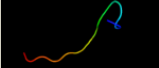
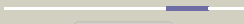


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	A6X977
Date	Sun Jul 8 11:45:02 BST 2012
Unique Job ID	8d1e852689dc89f1

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1m5za_	 Alignment		10.7	33	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
2	d2fyuk1	 Alignment		10.2	17	Fold: Single transmembrane helix Superfamily: Subunit XI (6.4 kDa protein) of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase) Family: Subunit XI (6.4 kDa protein) of cytochrome bc1 complex (Ubiquinol-cytochrome c reductase)
3	d1zu1a2	 Alignment		5.8	38	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: HkH motif-containing C2H2 finger