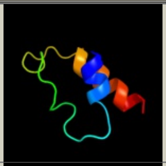
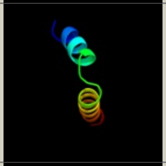
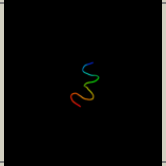
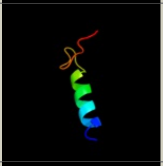
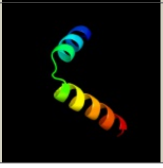
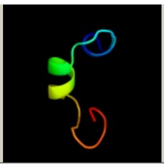
Phyre^2

Email	l.a.kelley@imperial.ac.uk
Description	P11289
Date	Thu Jan 5 11:32:32 GMT 2012
Unique Job ID	87303bd1c7a2df11

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2jrmA_	Alignment		68.1	15	**PDB header:**structural genomics, unknown function **Chain:** A: **PDB Molecule:**ribosome modulation factor; **PDBTitle:** solution nmr structure of ribosome modulation factor vp1593 from2 vibrio parahaemolyticus. northeast structural genomics target vpr55
2	c2ae3A_	Alignment		11.9	16	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**glutaryl 7-aminocephalosporanic acid acylase; **PDBTitle:** glutaryl 7-aminocephalosporanic acid acylase: mutational study of2 activation mechanism
3	c2khgA_	Alignment		7.5	38	**PDB header:**antimicrobial protein **Chain:** A: **PDB Molecule:**plnj; **PDBTitle:** plantaricin j in tfe
4	d2ijra1	Alignment		6.8	15	**Fold:**Api92-like **Superfamily:**Api92-like **Family:**Api92-like
5	c2wybA_	Alignment		6.0	16	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**acyl-homoserine lactone acylase pvdq subunit **PDBTitle:** the quorum quenching n-acyl homoserine lactone acylase pvdq2 with a covalently bound dodecanoic acid

6	d2fxta1	Alignment		5.6	23	**Fold:**Cystatin-like **Superfamily:**NTF2-like **Family:**TIM44-like