
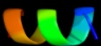
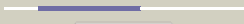







# Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P0ACX5
Date	Thu Jan 5 11:19:21 GMT 2012
Unique Job ID	12d21c351499129b

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3fxhA_</a>	 Alignment		13.8	55	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> integron gene cassette protein hfx_cass2; <b>PDBTitle:</b> crystal structure from the mobile metagenome of halifax2 harbour sewage outfall: integron cassette protein hfx_cass2
2	<a href="#">c2wl8D_</a>	 Alignment		11.3	21	<b>PDB header:</b> protein transport <b>Chain:</b> D: <b>PDB Molecule:</b> peroxisomal biogenesis factor 19; <b>PDBTitle:</b> x-ray crystal structure of pex19p
3	<a href="#">dls4na_</a>	 Alignment		8.9	75	<b>Fold:</b> Nucleotide-diphospho-sugar transferases <b>Superfamily:</b> Nucleotide-diphospho-sugar transferases <b>Family:</b> Glycolipid 2-alpha-mannosyltransferase
4	<a href="#">c1xrxD_</a>	 Alignment		5.9	67	<b>PDB header:</b> replication inhibitor <b>Chain:</b> D: <b>PDB Molecule:</b> seqa protein; <b>PDBTitle:</b> crystal structure of a dna-binding protein