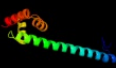
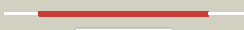
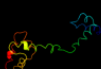


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2zjrn1	 Alignment		100.0	55	Fold: PABP domain-like Superfamily: Ribosomal protein L20 Family: Ribosomal protein L20
2	d2j01u1	 Alignment		100.0	59	Fold: PABP domain-like Superfamily: Ribosomal protein L20 Family: Ribosomal protein L20
3	c3bboS_	 Alignment		100.0	47	PDB header: ribosome Chain: S: PDB Molecule: ribosomal protein l20; PDBTitle: homology model for the spinach chloroplast 50s subunit2 fitted to 9.4a cryo-em map of the 70s chlororibosome
4	c2ghjD_	 Alignment		100.0	64	PDB header: structural protein Chain: D: PDB Molecule: 50s ribosomal protein l20; PDBTitle: crystal structure of folded and partially unfolded forms of2 aquifex aeolicus ribosomal protein l20
5	c2ftcL_	 Alignment		100.0	35	PDB header: ribosome Chain: L: PDB Molecule: mrpl20 protein; PDBTitle: structural model for the large subunit of the mammalian mitochondrial2 ribosome
6	d2gyco1	 Alignment		100.0	100	Fold: PABP domain-like Superfamily: Ribosomal protein L20 Family: Ribosomal protein L20
7	d1gyza_	 Alignment		99.9	66	Fold: PABP domain-like Superfamily: Ribosomal protein L20 Family: Ribosomal protein L20
8	d2coba1	 Alignment		48.8	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Psq domain
9	c2k53A_	 Alignment		38.6	27	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: a3dk08 protein; PDBTitle: nmr solution structure of a3dk08 protein from clostridium2 thermocellum: northeast structural genomics consortium3 target cmr9
10	d3deua1	 Alignment		31.5	8	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
11	d1fipa_	 Alignment		29.3	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like

12	c3e7lD_	Alignment		26.6	19	PDB header: transcription regulator Chain: D: PDB Molecule: transcriptional regulator (ntrc family); PDBTitle: crystal structure of sigma54 activator ntrc4's dna binding2 domain
13	c3fm5D_	Alignment		24.4	12	PDB header: transcription regulator Chain: D: PDB Molecule: transcriptional regulator; PDBTitle: x-ray crystal structure of transcriptional regulator (marr family)2 from rhodococcus sp. rha1
14	c2p9xB_	Alignment		20.6	31	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: hypothetical protein ph0832; PDBTitle: crystal structure of ph0832 from pyrococcus horikoshii ot3
15	d1ku9a_	Alignment		20.5	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding protein Mj223
16	c3deuB_	Alignment		18.6	8	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator slyA; PDBTitle: crystal structure of transcription regulatory protein slyA2 from salmonella typhimurium in complex with salicylate3 ligands
17	c2k5eA_	Alignment		18.6	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: solution structure of putative uncharacterized protein2 gsu1278 from methanocaldococcus jannaschii, northeast3 structural genomics consortium (nesg) target gsr195
18	d1etob_	Alignment		17.3	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
19	d1ntca_	Alignment		17.1	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
20	d2cg4a1	Alignment		16.2	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
21	d2cyya1	Alignment	not modelled	15.1	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
22	d1etxa_	Alignment	not modelled	14.3	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
23	c2ijlB_	Alignment	not modelled	14.1	29	PDB header: transcription Chain: B: PDB Molecule: molybdenum-binding transcriptional repressor; PDBTitle: the structure of a putative mode from agrobacterium tumefaciens.
24	d2cfxa1	Alignment	not modelled	12.9	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
25	c3g3rA_	Alignment	not modelled	12.5	23	PDB header: biosynthetic protein Chain: A: PDB Molecule: vacuolar transporter chaperone 4; PDBTitle: crystal structure of a eukaryotic polyphosphate polymerase2 in complex with apnhp-mn2+
26	d1y0ua_	Alignment	not modelled	11.5	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: ArsR-like transcriptional regulators
27	c2l4aA_	Alignment	not modelled	10.9	26	PDB header: dna binding protein Chain: A: PDB Molecule: leucine responsive regulatory protein; PDBTitle: nmr structure of the dna-binding domain of e.coli lrp
28	c3jw4C_	Alignment	not modelled	10.2	14	PDB header: transcription regulator Chain: C: PDB Molecule: transcriptional regulator, marr/emrr family; PDBTitle: the structure of a putative marr family transcriptional regulator from2 clostridium acetobutylicum

29	c3g3oA_	Alignment	not modelled	9.8	20	PDB header: biosynthetic protein Chain: A: PDB Molecule: vacuolar transporter chaperone 2; PDBTitle: crystal structure of the cytoplasmic tunnel domain in yeast2 vtc2p
30	d1l1ga1	Alignment	not modelled	9.3	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
31	c2ee7A_	Alignment	not modelled	8.9	12	PDB header: structural protein Chain: A: PDB Molecule: sperm flagellar protein 1; PDBTitle: solution structure of the ch domain from human sperm2 flagellar protein 1
32	d1umqa_	Alignment	not modelled	8.4	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
33	c1umqA_	Alignment	not modelled	8.4	21	PDB header: dna-binding protein Chain: A: PDB Molecule: photosynthetic apparatus regulatory protein; PDBTitle: solution structure and dna binding of the effector domain2 from the global regulator prrA(rega) from r. sphaeroides:3 insights into dna binding specificity
34	d1g2ha_	Alignment	not modelled	8.0	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: FIS-like
35	d1pc6a_	Alignment	not modelled	8.0	25	Fold: NinB Superfamily: NinB Family: NinB
36	d2hepa1	Alignment	not modelled	7.3	67	Fold: Long alpha-hairpin Superfamily: YnzC-like Family: YznC-like
37	c2hepA_	Alignment	not modelled	7.3	67	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: upf0291 protein ynzC; PDBTitle: solution nmr structure of the upf0291 protein ynzC from2 bacillus subtilis. northeast structural genomics target3 sr384.
38	d1b9ma1	Alignment	not modelled	6.9	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: N-terminal domain of molybdate-dependent transcriptional regulator ModE
39	c3cdhB_	Alignment	not modelled	6.2	12	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of the marr family transcriptional regulator spo14532 from silicibacter pomeroyi dss-3
40	d2d5ra1	Alignment	not modelled	5.9	19	Fold: Ribonuclease H-like motif Superfamily: Ribonuclease H-like Family: CAF1-like ribonuclease
41	d2cpta1	Alignment	not modelled	5.6	19	Fold: Spectrin repeat-like Superfamily: MIT domain Family: MIT domain
42	d2fi0a1	Alignment	not modelled	5.4	15	Fold: SP0561-like Superfamily: SP0561-like Family: SP0561-like
43	d1qpza1	Alignment	not modelled	5.4	24	Fold: lambda repressor-like DNA-binding domains Superfamily: lambda repressor-like DNA-binding domains Family: GalR/LacI-like bacterial regulator
44	c3s2wB_	Alignment	not modelled	5.2	18	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: the crystal structure of a marr transcriptional regulator from2 methanosarcina mazei go1
45	d1lcda_	Alignment	not modelled	5.2	13	Fold: lambda repressor-like DNA-binding domains Superfamily: lambda repressor-like DNA-binding domains Family: GalR/LacI-like bacterial regulator