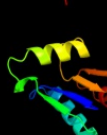
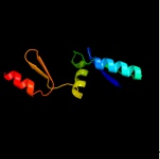
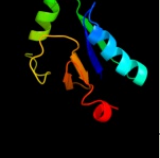
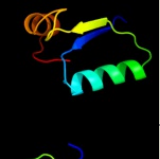
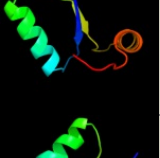
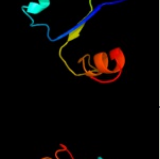
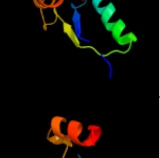
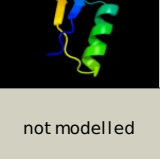


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1j9ba_	 Alignment		100.0	93	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: ArsC-like
2	c3f0iA_	 Alignment		100.0	46	PDB header: oxidoreductase Chain: A: PDB Molecule: arsenate reductase; PDBTitle: arsenate reductase from vibrio cholerae.
3	c3l78A_	 Alignment		100.0	18	PDB header: transcription Chain: A: PDB Molecule: regulatory protein spx; PDBTitle: the crystal structure of smu.1142c from streptococcus mutans ua159
4	c3rdwB_	 Alignment		100.0	44	PDB header: oxidoreductase Chain: B: PDB Molecule: putative arsenate reductase; PDBTitle: putative arsenate reductase from yersinia pestis
5	c3gkxB_	 Alignment		100.0	23	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: putative arsc family related protein; PDBTitle: crystal structure of putative arsc family related protein from2 bacteroides fragilis
6	d1z3ea1	 Alignment		100.0	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: ArsC-like
7	d1rw1a_	 Alignment		100.0	17	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: ArsC-like
8	c2kokA_	 Alignment		100.0	15	PDB header: oxidoreductase Chain: A: PDB Molecule: arsenate reductase; PDBTitle: solution structure of an arsenate reductase (arsc) related protein2 from brucella melitensis. seattle structural genomics center for3 infectious disease target braba.00007.a.
9	c3fz4A_	 Alignment		100.0	18	PDB header: oxidoreductase Chain: A: PDB Molecule: putative arsenate reductase; PDBTitle: the crystal structure of a possible arsenate reductase from2 streptococcus mutans ua159
10	d1h75a_	 Alignment		98.9	21	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
11	c3ic4A_	 Alignment		98.6	21	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin (grx-1); PDBTitle: the crystal structure of the glutaredoxin(grx-1) from archaeoglobus2 fulgidus

12	c3nznA_	Alignment		98.6	12	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin; PDBTitle: the crystal structure of the glutaredoxin from methanosarcina mazei2 go1
13	dlr7ha_	Alignment		98.5	21	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
14	c2khpA_	Alignment		98.1	22	PDB header: electron transport Chain: A: PDB Molecule: glutaredoxin; PDBTitle: solution structure of glutaredoxin from brucella melitensis
15	c3lqcA_	Alignment		98.1	12	PDB header: unknown function Chain: A: PDB Molecule: glutaredoxin 1; PDBTitle: crystal structure of glutaredoxin 1 from francisella2 tularensis
16	c1nm3B_	Alignment		98.0	15	PDB header: electron transport Chain: B: PDB Molecule: protein hi0572; PDBTitle: crystal structure of heamophilus influenza hybrid-prx5
17	dlfova_	Alignment		98.0	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
18	c2klxA_	Alignment		97.9	18	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin; PDBTitle: solution structure of glutaredoxin from bartonella henselae str.2 houston
19	dlnm3a1	Alignment		97.9	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
20	dltlva_	Alignment		97.8	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: SH3BGR (SH3-binding, glutamic acid-rich protein-like)
21	c3qmxA_	Alignment	not modelled	97.6	18	PDB header: electron transport Chain: A: PDB Molecule: glutaredoxin a; PDBTitle: x-ray crystal structure of synechocystis sp. pcc 6803 glutaredoxin a
22	c3fzaA_	Alignment	not modelled	97.6	12	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin; PDBTitle: crystal structure of poplar glutaredoxin s12 in complex with2 glutathione and beta-mercaptoethanol
23	c2ht9A_	Alignment	not modelled	97.6	9	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin-2; PDBTitle: the structure of dimeric human glutaredoxin 2
