

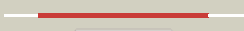


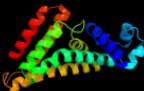


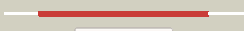










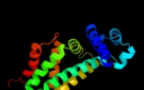

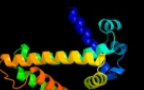




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3qbmA_</a>	 Alignment		99.9	21	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> tetr transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr transcriptional regulator (caur_2221) from2 chloroflexus aurantiacus j-10-fl at 1.80 a resolution
2	<a href="#">c3bruA_</a>	 Alignment		99.9	26	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> regulatory protein, tetr family; <b>PDBTitle:</b> crystal structure of regulatory protein tetr from rhodobacter2 sphaeroides
3	<a href="#">c2hyjA_</a>	 Alignment		99.9	17	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> the crystal structure of a tetr-family transcriptional regulator from2 streptomyces coelicolor
4	<a href="#">c3eupA_</a>	 Alignment		99.9	19	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> the crystal structure of the transcriptional regulator, tetr family2 from cytophaga hutchinsonii
5	<a href="#">c2g7sA_</a>	 Alignment		99.9	16	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> the crystal structure of transcriptional regulator, tetr family, from2 agrobacterium tumefaciens
6	<a href="#">c2pbxB_</a>	 Alignment		99.9	12	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> hemagglutinin/protease regulatory protein; <b>PDBTitle:</b> vibrio cholerae hapr
7	<a href="#">c2nx4A_</a>	 Alignment		99.9	17	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family protein; <b>PDBTitle:</b> the crystal structure of a the putative tetr-family transcriptional2 regulator rha06780 from rhodococcus sp. rha1.
8	<a href="#">c2uxoB_</a>	 Alignment		99.9	15	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> hth-type transcriptional regulator ttgr; <b>PDBTitle:</b> ttgr in complex with tetracycline
9	<a href="#">c3e7qB_</a>	 Alignment		99.9	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> the crystal structure of the putative transcriptional regulator from2 pseudomonas aeruginosa pao1
10	<a href="#">c3lwjA_</a>	 Alignment		99.9	19	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of putative tetr-family transcriptional regulator2 (yp_752756.1) from syntrophomonas wolfei str. goettingen at 2.07 a3 resolution
11	<a href="#">c1rktA_</a>	 Alignment		99.9	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> protein yfir; <b>PDBTitle:</b> crystal structure of yfir, a putative transcriptional regulator from2 bacillus subtilis

12	<a href="#">c2f07A_</a>	Alignment		99.9	18	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> yvdt; <b>PDBTitle:</b> crystal structure of yvdt from bacillus subtilis
13	<a href="#">c3on4D_</a>	Alignment		99.9	19	<b>PDB header:</b> transcription <b>Chain:</b> D: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of tetr transcriptional regulator from legionella2 pneumophila
14	<a href="#">c2g3bB_</a>	Alignment		99.9	16	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of putative tetr-family transcriptional regulator2 from rhodococcus sp.
15	<a href="#">c3bcgA_</a>	Alignment		99.9	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> hth-type transcriptional regulator acrr; <b>PDBTitle:</b> conformational changes of the acrr regulator reveal a2 mechanism of induction
16	<a href="#">c3knwB_</a>	Alignment		99.9	42	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative transcriptional regulator (tetr/acrr family); <b>PDBTitle:</b> crystal structure of a putative transcriptional regulator (tetr/acrr2 family member) from putative transcriptional regulator (tetr/acrr3 family)
17	<a href="#">c2zb9A_</a>	Alignment		99.9	14	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr family transcription regulator2 sco0332
18	<a href="#">c2i10A_</a>	Alignment		99.9	17	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr transcriptional regulator; <b>PDBTitle:</b> putative tetr transcriptional regulator from rhodococcus sp. rha1
19	<a href="#">c1sgmA_</a>	Alignment		99.9	22	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative hth-type transcriptional regulator yxaf; <b>PDBTitle:</b> crystal structure of hypothetical protein yxaf
20	<a href="#">c1vi0B_</a>	Alignment		99.9	12	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of a transcriptional regulator
21	<a href="#">c3nrgA_</a>	Alignment	not modelled	99.9	15	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetr family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr family transcriptional regulator2 (caur_2714) from chloroflexus aurantiacus j-10-fl at 2.56 a3 resolution
22	<a href="#">c3bhqB_</a>	Alignment	not modelled	99.9	13	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of a putative tetr-family transcriptional regulator2 (mlr_4833) from mesorhizobium loti maff303099 at 1.54 a resolution
23	<a href="#">c3anpD_</a>	Alignment	not modelled	99.9	18	<b>PDB header:</b> transcription <b>Chain:</b> D: <b>PDB Molecule:</b> transcriptional repressor, tetr family; <b>PDBTitle:</b> crystal structure of thermus thermophilus fadr, a tetr family2 transcriptional repressor, in complex with lauroyl-coa.
