




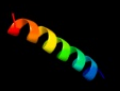


# Phyre2

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Description	Q46755
Date	Thu Jan 5 12:34:05 GMT 2012
Unique Job ID	ebfe6f51e9a2e92f

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1fftb2</a>	 Alignment		5.7	13	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Cytochrome c oxidase subunit II-like, transmembrane region <b>Family:</b> Cytochrome c oxidase subunit II-like, transmembrane region
2	<a href="#">dls7ba_</a>	 Alignment		5.4	7	<b>Fold:</b> Multidrug resistance efflux transporter EmrE <b>Superfamily:</b> Multidrug resistance efflux transporter EmrE <b>Family:</b> Multidrug resistance efflux transporter EmrE
3	<a href="#">c3dl8D_</a>	 Alignment		5.3	39	<b>PDB header:</b> protein transport <b>Chain:</b> D: <b>PDB Molecule:</b> sece; <b>PDB Title:</b> structure of the complex of aquifex aeolicus secyeg and2 bacillus subtilis seca