









# Phyre2

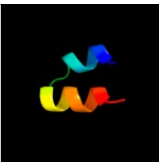
Email	l.a.kelley@imperial.ac.uk
Description	P76548
Date	Thu Jan 5 12:24:25 GMT 2012
Unique Job ID	7c183727ba9db0b5

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1kjna_</a>	 Alignment		7.2	30	<b>Fold:</b> Hypothetical protein MTH777 (MT0777) <b>Superfamily:</b> Hypothetical protein MTH777 (MT0777) <b>Family:</b> Hypothetical protein MTH777 (MT0777)
2	<a href="#">d2o8sa1</a>	 Alignment		6.5	29	<b>Fold:</b> AGR C 984p-like <b>Superfamily:</b> AGR C 984p-like <b>Family:</b> AGR C 984p-like
3	<a href="#">d2ab0a1</a>	 Alignment		6.2	19	<b>Fold:</b> Flavodoxin-like <b>Superfamily:</b> Class I glutamine amidotransferase-like <b>Family:</b> DJ-1/Pfpl
4	<a href="#">c2r5oA_</a>	 Alignment		6.0	14	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> putative atp binding component of abc- <b>PDBTitle:</b> crystal structure of the c-terminal domain of wzt

5 [d2izva1](#)

Alignment



5.0

38

**Fold:**SOCS box-like  
**Superfamily:**SOCS box-like  
**Family:**SOCS box-like