

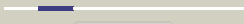









# Phyre2

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Description	P0AET5
Date	Thu Jan 5 11:24:11 GMT 2012
Unique Job ID	8a4c3531e3266136

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1s7ba_</a>	 Alignment		16.3	12	<b>Fold:</b> Multidrug resistance efflux transporter EmrE <b>Superfamily:</b> Multidrug resistance efflux transporter EmrE <b>Family:</b> Multidrug resistance efflux transporter EmrE
2	<a href="#">d1r0ka3</a>	 Alignment		9.9	30	<b>Fold:</b> FwdE/GAPDH domain-like <b>Superfamily:</b> Glyceraldehyde-3-phosphate dehydrogenase-like, C-terminal domain <b>Family:</b> Dihydrodipicolinate reductase-like
3	<a href="#">d1q0qa3</a>	 Alignment		9.0	25	<b>Fold:</b> FwdE/GAPDH domain-like <b>Superfamily:</b> Glyceraldehyde-3-phosphate dehydrogenase-like, C-terminal domain <b>Family:</b> Dihydrodipicolinate reductase-like
4	<a href="#">c3m92B_</a>	 Alignment		8.5	27	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> protein ycin; <b>PDBTitle:</b> the structure of ycin, an uncharacterized protein from shigella2 flexneri.
5	<a href="#">c1pk1B_</a>	 Alignment		6.6	28	<b>PDB header:</b> transcription repression <b>Chain:</b> B: <b>PDB Molecule:</b> sex comb on midleg cg9495-pa; <b>PDBTitle:</b> hetero sam domain structure of ph and scm.

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[dlpk1c1](#)

Alignment



6.5

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**Fold:**SAM domain-like  
**Superfamily:**SAM/Pointed domain  
**Family:**SAM (sterile alpha motif) domain