24	<a href="#">c3dpjB_</a>	Alignment	not modelled	99.9	16	<b>PDB header:</b> dna binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> transcription regulator, tetr family; <b>PDBTitle:</b> the crystal structure of a tetr transcription regulator2 from silicibacter pomeroyi dss
25	<a href="#">c3s5rA_</a>	Alignment	not modelled	99.9	13	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator tetr family; <b>PDBTitle:</b> crystal structure of a putative transcriptional regulator of the tetr2 family (syn_02108) from syntrophus aciditrophicus at 2.60 a3 resolution
26	<a href="#">c3on2C_</a>	Alignment	not modelled	99.9	18	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> C: <b>PDB Molecule:</b> probable transcriptional regulator; <b>PDBTitle:</b> structure of a protein with unknown function from rhodococcus sp. rha1
27	<a href="#">c2gfnA_</a>	Alignment	not modelled	99.9	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> hth-type transcriptional regulator pksa related protein; <b>PDBTitle:</b> crystal structure of hth-type transcriptional regulator pksa related2 protein from rhodococcus sp. rha1

28	<a href="#">c1u9nA</a>	 Alignment	not modelled	99.9	19	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional repressor ethr; <b>PDBTitle:</b> crystal structure of the transcriptional regulator ethr in2 a ligand bound conformation opens therapeutic perspectives3 against tuberculosis and leprosy
29	<a href="#">c2hkuB</a>	 Alignment	not modelled	99.9	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> a putative transcriptional regulator; <b>PDBTitle:</b> structural genomics, the crystal structure of a putative2 transcriptional regulator from rhodococcus sp. rha1
30	<a href="#">c3himA</a>	 Alignment	not modelled	99.9	12	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> probable transcriptional regulator; <b>PDBTitle:</b> the crystal structure of a bacterial regulatory protein in the tet2 family from rhodococcus rha1 to 2.2a
31	<a href="#">c3gziA</a>	 Alignment	not modelled	99.9	12	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> crystal structure of a transcriptional regulator of the tet family2 (shew_3567) from shewanella loihica pv-4 at 2.05 a resolution
32	<a href="#">c3pasA</a>	 Alignment	not modelled	99.9	12	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> tetr family transcription regulator; <b>PDBTitle:</b> crystal structure of a tet family transcription regulator (magu_1417)2 from marinobacter aquaeolei vt8 at 1.90 a resolution
33	<a href="#">c2rasB</a>	 Alignment	not modelled	99.9	12	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> crystal structure of a putative tet/acrr family transcriptional2 regulator (saro_0558) from novosphingobium aromaticivorans dsm at3 1.80 a resolution
34	<a href="#">c2genA</a>	 Alignment	not modelled	99.9	11	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> probable transcriptional regulator; <b>PDBTitle:</b> structural genomics, the crystal structure of a probable2 transcriptional regulator from pseudomonas aeruginosa pao1
35	<a href="#">c1zkgB</a>	 Alignment	not modelled	99.9	13	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> crystal structure of transcriptional regulator, tet family (tm1030)2 from thermotoga maritima at 2.30 a resolution
36	<a href="#">c3ppbB</a>	 Alignment	not modelled	99.9	14	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative tet family transcription regulator; <b>PDBTitle:</b> crystal structure of a putative tet family transcription regulator2 (shew_3104) from shewanella sp. pv-4 at 2.10 a resolution
37	<a href="#">c3g7rB</a>	 Alignment	not modelled	99.9	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative transcriptional regulator; <b>PDBTitle:</b> crystal structure of sco4454, a tet-family transcriptional2 regulator from streptomyces coelicolor
