



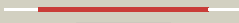




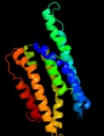



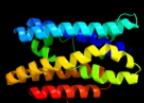





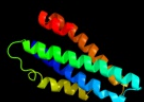





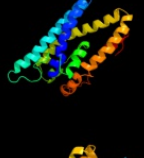
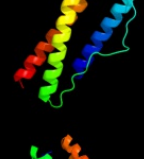


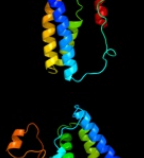
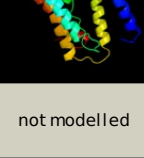


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1otka_</a>	 Alignment		100.0	100	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
2	<a href="#">c3pw1A_</a>	 Alignment		100.0	18	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> phenylacetic acid degradation protein paaa; <b>PDBTitle:</b> the phenylacetyl-coa monooxygenase paaac subcomplex with phenylacetyl-2 coa
3	<a href="#">c3pm5B_</a>	 Alignment		100.0	18	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> benzoyl-coa oxygenase component b; <b>PDBTitle:</b> crystal structure of boxb in mixed valent state with bound benzoyl-coa
4	<a href="#">d1mtyd_</a>	 Alignment		98.2	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
5	<a href="#">c3dhiA_</a>	 Alignment		98.1	15	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> toluene 4-monooxygenase hydroxylase alpha subunit; <b>PDBTitle:</b> crystal structure of reduced toluene 4-monooxygenase hydroxylase2 complexed with effector protein
6	<a href="#">d2inca1</a>	 Alignment		98.1	12	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
7	<a href="#">c2inpD_</a>	 Alignment		97.7	17	<b>PDB header:</b> oxidoreductase <b>Chain:</b> D: <b>PDB Molecule:</b> phenol hydroxylase component phl; <b>PDBTitle:</b> structure of the phenol hydroxylase-regulatory protein2 complex
8	<a href="#">d2oc5a1</a>	 Alignment		97.7	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> PMT1231-like
9	<a href="#">d2fzfa1</a>	 Alignment		97.6	8	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
10	<a href="#">c3ez0D_</a>	 Alignment		97.3	10	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> D: <b>PDB Molecule:</b> uncharacterized protein with ferritin-like fold; <b>PDBTitle:</b> crystal structure of protein of unknown function with ferritin-like2 fold (yp_832262.1) from arthrobacter sp. fb24 at 2.33 a resolution
11	<a href="#">c3dhgB_</a>	 Alignment		97.1	15	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> toluene 4-monooxygenase hydroxylase beta subunit; <b>PDBTitle:</b> crystal struture of toluene 4-monooxygenase hydroxylase

12	<a href="#">d1mhyd_</a>	Alignment		96.9	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
13	<a href="#">d2incb1</a>	Alignment		96.8	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
14	<a href="#">d1mhyb_</a>	Alignment		96.8	12	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
15	<a href="#">c2innA_</a>	Alignment		96.5	11	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> phenol hydroxylase component phn; <b>PDBTitle:</b> structure of the phenol hydroxylase-regulatory protein2 complex
16	<a href="#">d1jgca_</a>	Alignment		96.0	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
17	<a href="#">d2oh3a1</a>	Alignment		95.7	9	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> AMB4284-like
18	<a href="#">c3bknB_</a>	Alignment		95.5	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
19	<a href="#">c3ee4A_</a>	Alignment		95.3	11	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> probable ribonucleoside-diphosphate reductase; <b>PDBTitle:</b> r2-like ligand binding mn/fe oxidase from m. tuberculosis
20	<a href="#">c2o1zA_</a>	Alignment		95.2	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ribonucleotide reductase subunit r2; <b>PDBTitle:</b> plasmodium vivax ribonucleotide reductase subunit r22 (pv086155)
21	<a href="#">d1rcwa_</a>	Alignment	not modelled	94.8	11	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> PqqC-like
22	<a href="#">d1mxra_</a>	Alignment	not modelled	94.8	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
23	<a href="#">c2vzbA_</a>	Alignment	not modelled	94.8	9	<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
24	<a href="#">d1vjxa_</a>	Alignment	not modelled	94.6	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
25	<a href="#">c2vuxB_</a>	Alignment	not modelled	94.3	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase subunit m2 b; <b>PDBTitle:</b> human ribonucleotide reductase, subunit m2 b
26	<a href="#">d1j30a_</a>	Alignment	not modelled	94.3	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
27	<a href="#">c2rccC_</a>	Alignment	not modelled	94.1	8	<b>PDB header:</b> oxidoreductase <b>Chain:</b> C: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase subunit beta; <b>PDBTitle:</b> crystal structure of putative class i ribonucleotide reductase2 (np_241368.1) from bacillus halodurans at 1.90 a resolution
						<b>PDB header:</b> unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> tena/thi-4 protein, domain of unknown function

