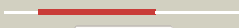
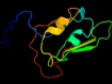
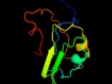
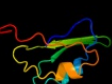
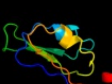
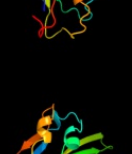
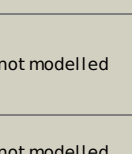
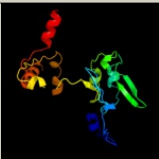


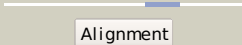
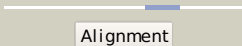
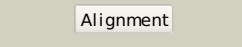
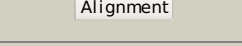
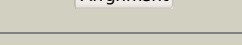
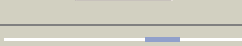
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2fug91	 Alignment		99.9	38	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
2	c2fugG_	 Alignment		99.9	38	PDB header: oxidoreductase Chain: G: PDB Molecule: nadh-quinone oxidoreductase chain 9; PDBTitle: crystal structure of the hydrophilic domain of respiratory complex i2 from thermus thermophilus
3	dlxera_	 Alignment		99.6	27	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Archaeal ferredoxins
4	c1gthD_	 Alignment		99.6	21	PDB header: oxidoreductase Chain: D: PDB Molecule: dihydropyrimidine dehydrogenase; PDBTitle: dihydropyrimidine dehydrogenase (dpd) from pig, ternary2 complex with nadph and 5-iodouracil
5	dlgtea5	 Alignment		99.6	20	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
6	c2gmhA_	 Alignment		99.6	20	PDB header: oxidoreductase Chain: A: PDB Molecule: electron transfer flavoprotein-ubiquinone PDBTitle: structure of porcine electron transfer flavoprotein-2 ubiquinone oxidoreductase in complexed with ubiquinone
7	c3gyxl_	 Alignment		99.5	28	PDB header: oxidoreductase Chain: J: PDB Molecule: adenylsulfate reductase; PDBTitle: crystal structure of adenylylsulfate reductase from2 desulfovibrio gigas
8	d2fug34	 Alignment		99.5	16	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
9	dljb0c_	 Alignment		99.5	26	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
10	c1kqfB_	 Alignment		99.5	32	PDB header: oxidoreductase Chain: B: PDB Molecule: formate dehydrogenase, nitrate-inducible, iron-sulfur PDBTitle: formate dehydrogenase n from e. coli
11	c1hfeL_	 Alignment		99.5	18	PDB header: hydrogenase Chain: L: PDB Molecule: protein (fe-only hydrogenase (e.c.1.18.99.1) PDBTitle: 1.6 a resolution structure of the fe-only hydrogenase from2 desulfovibrio desulfuricans

12	d2gmha3	Alignment		99.5	18	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: ETF-QO domain-like
13	c2c3yA_	Alignment		99.5	30	PDB header: oxidoreductase Chain: A: PDB Molecule: pyruvate-ferredoxin oxidoreductase; PDBTitle: crystal structure of the radical form of2 pyruvate:ferredoxin oxidoreductase from desulfovibrio3 africanus
14	d1hfel2	Alignment		99.5	25	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
15	d7fd1a_	Alignment		99.5	27	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
16	d2c42a5	Alignment		99.5	26	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
17	d1h98a_	Alignment		99.4	26	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
18	d1jnrb_	Alignment		99.4	25	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
19	d1bc6a_	Alignment		99.4	34	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: 7-Fe ferredoxin
20	d1rgva_	Alignment		99.4	24	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
21	d1kqfb1	Alignment	not modelled	99.4	32	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
22	c2zvsB_	Alignment	not modelled	99.4	22	PDB header: electron transport Chain: B: PDB Molecule: uncharacterized ferredoxin-like protein yfhI; PDBTitle: crystal structure of the 2[4fe-4s] ferredoxin from escherichia coli
