


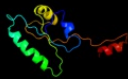
























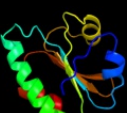




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2hfda1	 Alignment		100.0	100	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: HyaE-like
2	c2qsiB_	 Alignment		99.9	28	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: putative hydrogenase expression/formation protein hupg; PDBTitle: crystal structure of putative hydrogenase expression/formation protein2 hupg from rhodopseudomonas palustris cga009
3	d2es7a1	 Alignment		99.9	74	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: HyaE-like
4	c3p2aA_	 Alignment		99.8	17	PDB header: oxidoreductase Chain: A: PDB Molecule: putative thioredoxin-like protein; PDBTitle: crystal structure of thioredoxin 2 from yersinia pestis
5	c3qdnA_	 Alignment		99.8	19	PDB header: oxidoreductase Chain: A: PDB Molecule: putative thioredoxin protein; PDBTitle: putative thioredoxin protein from salmonella typhimurium
6	c3p2aB_	 Alignment		99.8	17	PDB header: oxidoreductase Chain: B: PDB Molecule: putative thioredoxin-like protein; PDBTitle: crystal structure of thioredoxin 2 from yersinia pestis
7	c2pptA_	 Alignment		99.8	16	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin-2; PDBTitle: crystal structure of thioredoxin-2
8	c3gnjD_	 Alignment		99.8	17	PDB header: structural genomics, unknown function Chain: D: PDB Molecule: thioredoxin domain protein; PDBTitle: the crystal structure of a thioredoxin-related protein from2 desulfitobacterium hafniense dcb
9	c3dxbE_	 Alignment		99.8	13	PDB header: splicing, transcription Chain: E: PDB Molecule: thioredoxin n-terminally fused to puf60(uhm); PDBTitle: structure of the uhm domain of puf60 fused to thioredoxin
10	d1thxa_	 Alignment		99.8	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
11	d1qgva_	 Alignment		99.8	8	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: spliceosomal protein U5-15Kd

12	c1xbsA_	Alignment		99.8	17	PDB header: transcription, cell cycle Chain: A: PDB Molecule: dim1-like protein; PDBTitle: crystal structure of human dim2: a dim1-like protein
13	c2l6dA_	Alignment		99.8	16	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin; PDBTitle: solution structure of desulfothioredoxin from desulfovibrio vulgaris2 hildenborough in its reduced form
14	d1dbya_	Alignment		99.8	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
15	c2r2jA_	Alignment		99.8	16	PDB header: chaperone Chain: A: PDB Molecule: thioredoxin domain-containing protein 4; PDBTitle: crystal structure of human erp44
16	c1a8yA_	Alignment		99.8	14	PDB header: calcium-binding protein Chain: A: PDB Molecule: calsequestrin; PDBTitle: crystal structure of calsequestrin from rabbit skeletal muscle2 sarcoplasmic reticulum at 2.4 a resolution
17	c3tcoA_	Alignment		99.8	14	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin (trxa-1); PDBTitle: crystallographic and spectroscopic characterization of sulfobobus2 solfataricus trxa1 provide insights into the determinants of3 thioredoxin fold stability
18	c3hz4A_	Alignment		99.8	13	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin; PDBTitle: crystal structure of thioredoxin from methanosarcina mazei
19	c2b5eA_	Alignment		99.8	8	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase; PDBTitle: crystal structure of yeast protein disulfide isomerase
20	d1fb6a_	Alignment		99.8	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
21	d2trxa_	Alignment	not modelled	99.8	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
22	d1gh2a_	Alignment	not modelled	99.8	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
23	c1t00A_	Alignment	not modelled	99.8	13	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: the structure of thioredoxin from s. coelicolor
24	c2e0qA_	Alignment	not modelled	99.8	18	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: crystal structure of k53e thioredoxin from sulfobobus2 tokodaii strain7
25	d1a8ya1	Alignment	not modelled	99.8	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Calsequestrin
26	c3hypB_	Alignment	not modelled	99.7	11	PDB header: electron transport Chain: B: PDB Molecule: thioredoxin; PDBTitle: crystal structure of bacteroides fragilis trxp_s105g mutant