24	c2e7pC_	Alignment	not modelled	97.5	11	PDB header: electron transport Chain: C: PDB Molecule: glutaredoxin; PDBTitle: crystal structure of the holo form of glutaredoxin c1 from populus2 tremula x tremuloides
25	c3h8qB_	Alignment	not modelled	97.5	15	PDB header: oxidoreductase Chain: B: PDB Molecule: thioredoxin reductase 3; PDBTitle: crystal structure of glutaredoxin domain of human thioredoxin2 reductase 3
26	dlabaa_	Alignment	not modelled	97.4	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
27	dllegoa_	Alignment	not modelled	97.2	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
28	dlwika_	Alignment	not modelled	96.9	26	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
						Fold: Thioredoxin fold

29	d1z9ha2	Alignment	not modelled	96.8	15	Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
30	d1k0da2	Alignment	not modelled	96.8	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
31	d1ktea_	Alignment	not modelled	96.7	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
32	cloyjC_	Alignment	not modelled	96.6	18	PDB header: transferase Chain: C: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure solution of rice gst1 (osgstu1) in complex with2 glutathione.
33	c2vo4A_	Alignment	not modelled	96.6	18	PDB header: transferase Chain: A: PDB Molecule: 2,4-d inducible glutathione s-transferase; PDBTitle: glutathione transferase from glycine max
34	clykaA_	Alignment	not modelled	96.6	25	PDB header: electron transport Chain: A: PDB Molecule: monothiol glutaredoxin ydhc; PDBTitle: solution structure of grx4, a monothiol glutaredoxin from2 e. coli.
35	d1tza_	Alignment	not modelled	96.5	16	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
36	c1k0dB_	Alignment	not modelled	96.5	9	PDB header: gene regulation Chain: B: PDB Molecule: ure2 protein; PDBTitle: ure2p in complex with glutathione
37	c1g7oA_	Alignment	not modelled	96.5	12	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin 2; PDBTitle: nmr solution structure of reduced e. coli glutaredoxin 2
38	d1eema2	Alignment	not modelled	96.4	21	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
39	c2hzfA_	Alignment	not modelled	96.4	19	PDB header: electron transport, oxidoreductase Chain: A: PDB Molecule: glutaredoxin-1; PDBTitle: crystal structures of a poxviral glutaredoxin in the oxidized and2 reduced states show redox-correlated structural changes
40	d1k0dd2	Alignment	not modelled	96.3	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
41	c2ct6A_	Alignment	not modelled	96.3	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain-binding glutamic acid-rich-like PDBTitle: solution structure of the sh3 domain-binding glutamic acid-2 rich-like protein 2
42	d1ljra2	Alignment	not modelled	96.3	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
43	d1jhba_	Alignment	not modelled	96.3	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
44	c1yy7A_	Alignment	not modelled	96.2	16	PDB header: transcription Chain: A: PDB Molecule: stringent starvation protein a; PDBTitle: crystal structure of stringent starvation protein a (sspa),2 an rna polymerase-associated transcription factor
45	c1oktA_	Alignment	not modelled	96.2	19	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: x-ray structure of glutathione s-transferase from the2 malarial parasite plasmodium falciparum
46	d1oyja2	Alignment	not modelled	96.2	7	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
47	c1ljrB_	Alignment	not modelled	96.1	12	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: glutathione transferase (hgst t2-2) from human
48	c3l4nA_	Alignment	not modelled	95.9	19	PDB header: oxidoreductase Chain: A: PDB Molecule: monothiol glutaredoxin-6; PDBTitle: crystal structure of yeast monothiol glutaredoxin grx6
49	c2jl4A_	Alignment	not modelled	95.8	16	PDB header: isomerase Chain: A: PDB Molecule: maleylpyruvate isomerase; PDBTitle: holo structure of maleyl pyruvate isomerase, a bacterial2 glutathione-s-transferase in zeta class
50	c3m3mA_	Alignment	not modelled	95.8	9	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of glutathione s-transferase from pseudomonas2 fluorescens [pf-5]
