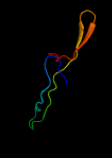
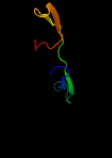


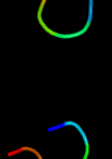

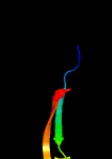

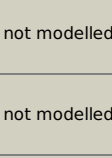


# Phyre2

Email [mdejesus@rockefeller.edu](mailto:mdejesus@rockefeller.edu)  
 Description RVBD2975A\_(RVBD2975A)\_3331864\_3332058  
 Date Thu Aug 8 16:20:14 BST 2019  
 Unique Job ID 2ca83813020035ac

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c5o60Y_</a>	Alignment		100.0	86	<b>PDB header:</b> ribosome <b>Chain:</b> Y; <b>PDB Molecule:</b> 50s ribosomal protein l28; <b>PDBTitle:</b> structure of the 50s large ribosomal subunit from mycobacterium2 smegmatis
2	<a href="#">d2jz6a1</a>	Alignment		100.0	46	<b>Fold:</b> L28p-like <b>Superfamily:</b> L28p-like <b>Family:</b> Ribosomal protein L28
3	<a href="#">c6dzpy_</a>	Alignment		99.9	53	<b>PDB header:</b> ribosome <b>Chain:</b> Y; <b>PDB Molecule:</b> <b>PDBTitle:</b> cryo-em structure of mycobacterium smegmatis c(minus) 50s ribosomal2 subunit
4	<a href="#">c6ddgj_</a>	Alignment		99.9	32	<b>PDB header:</b> ribosome/antibiotic <b>Chain:</b> J; <b>PDB Molecule:</b> 50s ribosomal protein l28; <b>PDBTitle:</b> structure of the 50s ribosomal subunit from methicillin resistant2 staphylococcus aureus in complex with the oxazolidinone antibiotic3 lzd-6
5	<a href="#">c3bboY_</a>	Alignment		99.9	26	<b>PDB header:</b> ribosome <b>Chain:</b> Y; <b>PDB Molecule:</b> ribosomal protein l28; <b>PDBTitle:</b> homology model for the spinach chloroplast 50s subunit fitted to 9.4a2 cryo-em map of the 70s chlororibosome
6	<a href="#">d2qamz1</a>	Alignment		99.9	25	<b>Fold:</b> L28p-like <b>Superfamily:</b> L28p-like <b>Family:</b> Ribosomal protein L28
7	<a href="#">c2j 81_</a>	Alignment		99.8	30	<b>PDB header:</b> ribosome <b>Chain:</b> 1; <b>PDB Molecule:</b> 50s ribosomal protein l28; <b>PDBTitle:</b> insights into translational termination from the structure2 of rf2 bound to the ribosome (part 4 of 4).3 this file contains the 50s subunit.
