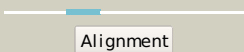

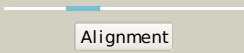

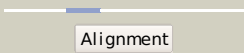

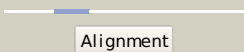

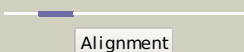

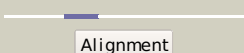



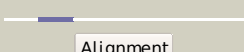

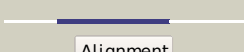



#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2k89A_</a>	 Alignment		38.7	18	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> phospholipase a-2-activating protein; <b>PDBTitle:</b> solution structure of a novel ubiquitin-binding domain from2 human plaa (pfuc, gly76-pro77 cis isomer)
2	<a href="#">c3pstA_</a>	 Alignment		30.3	33	<b>PDB header:</b> nuclear protein <b>Chain:</b> A: <b>PDB Molecule:</b> protein doa1; <b>PDBTitle:</b> crystal structure of pul and pfu(mutate) domain
3	<a href="#">c3l3fX_</a>	 Alignment		27.8	33	<b>PDB header:</b> protein binding <b>Chain:</b> X: <b>PDB Molecule:</b> protein doa1; <b>PDBTitle:</b> crystal structure of a pfu-pul domain pair of saccharomyces cerevisiae2 doa1/ufd3
4	<a href="#">d3c0na1</a>	 Alignment		25.9	32	<b>Fold:</b> C-type lectin-like <b>Superfamily:</b> C-type lectin-like <b>Family:</b> Aerolysin/Pertussis toxin (APT) domain
5	<a href="#">c2e76D_</a>	 Alignment		18.7	7	<b>PDB header:</b> photosynthesis <b>Chain:</b> D: <b>PDB Molecule:</b> cytochrome b6-f complex iron-sulfur subunit; <b>PDBTitle:</b> crystal structure of the cytochrome b6f complex with tridecyl-2 stigmatellin (tds) from m.laminosus
6	<a href="#">d1oe1a2</a>	 Alignment		14.1	19	<b>Fold:</b> Cupredoxin-like <b>Superfamily:</b> Cupredoxins <b>Family:</b> Multi domain cupredoxins
7	<a href="#">d2cs7a1</a>	 Alignment		11.4	23	<b>Fold:</b> IL8-like <b>Superfamily:</b> PhtA domain-like <b>Family:</b> PhtA domain-like
8	<a href="#">c2pq4B_</a>	 Alignment		10.8	39	<b>PDB header:</b> chaperone/oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> periplasmic nitrate reductase precursor; <b>PDBTitle:</b> nmr solution structure of napd in complex with napa1-352 signal peptide
9	<a href="#">d1pgl11</a>	 Alignment		7.0	10	<b>Fold:</b> Nucleoplasmin-like/VP (viral coat and capsid proteins) <b>Superfamily:</b> Positive stranded ssRNA viruses <b>Family:</b> Comoviridae-like VP

10	<a href="#">d1hq0a_</a>	Alignment		5.3	67	<b>Fold:</b> CNF1/YfiH-like putative cysteine hydrolases <b>Superfamily:</b> CNF1/YfiH-like putative cysteine hydrolases <b>Family:</b> Type 1 cytotoxic necrotizing factor, catalytic domain
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