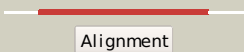

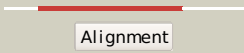


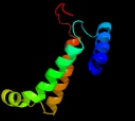

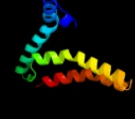


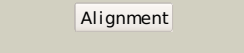

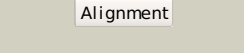

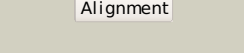



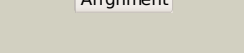

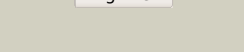









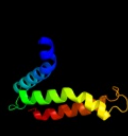


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1pb6a2	 Alignment		99.8	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
2	d2fx0a2	 Alignment		99.7	11	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
3	c3npiB_	 Alignment		99.4	21	PDB header: gene regulation Chain: B: PDB Molecule: tetr family regulatory protein; PDBTitle: crystal structure of a tetr family regulatory protein (dip1788) from <i>corynebacterium diphtheriae</i> at 2.96 a resolution
4	c2qtqB_	 Alignment		98.8	12	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of a predicted dna-binding transcriptional regulator2 (saro_1072) from <i>novosphingobium aromaticivorans</i> dsm at 1.85 a3 resolution
5	c1pb6D_	 Alignment		98.5	15	PDB header: transcription Chain: D: PDB Molecule: hypothetical transcriptional regulator ycdc; PDBTitle: crystal structure of hypothetical transcriptional regulator ycdc
6	d1vi0a2	 Alignment		98.2	17	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
7	c3gziA_	 Alignment		98.1	14	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of a transcriptional regulator of the tetr family2 (shew_3567) from <i>shewanella loihica</i> pv-4 at 2.05 a resolution
8	d2d6ya2	 Alignment		97.7	10	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
9	c3s5rA_	 Alignment		97.6	15	PDB header: dna binding protein Chain: A: PDB Molecule: transcriptional regulator tetr family; PDBTitle: crystal structure of a putative transcriptional regulator of the tetr2 family (syn_02108) from <i>syntrophus aciditrophicus</i> at 2.60 a3 resolution
10	d2id3a2	 Alignment		97.6	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
11	d2o7ta2	 Alignment		97.5	16	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain

12	d2fq4a2	Alignment		97.5	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
13	c2d6yA	Alignment		97.5	9	PDB header: gene regulation Chain: A: PDB Molecule: putative tetr family regulatory protein; PDBTitle: crystal structure of transcriptional factor sco4008 from streptomyces2 coelicolor a3(2)
14	c3dewA	Alignment		97.4	13	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the structure of a putative tetr family transcriptional regulator from2 geobacter sulfurreducens pca.
15	d1rkta2	Alignment		97.3	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
16	d2g3ba2	Alignment		97.3	12	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
17	c2id3A	Alignment		97.2	13	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator sco5951 from2 streptomyces coelicolor a3(2)
18	c2f07A	Alignment		97.1	14	PDB header: transcription Chain: A: PDB Molecule: yvdt; PDBTitle: crystal structure of yvdt from bacillus subtilis
19	c2hyiA	Alignment		97.0	15	PDB header: transcription Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: the crystal structure of a tetr-family transcriptional regulator from2 streptomyces coelicolor
20	c3nrgA	Alignment		96.8	11	PDB header: transcription Chain: A: PDB Molecule: tetr family transcriptional regulator; PDBTitle: crystal structure of a tetr family transcriptional regulator2 (caur_2714) from chloroflexus aurantiacus j-10-fl at 2.56 a3 resolution
21	c3f0cA	Alignment	not modelled	96.6	7	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator from cytophaga2 hutchinsonii atcc 33406
22	d1t33a2	Alignment	not modelled	96.6	10	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
23	c2rekA	Alignment	not modelled	96.6	10	PDB header: transcription Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator
24	d2fbqa2	Alignment	not modelled	96.5	11	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
25	c2pz9A	Alignment	not modelled	96.5	13	PDB header: transcription Chain: A: PDB Molecule: putative regulatory protein; PDBTitle: crystal structure of putative transcriptional regulator sco4942 from2 streptomyces coelicolor
26	c1t33B	Alignment	not modelled	96.5	11	PDB header: transcription Chain: B: PDB Molecule: putative transcriptional repressor (tetr/acrr family); PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional repressor (tetr/acrr family) from salmonella3 typhimurim lt2
27	c2g3bB	Alignment	not modelled	96.5	12	PDB header: transcription Chain: B: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of putative tetr-family transcriptional regulator2 from rhodococcus sp.
28	c2gwtA	Alignment	not modelled	96.5	13	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family;

