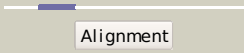
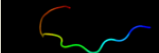
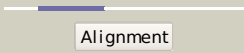

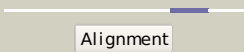



# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	A0A183
Date	Tue Apr 3 14:37:07 BST 2012
Unique Job ID	d0ceb6f85ab3388e

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2xskA_</a>	 Alignment		19.7	29	<b>PDB header:</b> chaperone <b>Chain:</b> A: <b>PDB Molecule:</b> csgc; <b>PDBTitle:</b> e. coli curli protein csgc - secys
2	<a href="#">c3r67A_</a>	 Alignment		12.0	25	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> putative glycosidase; <b>PDBTitle:</b> crystal structure of a putative glycosidase (bt_4094) from bacteroides2 thetaiotaomicron vpi-5482 at 2.30 a resolution
3	<a href="#">d1a7ia1</a>	 Alignment		10.8	43	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LIM domain