



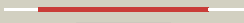



















#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1f2eB_	 Alignment		100.0	42	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: structure of sphingomonad, glutathione s-transferase complexed with2 glutathione
2	c2pmtA_	 Alignment		100.0	54	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase; PDBTitle: glutathione transferase from proteus mirabilis
3	c2gdrA_	 Alignment		100.0	49	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of a bacterial glutathione transferase
4	c3uarA_	 Alignment		100.0	49	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of glutathione transferase (target efi-501774) from2 methylococcus capsulatus str. bath with gsh bound
5	c2ntoA_	 Alignment		100.0	39	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: structure of the glutathione transferase from ochrobactrum anthropi in2 complex with glutathione
6	c3c8eB_	 Alignment		100.0	19	PDB header: transferase Chain: B: PDB Molecule: yghu, glutathione s-transferase homologue; PDBTitle: crystal structure analysis of yghu from e. coli
7	c1n2aB_	 Alignment		100.0	100	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of a bacterial glutathione transferase2 from escherichia coli with glutathione sulfonate in the3 active site
8	c2x64A_	 Alignment		100.0	37	PDB header: transferase Chain: A: PDB Molecule: glutathione-s-transferase; PDBTitle: glutathione-s-transferase from xylella fastidiosa
9	c1l1rB_	 Alignment		100.0	25	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: glutathione transferase (hgst t2-2) from human
10	c1c72A_	 Alignment		100.0	17	PDB header: transferase Chain: A: PDB Molecule: protein (glutathione s-transferase); PDBTitle: tyr115, gln165 and trp209 contribute to the 1,2-epoxy-3-(p-2 nitrophenoxy)propane conjugating activities of glutathione3 s-transferase cgstm1-1
11	c3m0fA_	 Alignment		100.0	16	PDB header: transferase Chain: A: PDB Molecule: uncharacterized protein gst_n; PDBTitle: crystal structure of glutathione s transferase in complex2 with glutathione from pseudomonas fluorescens

12	c3gx0A_	Alignment		100.0	18	PDB header: transferase Chain: A: PDB Molecule: gst-like protein yfcg; PDBTitle: crystal structure of gsh-dependent disulfide bond2 oxidoreductase
13	c2fheA_	Alignment		100.0	16	PDB header: transferase/substrate Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: fasciola hepatica glutathione s-transferase isoform 1 in complex with2 glutathione
14	c1gtuB_	Alignment		100.0	16	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: ligand-free human glutathione s-transferase m1a-1a
15	c1b8xA_	Alignment		100.0	14	PDB header: signal protein Chain: A: PDB Molecule: protein (aml-1b); PDBTitle: glutathione s-transferase fused with the nuclear matrix targeting2 signal of the transcription factor aml-1
16	c3ergA_	Alignment		100.0	26	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 2; PDBTitle: crystal structure of gtt2 from saccharomyces cerevisiae in complex2 with glutathione sulfnate
17	c3lykA_	Alignment		100.0	23	PDB header: transport protein Chain: A: PDB Molecule: stringent starvation protein a homolog; PDBTitle: structure of stringent starvation protein a homolog from2 haemophilus influenzae
18	c3csiA_	Alignment		100.0	15	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase p; PDBTitle: crystal structure of the glutathione transferase pi allelic variant*c,2 i104v/a113v, in complex with the chlorambucil-glutathione conjugate
19	c3lszA_	Alignment		100.0	23	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of glutathione s-transferase from2 rhodobacter sphaeroides
20	c1ua5A_	Alignment		100.0	15	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: non-fusion gst from s. japonicum in complex with glutathione
21	c1zl9A_	Alignment	not modelled	100.0	16	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 5; PDBTitle: crystal structure of a major nematode c.elegans specific gst (ce01613)
