

					Family: Ferredoxin domains from multidomain proteins
105	c1y56A_	Alignment	not modelled	97.4	PDB header: oxidoreductase Chain: A: PDB Molecule: hypothetical protein ph1363; PDBTitle: crystal structure of l-proline dehydrogenase from p.horikoshii
106	d1l5pa_	Alignment	not modelled	97.4	Fold: beta-Grasp (ubiquitin-like) Superfamily: 2Fe-2S ferredoxin-like Family: 2Fe-2S ferredoxin-related
107	c2piaA_	Alignment	not modelled	97.4	PDB header: reductase Chain: A: PDB Molecule: phthalate dioxygenase reductase; PDBTitle: phthalate dioxygenase reductase: a modular structure for electron2 transfer from pyridine nucleotides to [2fe-2s]
108	c4p6vF_	Alignment	not modelled	97.4	PDB header: oxidoreductase Chain: F: PDB Molecule: na(+) -translocating nadh-quinone reductase subunit f; PDBTitle: crystal structure of the na+ -translocating nadh: ubiquinone2 oxidoreductase from vibrio cholerae
109	d1blua_	Alignment	not modelled	97.3	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
110	d2gma3	Alignment	not modelled	97.3	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: ETF-QO domain-like
111	d2bt6a1	Alignment	not modelled	97.3	Fold: beta-Grasp (ubiquitin-like) Superfamily: 2Fe-2S ferredoxin-like Family: 2Fe-2S ferredoxin-related
112	c2ivfB_	Alignment	not modelled	97.3	PDB header: oxidoreductase Chain: B: PDB Molecule: ethylbenzene dehydrogenase beta-subunit; PDBTitle: ethylbenzene dehydrogenase from aromatoleum aromaticum
113	d7fd1a_	Alignment	not modelled	97.2	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
114	c6f0kB_	Alignment	not modelled	97.2	PDB header: membrane protein Chain: B: PDB Molecule: fe-s-cluster-containing hydrogenase; PDBTitle: alternative complex iii
115	c6btmB_	Alignment	not modelled	97.2	PDB header: membrane protein Chain: B: PDB Molecule: alternative complex iii subunit b; PDBTitle: structure of alternative complex iii from flavobacterium johnsoniae2 (wild type)
116	c3lxvfC_	Alignment	not modelled	97.1	PDB header: metal binding protein Chain: C: PDB Molecule: ferredoxin; PDBTitle: crystal structure of [2fe-2s] ferredoxin arx from novosphingobium2 aromaticivorans
117	d1clfa_	Alignment	not modelled	97.1	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
118	d1bc6a_	Alignment	not modelled	97.1	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
119	d2fdna_	Alignment	not modelled	97.1	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
120	c4ylfD_	Alignment	not modelled	97.1	PDB header: oxidoreductase Chain: D: PDB Molecule: dihydropyrimidine dehydrogenase subunit a; PDBTitle: insights into flavin-based electron bifurcation via the nadh-dependent2 reduced ferredoxin-nadp oxidoreductase structure