28	<a href="#">c3ddeB_</a>	Alignment	not modelled	93.9	12	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
29	<a href="#">c2p1iA_</a>	Alignment	not modelled	93.8	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ribonucleotide reductase, small chain; <b>PDBTitle:</b> plasmodium yoelii ribonucleotide reductase subunit r2 (py03671)
30	<a href="#">c2clbP_</a>	Alignment	not modelled	93.7	10	<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
31	<a href="#">dlw68a_</a>	Alignment	not modelled	93.4	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
32	<a href="#">c1h0oA_</a>	Alignment	not modelled	93.4	11	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase; <b>PDBTitle:</b> cobalt substitution of mouse r2 ribonucleotide reductase to2 model the reactive differrous state
33	<a href="#">dlmsa_</a>	Alignment	not modelled	93.2	9	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
34	<a href="#">c3hf1B_</a>	Alignment	not modelled	93.1	11	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase subunit m2 <b>PDBTitle:</b> crystal structure of human p53r2
35	<a href="#">c3fvbB_</a>	Alignment	not modelled	92.9	11	<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
36	<a href="#">c3gvyC_</a>	Alignment	not modelled	92.3	11	<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
37	<a href="#">c1yfdA_</a>	Alignment	not modelled	92.2	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase 1 beta <b>PDBTitle:</b> crystal structure of the y122h mutant of ribonucleotide2 reductase r2 protein from e. coli
38	<a href="#">d2htna1</a>	Alignment	not modelled	92.2	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
39	<a href="#">dl1koa1</a>	Alignment	not modelled	92.1	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
40	<a href="#">d2fkza1</a>	Alignment	not modelled	92.0	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
41	<a href="#">c3bjdA_</a>	Alignment	not modelled	91.8	15	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> putative 3-oxoacyl-(acyl-carrier-protein) synthase; <b>PDBTitle:</b> crystal structure of putative 3-oxoacyl-(acyl-carrier-protein)2 synthase from pseudomonas aeruginosa
42	<a href="#">dlnf4a_</a>	Alignment	not modelled	90.3	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
43	<a href="#">c3qhbA_</a>	Alignment	not modelled	90.3	15	<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
44	<a href="#">dlyv1a1</a>	Alignment	not modelled	90.2	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
45	<a href="#">c1dvbA_</a>	Alignment	not modelled	89.7	8	<b>PDB header:</b> electron transport <b>Chain:</b> A: <b>PDB Molecule:</b> rubrerythrin; <b>PDBTitle:</b> rubrerythrin
46	<a href="#">dlsyya_</a>	Alignment	not modelled	88.3	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
47	<a href="#">clsyyA_</a>	Alignment	not modelled	88.3	11	<b>PDB header:</b> replication,oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase beta chain; <b>PDBTitle:</b> crystal structure of the r2 subunit of ribonucleotide reductase from2 chlamydia trachomatis
48	<a href="#">dlza0a1</a>	Alignment	not modelled	88.3	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
49	<a href="#">c1r2fB_</a>	Alignment	not modelled	87.7	12	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> protein (ribonucleotide reductase r2); <b>PDBTitle:</b> ribonucleotide reductase r2f protein from salmonella2 typhimurium
50	<a href="#">dljkva_</a>	Alignment	not modelled	87.6	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Manganese catalase (T-catalase)
51	<a href="#">dlr2fa_</a>	Alignment	not modelled	87.5	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
52	<a href="#">dluzra_</a>	Alignment	not modelled	87.4	12	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
53	<a href="#">c3oqlB_</a>	Alignment	not modelled	87.0	10	<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

