

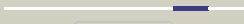
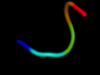

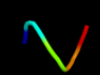

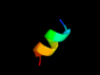


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	C1P5Z8
Date	Thu Jan 5 10:55:48 GMT 2012
Unique Job ID	f12d8f645ffc14bc

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2ovcA_	 Alignment		27.3	64	PDB header: transport protein Chain: A: PDB Molecule: potassium voltage-gated channel subfamily kqt member 4; PDBTitle: crystal structure of a coiled-coil tetramerization domain from kv7.42 channels
2	c1bdeA_	 Alignment		7.2	80	PDB header: aids Chain: A: PDB Molecule: vpr protein; PDBTitle: helical structure of polypeptides from the c-terminal half2 of hiv-1 vpr, nmr, 20 structures
3	c1x9vA_	 Alignment		6.2	80	PDB header: viral protein Chain: A: PDB Molecule: vpr protein; PDBTitle: dimeric structure of the c-terminal domain of vpr
4	c1vryA_	 Alignment		5.8	55	PDB header: membrane protein Chain: A: PDB Molecule: glycine receptor alpha-1 chain; PDBTitle: second and third transmembrane domains of the alpha-12 subunit of human glycine receptor