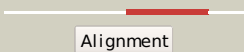

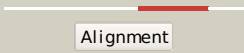


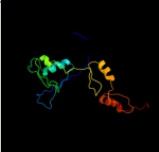
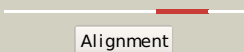

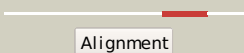
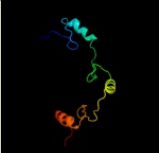
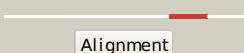
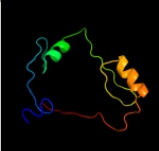


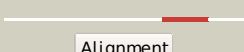

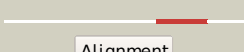
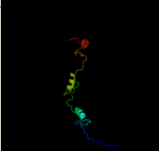
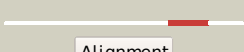

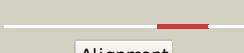
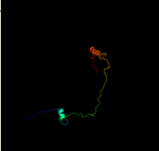
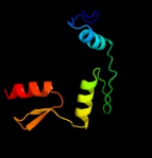
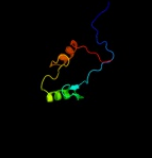


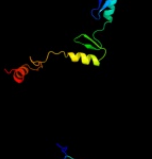
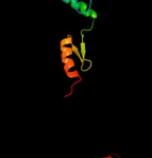
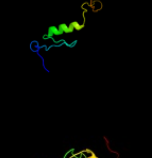

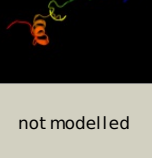


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	A0PJY2
Date	Fri May 25 09:29:34 BST 2012
Unique Job ID	993c879d1f628d04

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1tf6D_	 Alignment		99.9	20	PDB header: transcription/dna Chain: D; PDB Molecule: protein (transcription factor iiiia); PDBTitle: co-crystal structure of xenopus tiiiia zinc finger domain2 bound to the 5s ribosomal rna gene internal control region
2	c2wbtA_	 Alignment		99.9	22	PDB header: dna-binding protein Chain: A; PDB Molecule: b-129; PDBTitle: the structure of a double c2h2 zinc finger protein from a2 hyperthermophilic archaeal virus in the absence of dna
3	c2rpcA_	 Alignment		99.9	25	PDB header: transcription Chain: A; PDB Molecule: zinc finger protein zic 3; PDBTitle: solution structure of the tandem zf-c2h2 domains from the2 human zinc finger protein zic 3
4	c2cshA_	 Alignment		99.9	29	PDB header: transcription Chain: A; PDB Molecule: zinc finger protein 297b; PDBTitle: solution structure of tandem repeat of the zf-c2h2 domains2 of human zinc finger protein 297b
5	c2dmdA_	 Alignment		99.8	35	PDB header: transcription Chain: A; PDB Molecule: zinc finger protein 64, isoforms 1 and 2; PDBTitle: solution structure of the n-terminal c2h2 type zinc-binding2 domain of the zinc finger protein 64, isoforms 1 and 2
6	c1x6hA_	 Alignment		99.8	25	PDB header: dna binding protein Chain: A; PDB Molecule: transcriptional repressor ctcf; PDBTitle: solution structures of the c2h2 type zinc finger domain of2 human transcriptional repressor ctcf
7	c2i13A_	 Alignment		99.8	43	PDB header: dna binding protein/dna Chain: A; PDB Molecule: aart; PDBTitle: aart, a six finger zinc finger designed to recognize ann triplets
8	c1wjpa_	 Alignment		99.8	16	PDB header: metal binding protein Chain: A; PDB Molecule: zinc finger protein 295; PDBTitle: solution structure of zf-c2h2 domains from human zinc2 finger protein 295
9	c2ee8A_	 Alignment		99.8	31	PDB header: gene regulation Chain: A; PDB Molecule: protein odd-skipped-related 2; PDBTitle: solution structure of three zf-c2h2 domains from mouse2 protein odd-skipped-related 2 splicing isoform 2
10	c1g2dC_	 Alignment		99.8	36	PDB header: transcription/dna Chain: C; PDB Molecule: tata box zinc finger protein; PDBTitle: structure of a cys2his2 zinc finger/tata box complex (clone2 #2)
11	c2dmiA_	 Alignment		99.8	17	PDB header: transcription Chain: A; PDB Molecule: teashirt homolog 3; PDBTitle: solution structure of the first and the second zf-c2h2 like2 domains of human teashirt homolog 3

