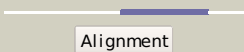
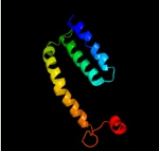
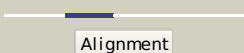

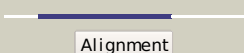
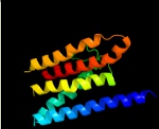




# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0AAC4
Date	Thu Jan 5 11:12:32 GMT 2012
Unique Job ID	79fd1a2c5c4fbc6

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d2r6gf1</a>	 Alignment		13.6	11	<b>Fold:</b> MalF N-terminal region-like <b>Superfamily:</b> MalF N-terminal region-like <b>Family:</b> MalF N-terminal region-like
2	<a href="#">c2k9pA_</a>	 Alignment		6.7	14	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> pheromone alpha factor receptor; <b>PDBTitle:</b> structure of tm1_tm2 in lppg micelles
3	<a href="#">c3k07A_</a>	 Alignment		6.3	9	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> cation efflux system protein cusa; <b>PDBTitle:</b> crystal structure of cusa
4	<a href="#">c2h3oA_</a>	 Alignment		5.4	25	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> merf; <b>PDBTitle:</b> structure of merft, a membrane protein with two trans-2 membrane helices