38	<a href="#">c2qtqB</a>	 Alignment	not modelled	99.9	12	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> crystal structure of a predicted dna-binding transcriptional regulator2 (saro_1072) from novosphingobium aromaticivorans dsm at 1.85 a3 resolution
39	<a href="#">c1pb6D</a>	 Alignment	not modelled	99.9	18	<b>PDB header:</b> transcription <b>Chain:</b> D: <b>PDB Molecule:</b> hypothetical transcriptional regulator ycdc; <b>PDBTitle:</b> crystal structure of hypothetical transcriptional regulator ycdc
40	<a href="#">c3rh2A</a>	 Alignment	not modelled	99.9	14	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> hypothetical tet-like transcriptional regulator; <b>PDBTitle:</b> crystal structure of a hypothetical tet-like transcriptional2 regulator (sama_0099) from shewanella amazonensis sb2b at 2.42 a3 resolution
41	<a href="#">c2iaiA</a>	 Alignment	not modelled	99.9	16	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> putative transcriptional regulator sco3833; <b>PDBTitle:</b> crystal structure of sco3833, a member of the tet transcriptional2 regulator family from streptomyces coelicolor a3
42	<a href="#">c3cwrA</a>	 Alignment	not modelled	99.9	11	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> crystal structure of transcriptional regulator of tet2 family (yp_425770.1) from rhodospirillum rubrum atcc 111703 at 1.50 a resolution
43	<a href="#">c1jumB</a>	 Alignment	not modelled	99.9	13	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> hypothetical transcriptional regulator in qaca <b>PDBTitle:</b> crystal structure of the multidrug binding transcriptional2 repressor qacr bound to the natural drug berberine
44	<a href="#">c3dcfB</a>	 Alignment	not modelled	99.9	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator of the tet/acrr <b>PDBTitle:</b> crystal structure of transcriptional regulator of the2 tet/acrr family (yp_290855.1) from thermobifida fusca yx-3 er1 at 2.50 a resolution
45	<a href="#">c3f0cA</a>	 Alignment	not modelled	99.9	14	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of transcriptional regulator from cytophaga2 hutchinsonii atcc 33406
46	<a href="#">c2id3A</a>	 Alignment	not modelled	99.9	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative transcriptional regulator; <b>PDBTitle:</b> crystal structure of transcriptional regulator sco5951 from2 streptomyces coelicolor a3(2)
47	<a href="#">c3dewA</a>	 Alignment	not modelled	99.9	20	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> the structure of a putative tet family transcriptional regulator from2 geobacter sulfurreducens pca.
48	<a href="#">c2wuiA</a>	 Alignment	not modelled	99.9	17	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of mexz, a key repressor responsible for2 antibiotic resistance in pseudomonas aeruginosa.
49	<a href="#">c2fq4A</a>	 Alignment	not modelled	99.9	17	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tet family; <b>PDBTitle:</b> the crystal structure of the transcriptional regulator (tet family)2 from bacillus cereus
50	<a href="#">c2hvtA</a>	 Alignment	not modelled	99.9	16	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetr-family transcriptional regulator;

50	<a href="#">c2lyvA</a>	Alignment	not modelled	99.8	10	<b>PDBTitle:</b> crystal structure of a tetr-family transcriptional regulator (eca1819)2 from <i>pectobacterium atrosepticum</i> at 1.64 a resolution <b>PDB header:</b> transcription regulator
51	<a href="#">c3he0A</a>	Alignment	not modelled	99.8	18	<b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> the structure of a putative transcriptional regulator tetr family2 protein from <i>vibrio parahaemolyticus</i> .
