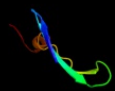
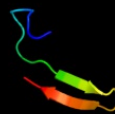
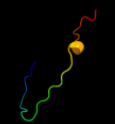
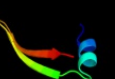
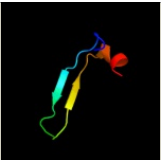


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2e5qA_	 Alignment		23.9	31	PDB header: transcription Chain: A: PDB Molecule: phd finger protein 19; PDBTitle: solution structure of the tudor domain of phd finger2 protein 19, isoform b [homo sapiens]
2	c2e5pA_	 Alignment		13.9	26	PDB header: transcription Chain: A: PDB Molecule: phd finger protein 1; PDBTitle: solution structure of the tudor domain of phd finger2 protein 1 (phf1 protein)
3	d2gycm1	 Alignment		9.7	14	Fold: Ribonuclease H-like motif Superfamily: Translational machinery components Family: Ribosomal protein L18 and S11
4	d1ovya_	 Alignment		9.1	22	Fold: Ribonuclease H-like motif Superfamily: Translational machinery components Family: Ribosomal protein L18 and S11
5	d1y4oa1	 Alignment		7.6	22	Fold: Profilin-like Superfamily: Roadblock/LC7 domain Family: Roadblock/LC7 domain
6	c2eqjA_	 Alignment		6.8	19	PDB header: transcription Chain: A: PDB Molecule: metal-response element-binding transcription PDBTitle: solution structure of the tudor domain of metal-response2 element-binding transcription factor 2
7	c3lznA_	 Alignment		6.1	47	PDB header: transport protein Chain: A: PDB Molecule: p19 protein; PDBTitle: crystal structure analysis of the apo p19 protein from campylobacter2 jejuni at 1.59 a at ph 9
8	c2i9wA_	 Alignment		5.7	47	PDB header: metal binding protein Chain: A: PDB Molecule: hypothetical protein; PDBTitle: crystal structure of a sec-c motif containing protein (psyc_2064) from2 psychrobacter arcticus at 1.75 a resolution
9	d2pi6a2	 Alignment		5.7	25	Fold: FKBP-like Superfamily: Chitinase insertion domain Family: Chitinase insertion domain

10

[d1zhva2](#)

Alignment



5.5

25

Fold: Ferredoxin-like
Superfamily: ACT-like
Family: Atu0741-like