8	<a href="#">c1vw4S_</a>	Alignment		99.7	19	<b>PDB header:</b> ribosome <b>Chain:</b> S; <b>PDB Molecule:</b> 54s ribosomal protein l24, mitochondrial; <b>PDBTitle:</b> structure of the yeast mitochondrial large ribosomal subunit
9	<a href="#">c4wfaU_</a>	Alignment		99.5	32	<b>PDB header:</b> ribosome <b>Chain:</b> U; <b>PDB Molecule:</b> 50s ribosomal protein l28; <b>PDBTitle:</b> the crystal structure of the large ribosomal subunit of staphylococcus2 aureus in complex with linezolid
10	<a href="#">c4v191_</a>	Alignment		99.3	12	<b>PDB header:</b> ribosome <b>Chain:</b> 1; <b>PDB Molecule:</b> mitoribosomal protein bl28m, mrpl28; <b>PDBTitle:</b> structure of the large subunit of the mammalian mitoribosome, part 12 of 2
11	<a href="#">d2zjru1</a>	Alignment		99.2	16	<b>Fold:</b> L28p-like <b>Superfamily:</b> L28p-like <b>Family:</b> Ribosomal protein L28

12	<a href="#">d2j0111</a>	Alignment		98.0	29	<b>Fold:</b> L28p-like <b>Superfamily:</b> L28p-like <b>Family:</b> Ribosomal protein L28
13	<a href="#">c4ce41_</a>	Alignment		92.6	15	<b>PDB header:</b> ribosome <b>Chain:</b> 1: <b>PDB Molecule:</b> mrpl28; <b>PDBTitle:</b> 39s large subunit of the porcine mitochondrial ribosome
14	<a href="#">d2cu8a1</a>	Alignment		57.4	42	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LIM domain
15	<a href="#">c2k5cA_</a>	Alignment		33.0	75	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein pf0385; <b>PDBTitle:</b> nmr structure for pf0385
16	<a href="#">c2elrA_</a>	Alignment		26.5	71	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein 406; <b>PDBTitle:</b> solution structure of the 15th c2h2 zinc finger of human2 zinc finger protein 406
17	<a href="#">d2yt9a2</a>	Alignment		25.2	83	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
18	<a href="#">c4n0oC_</a>	Alignment		19.4	50	<b>PDB header:</b> hydrolase/dna <b>Chain:</b> C: <b>PDB Molecule:</b> replicase polyprotein 1ab; <b>PDBTitle:</b> complex structure of arterivirus nonstructural protein 10 (helicase)2 with dna
19	<a href="#">c2uz0B_</a>	Alignment		16.8	18	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> tributyryn esterase; <b>PDBTitle:</b> the crystal structure of the esta protein, a2 virulence factor esta protein from streptococcus pneumonia
20	<a href="#">d2dmda2</a>	Alignment		13.6	57	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
21	<a href="#">c5zwnY_</a>	Alignment	not modelled	12.1	57	<b>PDB header:</b> splicing <b>Chain:</b> Y: <b>PDB Molecule:</b> protein luc7; <b>PDBTitle:</b> cryo-em structure of the yeast pre-b complex at an average resolution2 of 3.3 angstrom (part ii: u1 snrnp region)
22	<a href="#">c3iufA_</a>	Alignment	not modelled	11.4	86	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein ubi-d4; <b>PDBTitle:</b> crystal structure of the c2h2-type zinc finger domain of human ubi-d4
23	<a href="#">d1dxga_</a>	Alignment	not modelled	11.4	50	<b>Fold:</b> Rubredoxin-like <b>Superfamily:</b> Rubredoxin-like <b>Family:</b> Desulfiredoxin
24	<a href="#">c2n94A_</a>	Alignment	not modelled	10.2	42	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> box c/d snorna protein 1; <b>PDBTitle:</b> nmr structure of yeast bcd1 protein zinc finger
25	<a href="#">c3mhsE_</a>	Alignment	not modelled	9.8	29	<b>PDB header:</b> hydrolase/transcription regulator/protei <b>Chain:</b> E: <b>PDB Molecule:</b> saga-associated factor 73; <b>PDBTitle:</b> structure of the saga ubp8/sgf11/sus1/sgf73 dub module bound to2 ubiquitin aldehyde
26	<a href="#">d1lbia2</a>	Alignment	not modelled	9.5	31	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LIM domain
