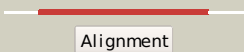
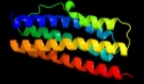
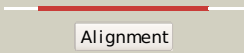
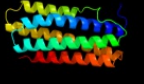
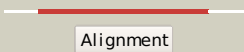
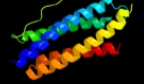


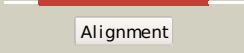
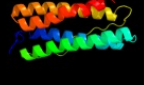




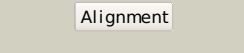
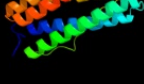


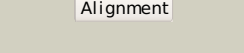
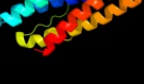
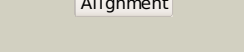

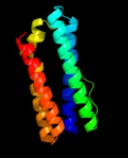







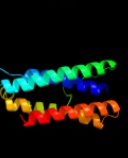


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d2gs4a1</a>	 Alignment		100.0	98	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> YciF-like
2	<a href="#">d2gyqa1</a>	 Alignment		100.0	46	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> YciF-like
3	<a href="#">c3hiuB_</a>	 Alignment		100.0	23	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> uncharacterized protein; <b>PDBTitle:</b> the crystal structure of protein (xcc3681) from xanthomonas2 campestris pv. campestris str. atcc 33913
4	<a href="#">c3oghB_</a>	 Alignment		99.9	22	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> protein ycie; <b>PDBTitle:</b> crystal structure of ycie protein from e. coli cft073, a member of2 ferritin-like superfamily of diiron-containing four-helix-bundle3 proteins
5	<a href="#">c3r2rA_</a>	 Alignment		98.7	18	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> bacterioferritin; <b>PDBTitle:</b> 1.65a resolution structure of iron soaked ftna from pseudomonas2 aeruginosa (ph 6.0)
6	<a href="#">c2qqyA_</a>	 Alignment		98.6	16	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> sigma b operon; <b>PDBTitle:</b> crystal structure of ferritin like, diiron-carboxylate proteins from2 bacillus anthracis str. ames
7	<a href="#">d2htna1</a>	 Alignment		98.6	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
8	<a href="#">c3bknB_</a>	 Alignment		98.6	15	<b>PDB header:</b> metal binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> bacterioferritin; <b>PDBTitle:</b> the structure of mycobacterial bacterioferritin
9	<a href="#">c2vzbA_</a>	 Alignment		98.6	17	<b>PDB header:</b> metal transport <b>Chain:</b> A: <b>PDB Molecule:</b> putative bacterioferritin-related protein; <b>PDBTitle:</b> a dodecameric thioferritin in the bacterial domain, characterization2 of the bacterioferritin-related protein from bacteroides fragilis
10	<a href="#">c3gvvC_</a>	 Alignment		98.5	13	<b>PDB header:</b> metal binding protein <b>Chain:</b> C: <b>PDB Molecule:</b> bacterioferritin; <b>PDBTitle:</b> crystal structure of bacterioferritin from r.sphaeroides
11	<a href="#">d2fkza1</a>	 Alignment		98.5	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin

