




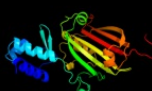









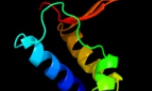
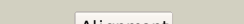

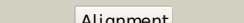

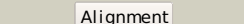






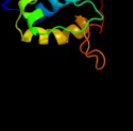

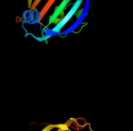
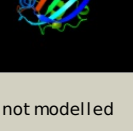


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3f8mA	 Alignment		100.0	22	PDB header: transcription Chain: A: PDB Molecule: gntr-family protein transcriptional regulator; PDBTitle: crystal structure of phnf from mycobacterium smegmatis
2	c3bwgA	 Alignment		100.0	19	PDB 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
3	c3edpB	 Alignment		100.0	22	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
4	c2wv0H	 Alignment		100.0	24	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
5	c3eetA	 Alignment		100.0	15	PDB header: transcription regulator Chain: A: PDB Molecule: putative gntr-family transcriptional regulator; PDBTitle: crystal structure of putative gntr-family transcriptional2 regulator
6	c3by6C	 Alignment		99.9	28	PDB header: transcription regulator Chain: C: PDB Molecule: predicted transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator from oenococcus oeni
7	c3neuA	 Alignment		99.9	19	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
8	c2du9A	 Alignment		99.9	30	PDB header: transcription Chain: A: PDB Molecule: predicted transcriptional regulators; PDBTitle: crystal structure of the transcriptional factor from c.glutamicum
9	d2oia1	 Alignment		99.9	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
10	c3f8IC	 Alignment		99.9	19	PDB header: transcription Chain: C: PDB Molecule: hth-type transcriptional repressor phnf; PDBTitle: crystal structure of the effector domain of phnf from mycobacterium2 smegmatis
11	d2oga1	 Alignment		99.9	12	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	d2fa1a1	Alignment		99.9	15	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
14	d2ikka1	Alignment		99.9	29	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
15	c3hfiA_	Alignment		99.9	15	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative regulator; PDBTitle: the crystal structure of the putative regulator from escherichia coli2 cft073
16	d3bwga2	Alignment		99.8	17	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
17	dlv4ra1	Alignment		99.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
18	d2p19a1	Alignment		99.8	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
19	c2ra5A_	Alignment		99.8	18	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of the putative transcriptional regulator2 from streptomyces coelicolor
20	d2ra5a1	Alignment		99.8	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
21	d2pkha1	Alignment	not modelled	99.8	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
22	c2di3A_	Alignment	not modelled	99.8	23	PDB header: transcription Chain: A: PDB Molecule: bacterial regulatory proteins, gntR family; PDBTitle: crystal structure of the transcriptional factor cgl29152 from corynebacterium glutamicum
23	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
24	c3c7jA_	Alignment	not modelled	99.8	25	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
25	c3fmsA_	Alignment	not modelled	99.8	22	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, gntR family; PDBTitle: crystal structure of tm0439, a gntR transcriptional2 regulator
26	d3bwga1	Alignment	not modelled	99.8	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
27	d1hw1a1	Alignment	not modelled	99.8	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
28	c3ihuA_	Alignment	not modelled	99.8	22	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
						PDB header: structural genomics, unknown function