23	d2fdna_	Alignment	not modelled	99.4	40	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
24	c2vpyB_	Alignment	not modelled	99.4	23	PDB header: oxidoreductase Chain: B: PDB Molecule: nrfc protein; PDBTitle: polysulfide reductase with bound quinone inhibitor,2 pentachlorophenol (pcp)
25	d1dura_	Alignment	not modelled	99.4	36	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
26	d3c8ya3	Alignment	not modelled	99.4	22	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
27	d1blua_	Alignment	not modelled	99.4	22	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
28	c1gx7A_	Alignment	not modelled	99.4	27	PDB header: oxidoreductase Chain: A: PDB Molecule: periplasmic [fe] hydrogenase large subunit; PDBTitle: best model of the electron transfer complex between2 cytochrome c3 and [fe]-hydrogenase

29	c1ti2F_	Alignment	not modelled	99.4	14	PDB header: oxidoreductase Chain: F: PDB Molecule: pyrogallol hydroxytransferase small subunit; PDBTitle: crystal structure of pyrogallol-phloroglucinol2 transhydroxylase from pelobacter acidigallici
30	d1vjwa_	Alignment	not modelled	99.4	22	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Single 4Fe-4S cluster ferredoxin
31	c1c4cA_	Alignment		99.3	16	PDB header: oxidoreductase Chain: A: PDB Molecule: protein (fe-only hydrogenase); PDBTitle: binding of exogenously added carbon monoxide at the active2 site of the fe-only hydrogenase (cpi) from clostridium3 pasteurianum
32	d1fcaa_	Alignment	not modelled	99.3	36	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
33	d1h0hb_	Alignment	not modelled	99.3	25	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
34	c2fgoA_	Alignment	not modelled	99.3	26	PDB header: electron transport Chain: A: PDB Molecule: ferredoxin; PDBTitle: structure of the 2[4fe-4s] ferredoxin from pseudomonas2 aeruginosa
35	d1vlfm2	Alignment	not modelled	99.3	15	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
36	c2fugC_	Alignment	not modelled	99.3	14	PDB header: oxidoreductase Chain: C: PDB Molecule: nadh-quinone oxidoreductase chain 3; PDBTitle: crystal structure of the hydrophilic domain of respiratory complex i2 from thermus thermophilus
37	d1y5b1	Alignment	not modelled	99.3	24	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
38	d1clfa_	Alignment	not modelled	99.3	36	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Short-chain ferredoxins
39	c2ivfB_	Alignment	not modelled	99.3	28	PDB header: oxidoreductase Chain: B: PDB Molecule: ethylbenzene dehydrogenase beta-subunit; PDBTitle: ethylbenzene dehydrogenase from aromatoleum aromaticum
40	d1iqza_	Alignment	not modelled	99.3	23	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Single 4Fe-4S cluster ferredoxin
41	d1sj1a_	Alignment	not modelled	99.2	21	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Single 4Fe-4S cluster ferredoxin
42	c2v2kB_	Alignment	not modelled	99.2	25	PDB header: electron transport Chain: B: PDB Molecule: ferredoxin; PDBTitle: the crystal structure of fdxa, a 7fe ferredoxin from2 mycobacterium smegmatis
43	c3c7bE_	Alignment	not modelled	99.1	26	PDB header: oxidoreductase Chain: E: PDB Molecule: sulfite reductase, dissimilatory-type subunit beta; PDBTitle: structure of the dissimilatory sulfite reductase from archaeoglobus2 fulgidus
44	d1fxra_	Alignment	not modelled	99.1	16	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Single 4Fe-4S cluster ferredoxin
45	c1dw1A_	Alignment	not modelled	99.1	19	PDB header: electron transfer Chain: A: PDB Molecule: ferredoxin i; PDBTitle: the ferredoxin-cytochrome complex using heteronuclear nmr2 and docking simulation
