
















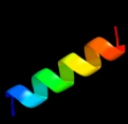




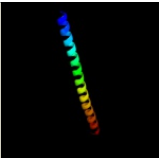
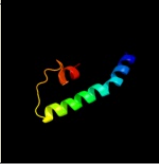


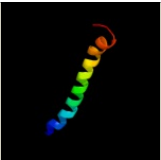
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2joka1	 Alignment		10.7	32	Fold: SopE-like GEF domain Superfamily: SopE-like GEF domain Family: SopE-like GEF domain
2	c2kelB_	 Alignment		9.7	50	PDB header: transcription repressor Chain: B: PDB Molecule: uncharacterized protein 56b; PDBTitle: structure of the transcription regulator svtr from the2 hyperthermophilic archaeal virus sirv1
3	c1sg7A_	 Alignment		8.4	19	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative cation transport regulator chab; PDBTitle: nmr solution structure of the putative cation transport2 regulator chab
4	d1sg7a1	 Alignment		8.4	19	Fold: ChaB-like Superfamily: ChaB-like Family: ChaB-like
5	c3swfA_	 Alignment		7.5	23	PDB header: transport protein Chain: A: PDB Molecule: cgmp-gated cation channel alpha-1; PDBTitle: cnga1 621-690 containing clz domain
6	c1wnkA_	 Alignment		7.3	53	PDB header: transcription Chain: A: PDB Molecule: fibroin-modulator-binding-protein-1; PDBTitle: nmr structure of fmbp-1 tandem repeat 3 in 30%(v/v) tfe2 solution
7	d1s35a2	 Alignment		7.0	9	Fold: Spectrin repeat-like Superfamily: Spectrin repeat Family: Spectrin repeat
8	d1ka8a_	 Alignment		6.6	41	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: P4 origin-binding domain-like
9	c3t38B_	 Alignment		6.4	24	PDB header: oxidoreductase Chain: B: PDB Molecule: arsenate reductase; PDBTitle: corynebacterium glutamicum thioredoxin-dependent arsenate reductase2 cg_arsc1'
10	d1u5pa1	 Alignment		6.3	16	Fold: Spectrin repeat-like Superfamily: Spectrin repeat Family: Spectrin repeat
11	d1u5pa2	 Alignment		6.2	19	Fold: Spectrin repeat-like Superfamily: Spectrin repeat Family: Spectrin repeat

12	d2spca_	Alignment		6.1	8	Fold: Spectrin repeat-like Superfamily: Spectrin repeat Family: Spectrin repeat
13	dlhcia4	Alignment		5.9	11	Fold: Spectrin repeat-like Superfamily: Spectrin repeat Family: Spectrin repeat

14

[c1wlxA](#)

Alignment



5.3

6

PDB header:protein binding
Chain: A: **PDB Molecule:**alpha-actinin 4;
PDBTitle: solution structure of the third spectrin repeat of alpha-2 actinin-4