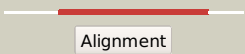
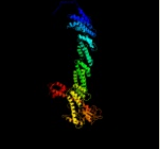

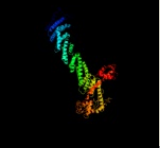
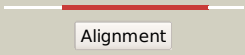

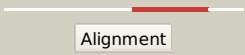

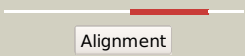
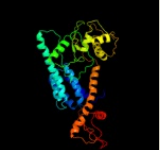
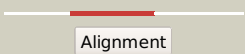

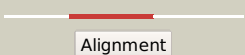

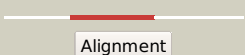

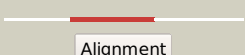








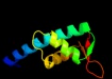






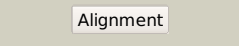
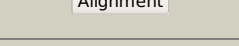
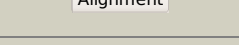
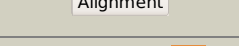


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2hyeC_	 Alignment		100.0	85	PDB header: protein binding Chain: C: PDB Molecule: cullin-4a; PDBTitle: crystal structure of the ddb1-cul4a-rbx1-sv5v complex
2	c4a0IH_	 Alignment		100.0	98	PDB header: ligase/dna-binding protein/dna Chain: H: PDB Molecule: cullin-4b; PDBTitle: structure of ddb1-ddb2-cul4b-rbx1 bound to a 12 bp abasic2 site containing dna-duplex
3	c1u6gA_	 Alignment		100.0	33	PDB header: ligase Chain: A: PDB Molecule: cullin homolog 1; PDBTitle: crystal structure of the cand1-cul1-roc1 complex
4	c3dplC_	 Alignment		100.0	32	PDB header: ligase Chain: C: PDB Molecule: cullin-5; PDBTitle: structural insights into nedd8 activation of cullin-ring2 ligases: conformational control of conjugation.
5	c1ldkB_	 Alignment		100.0	37	PDB header: ligase Chain: B: PDB Molecule: cullin homolog; PDBTitle: structure of the cul1-rbx1-skp1-f boxskp2 scf ubiquitin2 ligase complex
6	d2hyec2	 Alignment		100.0	81	Fold: alpha-alpha superhelix Superfamily: Cullin repeat-like Family: Cullin repeat
7	c4apfB_	 Alignment		100.0	30	PDB header: cell cycle Chain: B: PDB Molecule: cullin 3; PDBTitle: crystal structure of the human klhl11-cul3 complex at 3.1a2 resolution
8	c2wzkA_	 Alignment		100.0	21	PDB header: protein binding Chain: A: PDB Molecule: cullin-5; PDBTitle: structure of the cul5 n-terminal domain at 2.05a resolution.
9	d1ldja2	 Alignment		100.0	28	Fold: alpha-alpha superhelix Superfamily: Cullin repeat-like Family: Cullin repeat
10	d2hyec3	 Alignment		100.0	88	Fold: Cullin homology domain Superfamily: Cullin homology domain Family: Cullin homology domain
11	c4eozD_	 Alignment		100.0	30	PDB header: protein binding Chain: D: PDB Molecule: cullin-3; PDBTitle: crystal structure of the spop btb domain complexed with the cul3 n-2 terminal domain