22	c1k3yB_	Alignment	not modelled	100.0	16	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase a1; PDBTitle: crystal structure analysis of human glutathione s-transferase with s-2 hexyl glutathione and glycerol at 1.3 angstrom
23	c3qagA_	Alignment	not modelled	100.0	18	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase omega-2; PDBTitle: human glutathione transferase o2 with glutathione -new crystal form
24	c3isoB_	Alignment	not modelled	100.0	17	PDB header: transferase Chain: B: PDB Molecule: putative glutathione transferase; PDBTitle: crystal structure of 26 kda gst of clonorchis sinensis in p32212 symmetry
25	c1tu8A_	Alignment	not modelled	100.0	17	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 2; PDBTitle: structure of onchoverca volvulus pi-class glutathione s-2 transferase with its kompetitive inhibitor s-hexyl-gsh
26	c1jlvA_	Alignment	not modelled	100.0	21	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase gst1-3; PDBTitle: anopheles dirus species b glutathione s-transferases 1-3
27	c3m3mA_	Alignment	not modelled	100.0	22	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of glutathione s-transferase from pseudomonas2 fluorescens [pf-5]
28	c1yq1A_	Alignment	not modelled	100.0	18	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: structural genomics of caenorhabditis elegans: glutathione2 s-transferase
						PDB header: isomerase

29	c2jl4A_	Alignment	not modelled	100.0	21	Chain: A: PDB Molecule: maleyl pyruvate isomerase; PDBTitle: holo structure of maleyl pyruvate isomerase, a bacterial2 glutathione-s-transferase in zeta class PDB header: isomerase
30	c1iviA_	Alignment	not modelled	100.0	14	Chain: A: PDB Molecule: hematopoietic prostaglandin d synthase; PDBTitle: crystal structure of hematopoietic prostaglandin d synthase PDB header: transferase
31	c2gsqA_	Alignment	not modelled	100.0	18	Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: glutathione s-transferase from squid digestive gland complexed with s-2 (3-iodobenzyl)glutathione PDB header: transferase
32	c3h1nA_	Alignment	not modelled	100.0	18	Chain: A: PDB Molecule: probable glutathione s-transferase; PDBTitle: crystal structure of probable glutathione s-transferase from2 bordetella bronchiseptica rb50 PDB header: translation
33	c1nhyA_	Alignment	not modelled	100.0	16	Chain: A: PDB Molecule: elongation factor 1-gamma 1; PDBTitle: crystal structure of the gst-like domain of elongation2 factor 1-gamma from saccharomyces cerevisiae. PDB header: transferase
34	c3f6fA_	Alignment	not modelled	100.0	19	Chain: A: PDB Molecule: cg18548-pa (ip02196p) (ip02193p); PDBTitle: crystal structure of glutathione transferase dmgstd10 from2 drosophila melanogaster PDB header: transferase
35	c1m0uB_	Alignment	not modelled	100.0	14	Chain: B: PDB Molecule: gst2 gene product; PDBTitle: crystal structure of the drosophila glutathione s-2 transferase-2 in complex with glutathione PDB header: transcription
36	c1yy7A_	Alignment	not modelled	100.0	23	Chain: A: PDB Molecule: stringent starvation protein a; PDBTitle: crystal structure of stringent starvation protein a (sspa),2 an rna polymerase-associated transcription factor PDB header: transferase
37	c1gumA_	Alignment	not modelled	100.0	16	Chain: A: PDB Molecule: protein (glutathione transferase a4-4); PDBTitle: human glutathione transferase a4-4 without ligands PDB header: transferase
38	c1vf1A_	Alignment	not modelled	100.0	19	Chain: A: PDB Molecule: glutathione s-transferase 3; PDBTitle: cgsta1-1 in complex with glutathione PDB header: transcription
39	c3lypA_	Alignment	not modelled	100.0	23	Chain: A: PDB Molecule: stringent starvation protein a; PDBTitle: structure of stringent starvation protein a homolog from pseudomonas2 fluorescens PDB header: isomerase
40	c3nivD_	Alignment	not modelled	100.0	17	Chain: D: PDB Molecule: glutathione s-transferase; PDBTitle: the crystal structure of glutathione s-transferase from legionella2 pneumophila PDB header: transferase