12	<a href="#">dlnf4a_</a>	Alignment		98.5	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
13	<a href="#">c3qhbA_</a>	Alignment		98.4	16	<b>PDB header:</b> metal transport <b>Chain:</b> A: <b>PDB Molecule:</b> symerythrin; <b>PDBTitle:</b> crystal structure of oxidized symerythrin from cyanophora paradoxa
14	<a href="#">c3fvbB_</a>	Alignment		98.4	16	<b>PDB header:</b> metal binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> bacterioferritin; <b>PDBTitle:</b> crystal structure of ferritin (bacterioferritin) from2 brucella melitensis
15	<a href="#">dlyv1a1</a>	Alignment		98.3	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
16	<a href="#">d1lkoa1</a>	Alignment		98.2	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
17	<a href="#">c2clbP_</a>	Alignment		98.2	12	<b>PDB header:</b> metal binding protein <b>Chain:</b> P: <b>PDB Molecule:</b> dps-like protein; <b>PDBTitle:</b> the structure of the dps-like protein from sulfobolus2 solfataricus reveals a bacterioferritin-like di-metal3 binding site within a dps-like dodecameric assembly
18	<a href="#">d1jgca_</a>	Alignment		98.2	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
19	<a href="#">d1j30a_</a>	Alignment		98.2	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
20	<a href="#">d2oh3a1</a>	Alignment		97.9	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> AMB4284-like
21	<a href="#">d2fzfa1</a>	Alignment	not modelled	97.9	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
22	<a href="#">c1dvbA_</a>	Alignment	not modelled	97.9	17	<b>PDB header:</b> electron transport <b>Chain:</b> A: <b>PDB Molecule:</b> rubrerythrin; <b>PDBTitle:</b> rubrerythrin
23	<a href="#">d1yuza1</a>	Alignment	not modelled	97.8	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
24	<a href="#">d1nnqa1</a>	Alignment	not modelled	97.7	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
25	<a href="#">c2c41K_</a>	Alignment	not modelled	97.5	14	<b>PDB header:</b> iron-binding/oxidation protein <b>Chain:</b> K: <b>PDB Molecule:</b> dps family dna-binding stress response protein; <b>PDBTitle:</b> x-ray structure of dps from thermosynechococcus elongatus
26	<a href="#">c1yuzB_</a>	Alignment	not modelled	97.5	17	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> nigerythrin; <b>PDBTitle:</b> partially reduced state of nigerythrin
27	<a href="#">c3e6sD_</a>	Alignment	not modelled	97.4	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> D: <b>PDB Molecule:</b> ferritin; <b>PDBTitle:</b> crystal structure of ferritin soaked with iron from pseudo-2 nitzschia multiseris
28	<a href="#">c2hr5B_</a>	Alignment	not modelled	97.2	13	<b>PDB header:</b> metal binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> rubrerythrin; <b>PDBTitle:</b> pf1283- rubrerythrin from pyrococcus furiosus iron bound form