27	<a href="#">c2c4iA_</a>	Alignment	not modelled	9.2	44	<b>PDB header:</b> glycoprotein <b>Chain:</b> A: <b>PDB Molecule:</b> avidin; <b>PDBTitle:</b> crystal structure of engineered avidin
28	<a href="#">d1llmc2</a>	Alignment	not modelled	9.1	63	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
						<b>PDB header:</b> metal binding protein

29	<a href="#">c2jtgA</a>	Alignment	not modelled	9.1	30	<b>Chain:</b> A: <b>PDB Molecule:</b> thap domain-containing protein 1; <b>PDBTitle:</b> solution structure of the thap-zinc finger of thap1
30	<a href="#">d2cota1</a>	Alignment	not modelled	9.0	71	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
31	<a href="#">c1ardA</a>	Alignment	not modelled	8.9	38	<b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> yeast transcription factor adr1; <b>PDBTitle:</b> structures of dna-binding mutant zinc finger domains: implications for2 dna binding
32	<a href="#">d1u85a1</a>	Alignment	not modelled	8.9	29	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
33	<a href="#">d2adra1</a>	Alignment	not modelled	8.8	38	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
34	<a href="#">d1kpta</a>	Alignment	not modelled	8.8	47	<b>Fold:</b> Yeast killer toxins <b>Superfamily:</b> Yeast killer toxins <b>Family:</b> Virally encoded KP4 toxin
35	<a href="#">d2cota2</a>	Alignment	not modelled	8.6	63	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
36	<a href="#">c1meyG</a>	Alignment	not modelled	8.4	57	<b>PDB header:</b> transferase/dna <b>Chain:</b> G: <b>PDB Molecule:</b> consensus zinc finger; <b>PDBTitle:</b> crystal structure of a designed zinc finger protein bound2 to dna
37	<a href="#">c3hi2C</a>	Alignment	not modelled	8.4	38	<b>PDB header:</b> dna binding protein/toxin <b>Chain:</b> C: <b>PDB Molecule:</b> hth-type transcriptional regulator mqa2(ygit); <b>PDBTitle:</b> structure of the n-terminal domain of the e. coli antitoxin mqa2 (ygit/b3021) in complex with the e. coli toxin mqsr (ygiu/b3022)
38	<a href="#">c1va3A</a>	Alignment	not modelled	8.3	38	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcription factor sp1; <b>PDBTitle:</b> solution structure of transcription factor sp1 dna binding2 domain (zinc finger 3)
39	<a href="#">d1sp1a</a>	Alignment	not modelled	8.3	38	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
40	<a href="#">c1sp1A</a>	Alignment	not modelled	8.3	38	<b>PDB header:</b> zinc finger <b>Chain:</b> A: <b>PDB Molecule:</b> sp1f3; <b>PDBTitle:</b> nmr structure of a zinc finger domain from transcription2 factor sp1f3, minimized average structure
41	<a href="#">d1x6ea2</a>	Alignment	not modelled	8.2	57	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
42	<a href="#">d2b8ta2</a>	Alignment	not modelled	8.0	31	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> Type II thymidine kinase zinc finger
43	<a href="#">c1xf7A</a>	Alignment	not modelled	8.0	29	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> wilms' tumor protein; <b>PDBTitle:</b> high resolution nmr structure of the wilms' tumor2 suppressor protein (wt1) finger 3
44	<a href="#">d1xf7a</a>	Alignment	not modelled	8.0	29	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
45	<a href="#">c1z2qA</a>	Alignment	not modelled	8.0	21	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> lm5-1; <b>PDBTitle:</b> high-resolution solution structure of the lm5-1 fyve domain2 from leishmania major
46	<a href="#">c3jyvN</a>	Alignment	not modelled	8.0	20	<b>PDB header:</b> ribosome <b>Chain:</b> N: <b>PDB Molecule:</b> 40s ribosomal protein s29(a); <b>PDBTitle:</b> structure of the 40s rna and proteins and p/e trna for eukaryotic2 ribosome based on cryo-em map of thermomyces lanuginosus ribosome at3 8.9a resolution
