

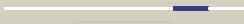
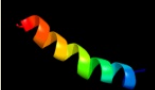

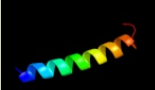




# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P75920
Date	Thu Jan 5 12:16:04 GMT 2012
Unique Job ID	b4d77cebf94ce076

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2wwbB_</a>	 Alignment		16.7	22	<b>PDB header:</b> ribosome <b>Chain:</b> B: <b>PDB Molecule:</b> protein transport protein sec61 subunit gamma; <b>PDBTitle:</b> cryo-em structure of the mammalian sec61 complex bound to the2 actively translating wheat germ 80s ribosome
2	<a href="#">d1rhzb_</a>	 Alignment		7.7	5	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Preprotein translocase SecE subunit <b>Family:</b> Preprotein translocase SecE subunit
3	<a href="#">d1ijdb_</a>	 Alignment		5.9	14	<b>Fold:</b> Light-harvesting complex subunits <b>Superfamily:</b> Light-harvesting complex subunits <b>Family:</b> Light-harvesting complex subunits
4	<a href="#">c3a98B_</a>	 Alignment		5.5	19	<b>PDB header:</b> signaling protein <b>Chain:</b> B: <b>PDB Molecule:</b> engulfment and cell motility protein 1; <b>PDBTitle:</b> crystal structure of the complex of the interacting regions of dock22 and elmo1