28	c2qwaA	Alignment	not modelled	96.3	13	PDBTitle: crystal structure of the tetr transcription regulatory2 protein from mycobacterium vanbaalenii
29	d2gfa2	Alignment	not modelled	96.5	10	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
30	c2wuiA	Alignment	not modelled	96.4	11	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of mexz, a key repressor responsible for2 antibiotic resistance in pseudomonas aeruginosa.
31	c2fq4A	Alignment	not modelled	96.4	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the crystal structure of the transcriptional regulator (tetr family)2 from bacillus cereus
32	c3bhqB	Alignment	not modelled	96.4	14	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a putative tetr-family transcriptional regulator2 (mlr_4833) from mesorhizobium loti maff303099 at 1.54 a resolution
33	c2jk3A	Alignment	not modelled	96.4	12	PDB header: transcription Chain: A: PDB Molecule: hemolysin ii regulatory protein; PDBTitle: crystal structure of the hlyiir mutant protein with2 residues 169-186 substituted by gssgssg linker
34	c2zb9A	Alignment	not modelled	96.3	15	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of tetr family transcription regulator2 sco0332
35	c3qbmA	Alignment	not modelled	96.2	14	PDB header: transcription regulator Chain: A: PDB Molecule: tetr transcriptional regulator; PDBTitle: crystal structure of a tetr transcriptional regulator (caur_2221) from2 chloroflexus aurantiacus j-10-fl at 1.80 a resolution
36	c2o7tA	Alignment	not modelled	96.0	16	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a tetr family transcriptional regulator2 (ncgl1578, cgl1640) from corynebacterium glutamicum at 2.10 a3 resolution
37	d1ui5a2	Alignment	not modelled	96.0	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
38	c2iaiA	Alignment	not modelled	96.0	13	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative transcriptional regulator sco3833; PDBTitle: crystal structure of sco3833, a member of the tetr transcriptional2 regulator family from streptomyces coelicolor a3
39	c1ui6B	Alignment	not modelled	95.9	12	PDB header: antibiotic Chain: B: PDB Molecule: a-factor receptor homolog; PDBTitle: crystal structure of gamma-butyrolactone receptor (arpa-like protein)
40	c1jumB	Alignment	not modelled	95.9	12	PDB header: transcription Chain: B: PDB Molecule: hypothetical transcriptional regulator in qaca PDBTitle: crystal structure of the multidrug binding transcriptional2 repressor qacr bound to the natural drug berberine
41	c3rh2A	Alignment	not modelled	95.8	17	PDB header: dna binding protein Chain: A: PDB Molecule: hypothetical tetr-like transcriptional regulator; PDBTitle: crystal structure of a hypothetical tetr-like transcriptional2 regulator (sama_0099) from shewanella amazonensis sb2b at 2.42 a3 resolution
42	c2uxoB	Alignment	not modelled	95.8	13	PDB header: transcription Chain: B: PDB Molecule: hth-type transcriptional regulator ttgr; PDBTitle: ttgr in complex with tetracycline
43	c3cdlA	Alignment	not modelled	95.8	15	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator aefr; PDBTitle: crystal structure of a tetr family transcriptional regulator from2 pseudomonas syringae pv. tomato str. dc3000
44	c3bcgA	Alignment	not modelled	95.7	12	PDB header: transcription regulator Chain: A: PDB Molecule: hth-type transcriptional regulator accrr; PDBTitle: conformational changes of the accrr regulator reveal a2 mechanism of induction
45	d2hya2	Alignment	not modelled	95.6	15	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
46	c2q24A	Alignment	not modelled	95.6	14	PDB header: transcription Chain: A: PDB Molecule: putative tetr family transcriptional regulator; PDBTitle: crystal structure of tetr transcriptional regulator sco0520 from2 streptomyces coelicolor
47	c2w53B	Alignment	not modelled	95.6	13	PDB header: transcription Chain: B: PDB Molecule: repressor; PDBTitle: structure of smet, the repressor of the stenotrophomonas2 maltophilia multidrug efflux pump smedef.
48	c2ibdB	Alignment	not modelled	95.5	19	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: possible transcriptional regulator; PDBTitle: crystal structure of probable transcriptional regulatory protein2 rha5900
49	c3on2C	Alignment	not modelled	95.4	13	PDB header: structural genomics, unknown function Chain: C: PDB Molecule: probable transcriptional regulator; PDBTitle: structure of a protein with unknown function from rhodococcus sp. rha1
50	c3anpD	Alignment	not modelled	95.3	12	PDB header: transcription Chain: D: PDB Molecule: transcriptional repressor, tetr family; PDBTitle: crystal structure of thermus thermophilus fadr, a tetr family2 transcriptional repressor, in complex with lauroyl-coa.
51	c3c2bA	Alignment	not modelled	95.1	15	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of tetr transcriptional regulator from agrobacterium2 tumefaciens
52	c3himA	Alignment	not modelled	95.0	13	PDB header: transcription regulator Chain: A: PDB Molecule: probable transcriptional regulator;

