
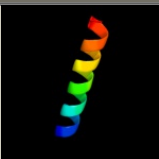






# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P31433
Date	Thu Jan 5 11:47:23 GMT 2012
Unique Job ID	841ddba48408f054

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1m56d_</a>	 Alignment		8.1	14	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Bacterial aa3 type cytochrome c oxidase subunit IV <b>Family:</b> Bacterial aa3 type cytochrome c oxidase subunit IV
2	<a href="#">c2latA_</a>	 Alignment		7.5	13	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> dolichyl-diphosphooligosaccharide--protein <b>PDBTitle:</b> solution structure of a human minimembrane protein ost4
3	<a href="#">c2hg5D_</a>	 Alignment		5.5	14	<b>PDB header:</b> membrane protein <b>Chain:</b> D: <b>PDB Molecule:</b> kcsa channel; <b>PDBTitle:</b> cs+ complex of a k channel with an amide to ester substitution in the2 selectivity filter