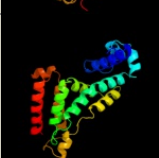
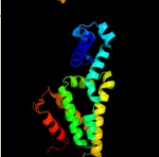
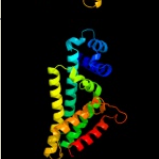
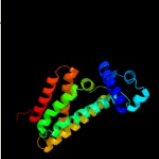


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1pb6D_	 Alignment		100.0	100	PDB header: transcription Chain: D: PDB Molecule: hypothetical transcriptional regulator ycdc; PDBTitle: crystal structure of hypothetical transcriptional regulator ycdc
2	c2qtqB_	 Alignment		99.9	19	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of a predicted dna-binding transcriptional regulator2 (saro_1072) from novosphingobium aromaticivorans dsm at 1.85 a3 resolution
3	c3e7qB_	 Alignment		99.9	14	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: the crystal structure of the putative transcriptional regulator from2 pseudomonas aeruginosa pao1
4	c2pbxB_	 Alignment		99.9	13	PDB header: transcription Chain: B: PDB Molecule: hemagglutinin/protease regulatory protein; PDBTitle: vibrio cholerae hapr
5	c3nrgA_	 Alignment		99.9	17	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
6	c3gziA_	 Alignment		99.9	18	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 shewanella loihica pv-4 at 2.05 a resolution
7	c3dewA_	 Alignment		99.9	20	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.
8	c3himA_	 Alignment		99.9	15	PDB header: transcription regulator Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: the crystal structure of a bacterial regulatory protein in the tetr2 family from rhodococcus rha1 to 2.2a
9	c2rasB_	 Alignment		99.9	15	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of a putative tetr/acrr family transcriptional2 regulator (saro_0558) from novosphingobium aromaticivorans dsm at3 1.80 a resolution
10	c3lwiA_	 Alignment		99.9	16	PDB header: transcription regulator Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of putative tetr-family transcriptional regulator2 (yp_752756.1) from syntrophomonas wolfei str. goettingen at 2.07 a3 resolution
11	c3ppbB_	 Alignment		99.9	19	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

12	c2iaiA_	Alignment		99.9	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative transcriptional regulator sco3833; PDBTitle: crystal structure of sco3833, a member of the tet ^r transcriptional2 regulator family from streptomyces coelicolor a3
13	c3dcfB_	Alignment		99.9	22	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator of the tet ^r /acrr PDBTitle: crystal structure of a transcriptional regulator of the2 tet ^r /acrr family (yp_290855.1) from thermobifida fusca yx-3 er1 at 2.50 a resolution
14	c3rh2A_	Alignment		99.9	17	PDB header: dna binding protein Chain: A: PDB Molecule: hypothetical tet ^r -like transcriptional regulator; PDBTitle: crystal structure of a hypothetical tet ^r -like transcriptional2 regulator (sama_0099) from shewanella amazonensis sb2b at 2.42 a3 resolution
15	c3bhqB_	Alignment		99.9	17	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a putative tet ^r -family transcriptional regulator2 (mlr_4833) from mesorhizobium loti maff303099 at 1.54 a resolution
16	c1vi0B_	Alignment		99.9	16	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator
17	c3pasA_	Alignment		99.9	17	PDB header: transcription regulator Chain: A: PDB Molecule: tet ^r family transcription regulator; PDBTitle: crystal structure of a tet ^r family transcription regulator (maqu_1417)2 from marinobacter aquaeolei vt8 at 1.90 a resolution
18	c3f0cA_	Alignment		99.9	17	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator from cytophaga2 hutchinsonii atcc 33406
19	c2zb9A_	Alignment		99.9	21	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of tet ^r family transcription regulator2 sco0332
20	c2nx4A_	Alignment		99.9	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: transcriptional regulator, tet ^r family protein; PDBTitle: the crystal structure of a putative tet ^r -family transcriptional2 regulator rha06780 from rhodococcus sp. rha1.