47	<a href="#">d1srka</a>	Alignment	not modelled	8.0	57	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
48	<a href="#">c1areA</a>	Alignment	not modelled	7.9	38	<b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> yeast transcription factor adr1; <b>PDBTitle:</b> structures of dna-binding mutant zinc finger domains: implications for2 dna binding
49	<a href="#">c3uk3D</a>	Alignment	not modelled	7.8	44	<b>PDB header:</b> dna/metal binding protein <b>Chain:</b> D: <b>PDB Molecule:</b> zinc finger protein 217; <b>PDBTitle:</b> crystal structure of znf217 bound to dna
50	<a href="#">c1rikA</a>	Alignment	not modelled	7.7	43	<b>PDB header:</b> de novo protein <b>Chain:</b> A: <b>PDB Molecule:</b> e6apc1 peptide; <b>PDBTitle:</b> e6-binding zinc finger (e6apc1)
51	<a href="#">c5xxud</a>	Alignment	not modelled	7.7	20	<b>PDB header:</b> ribosome <b>Chain:</b> D: <b>PDB Molecule:</b> ribosomal protein us3; <b>PDBTitle:</b> small subunit of toxoplasma gondii ribosome
52	<a href="#">d1a1ia2</a>	Alignment	not modelled	7.6	43	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
53	<a href="#">c3j20P</a>	Alignment	not modelled	7.5	40	<b>PDB header:</b> ribosome <b>Chain:</b> P: <b>PDB Molecule:</b> 30s ribosomal protein s14p type z; <b>PDBTitle:</b> promiscuous behavior of proteins in archaeal ribosomes revealed by2 cryo-em: implications for evolution of eukaryotic ribosomes (30s3 ribosomal subunit)
54	<a href="#">c2xznN</a>	Alignment	not modelled	7.5	30	<b>PDB header:</b> ribosome <b>Chain:</b> N: <b>PDB Molecule:</b> rps29e; <b>PDBTitle:</b> crystal structure of the eukaryotic 40s ribosomal2 subunit in complex with initiation factor 1. this file3 contains the 40s subunit and initiation factor for4 molecule 2

55	<a href="#">d1a1a3</a>	Alignment	not modelled	7.5	63	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
56	<a href="#">c6az1S_</a>	Alignment	not modelled	7.5	40	<b>PDB header:</b> ribosome/antibiotic <b>Chain:</b> S: <b>PDB Molecule:</b> ribosomal protein s14; <b>PDBTitle:</b> cryo-em structure of the small subunit of leishmania ribosome bound to2 paromomycin
57	<a href="#">d2epra1</a>	Alignment	not modelled	7.4	71	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
58	<a href="#">c5z58w_</a>	Alignment	not modelled	7.3	42	<b>PDB header:</b> splicing <b>Chain:</b> W: <b>PDB Molecule:</b> <b>PDBTitle:</b> cryo-em structure of a human activated spliceosome (early bact) at 4.92 angstrom.
59	<a href="#">c5o60d_</a>	Alignment	not modelled	7.2	33	<b>PDB header:</b> ribosome <b>Chain:</b> D: <b>PDB Molecule:</b> 50s ribosomal protein l3; <b>PDBTitle:</b> structure of the 50s large ribosomal subunit from mycobacterium2 smegmatis
60	<a href="#">c2lvrA_</a>	Alignment	not modelled	7.2	43	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger and btb domain-containing protein 17; <b>PDBTitle:</b> solution structure of miz-1 zinc finger 8
61	<a href="#">c2l8eA_</a>	Alignment	not modelled	7.2	57	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> polyhomeotic-like protein 1; <b>PDBTitle:</b> solution nmr structure of fcs domain of human polyhomeotic homolog 12 (hph1)
62	<a href="#">c1arfA_</a>	Alignment	not modelled	7.1	43	<b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> yeast transcription factor adr1; <b>PDBTitle:</b> structures of dna-binding mutant zinc finger domains: implications for2 dna binding
63	<a href="#">c5yxiA_</a>	Alignment	not modelled	7.0	33	<b>PDB header:</b> de novo protein <b>Chain:</b> A: <b>PDB Molecule:</b> drafx6; <b>PDBTitle:</b> designed protein drafx6