51	d1gwca2	Alignment	not modelled	95.8	5	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
52	c1z9hB_	Alignment	not modelled	95.8	9	PDB header: isomerase Chain: B: PDB Molecule: membrane-associated prostaglandin e synthase-2; PDBTitle: microsomal prostaglandin e synthase type-2
53	c3qagA_	Alignment	not modelled	95.8	18	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase omega-2; PDBTitle: human glutathione transferase o2 with glutathione -new crystal form
54	c1gwcC_	Alignment	not modelled	95.7	5	PDB header: transferase Chain: C: PDB Molecule: glutathione s-transferase tsi-1; PDBTitle: the structure of a tau class glutathione s-transferase from2 wheat, active in herbicide detoxification
55	d1gnwa2	Alignment	not modelled	95.7	19	Fold: Thioredoxin fold Superfamily: Thioredoxin-like

					Family: Glutathione S-transferase (GST), N-terminal domain
56	c1eemA	Alignment	not modelled	95.7	21 PDB header: transferase Chain: A: PDB Molecule: glutathione-s-transferase; PDBTitle: glutathione transferase from homo sapiens
57	c3cbuB	Alignment	not modelled	95.6	3 PDB header: transferase Chain: B: PDB Molecule: probable gst-related protein; PDBTitle: crystal structure of a putative glutathione s-transferase (reut_a1011)2 from ralstonia eutropha jmp134 at 2.05 a resolution
58	c3ay8A	Alignment	not modelled	95.6	33 PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: glutathione s-transferase unclassified 2 from bombyx mori
59	c3d5jB	Alignment	not modelled	95.6	15 PDB header: oxidoreductase Chain: B: PDB Molecule: glutaredoxin-2, mitochondrial; PDBTitle: structure of yeast grx2-c30s mutant with glutathionyl mixed2 disulfide
60	c2imiA	Alignment	not modelled	95.6	18 PDB header: transferase Chain: A: PDB Molecule: epsilon-class glutathione s-transferase; PDBTitle: structures of an insect epsilon-class glutathione s-2 transferase from the malaria vector anopheles gambiae:3 evidence for high ddt-detoxifying activity
61	c3ipzA	Alignment	not modelled	95.6	17 PDB header: electron transport, oxidoreductase Chain: A: PDB Molecule: monothiol glutaredoxin-s14, chloroplastic; PDBTitle: crystal structure of arabidopsis monothiol glutaredoxin atgrxcp
62	c3c1sA	Alignment	not modelled	95.5	11 PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin-1; PDBTitle: crystal structure of grx1 in glutathionylated form
63	d1v2aa2	Alignment	not modelled	95.5	8 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
64	d1wjka	Alignment	not modelled	95.5	11 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
65	c2on7A	Alignment	not modelled	95.4	21 PDB header: transferase Chain: A: PDB Molecule: na glutathione s-transferase 1; PDBTitle: structure of nagst-1
66	d1e6ba2	Alignment	not modelled	95.4	9 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
67	d1jlva2	Alignment	not modelled	95.3	16 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
68	c3lypA	Alignment	not modelled	95.3	22 PDB header: transcription Chain: A: PDB Molecule: stringent starvation protein a; PDBTitle: structure of stringent starvation protein a homolog from pseudomonas2 fluorescens
69	c3lykA	Alignment	not modelled	95.2	10 PDB header: transport protein Chain: A: PDB Molecule: stringent starvation protein a homolog; PDBTitle: structure of stringent starvation protein a homolog from2 haemophilus influenzae
70	d1aw9a2	Alignment	not modelled	95.2	12 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
71	c2wulB	Alignment	not modelled	95.2	17 PDB header: oxidoreductase Chain: B: PDB Molecule: glutaredoxin related protein 5; PDBTitle: crystal structure of the human glutaredoxin 5 with bound2 glutathione in an fes cluster
72	c1e6bA	Alignment	not modelled	95.1	9 PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of a zeta class glutathione s-transferase2 from arabidopsis thaliana
73	c3bbyA	Alignment	not modelled	95.1	14 PDB header: transferase Chain: A: PDB Molecule: uncharacterized gst-like protein yfcf; PDBTitle: crystal structure of glutathione s-transferase (np_416804.1) from2 escherichia coli k12 at 1.85 a resolution