29	c3ic7A_	Alignment	not modelled	99.8	29	Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of putative transcriptional regulator of gnt ^r family2 from bacteroides thetaiotaomicron
30	c2hs5A_	Alignment	not modelled	99.8	28	PDB header: transcription regulator Chain: A: PDB Molecule: putative transcriptional regulator gnt ^r ; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator gnt ^r from rhodococcus sp. rha1
31	c3tqnC_	Alignment	not modelled	99.8	26	PDB header: transcription Chain: C: PDB Molecule: transcriptional regulator, gnt ^r family; PDBTitle: structure of the transcriptional regulator of the gnt ^r family, from2 coxiella burnetii.
32	d2hs5a1	Alignment	not modelled	99.8	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Gnt ^r -like transcriptional regulators
33	c3lheA_	Alignment	not modelled	99.7	14	PDB header: transcription regulator Chain: A: PDB Molecule: gnt ^r family transcriptional regulator; PDBTitle: the crystal structure of the c-terminal domain of a gnt ^r 2 family transcriptional regulator from bacillus anthracis3 str. Sterne
34	d3ddva1	Alignment	not modelled	99.7	16	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
35	c2h09A_	Alignment	not modelled	98.3	16	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	98.1	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
37	c1f5tA_	Alignment	not modelled	97.9	12	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	97.9	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Heat-inducible transcription repressor HrcA, N-terminal domain
39	c2it0A_	Alignment	not modelled	97.8	16	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
40	d1jhfa1	Alignment	not modelled	97.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: LexA repressor, N-terminal DNA-binding domain
41	d1g3wa1	Alignment	not modelled	97.6	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
42	c3hruA_	Alignment	not modelled	97.6	22	PDB header: transcription Chain: A: PDB Molecule: metalloregulator scar; PDBTitle: crystal structure of scar with bound zn2+
43	c3ctaA_	Alignment	not modelled	97.5	15	PDB header: transferase Chain: A: PDB Molecule: riboflavin kinase; PDBTitle: crystal structure of riboflavin kinase from thermoplasma2 acidophilum
44	d2isya1	Alignment	not modelled	97.5	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
45	c2ewnA_	Alignment	not modelled	97.4	18	PDB header: ligase, transcription Chain: A: PDB Molecule: bira bifunctional protein; PDBTitle: ecoli biotin repressor with co-repressor analog
46	d1i5za1	Alignment	not modelled	97.4	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
47	d1j5ya1	Alignment	not modelled	97.4	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
48	c1g3wA_	Alignment	not modelled	97.3	12	PDB header: gene regulation Chain: A: PDB Molecule: diphtheria toxin repressor; PDBTitle: cd-cys102ser dtxr
49	c1j5yA_	Alignment	not modelled	97.3	19	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	22	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator; PDBTitle: possible transcriptional regulator from methanosarcina mazei go1 (gi2 21227196)
51	c3k69A_	Alignment	not modelled	97.2	21	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
52	c2v79B_	Alignment	not modelled	97.0	15	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
53	c2y75F_	Alignment	not modelled	97.0	17	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	d2h6ca1	Alignment	not modelled	96.9	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
55	c3lwfD_	Alignment	not modelled	96.9	15	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
56	c1fx7C_	Alignment	not modelled	96.9	17	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	d2gaua1	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
58	c1zybA_	Alignment	not modelled	96.7	12	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
59	c2x4hA_	Alignment	not modelled	96.7	13	PDB header: transcription Chain: A: PDB Molecule: hypothetical protein sso2273; PDBTitle: crystal structure of the hypothetical protein sso2273 from2 sulfolobus solfataricus
60	d2oz6a1	Alignment	not modelled	96.7	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
61	d1xd7a_	Alignment	not modelled	96.7	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
62	d2coha1	Alignment	not modelled	96.5	31	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
63	c3d0sA_	Alignment	not modelled	96.5	25	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulatory protein; PDBTitle: camp receptor protein from m.tuberculosis, camp-free form
64	d2d1ha1	Alignment	not modelled	96.5	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
65	d1ft9a1	Alignment	not modelled	96.5	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
66	c2gauA_	Alignment	not modelled	96.5	19	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, mcs9
67	c2oz6A_	Alignment	not modelled	96.5	17	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
68	c1ft9A_	Alignment	not modelled	96.4	19	PDB header: transcription Chain: A: PDB Molecule: carbon monoxide oxidation system PDBTitle: structure of the reduced (feii) co-sensing protein from r.2 rubrum
69	c2ev5B_	Alignment	not modelled	96.3	16	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator mntr; PDBTitle: bacillus subtilis manganese transport regulator (mntr)2 bound to calcium
70	d1sfxa_	Alignment	not modelled	96.3	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
71	d1ku9a_	Alignment	not modelled	96.3	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding protein Mj223
72	c3e97A_	Alignment	not modelled	96.3	18	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
73	c3e6dA_	Alignment	not modelled	96.3	14	PDB header: transcription regulation Chain: A: PDB Molecule: cyclic nucleotide-binding protein; PDBTitle: crystal structure of cprk c200s
74	d2bgca1	Alignment	not modelled	96.2	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
75	d3e5ua1	Alignment	not modelled	96.2	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
76	d2ev0a1	Alignment	not modelled	96.2	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
77	d1ylfa1	Alignment	not modelled	96.2	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
78	c2zy3A_	Alignment	not modelled	96.2	10	PDB header: lyase Chain: A: PDB Molecule: l-aspartate beta-decarboxylase; PDBTitle: dodecameric l-aspartate beta-decarboxylase
79	c3kccA_	Alignment	not modelled	96.1	17	PDB header: transcription Chain: A: PDB Molecule: catabolite gene activator; PDBTitle: crystal structure of d138l mutant of catabolite gene activator protein
80	c3iwzB_	Alignment	not modelled	96.1	20	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
81	c2l4aA_	Alignment	not modelled	96.1	19	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
82	c3dv8A_	Alignment	not modelled	96.1	17	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
83	d3broa1	Alignment	not modelled	96.1	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
84	c2zcwA_	Alignment	not modelled	96.1	31	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
85	c2vn2B_	Alignment	not modelled	96.1	19	PDB header: replication Chain: B: PDB Molecule: chromosome replication initiation protein; PDBTitle: crystal structure of the n-terminal domain of dnaD protein2 from geobacillus kaustophilus hta426
86	c2fmyB_	Alignment	not modelled	96.0	17	PDB header: dna binding protein Chain: B: PDB Molecule: carbon monoxide oxidation system transcription regulator PDBTitle: co-dependent transcription factor cooA from carboxydothermus2 hydrogenoformans (imidazole-bound form)
87	d2fxaa1	Alignment	not modelled	96.0	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
88	c1zreB_	Alignment	not modelled	96.0	17	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
89	d3ctaa1	Alignment	not modelled	96.0	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
90	c1stzB_	Alignment	not modelled	96.0	14	PDB header: transcription Chain: B: PDB Molecule: heat-inducible transcription repressor hrca homolog; PDBTitle: crystal structure of a hypothetical protein at 2.2 a resolution
91	d2nwia1	Alignment	not modelled	95.9	15	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: AF1396-like
92	c2zdbA_	Alignment	not modelled	95.9	17	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
93	c3k2zA_	Alignment	not modelled	95.9	15	PDB header: hydrolase Chain: A: PDB Molecule: lexa repressor; PDBTitle: crystal structure of a lexa protein from thermotoga maritima
94	d2b0la1	Alignment	not modelled	95.8	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CodY HTH domain
95	c3bj6B_	Alignment	not modelled	95.8	15	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of marr family transcription regulator sp03579
96	d2cg4a1	Alignment	not modelled	95.8	26	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
97	c3nqoB_	Alignment	not modelled	95.8	15	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
98	c3tgnA_	Alignment	not modelled	95.7	38	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
99	d2bv6a1	Alignment	not modelled	95.7	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
100	c2qwwB_	Alignment	not modelled	95.7	9	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
101	c1r22B_	Alignment	not modelled	95.7	21	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
102	c2gxgA_	Alignment	not modelled	95.6	22	PDB header: transcription Chain: A: PDB Molecule: 146aa long hypothetical transcriptional regulator; PDBTitle: crystal structure of emrr homolog from hyperthermophilic archaea2 sulfobolus tokodaii strain7
103	d1r1ua_	Alignment	not modelled	95.6	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: ArsR-like transcriptional regulators
104	c3g3zA_	Alignment	not modelled	95.6	14	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, marr family; PDBTitle: the structure of nmb1585, a marr family regulator from neisseria2 meningitidis
105	c3la2A_	Alignment	not modelled	95.6	14	PDB header: transcription Chain: A: PDB Molecule: global nitrogen regulator; PDBTitle: crystal structure of ntca in complex with 2-oxoglutarate
						Fold: DNA/RNA-binding 3-helical bundle