52	<a href="#">c1ui6B</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> antibiotic <b>Chain:</b> B: <b>PDB Molecule:</b> a-factor receptor homolog; <b>PDBTitle:</b> crystal structure of gamma-butyrolactone receptor (arpa-like protein)
53	<a href="#">c2ibdB</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> B: <b>PDB Molecule:</b> possible transcriptional regulator; <b>PDBTitle:</b> crystal structure of probable transcriptional regulatory protein2 rha5900
54	<a href="#">c3jsjC</a>	Alignment	not modelled	99.8	23	<b>PDB header:</b> transcription <b>Chain:</b> C: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a putative tetr-transcriptional regulator2 (sav143) from <i>streptomyces avermitilis</i> ma-4680 at 2.10 a resolution
55	<a href="#">c2fbgA</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> probable transcriptional regulator; <b>PDBTitle:</b> the crystal structure of transcriptional regulator pa3006
56	<a href="#">c3mnlA</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulatory protein (probably tetr-family); <b>PDBTitle:</b> the crystal structure of kstr (rv3574) from <i>mycobacterium tuberculosis</i> 2 h37rv
57	<a href="#">c2raeA</a>	Alignment	not modelled	99.8	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, acrr family protein; <b>PDBTitle:</b> crystal structure of a tetr/acrr family transcriptional regulator from2 <i>rhodococcus</i> sp. rha1
58	<a href="#">c2np5A</a>	Alignment	not modelled	99.8	22	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of a transcriptional regulator (rha1_ro04179) from2 <i>rhodococcus</i> sp. rha1.
59	<a href="#">c2dg8D</a>	Alignment	not modelled	99.8	18	<b>PDB header:</b> gene regulation <b>Chain:</b> D: <b>PDB Molecule:</b> putative tetr-family transcriptional regulatory protein; <b>PDBTitle:</b> crystal structure of the putative transcriptional regulator sco75182 from <i>streptomyces coelicolor</i> a3(2)
60	<a href="#">c3npiB</a>	Alignment	not modelled	99.8	10	<b>PDB header:</b> gene regulation <b>Chain:</b> B: <b>PDB Molecule:</b> tetr family regulatory protein; <b>PDBTitle:</b> crystal structure of a tetr family regulatory protein (dip1788) from2 <i>corynebacterium diphtheriae</i> at 2.96 a resolution
61	<a href="#">c2fd5A</a>	Alignment	not modelled	99.8	23	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> the crystal structure of a transcriptional regulator from <i>pseudomonas aeruginosa</i> pao1
62	<a href="#">c2o7tA</a>	Alignment	not modelled	99.8	10	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr family transcriptional regulator2 (ncgl1578, cgl1640) from <i>corynebacterium glutamicum</i> at 2.10 a3 resolution
63	<a href="#">c3g56A</a>	Alignment	not modelled	99.8	17	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> regulator of macrolide 2'-phosphotransferase i; <b>PDBTitle:</b> structure of the macrolide biosensor protein, mphr(a)
64	<a href="#">c2qwtA</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of the tetr transcription regulatory2 protein from <i>mycobacterium vanbaalenii</i>
65	<a href="#">c3ccvA</a>	Alignment	not modelled	99.8	19	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr-family transcriptional regulator from2 <i>bordetella parapertussis</i> 12822
66	<a href="#">c3mvpA</a>	Alignment	not modelled	99.8	12	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> tetr/acrr transcriptional regulator; <b>PDBTitle:</b> the crystal structure of a tetr/acrr transcriptional regulator from2 <i>streptococcus mutans</i> to 1.85a
67	<a href="#">c2qibA</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr-family transcriptional regulator from2 <i>streptomyces coelicolor</i>
68	<a href="#">c3nnrA</a>	Alignment	not modelled	99.8	11	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of a tetr-family transcriptional regulator2 (maqu_3571) from <i>marinobacter aquaeolei</i> vt8 at 2.49 a resolution
69	<a href="#">c3f1bA</a>	Alignment	not modelled	99.8	14	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> tetr-like transcriptional regulator; <b>PDBTitle:</b> the crystal structure of a tetr-like transcriptional regulator from2 <i>rhodococcus</i> sp. rha1.