46	c2v4jE_	Alignment	not modelled	99.1	26	PDB header: oxidoreductase Chain: E: PDB Molecule: sulfite reductase, dissimilatory-type subunit PDBTitle: the crystal structure of desulfovibrio vulgaris2 dissimilatory sulfite reductase bound to dsrC provides3 novel insights into the mechanism of sulfate respiration
47	d3c7bb1	Alignment	not modelled	99.0	28	Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
48	c3c7bA_	Alignment	not modelled	98.8	14	PDB header: oxidoreductase Chain: A: PDB Molecule: sulfite reductase, dissimilatory-type subunit alpha; PDBTitle: structure of the dissimilatory sulfite reductase from archaeoglobus2 fulgidus
49	c2v4jA_	Alignment	not modelled	98.7	13	PDB header: oxidoreductase Chain: A: PDB Molecule: sulfite reductase, dissimilatory-type subunit PDBTitle: the crystal structure of desulfovibrio vulgaris2 dissimilatory sulfite reductase bound to dsrC provides3 novel insights into the mechanism of sulfate respiration
50	c3bk7A_	Alignment	not modelled	98.3	29	PDB header: hydrolase/translation Chain: A: PDB Molecule: abc transporter atp-binding protein; PDBTitle: structure of the complete abce1/rnaase-l inhibitor protein2 from pyrococcus abyssi
51	d2bs2b1	Alignment	not modelled	98.1	15	Fold: Globin-like Superfamily: alpha-helical ferredoxin Family: Fumarate reductase/Succinate dehydrogenase iron-sulfur protein, C-terminal domain
52	c2b76N_	Alignment	not modelled	98.0	26	PDB header: oxidoreductase Chain: N: PDB Molecule: fumarate reductase iron-sulfur protein;

					PDBTitle: e. coli quinol fumarate reductase frda e49q mutation
53	c2bs2E	Alignment	not modelled	98.0	24 PDB header: oxidoreductase Chain: E: PDB Molecule: quinol-fumarate reductase iron-sulfur subunit b; PDBTitle: quinol:fumarate reductase from wolinnella succinogenes
54	c1nekB	Alignment	not modelled	98.0	23 PDB header: oxidoreductase/electron transport Chain: B: PDB Molecule: succinate dehydrogenase iron-sulfur protein; PDBTitle: complex ii (succinate dehydrogenase) from e. coli with2 ubiquinone bound
55	d1kf6b1	Alignment	not modelled	97.9	36 Fold: Globin-like Superfamily: alpha-helical ferredoxin Family: Fumarate reductase/Succinate dehydrogenase iron-sulfur protein, C-terminal domain
56	c2h89B	Alignment	not modelled	97.9	25 PDB header: oxidoreductase Chain: B: PDB Molecule: succinate dehydrogenase ip subunit; PDBTitle: avian respiratory complex ii with malonate bound
57	d1nekbl	Alignment	not modelled	97.8	23 Fold: Globin-like Superfamily: alpha-helical ferredoxin Family: Fumarate reductase/Succinate dehydrogenase iron-sulfur protein, C-terminal domain
58	d2v4jb1	Alignment	not modelled	97.5	33 Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
59	c3cf4A	Alignment	not modelled	97.4	34 PDB header: oxidoreductase Chain: A: PDB Molecule: acetyl-coa decarboxylase/synthase alpha subunit; PDBTitle: structure of the codh component of the m. barkeri acds complex
60	d3c7ba1	Alignment	not modelled	91.7	19 Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
61	d2v4ja1	Alignment	not modelled	91.2	18 Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Ferredoxin domains from multidomain proteins
62	d1fxda	Alignment	not modelled	84.4	20 Fold: Ferredoxin-like Superfamily: 4Fe-4S ferredoxins Family: Single 4Fe-4S cluster ferredoxin
63	c2vdcl	Alignment	not modelled	78.4	35 PDB header: oxidoreductase Chain: I: PDB Molecule: glutamate synthase [nadph] small chain; PDBTitle: the 9.5 a resolution structure of glutamate synthase from2 cryo-electron microscopy and its oligomerization behavior3 in solution: functional implications.
