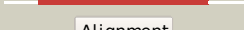
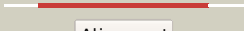


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3bwgA_	 Alignment		100.0	19	PDB header: transcription regulator Chain: A: PDB Molecule: uncharacterized hth-type transcriptional regulator yydk; PDBTitle: the crystal structure of possible transcriptional regulator yydk from2 bacillus subtilis subsp. subtilis str. 168
2	c3edpB_	 Alignment		100.0	19	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: lin2111 protein; PDBTitle: the crystal structure of the protein lin2111 (functionally unknown)2 from listeria innocua clip11262
3	c3f8mA_	 Alignment		100.0	24	PDB header: transcription Chain: A: PDB Molecule: gntr-family protein transcriptional regulator; PDBTitle: crystal structure of phnf from mycobacterium smegmatis
4	c3eetA_	 Alignment		100.0	19	PDB header: transcription regulator Chain: A: PDB Molecule: putative gntr-family transcriptional regulator; PDBTitle: crystal structure of putative gntr-family transcriptional2 regulator
5	c2wv0H_	 Alignment		100.0	25	PDB header: transcription Chain: H: PDB Molecule: hth-type transcriptional repressor yvoa; PDBTitle: crystal structure of the gntr-hutc family member yvoa from2 bacillus subtilis
6	c3f8lC_	 Alignment		99.9	24	PDB header: transcription Chain: C: PDB Molecule: hth-type transcriptional repressor phnf; PDBTitle: crystal structure of the effector domain of phnf from mycobacterium2 smegmatis
7	c3by6C_	 Alignment		99.9	27	PDB header: transcription regulator Chain: C: PDB Molecule: predicted transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator from oenococcus oeni
8	c3neuA_	 Alignment		99.9	25	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lin1836 protein; PDBTitle: the crystal structure of a functionally-unknown protein lin1836 from2 listeria innocua clip11262
9	c2du9A_	 Alignment		99.9	21	PDB header: transcription Chain: A: PDB Molecule: predicted transcriptional regulators; PDBTitle: crystal structure of the transcriptional factor from c. glutamicum
10	d2fa1a1	 Alignment		99.9	25	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
11	d2p19a1	 Alignment		99.9	25	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain

12	d3cnva1	Alignment		99.9	19	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
13	c2ra5A	Alignment		99.9	23	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of the putative transcriptional regulator2 from streptomyces coelicolor
14	d2ra5a1	Alignment		99.9	23	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
15	d2ooia1	Alignment		99.9	20	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
16	d1v4ra1	Alignment		99.8	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
17	d3bwga2	Alignment		99.8	19	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
18	c2di3A	Alignment		99.8	30	PDB header: transcription Chain: A: PDB Molecule: bacterial regulatory proteins, gntR family; PDBTitle: crystal structure of the transcriptional factor cgl29152 from corynebacterium glutamicum
19	d2ikka1	Alignment		99.8	20	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
20	d2ogga1	Alignment		99.8	17	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
21	c3c7jA	Alignment	not modelled	99.8	23	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, gntR family; PDBTitle: crystal structure of transcriptional regulator (gntR family member)2 from pseudomonas syringae pv. tomato str. dc3000
22	c1e2xA	Alignment	not modelled	99.8	28	PDB header: transcriptional regulation Chain: A: PDB Molecule: fatty acid metabolism regulator protein; PDBTitle: fadr, fatty acid responsive transcription factor from e.2 coli
23	c3fmsA	Alignment	not modelled	99.8	24	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, gntR family; PDBTitle: crystal structure of tm0439, a gntR transcriptional2 regulator
24	c3ihuA	Alignment	not modelled	99.8	24	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, gntR family; PDBTitle: crystal structure of dna binding protein (yp_298823.1) from ralstonia2 eutropha jmp134 at 1.92 a resolution
25	c3hfiA	Alignment	not modelled	99.8	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative regulator; PDBTitle: the crystal structure of the putative regulator from escherichia coli2 cft073
26	d3bwga1	Alignment	not modelled	99.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
27	c2hs5A	Alignment	not modelled	99.8	26	PDB header: transcription regulator Chain: A: PDB Molecule: putative transcriptional regulator gntR; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator gntR from rhodococcus sp. rha1
28	c3ic7A	Alignment	not modelled	99.8	19	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of putative transcriptional regulator of gntR family2 from bacteroides thetaiotaomicron PDB header: transcription regulator

29	c3lheA_	Alignment	not modelled	99.8	21	Chain: A: PDB Molecule: gntr family transcriptional regulator; PDBTitle: the crystal structure of the c-terminal domain of a gntr2 family transcriptional regulator from bacillus anthracis3 str. sterne
30	dlhw1a1	Alignment	not modelled	99.8	33	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GnTr-like transcriptional regulators
31	d2pkha1	Alignment	not modelled	99.8	19	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
32	c3tqnC_	Alignment	not modelled	99.8	22	PDB header: transcription Chain: C: PDB Molecule: transcriptional regulator, gntr family; PDBTitle: structure of the transcriptional regulator of the gntr family, from2 coxiella burnetii.
