

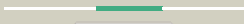






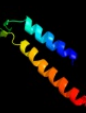




Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P46119
Date	Wed Jan 25 15:20:56 GMT 2012
Unique Job ID	8e735132a8d099dc

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2y69Z_	 Alignment		48.1	30	PDB header: electron transport Chain: Z: PDB Molecule: cytochrome c oxidase polypeptide 8h; PDBTitle: bovine heart cytochrome c oxidase re-refined with molecular2 oxygen
2	d1v54m_	 Alignment		40.3	30	Fold: Single transmembrane helix Superfamily: Mitochondrial cytochrome c oxidase subunit VIIIb (aka IX) Family: Mitochondrial cytochrome c oxidase subunit VIIIb (aka IX)
3	c2jwaA_	 Alignment		10.2	29	PDB header: transferase Chain: A: PDB Molecule: receptor tyrosine-protein kinase erbb-2; PDBTitle: erbb2 transmembrane segment dimer spatial structure
4	c3fhgA_	 Alignment		9.8	10	PDB header: dna repair, hydrolase, lyase Chain: A: PDB Molecule: n-glycosylase/dna lyase; PDBTitle: crystal structure of sulfolobus solfataricus 8-oxoguanine dna2 glycosylase (ssogg)
5	c3dwwA_	 Alignment		7.9	17	PDB header: isomerase Chain: A: PDB Molecule: prostaglandin e synthase; PDBTitle: electron crystallographic structure of human microsomal2 prostaglandin e synthase 1
6	c1wu0A_	 Alignment		6.0	5	PDB header: hydrolase Chain: A: PDB Molecule: atp synthase c chain; PDBTitle: solution structure of subunit c of f1fo-atp synthase from2 the thermophilic bacillus ps3

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[c2wpdP_](#)

Alignment



5.8

29

PDB header:hydrolase
Chain: P: **PDB Molecule:**atp synthase subunit 9, mitochondrial;
PDBTitle: the mg.adp inhibited state of the yeast f1c10 atp synthase