54	<a href="#">d1afra_</a>	<div><div></div></div> Alignment	not modelled	85.7	12	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
55	<a href="#">c3oghB_</a>	<div><div></div></div> Alignment	not modelled	85.3	7	<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
56	<a href="#">d1nnqa1</a>	<div><div></div></div> Alignment	not modelled	85.0	17	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
57	<a href="#">c3e6sD_</a>	<div><div></div></div> Alignment	not modelled	84.5	11	<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
58	<a href="#">d3dhza1</a>	<div><div></div></div> Alignment	not modelled	84.3	10	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
59	<a href="#">c3r2rA_</a>	<div><div></div></div> Alignment	not modelled	83.4	13	<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)
60	<a href="#">c2qqyA_</a>	<div><div></div></div> Alignment	not modelled	83.0	11	<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
61	<a href="#">c2rd3A_</a>	<div><div></div></div> Alignment	not modelled	82.8	12	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of tena homologue (hp1287) from2 helicobacter pylori
62	<a href="#">d2itba1</a>	<div><div></div></div> Alignment	not modelled	81.6	14	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> MiaE-like
63	<a href="#">d1jk0a_</a>	<div><div></div></div> Alignment	not modelled	80.0	9	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
64	<a href="#">c1smqD_</a>	<div><div></div></div> Alignment	not modelled	79.9	9	<b>PDB header:</b> oxidoreductase <b>Chain:</b> D: <b>PDB Molecule:</b> ribonucleoside-diphosphate reductase small chain <b>PDBTitle:</b> structure of the ribonucleotide reductase rnn2 homodimer2 from saccharomyces cerevisiae
65	<a href="#">d1vlga_</a>	<div><div></div></div> Alignment	not modelled	79.1	16	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
66	<a href="#">c3q4nA_</a>	<div><div></div></div> Alignment	not modelled	77.4	17	<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
67	<a href="#">c2hr5B_</a>	<div><div></div></div> Alignment	not modelled	77.2	17	<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
68	<a href="#">d1yza1</a>	<div><div></div></div> Alignment	not modelled	75.6	10	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
69	<a href="#">c2qcxA_</a>	<div><div></div></div> Alignment	not modelled	73.8	14	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional activator tena; <b>PDBTitle:</b> crystal structure of bacillus subtilis tena y112f mutant complexed2 with formyl aminomethyl pyrimidine
70	<a href="#">d1jk0b_</a>	<div><div></div></div> Alignment	not modelled	73.0	10	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
71	<a href="#">d1oquc_</a>	<div><div></div></div> Alignment	not modelled	69.5	11	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
72	<a href="#">c3bykC_</a>	<div><div></div></div> Alignment	not modelled	68.0	11	<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
73	<a href="#">d1z72a1</a>	<div><div></div></div> Alignment	not modelled	66.8	17	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
74	<a href="#">d1euma_</a>	<div><div></div></div> Alignment	not modelled	66.5	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
75	<a href="#">d1krqa_</a>	<div><div></div></div> Alignment	not modelled	52.6	15	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
76	<a href="#">c3oqlA_</a>	<div><div></div></div> Alignment	not modelled	46.8	11	<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
77	<a href="#">d1otva_</a>	<div><div></div></div> Alignment	not modelled	45.8	10	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> PqqC-like
78	<a href="#">d2hgsa4</a>	<div><div></div></div> Alignment	not modelled	44.4	22	<b>Fold:</b> ATP-grasp <b>Superfamily:</b> Glutathione synthetase ATP-binding domain-like <b>Family:</b> Eukaryotic glutathione synthetase ATP-binding domain
79	<a href="#">c3bt5A_</a>	<div><div></div></div> Alignment	not modelled	42.0	14	<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></div></div>				<b>Fold:</b> Ferritin-like