33	d3ddva1	Alignment	not modelled	99.8	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
34	d2hs5a1	Alignment	not modelled	99.8	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GnTr-like transcriptional regulators
35	c2h09A_	Alignment	not modelled	98.2	21	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator mntr; PDBTitle: crystal structure of diphtheria toxin repressor like protein2 from e. coli
36	d1biaa1	Alignment	not modelled	98.2	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
37	c1f5tA_	Alignment	not modelled	97.9	20	PDB header: transcription/dna Chain: A: PDB Molecule: diphtheria toxin repressor; PDBTitle: diphtheria tox repressor (c102d mutant) complexed with2 nickel and dtxr consensus binding sequence
38	d1stza1	Alignment	not modelled	97.8	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Heat-inducible transcription repressor HrcA, N-terminal domain
39	d1jhfa1	Alignment	not modelled	97.8	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: LexA repressor, N-terminal DNA-binding domain
40	c2it0A_	Alignment	not modelled	97.7	15	PDB header: transcription/dna Chain: A: PDB Molecule: iron-dependent repressor ider; PDBTitle: crystal structure of a two-domain ider-dna complex crystal2 form ii
41	c3hruA_	Alignment	not modelled	97.6	22	PDB header: transcription Chain: A: PDB Molecule: metalloregulator scar; PDBTitle: crystal structure of scar with bound zn2+
42	c3ctaA_	Alignment	not modelled	97.5	20	PDB header: transferase Chain: A: PDB Molecule: riboflavin kinase; PDBTitle: crystal structure of riboflavin kinase from thermoplasma2 acidophilum
43	d1g3wa1	Alignment	not modelled	97.4	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
44	d1j5ya1	Alignment	not modelled	97.4	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
45	c1g3wA_	Alignment	not modelled	97.4	20	PDB header: gene regulation Chain: A: PDB Molecule: diphtheria toxin repressor; PDBTitle: cd-cys102ser dtxr
46	d1i5za1	Alignment	not modelled	97.3	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
47	d2isya1	Alignment	not modelled	97.3	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
48	c2ewnA_	Alignment	not modelled	97.3	19	PDB header: ligase, transcription Chain: A: PDB Molecule: bira bifunctional protein; PDBTitle: ecoli biotin repressor with co-repressor analog
49	c1j5yA_	Alignment	not modelled	97.2	16	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, biotin repressor family; PDBTitle: crystal structure of transcriptional regulator (tm1602) from2 thermotoga maritima at 2.3 a resolution
50	c3r0aB_	Alignment	not modelled	97.2	24	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator; PDBTitle: possible transcriptional regulator from methanosarcina mazei go1 (gi2 21227196)
51	c1zybA_	Alignment	not modelled	97.1	15	PDB header: transcription regulator Chain: A: PDB Molecule: transcription regulator, crp family; PDBTitle: crystal structure of transcription regulator from bacteroides2 thetaiotaomicron vpi-5482 at 2.15 a resolution
52	c3k69A_	Alignment	not modelled	97.1	16	PDB header: transcription Chain: A: PDB Molecule: putative transcription regulator; PDBTitle: crystal structure of a putative transcriptional regulator (lp_0360)2 from lactobacillus plantarum at 1.95 a resolution
53	c2y75F_	Alignment	not modelled	97.1	20	PDB header: transcription Chain: F: PDB Molecule: hth-type transcriptional regulator cymr; PDBTitle: the structure of cymr (yrzc) the global cysteine regulator2 of b. subtilis
54	c2v79B_	Alignment	not modelled	96.9	12	PDB header: dna-binding protein Chain: B: PDB Molecule: dna replication protein dnad; PDBTitle: crystal structure of the n-terminal domain of dnad from2 bacillus subtilis
55	d2h6ca1	Alignment	not modelled	96.8	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain

						Family: CAP C-terminal domain-like
56	c1fx7C	Alignment	not modelled	96.8	14	PDB header: signaling protein Chain: C: PDB Molecule: iron-dependent repressor ider; PDBTitle: crystal structure of the iron-dependent regulator (ider)2 from mycobacterium tuberculosis