12	c2kmkA	Alignment		99.7	36	PDB header: dna binding protein/dna Chain: A: PDB Molecule: zinc finger protein gfi-1; PDBTitle: gfi-1 zinc fingers 3-5 complexed with dna
13	c2cotA	Alignment		99.7	31	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 435; PDBTitle: solution structure of the first and second zf-c2h2 domain2 of zinc finger protein 435
14	c2dlqA	Alignment		99.7	32	PDB header: transcription Chain: A: PDB Molecule: gli-kruppel family member hkr3; PDBTitle: solution structure of the tandem four zf-c2h2 domain2 repeats of murine gli-kruppel family member hkr3
15	c1llmC	Alignment		99.7	28	PDB header: transcription/dna Chain: C: PDB Molecule: chimera of zif23-gcn4; PDBTitle: crystal structure of a zif23-gcn4 chimera bound to dna
16	c2yt9A	Alignment		99.7	36	PDB header: transcription Chain: A: PDB Molecule: zinc finger-containing protein 1; PDBTitle: solution structure of c2h2 type zinc finger domain 345 in2 zinc finger protein 278
17	c2lceA	Alignment		99.7	39	PDB header: transcription regulator Chain: A: PDB Molecule: b-cell lymphoma 6 protein; PDBTitle: chemical shift assignment of hr4436b from homo sapiens, northeast2 structural genomics consortium
18	c1x6eA	Alignment		99.7	36	PDB header: dna binding protein Chain: A: PDB Molecule: zinc finger protein 24; PDBTitle: solution structures of the c2h2 type zinc finger domain of2 human zinc finger protein 24
19	c2d9hA	Alignment		99.7	28	PDB header: dna binding protein Chain: A: PDB Molecule: zinc finger protein 692; PDBTitle: solution structure of the forth and fifth zf-c2h2 domains2 of zinc finger protein 692
20	c1x5wA	Alignment		99.7	25	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 64, isoforms 1; PDBTitle: solution structure of the c2h2 type zinc-binding domain of2 human zinc finger protein 64, isoforms 1 and 2
21	c2adrA	Alignment	not modelled	99.6	34	PDB header: transcription regulation Chain: A: PDB Molecule: adr1; PDBTitle: adr1 dna-binding domain from saccharomyces cerevisiae, nmr,2 25 structures
22	c2gliA	Alignment	not modelled	99.6	17	PDB header: transcription/dna Chain: A: PDB Molecule: protein (five-finger gli); PDBTitle: five-finger gli/dna complex
23	c2dlkA	Alignment	not modelled	99.6	26	PDB header: dna binding protein Chain: A: PDB Molecule: novel protein; PDBTitle: solution structure of the first and the second zf-c2h22 domains of zinc finger protein 692
24	c1tf3A	Alignment	not modelled	99.6	35	PDB header: transcription/dna Chain: A: PDB Molecule: transcription factor iiiia; PDBTitle: tfiiiia finger 1-3 bound to dna, nmr, 22 structures
25	c2ebtA	Alignment	not modelled	99.6	32	PDB header: transcription Chain: A: PDB Molecule: krueppel-like factor 5; PDBTitle: solution structure of three tandem repeats of zf-c2h22 domains from human kruppel-like factor 5
26	c2ej4A	Alignment	not modelled	99.6	19	PDB header: gene regulation Chain: A: PDB Molecule: zinc finger protein zic 3; PDBTitle: functional and structural basis of nuclear localization2 signal in zic3 zinc finger domain: a role of conserved3 tryptophan residue in the zinc finger domain
27	c2drpD	Alignment	not modelled	99.6	29	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
28	c2ctiA	Alignment	not modelled	99.6	37	PDB header: transcription Chain: A: PDB Molecule: transcriptional repressor ctcf; PDBTitle: solution structure of the zinc finger domain of2 transcriptional repressor ctcf protein

