



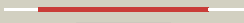




























#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3bwgA_	 Alignment		100.0	16	PDB header: transcription regulator Chain: A: PDB Molecule: uncharacterized hth-type transcriptional regulator yydk; PDBTitle: the crystal structure of possible transcriptional regulator yydk from2 bacillus subtilis subsp. subtilis str. 168
2	c3edpB_	 Alignment		100.0	13	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	18	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	12	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	22	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	d2ooa1	 Alignment		99.9	10	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
7	d2fa1a1	 Alignment		99.9	10	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
8	c3f8lC_	 Alignment		99.9	15	PDB header: transcription Chain: C: PDB Molecule: hth-type transcriptional repressor phnf; PDBTitle: crystal structure of the effector domain of phnf from mycobacterium2 smegmatis
9	d3cnva1	 Alignment		99.9	12	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
10	d2p19a1	 Alignment		99.9	16	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
11	c2ra5A_	 Alignment		99.9	18	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of the putative transcriptional regulator2 from streptomyces coelicolor

12	d2ra5a1	Alignment		99.9	18	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
13	d2ogga1	Alignment		99.9	8	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
14	d3bwga2	Alignment		99.9	14	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
15	d2ikka1	Alignment		99.8	13	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
16	c3hfiA	Alignment		99.8	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
17	c3by6C	Alignment		99.8	22	PDB header: transcription regulator Chain: C: PDB Molecule: predicted transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator from oenococcus oeni
18	c2du9A	Alignment		99.8	22	PDB header: transcription Chain: A: PDB Molecule: predicted transcriptional regulators; PDBTitle: crystal structure of the transcriptional factor from c. glutamicum
19	c1e2xA	Alignment		99.8	29	PDB header: transcriptional regulation Chain: A: PDB Molecule: fatty acid metabolism regulator protein; PDBTitle: fadr, fatty acid responsive transcription factor from e.2 coli
20	c3neuA	Alignment		99.8	21	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
21	c2di3A	Alignment	not modelled	99.8	25	PDB header: transcription Chain: A: PDB Molecule: bacterial regulatory proteins, gntr family; PDBTitle: crystal structure of the transcriptional factor cgl29152 from corynebacterium glutamicum
22	d2pkha1	Alignment	not modelled	99.8	13	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
23	d1v4ra1	Alignment	not modelled	99.8	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Gntr-like transcriptional regulators
24	c3c7jA	Alignment	not modelled	99.8	31	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	c3ihuA	Alignment	not modelled	99.8	25	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
26	d1hw1a1	Alignment	not modelled	99.8	34	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Gntr-like transcriptional regulators
27	c3fmsA	Alignment	not modelled	99.8	32	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, gntr family; PDBTitle: crystal structure of tm0439, a gntr transcriptional2 regulator
28	c3lhaA	Alignment	not modelled	99.8	14	PDB header: transcription regulator 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
						PDB header: transcription regulator

29	c2hs5A	Alignment	not modelled	99.8	28	Chain: A: PDB Molecule: putative transcriptional regulator gntR; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator gntR from rhodococcus sp. rha1
30	d3bwga1	Alignment	not modelled	99.8	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
31	d3ddva1	Alignment	not modelled	99.7	12	Fold: Chorismate lyase-like Superfamily: Chorismate lyase-like Family: UTRA domain
32	c3ic7A	Alignment	not modelled	99.7	20	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
33	d2hs5a1	Alignment	not modelled	99.7	28	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: GntR-like transcriptional regulators
34	c3tqnC	Alignment	not modelled	99.7	21	PDB header: transcription Chain: C: PDB Molecule: transcriptional regulator, gntR family; PDBTitle: structure of the transcriptional regulator of the gntR family, from2 coxiella burnetii.
35	c2h09A	Alignment	not modelled	98.1	20	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	17	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	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
39	d1stza1	Alignment	not modelled	97.8	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Heat-inducible transcription repressor HrcA, N-terminal domain
40	d1jhfa1	Alignment	not modelled	97.7	15	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	97.5	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
42	c3hruA	Alignment	not modelled	97.5	18	PDB header: transcription Chain: A: PDB Molecule: metalloregulator scar; PDBTitle: crystal structure of scar with bound zn2+
43	c3ctaA	Alignment	not modelled	97.4	12	PDB header: transferase Chain: A: PDB Molecule: riboflavin kinase; PDBTitle: crystal structure of riboflavin kinase from thermoplasma2 acidophilum
44	d2isya1	Alignment	not modelled	97.4	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
45	d1j5ya1	Alignment	not modelled	97.3	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
46	c3k69A	Alignment	not modelled	97.3	19	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
47	c2ewnA	Alignment	not modelled	97.2	19	PDB header: ligase, transcription Chain: A: PDB Molecule: bira bifunctional protein; PDBTitle: ecoli biotin repressor with co-repressor analog
48	c1fx7C	Alignment	not modelled	97.2	21	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
49	c2y75F	Alignment	not modelled	97.2	10	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
50	c1g3wA	Alignment	not modelled	97.2	17	PDB header: gene regulation Chain: A: PDB Molecule: diphtheria toxin repressor; PDBTitle: cd-cys102ser dtxr
51	c3lwfD	Alignment	not modelled	97.2	12	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
52	c1j5yA	Alignment	not modelled	97.2	13	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
53	d1i5za1	Alignment	not modelled	97.1	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
54	d1xd7a	Alignment	not modelled	97.0	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
55	c3r0aB	Alignment	not modelled	97.0	18	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator; PDBTitle: possible transcriptional regulator from methanosarcina