12	d1ldja3	Alignment		100.0	36	Fold: Cullin homology domain Superfamily: Cullin homology domain Family: Cullin homology domain
13	d2hyec1	Alignment		99.9	92	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: SCF ubiquitin ligase complex WHB domain
14	d1ldja1	Alignment		99.9	42	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: SCF ubiquitin ligase complex WHB domain
15	c3o2pE_	Alignment		99.8	33	PDB header: ligase, cell cycle Chain: E; PDB Molecule: cell division control protein 53; PDBTitle: a dual e3 mechanism for rub1 ligation to cdc53: dcn1(p)-cdc53(whb)
16	d1iuya_	Alignment		99.8	46	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: SCF ubiquitin ligase complex WHB domain
17	c2xv4S_	Alignment		95.5	14	PDB header: transcription Chain: S; PDB Molecule: dna-directed rna polymerase iii subunit rpc3; PDBTitle: structure of human rpc62 (partial)
18	c3lmmB_	Alignment		92.2	9	PDB header: structural genomics, unknown function Chain: B; PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of the dip2311 protein from2 corynebacterium diphtheriae, northeast structural genomics3 consortium target cdr35
19	d1p6ra_	Alignment		90.9	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Penicillinase repressor
20	c3g3zA_	Alignment		90.6	11	PDB header: transcription Chain: A; PDB Molecule: transcriptional regulator, marr family; PDBTitle: the structure of nmb1585, a marr family regulator from neisseria2 meningitidis
21	d1wi9a_	Alignment	not modelled	90.5	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: PCI domain (PINT motif)
22	c3cuoB_	Alignment	not modelled	89.3	14	PDB header: transcription regulator Chain: B; PDB Molecule: uncharacterized hth-type transcriptional regulator ygav; PDBTitle: crystal structure of the predicted dna-binding transcriptional2 regulator from e. coli
23	c3cjnA_	Alignment	not modelled	89.1	13	PDB header: transcription regulator Chain: A; PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of transcriptional regulator, marr family, from2 silicibacter pomeroyi
24	c3bpxB_	Alignment	not modelled	89.0	16	PDB header: transcription regulator Chain: B; PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of marr
25	c2qwwB_	Alignment	not modelled	88.4	13	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
26	d2etha1	Alignment	not modelled	88.2	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
27	d1mkma1	Alignment	not modelled	88.1	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator lclR, N-terminal domain
28	c1r22B_	Alignment	not modelled	88.0	22	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

29	c2k4bA	 Alignment	not modelled	87.9	16	PDB header: dna binding protein Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: copr repressor structure
30	d1ylfa1	 Alignment	not modelled	87.8	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
31	c4em1A	 Alignment	not modelled	87.8	24	PDB header: transcription Chain: A: PDB Molecule: uncharacterized hth-type transcriptional regulator sar2349; PDBTitle: staphylococcus aureus marr native
32	d1okra	 Alignment	not modelled	87.5	25	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Penicillinase repressor
33	c3nrvC	 Alignment	not modelled	87.3	15	PDB header: transcription regulator Chain: C: PDB Molecule: putative transcriptional regulator (marr/emrr family); PDBTitle: crystal structure of marr/emrr family transcriptional regulator from2 acinetobacter sp. adp1
34	c4fhtA	 Alignment	not modelled	87.2	16	PDB header: transcription Chain: A: PDB Molecule: pcav transcriptional regulator; PDBTitle: crystal structure of the pcav transcriptional regulator from2 streptomyces coelicolor in complex with its natural ligand
35	c2rdpA	 Alignment	not modelled	86.9	9	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator marr; PDBTitle: the structure of a marr family protein from bacillus2 stearothermophilus
36	d1lnwa	 Alignment	not modelled	86.8	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
37	c2oqgA	 Alignment	not modelled	86.7	11	PDB header: transcription Chain: A: PDB Molecule: possible transcriptional regulator, arsr family protein; PDBTitle: arsr-like transcriptional regulator from rhodococcus sp. rha1
38	c2nyxB	 Alignment	not modelled	86.7	15	PDB header: transcription Chain: B: PDB Molecule: probable transcriptional regulatory protein, rv1404; PDBTitle: crystal structure of rv1404 from mycobacterium tuberculosis
39	c3e6mD	 Alignment	not modelled	86.4	15	PDB header: transcription regulator Chain: D: PDB Molecule: marr family transcriptional regulator; PDBTitle: the crystal structure of a marr family transcriptional2 regulator from silicibacter pomeroyi dss.
40	d2frha1	 Alignment	not modelled	86.3	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
41	d2fbia1	 Alignment	not modelled	86.2	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
42	d1p4xa2	 Alignment	not modelled	86.2	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
43	c1p4xA	 Alignment	not modelled	86.0	18	PDB header: transcription Chain: A: PDB Molecule: staphylococcal accessory regulator a homologue; PDBTitle: crystal structure of sars protein from staphylococcus aureus
44	c3t8tA	 Alignment	not modelled	85.9	22	PDB header: unknown function Chain: A: PDB Molecule: staphylococcus aureus cymr (oxidized form); PDBTitle: crystal structure of staphylococcus aureus cymr oxidized form
45	c2xroE	 Alignment	not modelled	85.8	13	PDB header: dna-binding protein/dna Chain: E: PDB Molecule: hth-type transcriptional regulator ttgv; PDBTitle: crystal structure of ttgv in complex with its dna operator
46	d1z91a1	 Alignment	not modelled	85.5	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
47	c3pqkD	 Alignment	not modelled	85.4	25	PDB header: transcription Chain: D: PDB Molecule: biofilm growth-associated repressor; PDBTitle: crystal structure of the transcriptional repressor bigr from xylella2 fastidiosa
48	d1r1ta	 Alignment	not modelled	85.2	19	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: ArsR-like transcriptional regulators
49	c2jscB	 Alignment	not modelled	85.2	13	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator rv1994c/mt2050; PDBTitle: nmr structure of the cadmium metal-sensor cmtr from mycobacterium2 tuberculosis
50	c3bjaA	 Alignment	not modelled	85.1	13	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, marr family, putative; PDBTitle: crystal structure of putative marr-like transcription regulator2 (np_978771.1) from bacillus cereus atcc 10987 at 2.38 a resolution
51	c2nnnB	 Alignment	not modelled	85.1	11	PDB header: transcription Chain: B: PDB Molecule: probable transcriptional regulator; PDBTitle: crystal structure of probable transcriptional regulator from2 pseudomonas aeruginosa
52	c3oopA	 Alignment	not modelled	85.0	20	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
53	d1ub9a	 Alignment	not modelled	84.9	9	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators

54	c2fa5B_		Alignment	not modelled	84.9	12	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator marr/emrr family; PDBTitle: the crystal structure of an unliganded multiple antibiotic-2 resistance repressor (marr) from xanthomonas campestris
55	d1sfxa_		Alignment	not modelled	84.8	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
56	d2bv6a1		Alignment	not modelled	84.6	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
57	c3nqoB_		Alignment	not modelled	84.5	22	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
58	d2a61a1		Alignment	not modelled	84.4	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
59	c2kkoB_		Alignment	not modelled	84.0	24	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.
60	d1s3ja_		Alignment	not modelled	83.8	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
61	c1mkmA_		Alignment	not modelled	83.7	10	PDB header: transcription Chain: A: PDB Molecule: iclr transcriptional regulator; PDBTitle: crystal structure of the thermotoga maritima iclr
62	c2o0yB_		Alignment	not modelled	83.7	22	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of putative transcriptional regulator rha1_ro069532 (iclr-family) from rhodococcus sp.
63	c2fxaB_		Alignment	not modelled	83.5	13	PDB header: transcription Chain: B: PDB Molecule: protease production regulatory protein hpr; PDBTitle: structure of the protease production regulatory protein hpr from2 bacillus subtilis.
64	d1hsja1		Alignment	not modelled	83.5	20	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
65	c3f6vA_		Alignment	not modelled	83.4	12	PDB header: transcription regulator Chain: A: PDB Molecule: possible transcriptional regulator, arsr family PDBTitle: crystal structure of possible transcriptional regulator for2 arsenical resistance
66	c3bj6B_		Alignment	not modelled	82.9	15	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of marr family transcription regulator sp03579
67	c2zkzC_		Alignment	not modelled	82.7	14	PDB header: transcription Chain: C: PDB Molecule: transcriptional repressor pagr; PDBTitle: crystal structure of the transcriptional repressor pagr of bacillus2 anthracis
68	c3bddD_		Alignment	not modelled	82.7	14	PDB header: transcription Chain: D: PDB Molecule: regulatory protein marr; PDBTitle: crystal structure of a putative multiple antibiotic-resistance2 repressor (ssu05_1136) from streptococcus suis 89/1591 at 2.20 a3 resolution
69	c3k0IA_		Alignment	not modelled	82.3	20	PDB header: transcription regulator Chain: A: PDB Molecule: repressor protein; PDBTitle: crystal structure of putative marr family transcriptional2 regulator from acinetobacter sp. adp
70	d2bbya_		Alignment	not modelled	82.3	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding domain from rap30
71	c3jthA_		Alignment	not modelled	82.2	12	PDB header: transcription Chain: A: PDB Molecule: transcription activator hlyu; PDBTitle: crystal structure of a transcriptional regulator hlyu from2 vibrio vulnificus cmcp6
72	d3broa1		Alignment	not modelled	82.1	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
73	c2wteB_		Alignment	not modelled	82.1	15	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.
74	c3k69A_		Alignment	not modelled	82.1	11	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
75	d1jgsa_		Alignment	not modelled	82.0	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
76	d1ku9a_		Alignment	not modelled	81.9	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: DNA-binding protein Mj223
77	d2d1ha1		Alignment	not modelled	81.8	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: TrmB-like
78	c3r4kD_		Alignment	not modelled	81.3	21	PDB header: dna binding protein Chain: D: PDB Molecule: transcriptional regulator, iclr family; PDBTitle: crystal structure of a putative iclr transcriptional regulator2 (tm1040_3717) from silicibacter sp. tm1040 at 2.46 a