57	c3lwFD	Alignment	not modelled	96.8	20	PDB header: transcription regulator Chain: D: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of putative transcriptional regulator (np_470886.1)2 from listeria innocua at 2.06 a resolution
58	c2gauA	Alignment	not modelled	96.7	10	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, crp/fnr family; PDBTitle: crystal structure of transcriptional regulator, crp/fnr family from2 porphyromonas gingivalis (apc80792), structural genomics, mcsg
59	d1xd7a	Alignment	not modelled	96.7	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
60	c2oz6A	Alignment	not modelled	96.7	14	PDB header: dna binding protein Chain: A: PDB Molecule: virulence factor regulator; PDBTitle: crystal structure of virulence factor regulator from pseudomonas2 aeruginosa in complex with camp
61	d1sfxa	Alignment	not modelled	96.6	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
62	d2gaua1	Alignment	not modelled	96.6	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
63	c2x4hA	Alignment	not modelled	96.6	18	PDB header: transcription Chain: A: PDB Molecule: hypothetical protein sso2273; PDBTitle: crystal structure of the hypothetical protein sso2273 from2 sulfobolus solfataricus
64	c3iwzB	Alignment	not modelled	96.6	10	PDB header: transcription Chain: B: PDB Molecule: catabolite activation-like protein; PDBTitle: the c-di-gmp responsive global regulator clp links cell-cell signaling2 to virulence gene expression in xanthomonas campestris
65	c3dv8A	Alignment	not modelled	96.5	14	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, crp/fnr family; PDBTitle: crystal structure of a putative transcriptional regulator of the2 crp/fnr family (eubrec_1222) from eubacterium rectale atcc 33656 at3 2.55 a resolution
66	d3broa1	Alignment	not modelled	96.5	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
67	d2d1ha1	Alignment	not modelled	96.5	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
68	c1zreB	Alignment	not modelled	96.4	16	PDB header: gene regulation/dna Chain: B: PDB Molecule: catabolite gene activator; PDBTitle: 4 crystal structures of cap-dna with all base-pair2 substitutions at position 6, cap-[6g;17c]icap38 dna
69	d1ft9a1	Alignment	not modelled	96.4	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
70	c3bj6B	Alignment	not modelled	96.4	14	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of marr family transcription regulator sp03579
71	d2oz6a1	Alignment	not modelled	96.4	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
72	c3fx3A	Alignment	not modelled	96.4	16	PDB header: camp-binding protein Chain: A: PDB Molecule: cyclic nucleotide-binding protein; PDBTitle: structure of a putative camp-binding regulatory protein from2 silicibacter pomeroyi dss-3
73	c3d0sA	Alignment	not modelled	96.4	29	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulatory protein; PDBTitle: camp receptor protein from m.tuberculosis, camp-free form
74	c3e97A	Alignment	not modelled	96.3	19	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, crp/fnr family; PDBTitle: crystal structure of transcriptional regulator of crp/fnr2 family (yp_604437.1) from deinococcus geothermalis dsm3 11300 at 1.86 a resolution
75	d2coha1	Alignment	not modelled	96.2	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
76	c1ft9A	Alignment	not modelled	96.2	23	PDB header: transcription Chain: A: PDB Molecule: carbon monoxide oxidation system transcription PDBTitle: structure of the reduced (feii) co-sensing protein from r.2 rubrum
77	c3kccA	Alignment	not modelled	96.2	15	PDB header: transcription Chain: A: PDB Molecule: catabolite gene activator; PDBTitle: crystal structure of d138l mutant of catabolite gene activator protein
78	d3e5ua1	Alignment	not modelled	96.2	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
79	d2fbha1	Alignment	not modelled	96.1	23	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
80	d2fxaa1	Alignment	not modelled	96.1	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators

