








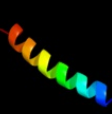

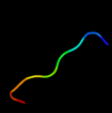












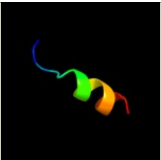
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3fvA_	 Alignment		26.0	40	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized novel protein; PDBTitle: crystal structure of novel protein of unknown function (yp_111841.1)2 from burkholderia pseudomallei k96243 at 1.90 a resolution
2	c3kikF_	 Alignment		17.8	48	PDB header: transcription Chain: F: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
3	c3kikG_	 Alignment		17.3	48	PDB header: transcription Chain: G: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
4	c3kikH_	 Alignment		17.2	48	PDB header: transcription Chain: H: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
5	c3kikE_	 Alignment		17.2	48	PDB header: transcription Chain: E: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
6	c2lk0A_	 Alignment		16.3	44	PDB header: rna binding protein Chain: A: PDB Molecule: rna-binding protein 5; PDBTitle: solution structure and binding studies of the ranbp2-type zinc finger2 of rbm5
7	d2ftua1	 Alignment		14.1	54	Fold: RAP domain-like Superfamily: RAP domain-like Family: RAP domain
8	d2fcwa1	 Alignment		13.8	54	Fold: RAP domain-like Superfamily: RAP domain-like Family: RAP domain
9	c3kijF_	 Alignment		7.7	48	PDB header: transcription Chain: F: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
10	c3kijH_	 Alignment		7.0	48	PDB header: transcription Chain: H: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex
11	c3kijE_	 Alignment		7.0	48	PDB header: transcription Chain: E: PDB Molecule: saga-associated factor 11; PDBTitle: sgf11:sus1 complex

12	c1htrP_	<div>Alignment</div>		6.9	63	<p>PDB header:aspartyl protease Chain: P: PDB Molecule:progastricsin (pro segment); PDBTitle: crystal and molecular structures of human progastricsin at 1.622 angstroms resolution</p>
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[d2fcta1](#)

Alignment



6.0

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Fold:Double-stranded beta-helix
Superfamily:Clavamate synthase-like
Family:PhyH-like