52	c3mna	Alignment	not modelled	93.9	13	PDBTitle: the crystal structure of a bacterial regulatory protein in the tet2 family from rhodococcus rha1 to 2.2a PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator, tet family; PDBTitle: crystal structure of a putative tet/acrr family transcriptional2 regulator (saro_0558) from novosphingobium aromaticivorans dsm at3 1.80 a resolution PDB header: transcription Chain: B: PDB Molecule: hemagglutinin/protease regulatory protein; PDBTitle: vibrio cholerae hapr
53	c2rasB	Alignment	not modelled	94.9	8	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
54	c2pbxB	Alignment	not modelled	94.8	8	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
55	d2g7sa2	Alignment	not modelled	94.7	7	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator of the tet/acrr PDBTitle: crystal structure of transcriptional regulator of the2 tet/acrr family (yp_290855.1) from thermobifida fusca yx-3 er1 at 2.50 a resolution
56	d2gena2	Alignment	not modelled	94.7	15	PDB header: transcription Chain: A: PDB Molecule: putative tet transcriptional regulator; PDBTitle: putative tet transcriptional regulator from rhodococcus sp. rha1
57	c3dcfB	Alignment	not modelled	94.6	11	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator
58	c2i10A	Alignment	not modelled	94.2	11	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
59	c1vi0B	Alignment	not modelled	94.2	19	PDB header: transcription Chain: A: PDB Molecule: protein yfir; PDBTitle: crystal structure of yfir, a putative transcriptional regulator from2 bacillus subtilis
60	d1jt6a2	Alignment	not modelled	94.1	12	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tet family; PDBTitle: the crystal structure of the transcriptional regulator, tet family2 from cytophaga hutchinsonii
61	c1rktA	Alignment	not modelled	94.1	17	PDB header: transcription Chain: A: PDB Molecule: hth-type transcriptional regulator pksa related protein; PDBTitle: crystal structure of hth-type transcriptional regulator pksa related2 protein from rhodococcus sp. rha1
62	c3eupA	Alignment	not modelled	93.9	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tet family; PDBTitle: crystal structure of transcriptional regulator of tet2 family (yp_425770.1) from rhodospirillum rubrum atcc 111703 at 1.50 a resolution
63	c2gfnA	Alignment	not modelled	93.8	9	PDB header: transcription Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: structural genomics, the crystal structure of a probable2 transcriptional regulator from pseudomonas aeruginosa pao1
64	c3cwrA	Alignment	not modelled	93.5	9	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, tet family; PDBTitle: crystal structure of transcriptional regulator, tet family (tm1030)2 from thermotoga maritima at 2.30 a resolution
65	c2genA	Alignment	not modelled	93.4	14	PDB header: transcription Chain: A: PDB Molecule: regulatory protein, tet family; PDBTitle: crystal structure of regulatory protein tet from rhodobacter2 sphaeroides
66	c1zkgB	Alignment	not modelled	93.1	10	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: the crystal structure of the putative transcriptional regulator from2 pseudomonas aeruginosa pao1
67	c3bruA	Alignment	not modelled	92.7	12	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
68	c3e7qB	Alignment	not modelled	92.3	13	PDB header: transcription regulator Chain: A: PDB Molecule: putative tet-family transcriptional regulator; PDBTitle: crystal structure of putative tet-family transcriptional regulator2 (yp_752756.1) from syntrophomonas wolfei str. goettingen at 2.07 a3 resolution
69	d2i10a2	Alignment	not modelled	92.0	15	PDB header: transcription Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: the crystal structure of transcriptional regulator pa3006
70	c3lwiA	Alignment	not modelled	91.7	16	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tet family; PDBTitle: the crystal structure of transcriptional regulator, tet family, from2 agrobacterium tumefaciens
71	c2fbgA	Alignment	not modelled	91.6	11	PDB header: transcription regulator Chain: C: PDB Molecule: transcription regulator; PDBTitle: crystal structure of a tet transcription regulator from haloarcula2 marismortui atcc 43049
72	c2g7sA	Alignment	not modelled	91.3	11	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tet family; PDBTitle: the structure of a putative transcriptional regulator tet family2 protein from vibrio parahaemolyticus.
73	c3crjC	Alignment	not modelled	90.3	16	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, acrr family; PDBTitle: crystal structure of predicted dna-binding transcriptional regulator2 of tet/acrr family (np_350189.1) from clostridium acetobutylicum at3 2.10 a resolution
74	c3he0A	Alignment	not modelled	89.2	18	PDB header: dna binding protein Chain: A: PDB Molecule: transcriptional repressor ethr; PDBTitle: crystal structure of the transcriptional regulator ethr in2 a ligand bound conformation opens therapeutic perspectives3
75	c3b81A	Alignment	not modelled	88.9	19	
76	c1u9nA	Alignment	not modelled	88.4	4	

