Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2283_(-)_2555951_2556145
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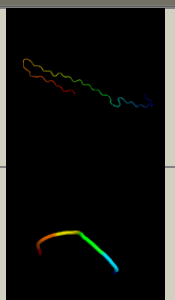
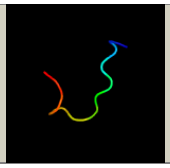
Unique Job ID	d3e8c78e44d43a12

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c6gx5A_	Alignment		23.0	23	**PDB header:**protein fibril **Chain:** A: **PDB Molecule:**microtubule-associated protein tau; **PDBTitle:** narrow pick filament from pick's disease brain
2	c2mp6A_	Alignment		18.6	100	**PDB header:**signaling protein **Chain:** A: **PDB Molecule:**suppressor of cytokine signaling 5; **PDBTitle:** structure and function of the jak interaction region in the2 intrinsically disordered n-terminus of socs5
3	c5u1dX_	Alignment		12.5	45	**PDB header:**transport protein **Chain:** X: **PDB Molecule:**tap transporter inhibitor icp47; **PDBTitle:** cryo-em structure of the human tap atp-binding cassette transporter
4	c5t5mB_	Alignment		6.1	18	**PDB header:**oxidoreductase **Chain:** B: **PDB Molecule:**tungsten formylmethanofuran dehydrogenase subunit fwdb; **PDBTitle:** tungsten-containing formylmethanofuran dehydrogenase from2 methanothermobacter wolfeii, trigonal form at 2.5 a.
5	c1x9vA_	Alignment		5.9	43	**PDB header:**viral protein **Chain:** A: **PDB Molecule:**vpr protein; **PDBTitle:** dimeric structure of the c-terminal domain of vpr
6	d1x5wa1	Alignment		5.3	50	**Fold:**beta-beta-alpha zinc fingers **Superfamily:**beta-beta-alpha zinc fingers **Family:**Classic zinc finger, C2H2

7	c2eo2A_	Alignment		5.1	54	**PDB header:**oxidoreductase **Chain:** A: **PDB Molecule:**adult male hypothalamus cdna, riken full-length **PDBTitle:** solution structure of the insertion region (510-573) of2 fthfs domain from mouse methylenetetrahydrofolate3 dehydrogenase (nadp+ dependent) 1-like protein