					resolution
79	c2pexA_	Alignment	not modelled	81.3	23 PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator ohrr; PDBTitle: structure of reduced c22s ohrr from xanthomonas campestris
80	d2fbha1	Alignment	not modelled	81.1	22 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
81	c3tgnA_	Alignment	not modelled	81.1	16 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
82	c3s2wB_	Alignment	not modelled	80.6	14 PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: the crystal structure of a marr transcriptional regulator from2 methanosarcina mazel go1
83	c2l02B_	Alignment	not modelled	80.6	27 PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein; PDBTitle: solution nmr structure of protein bt2368 from bacteroides2 thetaiotaomicron, northeast structural genomics consortium target3 btr375
84	c3cdhB_	Alignment	not modelled	80.5	7 PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of the marr family transcriptional regulator spo14532 from silicibacter pomeroyi dss-3
85	c4hblA_	Alignment	not modelled	80.4	18 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, marr family; PDBTitle: crystal structure of abfr of staphylococcus epidermidis
86	c2y75F_	Alignment	not modelled	79.8	19 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
87	d1biaa1	Alignment	not modelled	79.8	19 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Biotin repressor-like
88	c3zmdD_	Alignment	not modelled	79.6	11 PDB header: transcription Chain: D: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of absc, a marr family transcriptional2 regulator from streptomyces coelicolor
89	d1sd4a_	Alignment	not modelled	79.4	22 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Penicillinase repressor
90	c2g7uB_	Alignment	not modelled	79.0	29 PDB header: structural genomics, unknown function Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: 2.3 a structure of putative catechol degradative operon regulator from2 rhodococcus sp. rha1
91	d1fy7a_	Alignment	not modelled	78.7	22 Fold: Acyl-CoA N-acyltransferases (Nat) Superfamily: Acyl-CoA N-acyltransferases (Nat) Family: N-acetyl transferase, NAT
92	c3f6oB_	Alignment	not modelled	78.6	11 PDB header: transcription regulator Chain: B: PDB Molecule: probable transcriptional regulator, arsr family PDBTitle: crystal structure of arsr family transcriptional regulator,2 rha00566
93	c3f3xA_	Alignment	not modelled	78.3	13 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, marr family, putative; PDBTitle: crystal structure of the transcriptional regulator bldr2 from sulfolobus solfataricus
94	c2lkpB_	Alignment	not modelled	77.8	19 PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, arsr family; PDBTitle: solution structure of apo-nmtr
95	d1sxje1	Alignment	not modelled	77.7	19 Fold: post-AAA+ oligomerization domain-like Superfamily: post-AAA+ oligomerization domain-like Family: DNA polymerase III clamp loader subunits, C-terminal domain
96	c3hrmA_	Alignment	not modelled	77.2	16 PDB header: transcription regulator Chain: A: PDB Molecule: hth-type transcriptional regulator sarz; PDBTitle: crystal structure of staphylococcus aureus protein sarz in sulfenic2 acid form
97	d1rlua_	Alignment	not modelled	76.9	14 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: ArsR-like transcriptional regulators
98	d1lj9a_	Alignment	not modelled	76.1	14 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
99	d2cg4a1	Alignment	not modelled	76.1	15 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
100	d3deua1	Alignment	not modelled	75.8	22 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
101	c3lwfD_	Alignment	not modelled	75.5	13 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
102	c2vn2B_	Alignment	not modelled	75.2	23 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
103	d2fxaa1	Alignment	not modelled	74.5	12 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: MarR-like transcriptional regulators
104	d1p4xa1	Alignment	not modelled	74.2	9 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain

