


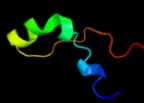













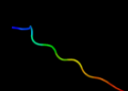


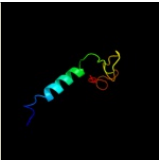


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3cz6A_	 Alignment		14.6	28	PDB header: protein binding Chain: A: PDB Molecule: dna-binding protein rap1; PDBTitle: crystal structure of the rap1 c-terminus
2	c2qqhA_	 Alignment		12.6	25	PDB header: immune system, membrane protein Chain: A: PDB Molecule: complement component c8 alpha chain; PDBTitle: structure of c8a-macpf reveals mechanism of membrane attack2 in complement immune defense
3	d1bg6a1	 Alignment		12.1	23	Fold: 6-phosphogluconate dehydrogenase C-terminal domain-like Superfamily: 6-phosphogluconate dehydrogenase C-terminal domain-like Family: N-(1-D-carboxylethyl)-L-norvaline dehydrogenase
4	d1es9a_	 Alignment		9.8	20	Fold: Flavodoxin-like Superfamily: SGNH hydrolase Family: Acetylhydrolase
5	c3h0gE_	 Alignment		7.6	38	PDB header: transcription Chain: E: PDB Molecule: dna-directed rna polymerases i, ii, and iii PDBTitle: rna polymerase ii from schizosaccharomyces pombe
6	d1dzfa2	 Alignment		7.4	45	Fold: RPB5-like RNA polymerase subunit Superfamily: RPB5-like RNA polymerase subunit Family: RPB5
7	d1n8ia_	 Alignment		7.3	36	Fold: TIM beta/alpha-barrel Superfamily: Malate synthase G Family: Malate synthase G
8	d2f22a1	 Alignment		6.6	50	Fold: DinB/YfiT-like putative metalloenzymes Superfamily: DinB/YfiT-like putative metalloenzymes Family: DinB-like
9	d1hmja_	 Alignment		6.3	55	Fold: RPB5-like RNA polymerase subunit Superfamily: RPB5-like RNA polymerase subunit Family: RPB5
10	c2lbfA_	 Alignment		5.6	29	PDB header: ribosomal protein Chain: A: PDB Molecule: 60s acidic ribosomal protein p1; PDBTitle: solution structure of the dimerization domain of human ribosomal2 protein p1/p2 heterodimer

11 [d2axta1](#)

Alignment



5.4

24

Fold:Bacterial photosystem II reaction centre, L and M subunits
Superfamily:Bacterial photosystem II reaction centre, L and M subunits
Family:Bacterial photosystem II reaction centre, L and M subunits