




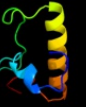











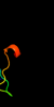

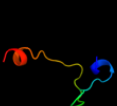




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dli94r_	 Alignment		100.0	42	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Ribosomal protein S18 Family: Ribosomal protein S18
2	d2uubr1	 Alignment		99.9	43	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Ribosomal protein S18 Family: Ribosomal protein S18
3	d2qalr1	 Alignment		99.9	100	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Ribosomal protein S18 Family: Ribosomal protein S18
4	c3df1R_	 Alignment		99.9	100	PDB header: ribosome Chain: R: PDB Molecule: 30s ribosomal protein s18; PDBTitle: crystal structure of the bacterial ribosome from escherichia2 coli in complex with hygromycin b. this file contains the3 30s subunit of the first 70s ribosome, with hygromycin b4 bound. the entire crystal structure contains two 70s5 ribosomes.
5	c3bbnR_	 Alignment		99.9	39	PDB header: ribosome Chain: R: PDB Molecule: ribosomal protein s18; PDBTitle: homology model for the spinach chloroplast 30s subunit2 fitted to 9.4a cryo-em map of the 70s chlororibosome.
6	dlg1xc_	 Alignment		99.8	52	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Ribosomal protein S18 Family: Ribosomal protein S18
7	c2pmzG_	 Alignment		17.6	17	PDB header: translation, transferase Chain: G: PDB Molecule: dna-directed rna polymerase subunit a"; PDBTitle: archaeal rna polymerase from sulfolobus solfataricus
8	c2ectA_	 Alignment		17.2	5	PDB header: metal binding protein Chain: A: PDB Molecule: ring finger protein 126; PDBTitle: solution structure of the zinc finger, c3hc4 type (ring2 finger) domain of ring finger protein 126
9	c2d8tA_	 Alignment		15.2	18	PDB header: metal binding protein Chain: A: PDB Molecule: ring finger protein 146; PDBTitle: solution structure of the ring domain of the human ring2 finger protein 146
10	c2csyA_	 Alignment		15.0	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: zinc finger protein 183-like 1; PDBTitle: solution structure of the ring domain of the zinc finger2 protein 183-like 1
11	c3h0gA_	 Alignment		13.4	21	PDB header: transcription Chain: A: PDB Molecule: dna-directed rna polymerase ii subunit rpb1; PDBTitle: rna polymerase ii from schizosaccharomyces pombe

29	d1z84a1	Alignment	not modelled	5.5	13	Superfamily: HIT-like Family: Hexose-1-phosphate uridylyltransferase
30	d1g4yb	Alignment	not modelled	5.4	35	Fold: Small-conductance potassium channel Superfamily: Small-conductance potassium channel Family: Small-conductance potassium channel
31	c3ktsA	Alignment	not modelled	5.4	40	PDB header: transcriptional regulator Chain: A: PDB Molecule: glycerol uptake operon antiterminator regulatory protein; PDBTitle: crystal structure of glycerol uptake operon antiterminator regulatory2 protein from listeria monocytogenes str. 4b f2365
32	c2kizA	Alignment	not modelled	5.3	10	PDB header: metal binding protein Chain: A: PDB Molecule: e3 ubiquitin-protein ligase arkadia; PDBTitle: solution structure of arkadia ring-h2 finger domain