70	<a href="#">c3b81A</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, acrr family; <b>PDBTitle:</b> crystal structure of predicted dna-binding transcriptional regulator2 of tetr/acrr family (np_350189.1) from <i>clostridium acetobutylicum</i> at3 2.10 a resolution
71	<a href="#">c1t33B</a>	Alignment	not modelled	99.8	12	<b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> putative transcriptional repressor (tetr/acrr family); <b>PDBTitle:</b> structural genomics, the crystal structure of a putative2 transcriptional repressor (tetr/acrr family) from <i>salmonella</i> 3 typhimurim lt2
72	<a href="#">c3bjbE</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> transcription regulator <b>Chain:</b> E: <b>PDB Molecule:</b> probable transcriptional regulator, tetr family protein; <b>PDBTitle:</b> crystal structure of a tetr transcriptional regulator from <i>rhodococcus</i> 2 sp. rha1
						<b>PDB header:</b> transcription

73	<a href="#">c2rekA</a>	Alignment	not modelled	99.8	9	<b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr-family transcriptional regulator
74	<a href="#">c3c07B</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr family transcriptional regulator from2 streptomyces coelicolor a3(2)
75	<a href="#">c2q24A</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr family transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr transcriptional regulator sco0520 from2 streptomyces coelicolor
76	<a href="#">c3cjdB</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of putative tetr transcriptional regulator2 (yp_510936.1) from jannaschia sp. ccs1 at 1.79 a resolution
77	<a href="#">c3aqsD</a>	Alignment	not modelled	99.8	18	<b>PDB header:</b> transcription regulator <b>Chain:</b> D: <b>PDB Molecule:</b> bacterial regulatory proteins, tetr family; <b>PDBTitle:</b> crystal structure of rolr (ncgl1110) without ligand
78	<a href="#">c3c2bA</a>	Alignment	not modelled	99.8	13	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of tetr transcriptional regulator from agrobacterium2 tumefaciens
79	<a href="#">c2qcoA</a>	Alignment	not modelled	99.8	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> cmer; <b>PDBTitle:</b> crystal structure of the transcriptional regulator cmer from2 campylobacter jejuni
80	<a href="#">c2oi8A</a>	Alignment	not modelled	99.8	12	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative regulatory protein sco4313; <b>PDBTitle:</b> crystal structure of putative regulatory protein sco4313
81	<a href="#">c3qkxB</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> uncharacterized hth-type transcriptional regulator hi_0893; <b>PDBTitle:</b> crystal structure of a tetr-family transcriptional regulator (hi0893)2 from haemophilus influenzae rd at 2.35 a resolution
82	<a href="#">c2d6yA</a>	Alignment	not modelled	99.8	14	<b>PDB header:</b> gene regulation <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr family regulatory protein; <b>PDBTitle:</b> crystal structure of transcriptional factor sco4008 from streptomyces2 coelicolor a3(2)
83	<a href="#">c3iuvA</a>	Alignment	not modelled	99.8	17	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized tetr family protein; <b>PDBTitle:</b> the structure of a member of tetr family (sco1917) from2 streptomyces coelicolor a3
84	<a href="#">c2jk3A</a>	Alignment	not modelled	99.8	14	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> hemolysin ii regulatory protein; <b>PDBTitle:</b> crystal structure of the hlyiir mutant protein with2 residues 169-186 substituted by gssgssg linker
85	<a href="#">c2dg7A</a>	Alignment	not modelled	99.8	14	<b>PDB header:</b> gene regulation <b>Chain:</b> A: <b>PDB Molecule:</b> putative transcriptional regulator; <b>PDBTitle:</b> crystal structure of the putative transcriptional regulator sco03372 from streptomyces coelicolor a3(2)
86	<a href="#">c3lsjA</a>	Alignment	not modelled	99.8	18	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> dest; <b>PDBTitle:</b> crystal structure of dest in complex with palmitoyl-coa
87	<a href="#">c2of7A</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> structural genomics, the crystal structure of a tetr-family2 transcriptional regulator from streptomyces coelicolor a3
88	<a href="#">c3crjC</a>	Alignment	not modelled	99.8	14	<b>PDB header:</b> transcription regulator <b>Chain:</b> C: <b>PDB Molecule:</b> transcription regulator; <b>PDBTitle:</b> crystal structure of a tetr transcription regulator from haloarcula2 marismortui atcc 43049
