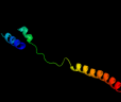
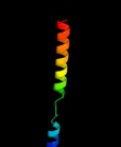


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P68206
Date	Thu Jan 5 12:11:00 GMT 2012
Unique Job ID	c36e066ba9e468ff

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1ryka_	 Alignment		100.0	100	Fold: SAM domain-like Superfamily: Hypothetical protein Yjbj Family: Hypothetical protein Yjbj
2	c1ywwA_	 Alignment		99.9	55	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein pa4738; PDBTitle: nmr structure of p. aeruginosa protein pa4738: northeast2 structural genomics consortium target pap2
3	c2l9qA_	 Alignment		16.1	19	PDB header: chaperone Chain: A: PDB Molecule: 12 kda heat shock protein; PDBTitle: structural characterization of small heat shock protein (hsp12)
4	c2js5B_	 Alignment		9.8	16	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: nmr structure of protein q60c73_metca. northeast structural2 genomics consortium target mcr1
5	c2k88A_	 Alignment		8.4	21	PDB header: hydrolase Chain: A: PDB Molecule: vacuolar proton pump subunit g; PDBTitle: association of subunit d (vma6p) and e (vma4p) with g2 (vma10p) and the nmr solution structure of subunit g (g1-3 59) of the saccharomyces cerevisiae v1vo atpase

6

[c3lf9A_](#)

Alignment



5.7

11

PDB header:immune system
Chain: A: **PDB Molecule:**4e10_d0_lis1a_001_c (t161);
PDBTitle: crystal structure of hiv epitope-scaffold
4e10_d0_lis1a_001_c