









Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P56100
Date	Thu Jan 5 12:06:17 GMT 2012
Unique Job ID	f7568aaf6bb7901a

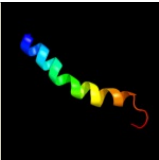
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2e74b1	 Alignment		11.5	30	Fold: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Superfamily: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Family: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase)
2	d1q90d	 Alignment		11.3	40	Fold: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Superfamily: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Family: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase)
3	d1vf5b	 Alignment		7.2	30	Fold: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Superfamily: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Family: a domain/subunit of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase)
4	c2nr1A	 Alignment		5.6	46	PDB header: receptor Chain: A: PDB Molecule: nr1 m2; PDBTitle: transmembrane segment 2 of nmda receptor nr1, nmr, 102 structures

5

[c3mk7B_](#)

Alignment



5.5

21

PDB header:oxidoreductase
Chain: B: **PDB Molecule:**cytochrome c oxidase, cbb3-type, subunit
o;
PDBTitle: the structure of cbb3 cytochrome oxidase