


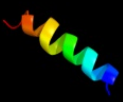
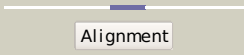

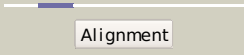

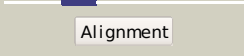



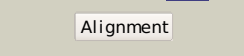
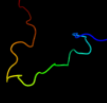
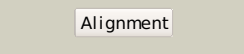

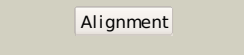

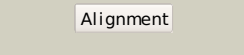

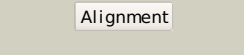



Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P27375
Date	Thu Jan 5 11:43:57 GMT 2012
Unique Job ID	1f515025f65cfa93

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1kl9a1	 Alignment		37.4	13	Fold: SAM domain-like Superfamily: eIF2alpha middle domain-like Family: eIF2alpha middle domain-like
2	d1q46a1	 Alignment		27.2	20	Fold: SAM domain-like Superfamily: eIF2alpha middle domain-like Family: eIF2alpha middle domain-like
3	c2vtyA	 Alignment		18.8	25	PDB header: apoptosis Chain: A: PDB Molecule: protein f1; PDBTitle: vaccinia virus anti-apoptotic f1l is a novel bcl-2-like2 domain swapped dimer
4	d1u84a	 Alignment		12.7	26	Fold: YugE-like Superfamily: YugE-like Family: YugE-like
5	d1fcda3	 Alignment		9.4	25	Fold: CO dehydrogenase flavoprotein C-domain-like Superfamily: FAD/NAD-linked reductases, dimerisation (C-terminal) domain Family: FAD/NAD-linked reductases, dimerisation (C-terminal) domain
6	c3doaA	 Alignment		7.5	14	PDB header: protein binding Chain: A: PDB Molecule: fibrinogen binding protein; PDBTitle: the crystal structure of the fibrinogen binding protein from2 staphylococcus aureus
7	c2q8aA	 Alignment		7.4	12	PDB header: immune system Chain: A: PDB Molecule: apical membrane antigen 1; PDBTitle: structure of the malaria antigen ama1 in complex with a growth-2 inhibitory antibody
8	d1sqwa1	 Alignment		7.0	23	Fold: PUA domain-like Superfamily: PUA domain-like Family: PUA domain
9	d1cz5a1	 Alignment		6.1	18	Fold: Double psi beta-barrel Superfamily: ADC-like Family: Cdc48 N-terminal domain-like
10	d1mv8a1	 Alignment		5.9	30	Fold: 6-phosphogluconate dehydrogenase C-terminal domain-like Superfamily: 6-phosphogluconate dehydrogenase C-terminal domain-like Family: UDP-glucose/GDP-mannose dehydrogenase dimerisation domain
11	d1dzfa2	 Alignment		5.7	24	Fold: RPB5-like RNA polymerase subunit Superfamily: RPB5-like RNA polymerase subunit Family: RPB5

12

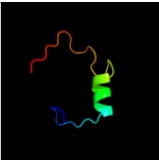
d1tp6a_

Alignment

5.6

16

Fold:Cystatin-like
Superfamily:NTF2-like
Family:PA1314-like



13 [dlz67a1](#)

Alignment



5.6

44

Fold:YidB-like
Superfamily:YidB-like
Family:YidB-like