81	d1ku9a	Alignment	not modelled	96.0	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding protein Mj223
82	c2vn2B	Alignment	not modelled	96.0	19	PDB header: replication Chain: B: PDB Molecule: chromosome replication initiation protein; PDBTitle: crystal structure of the n-terminal domain of dna protein2 from geobacillus kaustophilus hta426
83	d3ctaa1	Alignment	not modelled	96.0	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
84	d2cg4a1	Alignment	not modelled	96.0	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
85	d2bgca1	Alignment	not modelled	96.0	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
86	c3e6dA	Alignment	not modelled	96.0	20	PDB header: transcription regulation Chain: A: PDB Molecule: cyclic nucleotide-binding protein; PDBTitle: crystal structure of cprk c200s
87	d2ev0a1	Alignment	not modelled	95.9	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
88	d1ilga1	Alignment	not modelled	95.9	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
89	d1dpua	Alignment	not modelled	95.9	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: C-terminal domain of RPA32
90	c1dpuA	Alignment	not modelled	95.9	22	PDB header: dna binding protein Chain: A: PDB Molecule: replication protein a (rpa32) c-terminal domain; PDBTitle: solution structure of the c-terminal domain of human rpa322 complexed with ung2(73-88)
91	c3la2A	Alignment	not modelled	95.9	14	PDB header: transcription Chain: A: PDB Molecule: global nitrogen regulator; PDBTitle: crystal structure of ntca in complex with 2-oxoglutarate
92	c2fmyB	Alignment	not modelled	95.9	14	PDB header: dna binding protein Chain: B: PDB Molecule: carbon monoxide oxidation system transcription regulator PDBTitle: co-dependent transcription factor coa from carboxydotherrmus2 hydrogenoformans (imidazole-bound form)
93	c2zwcA	Alignment	not modelled	95.8	28	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, fnr/crp family; PDBTitle: crystal structure of ttha1359, a transcriptional regulator,2 crp/fnr family from thermus thermophilus hb8
94	c2ev5B	Alignment	not modelled	95.8	16	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator mntr; PDBTitle: bacillus subtilis manganese transport regulator (mntr)2 bound to calcium
95	c2bgcA	Alignment	not modelled	95.8	11	PDB header: transcription Chain: A: PDB Molecule: prfa; PDBTitle: prfa-g145s, a constitutive active mutant of the2 transcriptional regulator in l.monocytogenes
96	d1ylfa1	Alignment	not modelled	95.7	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
97	c2l4aA	Alignment	not modelled	95.6	21	PDB header: dna binding protein Chain: A: PDB Molecule: leucine responsive regulatory protein; PDBTitle: nmr structure of the dna-binding domain of e.coli lrp
98	d1mkma1	Alignment	not modelled	95.6	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator lclR, N-terminal domain
99	c2gxgA	Alignment	not modelled	95.6	20	PDB header: transcription Chain: A: PDB Molecule: 146aa long hypothetical transcriptional regulator; PDBTitle: crystal structure of emrr homolog from hyperthermophilic archaea2 sulfobolus tokodaii strain7
100	c3k2zA	Alignment	not modelled	95.6	15	PDB header: hydrolase Chain: A: PDB Molecule: lexa repressor; PDBTitle: crystal structure of a lexa protein from thermotoga maritima
101	d2cfxa1	Alignment	not modelled	95.5	23	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
102	c2zdbA	Alignment	not modelled	95.5	19	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, crp family; PDBTitle: crystal structure of tthb099, a transcriptional regulator crp family2 from thermus thermophilus hb8
103	d2cyya1	Alignment	not modelled	95.5	26	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
104	d3deua1	Alignment	not modelled	95.5	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
105	c3dkwB	Alignment	not modelled	95.4	11	PDB header: transcription regulator Chain: B: PDB Molecule: dnr protein; PDBTitle: crystal structure of dnr from pseudomonas aeruginosa.