74	d1okta2	Alignment	not modelled	95.1	19 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
75	c3lg6B	Alignment	not modelled	95.1	15 PDB header: transferase Chain: B: PDB Molecule: putative glutathione transferase; PDBTitle: crystal structure of putative glutathione transferase from2 coccidioides immitis
76	c1zl9A	Alignment	not modelled	95.1	9 PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 5; PDBTitle: crystal structure of a major nematode c.elegans specific gst (ce01613)
77	c2jacA	Alignment	not modelled	95.1	13 PDB header: electron transport Chain: A: PDB Molecule: glutaredoxin-1; PDBTitle: glutaredoxin grx1p c30s mutant from yeast
78	d1jlwa2	Alignment	not modelled	95.0	16 Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
79	c3nivD	Alignment	not modelled	95.0	9 PDB header: isomerase Chain: D: PDB Molecule: glutathione s-transferase; PDBTitle: the crystal structure of glutathione s-transferase from legionella2 pneumophila
80	c3m8nA	Alignment	not modelled	94.9	10 PDB header: transferase Chain: A: PDB Molecule: possible glutathione s-transferase; PDBTitle: crystal structure of a possible gutathione s-tranferase from2 rhodopseudomonas palustris
81	c2c3nB	Alignment	not modelled	94.9	6 PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase theta 1; PDBTitle: human glutathione-s-transferase t1-1, apo form

82	d1pn9a2	Alignment	not modelled	94.9	16	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
83	d1axda2	Alignment	not modelled	94.8	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
84	c3lxzD	Alignment	not modelled	94.8	6	PDB header: transferase Chain: D: PDB Molecule: glutathione s-transferase family protein; PDBTitle: structure of probable glutathione s-transferase(pp0183) from2 pseudomonas putida
85	c2ws2B	Alignment	not modelled	94.7	9	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: the 2 angstrom structure of a nu-class gst from haemonchus contortus
86	c1v2aD	Alignment	not modelled	94.7	6	PDB header: transferase Chain: D: PDB Molecule: glutathione transferase gst1-6; PDBTitle: glutathione s-transferase 1-6 from anopheles dirus species b
87	d1g7oa2	Alignment	not modelled	94.7	8	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
88	d1lloa	Alignment	not modelled	94.7	20	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
89	d1pd212	Alignment	not modelled	94.6	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
90	c3touB	Alignment	not modelled	94.5	6	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase protein; PDBTitle: crystal structure of glutathione transferase (target efi-501058) from2 ralstonia solanacearum gmi1000 with gsh bound
91	c3rbtD	Alignment	not modelled	94.5	7	PDB header: transferase Chain: D: PDB Molecule: glutathione transferase o1; PDBTitle: crystal structure of glutathione s-transferase omega 3 from the2 silkworm bombyx mori
92	c3gx8A	Alignment	not modelled	94.5	18	PDB header: electron transport Chain: A: PDB Molecule: monothiol glutaredoxin-5, mitochondrial; PDBTitle: structural and biochemical characterization of yeast2 monothiol glutaredoxin grx5
93	c1jlvA	Alignment	not modelled	94.4	16	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase gst1-3; PDBTitle: anopheles dirus species b glutathione s-transferases 1-3
94	d1fw1a2	Alignment	not modelled	94.4	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
95	d1r5aa2	Alignment	not modelled	94.4	22	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
96	c1gnwA	Alignment	not modelled	94.3	19	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: structure of glutathione s-transferase
97	c3fy7B	Alignment	not modelled	94.3	17	PDB header: transport protein Chain: B: PDB Molecule: chloride intracellular channel protein 3; PDBTitle: crystal structure of homo sapiens clic3
98	c3f6fA	Alignment	not modelled	94.2	14	PDB header: transferase Chain: A: PDB Molecule: cg18548-pa (ip02196p) (ip02193p); PDBTitle: crystal structure of glutathione transferase dmgstd10 from2 drosophila melanogaster
