






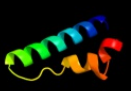







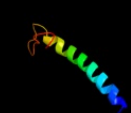

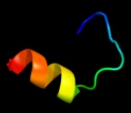

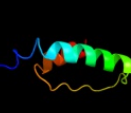




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1ej7l2	 Alignment		19.6	15	Fold: Ferredoxin-like Superfamily: RuBisCO, large subunit, small (N-terminal) domain Family: Ribulose 1,5-bisphosphate carboxylase-oxygenase
2	c1onvB_	 Alignment		17.4	44	PDB header: transcription Chain: B: PDB Molecule: serine phosphatase fcp1a; PDBTitle: nmr structure of a complex containing the tiiif subunit2 rap74 and the rnap ii ctd phosphatase fcp1
3	c2kouA_	 Alignment		16.9	15	PDB header: hydrolase Chain: A: PDB Molecule: dicer-like protein 4; PDBTitle: dicer like protein
4	c2xpnA_	 Alignment		15.6	13	PDB header: transcription Chain: A: PDB Molecule: iws1; PDBTitle: crystal structure of a spt6-iws1(spn1) complex from2 encephalitozoon cuniculi, form i
5	d1in0a2	 Alignment		11.1	19	Fold: Ferredoxin-like Superfamily: YajQ-like Family: YajQ-like
6	c3h05A_	 Alignment		10.4	16	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein vpa0413; PDBTitle: the crystal structure of a putative nicotinate-nucleotide2 adenylyltransferase from vibrio parahaemolyticus
7	d1wdda2	 Alignment		8.4	15	Fold: Ferredoxin-like Superfamily: RuBisCO, large subunit, small (N-terminal) domain Family: Ribulose 1,5-bisphosphate carboxylase-oxygenase
8	c2a7oA_	 Alignment		8.1	27	PDB header: transcription Chain: A: PDB Molecule: huntingtin interacting protein b; PDBTitle: solution structure of the hset2/hypb sri domain
9	d2r7ga1	 Alignment		7.6	17	Fold: Cyclin-like Superfamily: Cyclin-like Family: Retinoblastoma tumor suppressor domains
10	d5mdha1	 Alignment		6.3	11	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: LDH N-terminal domain-like
11	d7mdha1	 Alignment		5.6	21	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: LDH N-terminal domain-like

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[c1q90G](#)

Alignment



5.6

50

PDB header:photosynthesis
Chain: G: **PDB Molecule:**cytochrome b6f complex subunit petg;
PDBTitle: structure of the cytochrome b6f (plastoquinone :
plastocyanin2 oxidoreductase) from chlamydomonas reinhardtii

13	d1q90g_	Alignment		5.6	50	Fold: Single transmembrane helix Superfamily: PetG subunit of the cytochrome b6f complex Family: PetG subunit of the cytochrome b6f complex
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