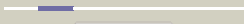


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P76158
Date	Thu Jan 5 12:19:50 GMT 2012
Unique Job ID	431a3b0355678bca

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kpeB_	 Alignment		10.4	43	PDB header: membrane protein Chain: B: PDB Molecule: glycophorin-a; PDBTitle: refined structure of glycophorin a transmembrane segment dimer in dpc2 micelles
2	c2kpeA_	 Alignment		10.4	43	PDB header: membrane protein Chain: A: PDB Molecule: glycophorin-a; PDBTitle: refined structure of glycophorin a transmembrane segment dimer in dpc2 micelles
3	d2ay0a1	 Alignment		7.9	32	Fold: Ribbon-helix-helix Superfamily: Ribbon-helix-helix Family: PutA pre-N-terminal region-like
4	c3gmbB_	 Alignment		6.0	13	PDB header: oxidoreductase Chain: B: PDB Molecule: 2-methyl-3-hydroxypyridine-5-carboxylic acid PDBTitle: crystal structure of 2-methyl-3-hydroxypyridine-5-carboxylic2 acid oxygenase