99	c1jlvA	Alignment	not modelled	94.2	16	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase gst1-4; PDBTitle: anopheles dirus species b glutathione s-transferases 1-4
100	d1v58a1	Alignment	not modelled	94.1	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: DsbC/DsbG C-terminal domain-like
101	c1v57A	Alignment	not modelled	94.1	12	PDB header: isomerase Chain: A: PDB Molecule: thiol:disulfide interchange protein dsbg; PDBTitle: crystal structure of the disulfide bond isomerase dsbg
102	c2cz2A	Alignment	not modelled	94.1	13	PDB header: isomerase, transferase Chain: A: PDB Molecule: maleylacetoacetate isomerase; PDBTitle: crystal structure of glutathione transferase zeta 1-12 (maleylacetoacetate isomerase) from mus musculus (form-1 crystal)
103	c1r5aA	Alignment	not modelled	94.0	22	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase; PDBTitle: glutathione s-transferase
104	c3gx0A	Alignment	not modelled	94.0	13	PDB header: transferase Chain: A: PDB Molecule: gst-like protein yfcg; PDBTitle: crystal structure of gsh-dependent disulfide bond2 oxidoreductase
105	c3ic8D	Alignment	not modelled	93.8	11	PDB header: structural genomics, unknown function Chain: D: PDB Molecule: uncharacterized gst-like proteinprotein; PDBTitle: the crystal structure of a gst-like protein from pseudomonas syringae2 to 2.4a
106	c2x64A	Alignment	not modelled	93.8	18	PDB header: transferase Chain: A: PDB Molecule: glutathione-s-transferase; PDBTitle: glutathione-s-transferase from xylella fastidiosa
107	d1tw9a2	Alignment	not modelled	93.8	6	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
108	c1byeA	Alignment	not modelled	93.8	10	PDB header: transferase Chain: A: PDB Molecule: protein (glutathione s-transferase); PDBTitle: glutathione s-transferase i from mais in complex with2

						atrazine glutathione conjugate
109	clu6tA_	Alignment	not modelled	93.7	12	PDB header: protein binding, signaling protein Chain: A: PDB Molecule: sh3 domain-binding glutamic acid-rich-like PDBTitle: crystal structure of the human sh3 binding glutamic-rich2 protein like
110	claw9A_	Alignment	not modelled	93.7	13	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase iii; PDBTitle: structure of glutathione s-transferase iii in apo form
111	c2on5E_	Alignment	not modelled	93.5	12	PDB header: transferase Chain: E: PDB Molecule: na glutathione s-transferase 2; PDBTitle: structure of nagst-2
112	clk0nB_	Alignment	not modelled	93.5	9	PDB header: metal transport Chain: B: PDB Molecule: chloride intracellular channel protein 1; PDBTitle: chloride intracellular channel 1 (clic1) complexed with glutathione
113	clyq1A_	Alignment	not modelled	92.9	7	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: structural genomics of caenorhabditis elegans: glutathione2 s-transferase
114	c3ergA_	Alignment	not modelled	92.8	8	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 2; PDBTitle: crystal structure of gtt2 from saccharomyces cerevisiae in complex2 with glutathione sulfinate
115	c2r4vA_	Alignment	not modelled	92.8	14	PDB header: transport protein Chain: A: PDB Molecule: chloride intracellular channel protein 2; PDBTitle: structure of human clic2, crystal form a
116	d1tu7a2	Alignment	not modelled	92.6	9	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
117	d1n2aa2	Alignment	not modelled	92.6	20	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
118	c2aheA_	Alignment	not modelled	92.4	11	PDB header: metal transport Chain: A: PDB Molecule: chloride intracellular channel protein 4; PDBTitle: crystal structure of a soluble form of clic4. intercellular2 chloride ion channel
119	d1k0ma2	Alignment	not modelled	92.2	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
120	c2fgxA_	Alignment	not modelled	92.1	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative thioredoxin; PDBTitle: solution nmr structure of protein ne2328 from nitrosomonas2 europaea. northeast structural genomics consortium target3 net3.