106	dli1ga1	Alignment	not modelled	95.6	21	Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
107	c3f6vA	Alignment	not modelled	95.5	32	PDB header: transcription regulator Chain: A: PDB Molecule: possible transcriptional regulator, arsr family PDBTitle: crystal structure of possible transcriptional regulator for2 arsenical resistance
108	d1p4xa2	Alignment	not modelled	95.5	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
109	c2kkoB	Alignment	not modelled	95.5	20	PDB header: dna binding protein Chain: B: PDB Molecule: possible transcriptional regulatory protein PDBTitle: solution nmr structure of the homodimeric winged helix-turn-2 helix dna-binding domain (fragment 1-100) mb0332 from3 mycobacterium bovis, a possible arsr-family transcriptional4 regulator. northeast structural genomics consortium target5 mbr242e.
110	d2cyya1	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
111	d2fbha1	Alignment	not modelled	95.5	24	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
112	d2hr3a1	Alignment	not modelled	95.5	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
113	c2fxaB	Alignment	not modelled	95.5	14	PDB header: transcription Chain: B: PDB Molecule: protease production regulatory protein hpr; PDBTitle: structure of the protease production regulatory protein hpr from2 bacillus subtilis.
114	d1pdnc	Alignment	not modelled	95.5	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Paired domain
115	c2oqgA	Alignment	not modelled	95.4	29	PDB header: transcription Chain: A: PDB Molecule: possible transcriptional regulator, arsr family protein; PDBTitle: arsr-like transcriptional regulator from rhodococcus sp. rha1
116	c2w48D	Alignment	not modelled	95.4	17	PDB header: transcription Chain: D: PDB Molecule: sorbitol operon regulator; PDBTitle: crystal structure of the full-length sorbitol operon2 regulator sorc from klebsiella pneumoniae
117	d1ub9a	Alignment	not modelled	95.4	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
118	c2wteB	Alignment	not modelled	95.3	24	PDB header: antiviral protein Chain: B: PDB Molecule: csa3; PDBTitle: the structure of the crispr-associated protein, csa3, from2 sulfolobus solfataricus at 1.8 angstrom resolution.
119	c2rdpA	Alignment	not modelled	95.2	16	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator marr; PDBTitle: the structure of a marr family protein from bacillus2 stearothermophilus
120	c2bgcA	Alignment	not modelled	95.2	18	PDB header: transcription Chain: A: PDB Molecule: prfa; PDBTitle: prfa-g145s, a constitutive active mutant of the2 transcriptional regulator in l.monocytogenes