

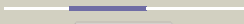





Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0ADM4
Date	Thu Jan 5 11:21:21 GMT 2012
Unique Job ID	96ec33ffed8c9a90

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2pxgA_	 Alignment		14.2	12	PDB header: membrane protein Chain: A: PDB Molecule: outer membrane protein; PDBTitle: nmr solution structure of omla
2	c3sqgG_	 Alignment		11.2	31	PDB header: transferase Chain: G: PDB Molecule: methyl coenzyme m reductase, alpha subunit; PDBTitle: crystal structure of a methyl-coenzyme m reductase purified from black2 sea mats
3	dlzkda1	 Alignment		9.9	12	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: RPA4359-like