
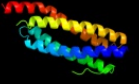


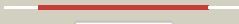
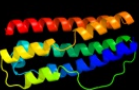
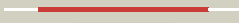


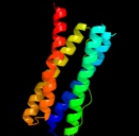

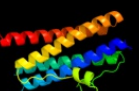



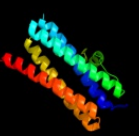

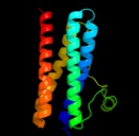



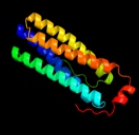


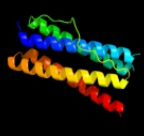

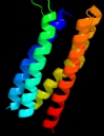
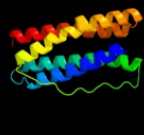





#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3hiuB_</a>	 Alignment		100.0	38	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> uncharacterized protein; <b>PDBTitle:</b> the crystal structure of protein (xcc3681) from xanthomonas2 campestris pv. campestris str. atcc 33913
2	<a href="#">d2gyqa1</a>	 Alignment		100.0	23	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> YciF-like
3	<a href="#">d2gs4a1</a>	 Alignment		100.0	19	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> YciF-like
4	<a href="#">c3oghB_</a>	 Alignment		100.0	94	<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		99.2	16	<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="#">d2htna1</a>	 Alignment		99.2	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
7	<a href="#">c2qqyA_</a>	 Alignment		99.1	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
8	<a href="#">c3gvvC_</a>	 Alignment		99.1	12	<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
9	<a href="#">c3bknB_</a>	 Alignment		99.1	12	<b>PDB header:</b> metal binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> bacterioferritin; <b>PDBTitle:</b> the structure of mycobacterial bacterioferritin
10	<a href="#">c2vzbA_</a>	 Alignment		99.1	15	<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
11	<a href="#">d1nf4a_</a>	 Alignment		99.1	20	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin

12	<a href="#">d2fkza1</a>	Alignment		99.0	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
13	<a href="#">c3fvbB_</a>	Alignment		99.0	14	<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
14	<a href="#">d1jgca_</a>	Alignment		98.9	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
15	<a href="#">d1j30a_</a>	Alignment		98.9	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
16	<a href="#">d1lkoa1</a>	Alignment		98.9	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
17	<a href="#">d1yv1a1</a>	Alignment		98.8	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
18	<a href="#">c3qhbA_</a>	Alignment		98.8	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
19	<a href="#">c2clbP_</a>	Alignment		98.6	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 sulfobus2 solfataricus reveals a bacterioferritin-like di-metal3 binding site within a dps-like dodecameric assembly
20	<a href="#">d1yuza1</a>	Alignment		98.6	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
21	<a href="#">c1dvbA_</a>	Alignment	not modelled	98.6	15	<b>PDB header:</b> electron transport <b>Chain:</b> A: <b>PDB Molecule:</b> rubrerythrin; <b>PDBTitle:</b> rubrerythrin
22	<a href="#">c3e6sD_</a>	Alignment	not modelled	98.5	14	<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
23	<a href="#">d2oh3a1</a>	Alignment	not modelled	98.5	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> AMB4284-like
24	<a href="#">d1nnqa1</a>	Alignment	not modelled	98.4	18	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
25	<a href="#">c1yuzB_</a>	Alignment	not modelled	98.4	15	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> nigerythrin; <b>PDBTitle:</b> partially reduced state of nigerythrin
26	<a href="#">d2fzfa1</a>	Alignment	not modelled	98.4	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
27	<a href="#">d1vlga_</a>	Alignment	not modelled	98.3	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
28	<a href="#">c2hr5B_</a>	Alignment	not modelled	98.3	19	<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="#">d1jiga_</a>	Alignment	not modelled	98.3	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like