64	<a href="#">d2d8ya1</a>	Alignment	not modelled	6.9	25	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LIM domain
65	<a href="#">c3zey8_</a>	Alignment	not modelled	6.8	30	<b>PDB header:</b> ribosome <b>Chain:</b> 8: <b>PDB Molecule:</b> ribosomal protein s29, putative; <b>PDBTitle:</b> high-resolution cryo-electron microscopy structure of the trypanosoma2 brucei ribosome
66	<a href="#">c2ko5A_</a>	Alignment	not modelled	6.8	33	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> ring finger protein z; <b>PDBTitle:</b> nmr solution structure of lfv-z
67	<a href="#">d2csha2</a>	Alignment	not modelled	6.8	40	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
68	<a href="#">c1dvpA_</a>	Alignment	not modelled	6.8	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> hepatocyte growth factor-regulated tyrosine <b>PDBTitle:</b> crystal structure of the vhs and fyve tandem domains of hrs,2 a protein involved in membrane trafficking and signal3 transduction
69	<a href="#">c5uxmA_</a>	Alignment	not modelled	6.7	24	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> phospho-2-dehydro-3-deoxyheptonate aldolase; <b>PDBTitle:</b> type ii dah7ps from pseudomonas aeruginosa with trp bound
70	<a href="#">c2ytbA_</a>	Alignment	not modelled	6.7	63	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein 32; <b>PDBTitle:</b> solution structure of c2h2 type zinc finger domain 5 in2 zinc finger protein 32
71	<a href="#">c2m0fA_</a>	Alignment	not modelled	6.6	57	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger and btb domain-containing protein 17; <b>PDBTitle:</b> solution structure of miz-1 zinc finger 7
72	<a href="#">c2zqkn_</a>	Alignment	not modelled	6.6	25	<b>PDB header:</b> ribosomal protein/rna <b>Chain:</b> N: <b>PDB Molecule:</b> <b>PDBTitle:</b> structure of a mammalian ribosomal 40s subunit within an 80s complex2 obtained by docking homology models of the rna and proteins into an3 8.7 a cryo-em map
73	<a href="#">c5hudA_</a>	Alignment	not modelled	6.6	24	<b>PDB header:</b> transferase/isomerase <b>Chain:</b> A: <b>PDB Molecule:</b> 3-deoxy-d-arabino-heptulosonate 7-phosphate (dahp) <b>PDBTitle:</b> non-covalent complex of and dahp synthase and chorismate mutase from2 corynebacterium glutamicum with bound transition state analog
74	<a href="#">c6g90T_</a>	Alignment	not modelled	6.6	33	<b>PDB header:</b> splicing <b>Chain:</b> T: <b>PDB Molecule:</b> pre-mrna-splicing factor prp9; <b>PDBTitle:</b> prespliceosome structure provides insight into spliceosome assembly2 and regulation (map a2)
75	<a href="#">c2lvtA_</a>	Alignment	not modelled	6.6	57	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger and btb domain-containing protein 17; <b>PDBTitle:</b> solution structure of miz-1 zinc finger 9
76	<a href="#">c1wbiD_</a>	Alignment	not modelled	6.5	38	<b>PDB header:</b> avidin-related protein <b>Chain:</b> D: <b>PDB Molecule:</b> avidin-related protein 2; <b>PDBTitle:</b> avr2
77	<a href="#">c6bmcA_</a>	Alignment	not modelled	6.5	18	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> phospho-2-dehydro-3-deoxyheptonate aldolase; <b>PDBTitle:</b> the structure of a dimeric type ii dah7ps associated with pyocyanin2 biosynthesis in pseudomonas aeruginosa
78	<a href="#">d1zfoa_</a>	Alignment	not modelled	6.5	42	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LASP-1
79	<a href="#">c2lvuA_</a>	Alignment	not modelled	6.5	63	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger and btb domain-containing protein 17;

						<b>PDBTitle:</b> solution structure of miz-1 zinc finger 10
80	<a href="#">d2dipa1</a>	Alignment	not modelled	6.4	42	<b>Fold:</b> RING/U-box <b>Superfamily:</b> RING/U-box <b>Family:</b> ZZ domain
