


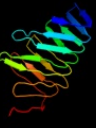

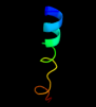


# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0ADV9
Date	Wed Jan 25 15:20:29 GMT 2012
Unique Job ID	49bc9337c2dc215c

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3my2A_</a>	 Alignment		100.0	98	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> lipopolysaccharide export system protein lptc; <b>PDBTitle:</b> crystal structure of lptc
2	<a href="#">c2r1aD_</a>	 Alignment		98.6	17	<b>PDB header:</b> transport protein <b>Chain:</b> D: <b>PDB Molecule:</b> protein yhbn; <b>PDBTitle:</b> crystal structure of the periplasmic lipopolysaccharide transport2 protein lpta (yhbn), trigonal form
3	<a href="#">c2voyE_</a>	 Alignment		8.2	21	<b>PDB header:</b> hydrolase <b>Chain:</b> E: <b>PDB Molecule:</b> sarcoplasmic/endoplasmic reticulum calcium <b>PDBTitle:</b> cryoem model of copa, the copper transporting atpase from2 archaeoglobus fulgidus