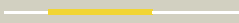
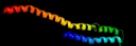
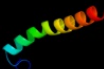
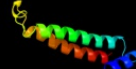
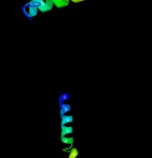
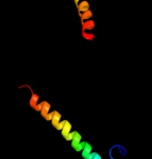
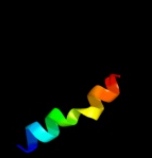
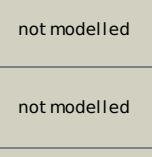


Phyre2

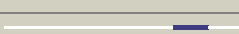
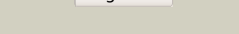
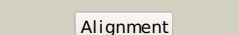
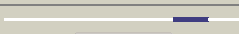
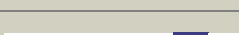
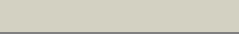
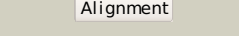
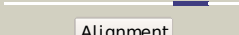
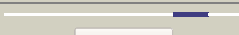
Email	I.a.kelley@imperial.ac.uk
Description	P0ABU7
Date	Thu Jan 5 11:16:26 GMT 2012
Unique Job ID	b7def0568a8a0d97

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2bbjB_	 Alignment		72.1	12	PDB header: metal transport/membrane protein Chain: B: PDB Molecule: divalent cation transport-related protein; PDBTitle: crystal structure of the cora mg2+ transporter
2	c1ciiA_	 Alignment		63.0	15	PDB header: transmembrane protein Chain: A: PDB Molecule: colicin ia; PDBTitle: colicin ia
3	d1t33a1	 Alignment		40.6	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
4	d1vi0a1	 Alignment		38.5	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
5	d2fbqa1	 Alignment		38.4	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
6	c2ap7A_	 Alignment		33.9	46	PDB header: antibiotic Chain: A: PDB Molecule: bombinin h2; PDBTitle: solution structure of bombinin h2 in dpc micelles
7	c2fbqA_	 Alignment		33.5	10	PDB header: transcription Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: the crystal structure of transcriptional regulator pa3006
8	d1o5ha_	 Alignment		33.4	9	Fold: Methenyltetrahydrofolate cyclohydrolase-like Superfamily: Methenyltetrahydrofolate cyclohydrolase-like Family: Methenyltetrahydrofolate cyclohydrolase-like
9	c2ap8A_	 Alignment		29.4	46	PDB header: antibiotic Chain: A: PDB Molecule: bombinin h4; PDBTitle: solution structure of bombinin h4 in dpc micelles
10	d1ji6a3	 Alignment		25.2	14	Fold: Toxins' membrane translocation domains Superfamily: delta-Endotoxin (insecticide), N-terminal domain Family: delta-Endotoxin (insecticide), N-terminal domain
11	d2o7ta1	 Alignment		25.1	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain

