

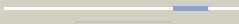




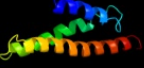

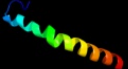







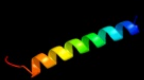

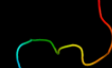

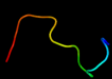


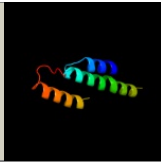
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2azea1	 Alignment		28.4	30	Fold: E2F-DP heterodimerization region Superfamily: E2F-DP heterodimerization region Family: DP dimerization segment
2	c2l2tA	 Alignment		25.4	21	PDB header: membrane protein Chain: A: PDB Molecule: receptor tyrosine-protein kinase erbb-4; PDBTitle: solution nmr structure of the erbb4 dimeric membrane domain
3	c2kncB	 Alignment		23.8	29	PDB header: cell adhesion Chain: B: PDB Molecule: integrin beta-3; PDBTitle: platelet integrin alfaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
4	d1iwga8	 Alignment		12.0	9	Fold: Multidrug efflux transporter AcrB transmembrane domain Superfamily: Multidrug efflux transporter AcrB transmembrane domain Family: Multidrug efflux transporter AcrB transmembrane domain
5	c2jp3A	 Alignment		11.7	14	PDB header: transcription Chain: A: PDB Molecule: fxyd domain-containing ion transport regulator 4; PDBTitle: solution structure of the human fxyd4 (chif) protein in sds2 micelles
6	c2jo1A	 Alignment		11.6	19	PDB header: hydrolase regulator Chain: A: PDB Molecule: phospholemman; PDBTitle: structure of the na,k-atpase regulatory protein fxyd1 in2 micelles
7	d2gdwa1	 Alignment		9.5	25	Fold: Acyl carrier protein-like Superfamily: ACP-like Family: Peptidyl carrier domain
8	d2pnga1	 Alignment		9.0	21	Fold: Acyl carrier protein-like Superfamily: ACP-like Family: Acyl-carrier protein (ACP)
9	c2rmzA	 Alignment		8.5	28	PDB header: cell adhesion Chain: A: PDB Molecule: integrin beta-3; PDBTitle: bicelle-embedded integrin beta3 transmembrane segment
10	c1ywua	 Alignment		6.5	18	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein pa4608; PDBTitle: solution nmr structure of pseudomonas aeruginosa protein pa4608.2 northeast structural genomics target pat7
11	d1ywua1	 Alignment		6.5	18	Fold: Split barrel-like Superfamily: PilZ domain-like Family: PilZ domain

12

[c3rkoK_](#)

Alignment



5.2

19

PDB header:oxidoreductase**Chain:** K: **PDB Molecule:**nadh-quinone oxidoreductase subunit k;
PDBTitle: crystal structure of the membrane domain of respiratory complex i from2 e. coli at 3.0 angstrom resolution