

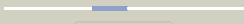













Phyre2

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Description	Q47710
Date	Thu Jan 5 12:37:08 GMT 2012
Unique Job ID	c2a4a1f63d32abaa

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3cazA_	 Alignment		38.1	32	PDB header: signaling protein Chain: A: PDB Molecule: bar protein; PDBTitle: crystal structure of a bar protein from galdieria sulphuraria
2	c3sqgF_	 Alignment		28.7	29	PDB header: transferase Chain: F: PDB Molecule: methyl-coenzyme m reductase, gamma subunit; PDBTitle: crystal structure of a methyl-coenzyme m reductase purified from black2 sea mats
3	c2yh5A_	 Alignment		16.9	30	PDB header: lipid binding protein Chain: A: PDB Molecule: dapx protein; PDBTitle: structure of the c-terminal domain of bamc
4	c3gn4A_	 Alignment		9.8	16	PDB header: motor protein Chain: A: PDB Molecule: myosin-vi; PDBTitle: myosin lever arm
5	c3s6iA_	 Alignment		9.5	14	PDB header: hydrolase/dna Chain: A: PDB Molecule: dna-3-methyladenine glycosylase 1; PDBTitle: schizosaccharomyces pombe 3-methyladenine dna glycosylase (mag1) in2 complex with abasic-dna.
6	c1b0nB_	 Alignment		7.8	19	PDB header: transcription regulator Chain: B: PDB Molecule: protein (sini protein); PDBTitle: sinr protein/sini protein complex
7	d1b0nb_	 Alignment		7.8	19	Fold: Dimerisation interlock Superfamily: SinR repressor dimerisation domain-like Family: SinR repressor dimerisation domain-like

<div>8</div> <div>c2yg8B_</div> <div>Alignment</div>		<div>7.8</div>	<div>12</div> <div>PDB header:hydrolase Chain: B: PDB Molecule:dna-3-methyladenine glycosidase ii, putative; PDBTitle: structure of an unusual 3-methyladenine dna glycosylase ii (2 alka) from deinococcus radiodurans</div>
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