29	<a href="#">d1o9ra_</a>	<div>Alignment</div>	not modelled	97.0	6	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
30	<a href="#">c2chpC_</a>	<div>Alignment</div>	not modelled	97.0	13	<b>PDB header:</b> dna-binding protein <b>Chain:</b> C: <b>PDB Molecule:</b> metalloregulation dna-binding stress protein; <b>PDBTitle:</b> crystal structure of the dodecameric ferritin mrga from2 b.subtilis 168
31	<a href="#">d1jiga_</a>	<div>Alignment</div>	not modelled	97.0	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
32	<a href="#">c3iq1A_</a>	<div>Alignment</div>	not modelled	97.0	11	<b>PDB header:</b> metal transport <b>Chain:</b> A: <b>PDB Molecule:</b> dps family protein; <b>PDBTitle:</b> crystal structure of dps protein from vibrio cholerae o1, a member of2 a broad superfamily of ferritin-like diiron-carboxylate proteins
33	<a href="#">c2d5kC_</a>	<div>Alignment</div>	not modelled	96.9	7	<b>PDB header:</b> metal binding protein <b>Chain:</b> C: <b>PDB Molecule:</b> dps family protein; <b>PDBTitle:</b> crystal structure of dps from staphylococcus aureus
34	<a href="#">d1ji5a_</a>	<div>Alignment</div>	not modelled	96.9	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
35	<a href="#">d1n1ga_</a>	<div>Alignment</div>	not modelled	96.9	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
36	<a href="#">c3bvkC_</a>	<div>Alignment</div>	not modelled	96.9	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> C: <b>PDB Molecule:</b> ferritin; <b>PDBTitle:</b> structural basis for the iron uptake mechanism of helicobacter pylori2 ferritin
37	<a href="#">c3qz3A_</a>	<div>Alignment</div>	not modelled	96.8	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ferritin; <b>PDBTitle:</b> the crystal structure of ferritin from vibrio cholerae o1 biovar el2 tor str. n16961
38	<a href="#">d1vlga_</a>	<div>Alignment</div>	not modelled	96.8	17	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
39	<a href="#">d1vjxa_</a>	<div>Alignment</div>	not modelled	96.8	18	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
40	<a href="#">c2vxxB_</a>	<div>Alignment</div>	not modelled	96.8	14	<b>PDB header:</b> dna-binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> starvation induced dna binding protein; <b>PDBTitle:</b> x-ray structure of dpsa from thermosynechococcus elongatus
41	<a href="#">d1euma_</a>	<div>Alignment</div>	not modelled	96.7	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
42	<a href="#">c3kwoA_</a>	<div>Alignment</div>	not modelled	96.7	14	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> putative bacterioferritin; <b>PDBTitle:</b> crystal structure of putative bacterioferritin from2 campylobacter jejuni
43	<a href="#">d1krqa_</a>	<div>Alignment</div>	not modelled	96.6	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
44	<a href="#">c3fseB_</a>	<div>Alignment</div>	not modelled	96.5	15	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> two-domain protein containing dj-1/thij/pfpi-like and <b>PDBTitle:</b> crystal structure of a two-domain protein containing dj-1/thij/pfpi-2 like and ferritin-like domains (ava_4496) from anabaena variabilis3 atcc 29413 at 1.90 a resolution
45	<a href="#">c1u7mB_</a>	<div>Alignment</div>	not modelled	96.5	18	<b>PDB header:</b> de novo protein <b>Chain:</b> B: <b>PDB Molecule:</b> four-helix bundle model; <b>PDBTitle:</b> solution structure of a diiron protein model: due ferri(ii)2 turn mutant
46	<a href="#">d1s3qa1</a>	<div>Alignment</div>	not modelled	96.4	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
47	<a href="#">c2c6rA_</a>	<div>Alignment</div>	not modelled	96.3	13	<b>PDB header:</b> dna-binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> dna-binding stress response protein, dps family; <b>PDBTitle:</b> fe-soaked crystal structure of the dps92 from deinococcus2 radiodurans
48	<a href="#">d1tjoa_</a>	<div>Alignment</div>	not modelled	96.3	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
49	<a href="#">c2xgwA_</a>	<div>Alignment</div>	not modelled	96.0	10	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> peroxide resistance protein; <b>PDBTitle:</b> zinc-bound crystal structure of streptococcus pyogenes dpr
50	<a href="#">d1vela_</a>	<div>Alignment</div>	not modelled	95.6	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
51	<a href="#">d2ib0a1</a>	<div>Alignment</div>	not modelled	95.6	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Rv2844-like
52	<a href="#">c2ib0A_</a>	<div>Alignment</div>	not modelled	95.6	11	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> conserved hypothetical alanine rich protein; <b>PDBTitle:</b> crystal structure of a conserved hypothetical protein, rv2844, from2 mycobacterium tuberculosis
53	<a href="#">c2jd8C_</a>	<div>Alignment</div>	not modelled	95.5	19	<b>PDB header:</b> metal transport <b>Chain:</b> C: <b>PDB Molecule:</b> ferritin homolog; <b>PDBTitle:</b> crystal structure of the zn-soaked ferritin from the2 hyperthermophilic archaeal anaerobe pyrococcus furiosus
		<div></div>				<b>Fold:</b> Ferritin-like

