
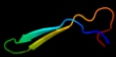








# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P76164
Date	Thu Jan 5 12:19:59 GMT 2012
Unique Job ID	f696e157d35d9074

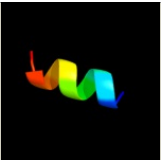
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3glaA_</a>	 Alignment		20.4	22	<b>PDB header:</b> chaperone <b>Chain:</b> A: <b>PDB Molecule:</b> low molecular weight heat shock protein; <b>PDBTitle:</b> crystal structure of the hspa from xanthomonas axonopodis
2	<a href="#">d1bdfa1</a>	 Alignment		13.7	21	<b>Fold:</b> DCoH-like <b>Superfamily:</b> RBP11-like subunits of RNA polymerase <b>Family:</b> RNA polymerase alpha subunit dimerisation domain
3	<a href="#">c2kcdA_</a>	 Alignment		12.7	27	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein ssp0047; <b>PDBTitle:</b> solution nmr structure of ssp0047 from staphylococcus2 saprophyticus. northeast structural genomics consortium3 target syr6.
4	<a href="#">c2pq4B_</a>	 Alignment		6.6	67	<b>PDB header:</b> chaperone/oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> periplasmic nitrate reductase precursor; <b>PDBTitle:</b> nmr solution structure of napd in complex with napa1-352 signal peptide

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[clavyA](#)

Alignment



6.2

73

**PDB header:**coiled coil  
**Chain:** A: **PDB Molecule:**fibrin;  
**PDBTitle:** fibrin deletion mutant m (bacteriophage t4)