

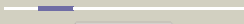

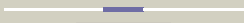
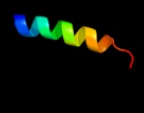








Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0AAP7
Date	Thu Jan 5 11:13:27 GMT 2012
Unique Job ID	057bef0fd1389fe5

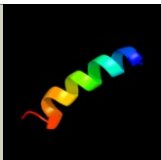
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1b9uA_	 Alignment		40.2	56	PDB header: hydrolase Chain: A: PDB Molecule: protein (atp synthase); PDBTitle: membrane domain of the subunit b of the e.coli atp synthase
2	d1q90n_	 Alignment		13.3	40	Fold: Single transmembrane helix Superfamily: PetN subunit of the cytochrome b6f complex Family: PetN subunit of the cytochrome b6f complex
3	c3mq1A_	 Alignment		10.5	32	PDB header: allergen Chain: A: PDB Molecule: mite allergen der p 5; PDBTitle: crystal structure of dust mite allergen der p 5
4	c2jmhA_	 Alignment		9.5	21	PDB header: allergen Chain: A: PDB Molecule: mite allergen blo t 5; PDBTitle: nmr solution structure of blo t 5, a major mite allergen2 from blomia tropicalis
5	d2i7na2	 Alignment		7.7	53	Fold: Ribonuclease H-like motif Superfamily: Actin-like ATPase domain Family: Fumble-like
6	d2axtm1	 Alignment		5.6	50	Fold: Single transmembrane helix Superfamily: Photosystem II reaction center protein M, PsbM Family: PsbM-like

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[c2rocB_](#)

Alignment



5.4

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PDB header:apoptosis

Chain: B: **PDB Molecule:**bcl-2-binding component 3;

PDBTitle: solution structure of mcl-1 complexed with puma