29	d2csha1	Alignment	not modelled	99.6	30	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
30	c2ctdA_	Alignment	not modelled	99.6	15	PDB header: metal binding protein Chain: A: PDB Molecule: zinc finger protein 512; PDBTitle: solution structure of two zf-c2h2 domains from human zinc2 finger protein 512
31	c1un6D_	Alignment	not modelled	99.6	22	PDB header: rna-binding protein/rna Chain: D: PDB Molecule: transcription factor iiiia; PDBTitle: the crystal structure of a zinc finger - rna complex2 reveals two modes of molecular recognition
32	c1f2i1_	Alignment	not modelled	99.6	27	PDB header: transcription/dna Chain: I: PDB Molecule: fusion of n-terminal 17-mer peptide extension to PDBTitle: cocrystal structure of selected zinc finger dimer bound to2 dna
33	c1bboA_	Alignment	not modelled	99.5	27	PDB header: dna-binding protein Chain: A: PDB Molecule: human enhancer-binding protein mbp-1; PDBTitle: high-resolution solution structure of the double cys2*his22 zinc finger from the human enhancer binding protein mbp-1
34	c2epaA_	Alignment	not modelled	99.5	27	PDB header: transcription Chain: A: PDB Molecule: krueppel-like factor 10; PDBTitle: solution structure of the first and second zf-c2h2 domains2 from human krueppel-like factor 10
35	c2prtA_	Alignment	not modelled	99.5	33	PDB header: transcription/dna Chain: A: PDB Molecule: wilms tumor 1; PDBTitle: structure of the wilms tumor suppressor protein zinc finger2 domain bound to dna
36	c2ghfA_	Alignment	not modelled	99.4	20	PDB header: transcription, metal binding protein Chain: A: PDB Molecule: zinc fingers and homeoboxes protein 1; PDBTitle: solution structure of the complete zinc-finger region of2 human zinc-fingers and homeoboxes 1 (zhx1)
37	c2emcA_	Alignment	not modelled	99.4	32	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 641-2 673) of human zinc finger protein 473
38	c2eomA_	Alignment	not modelled	99.4	29	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 95 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 341-2 373) of human zinc finger protein 95 homolog
39	c2en6A_	Alignment	not modelled	99.4	38	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
40	c2ythA_	Alignment	not modelled	99.4	36	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
41	c2epzA_	Alignment	not modelled	99.4	34	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
42	c1ubdC_	Alignment	not modelled	99.4	31	PDB header: transcription/dna Chain: C: PDB Molecule: protein (yy1 zinc finger domain); PDBTitle: co-crystal structure of human yy1 zinc finger domain bound2 to the adeno-associated virus p5 initiator element
43	c2ytjA_	Alignment	not modelled	99.4	34	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
44	c2ytmA_	Alignment	not modelled	99.4	31	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
45	c2em3A_	Alignment	not modelled	99.4	36	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 640-2 672) of human zinc finger protein 28 homolog
46	c2emlA_	Alignment	not modelled	99.4	33	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
47	c2gqjA_	Alignment	not modelled	99.4	18	PDB header: gene regulation Chain: A: PDB Molecule: zinc finger protein kiaa1196; PDBTitle: solution structure of the two zf-c2h2 like domains(493-575)2 of human zinc finger protein kiaa1196
48	c2epsA_	Alignment	not modelled	99.4	24	PDB header: transcription Chain: A: PDB Molecule: poz-, at hook-, and zinc finger-containing PDBTitle: solution structure of the 4th zinc finger domain of zinc2 finger protein 278
49	c1zu1A_	Alignment	not modelled	99.3	12	PDB header: rna binding protein Chain: A: PDB Molecule: rna binding protein zfa; PDBTitle: solution structure of the n-terminal zinc fingers of the2 xenopus laevis double stranded rna binding protein zfa
50	c2eovA_	Alignment	not modelled	99.3	29	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
51	c2em0A_	Alignment	not modelled	99.3	34	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the 18th zf-c2h2 domain from human2 zinc finger protein 224
52	c2emwA_	Alignment	not modelled	99.3	33	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
53	c2epwA_	Alignment	not modelled	99.3	36	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 268; PDBTitle: solution structure of the 24th c2h2 type zinc finger domain2 of zinc finger protein 268
						PDB header: transcription