27	c2diyA_	Alignment	not modelled	99.7	15	PDB header: protein binding Chain: A: PDB Molecule: thioredoxin-like protein 2; PDBTitle: the solution structure of the thioredoxin domain of human2 thioredoxin-like protein 2
28	c2albA_	Alignment	not modelled	99.7	11	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a3; PDBTitle: nmr structure of the n-terminal domain a of the2 glycoprotein chaperone erp57
						PDB header: oxidoreductase

29	c2i4aA	Alignment	not modelled	99.7	18	Chain: A: PDB Molecule: thioredoxin; PDBTitle: crystal structure of thioredoxin from the acidophile2 acetobacter aceti
30	c2wz9A	Alignment	not modelled	99.7	15	PDB header: protein binding Chain: A: PDB Molecule: glutaredoxin-3; PDBTitle: crystal structure of the thioredoxin domain of human txn2
31	d1f9ma	Alignment	not modelled	99.7	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
32	c2yzuA	Alignment	not modelled	99.7	20	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: crystal structure of oxidized thioredoxin from thermus thermophilus2 hb8
33	c2i1uA	Alignment	not modelled	99.7	15	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: mycobacterium tuberculosis thioredoxin c
34	d1a8la2	Alignment	not modelled	99.7	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
35	d1xwaa	Alignment	not modelled	99.7	19	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
36	c2vocA	Alignment	not modelled	99.7	19	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: thioredoxin a active site mutants form mixed disulfide2 dimers that resemble enzyme-substrate reaction3 intermediate
37	c3f8uA	Alignment	not modelled	99.7	11	PDB header: immune system/isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a3erp57; PDBTitle: tapasin/erp57 heterodimer
38	c1w89E	Alignment	not modelled	99.7	13	PDB header: electron transport Chain: E: PDB Molecule: thioredoxin; PDBTitle: structure of the reduced form of human thioredoxin 2
39	c2dj0A	Alignment	not modelled	99.7	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: thioredoxin-related transmembrane protein 2; PDBTitle: the solution structure of the thioredoxin domain of human2 thioredoxin-related transmembrane protein 2
40	d1nw2a	Alignment	not modelled	99.7	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
41	c2vimA	Alignment	not modelled	99.7	13	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin; PDBTitle: x-ray structure of fasciola hepatica thioredoxin
42	c2dj1A	Alignment	not modelled	99.7	13	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a4; PDBTitle: the solution structure of the first thioredoxin domain of2 mouse protein disulfide-isomerase a4
43	c1x5dA	Alignment	not modelled	99.7	12	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a6; PDBTitle: the solution structure of the second thioredoxin-like2 domain of human protein disulfide-isomerase a6
44	c2o7kA	Alignment	not modelled	99.7	14	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin; PDBTitle: s. aureus thioredoxin
45	c2xbqB	Alignment	not modelled	99.7	13	PDB header: oxidoreductase Chain: B: PDB Molecule: thioredoxin; PDBTitle: crystal structure of reduced schistosoma mansoni2 thioredoxin pre-protein at 1.7 angstrom
46	d2ifqa1	Alignment	not modelled	99.7	16	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
47	c2l5lA	Alignment	not modelled	99.7	12	PDB header: transport protein Chain: A: PDB Molecule: thioredoxin; PDBTitle: solution structure of thioredoxin from bacteroides vulgatus
48	c2vafA	Alignment	not modelled	99.7	11	PDB header: metal-binding protein Chain: A: PDB Molecule: calsequestrin-2; PDBTitle: crystal structure of human cardiac calsequestrin
49	c1v98A	Alignment	not modelled	99.7	17	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin; PDBTitle: crystal structure analysis of thioredoxin from thermus thermophilus
50	d1r26a	Alignment	not modelled	99.7	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
51	d1j08a2	Alignment	not modelled	99.7	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
52	c3emxB	Alignment	not modelled	99.7	16	PDB header: oxidoreductase Chain: B: PDB Molecule: thioredoxin; PDBTitle: crystal structure of thioredoxin from aeropyrum pernix
