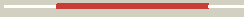
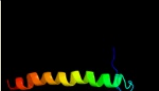
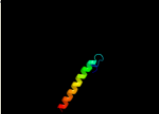


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P26647
Date	Thu Jan 5 11:43:05 GMT 2012
Unique Job ID	f19bd30012d292e8

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dliuqa_	 Alignment		98.6	16	Fold: Glycerol-3-phosphate (1)-acyltransferase Superfamily: Glycerol-3-phosphate (1)-acyltransferase Family: Glycerol-3-phosphate (1)-acyltransferase
2	c2ke4A_	 Alignment		16.7	26	PDB header: membrane protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: the nmr structure of the tc10 and cdc42 interacting domain2 of cip4
3	c1ybxA_	 Alignment		7.6	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: conserved hypothetical protein; PDBTitle: conserved hypothetical protein cth-383 from clostridium thermocellum