PDB header: membrane protein Chain: B: PDB Molecule: bcl2/adenovirus e1b 19 kda protein-interacting PDBTitle: solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles
17	c2ka2A_	Alignment		7.6	12	PDB header: membrane protein Chain: A: PDB Molecule: bcl2/adenovirus e1b 19 kda protein-interacting PDBTitle: solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles with his173-ser172 intermonomer3 hydrogen bond restraints
18	d2auwa1	Alignment		6.9	27	Fold: lambda repressor-like DNA-binding domains Superfamily: lambda repressor-like DNA-binding domains Family: NE0471 C-terminal domain-like
19	c2cw1A_	Alignment		6.8	45	PDB header: de novo protein Chain: A: PDB Molecule: sn4m; PDBTitle: solution structure of the de novo-designed lambda cro fold2 protein
20	d2jrxa1	Alignment		6.7	13	Fold: YejL-like Superfamily: YejL-like Family: YejL-like
21	d2jpqa1	Alignment	not modelled	6.7	21	Fold: YejL-like Superfamily: YejL-like Family: YejL-like
22	c3n9iA_	Alignment	not modelled	6.5	14	PDB header: ligase Chain: A: PDB Molecule: tryptophanyl-trna synthetase; PDBTitle: crystal structure of tryptophanyl-trna synthetase from yersinia pestis2 co92
23	c2j5dA_	Alignment	not modelled	6.3	12	PDB header: membrane protein Chain: A: PDB Molecule: bcl2/adenovirus e1b 19 kda protein-interacting PDBTitle: nmr structure of bnip3 transmembrane domain in lipid2 bicelles
24	c1xl6B_	Alignment	not modelled	5.8	17	PDB header: metal transport Chain: B: PDB Molecule: inward rectifier potassium channel; PDBTitle: intermediate gating structure 2 of the inwardly rectifying k+ channel2 kirbac3.1
25	d2qtia1	Alignment	not modelled	5.2	13	Fold: YejL-like Superfamily: YejL-like Family: YejL-like