64	d1xpaa2	Alignment	not modelled	67.3	25 Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: DNA repair factor XPA DNA- and RPA-binding domain, N-terminal subdomain
65	c2drpD	Alignment	not modelled	41.9	19 PDB header: transcription/dna Chain: D: PDB Molecule: protein (tramtrack dna-binding domain); PDBTitle: the crystal structure of a two zinc-finger peptide reveals2 an extension to the rules for zinc-finger/dna recognition
66	c1meyG	Alignment	not modelled	39.0	14 PDB header: transferase/dna Chain: G: PDB Molecule: consensus zinc finger; PDBTitle: crystal structure of a designed zinc finger protein bound2 to dna
67	d2cota1	Alignment	not modelled	36.6	15 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
68	d1x6ea2	Alignment	not modelled	36.2	15 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
69	d1llmc2	Alignment	not modelled	35.1	24 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
70	c2k5cA	Alignment	not modelled	34.4	21 PDB header: metal binding protein Chain: A: PDB Molecule: uncharacterized protein pf0385; PDBTitle: nmr structure for pf0385
71	c2xigA	Alignment	not modelled	34.2	15 PDB header: transcription Chain: A: PDB Molecule: ferric uptake regulation protein; PDBTitle: the structure of the helicobacter pylori ferric uptake2 regulator fur reveals three functional metal binding sites
72	d2cota2	Alignment	not modelled	33.8	11 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
73	d1x6ha2	Alignment	not modelled	31.3	19 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
74	c1g8jC	Alignment	not modelled	31.2	13 PDB header: oxidoreductase Chain: C: PDB Molecule: arsenite oxidase; PDBTitle: crystal structure analysis of arsenite oxidase from2 alcaligenes faecalis
75	d2ct1a2	Alignment	not modelled	31.2	19 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
76	d1u85a1	Alignment	not modelled	31.1	24 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
77	c2eoqA	Alignment	not modelled	27.9	12 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the c2h2 type zinc finger (region 283-2 315) of human zinc finger protein 224
78	c2ytmA	Alignment	not modelled	27.8	11 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region

					696-2 728) of human zinc finger protein 28 homolog
79	d1x6ea1	Alignment	not modelled	27.1	12 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
80	c2ytbA	Alignment	not modelled	26.7	15 PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger protein 32; PDBTitle: solution structure of c2h2 type zinc finger domain 5 in2 zinc finger protein 32
81	c2elsA	Alignment	not modelled	26.5	13 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 406; PDBTitle: solution structure of the 2nd c2h2 zinc finger of human2 zinc finger protein 406
82	c2yu5A	Alignment	not modelled	26.3	24 PDB header: rna binding protein Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the zf-c2h2 domain (669-699aa) in2 zinc finger protein 473
83	c3tsuA	Alignment	not modelled	26.0	24 PDB header: transferase Chain: A: PDB Molecule: transcriptional regulatory protein; PDBTitle: crystal structure of e. coli hypf with amp-pnp and carbamoyl phosphate
84	c2elta	Alignment	not modelled	25.5	13 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 406; PDBTitle: solution structure of the 3rd c2h2 zinc finger of human2 zinc finger protein 406
85	d1a1ia3	Alignment	not modelled	25.3	24 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
86	d2epsa1	Alignment	not modelled	24.8	13 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
87	d1xf7a	Alignment	not modelled	24.7	20 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
88	c1xf7A	Alignment	not modelled	24.7	20 PDB header: transcription Chain: A: PDB Molecule: wilms' tumor protein; PDBTitle: high resolution nmr structure of the wilms' tumor2 suppressor protein (wt1) finger 3
89	c2eq4A	Alignment	not modelled	24.5	15 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the 11th c2h2 type zinc finger domain2 of zinc finger protein 224