53	d2b5ea4	Alignment	not modelled	99.7	7	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
54	c3ed3A	Alignment	not modelled	99.7	11	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase mpd1; PDBTitle: crystal structure of the yeast dithiol/disulfide2 oxidoreductase mpd1p
55	d1m7ta	Alignment	not modelled	99.7	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase

56	c2j23A_	Alignment	not modelled	99.7	10	PDB header: immune protein Chain: A: PDB Molecule: thioredoxin; PDBTitle: cross-reactivity and crystal structure of malassezia2 sympodialis thioredoxin (mala s 13), a member of a new pan-3 allergen family
57	d1syra_	Alignment	not modelled	99.7	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
58	c1sjjA_	Alignment	not modelled	99.7	10	PDB header: metal binding protein Chain: A: PDB Molecule: calsequestrin, cardiac muscle isoform; PDBTitle: comparing skeletal and cardiac calsequestrin structures and2 their calcium binding: a proposed mechanism for coupled3 calcium binding and protein polymerization
59	c2dmlA_	Alignment	not modelled	99.7	12	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a6; PDBTitle: the solution structure of the first thioredoxin domain of2 mouse protein disulfide-isomerase a6
60	c3apsA_	Alignment	not modelled	99.7	10	PDB header: oxidoreductase Chain: A: PDB Molecule: dnaj homolog subfamily c member 10; PDBTitle: crystal structure of trx4 domain of erdj5
61	c2oe0B_	Alignment	not modelled	99.7	16	PDB header: electron transport Chain: B: PDB Molecule: thioredoxin-3; PDBTitle: crystal structure of mitochondrial thioredoxin 3 from2 saccharomyces cerevisiae
62	c3d22A_	Alignment	not modelled	99.7	15	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin h-type; PDBTitle: crystal structure of a poplar thioredoxin h mutant,2 ptxrh4c61s
63	d1ep7a_	Alignment	not modelled	99.7	16	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
64	d1xfla_	Alignment	not modelled	99.7	10	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
65	c2v1vA_	Alignment	not modelled	99.7	16	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin h isoform 2.; PDBTitle: crystal structure of barley thioredoxin h isoform 2 in2 partially radiation-reduced state
66	c2dizA_	Alignment	not modelled	99.7	15	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin domain-containing protein 5; PDBTitle: the solution structure of the third thioredoxin domain of2 human thioredoxin domain-containing protein 5
67	d1ti3a_	Alignment	not modelled	99.7	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
68	c2vm2C_	Alignment	not modelled	99.7	13	PDB header: oxidoreductase Chain: C: PDB Molecule: thioredoxin h isoform 1.; PDBTitle: crystal structure of barley thioredoxin h isoform 12 crystallized using peg as precipitant
69	d1a8la1	Alignment	not modelled	99.7	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
70	d1oaza_	Alignment	not modelled	99.6	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
71	d1g7ea_	Alignment	not modelled	99.6	10	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: ERP29 N domain-like
72	d1j08a1	Alignment	not modelled	99.6	14	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
73	c3apoA_	Alignment	not modelled	99.6	12	PDB header: oxidoreductase Chain: A: PDB Molecule: dnaj homolog subfamily c member 10; PDBTitle: crystal structure of full-length erdj5
74	c1j08A_	Alignment	not modelled	99.6	12	PDB header: isomerase Chain: A: PDB Molecule: glutaredoxin-like protein; PDBTitle: crystal structure of glutaredoxin-like protein from2 pyrococcus horikoshii
75	c3ul3A_	Alignment	not modelled	99.6	15	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin; PDBTitle: structural insights into thioredoxin-2: a component of malaria2 parasite protein secretion machinery
76	d1mekA_	Alignment	not modelled	99.6	11	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
77	c1x5eA_	Alignment	not modelled	99.6	13	PDB header: electron transport Chain: A: PDB Molecule: thioredoxin domain containing protein 1; PDBTitle: the solution structure of the thioredoxin-like domain of2 human thioredoxin-related transmembrane protein
