Phyre2

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Description	Q6BF86
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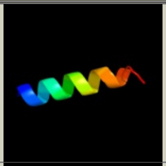
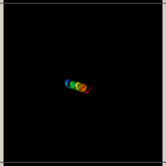
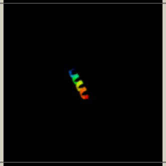
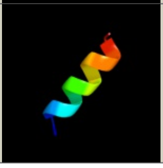
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2hg5D_	Alignment		10.7	11	**PDB header:**membrane protein **Chain:** D: **PDB Molecule:**kcsa channel; **PDBTitle:** cs+ complex of a k channel with an amide to ester substitution in the2 selectivity filter
2	d1no7a_	Alignment		9.7	35	**Fold:**Major capsid protein VP5 **Superfamily:**Major capsid protein VP5 **Family:**Major capsid protein VP5
3	c1no7A_	Alignment		9.7	35	**PDB header:**viral protein **Chain:** A: **PDB Molecule:**major capsid protein; **PDBTitle:** structure of the large protease resistant upper domain of2 vp5, the major capsid protein of herpes simplex virus-1
4	c3im4C_	Alignment		6.7	47	**PDB header:**structural protein, signaling protein **Chain:** C: **PDB Molecule:**dual specificity a kinase-anchoring protein 2; **PDBTitle:** crystal structure of camp-dependent protein kinase a2 regulatory subunit i alpha in complex with dual-specific a-3 kinase anchoring protein 2
5	c3t41B_	Alignment		5.6	36	**PDB header:**hydrolase **Chain:** B: **PDB Molecule:**epidermin leader peptide processing serine protease epip; **PDBTitle:** 1.95 angstrom resolution crystal structure of epidermin leader peptide2 processing serine protease (epip) s393a mutant from staphylococcus3 aureus

6	d1r0e_	Alignment		5.5	29	**Fold:**Subtilisin-like **Superfamily:**Subtilisin-like **Family:**Subtilases