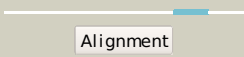

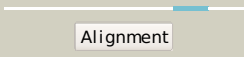

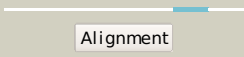

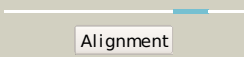

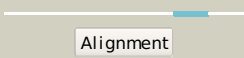

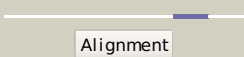

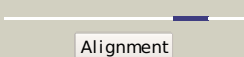
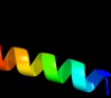
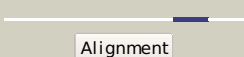



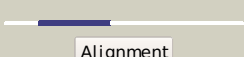

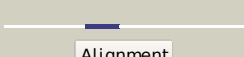


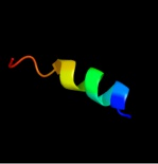


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1cksa_	 Alignment		35.9	67	Fold: Cell cycle regulatory proteins Superfamily: Cell cycle regulatory proteins Family: Cell cycle regulatory proteins
2	d2astc1	 Alignment		35.1	67	Fold: Cell cycle regulatory proteins Superfamily: Cell cycle regulatory proteins Family: Cell cycle regulatory proteins
3	d1puca_	 Alignment		33.3	56	Fold: Cell cycle regulatory proteins Superfamily: Cell cycle regulatory proteins Family: Cell cycle regulatory proteins
4	c1qb3B_	 Alignment		31.5	56	PDB header: cell cycle Chain: B: PDB Molecule: cyclin-dependent kinases regulatory subunit; PDBTitle: crystal structure of the cell cycle regulatory protein cks1
5	d1qb3a_	 Alignment		30.6	56	Fold: Cell cycle regulatory proteins Superfamily: Cell cycle regulatory proteins Family: Cell cycle regulatory proteins
6	d2nvna1	 Alignment		10.4	28	Fold: ssDNA-binding transcriptional regulator domain Superfamily: ssDNA-binding transcriptional regulator domain Family: PMN2A0962/syc2379c-like
7	d2it9a1	 Alignment		9.7	28	Fold: ssDNA-binding transcriptional regulator domain Superfamily: ssDNA-binding transcriptional regulator domain Family: PMN2A0962/syc2379c-like
8	c1wq6A_	 Alignment		8.0	36	PDB header: oncoprotein Chain: A: PDB Molecule: aml1-eto; PDBTitle: the tetramer structure of the nervy homology two (nhr2) domain of aml1-2 eto is critical for aml1-eto's activity
9	d1na6a1	 Alignment		7.7	28	Fold: DNA-binding pseudobarrel domain Superfamily: DNA-binding pseudobarrel domain Family: Type II restriction endonuclease effector domain
10	d3cx5c1	 Alignment		6.3	16	Fold: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Superfamily: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Family: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase)
11	d1d1da1	 Alignment		5.3	36	Fold: Acyl carrier protein-like Superfamily: Retrovirus capsid dimerization domain-like Family: Retrovirus capsid protein C-terminal domain

12	dlin1a_	Alignment		5.3	33	Fold: BRCT domain Superfamily: BRCT domain Family: DNA ligase
13	c1z9eA_	Alignment		5.3	7	PDB header: protein binding/transcription Chain: A: PDB Molecule: pc4 and sfrs1 interacting protein 2; PDBTitle: solution structure of the hiv-1 integrase-binding domain in2 ledgf/p75

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[c3c8iA_](#)

Alignment



5.1

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PDB header:membrane protein
Chain: A: **PDB Molecule:**putative membrane protein;
PDBTitle: crystal structure of a putative membrane protein from corynebacterium2 diphtheriae