54	<a href="#">d2bk6a1</a>	<div><div>Alignment</div></div>	not modelled	95.4	11	<b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
55	<a href="#">d1ji4a_</a>	<div><div>Alignment</div></div>	not modelled	95.3	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
56	<a href="#">c3q4nA_</a>	<div><div>Alignment</div></div>	not modelled	95.3	12	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein mj0754; <b>PDBTitle:</b> crystal structure of hypothetical protein mj0754 from methanococcus2 jannaschii dsm 2661
57	<a href="#">c3qd8M_</a>	<div><div>Alignment</div></div>	not modelled	95.3	14	<b>PDB header:</b> metal binding protein <b>Chain:</b> M: <b>PDB Molecule:</b> probable bacterioferritin bfrb; <b>PDBTitle:</b> crystal structure of mycobacterium tuberculosis bfrb
58	<a href="#">c2rbdB_</a>	<div><div>Alignment</div></div>	not modelled	95.0	9	<b>PDB header:</b> structural protein <b>Chain:</b> B: <b>PDB Molecule:</b> bh2358 protein; <b>PDBTitle:</b> crystal structure of a putative spore coat protein (bh2358) from2 bacillus halodurans c-125 at 1.54 a resolution
59	<a href="#">d2fjca1</a>	<div><div>Alignment</div></div>	not modelled	95.0	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
60	<a href="#">c2wlaA_</a>	<div><div>Alignment</div></div>	not modelled	94.4	10	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> dps-like peroxide resistance protein; <b>PDBTitle:</b> streptococcus pyogenes dpr
61	<a href="#">d1dpsa_</a>	<div><div>Alignment</div></div>	not modelled	94.0	7	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
62	<a href="#">c2f7nA_</a>	<div><div>Alignment</div></div>	not modelled	93.3	9	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> dna-binding stress response protein, dps family; <b>PDBTitle:</b> structure of d. radiodurans dps-1
63	<a href="#">d2yw6a1</a>	<div><div>Alignment</div></div>	not modelled	92.8	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
64	<a href="#">d1r03a_</a>	<div><div>Alignment</div></div>	not modelled	92.1	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
65	<a href="#">d1rcda_</a>	<div><div>Alignment</div></div>	not modelled	91.9	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
66	<a href="#">d2ceia1</a>	<div><div>Alignment</div></div>	not modelled	91.5	12	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
67	<a href="#">c1lt1G_</a>	<div><div>Alignment</div></div>	not modelled	91.2	15	<b>PDB header:</b> de novo protein <b>Chain:</b> G: <b>PDB Molecule:</b> l3g-df1; <b>PDBTitle:</b> sliding helix induced change of coordination geometry in a2 model di-mn(ii) protein
68	<a href="#">d1lb3a_</a>	<div><div>Alignment</div></div>	not modelled	90.9	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
69	<a href="#">c2pybC_</a>	<div><div>Alignment</div></div>	not modelled	89.2	8	<b>PDB header:</b> metal transport <b>Chain:</b> C: <b>PDB Molecule:</b> neutrophil activating protein; <b>PDBTitle:</b> napa protein from borrelia burgdorferi
70	<a href="#">d1mfra_</a>	<div><div>Alignment</div></div>	not modelled	89.1	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
71	<a href="#">c2hz8A_</a>	<div><div>Alignment</div></div>	not modelled	88.9	24	<b>PDB header:</b> de novo protein <b>Chain:</b> A: <b>PDB Molecule:</b> de novo designed diiron protein; <b>PDBTitle:</b> qm/mm structure refined from nmr-structure of a single2 chain diiron protein
72	<a href="#">d1jkva_</a>	<div><div>Alignment</div></div>	not modelled	88.5	9	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Manganese catalase (T-catalase)
73	<a href="#">d1umna_</a>	<div><div>Alignment</div></div>	not modelled	86.4	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
74	<a href="#">c2z90D_</a>	<div><div>Alignment</div></div>	not modelled	82.6	8	<b>PDB header:</b> dna binding protein <b>Chain:</b> D: <b>PDB Molecule:</b> starvation-inducible dna-binding protein or fine <b>PDBTitle:</b> crystal structure of the second dps from mycobacterium2 smegmatis
75	<a href="#">d1q06a_</a>	<div><div>Alignment</div></div>	not modelled	82.4	10	<b>Fold:</b> Putative DNA-binding domain <b>Superfamily:</b> Putative DNA-binding domain <b>Family:</b> DNA-binding N-terminal domain of transcription activators
76	<a href="#">c2yjkF_</a>	<div><div>Alignment</div></div>	not modelled	81.1	7	<b>PDB header:</b> metal-binding protein <b>Chain:</b> F: <b>PDB Molecule:</b> afp; <b>PDBTitle:</b> structure of dps from microbacterium arborescens in the2 high iron form
77	<a href="#">d2za7a1</a>	<div><div>Alignment</div></div>	not modelled	80.8	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
78	<a href="#">d1q08a_</a>	<div><div>Alignment</div></div>	not modelled	79.4	14	<b>Fold:</b> Putative DNA-binding domain <b>Superfamily:</b> Putative DNA-binding domain <b>Family:</b> DNA-binding N-terminal domain of transcription activators
79	<a href="#">d1zpya1</a>	<div><div>Alignment</div></div>	not modelled	65.7	10	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> half-ferritin
80	<a href="#">c3bt5A_</a>	<div><div>Alignment</div></div>	not modelled	59.5	12	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein duf305; <b>PDBTitle:</b> crystal structure of duf305 fragment from deinococcus2 radiodurans
		<div><div>Alignment</div></div>				<b>Fold:</b> Ferritin-like

