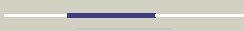
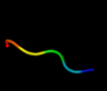
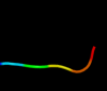
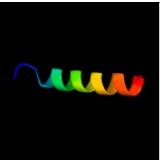


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P11866
Date	Wed Jan 25 15:20:38 GMT 2012
Unique Job ID	2f3efcea603826ea

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2rdcA_	 Alignment		12.2	36	PDB header: lipid binding protein Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a putative lipid binding protein (gsu0061) from2 geobacter sulfurreducens pca at 1.80 a resolution
2	d2ivxa1	 Alignment		9.0	18	Fold: Cyclin-like Superfamily: Cyclin-like Family: Cyclin
3	c1j73A_	 Alignment		8.2	50	PDB header: hormone/growth factor Chain: A: PDB Molecule: insulin a; PDBTitle: crystal structure of an unstable insulin analog with native activity.
4	c1j73C_	 Alignment		8.2	50	PDB header: hormone/growth factor Chain: C: PDB Molecule: insulin a; PDBTitle: crystal structure of an unstable insulin analog with native activity.
5	c3fq9A_	 Alignment		7.5	50	PDB header: hormone Chain: A: PDB Molecule: insulin; PDBTitle: design of an insulin analog with enhanced receptor-binding2 selectivity. rationale, structure, and therapeutic3 implications
6	c3fq9C_	 Alignment		7.5	50	PDB header: hormone Chain: C: PDB Molecule: insulin; PDBTitle: design of an insulin analog with enhanced receptor-binding2 selectivity. rationale, structure, and therapeutic3 implications
7	c3lemA_	 Alignment		7.2	17	PDB header: hydrolase Chain: A: PDB Molecule: fructosyltransferase; PDBTitle: crystal structure of fructosyltransferase (d191a) from a. japonicus in2 complex with nystose
8	c3kf5A_	 Alignment		6.2	67	PDB header: hydrolase Chain: A: PDB Molecule: invertase; PDBTitle: structure of invertase from schwanniomyces occidentalis
9	c3ugfB_	 Alignment		5.7	83	PDB header: transferase Chain: B: PDB Molecule: sucrose:(sucrose/fructan) 6-fructosyltransferase; PDBTitle: crystal structure of a 6-sst/6-sft from pachysandra terminalis
10	c3pijA_	 Alignment		5.4	67	PDB header: hydrolase Chain: A: PDB Molecule: beta-fructofuranosidase; PDBTitle: beta-fructofuranosidase from bifidobacterium longum - complex with2 fructose

11	c3qz3A	Alignment		5.1	45	PDB header: oxidoreductase Chain: A: PDB Molecule: ferritin; PDB Title: the crystal structure of ferritin from vibrio cholerae o1 biovar el2 tor str. n16961
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