					Family: MarR-like transcriptional regulators
105	c1hsjA_	Alignment	not modelled	73.9	25 PDB header: transcription/sugar binding protein Chain: A: PDB Molecule: fusion protein consisting of staphylococcus PDBTitle: sarr mbp fusion structure
106	c2fsgA_	Alignment	not modelled	73.6	14 PDB header: protein transport Chain: A: PDB Molecule: preprotein translocase seca subunit; PDBTitle: complex seca:atp from escherichia coli
107	d1ilga1	Alignment	not modelled	73.2	17 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Lrp/AsnC-like transcriptional regulator N-terminal domain
108	c4b8xB_	Alignment	not modelled	73.2	19 PDB header: transcription Chain: B: PDB Molecule: possible marr-transcriptional regulator; PDBTitle: near atomic resolution crystal structure of sco5413, a marr family2 transcriptional regulator from streptomyces coelicolor
109	c3zplE_	Alignment	not modelled	72.6	15 PDB header: transcription/dna Chain: E: PDB Molecule: putative marr-family transcriptional repressor; PDBTitle: crystal structure of sco3205, a marr family transcriptional2 regulator from streptomyces coelicolor, in complex with dna
110	d2g9wa1	Alignment	not modelled	72.4	7 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Penicillinase repressor
111	c3deuB_	Alignment	not modelled	72.3	22 PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator slyA; PDBTitle: crystal structure of transcription regulatory protein slyA2 from salmonella typhimurium in complex with salicylate3 ligands
112	c2l4aA_	Alignment	not modelled	71.2	12 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
113	c2gxgA_	Alignment	not modelled	70.9	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
114	c4ijaA_	Alignment	not modelled	70.3	15 PDB header: protein binding Chain: A: PDB Molecule: xylr protein; PDBTitle: structure of s. aureus methicillin resistance factor mecR2
115	c2ia2D_	Alignment	not modelled	70.2	22 PDB header: transcription Chain: D: PDB Molecule: putative transcriptional regulator; PDBTitle: the crystal structure of a putative transcriptional regulator rha061952 from rhodococcus sp. rha1
116	c3kp3B_	Alignment	not modelled	70.0	20 PDB header: transcription regulator/antibiotic Chain: B: PDB Molecule: transcriptional regulator tcar; PDBTitle: staphylococcus epidermidis in complex with ampicillin
117	c2z99A_	Alignment	not modelled	69.0	16 PDB header: cell cycle Chain: A: PDB Molecule: putative uncharacterized protein; PDBTitle: crystal structure of scpb from mycobacterium tuberculosis
118	d1xd7a_	Alignment	not modelled	68.7	17 Fold: DNA/RNA-binding 3-helical bundle Superfamily: "Winged helix" DNA-binding domain Family: Transcriptional regulator Rrf2
119	c3fm5D_	Alignment	not modelled	66.3	11 PDB header: transcription regulator Chain: D: PDB Molecule: transcriptional regulator; PDBTitle: x-ray crystal structure of transcriptional regulator (marr family)2 from rhodococcus sp. rha1
120	c4hf1B_	Alignment	not modelled	65.9	19 PDB header: transcription/dna Chain: B: PDB Molecule: hth-type transcriptional regulator iscr; PDBTitle: crystal structure of iscr bound to its promoter