


















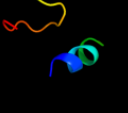


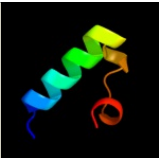
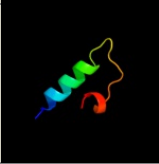


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1nnva_	 Alignment		100.0	48	Fold: Cystatin-like Superfamily: Putative dsDNA mimic Family: Putative dsDNA mimic
2	c1bjzA_	 Alignment		14.1	21	PDB header: transcription regulation Chain: A: PDB Molecule: tetracycline repressor; PDBTitle: tetracycline chelated mg2+-ion initiates helix unwinding for tet2 repressor induction
3	c1bmlD_	 Alignment		13.4	30	PDB header: blood clotting Chain: D: PDB Molecule: streptokinase; PDBTitle: complex of the catalytic domain of human plasmin and streptokinase
4	d2np5a1	 Alignment		13.4	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
5	d1c4pc_	 Alignment		12.1	26	Fold: beta-Grasp (ubiquitin-like) Superfamily: Staphylokinase/streptokinase Family: Staphylokinase/streptokinase
6	d1qgra_	 Alignment		11.2	26	Fold: beta-Grasp (ubiquitin-like) Superfamily: Staphylokinase/streptokinase Family: Staphylokinase/streptokinase
7	d2q7sa1	 Alignment		10.8	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
8	d2vkea1	 Alignment		9.3	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
9	d1x99a_	 Alignment		9.0	17	Fold: Cytolysin/lectin Superfamily: Cytolysin/lectin Family: Fungal fruit body lectin
10	c3htxA_	 Alignment		7.2	38	PDB header: transferase/rna Chain: A: PDB Molecule: hen1; PDBTitle: crystal structure of small rna methyltransferase hen1
11	c3o60A_	 Alignment		7.1	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lin0861 protein; PDBTitle: the crystal structure of lin0861 from listeria innocua to 2.8a

12	d1zk8a1	Alignment		6.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
13	c2hxiA_	Alignment		5.7	17	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator from streptomyces coelicolor a3(2)

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[c2np5A](#)

Alignment



5.3

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PDB header:transcription**Chain:** A: **PDB Molecule:**transcriptional regulator;**PDBTitle:** crystal structure of a transcriptional regulator (rha1_ro04179) from2 rhodococcus sp. rha1.