54	c2eq4A	Alignment	not modelled	99.3	28	Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the 11th c2h2 type zinc finger domain2 of zinc finger protein 224
55	c1b8tA	Alignment	not modelled	99.3	15	PDB header: contractile Chain: A: PDB Molecule: protein (crp1); PDBTitle: solution structure of the chicken crp1
56	c2epqA	Alignment	not modelled	99.3	27	PDB header: transcription Chain: A: PDB Molecule: poz-, at hook-, and zinc finger-containing PDBTitle: solution structure of the third zinc finger domain of zinc2 finger protein 278
57	c2eozA	Alignment	not modelled	99.3	32	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 809-2 841) of human zinc finger protein 473
58	c2em4A	Alignment	not modelled	99.3	26	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
59	c2eohA	Alignment	not modelled	99.3	26	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 28 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 780-2 812) of human zinc finger protein 28 homolog
60	c2epuA	Alignment	not modelled	99.3	28	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 32; PDBTitle: solution structure of the second c2h2 type zinc finger2 domain of zinc finger protein 32
61	c2eprA	Alignment	not modelled	99.2	33	PDB header: transcription Chain: A: PDB Molecule: poz-, at hook-, and zinc finger-containing PDBTitle: solution structure of the second zinc finger domain of2 zinc finger protein 278
62	c2eoaA	Alignment	not modelled	99.2	36	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
63	c2yttA	Alignment	not modelled	99.2	35	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
64	c2entA	Alignment	not modelled	99.2	29	PDB header: transcription Chain: A: PDB Molecule: krueppel-like factor 15; PDBTitle: solution structure of the second c2h2-type zinc finger2 domain from human krueppel-like factor 15
65	c2em5A	Alignment	not modelled	99.2	24	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 95 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 768-2 800) of human zinc finger protein 95 homolog
66	c2emzA	Alignment	not modelled	99.2	33	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 95 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 628-2 660) of human zinc finger protein 95 homolog
67	c2ytbA	Alignment	not modelled	99.2	33	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
68	c2epcA	Alignment	not modelled	99.1	25	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 32; PDBTitle: solution structure of zinc finger domain 7 in zinc finger2 protein 32
69	c2en3A	Alignment	not modelled	99.1	26	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 95 homolog; PDBTitle: solution structure of the c2h2 type zinc finger (region 796-2 828) of human zinc finger protein 95 homolog
70	c2en2A	Alignment	not modelled	99.1	33	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
71	d2ct1a2	Alignment	not modelled	99.1	28	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
72	c2ytaA	Alignment	not modelled	99.1	35	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
73	c2elzA	Alignment	not modelled	99.1	29	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the 17th zf-c2h2 domain from human2 zinc finger protein 224
74	d1x6ha2	Alignment	not modelled	99.1	34	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
75	d2epsa1	Alignment	not modelled	99.1	29	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
76	c2en8A	Alignment	not modelled	99.0	31	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 224; PDBTitle: solution structure of the c2h2 type zinc finger (region 171-2 203) of human zinc finger protein 224
77	c2el5A	Alignment	not modelled	99.0	32	PDB header: transcription Chain: A: PDB Molecule: zinc finger protein 268; PDBTitle: solution structure of the 18th zf-c2h2 domain from human2 zinc finger protein 268
78	c2yteA	Alignment	not modelled	99.0	28	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
79	d1p7aa	Alignment	not modelled	99.0	30	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
						PDB header: transcription