41	c1oktA_	Alignment	not modelled	100.0	13	Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: x-ray structure of glutathione s-transferase from the2 malarial parasite plasmodium falciparum PDB header: transferase
42	c1gwcC_	Alignment	not modelled	100.0	23	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 PDB header: transferase
43	c1v2aD_	Alignment	not modelled	100.0	15	Chain: D: PDB Molecule: glutathione transferase gst1-6; PDBTitle: glutathione s-transferase 1-6 from anopheles dirus species b PDB header: transferase
44	c3lxzD_	Alignment	not modelled	100.0	23	Chain: D: PDB Molecule: glutathione s-transferase family protein; PDBTitle: structure of probable glutathione s-transferase(pp0183) from2 pseudomonas putida PDB header: transferase
45	c2vo4A_	Alignment	not modelled	100.0	20	Chain: A: PDB Molecule: 2,4-d inducible glutathione s-transferase; PDBTitle: glutathione transferase from glycine max PDB header: metal transport
46	c2aheA_	Alignment	not modelled	100.0	19	Chain: A: PDB Molecule: chloride intracellular channel protein 4; PDBTitle: crystal structure of a soluble form of clic4. intercellular2 chloride ion channel PDB header: transferase
47	c1r5aA_	Alignment	not modelled	100.0	21	Chain: A: PDB Molecule: glutathione transferase; PDBTitle: glutathione s-transferase PDB header: transferase
48	c1oyiC_	Alignment	not modelled	100.0	20	Chain: C: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure solution of rice gst1 (osgst1) in complex with2 glutathione. PDB header: transferase
49	c2hnlB_	Alignment	not modelled	100.0	15	Chain: B: PDB Molecule: glutathione s-transferase 1; PDBTitle: structure of the prostaglandin d synthase from the parasitic nematode2 onchocerca volvulus PDB header: transferase
50	c2fnoB_	Alignment	not modelled	100.0	15	Chain: B: PDB Molecule: agr_pat_752p; PDBTitle: crystal structure of a glutathione s-transferase (atu5508) from2 agrobacterium tumefaciens str. c58 at 2.00 a resolution PDB header: transferase
51	c2c3nB_	Alignment	not modelled	100.0	24	Chain: B: PDB Molecule: glutathione s-transferase theta 1; PDBTitle: human glutathione-s-transferase t1-1, apo form PDB header: transferase
52	c2on7A_	Alignment	not modelled	100.0	17	Chain: A: PDB Molecule: na glutathione s-transferase 1; PDBTitle: structure of nagst-1 PDB header: transferase
53	c1gnwA_	Alignment	not modelled	100.0	23	Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: structure of glutathione s-transferase PDB header: transferase
54	c1eemA_	Alignment	not modelled	100.0	18	Chain: A: PDB Molecule: glutathione-s-transferase; PDBTitle: glutathione transferase from homo sapiens PDB header: transferase
55	c1hvaA_	Alignment	not modelled	100.0	23	Chain: A: PDB Molecule: protein (glutathione s-transferase); PDBTitle: glutathione s-transferase PDB header: transferase

55	c10yeA	Alignment	not modelled	100.0	22	PDBTitle: glutathione s-transferase i from mais in complex with2 atrazine glutathione conjugate
56	c1jlwA	Alignment	not modelled	100.0	22	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase gst1-4; PDBTitle: anopheles dirus species b glutathione s-transferases 1-4
57	c2on5E	Alignment	not modelled	100.0	17	PDB header: transferase Chain: E: PDB Molecule: na glutathione s-transferase 2; PDBTitle: structure of nagst-2
58	c1bg5A	Alignment	not modelled	100.0	14	PDB header: ankyrin binding Chain: A: PDB Molecule: fusion protein of alpha-na,k-tpase with PDBTitle: crystal structure of the ankyrin binding domain of alpha-na,2 k-tpase as a fusion protein with glutathione s-transferase
59	c1k0dB	Alignment	not modelled	100.0	22	PDB header: gene regulation Chain: B: PDB Molecule: ure2 protein; PDBTitle: ure2p in complex with glutathione
60	c1g7oA	Alignment	not modelled	100.0	17	PDB header: oxidoreductase Chain: A: PDB Molecule: glutaredoxin 2; PDBTitle: nmr solution structure of reduced e. coli glutaredoxin 2
61	c2cz2A	Alignment	not modelled	100.0	19	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)
