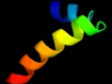
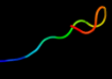
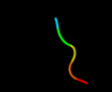
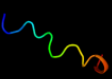
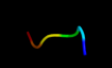


Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1jt6a2	 Alignment		33.6	29	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
2	d2idaa1	 Alignment		9.4	36	Fold: RING/U-box Superfamily: RING/U-box Family: Zf-UBP
3	d1ra0a1	 Alignment		7.4	38	Fold: Composite domain of metallo-dependent hydrolases Superfamily: Composite domain of metallo-dependent hydrolases Family: Cytosine deaminase
4	d2c1wa1	 Alignment		7.3	28	Fold: EndoU-like Superfamily: EndoU-like Family: Eukaryotic EndoU ribonuclease
5	d1eita_	 Alignment		6.5	83	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: omega toxin-like Family: Spider toxins
6	d2riga_	 Alignment		6.2	33	Fold: 4-helical cytokines Superfamily: 4-helical cytokines Family: Interferons/interleukin-10 (IL-10)
7	c2rigA_	 Alignment		6.2	33	PDB header: cytokine Chain: A: PDB Molecule: interferon-gamma; PDBTitle: crystal structure of recombinant rabbit interferon-gamma at 2.7-angstroms resolution
8	c1v91A_	 Alignment		6.2	67	PDB header: toxin Chain: A: PDB Molecule: delta-palutoxin it2; PDBTitle: solution structure of insectidal toxin delta-paluit2-nh2
9	d1dx5i3	 Alignment		6.0	44	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: EGF/Laminin Family: EGF-type module
10	c2owoA_	 Alignment		5.9	26	PDB header: ligase/dna Chain: A: PDB Molecule: dna ligase; PDBTitle: last stop on the road to repair: structure of e.coli dna ligase bound2 to nicked dna-adenylate
11	d1dgsa1	 Alignment		5.7	26	Fold: SAM domain-like Superfamily: RuvA domain 2-like Family: NAD+-dependent DNA ligase, domain 3

12	dlg1pa_	Alignment		5.6	41	Fold: Knottins (small inhibitors, toxins, lectins) Superfamily: omega toxin-like Family: Conotoxin
13	c2dj8A_	Alignment		5.1	40	PDB header: metal binding protein Chain: A: PDB Molecule: protein cbfa2t1; PDBTitle: solution structure of zf-mynd domain of protein cbfa2t2 (protein mtg8)

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Alignment



5.0

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PDB header:transcription
Chain: C: **PDB Molecule:**helicase protein mom1;
PDBTitle: structural basis of transcriptional gene silencing mediated by2 arabidopsis mom1