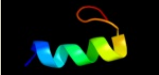
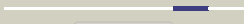


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P76460
Date	Thu Jan 5 12:23:05 GMT 2012
Unique Job ID	71592f38bb7fbec2

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3myrB_	 Alignment		7.1	16	PDB header: oxidoreductase Chain: B: PDB Molecule: nickel-dependent hydrogenase large subunit; PDBTitle: crystal structure of [nife] hydrogenase from allochromatium vinosum in2 its ni-a state
2	d1dv0a_	 Alignment		7.0	26	Fold: RuvA C-terminal domain-like Superfamily: UBA-like Family: UBA domain
3	c3d0wD_	 Alignment		6.4	16	PDB header: structural genomics, unknown function Chain: D: PDB Molecule: yflh protein; PDBTitle: crystal structure of yflh protein from bacillus subtilis.2 northeast structural genomics consortium target sr326