80	c2eoxA	Alignment	not modelled	99.0	29	Chain: A: PDB Molecule: zinc finger protein 473; PDBTitle: solution structure of the c2h2 type zinc finger (region 315-2 345) of human zinc finger protein 473
81	d2cota2	Alignment	not modelled	99.0	24	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
82	d2epra1	Alignment	not modelled	98.9	42	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
83	c2yu5A	Alignment	not modelled	98.9	32	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
84	c2elsA	Alignment	not modelled	98.9	34	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
85	c2elta	Alignment	not modelled	98.9	31	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
86	d2dlka2	Alignment	not modelled	98.9	33	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
87	c2elqA	Alignment	not modelled	98.8	37	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
88	d1srka	Alignment	not modelled	98.8	24	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
89	c1zw8A	Alignment	not modelled	98.8	19	PDB header: transcription Chain: A: PDB Molecule: zinc-responsive transcriptional regulator zap1; PDBTitle: solution structure of a zap1 zinc-responsive domain2 provides insights into metalloregulatory transcriptional3 repression in saccharomyces cerevisiae
90	d1x6ea1	Alignment	not modelled	98.8	39	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
91	d1ncsa	Alignment	not modelled	98.7	29	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
92	c1xf7A	Alignment	not modelled	98.7	34	PDB header: transcription Chain: A: PDB Molecule: wilms' tumor protein; PDBTitle: high resolution nmr structure of the wilms' tumor2 suppressor protein (wt1) finger 3
93	d1xf7a	Alignment	not modelled	98.7	34	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
94	d1u85a1	Alignment	not modelled	98.7	36	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
95	c1ardA	Alignment	not modelled	98.7	31	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
96	c1areA	Alignment	not modelled	98.7	31	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
97	c1arfA	Alignment	not modelled	98.6	31	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
98	d2adra1	Alignment	not modelled	98.6	31	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
99	c1meyG	Alignment	not modelled	98.6	43	PDB header: transferase/dna Chain: G: PDB Molecule: consensus zinc finger; PDBTitle: crystal structure of a designed zinc finger protein bound2 to dna
100	c1va3A	Alignment	not modelled	98.6	41	PDB header: transcription Chain: A: PDB Molecule: transcription factor sp1; PDBTitle: solution structure of transcription factor sp1 dna binding2 domain (zinc finger 3)
101	c1sp1A	Alignment	not modelled	98.6	41	PDB header: zinc finger Chain: A: PDB Molecule: sp1f3; PDBTitle: nmr structure of a zinc finger domain from transcription2 factor sp1f3, minimized average structure
102	d1sp1a	Alignment	not modelled	98.6	41	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
103	d1f2ig1	Alignment	not modelled	98.6	24	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
104	c1va1A	Alignment	not modelled	98.5	26	PDB header: transcription Chain: A: PDB Molecule: transcription factor sp1; PDBTitle: solution structure of transcription factor sp1 dna binding2 domain (zinc finger 1)
105	d1alia2	Alignment	not modelled	98.5	39	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
						PDB header: de novo protein

106	c1rika_	Alignment	not modelled	98.5	39	Chain: A; PDB Molecule: e6apc1 peptide; PDBTitle: e6-binding zinc finger (e6apc1)
107	d2glia3	Alignment	not modelled	98.5	32	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
108	d1paaa_	Alignment	not modelled	98.4	33	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
109	d1ubdc3	Alignment	not modelled	98.4	36	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
110	c2d9kA_	Alignment	not modelled	98.4	17	PDB header: immune system Chain: A; PDB Molecule: fln29 gene product; PDBTitle: solution structure of the zf-traf domain of fln29 gene2 product
111	d2cota1	Alignment	not modelled	98.4	41	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
112	d1llmc2	Alignment	not modelled	98.4	41	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
113	d1sp2a_	Alignment	not modelled	98.4	34	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
114	d1u86a1	Alignment	not modelled	98.4	32	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
115	d1tf3a2	Alignment	not modelled	98.4	39	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
116	c3hcsA_	Alignment	not modelled	98.4	14	PDB header: signaling protein Chain: A; PDB Molecule: tnf receptor-associated factor 6; PDBTitle: crystal structure of the n-terminal domain of traf6
117	d1a1ia3	Alignment	not modelled	98.3	38	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
118	d1x6ea2	Alignment	not modelled	98.3	42	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2
119	c2xqnT_	Alignment	not modelled	98.3	14	PDB header: metal-binding protein Chain: T; PDB Molecule: testin; PDBTitle: complex of the 2nd and 3rd lim domains of tes with the evh12 domain of mena and the n-terminal domain of actin-like protein3 arp7a
120	d1aaya1	Alignment	not modelled	98.2	26	Fold: beta-beta-alpha zinc fingers Superfamily: beta-beta-alpha zinc fingers Family: Classic zinc finger, C2H2