89	<a href="#">c2zcxA</a>	Alignment	not modelled	99.8	13	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr family transcriptional regulator2 sco7815
90	<a href="#">c3geuC</a>	Alignment	not modelled	99.8	15	<b>PDB header:</b> cell adhesion <b>Chain:</b> C: <b>PDB Molecule:</b> intercellular adhesion protein r; <b>PDBTitle:</b> crystal structure of icar from staphylococcus aureus, a member of the2 tetracycline repressor protein family
91	<a href="#">c3colB</a>	Alignment	not modelled	99.8	11	<b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative transcription regulator; <b>PDBTitle:</b> crystal structure of putative transcription regulator from2 lactobacillus plantarum
92	<a href="#">c3cdlA</a>	Alignment	not modelled	99.8	17	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator aeFr; <b>PDBTitle:</b> crystal structure of a tetr family transcriptional regulator from2 pseudomonas syringae pv. tomato str. dc3000
93	<a href="#">c1zk8A</a>	Alignment	not modelled	99.8	17	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of transcriptional regulator from bacillus cereus2 atcc 14579
94	<a href="#">c2yvH A</a>	Alignment	not modelled	99.8	12	<b>PDB header:</b> transcription/dna <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of the operator-binding form of the multi-drug2 binding transcriptional repressor cgmR
95	<a href="#">c2eh3A</a>	Alignment	not modelled	99.8	20	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> crystal structure of aq_1058, a transcriptional regulator (terr/acrr2 family) from aquifex aeolicus vf5
96	<a href="#">c3nxC A</a>	Alignment	not modelled	99.8	16	<b>PDB header:</b> dna binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> hth-type protein slma; <b>PDBTitle:</b> molecular mechanism by which the escherichia coli nucleoid occlusion2 factor, slma, keeps cytokinesis in check
97	<a href="#">c3bniA</a>	Alignment	not modelled	99.8	18	<b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of tetr-family transcriptional regulator



					from2 streptomyces coelicolor
98	<a href="#">c2pz9A_</a>	Alignment	not modelled	99.8	19 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative regulatory protein; <b>PDBTitle:</b> crystal structure of putative transcriptional regulator sco4942 from2 streptomyces coelicolor
99	<a href="#">c2zcmB_</a>	Alignment	not modelled	99.8	13 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> biofilm operon icaabcd hth-type negative transcriptional <b>PDBTitle:</b> crystal structure of icar, a repressor of the tetr family
100	<a href="#">c3kkcB_</a>	Alignment	not modelled	99.8	12 <b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> tetr family transcriptional regulator; <b>PDBTitle:</b> the crystal structure of tetr transcriptional regulator from2 streptococcus agalactiae 2603v
101	<a href="#">c2iu5A_</a>	Alignment	not modelled	99.7	13 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> hth-type dhaklm operon transcriptional activator dhas; <b>PDBTitle:</b> dihydroxyacetone kinase operon activator dhas
102	<a href="#">c3ni7A_</a>	Alignment	not modelled	99.7	12 <b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> bacterial regulatory proteins, tetr family; <b>PDBTitle:</b> crystal structure of the tetr transcriptional regulator from2 nitrosomonas europaea atcc 19718
103	<a href="#">c3kkdA_</a>	Alignment	not modelled	99.7	15 <b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator; <b>PDBTitle:</b> structure of a putative tetr transcriptional regulator (pa3699) from2 pseudomonas aeruginosa pa01
104	<a href="#">c3egqB_</a>	Alignment	not modelled	99.7	14 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> tetr family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a tetr-family transcriptional regulator (af_1817)2 from archaeoglobus fulgidus at 2.55 a resolution
105	<a href="#">c3o60A_</a>	Alignment	not modelled	99.7	11 <b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> lin0861 protein; <b>PDBTitle:</b> the crystal structure of lin0861 from listeria innocua to 2.8a
106	<a href="#">c1z0xA_</a>	Alignment	not modelled	99.7	13 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional regulator, tetr family; <b>PDBTitle:</b> crystal structure of transcriptional regulator, tetr family from2 enterococcus faecalis v583
107	<a href="#">c3bqyA_</a>	Alignment	not modelled	99.7	13 <b>PDB header:</b> transcription regulator <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr family transcriptional regulator; <b>PDBTitle:</b> crystal structure of a possible tetr family transcriptional regulator2 from streptomyces coelicolor a3(2).
