

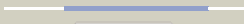
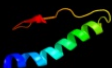

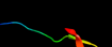

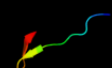










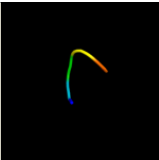


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dly4oa1	 Alignment		27.0	17	Fold: Profilin-like Superfamily: Roadblock/LC7 domain Family: Roadblock/LC7 domain
2	d2hz5a1	 Alignment		25.7	18	Fold: Profilin-like Superfamily: Roadblock/LC7 domain Family: Roadblock/LC7 domain
3	d1prtb1	 Alignment		11.8	41	Fold: OB-fold Superfamily: Bacterial enterotoxins Family: Bacterial AB5 toxins, B-subunits
4	d1prtc1	 Alignment		9.2	41	Fold: OB-fold Superfamily: Bacterial enterotoxins Family: Bacterial AB5 toxins, B-subunits
5	c1bcph_	 Alignment		7.1	41	PDB header: toxin Chain: H: PDB Molecule: pertussis toxin; PDBTitle: binary complex of pertussis toxin and atp
6	c1u7mb_	 Alignment		7.1	44	PDB header: de novo protein Chain: B: PDB Molecule: four-helix bundle model; PDBTitle: solution structure of a diiron protein model: due ferri(ii)2 turn mutant
7	c2jugB_	 Alignment		6.9	11	PDB header: biosynthetic protein Chain: B: PDB Molecule: tubc protein; PDBTitle: multienzyme docking in hybrid megasynthetases
8	c2rpiA_	 Alignment		6.6	29	PDB header: hydrolase Chain: A: PDB Molecule: ribonuclease h; PDBTitle: the nmr structure of the submillisecond folding2 intermediate of the thermus thermophilus ribonuclease h
9	c1zllE_	 Alignment		5.9	24	PDB header: membrane protein/signaling protein Chain: E: PDB Molecule: cardiac phospholamban; PDBTitle: nmr structure of unphosphorylated human phospholamban2 pentamer

10 [dlehsa_](#)

Alignment



5.4

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Fold:Toxic hairpin
Superfamily:Heat-stable enterotoxin B
Family:Heat-stable enterotoxin B