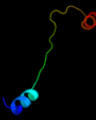


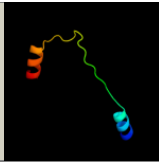
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlwiha_	 Alignment		10.4	27	Fold: RRF/tRNA synthetase additional domain-like Superfamily: Ribosome recycling factor, RRF Family: Ribosome recycling factor, RRF
2	dlxcoa_	 Alignment		9.9	29	Fold: Isocitrate/Isopropylmalate dehydrogenase-like Superfamily: Isocitrate/Isopropylmalate dehydrogenase-like Family: Phosphotransacetylase
3	dlr5ja_	 Alignment		9.8	29	Fold: Isocitrate/Isopropylmalate dehydrogenase-like Superfamily: Isocitrate/Isopropylmalate dehydrogenase-like Family: Phosphotransacetylase
4	c3ke2A_	 Alignment		9.3	52	PDB header: unknown function Chain: A: PDB Molecule: uncharacterized protein yp_928783.1; PDBTitle: crystal structure of a duf2131 family protein (sama_2911) from <i>Shewanella amazonensis</i> sb2b at 2.50 Å resolution
5	c2k9iB_	 Alignment		8.9	65	PDB header: dna binding protein Chain: B: PDB Molecule: plasmid prn1, complete sequence; PDBTitle: nmr structure of plasmid copy control protein orf56 from <i>Sulfolobus islandicus</i>
6	c3ogiC_	 Alignment		8.0	50	PDB header: structural genomics, unknown function Chain: C: PDB Molecule: putative esat-6-like protein 6; PDBTitle: crystal structure of the mycobacterium tuberculosis h37rv esxop2 complex (rv2346c-rv2347c)
7	dlwqga1	 Alignment		7.9	27	Fold: RRF/tRNA synthetase additional domain-like Superfamily: Ribosome recycling factor, RRF Family: Ribosome recycling factor, RRF
8	dljqna_	 Alignment		7.7	13	Fold: TIM beta/alpha-barrel Superfamily: Phosphoenolpyruvate/pyruvate domain Family: Phosphoenolpyruvate carboxylase
9	c2hzsL_	 Alignment		7.1	70	PDB header: transcription Chain: L: PDB Molecule: rna polymerase ii mediator complex subunit 8; PDBTitle: structure of the mediator head submodule med8c/18/20
10	c2hzsJ_	 Alignment		7.1	70	PDB header: transcription Chain: J: PDB Molecule: rna polymerase ii mediator complex subunit 8; PDBTitle: structure of the mediator head submodule med8c/18/20
11	dlvl7a1	 Alignment		6.9	33	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain

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[cljqoA_](#)



Alignment



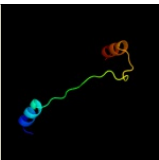
6.9

26

PDB header: lyase
Chain: A: **PDB Molecule:** phosphoenolpyruvate carboxylase;
PDB Title: crystal structure of c4-form phosphoenolpyruvate
carboxylase from2 maize

13 [dljqoa_](#)

Alignment



6.9

26

Fold:TIM beta/alpha-barrel
Superfamily:Phosphoenolpyruvate/pyruvate domain
Family:Phosphoenolpyruvate carboxylase