					mazei go1 (gi2 21227196)
56	c2v79B_	Alignment	not modelled	96.8	17 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
57	d2gaua1	Alignment	not modelled	96.6	16 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
58	c1ft9A_	Alignment	not modelled	96.5	19 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
59	c2x4hA_	Alignment	not modelled	96.4	16 PDB header: transcription Chain: A: PDB Molecule: hypothetical protein sso2273; PDBTitle: crystal structure of the hypothetical protein sso2273 from2 sulfobolbus solfataricus
60	c1zybA_	Alignment	not modelled	96.4	11 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
61	d2h6ca1	Alignment	not modelled	96.4	28 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
62	d1ft9a1	Alignment	not modelled	96.4	20 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
63	d2oz6a1	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
64	c2gauA_	Alignment	not modelled	96.4	15 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
65	c3d0sA_	Alignment	not modelled	96.3	22 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulatory protein; PDBTitle: camp receptor protein from m.tuberculosis, camp-free form
66	c2ev5B_	Alignment	not modelled	96.3	10 PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator mntr; PDBTitle: bacillus subtilis manganese transport regulator (mntr)2 bound to calcium
67	c2oz6A_	Alignment	not modelled	96.2	23 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	c2l4aA_	Alignment	not modelled	96.2	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
69	d2coha1	Alignment	not modelled	96.2	25 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
70	d1ylfa1	Alignment	not modelled	96.1	9 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
71	c3e97A_	Alignment	not modelled	96.1	17 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
72	d2cg4a1	Alignment	not modelled	96.1	16 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
73	d2d1ha1	Alignment	not modelled	96.1	16 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
74	d1sfxa_	Alignment	not modelled	96.0	15 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
75	c3e6dA_	Alignment	not modelled	96.0	29 PDB header: transcription regulation Chain: A: PDB Molecule: cyclic nucleotide-binding protein; PDBTitle: crystal structure of cprk c200s
76	d3e5ua1	Alignment	not modelled	95.9	17 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
77	c2zcwA_	Alignment	not modelled	95.9	25 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
78	c3kcca_	Alignment	not modelled	95.9	25 PDB header: transcription Chain: A: PDB Molecule: catabolite gene activator; PDBTitle: crystal structure of d138l mutant of catabolite gene activator protein
79	d2bgca1	Alignment	not modelled	95.9	25 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: CAP C-terminal domain-like
80	c2vn2B_	Alignment	not modelled	95.8	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
					Fold: DNA/RNA-binding 3-helical bundle

