

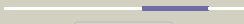


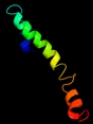








Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P60632
Date	Thu Jan 5 12:06:59 GMT 2012
Unique Job ID	c188e210a7e5b39b

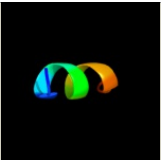
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kncA_	 Alignment		28.1	15	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iiB; PDBTitle: platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
2	d1fftb2	 Alignment		15.0	9	Fold: Transmembrane helix hairpin Superfamily: Cytochrome c oxidase subunit II-like, transmembrane region Family: Cytochrome c oxidase subunit II-like, transmembrane region
3	d1iwga8	 Alignment		12.3	13	Fold: Multidrug efflux transporter AcrB transmembrane domain Superfamily: Multidrug efflux transporter AcrB transmembrane domain Family: Multidrug efflux transporter AcrB transmembrane domain
4	d1veka_	 Alignment		10.2	25	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: UBA domain
5	c2l2tA_	 Alignment		9.8	9	PDB header: membrane protein Chain: A: PDB Molecule: receptor tyrosine-protein kinase erbb-4; PDBTitle: solution nmr structure of the erbb4 dimeric membrane domain
6	d2oara1	 Alignment		7.3	13	Fold: Gated mechanosensitive channel Superfamily: Gated mechanosensitive channel Family: Gated mechanosensitive channel

7

[d2cpwa1](#)

Alignment



6.1

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Fold:RuvA C-terminal domain-like
Superfamily:UBA-like
Family:UBA domain