

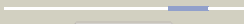














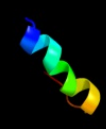







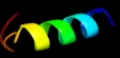
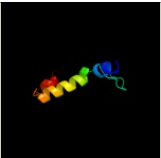


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3h9pB_	 Alignment		100.0	18	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: putative triphosphoribosyl-dephospho-coa synthase; PDBTitle: crystal structure of putative triphosphoribosyl-dephospho-coa synthase2 from archaeoglobus fulgidus
2	c2xc7A_	 Alignment		28.5	28	PDB header: rna binding protein Chain: A: PDB Molecule: phosphorylated adapter rna export protein; PDBTitle: solution structure of phax-rbd in complex with ssrna
3	c2w4sD_	 Alignment		24.6	25	PDB header: structural protein Chain: D: PDB Molecule: ankyrin-repeat protein; PDBTitle: novel rna-binding domain in cryptosporidium parvum at 2.52 angstrom resolution
4	c1p58F_	 Alignment		18.9	22	PDB header: virus Chain: F: PDB Molecule: envelope protein m; PDBTitle: complex organization of dengue virus membrane proteins as revealed by 2 9.5 angstrom cryo-em reconstruction
5	c3c6aA_	 Alignment		16.5	17	PDB header: viral protein Chain: A: PDB Molecule: terminase large subunit; PDBTitle: crystal structure of the rb49 gp17 nuclease domain
6	c3omdB_	 Alignment		15.7	17	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of unknown function protein from leptospirillum2 rubarum
7	d2f05a1	 Alignment		12.5	12	Fold: PAH2 domain Superfamily: PAH2 domain Family: PAH2 domain
8	d1w53a_	 Alignment		10.6	23	Fold: KaiA/RbsU domain Superfamily: KaiA/RbsU domain Family: Phosphoserine phosphatase RsbU, N-terminal domain
9	d1o22a_	 Alignment		9.8	22	Fold: Hypothetical protein TM0875 Superfamily: Hypothetical protein TM0875 Family: Hypothetical protein TM0875
10	d1q3qa3	 Alignment		8.6	30	Fold: GroEL-intermediate domain like Superfamily: GroEL-intermediate domain like Family: Group II chaperonin (CCT, TRIC), intermediate domain
11	d1au7a2	 Alignment		7.6	30	Fold: lambda repressor-like DNA-binding domains Superfamily: lambda repressor-like DNA-binding domains Family: POU-specific domain

12	d1a6db3	Alignment		7.3	22	Fold: GroEL-intermediate domain like Superfamily: GroEL-intermediate domain like Family: Group II chaperonin (CCT, TRIC), intermediate domain
13	d1a6da3	Alignment		6.8	22	Fold: GroEL-intermediate domain like Superfamily: GroEL-intermediate domain like Family: Group II chaperonin (CCT, TRIC), intermediate domain
14	d1s5qb_	Alignment		6.2	12	Fold: PAH2 domain Superfamily: PAH2 domain Family: PAH2 domain
15	c2l9vA_	Alignment		6.1	15	PDB header: rna binding protein Chain: A; PDB Molecule: pre-mrna-processing factor 40 homolog a; PDBTitle: nmr structure of the ff domain l24a mutant's folding transition state

16	d1uoua2	Alignment		5.2	24	<p>Fold: Nucleoside phosphorylase/phosphoribosyltransferase catalytic domain</p> <p>Superfamily: Nucleoside phosphorylase/phosphoribosyltransferase catalytic domain</p> <p>Family: Nucleoside phosphorylase/phosphoribosyltransferase catalytic domain</p>
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