81	<a href="#">d1zuja1</a>	Alignment	not modelled	48.4	11	<b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
82	<a href="#">d1z72a1</a>	Alignment	not modelled	47.4	9	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
83	<a href="#">d1udda_</a>	Alignment	not modelled	44.1	17	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
84	<a href="#">d1rkta2</a>	Alignment	not modelled	40.4	13	<b>Fold:</b> Tetracyclin repressor-like, C-terminal domain <b>Superfamily:</b> Tetracyclin repressor-like, C-terminal domain <b>Family:</b> Tetracyclin repressor-like, C-terminal domain
85	<a href="#">d1cnt1_</a>	Alignment	not modelled	40.3	16	<b>Fold:</b> 4-helical cytokines <b>Superfamily:</b> 4-helical cytokines <b>Family:</b> Long-chain cytokines
86	<a href="#">c1o06A_</a>	Alignment	not modelled	36.6	22	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> vacuolar protein sorting-associated protein vps27; <b>PDBTitle:</b> crystal structure of the vps27p ubiquitin interacting motif (uim)
87	<a href="#">d2cwlal</a>	Alignment	not modelled	31.5	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Manganese catalase (T-catalase)
88	<a href="#">d2g3ba2</a>	Alignment	not modelled	29.8	8	<b>Fold:</b> Tetracyclin repressor-like, C-terminal domain <b>Superfamily:</b> Tetracyclin repressor-like, C-terminal domain <b>Family:</b> Tetracyclin repressor-like, C-terminal domain
89	<a href="#">c3a9qR_</a>	Alignment	not modelled	27.6	16	<b>PDB header:</b> oxidoreductase <b>Chain:</b> R: <b>PDB Molecule:</b> <b>PDBTitle:</b> crystal structure analysis of e173a variant of the soybean2 ferritin sfer4
90	<a href="#">c2qf9B_</a>	Alignment	not modelled	27.2	11	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> putative secreted protein; <b>PDBTitle:</b> crystal structure of putative secreted protein duf305 from2 streptomyces coelicolor
91	<a href="#">c3oqlB_</a>	Alignment	not modelled	26.4	9	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> tena homolog; <b>PDBTitle:</b> crystal structure of a tena homolog (pspto1738) from pseudomonas2 syringae pv. tomato str. dc3000 at 2.54 a resolution
92	<a href="#">d1zs3a1</a>	Alignment	not modelled	22.4	10	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
93	<a href="#">c3t8K_</a>	Alignment	not modelled	22.0	17	<b>PDB header:</b> transferase/ligase <b>Chain:</b> K: <b>PDB Molecule:</b> effector protein hopab2; <b>PDBTitle:</b> the avrptob-bak1 complex reveals two structurally similar2 kinaseinteracting domains in a single type iii effector
94	<a href="#">c2l5gA_</a>	Alignment	not modelled	21.7	33	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> g protein pathway suppressor 2; <b>PDBTitle:</b> co-ordinates and 1h, 13c and 15n chemical shift assignments for the2 complex of gps2 53-90 and smrt 167-207
95	<a href="#">d1uzra_</a>	Alignment	not modelled	20.0	8	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
96	<a href="#">c3oqlA_</a>	Alignment	not modelled	19.7	9	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tena homolog; <b>PDBTitle:</b> crystal structure of a tena homolog (pspto1738) from pseudomonas2 syringae pv. tomato str. dc3000 at 2.54 a resolution
97	<a href="#">c1uo0A_</a>	Alignment	not modelled	18.5	48	<b>PDB header:</b> four helix bundle <b>Chain:</b> A: <b>PDB Molecule:</b> general control protein gc4; <b>PDBTitle:</b> structure based engineering of internal molecular surfaces2 of four helix bundles
98	<a href="#">c1uo0B_</a>	Alignment	not modelled	18.5	48	<b>PDB header:</b> four helix bundle <b>Chain:</b> B: <b>PDB Molecule:</b> general control protein gc4; <b>PDBTitle:</b> structure based engineering of internal molecular surfaces2 of four helix bundles
99	<a href="#">c3e0zB_</a>	Alignment	not modelled	16.9	10	<b>PDB header:</b> unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> protein of unknown function; <b>PDBTitle:</b> crystal structure of a putative imidazole glycerol phosphate synthase2 homolog (eubrec_1070) from eubacterium rectale at 1.75 a resolution