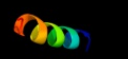


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0ADB4
Date	Thu Jan 5 11:20:31 GMT 2012
Unique Job ID	02548d1c05665d08

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2rh3a1	 Alignment		25.9	24	Fold: Ribbon-helix-helix Superfamily: Ribbon-helix-helix Family: VirC2-like
2	c1pyvA_	 Alignment		11.6	50	PDB header: hydrolase Chain: A: PDB Molecule: atp synthase beta chain, mitochondrial precursor; PDBTitle: nmr solution structure of the mitochondrial f1b presequence2 peptide from nicotiana plumbaginifolia
3	d1sp8a2	 Alignment		9.0	38	Fold: Glyoxalase/Bleomycin resistance protein/Dihydroxybiphenyl dioxygenase Superfamily: Glyoxalase/Bleomycin resistance protein/Dihydroxybiphenyl dioxygenase Family: Extradiol dioxygenases
4	c1debA_	 Alignment		7.4	35	PDB header: structural protein Chain: A: PDB Molecule: adenomatous polyposis coli protein; PDBTitle: crystal structure of the n-terminal coiled coil domain from2 apc

5

[d1nh9a_](#)

Alignment



6.7

43

Fold:IF3-like
Superfamily:AlbA-like
Family:DNA-binding protein AlbA