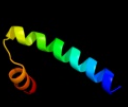
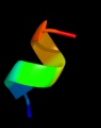


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2rkha_	 Alignment		14.0	30	PDB header: transcription Chain: A: PDB Molecule: putative apha-like transcription factor; PDBTitle: crystal structure of a putative apha-like transcription factor2 (zp_00208345.1) from magnetospirillum magnetotacticum ms-1 at 2.00 a3 resolution
2	c3bj6B_	 Alignment		13.3	12	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of marr family transcription regulator sp03579
3	c2voyG_	 Alignment		11.1	15	PDB header: hydrolase Chain: G: PDB Molecule: sarcoplasmic/endoplasmic reticulum calcium PDBTitle: cryoem model of copa, the copper transporting atpase from2 archaeoglobus fulgidus
4	d1gdha2	 Alignment		10.5	15	Fold: Flavodoxin-like Superfamily: Formate/glycerate dehydrogenase catalytic domain-like Family: Formate/glycerate dehydrogenases, substrate-binding domain
5	d2a61a1	 Alignment		8.7	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
6	c1yhuA_	 Alignment		8.0	14	PDB header: oxygen storage/transport Chain: A: PDB Molecule: hemoglobin a1 chain; PDBTitle: crystal structure of riftia pachytila c1 hemoglobin reveals novel2 assembly of 24 subunits.
7	c2d2mB_	 Alignment		7.5	16	PDB header: oxygen storage/transport Chain: B: PDB Molecule: giant hemoglobin, a2(a5) globin chain; PDBTitle: structure of an extracellular giant hemoglobin of the2 gutless beard worm oligobranchia mashikoi
8	c1t6fA_	 Alignment		6.7	38	PDB header: cell cycle Chain: A: PDB Molecule: geminin; PDBTitle: crystal structure of the coiled-coil dimerization motif of2 geminin

9

[dlq1fa_](#)

Alignment



6.4

17

Fold:Globin-like
Superfamily:Globin-like
Family:Globins