62	c1b48A	Alignment	not modelled	100.0	19	PDB header: transferase Chain: A: PDB Molecule: protein (glutathione s-transferase); PDBTitle: crystal structure of mgsta4-4 in complex with gsh conjugate2 of 4-hydroxynonenal in one subunit and gsh in the other:3 evidence of signaling across dimer interface in mgsta4-4
63	c3lg6B	Alignment	not modelled	100.0	18	PDB header: transferase Chain: B: PDB Molecule: putative glutathione transferase; PDBTitle: crystal structure of putative glutathione transferase from2 coccidioides immitis
64	c3lq7B	Alignment	not modelled	100.0	19	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of glutathione s-transferase from2 agrobacterium tumefaciens str. c58
65	c1aw9A	Alignment	not modelled	100.0	27	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase iii; PDBTitle: structure of glutathione s-transferase iii in apo form
66	c3bbyA	Alignment	not modelled	100.0	19	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
67	c2wb9A	Alignment	not modelled	100.0	18	PDB header: transferase Chain: A: PDB Molecule: glutathione transferase sigma class; PDBTitle: fasciola hepatica sigma class gst
68	c3ay8A	Alignment	not modelled	100.0	17	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: glutathione s-transferase unclassified 2 from bombyx mori
69	c2imiA	Alignment	not modelled	100.0	12	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
70	c3m8nA	Alignment	not modelled	100.0	20	PDB header: transferase Chain: A: PDB Molecule: possible glutathione s-transferase; PDBTitle: crystal structure of a possible glutathione s-tranferase from2 rhodopseudomonas palustris
71	c1k0nB	Alignment	not modelled	100.0	18	PDB header: metal transport Chain: B: PDB Molecule: chloride intracellular channel protein 1; PDBTitle: chloride intracellular channel 1 (cllc1) complexed with glutathione
72	c2ws2B	Alignment	not modelled	100.0	20	PDB header: transferase Chain: B: PDB Molecule: glutathione s-transferase; PDBTitle: the 2 angstrom structure of a nu-class gst from haemonchus contortus
73	c3cbuB	Alignment	not modelled	100.0	19	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
74	c3rbtD	Alignment	not modelled	100.0	15	PDB header: transferase Chain: D: PDB Molecule: glutathione transferase o1; PDBTitle: crystal structure of glutathione s-transferase omega 3 from the2 silkworm bombyx mori
75	c2c8uA	Alignment	not modelled	100.0	17	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase 28 kda; PDBTitle: structure of r21q mutant of sh28gst
76	c3touB	Alignment	not modelled	100.0	21	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
77	c1tw9C	Alignment	not modelled	100.0	17	PDB header: transferase Chain: C: PDB Molecule: glutathione s-transferase 2; PDBTitle: glutathione transferase-2, apo form, from the nematode heligmosomoides2 polygyrus
78	c2r4vA	Alignment	not modelled	100.0	17	PDB header: transport protein Chain: A: PDB Molecule: chloride intracellular channel protein 2; PDBTitle: structure of human clic2, crystal form a
79	c3ic8D	Alignment	not modelled	100.0	10	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
80	c3ppuB	Alignment	not modelled	100.0	19	PDB header: transferase Chain: B: PDB Molecule: glutathione-s-transferase; PDBTitle: crystal structure of the glutathione-s-transferase xi from2 phanerochaete chrysosporium
						PDB header: transport protein

81	c3fy7B_	Alignment	not modelled	100.0	19	Chain: B: PDB Molecule: chloride intracellular channel protein 3; PDBTitle: crystal structure of homo sapiens clic3
82	c2yv9B_	Alignment	not modelled	100.0	12	PDB header: metal transport Chain: B: PDB Molecule: chloride intracellular channel exc-4; PDBTitle: crystal structure of the clic homologue exc-4 from c.2 elegans
83	c2yv7A_	Alignment	not modelled	100.0	18	PDB header: metal transport Chain: A: PDB Molecule: cg10997-pa; PDBTitle: crystal structure of the clic homolog from drosophila2 melanogaster
84	c1e6bA_	Alignment	not modelled	100.0	23	PDB header: transferase Chain: A: PDB Molecule: glutathione s-transferase; PDBTitle: crystal structure of a zeta class glutathione s-transferase2 from arabidopsis thaliana