21	c3qbmA_	Alignment	not modelled	99.9	16	PDB header: transcription regulator Chain: A: PDB Molecule: tet ^r transcriptional regulator; PDBTitle: crystal structure of a tet ^r transcriptional regulator (caur_2221) from2 chloroflexus aurantiacus j-10-fl at 1.80 a resolution
22	c3eupA_	Alignment	not modelled	99.9	16	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tet ^r family; PDBTitle: the crystal structure of a tet ^r family regulatory protein, tet ^r family2 from cytophaga hutchinsonii
23	c2hyjA_	Alignment	not modelled	99.9	17	PDB header: transcription Chain: A: PDB Molecule: putative tet ^r -family transcriptional regulator; PDBTitle: the crystal structure of a tet ^r -family transcriptional regulator from2 streptomyces coelicolor
24	c2f07A_	Alignment	not modelled	99.9	15	PDB header: transcription Chain: A: PDB Molecule: yvdt; PDBTitle: crystal structure of yvdt from bacillus subtilis
25	c3npiB_	Alignment	not modelled	99.9	17	PDB header: gene regulation Chain: B: PDB Molecule: tet ^r family regulatory protein; PDBTitle: crystal structure of a tet ^r family regulatory protein (dip1788) from2 corynebacterium diphtheriae at 2.96 a resolution
26	c3on2C_	Alignment	not modelled	99.9	17	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
27	c3ccyA_	Alignment	not modelled	99.9	17	PDB header: transcription regulator Chain: A: PDB Molecule: putative tet ^r -family transcriptional regulator; PDBTitle: crystal structure of a tet ^r -family transcriptional regulator from2 bordetella parapertussis 12822
28	c1rktA_	Alignment	not modelled	99.9	16	PDB header: transcription Chain: A: PDB Molecule: protein yfir; PDBTitle: crystal structure of yfir, a putative transcriptional

					regulator from2 bacillus subtilis
29	c3lsjA_	Alignment	not modelled	99.9	17 PDB header: transcription Chain: A: PDB Molecule: dest; PDBTitle: crystal structure of dest in complex with palmitoyl-coa
30	c2genA_	Alignment	not modelled	99.9	22 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
31	c3anpD_	Alignment	not modelled	99.9	19 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.
32	c2g7sA_	Alignment	not modelled	99.9	19 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the crystal structure of transcriptional regulator, tetr family, from2 agrobacterium tumefaciens
33	c3s5rA_	Alignment	not modelled	99.9	20 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 syntrophus aciditrophicus at 2.60 a3 resolution
34	c1jumB_	Alignment	not modelled	99.9	19 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
35	c1t33B_	Alignment	not modelled	99.9	22 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
36	c2fq4A_	Alignment	not modelled	99.9	20 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
37	c2hytA_	Alignment	not modelled	99.9	19 PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of a tetr-family transcriptional regulator (eca1819)2 from pectobacterium atrosepticum at 1.64 a resolution
38	c2fbgA_	Alignment	not modelled	99.9	19 PDB header: transcription Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: the crystal structure of transcriptional regulator pa3006
39	c1u9nA_	Alignment	not modelled	99.9	17 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 against tuberculosis and leprosy
40	c2ibdB_	Alignment	not modelled	99.9	20 PDB header: structural genomics, unknown function Chain: B: PDB Molecule: possible transcriptional regulator; PDBTitle: crystal structure of probable transcriptional regulatory protein2 rha5900
41	c3nnrA_	Alignment	not modelled	99.9	14 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
42	c2q24A_	Alignment	not modelled	99.9	20 PDB header: transcription Chain: A: PDB Molecule: putative tetr family transcriptional regulator; PDBTitle: crystal structure of tetr transcriptional regulator sco0520 from2 streptomyces coelicolor
43	c3bcgA_	Alignment	not modelled	99.9	15 PDB header: transcription regulator Chain: A: PDB Molecule: hth-type transcriptional regulator acrr; PDBTitle: conformational changes of the acrr regulator reveal a2 mechanism of induction
44	c2uxoB_	Alignment	not modelled	99.9	16 PDB header: transcription Chain: B: PDB Molecule: hth-type transcriptional regulator ttgr; PDBTitle: ttgr in complex with tetracycline
45	c3f1bA_	Alignment	not modelled	99.9	20 PDB header: transcription regulator Chain: A: PDB Molecule: tetr-like transcriptional regulator; PDBTitle: the crystal structure of a tetr-like transcriptional regulator from2 rhodococcus sp. rha1.
46	c1zkqB_	Alignment	not modelled	99.9	21 PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator, tetr family (tm1030)2 from thermotoga maritima at 2.30 a resolution
47	c3bruA_	Alignment	not modelled	99.9	20 PDB header: transcription Chain: A: PDB Molecule: regulatory protein, tetr family; PDBTitle: crystal structure of regulatory protein tetr from rhodobacter2 sphaeroides
48	c2raeA_	Alignment	not modelled	99.9	16 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, acrr family protein; PDBTitle: crystal structure of a tetr/acrr family transcriptional regulator from2 rhodococcus sp. rha1
49	c2id3A_	Alignment	not modelled	99.9	18 PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator sco5951 from2 streptomyces coelicolor a3(2)
50	c2g3bB_	Alignment	not modelled	99.9	19 PDB header: transcription Chain: B: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of putative tetr-family transcriptional regulator2 from rhodococcus sp.