12	c1tt9B_	Alignment		23.3	12	PDB header: transferase, lyase Chain: B: PDB Molecule: formimidoyltransferase-cyclodeaminase PDBTitle: structure of the bifunctional and golgi associated2 formiminotransferase cyclodeaminase octamer
13	d1jt6a1	Alignment		22.3	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
14	d1dlca3	Alignment		22.2	10	Fold: Toxins' membrane translocation domains Superfamily: delta-Endotoxin (insectocide), N-terminal domain Family: delta-Endotoxin (insectocide), N-terminal domain
15	d1zk8a1	Alignment		20.1	9	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
16	d2gena1	Alignment		17.5	12	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
17	c2kncB_	Alignment		17.3	19	PDB header: cell adhesion Chain: B: PDB Molecule: integrin beta-3; PDBTitle: platelet integrin alfaIib-beta3 transmembrane-cytoplasmic2 heterocomplex
18	d2gfna1	Alignment		17.1	17	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
19	d2axtj1	Alignment		16.9	28	Fold: Single transmembrane helix Superfamily: Photosystem II reaction center protein J, Psbj Family: Psbj-like
20	c3f1bA_	Alignment		16.1	14	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.
21	d2fd5a1	Alignment	not modelled	14.9	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
22	c2kncA_	Alignment	not modelled	14.7	22	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iib; PDBTitle: platelet integrin alfaIib-beta3 transmembrane-cytoplasmic2 heterocomplex
23	d2np5a1	Alignment	not modelled	14.0	18	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
24	d2id6a1	Alignment	not modelled	13.9	11	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
25	d1rkta1	Alignment	not modelled	13.7	13	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
26	d3c07a1	Alignment	not modelled	12.7	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
27	c3gziA_	Alignment	not modelled	12.7	16	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
28	c3ppbB_	Alignment	not modelled	12.6	6	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
29	d1sgma1	Alignment	not modelled	12.2	15 Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
30	c2wgbB	Alignment	not modelled	12.2	19 PDB header: transcription Chain: B: PDB Molecule: tetr family transcriptional repressor Ifrr; PDBTitle: crystal structure of the tetr-like transcriptional2 regulator Ifrr from mycobacterium smegmatis
31	d2i10a1	Alignment	not modelled	11.9	15 Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
32	c3mn1A	Alignment	not modelled	11.6	17 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
33	c1t33B	Alignment	not modelled	11.3	12 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
34	c2rekA	Alignment	not modelled	11.2	16 PDB header: transcription Chain: A: PDB Molecule: putative tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator
35	c3jsjC	Alignment	not modelled	11.1	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
36	c1u9nA	Alignment	not modelled	11.0	15 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
37	c2g7sA	Alignment	not modelled	11.0	6 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the crystal structure of transcriptional regulator, tetr family, from2 agrobacterium tumefaciens
38	c2hyjA	Alignment	not modelled	10.8	8 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
39	c1ji6A	Alignment	not modelled	10.8	14 PDB header: toxin Chain: A: PDB Molecule: pesticidal crystal protein cry3bb; PDBTitle: crystal structure of the insecticidal bacterial del2 endotoxin cry3bb1 bacillus thuringiensis
40	c3aqsD	Alignment	not modelled	10.4	19 PDB header: transcription regulator Chain: D: PDB Molecule: bacterial regulatory proteins, tetr family; PDBTitle: crystal structure of rolr (ncgl1110) without ligand
41	c3iuvA	Alignment	not modelled	10.2	15 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
42	c3c07B	Alignment	not modelled	10.1	19 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)
43	c1vi0B	Alignment	not modelled	10.1	17 PDB header: transcription Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator
44	c2genA	Alignment	not modelled	9.6	14 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
45	c3djpB	Alignment	not modelled	9.5	11 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
46	c3knwB	Alignment	not modelled	9.5	14 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)
47	c1ui6B	Alignment	not modelled	9.5	19 PDB header: antibiotic Chain: B: PDB Molecule: a-factor receptor homolog; PDBTitle: crystal structure of gamma-butyrolactone receptor (arpa-like protein)
48	c3eupA	Alignment	not modelled	9.3	14 PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: the crystal structure of the transcriptional regulator, tetr family2 from cytophaga hutchinsonii
49	c2qibA	Alignment	not modelled	9.1	12 PDB header: transcription Chain: A: PDB Molecule: tetr-family transcriptional regulator; PDBTitle: crystal structure of tetr-family transcriptional regulator from2 streptomyces coelicolor
50	c2oerA	Alignment	not modelled	8.8	14 PDB header: transcription Chain: A: PDB Molecule: probable transcriptional regulator; PDBTitle: probable transcriptional regulator from pseudomonas2 aeruginosa
51	c3qbmA	Alignment	not modelled	8.7	19 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

