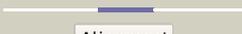
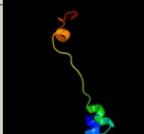
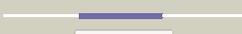


# Phyre2

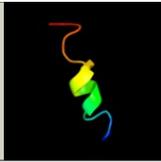
Email	l.a.kelley@imperial.ac.uk
Description	P75786
Date	Wed Jan 25 15:21:02 GMT 2012
Unique Job ID	02fed6107ac5e23d

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3lw5K_</a>	 Alignment		17.1	44	<b>PDB header:</b> photosynthesis <b>Chain:</b> K; <b>PDB Molecule:</b> photosystem i reaction center subunit x psak; <b>PDBTitle:</b> improved model of plant photosystem i
2	<a href="#">d1e8ga1</a>	 Alignment		15.1	36	<b>Fold:</b> Ferredoxin-like <b>Superfamily:</b> FAD-linked oxidases, C-terminal domain <b>Family:</b> Vanillyl-alcohol oxidase-like
3	<a href="#">d2hkva1</a>	 Alignment		11.8	28	<b>Fold:</b> DinB/YfiT-like putative metalloenzymes <b>Superfamily:</b> DinB/YfiT-like putative metalloenzymes <b>Family:</b> DinB-like
4	<a href="#">d2q66a2</a>	 Alignment		11.5	18	<b>Fold:</b> Nucleotidyltransferase <b>Superfamily:</b> Nucleotidyltransferase <b>Family:</b> Poly(A) polymerase, PAP, N-terminal domain
5	<a href="#">c3k13A_</a>	 Alignment		10.9	20	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> 5-methyltetrahydrofolate-homocysteine methyltransferase; <b>PDBTitle:</b> structure of the pterin-binding domain metr of 5-2 methyltetrahydrofolate-homocysteine methyltransferase from3 bacteroides thetaiotaomicron
6	<a href="#">c2cmpA_</a>	 Alignment		8.1	29	<b>PDB header:</b> terminase <b>Chain:</b> A; <b>PDB Molecule:</b> terminase small subunit; <b>PDBTitle:</b> crystal structure of the dna binding domain of g1p small2 terminase subunit from bacteriophage sf6
7	<a href="#">c1ahuB_</a>	 Alignment		6.9	36	<b>PDB header:</b> flavoenzyme <b>Chain:</b> B; <b>PDB Molecule:</b> vanillyl-alcohol oxidase; <b>PDBTitle:</b> structure of the octameric flavoenzyme vanillyl-alcohol2 oxidase in complex with p-cresol
8	<a href="#">c3c66B_</a>	 Alignment		6.3	18	<b>PDB header:</b> transferase <b>Chain:</b> B; <b>PDB Molecule:</b> poly(a) polymerase; <b>PDBTitle:</b> yeast poly(a) polymerase in complex with fip1 residues 80-105
9	<a href="#">c2g9lA_</a>	 Alignment		5.7	43	<b>PDB header:</b> antibiotic <b>Chain:</b> A; <b>PDB Molecule:</b> gaegurin-4; <b>PDBTitle:</b> the high-resolution solution conformation of an2 antimicrobial peptide gaegurin 4 and its mode of membrane3 interaction

10 [c2wsfG\\_](#)

Alignment



5.3

44

**PDB header:** photosynthesis  
**Chain:** G: **PDB Molecule:** photosystem i reaction center subunit v.  
**PDB Title:** improved model of plant photosystem i