78	c2dj2A_	Alignment	not modelled	99.6	15	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a4; PDBTitle: the solution structure of the second thioredoxin domain of2 mouse protein disulfide-isomerase a4
79	c2qc7A_	Alignment	not modelled	99.6	11	PDB header: chaperone Chain: A: PDB Molecule: endoplasmic reticulum protein erp29; PDBTitle: crystal structure of the protein-disulfide isomerase related chaperone2 erp29
80	d1zmaa1	Alignment	not modelled	99.6	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
81	c3idvA_	Alignment	not modelled	99.6	14	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a4; PDBTitle: crystal structure of the a0a fragment of erp72
						PDB header: transferase

82	c3g9bA	Alignment	not modelled	99.6	8	Chain: A: PDB Molecule: dolichyl-diphosphooligosaccharide-protein PDBTitle: crystal structure of reduced ost6l
83	c2l57A	Alignment	not modelled	99.6	11	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein; PDBTitle: solution structure of an uncharacterized thioredoxin-like protein from2 clostridium perfringens
84	c2i9hA	Alignment	not modelled	99.6	17	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin i; PDBTitle: nmr solution structure of the reduced form of thioredoxin 12 from yeast (trx1)
85	c1ovnA	Alignment	not modelled	99.6	18	PDB header: chaperone Chain: A: PDB Molecule: windbeutel; PDBTitle: crystal structure and functional analysis of drosophila2 wind-- a pdi-related protein
86	c3apqB	Alignment	not modelled	99.6	14	PDB header: oxidoreductase Chain: B: PDB Molecule: dnaj homolog subfamily c member 10; PDBTitle: crystal structure of j-trx1 fragment of erdj5
87	c3d6iB	Alignment	not modelled	99.6	14	PDB header: oxidoreductase Chain: B: PDB Molecule: monothiol glutaredoxin-3; PDBTitle: structure of the thioredoxin-like domain of yeast2 glutaredoxin 3
88	c3cxgA	Alignment	not modelled	99.6	9	PDB header: oxidoreductase Chain: A: PDB Molecule: putative thioredoxin; PDBTitle: crystal structure of plasmodium falciparum thioredoxin, pfi0790w
89	c2f51B	Alignment	not modelled	99.6	13	PDB header: electron transport Chain: B: PDB Molecule: thioredoxin; PDBTitle: structure of trichomonas vaginalis thioredoxin
90	c3h79A	Alignment	not modelled	99.5	14	PDB header: unknown function Chain: A: PDB Molecule: thioredoxin-like protein; PDBTitle: crystal structure of trypanosoma cruzi thioredoxin-like hypothetical2 protein q4dv70
91	c2aytB	Alignment	not modelled	99.5	14	PDB header: oxidoreductase Chain: B: PDB Molecule: glutaredoxin-like protein; PDBTitle: the crystal structure of a protein disulfide oxidoreductase from2 aquifex aeolicus
92	c1zypB	Alignment	not modelled	99.5	13	PDB header: oxidoreductase Chain: B: PDB Molecule: alkyl hydroperoxide reductase subunit f; PDBTitle: synchrotron reduced form of the n-terminal domain of2 salmonella typhimurium ahpf
93	d2b5ea1	Alignment	not modelled	99.5	17	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
94	c2hlsB	Alignment	not modelled	99.5	13	PDB header: oxidoreductase Chain: B: PDB Molecule: protein disulfide oxidoreductase; PDBTitle: the crystal structure of a protein disulfide oxidoreductase from2 aeropyrum pernix k1
95	c2dj3A	Alignment	not modelled	99.5	13	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a4; PDBTitle: the solution structure of the third thioredoxin domain of2 mouse protein disulfide-isomerase a4
96	d1v9wa	Alignment	not modelled	99.5	7	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Txn15-like
97	d1fo5a	Alignment	not modelled	99.5	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
98	d1woua	Alignment	not modelled	99.5	8	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Txn15-like
99	d2c0ga2	Alignment	not modelled	99.5	17	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: ERP29 N domain-like
100	c1x5cA	Alignment	not modelled	99.5	12	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase; PDBTitle: the solution structure of the second thioredoxin-like2 domain of human protein disulfide-isomerase
101	c2dbcA	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: unnamed protein product; PDBTitle: solution structure of the thioredoxin-like domain of2 phosducin-like protein 2(pdcl2)
102	c2fgxA	Alignment	not modelled	99.5	8	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.
