









# Phyre2

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Description	P58036
Date	Thu Jan 5 12:06:34 GMT 2012
Unique Job ID	ce588c9600b3e68b

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3fcnA_</a>	 Alignment		11.2	38	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> an alpha-helical protein of unknown function (pfam01724); <b>PDBTitle:</b> crystal structure of an alpha-helical protein of unknown function2 (rru_a3208) from rhodospirillum rubrum atcc 11170 at 1.45 a3 resolution
2	<a href="#">c3owtC_</a>	 Alignment		8.9	67	<b>PDB header:</b> protein binding <b>Chain:</b> C: <b>PDB Molecule:</b> regulatory protein sir3; <b>PDBTitle:</b> crystal structure of s. cerevisiae rap1-sir3 complex
3	<a href="#">d1hs7a_</a>	 Alignment		6.9	50	<b>Fold:</b> STAT-like <b>Superfamily:</b> t-snare proteins <b>Family:</b> t-snare proteins
4	<a href="#">c2ae8C_</a>	 Alignment		5.8	32	<b>PDB header:</b> lyase <b>Chain:</b> C: <b>PDB Molecule:</b> imidazoleglycerol-phosphate dehydratase; <b>PDBTitle:</b> crystal structure of imidazoleglycerol-phosphate dehydratase from2 staphylococcus aureus subsp. aureus n315