51	c3he0A_	Alignment	not modelled	99.9	18 PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the structure of a putative transcriptional regulator tetr family2 protein from vibrio parahaemolyticus.
52	c2rakA_	Alignment	not modelled	99.9	21 PDB header: transcription Chain: A: PDB Molecule: putative tetr-family transcriptional

52	c2t8A_	Alignment	not modelled	99.9	21	regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator
53	c1zk8A_	Alignment	not modelled	99.9	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator from bacillus cereus2 atcc 14579
54	c2qwtA_	Alignment	not modelled	99.9	18	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of the tetr transcription regulatory2 protein from mycobacterium vanbaalenii
55	c3mn1A_	Alignment	not modelled	99.9	16	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulatory protein (probably tetr-family); PDBTitle: the crystal structure of kstr (rv3574) from mycobacterium tuberculosis2 h37rv
56	c3qkxB_	Alignment	not modelled	99.9	15	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
57	c2np5A_	Alignment	not modelled	99.9	17	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator (rha1_ro04179) from2 rhodococcus sp. rha1.
58	c2o7tA_	Alignment	not modelled	99.9	15	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
59	c2jk3A_	Alignment	not modelled	99.9	21	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
60	c2qcoA_	Alignment	not modelled	99.9	17	PDB header: transcription Chain: A: PDB Molecule: cmer; PDBTitle: crystal structure of the transcriptional regulator cmer from2 campylobacter jejuni
61	c1ui6B_	Alignment	not modelled	99.9	16	PDB header: antibiotic Chain: B: PDB Molecule: a-factor receptor homolog; PDBTitle: crystal structure of gamma-butyrolactone receptor (arpa-like protein)
62	c3c2bA_	Alignment	not modelled	99.9	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of tetr transcriptional regulator from agrobacterium2 tumefaciens
63	c2gfnA_	Alignment	not modelled	99.9	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
64	c2zcxA_	Alignment	not modelled	99.9	18	PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr family transcriptional regulator2 sco7815
65	c3c07B_	Alignment	not modelled	99.9	16	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)
66	c3g7rB_	Alignment	not modelled	99.8	17	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of sco4454, a tetr-family transcriptional2 regulator from streptomyces coelicolor
67	c3cjdB_	Alignment	not modelled	99.8	13	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of putative tetr transcriptional regulator2 (yp_510936.1) from jannaschia sp. ccs1 at 1.79 a resolution
68	c3bjbE_	Alignment	not modelled	99.8	14	PDB header: transcription regulator Chain: E: PDB Molecule: probable transcriptional regulator, tetr family protein; PDBTitle: crystal structure of a tetr transcriptional regulator from rhodococcus2 sp. rha1
69	c3cwrA_	Alignment	not modelled	99.8	13	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator of tetr2 family (yp_425770.1) from rhodospirillum rubrum atcc 111703 at 1.50 a resolution
70	c2wuiA_	Alignment	not modelled	99.8	17	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of mexz, a key repressor responsible for2 antibiotic resistance in pseudomonas aeruginosa.