103	d2fwha1	Alignment	not modelled	99.5	16	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
104	c3dmlA	Alignment	not modelled	99.4	10	PDB header: oxidoreductase Chain: A: PDB Molecule: putative uncharacterized protein; PDBTitle: crystal structure of the periplasmic thioredoxin soxs from2 paracoccus pantotrophus (reduced form)
105	d1nhoa	Alignment	not modelled	99.4	18	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
106	d1hyua4	Alignment	not modelled	99.4	13	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
107	c2dmmA	Alignment	not modelled	99.4	12	PDB header: isomerase Chain: A: PDB Molecule: protein disulfide-isomerase a3; PDBTitle: the solution structure of the second thioredoxin domain of2 human protein disulfide-isomerase a3
108	c1h9vC	Alignment	not modelled	99.4	15	PDB header: signaling protein Chain: C: PDB Molecule: protein (phosducin);

108	c109xc_	Alignment	not modelled	99.4	13	PDBTitle: structural analysis of phosducin and its phosphorylation-2 regulated interaction with transducin
109	c3uemA_	Alignment	not modelled	99.4	14	PDB header: chaperone Chain: A: PDB Molecule: protein disulfide-isomerase; PDBTitle: crystal structure of human pdi bb'a' domains
110	c2kucA_	Alignment	not modelled	99.4	14	PDB header: isomerase Chain: A: PDB Molecule: putative disulphide-isomerase; PDBTitle: solution structure of a putative disulphide-isomerase from bacteroides2 thetaiotaomicron
111	c3kp8A_	Alignment	not modelled	99.4	16	PDB header: oxidoreductase Chain: A: PDB Molecule: vkorc1/thioredoxin domain protein; PDBTitle: the thioredoxin-like domain of a vkor homolog from2 synechococcus sp.
112	d1a0rp_	Alignment	not modelled	99.3	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Phosducin
113	d2djja1	Alignment	not modelled	99.3	12	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: PDI-like
114	d2trcp_	Alignment	not modelled	99.3	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Phosducin
115	c2hyxA_	Alignment	not modelled	99.2	18	PDB header: unknown function Chain: A: PDB Molecule: protein dipz; PDBTitle: structure of the c-terminal domain of dipz from mycobacterium2 tuberculosis
116	c2ju5A_	Alignment	not modelled	99.1	16	PDB header: oxidoreductase Chain: A: PDB Molecule: thioredoxin disulfide isomerase; PDBTitle: dsbh oxidoreductase
117	c3f9uA_	Alignment	not modelled	99.1	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative exported cytochrome c biogenesis-related protein; PDBTitle: crystal structure of c-terminal domain of putative exported cytochrome2 c biogenesis-related protein from bacteroides fragilis
118	d1z5ye1	Alignment	not modelled	99.1	19	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione peroxidase-like
119	d1wjka_	Alignment	not modelled	98.9	15	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Thioltransferase
120	d1knga_	Alignment	not modelled	98.9	19	Fold: Thioredoxin fold Superfamily: Thioredoxin-like Family: Glutathione peroxidase-like