						against tuberculosis and leprosy PDB header: transcription regulator Chain: A: PDB Molecule: tetr family transcription regulator; PDBTitle: crystal structure of a tetr family transcription regulator (maqu_1417)2 from marinobacter aquaeolei vt8 at 1.90 a resolution
77	c3pasA	Alignment	not modelled	88.4	9	PDB header: transcription regulator Chain: B: PDB Molecule: putative tetr family transcription regulator; PDBTitle: crystal structure of a putative tetr family transcription regulator2 (shew_3104) from shewanella sp. pv-4 at 2.10 a resolution
78	c3ppbB	Alignment	not modelled	87.8	9	PDB header: transcription regulator Chain: B: PDB Molecule: putative tetr family transcription regulator; PDBTitle: crystal structure of a putative tetr family transcription regulator2 (shew_3104) from shewanella sp. pv-4 at 2.10 a resolution
79	c2nx4A	Alignment	not modelled	87.4	11	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: transcriptional regulator, tetr family protein; PDBTitle: the crystal structure of a putative tetr-family transcriptional2 regulator rha06780 from rhodococcus sp. rha1.
80	c2hytA	Alignment	not modelled	85.7	17	PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of a tetr-family transcriptional regulator (eca1819)2 from peptobacterium atrosepticum at 1.64 a resolution
81	c3knwB	Alignment	not modelled	85.4	8	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator (tetr/acrr family); PDBTitle: crystal structure of a putative transcriptional regulator (tetr/acrr2 family member) from putative transcriptional regulator (tetr/acrr3 family)
82	d1t56a2	Alignment	not modelled	84.4	4	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
83	c2qcoA	Alignment	not modelled	82.6	5	PDB header: transcription Chain: A: PDB Molecule: cmer; PDBTitle: crystal structure of the transcriptional regulator cmer from2 campylobacter jejuni
84	c3ccyA	Alignment	not modelled	78.2	8	PDB header: transcription regulator Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of a tetr-family transcriptional regulator from2 bordetella parapertussis 12822
85	c3colB	Alignment	not modelled	67.3	8	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcription regulator; PDBTitle: crystal structure of putative transcription regulator from2 lactobacillus plantarum
86	d1zk8a2	Alignment	not modelled	60.1	11	Fold: Tetracyclin repressor-like, C-terminal domain Superfamily: Tetracyclin repressor-like, C-terminal domain Family: Tetracyclin repressor-like, C-terminal domain
87	c3qkxB	Alignment	not modelled	56.4	3	PDB header: transcription regulator Chain: B: PDB Molecule: uncharacterized hth-type transcriptional regulator hi_0893; PDBTitle: crystal structure of a tetr-family transcriptional regulator (hi0893)2 from haemophilus influenzae rd at 2.35 a resolution