81	d2ev0a1	Alignment	not modelled	95.8	10	Superfamily: "Winged helix" DNA-binding domain Family: Iron-dependent repressor protein
82	c3iwzB	Alignment	not modelled	95.8	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
83	d2cyya1	Alignment	not modelled	95.6	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
84	c2zy3A	Alignment	not modelled	95.6	12	PDB header: lyase Chain: A: PDB Molecule: l-aspartate beta-decarboxylase; PDBTitle: dodecameric l-aspartate beta-decarboxylase
85	c2fmyB	Alignment	not modelled	95.6	17	PDB header: dna binding protein Chain: B: PDB Molecule: carbon monoxide oxidation system transcription regulator PDBTitle: co-dependent transcription factor cooa from carboxydotherrmus2 hydrogenofomans (imidazole-bound form)
86	c2zdbA	Alignment	not modelled	95.6	22	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
87	c1zreB	Alignment	not modelled	95.6	25	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
88	c2e7xA	Alignment	not modelled	95.6	24	PDB header: transcription regulator Chain: A: PDB Molecule: 150aa long hypothetical transcriptional regulator; PDBTitle: structure of the lrp/asnc like transcriptional regulator from2 sulfobolus tokodaii 7 complexed with its cognate ligand
89	d1i1ga1	Alignment	not modelled	95.6	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
90	d1ku9a	Alignment	not modelled	95.6	23	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding protein Mj223
91	c3dv8A	Alignment	not modelled	95.6	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
92	c3bj6B	Alignment	not modelled	95.6	20	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of marr family transcription regulator sp03579
93	c2cg4B	Alignment	not modelled	95.5	17	PDB header: transcription Chain: B: PDB Molecule: regulatory protein asnc; PDBTitle: structure of e.coli asnc
94	c3i4pA	Alignment	not modelled	95.4	20	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, asnc family; PDBTitle: crystal structure of asnc family transcriptional regulator from2 agrobacterium tumefaciens
95	d3broa1	Alignment	not modelled	95.4	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
96	c2oqgA	Alignment	not modelled	95.4	18	PDB header: transcription Chain: A: PDB Molecule: possible transcriptional regulator, arsr family protein; PDBTitle: arsr-like transcriptional regulator from rhodococcus sp. rha1
97	d2bv6a1	Alignment	not modelled	95.4	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
98	c3k2zA	Alignment	not modelled	95.4	25	PDB header: hydrolase Chain: A: PDB Molecule: lexa repressor; PDBTitle: crystal structure of a lexa protein from thermotoga maritima
99	c1r22B	Alignment	not modelled	95.4	20	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
100	d2cfxa1	Alignment	not modelled	95.4	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
101	d3ctaa1	Alignment	not modelled	95.3	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
102	c3f6tA	Alignment	not modelled	95.3	15	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
103	d2fxaa1	Alignment	not modelled	95.3	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
104	c2e1cA	Alignment	not modelled	95.2	22	PDB header: transcription/dna Chain: A: PDB Molecule: putative hth-type transcriptional regulator ph1519; PDBTitle: structure of putative hth-type transcriptional regulator ph1519/dna2 complex
105	c3la2A	Alignment	not modelled	95.2	26	PDB header: transcription Chain: A: PDB Molecule: global nitrogen regulator; PDBTitle: crystal structure of ntca in complex with 2-oxoglutarate
						PDB header: transcription

106	c2qwwB_	Alignment	not modelled	95.2	15	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	c3tgnA_	Alignment	not modelled	95.1	18	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
108	c2p6tH_	Alignment	not modelled	95.1	22	PDB header: transcription Chain: H: PDB Molecule: transcriptional regulator, lrp/asnc family; PDBTitle: crystal structure of transcriptional regulator nmb0573 and l-leucine2 complex from neisseria meningitidis
109	c2dbbA_	Alignment	not modelled	95.1	24	PDB header: transcriptional regulator Chain: A: PDB Molecule: putative hth-type transcriptional regulator ph0061; PDBTitle: crystal structure of ph0061
110	c3nqoB_	Alignment	not modelled	95.1	13	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
111	c2gxgA_	Alignment	not modelled	95.1	11	PDB header: transcription Chain: A: PDB Molecule: 146aa long hypothetical transcriptional regulator; PDBTitle: crystal structure of emrr homolog from hyperthermophilic archaea2 sulfobolus tokodaii strain7
112	c2ia0A_	Alignment	not modelled	95.1	24	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative hth-type transcriptional regulator pf0864; PDBTitle: transcriptional regulatory protein pf0864 from pyrococcus furiosus a2 member of the asnc family (pf0864)
113	d1ub9a_	Alignment	not modelled	95.0	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
114	c2vbzA_	Alignment	not modelled	95.0	18	PDB header: dna-binding protein Chain: A: PDB Molecule: transcriptional regulatory protein; PDBTitle: feast or famine regulatory protein (rv3291c)from m.2 tuberculosis complexed with l-tryptophan
115	d2fbha1	Alignment	not modelled	95.0	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
116	d1p4xa2	Alignment	not modelled	95.0	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
117	d1r1ua_	Alignment	not modelled	95.0	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: ArsR-like transcriptional regulators
118	d1mkma1	Alignment	not modelled	95.0	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator lclR, N-terminal domain
119	c2cfxD_	Alignment	not modelled	95.0	22	PDB header: transcription Chain: D: PDB Molecule: hth-type transcriptional regulator lrpcc; PDBTitle: structure of b.subtilis lrpcc
120	c2fxaB_	Alignment	not modelled	94.9	17	PDB header: transcription Chain: B: PDB Molecule: protease production regulatory protein hpr; PDBTitle: structure of the protease production regulatory protein hpr from2 bacillus subtilis.