					<b>Family:</b> Ferritin
30	<a href="#">dlo9ra_</a>	Alignment	not modelled	98.2	9 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
31	<a href="#">c2c41K_</a>	Alignment	not modelled	98.1	11 <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
32	<a href="#">dlji5a_</a>	Alignment	not modelled	98.1	12 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
33	<a href="#">dltjoa_</a>	Alignment	not modelled	98.1	12 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
34	<a href="#">c3iq1A_</a>	Alignment	not modelled	98.0	14 <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
35	<a href="#">dlkrqa_</a>	Alignment	not modelled	98.0	17 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
36	<a href="#">c2d5kC_</a>	Alignment	not modelled	98.0	11 <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
37	<a href="#">c2vxxB_</a>	Alignment	not modelled	98.0	11 <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
38	<a href="#">c3qz3A_</a>	Alignment	not modelled	98.0	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
39	<a href="#">dlnlqa_</a>	Alignment	not modelled	97.9	14 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
40	<a href="#">c2c6rA_</a>	Alignment	not modelled	97.8	6 <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
41	<a href="#">c3bvkc_</a>	Alignment	not modelled	97.8	14 <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
42	<a href="#">c2chpC_</a>	Alignment	not modelled	97.8	10 <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
43	<a href="#">c3qd8M_</a>	Alignment	not modelled	97.8	15 <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
44	<a href="#">dleuma_</a>	Alignment	not modelled	97.8	12 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
45	<a href="#">c3kwoA_</a>	Alignment	not modelled	97.7	9 <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
46	<a href="#">c2jd8C_</a>	Alignment	not modelled	97.7	14 <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
47	<a href="#">dlrcda_</a>	Alignment	not modelled	97.7	13 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
48	<a href="#">dlji4a_</a>	Alignment	not modelled	97.7	13 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
49	<a href="#">dl1b3a_</a>	Alignment	not modelled	97.7	9 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
50	<a href="#">d2ceia1</a>	Alignment	not modelled	97.6	14 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
51	<a href="#">dl1dpsa_</a>	Alignment	not modelled	97.6	7 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
52	<a href="#">dlmfra_</a>	Alignment	not modelled	97.6	14 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
53	<a href="#">d2fjca1</a>	Alignment	not modelled	97.5	10 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
54	<a href="#">dlr03a_</a>	Alignment	not modelled	97.5	16 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
55	<a href="#">d2bk6a1</a>	Alignment	not modelled	97.5	10 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like

					<b>Family:</b> Ferritin
56	<a href="#">d1vela_</a>	Alignment	not modelled	97.5	9 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
57	<a href="#">d1vjxa_</a>	Alignment	not modelled	97.3	13 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
58	<a href="#">d1s3qa1</a>	Alignment	not modelled	97.3	11 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
59	<a href="#">c1u7mB_</a>	Alignment	not modelled	97.2	30 <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
60	<a href="#">d2yw6a1</a>	Alignment	not modelled	97.0	8 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
61	<a href="#">c2ib0A_</a>	Alignment	not modelled	96.9	13 <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
62	<a href="#">d2ib0a1</a>	Alignment	not modelled	96.9	13 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Rv2844-like
63	<a href="#">c2xgwA_</a>	Alignment	not modelled	96.9	15 <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
64	<a href="#">d2za7a1</a>	Alignment	not modelled	96.9	11 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
65	<a href="#">d1umna_</a>	Alignment	not modelled	96.9	8 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
66	<a href="#">c2f7nA_</a>	Alignment	not modelled	96.9	10 <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
67	<a href="#">c2wlaA_</a>	Alignment	not modelled	96.7	15 <b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> dps-like peroxide resistance protein; <b>PDBTitle:</b> streptococcus pyogenes dpr
68	<a href="#">c2z90D_</a>	Alignment	not modelled	96.5	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
69	<a href="#">c3fseB_</a>	Alignment	not modelled	96.0	23 <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
70	<a href="#">c2rbdB_</a>	Alignment	not modelled	95.9	10 <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
71	<a href="#">c2yjkF_</a>	Alignment	not modelled	95.6	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
72	<a href="#">c1lt1G_</a>	Alignment	not modelled	95.2	28 <b>PDB header:</b> de novo protein <b>Chain:</b> G: <b>PDB Molecule:</b> l13g-df1; <b>PDBTitle:</b> sliding helix induced change of coordination geometry in a2 model di-mn(ii) protein
73	<a href="#">c2pybC_</a>	Alignment	not modelled	95.1	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
74	<a href="#">c3a9qR_</a>	Alignment	not modelled	95.0	15 <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
75	<a href="#">c3q4nA_</a>	Alignment	not modelled	94.0	11 <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
76	<a href="#">c3bt5A_</a>	Alignment	not modelled	93.8	11 <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
77	<a href="#">c2hz8A_</a>	Alignment	not modelled	92.1	18 <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
78	<a href="#">d1z6om1</a>	Alignment	not modelled	90.8	14 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
79	<a href="#">c2qf9B_</a>	Alignment	not modelled	90.5	9 <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
80	<a href="#">c3ddeB_</a>	Alignment	not modelled	89.2	11 <b>PDB header:</b> unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> tena/thi-4 protein, domain of unknown function with a heme <b>PDBTitle:</b> crystal structure of a domain of unknown function with a heme2 oxygenase-like fold (sden_3740) from shewanella