81	<a href="#">c2yu5A</a>	Alignment	not modelled	6.3	29	<b>PDB header:</b> rna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein 473; <b>PDBTitle:</b> solution structure of the zf-c2h2 domain (669-699aa) in2 zinc finger protein 473
82	<a href="#">d2b7oa1</a>	Alignment	not modelled	6.2	24	<b>Fold:</b> TIM beta/alpha-barrel <b>Superfamily:</b> Aldolase <b>Family:</b> Class-II DAHP synthetase
83	<a href="#">d1wfka_</a>	Alignment	not modelled	6.2	20	<b>Fold:</b> FYVE/PHD zinc finger <b>Superfamily:</b> FYVE/PHD zinc finger <b>Family:</b> FYVE, a phosphatidylinositol-3-phosphate binding domain
84	<a href="#">c2qm0B_</a>	Alignment	not modelled	6.1	9	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> bes; <b>PDBTitle:</b> crystal structure of bes protein from bacillus cereus
85	<a href="#">c5xyid</a>	Alignment	not modelled	6.0	50	<b>PDB header:</b> ribosome <b>Chain:</b> D: <b>PDB Molecule:</b> ribosomal protein s3, putative; <b>PDBTitle:</b> small subunit of trichomonas vaginalis ribosome
86	<a href="#">d2yt9a1</a>	Alignment	not modelled	6.0	63	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
87	<a href="#">d1imla1</a>	Alignment	not modelled	5.8	31	<b>Fold:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Superfamily:</b> Glucocorticoid receptor-like (DNA-binding domain) <b>Family:</b> LIM domain
88	<a href="#">d2ct5a1</a>	Alignment	not modelled	5.7	25	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> BED zinc finger
89	<a href="#">c3j3v2_</a>	Alignment	not modelled	5.7	43	<b>PDB header:</b> ribosome <b>Chain:</b> 2: <b>PDB Molecule:</b> 50s ribosomal protein l34; <b>PDBTitle:</b> atomic model of the immature 50s subunit from bacillus subtilis (state2 i-a)
90	<a href="#">d2ct1a2</a>	Alignment	not modelled	5.6	43	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
91	<a href="#">c2djrA</a>	Alignment	not modelled	5.6	33	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger bed domain-containing protein 2; <b>PDBTitle:</b> solution structures of the c2h2 type zinc finger domain of2 human zinc finger bed domain containing protein 2
92	<a href="#">c2naxA</a>	Alignment	not modelled	5.6	56	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> protein pcf11; <b>PDBTitle:</b> structure of cchc zinc finger domain of pcf11
93	<a href="#">d1jb0d_</a>	Alignment	not modelled	5.6	33	<b>Fold:</b> Photosystem I subunit Psd <b>Superfamily:</b> Photosystem I subunit Psd <b>Family:</b> Photosystem I subunit Psd
94	<a href="#">d1x6ea1</a>	Alignment	not modelled	5.6	57	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
95	<a href="#">c2eltA</a>	Alignment	not modelled	5.5	29	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein 406; <b>PDBTitle:</b> solution structure of the 3rd c2h2 zinc finger of human2 zinc finger protein 406
96	<a href="#">d1p7aa</a>	Alignment	not modelled	5.5	25	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
97	<a href="#">c4qiwp_</a>	Alignment	not modelled	5.5	27	<b>PDB header:</b> transcription <b>Chain:</b> P: <b>PDB Molecule:</b> dna-directed rna polymerase subunit p; <b>PDBTitle:</b> crystal structure of euryarchaeal rna polymerase from thermococcus2 kodakarensis
98	<a href="#">d2epsa1</a>	Alignment	not modelled	5.4	63	<b>Fold:</b> beta-beta-alpha zinc fingers <b>Superfamily:</b> beta-beta-alpha zinc fingers <b>Family:</b> Classic zinc finger, C2H2
99	<a href="#">c2ytaA</a>	Alignment	not modelled	5.4	50	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger protein 32; <b>PDBTitle:</b> solution structure of c2h2 type zinc finger domain 3 in2 zinc finger protein 32