71	c2oi8A_	Alignment	not modelled	99.8	16	PDB header: transcription Chain: A: PDB Molecule: putative regulatory protein sco4313; PDBTitle: crystal structure of putative regulatory protein sco4313
72	c2hkuB_	Alignment	not modelled	99.8	15	PDB header: transcription regulator Chain: B: PDB Molecule: a putative transcriptional regulator; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator from rhodococcus sp. rha1
73	c2d6yA_	Alignment	not modelled	99.8	19	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)
74	c3cd1A_	Alignment	not modelled	99.8	23	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
75	c3colB_	Alignment	not modelled	99.8	13	PDB header: transcription regulator Chain: B: PDB Molecule: putative transcription regulator; PDBTitle: crystal structure of putative transcription regulator from2 lactobacillus plantarum
						PDB header: transcription

76	c2i10A_	Alignment	not modelled	99.8	15	Chain: A: PDB Molecule: putative tetr transcriptional regulator; PDBTitle: putative tetr transcriptional regulator from rhodococcus sp. rha1
77	c3on4D_	Alignment	not modelled	99.8	18	PDB header: transcription Chain: D: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of tetr transcriptional regulator from legionella2 pneumophila
78	c3mvpA_	Alignment	not modelled	99.8	19	PDB header: transcription regulator Chain: A: PDB Molecule: tetr/acrr transcriptional regulator; PDBTitle: the crystal structure of a tetr/acrr transcriptional regulator from2 streptococcus mutans to 1.85a
79	c2of7A_	Alignment	not modelled	99.8	18	PDB header: transcription Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: structural genomics, the crystal structure of a tetr-family2 transcriptional regulator from streptomyces coelicolor a3
80	c3b81A_	Alignment	not modelled	99.8	17	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, acrr family; PDBTitle: crystal structure of predicted dna-binding transcriptional regulator2 of tetr/acrr family (np_350189.1) from clostridium acetobutylicum at3 2.10 a resolution
81	c2qibA_	Alignment	not modelled	99.8	15	PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator from2 streptomyces coelicolor
82	c3geuC_	Alignment	not modelled	99.8	19	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
83	c3djpB_	Alignment	not modelled	99.8	20	PDB header: dna binding protein Chain: B: PDB Molecule: transcription regulator, tetr family; PDBTitle: the crystal structure of a tetr transcription regulator2 from silicibacter pomeroyi dss
84	c3knwB_	Alignment	not modelled	99.8	20	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)
85	c1sgmA_	Alignment	not modelled	99.8	18	PDB header: transcription Chain: A: PDB Molecule: putative hth-type transcriptional regulator yxaf; PDBTitle: crystal structure of hypothetical protein yxaf
86	c3aqsD_	Alignment	not modelled	99.8	15	PDB header: transcription regulator Chain: D: PDB Molecule: bacterial regulatory proteins, tetr family; PDBTitle: crystal structure of rolr (ncgl1110) without ligand
87	c2dg7A_	Alignment	not modelled	99.8	14	PDB header: gene regulation Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: crystal structure of the putative transcriptional regulator sco03372 from streptomyces coelicolor a3(2)
88	c1z0xA_	Alignment	not modelled	99.8	12	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of transcriptional regulator, tetr family from2 enterococcus faecalis v583
89	c2zcmB_	Alignment	not modelled	99.8	16	PDB header: transcription Chain: B: PDB Molecule: biofilm operon icaabcd hth-type negative transcriptional PDBTitle: crystal structure of icar, a repressor of the tetr family
90	c2eh3A_	Alignment	not modelled	99.8	18	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of aq_1058, a transcriptional regulator (terr/acrr2 family) from aquifex aeolicus vf5
91	c3fiwB_	Alignment	not modelled	99.8	16	PDB header: transcription regulator Chain: B: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: structure of sco0253, a tetr-family transcriptional regulator from2 streptomyces coelicolor
92	c3jsjC_	Alignment	not modelled	99.8	19	PDB header: transcription Chain: C: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of a putative tetr-transcriptional regulator2 (sav143) from streptomyces avermitilis ma-4680 at 2.10 a resolution
93	c3o60A_	Alignment	not modelled	99.8	10	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lin0861 protein; PDBTitle: the crystal structure of lin0861 from listeria innocua to 2.8a
94	c3g56A_	Alignment	not modelled	99.8	14	PDB header: dna binding protein Chain: A: PDB Molecule: regulator of macrolide 2'-phosphotransferase i; PDBTitle: structure of the macrolide biosensor protein, mphr(a)
95	c2dg8D_	Alignment	not modelled	99.8	17	PDB header: gene regulation Chain: D: PDB Molecule: putative tetr-family transcriptional regulatory protein; PDBTitle: crystal structure of the putative trasncriptional regulator sco75182 from streptomyces coelicolor a3(2)
96	c2y31A_	Alignment	not modelled	99.8	15	PDB header: transcription Chain: A: PDB Molecule: putative repressor simreg2; PDBTitle: simocyclinone c4 bound form of tetr-like repressor simr
97	c2pz9A_	Alignment	not modelled	99.8	24	PDB header: transcription Chain: A: PDB Molecule: putative regulatory protein; PDBTitle: crystal structure of putative transcriptional regulator sco4942 from2 streptomyces coelicolor
98	c3nxC_	Alignment	not modelled	99.8	14	PDB header: dna binding protein Chain: A: PDB Molecule: hth-type protein slma; PDBTitle: molecular mechanism by which the escherichia coli nucleoid occlusion2 factor, slma, keeps cytokinesis in check
99	c2yvha_	Alignment	not modelled	99.8	18	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
						PDB header: transcription

100	c2guhA	Alignment	not modelled	99.8	16	Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of the putative tetr-family transcriptional2 regulator from rhodococcus sp. rha1
101	c3iuvA	Alignment	not modelled	99.8	18	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized tetr family protein; PDBTitle: the structure of a member of tetr family (sco1917) from2 streptomyces coelicolor a3
102	c3b6aC	Alignment	not modelled	99.8	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
103	c2fd5A	Alignment	not modelled	99.8	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: the crystal structure of a transcriptional regulator from pseudomonas2 aeruginosa pao1
104	c3bqyA	Alignment	not modelled	99.8	18	PDB header: transcription regulator Chain: A: PDB Molecule: putative tetr family transcriptional regulator; PDBTitle: crystal structure of a possible tetr family transcriptional regulator2 from streptomyces coelicolor a3(2).
