

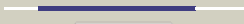
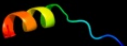
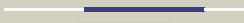





Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	C6Y4B1
Date	Sun Jul 8 11:45:06 BST 2012
Unique Job ID	a273d8081d00b8c3

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1wcga1	 Alignment		11.5	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
2	d1e4mm_	 Alignment		8.4	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
3	c2dzdB_	 Alignment		6.6	23	PDB header: ligase Chain: B: PDB Molecule: pyruvate carboxylase; PDBTitle: crystal structure of the biotin carboxylase domain of2 pyruvate carboxylase
4	c3gidB_	 Alignment		5.3	0	PDB header: ligase Chain: B: PDB Molecule: acetyl-coa carboxylase 2; PDBTitle: the biotin carboxylase (bc) domain of human acetyl-coa2 carboxylase 2 (acc2) in complex with soraphen a