

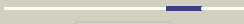
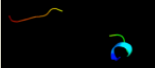



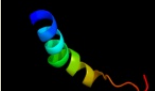




Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P64564
Date	Thu Jan 5 12:09:29 GMT 2012
Unique Job ID	d9d23adf9f05c858

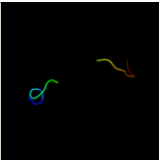
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlizna_	 Alignment		10.0	21	Fold: Subunits of heterodimeric actin filament capping protein Capz Superfamily: Subunits of heterodimeric actin filament capping protein Capz Family: Capz alpha-1 subunit
2	dlvkoa1	 Alignment		7.5	33	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain
3	c3rruA_	 Alignment		7.2	17	PDB header: signaling protein Chain: A: PDB Molecule: tom1l1 protein; PDBTitle: x-ray crystal structure of the vhs domain of human tom1-like protein,2 northeast structural genomics consortium target hr3050e
4	c3lvwA_	 Alignment		6.5	7	PDB header: ligase Chain: A: PDB Molecule: glutamate--cysteine ligase; PDBTitle: glutathione-inhibited scgcl
5	dlnp7a1	 Alignment		6.4	21	Fold: Cryptochrome/photolyase FAD-binding domain Superfamily: Cryptochrome/photolyase FAD-binding domain Family: Cryptochrome/photolyase FAD-binding domain

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[dlp1ja1](#)

Alignment



5.7

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Fold:NAD(P)-binding Rossmann-fold domains
Superfamily:NAD(P)-binding Rossmann-fold domains
Family:Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain