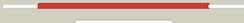
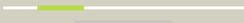
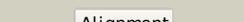
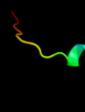


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0AAN1
Date	Thu Jan 5 11:13:20 GMT 2012
Unique Job ID	9dddbd37ed3e3992

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kc5A_	 Alignment		100.0	100	PDB header: chaperone Chain: A; PDB Molecule: hydrogenase-2 operon protein hybe; PDBTitle: solution structure of hybe from escherichia coli
2	d1e6va2	 Alignment		67.0	16	Fold: Ferredoxin-like Superfamily: Methyl-coenzyme M reductase subunits Family: Methyl-coenzyme M reductase alpha and beta chain N-terminal domain
3	d1hbna2	 Alignment		53.8	19	Fold: Ferredoxin-like Superfamily: Methyl-coenzyme M reductase subunits Family: Methyl-coenzyme M reductase alpha and beta chain N-terminal domain
4	c1hbuD_	 Alignment		24.1	19	PDB header: methanogenesis Chain: D; PDB Molecule: methyl-coenzyme m reductase i alpha subunit; PDBTitle: methyl-coenzyme m reductase in the mcr-red1-silent state in2 complex with coenzyme m
5	c1cn3F_	 Alignment		17.3	50	PDB header: viral protein Chain: F; PDB Molecule: fragment of coat protein vp2; PDBTitle: interaction of polyomavirus internal protein vp2 with major2 capsid protein vp1 and implications for participation of3 vp2 in viral entry
6	c2ideE_	 Alignment		7.4	20	PDB header: biosynthetic protein Chain: E; PDB Molecule: molybdenum cofactor biosynthesis protein c; PDBTitle: crystal structure of the molybdenum cofactor biosynthesis protein c2 (ttha1789) from thermus thermophilus hb8
7	c2eevA_	 Alignment		7.2	20	PDB header: biosynthetic protein Chain: A; PDB Molecule: molybdopterin biosynthesis; PDBTitle: structure of gk0241 protein from geobacillus kaustophilus
8	d1ekra_	 Alignment		6.6	13	Fold: Ferredoxin-like Superfamily: Molybdenum cofactor biosynthesis protein C, MoaC Family: Molybdenum cofactor biosynthesis protein C, MoaC

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[c3cmqA](#)

Alignment



6.5

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PDB header:ligase
Chain: A: **PDB Molecule:**phenylalanyl-trna synthetase, mitochondrial;
PDBTitle: crystal structure of human mitochondrial phenylalanine trna2 synthetase