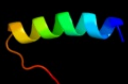
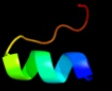


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2rbdB_	 Alignment		25.9	21	PDB header: structural protein Chain: B; PDB Molecule: bh2358 protein; PDBTitle: crystal structure of a putative spore coat protein (bh2358) from2 bacillus halodurans c-125 at 1.54 a resolution
2	c3d8lB_	 Alignment		22.8	20	PDB header: viral protein Chain: B; PDB Molecule: orf12; PDBTitle: crystal structure of orf12 from the lactococcus lactis bacteriophage2 p2
3	dluera1	 Alignment		18.6	25	Fold: Long alpha-hairpin Superfamily: Fe,Mn superoxide dismutase (SOD), N-terminal domain Family: Fe,Mn superoxide dismutase (SOD), N-terminal domain
4	dluz3a1	 Alignment		13.1	12	Fold: ENT-like Superfamily: ENT-like Family: Emsy N terminal (ENT) domain-like
5	dlzkea1	 Alignment		12.0	43	Fold: ROP-like Superfamily: HP1531-like Family: HP1531-like
6	c3iz5t_	 Alignment		11.5	16	PDB header: ribosome Chain: T; PDB Molecule: 60s ribosomal protein l19 (l19e); PDBTitle: localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
7	c2xozB_	 Alignment		10.1	14	PDB header: ligase Chain: B; PDB Molecule: e3 ubiquitin-protein ligase chfr; PDBTitle: c-terminal cysteine rich domain of human chfr bound to amp
8	dlplja1	 Alignment		9.4	13	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain
9	dlvkoa1	 Alignment		8.6	13	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain
10	clvkoA_	 Alignment		8.5	13	PDB header: isomerase Chain: A; PDB Molecule: inositol-3-phosphate synthase; PDBTitle: crystal structure of inositol-3-phosphate synthase (ce21227) from2 caenorhabditis elegans at 2.30 a resolution
11	dlhcua_	 Alignment		8.3	20	Fold: alpha/alpha toroid Superfamily: Seven-hairpin glycosidases Family: Class I alpha-1;2-mannosidase, catalytic domain

12	c1plhD_	Alignment		8.0	13	PDB header: isomerase Chain: D: PDB Molecule: inositol-3-phosphate synthase; PDBTitle: crystal structure of the 1l-myo-inositol/nad+ complex
13	dlma1a1	Alignment		7.7	25	Fold: Long alpha-hairpin Superfamily: Fe,Mn superoxide dismutase (SOD), N-terminal domain Family: Fe,Mn superoxide dismutase (SOD), N-terminal domain
14	dlidxsa_	Alignment		7.0	30	Fold: SAM domain-like Superfamily: SAM/Pointed domain Family: SAM (sterile alpha motif) domain
15	dlb77a1	Alignment		7.0	13	Fold: DNA clamp Superfamily: DNA clamp Family: DNA polymerase processivity factor
16	c2pqrD_	Alignment		6.9	33	PDB header: apoptosis Chain: D: PDB Molecule: wd repeat protein ykr036c; PDBTitle: crystal structure of yeast fis1 complexed with a fragment of yeast2 caf4
17	c3izct_	Alignment		6.6	17	PDB header: ribosome Chain: T: PDB Molecule: 60s ribosomal protein rpl19 (l19e); PDBTitle: localization of the large subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
18	c1n0fF_	Alignment		6.1	30	PDB header: biosynthetic protein Chain: F: PDB Molecule: protein mrar; PDBTitle: crystal structure of a cell division and cell wall2 biosynthesis protein upf0040 from mycoplasma pneumoniae:3 indication of a novel fold with a possible new conserved4 sequence motif
19	dlbsma1	Alignment		5.6	22	Fold: Long alpha-hairpin Superfamily: Fe,Mn superoxide dismutase (SOD), N-terminal domain Family: Fe,Mn superoxide dismutase (SOD), N-terminal domain
20	d2a5yb1	Alignment		5.6	38	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CED-4 C-terminal domain-like
21	dlug3a1	Alignment	not modelled	5.5	19	Fold: alpha-alpha superhelix Superfamily: ARM repeat Family: MIF4G domain-like