

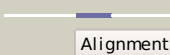
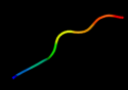
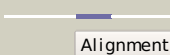





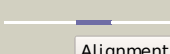

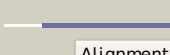

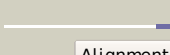

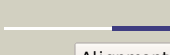







#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d2hafa1</a>	 Alignment		20.0	63	<b>Fold:</b> VC0467-like <b>Superfamily:</b> VC0467-like <b>Family:</b> VC0467-like
2	<a href="#">c2aj2A_</a>	 Alignment		17.7	63	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> hypothetical upf0301 protein vc0467; <b>PDBTitle:</b> x-ray crystal structure of protein vc0467 from vibrio2 cholerae. northeast structural genomics consortium target3 vcr8.
3	<a href="#">d2do8a1</a>	 Alignment		16.5	38	<b>Fold:</b> VC0467-like <b>Superfamily:</b> VC0467-like <b>Family:</b> VC0467-like
4	<a href="#">d2gs5a1</a>	 Alignment		15.7	25	<b>Fold:</b> VC0467-like <b>Superfamily:</b> VC0467-like <b>Family:</b> VC0467-like
5	<a href="#">d2ew0a1</a>	 Alignment		13.2	50	<b>Fold:</b> VC0467-like <b>Superfamily:</b> VC0467-like <b>Family:</b> VC0467-like
6	<a href="#">d2gzoa1</a>	 Alignment		12.7	50	<b>Fold:</b> VC0467-like <b>Superfamily:</b> VC0467-like <b>Family:</b> VC0467-like
7	<a href="#">c2q7hA_</a>	 Alignment		12.3	33	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> methylated-dna--protein-cysteine <b>PDBTitle:</b> structure of an o6-methylguanine dna methyltransferase from2 methanococcus jannaschii (mj1529)
8	<a href="#">c3izbT_</a>	 Alignment		11.9	36	<b>PDB header:</b> ribosome <b>Chain:</b> T: <b>PDB Molecule:</b> 40s ribosomal protein rps21 (s21e); <b>PDBTitle:</b> localization of the small subunit ribosomal proteins into a 6.1 a2 cryo-em map of saccharomyces cerevisiae translating 80s ribosome
9	<a href="#">d1jdga_</a>	 Alignment		9.4	25	<b>Fold:</b> lF3-like <b>Superfamily:</b> SirA-like <b>Family:</b> SirA-like
10	<a href="#">d1kn0a_</a>	 Alignment		5.6	71	<b>Fold:</b> dsRBD-like <b>Superfamily:</b> dsRNA-binding domain-like <b>Family:</b> The homologous-pairing domain of Rad52 recombinase
11	<a href="#">c3iz6T_</a>	 Alignment		5.6	45	<b>PDB header:</b> ribosome <b>Chain:</b> T: <b>PDB Molecule:</b> 40s ribosomal protein s21 (s21e); <b>PDBTitle:</b> localization of the small subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome

12	<a href="#">dlyvra1</a>	 Alignment		5.5	50	<b>Fold:</b> alpha-alpha superhelix <b>Superfamily:</b> TROVE domain-like <b>Family:</b> TROVE domain-like
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