					denitrificans os2173 at 2.30 a resolution
81	<a href="#">dlzs3a1</a>	Alignment	not modelled	88.7	9 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
82	<a href="#">dlzpya1</a>	Alignment	not modelled	87.6	18 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> half-ferritin
83	<a href="#">dlzuja1</a>	Alignment	not modelled	81.8	6 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
84	<a href="#">d2etda1</a>	Alignment	not modelled	75.8	22 <b>Fold:</b> Bromodomain-like <b>Superfamily:</b> LemA-like <b>Family:</b> LemA-like
85	<a href="#">dlq08a</a>	Alignment	not modelled	75.0	18 <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
86	<a href="#">dlbg7a</a>	Alignment	not modelled	73.1	15 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
87	<a href="#">dlrcwa</a>	Alignment	not modelled	72.3	14 <b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> PqqC-like
88	<a href="#">dljkva</a>	Alignment	not modelled	67.4	15 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Manganese catalase (T-catalase)
89	<a href="#">d2cwlal</a>	Alignment	not modelled	59.8	14 <b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Manganese catalase (T-catalase)
90	<a href="#">dlq06a</a>	Alignment	not modelled	59.2	8 <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
91	<a href="#">c2lchA</a>	Alignment	not modelled	51.0	10 <b>PDB header:</b> de novo protein <b>Chain:</b> A: <b>PDB Molecule:</b> protein or38; <b>PDBTitle:</b> solution nmr structure of a protein with a redesigned hydrophobic2 core, northeast structural genomics consortium target or38
92	<a href="#">c3oqlB</a>	Alignment	not modelled	42.6	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
93	<a href="#">dltgga</a>	Alignment	not modelled	32.1	11 <b>Fold:</b> Four-helical up-and-down bundle <b>Superfamily:</b> Histidine-containing phosphotransfer domain, HPT domain <b>Family:</b> Chemotaxis protein CheA P1 domain
94	<a href="#">dli5na</a>	Alignment	not modelled	32.0	10 <b>Fold:</b> Four-helical up-and-down bundle <b>Superfamily:</b> Histidine-containing phosphotransfer domain, HPT domain <b>Family:</b> Chemotaxis protein CheA P1 domain
95	<a href="#">dlz72a1</a>	Alignment	not modelled	31.5	10 <b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
96	<a href="#">c1o06A</a>	Alignment	not modelled	29.7	35 <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)
97	<a href="#">dludda</a>	Alignment	not modelled	26.6	14 <b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
98	<a href="#">c3gp4B</a>	Alignment	not modelled	25.9	13 <b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator, merr family; <b>PDBTitle:</b> crystal structure of putative merr family transcriptional regulator2 from listeria monocytogenes
99	<a href="#">c3oqlA</a>	Alignment	not modelled	25.5	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
100	<a href="#">c2xykB</a>	Alignment	not modelled	22.9	10 <b>PDB header:</b> oxygen storage/transport <b>Chain:</b> B: <b>PDB Molecule:</b> 2-on-2 hemoglobin; <b>PDBTitle:</b> group ii 2-on-2 hemoglobin from the plant pathogen2 agrobacterium tumefaciens