88	c3c07B	Alignment	not modelled	46.6	4	PDB header: transcription regulator Chain: B: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of a tetr family transcriptional regulator from2 streptomyces coelicolor a3(2)
89	c2np5A	Alignment	not modelled	44.5	21	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator (rha1_ro04179) from2 rhodococcus sp. rha1.
90	d1g8eb	Alignment	not modelled	42.6	25	Fold: Flagellar transcriptional activator FlhD Superfamily: Flagellar transcriptional activator FlhD Family: Flagellar transcriptional activator FlhD
91	d1g8ea	Alignment	not modelled	41.6	25	Fold: Flagellar transcriptional activator FlhD Superfamily: Flagellar transcriptional activator FlhD Family: Flagellar transcriptional activator FlhD
92	c1zk8A	Alignment	not modelled	41.6	12	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator from bacillus cereus2 atcc 14579
93	c3aqsD	Alignment	not modelled	36.9	8	PDB header: transcription regulator Chain: D: PDB Molecule: bacterial regulatory proteins, tetr family; PDBTitle: crystal structure of rolr (ncgl1110) without ligand
94	c2zcxA	Alignment	not modelled	34.8	11	PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr family transcriptional regulator2 sco7815
95	c1z0xA	Alignment	not modelled	32.3	9	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator, tetr family from2 enterococcus faecalis v583
96	c1sgmA	Alignment	not modelled	30.6	13	PDB header: transcription Chain: A: PDB Molecule: putative hth-type transcriptional regulator yxaf; PDBTitle: crystal structure of hypothetical protein yxaf
97	c2yvHA	Alignment	not modelled	30.5	13	PDB header: transcription/dna Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of the operator-binding form of the multi-drug2 binding transcriptional repressor cgmr
98	c3geuC	Alignment	not modelled	28.4	8	PDB header: cell adhesion Chain: C: PDB Molecule: intercellular adhesion protein r; PDBTitle: crystal structure of icaR from staphylococcus aureus, a member of the2 tetracycline repressor protein family
99	c3nnrA	Alignment	not modelled	28.3	10	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of a tetr-family transcriptional regulator2 (maqu_3571) from marinobacter aquaeolei vt8 at 2.49 a resolution
100	c3nn4D	Alignment	not modelled	23.5	20	PDB header: transcription Chain: D: PDB Molecule: transcriptional regulator, tetr family;

100	c3b6aC	Alignment	not modelled	23.5	20	PDBTitle: crystal structure of tetr transcriptional regulator from legionella2 pneumophila
101	c3b6aC	Alignment	not modelled	23.5	13	PDB header: transcription Chain: C: PDB Molecule: actr protein; PDBTitle: crystal structure of the streptomyces coelicolor tetr2 family protein actr in complex with actinorhodin