80	<a href="#">d1s3qa1</a>	Alignment	not modelled	40.7	13	<b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin
81	<a href="#">c3chtA</a>	Alignment	not modelled	40.3	10	<b>PDB header:</b> oxidoreductase <b>Chain:</b> A: <b>PDB Molecule:</b> p-aminobenzoate n-oxygenase; <b>PDBTitle:</b> crystal structure of di-iron aurf with partially bound ligand
82	<a href="#">c3qd8M</a>	Alignment	not modelled	38.3	8	<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
83	<a href="#">d1udda</a>	Alignment	not modelled	36.2	9	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
84	<a href="#">d2gm8a1</a>	Alignment	not modelled	34.0	12	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
85	<a href="#">c2y69R</a>	Alignment	not modelled	29.9	19	<b>PDB header:</b> electron transport <b>Chain:</b> R: <b>PDB Molecule:</b> cytochrome c oxidase subunit 5a; <b>PDBTitle:</b> bovine heart cytochrome c oxidase re-refined with molecular2 oxygen
86	<a href="#">c3no6B</a>	Alignment	not modelled	28.4	11	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional activator tena; <b>PDBTitle:</b> crystal structure of a putative thiaminase ii (se1693) from2 staphylococcus epidermidis atcc 12228 at 1.65 a resolution
87	<a href="#">d1v54e</a>	Alignment	not modelled	27.1	19	<b>Fold:</b> alpha-alpha superhelix <b>Superfamily:</b> Cytochrome c oxidase subunit E <b>Family:</b> Cytochrome c oxidase subunit E
88	<a href="#">d1zpya1</a>	Alignment	not modelled	26.4	17	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> half-ferritin
89	<a href="#">c3qz3A</a>	Alignment	not modelled	25.9	13	<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
90	<a href="#">c3fpvC</a>	Alignment	not modelled	24.7	35	<b>PDB header:</b> heme binding protein <b>Chain:</b> C: <b>PDB Molecule:</b> extracellular haem-binding protein; <b>PDBTitle:</b> crystal structure of hbps
91	<a href="#">c1to9A</a>	Alignment	not modelled	24.0	13	<b>PDB header:</b> biosynthetic protein <b>Chain:</b> A: <b>PDB Molecule:</b> thi-4 protein; <b>PDBTitle:</b> crystal structure of thi-4 protein from bacillus subtilis
92	<a href="#">d1to9a</a>	Alignment	not modelled	24.0	13	<b>Fold:</b> Heme oxygenase-like <b>Superfamily:</b> Heme oxygenase-like <b>Family:</b> TENA/THI-4
93	<a href="#">c2jd8C</a>	Alignment	not modelled	22.4	17	<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
94	<a href="#">c2yumA</a>	Alignment	not modelled	20.9	21	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> zinc finger zz-type-containing protein 3; <b>PDBTitle:</b> solution structure of the myb-like dna-binding domain of2 human zzz3 protein
95	<a href="#">c3h3mB</a>	Alignment	not modelled	20.6	14	<b>PDB header:</b> structural genomics <b>Chain:</b> B: <b>PDB Molecule:</b> flagellar protein flit; <b>PDBTitle:</b> crystal structure of flagellar protein flit from bordetella2 bronchiseptica
96	<a href="#">c1yuzB</a>	Alignment	not modelled	20.1	10	<b>PDB header:</b> oxidoreductase <b>Chain:</b> B: <b>PDB Molecule:</b> nigerythrin; <b>PDBTitle:</b> partially reduced state of nigerythrin
97	<a href="#">d1mtyb</a>	Alignment	not modelled	15.8	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ribonucleotide reductase-like
98	<a href="#">d2qdya1</a>	Alignment	not modelled	14.2	15	<b>Fold:</b> Nitrile hydratase alpha chain <b>Superfamily:</b> Nitrile hydratase alpha chain <b>Family:</b> Nitrile hydratase alpha chain
99	<a href="#">d1r03a</a>	Alignment	not modelled	14.0	13	<b>Fold:</b> Ferritin-like <b>Superfamily:</b> Ferritin-like <b>Family:</b> Ferritin