
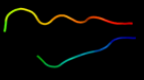



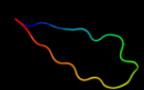



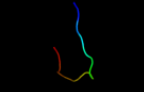



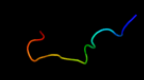

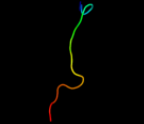



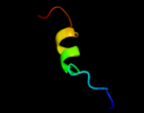

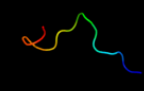








#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1axib1	 Alignment		28.3	44	Fold: Immunoglobulin-like beta-sandwich Superfamily: Fibronectin type III Family: Fibronectin type III
2	d1hm6a_	 Alignment		14.9	32	Fold: Annexin Superfamily: Annexin Family: Annexin
3	d1f6fb1	 Alignment		10.2	32	Fold: Immunoglobulin-like beta-sandwich Superfamily: Fibronectin type III Family: Fibronectin type III
4	d2ctea1	 Alignment		9.7	31	Fold: Eukaryotic type KH-domain (KH-domain type I) Superfamily: Eukaryotic type KH-domain (KH-domain type I) Family: Eukaryotic type KH-domain (KH-domain type I)
5	d1erna1	 Alignment		9.6	20	Fold: Immunoglobulin-like beta-sandwich Superfamily: Fibronectin type III Family: Fibronectin type III
6	d1viga_	 Alignment		8.9	21	Fold: Eukaryotic type KH-domain (KH-domain type I) Superfamily: Eukaryotic type KH-domain (KH-domain type I) Family: Eukaryotic type KH-domain (KH-domain type I)
7	c3hagA_	 Alignment		8.1	67	PDB header: virus Chain: A: PDB Molecule: capsid protein; PDBTitle: crystal structure of the hepatitis e virus-like particle
8	d1oe4a_	 Alignment		7.5	47	Fold: Uracil-DNA glycosylase-like Superfamily: Uracil-DNA glycosylase-like Family: Single-strand selective monofunctional uracil-DNA glycosylase SMUG1
9	c2ztnA_	 Alignment		7.3	67	PDB header: virus Chain: A: PDB Molecule: capsid protein; PDBTitle: hepatitis e virus orf2 (genotype 3)
10	c2k8fB_	 Alignment		7.1	26	PDB header: transferase/transcription Chain: B: PDB Molecule: cellular tumor antigen p53; PDBTitle: structural basis for the regulation of p53 function by p300
11	c2zzqA_	 Alignment		6.4	67	PDB header: virus Chain: A: PDB Molecule: protein orf3, capsid protein; PDBTitle: crystal structure analysis of the hev capsid protein, porf2

12	c1gp8A_		Alignment		6.4	78	PDB header: viral protein Chain: A: PDB Molecule: protein (scaffolding protein); PDBTitle: nmr solution structure of the coat protein-binding domain2 of bacteriophage p22 scaffolding protein
13	dlrh1a1		Alignment		6.3	45	Fold: Cloacin translocation domain Superfamily: Cloacin translocation domain Family: Cloacin translocation domain
14	c1rh1A_		Alignment		5.7	45	PDB header: antibiotic Chain: A: PDB Molecule: colicin b; PDBTitle: crystal structure of the cytotoxic bacterial protein2 colicin b at 2.5 a resolution

15

[d1f35a_](#)

Alignment



5.6

43

Fold:Olfactory marker protein
Superfamily:Olfactory marker protein
Family:Olfactory marker protein