90	d1p7aa	Alignment	not modelled	24.4	29 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
91	c2emwA	Alignment	not modelled	24.4	12 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 268; PDBTitle: solution structure of the c2h2 type zinc finger (region 301-2 331) of human zinc finger protein 268
92	d1srka	Alignment	not modelled	24.1	19 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
93	d1a1ia2	Alignment	not modelled	23.4	19 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
94	c2ythA	Alignment	not modelled	23.4	11 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the c2h2 type zinc finger (region 479-2 511) of human zinc finger protein 224
95	c2emlA	Alignment	not modelled	22.8	15 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 752-2 784) of human zinc finger protein 28 homolog
96	c1va3A	Alignment	not modelled	22.8	15 PDB header: transcription Chain: A: PDB Molecule: transcription factor sp1; PDBTitle: solution structure of transcription factor sp1 dna binding2 domain (zinc finger 3)
97	c2ytaA	Alignment	not modelled	22.5	19 PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger protein 32; PDBTitle: solution structure of c2h2 type zinc finger domain 3 in2 zinc finger protein 32
98	c2yttA	Alignment	not modelled	22.5	19 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 204-2 236) of human zinc finger protein 473
99	c3mwmA	Alignment	not modelled	22.4	15 PDB header: transcription Chain: A: PDB Molecule: putative metal uptake regulation protein; PDBTitle: graded expression of zinc-responsive genes through two regulatory2 zinc-binding sites in zur
100	c2en6A	Alignment	not modelled	22.3	11 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 268; PDBTitle: solution structure of the c2h2 type zinc finger (region 887-2 919) of human zinc finger protein 268
101	c2elqA	Alignment	not modelled	22.2	16 PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 406; PDBTitle: solution structure of the 14th c2h2 zinc finger of human2 zinc finger protein 406
102	d1sp1a	Alignment	not modelled	22.2	15 Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
103	c1sp1A	Alignment	not modelled	22.2	15 PDB header: zinc finger Chain: A: PDB Molecule: sp1f3; PDBTitle: nmr structure of a zinc finger domain from transcription2 factor sp1f3, minimized average structure
104	d1gtea1	Alignment	not modelled	22.2	35 Fold: Globin-like Superfamily: alpha-helical ferredoxin Family: Dihydropyrimidine dehydrogenase, N-terminal domain

105	c2kvfA_		not modelled	22.1	21	PDB header: transcription Chain: A: PDB Molecule: zinc finger and btb domain-containing protein 32; PDBTitle: structure of the three-cys2his2 domain of mouse testis zinc2 finger protein
106	c1areA_		not modelled	22.0	30	PDB header: transcription regulation Chain: A: PDB Molecule: yeast transcription factor adr1; PDBTitle: structures of dna-binding mutant zinc finger domains:2 implications for dna binding
107	c2eovA_		not modelled	21.9	15	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 484; PDBTitle: solution structure of the c2h2 type zinc finger (region 519-2 551) of human zinc finger protein 484
108	c2em4A_		not modelled	21.6	11	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 724-2 756) of human zinc finger protein 28 homolog
109	d1ubdc2		not modelled	21.4	19	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
110	c2epzA_		not modelled	21.0	11	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the 4th c2h2 type zinc finger domain2 of zinc finger protein 28 homolog
111	c2ytjA_		not modelled	21.0	11	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 484; PDBTitle: solution structure of the c2h2 type zinc finger (region 771-2 803) of human zinc finger protein 484
112	c2yteA_		not modelled	20.9	14	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 484-2 512) of human zinc finger protein 473
113	c2en2A_		not modelled	20.8	12	PDB header: transcription Chain: A: PDB Molecule: b-cell lymphoma 6 protein; PDBTitle: solution structure of the c2h2 type zinc finger (region 598-2 626) of human b-cell lymphoma 6 protein