106	c2qwwB	Alignment	not modelled	95.4	15	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of multiple antibiotic-resistance repressor (marr)2 (yp_013417.1) from listeria monocytogenes 4b

					f2365 at 2.07 a3 resolution
107	c2zy3A_	Alignment	not modelled	95.4	10 PDB header: lyase Chain: A: PDB Molecule: l-aspartate beta-decarboxylase; PDBTitle: dodecameric l-aspartate beta-decarboxylase
108	c2oqgA_	Alignment	not modelled	95.4	23 PDB header: transcription Chain: A: PDB Molecule: possible transcriptional regulator, arsr family protein; PDBTitle: arsr-like transcriptional regulator from rhodococcus sp. rha1
109	c3nqoB_	Alignment	not modelled	95.4	16 PDB header: transcription Chain: B: PDB Molecule: marr-family transcriptional regulator; PDBTitle: crystal structure of a marr family transcriptional regulator (cd1569)2 from clostridium difficile 630 at 2.20 a resolution
110	d2bv6a1	Alignment	not modelled	95.4	15 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
111	d1p4xa2	Alignment	not modelled	95.4	21 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
112	c3f6tA_	Alignment	not modelled	95.3	18 PDB header: transferase Chain: A: PDB Molecule: aspartate aminotransferase; PDBTitle: crystal structure of aspartate aminotransferase (e.c. 2.6.1.1)2 (yp_194538.1) from lactobacillus acidophilus ncfm at 2.15 a3 resolution
113	c3tgnA_	Alignment	not modelled	95.3	26 PDB header: transcription Chain: A: PDB Molecule: adc operon repressor adcr; PDBTitle: crystal structure of the zinc-dependent marr family transcriptional2 regulator adcr in the zn(ii)-bound state
114	d2fbia1	Alignment	not modelled	95.3	13 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
115	c3l09B_	Alignment	not modelled	95.3	20 PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of putative transcriptional regulator2 (jann_22dec04_contig27_revised_gene3569) from jannaschia sp. ccs1 at3 2.81 a resolution
116	d1jgsa_	Alignment	not modelled	95.3	18 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
117	d2nwia1	Alignment	not modelled	95.2	11 Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: AF1396-like
118	c3f6vA_	Alignment	not modelled	95.2	24 PDB header: transcription regulator Chain: A: PDB Molecule: possible transcriptional regulator, arsr family PDBTitle: crystal structure of possible transcriptional regulator for2 arsenical resistance
119	c1r22B_	Alignment	not modelled	95.2	24 PDB header: transcription repressor Chain: B: PDB Molecule: transcriptional repressor smtb; PDBTitle: crystal structure of the cyanobacterial metallothionein2 repressor smtb (c14s/c61s/c121s mutant) in the zn2alpha5-3 form
120	c3oopA_	Alignment	not modelled	95.2	15 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lin2960 protein; PDBTitle: the structure of a protein with unknown function from listeria innocua2 clip11262