105	c3kkdA	Alignment	not modelled	99.8	13	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: structure of a putative tetr transcriptional regulator (pa3699) from2 pseudomonas aeruginosa pa01
106	c1bjzA	Alignment	not modelled	99.7	14	PDB header: transcription regulation Chain: A: PDB Molecule: tetracycline repressor; PDBTitle: tetracycline chelated mg2+-ion initiates helix unwinding for tet2 repressor induction
107	c3bniA	Alignment	not modelled	99.7	17	PDB header: transcription regulator Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator from2 streptomyces coelicolor
108	c2hxoB	Alignment	not modelled	99.7	13	PDB header: transcription Chain: B: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: structure of the transcriptional regulator sco7222, a tetr from2 streptomyces coelicolor
109	c3kkcB	Alignment	not modelled	99.7	14	PDB header: transcription regulator Chain: B: PDB Molecule: tetr family transcriptional regulator; PDBTitle: the crystal structure of tetr transcriptional regulator from2 streptococcus agalactiae 2603v
110	c3crjC	Alignment	not modelled	99.7	15	PDB header: transcription regulator Chain: C: PDB Molecule: transcription regulator; PDBTitle: crystal structure of a tetr transcription regulator from haloarcula2 marismortui atcc 43049
111	c2iu5A	Alignment	not modelled	99.7	12	PDB header: transcription Chain: A: PDB Molecule: hth-type dhaklm operon transcriptional activator dhas; PDBTitle: dihydroxyacetone kinase operon activator dhas
112	c2g7lA	Alignment	not modelled	99.7	12	PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of putative transcription regulator sco7704 from2 streptomyces coelicor
113	c3ni7A	Alignment	not modelled	99.7	16	PDB header: transcription regulator Chain: A: PDB Molecule: bacterial regulatory proteins, tetr family; PDBTitle: crystal structure of the tetr transcriptional regulator from2 nitrosomonas europaea atcc 19718
114	c2hxiA	Alignment	not modelled	99.7	15	PDB header: transcription Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: structural genomics, the crystal structure of a putative2 transcriptional regulator from streptomyces coelicolor a3(2)
115	c3egqB	Alignment	not modelled	99.7	16	PDB header: transcription Chain: B: PDB Molecule: tetr family transcriptional regulator; PDBTitle: crystal structure of a tetr-family transcriptional regulator (af 1817)2 from archaeoglobus fulgidus at 2.55 a resolution
116	c2vprA	Alignment	not modelled	99.7	15	PDB header: transcription Chain: A: PDB Molecule: tetracycline resistance repressor protein; PDBTitle: tet repressor class h in complex with 5a,6-2 anhydrotetracycline-mg
117	c2qkoA	Alignment	not modelled	99.7	18	PDB header: transcription Chain: A: PDB Molecule: possible transcriptional regulator, tetr family protein; PDBTitle: crystal structure of transcriptional regulator rha06399 from2 rhodococcus sp. rha1
118	c2g7gA	Alignment	not modelled	99.7	19	PDB header: transcription Chain: A: PDB Molecule: rha04620, putative transcriptional regulator; PDBTitle: the crystal structure of the putative transcriptional regulator2 rha04620 from rhodococcus sp. rha1
119	d2oi8a1	Alignment	not modelled	99.6	22	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
120	c2w53B	Alignment	not modelled	99.6	16	PDB header: transcription Chain: B: PDB Molecule: repressor; PDBTitle: structure of smet, the repressor of the stenotrophomonas2 maltophilia multidrug efflux pump smedef.