108	<a href="#">c3fiwB_</a>	Alignment	not modelled	99.7	16 <b>PDB header:</b> transcription regulator <b>Chain:</b> B: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> structure of sco0253, a tetr-family transcriptional regulator from2 streptomyces coelicolor
109	<a href="#">c2qkoA_</a>	Alignment	not modelled	99.7	15 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> possible transcriptional regulator, tetr family protein; <b>PDBTitle:</b> crystal structure of transcriptional regulator rha06399 from2 rhodococcus sp. rha1
110	<a href="#">c1bjzA_</a>	Alignment	not modelled	99.7	16 <b>PDB header:</b> transcription regulation <b>Chain:</b> A: <b>PDB Molecule:</b> tetracycline repressor; <b>PDBTitle:</b> tetracycline chelated mg2+-ion initiates helix unwinding for tet2 repressor induction
111	<a href="#">c2hxoB_</a>	Alignment	not modelled	99.7	14 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> structure of the transcriptional regulator sco7222, a tetr from2 streptomyces coelicolor
112	<a href="#">c3b6aC_</a>	Alignment	not modelled	99.7	12 <b>PDB header:</b> transcription <b>Chain:</b> C: <b>PDB Molecule:</b> actr protein; <b>PDBTitle:</b> crystal structure of the streptomyces coelicolor tetr2 family protein actr in complex with actinorhodin
113	<a href="#">c2g7lA_</a>	Alignment	not modelled	99.7	15 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of putative transcription regulator sco7704 from2 streptomyces coelicolor
114	<a href="#">c2guhA_</a>	Alignment	not modelled	99.7	16 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative tetr-family transcriptional regulator; <b>PDBTitle:</b> crystal structure of the putative tetr-family transcriptional2 regulator from rhodococcus sp. rha1
115	<a href="#">c2y31A_</a>	Alignment	not modelled	99.7	10 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative repressor simreg2; <b>PDBTitle:</b> simocyclinone c4 bound form of tetr-like repressor simr
116	<a href="#">c2vprA_</a>	Alignment	not modelled	99.7	13 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> tetracycline resistance repressor protein; <b>PDBTitle:</b> tet repressor class h in complex with 5a,6-2 anhydrotetracycline-mg
117	<a href="#">c2hxiA_</a>	Alignment	not modelled	99.7	17 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> putative transcriptional regulator; <b>PDBTitle:</b> structural genomics, the crystal structure of a putative2 transcriptional regulator from streptomyces coelicolor a3(2)
118	<a href="#">c2w53B_</a>	Alignment	not modelled	99.6	18 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> repressor; <b>PDBTitle:</b> structure of smet, the repressor of the stenotrophomonas2 maltophilia multidrug efflux pump smedef.
119	<a href="#">c2wgbB_</a>	Alignment	not modelled	99.6	12 <b>PDB header:</b> transcription <b>Chain:</b> B: <b>PDB Molecule:</b> tetr family transcriptional repressor lfrr; <b>PDBTitle:</b> crystal structure of the tetr-like transcriptional2 regulator lfrr from mycobacterium smegmatis
120	<a href="#">c2oerA_</a>	Alignment	not modelled	99.6	19 <b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> probable transcriptional regulator; <b>PDBTitle:</b> probable transcriptional regulator from pseudomonas2 aeruginosa