85	c3m1gC_	Alignment	not modelled	99.9	18	PDB header: transferase Chain: C: PDB Molecule: putative glutathione s-transferase; PDBTitle: the structure of a putative glutathione s-transferase from2 corynebacterium glutamicum
86	c1z9hB_	Alignment	not modelled	99.9	22	PDB header: isomerase Chain: B: PDB Molecule: membrane-associated prostaglandin e synthase-2; PDBTitle: microsomal prostaglandin e synthase type-2
87	c2uz8A_	Alignment	not modelled	99.9	20	PDB header: rna-binding protein Chain: A: PDB Molecule: eukaryotic translation elongation factor 1 PDBTitle: the crystal structure of p18, human translation elongation2 factor 1 epsilon 1
88	d1n2aa1	Alignment	not modelled	99.9	100	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
89	d1f2ea1	Alignment	not modelled	99.9	32	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
90	d1pmta1	Alignment	not modelled	99.8	48	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
91	d1n2aa2	Alignment	not modelled	99.8	100	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
92	d1f2ea2	Alignment	not modelled	99.8	58	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
93	d1pmta2	Alignment	not modelled	99.8	63	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
94	c2hsmA_	Alignment	not modelled	99.8	14	PDB header: ligase/rna binding protein Chain: A: PDB Molecule: glutamyl-trna synthetase, cytoplasmic; PDBTitle: structural basis of yeast aminoacyl-trna synthetase complex2 formation revealed by crystal structures of two binary sub-3 complexes
95	d1jlwa2	Alignment	not modelled	99.8	28	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
96	d1v2aa2	Alignment	not modelled	99.8	21	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
97	d1pn9a2	Alignment	not modelled	99.7	30	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
98	d1axda1	Alignment	not modelled	99.7	22	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
99	d3gtub1	Alignment	not modelled	99.7	16	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
100	d1aw9a1	Alignment	not modelled	99.7	23	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
101	d2c4ja1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
102	d1b4pa1	Alignment	not modelled	99.7	17	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
103	d1xw6a1	Alignment	not modelled	99.7	17	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
104	d4gtua1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
105	d1gsua1	Alignment	not modelled	99.7	19	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
106	d2fhea1	Alignment	not modelled	99.7	14	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
107	d2gsta1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
						Fold: GST C-terminal domain-like

108	d1jlva1	Alignment	not modelled	99.7	16	Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
109	d1ljra2	Alignment	not modelled	99.7	29	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
110	d1nhya1	Alignment	not modelled	99.7	11	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
111	d1e6ba2	Alignment	not modelled	99.7	29	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
112	d1jlva2	Alignment	not modelled	99.7	29	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
113	d1duga1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
114	d1gnea1	Alignment	not modelled	99.7	14	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
115	d2gsra1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
116	d1lbka1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
117	d1gnwa2	Alignment	not modelled	99.7	32	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione S-transferase (GST), N-terminal domain
118	d2a2ra1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
119	d1gnwa1	Alignment	not modelled	99.7	15	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain
120	d1fwla1	Alignment	not modelled	99.7	16	Fold: GST C-terminal domain-like Superfamily: GST C-terminal domain-like Family: Glutathione S-transferase (GST), C-terminal domain