52	c2q24A	 Alignment	not modelled	8.7	14	PDB header: transcription Chain: A: PDB Molecule: putative tetr family transcriptional regulator; PDBTitle: crystal structure of tetr transcriptional regulator sco0520 from2 streptomyces coelicolor
53	c2rasB	 Alignment	not modelled	8.6	8	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
54	c2gfnA	 Alignment	not modelled	8.5	16	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
55	c2k1aA	 Alignment	not modelled	8.3	23	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iiib; PDBTitle: bicelle-embedded integrin alpha(iiib) transmembrane segment
56	d1q8da	 Alignment	not modelled	8.3	63	Fold: GDNF receptor-like Superfamily: GDNF receptor-like Family: GDNF receptor-like
57	c2hytA	 Alignment	not modelled	8.2	17	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
58	d1t56a1	 Alignment	not modelled	8.0	15	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
59	c2eh3A	 Alignment	not modelled	7.6	9	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
60	c1rktA	 Alignment	not modelled	7.5	11	PDB header: transcription Chain: A: PDB Molecule: protein yfir; PDBTitle: crystal structure of yfir, a putative transcriptional regulator from2 bacillus subtilis
61	c2pz9A	 Alignment	not modelled	7.5	9	PDB header: transcription Chain: A: PDB Molecule: putative regulatory protein; PDBTitle: crystal structure of putative transcriptional regulator sco4942 from2 streptomyces coelicolor
62	c2iaiA	 Alignment	not modelled	7.4	6	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
63	c3lwlA	 Alignment	not modelled	7.4	28	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
64	c2ibdB	 Alignment	not modelled	7.3	8	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: possible transcriptional regulator; PDBTitle: crystal structure of probable transcriptional regulatory protein2 rha5900
65	c3f0cA	 Alignment	not modelled	7.3	14	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of transcriptional regulator from cytophaga2 hutchinsonii atcc 33406
66	c2o7tA	 Alignment	not modelled	7.3	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
67	c2dg8D	 Alignment	not modelled	7.3	11	PDB header: gene regulation Chain: D: PDB Molecule: putative tetr-family transcriptional regulatory protein; PDBTitle: crystal structure of the putative transcripional regulator sco75182 from streptomyces coelicolor a3(2)
68	c3himA	 Alignment	not modelled	7.1	11	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
69	c3dcfB	 Alignment	not modelled	7.0	17	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator of the tetr/acrr PDBTitle: crystal structure of transcriptional regulator of the2 tetr/acrr family (yp_290855.1) from thermobifida fusca yx-3 er1 at 2.50 a resolution
70	d2g3ba1	 Alignment	not modelled	7.0	21	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
71	c3e7qB	 Alignment	not modelled	7.0	14	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulator; PDBTitle: the crystal structure of the putative transcriptional regulator from2 pseudomonas aeruginosa pao1
72	c1bttA	 Alignment	not modelled	7.0	45	PDB header: transmembrane protein Chain: A: PDB Molecule: band 3 anion transport protein; PDBTitle: the solution structures of the first and second2 transmembrane-spanning segments of band 3
73	c1btsA	 Alignment	not modelled	7.0	45	PDB header: transmembrane protein Chain: A: PDB Molecule: band 3 anion transport protein; PDBTitle: the solution structures of the first and second2 transmembrane-spanning segments of band 3
74	c3anpD	 Alignment	not modelled	6.9	11	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.
75	c2v5eA	 Alignment	not modelled	6.8	42	PDB header: receptor/glycoprotein complex Chain: A: PDB Molecule: gdnf family receptor alpha-1; PDBTitle: the structure of the gdnf:coreceptor complex: insights2

						into ret signalling and heparin binding.
76	c3fubA	Alignment	not modelled	6.8	42	PDB header: hormone Chain: A: PDB Molecule: gdnf family receptor alpha-1; PDBTitle: crystal structure of gdnf-gfralpha1 complex
77	d2oi8a1	Alignment	not modelled	6.7	9	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
78	c3nxcA	Alignment	not modelled	6.6	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
79	c1jumB	Alignment	not modelled	6.5	11	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
80	c3ccyA	Alignment	not modelled	6.5	14	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
81	d1l9bm	Alignment	not modelled	6.5	9	Fold: Bacterial photosystem II reaction centre, L and M subunits Superfamily: Bacterial photosystem II reaction centre, L and M subunits Family: Bacterial photosystem II reaction centre, L and M subunits
82	c3rh2A	Alignment	not modelled	6.5	14	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
83	c3nnrA	Alignment	not modelled	6.4	11	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
84	c3geuC	Alignment	not modelled	6.4	5	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
85	c3lsjA	Alignment	not modelled	6.3	14	PDB header: transcription Chain: A: PDB Molecule: dest; PDBTitle: crystal structure of dest in complex with palmitoyl-coa
86	c2l4gA	Alignment	not modelled	6.3	45	PDB header: viral protein Chain: A: PDB Molecule: haemagglutinin; PDBTitle: influenza haemagglutinin fusion peptide mutant g13a
87	c2np5A	Alignment	not modelled	6.2	14	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator; PDBTitle: crystal structure of a transcriptional regulator (rha1_ro04179) from2 rhodococcus sp. rha1.
88	d2d6ya1	Alignment	not modelled	6.2	16	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
89	c1zkgB	Alignment	not modelled	6.1	10	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
90	c3on4D	Alignment	not modelled	6.1	11	PDB header: transcription Chain: D: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of tetr transcriptional regulator from legionella2 pneumophila
91	d1pb6a1	Alignment	not modelled	6.0	14	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
92	c3crjC	Alignment	not modelled	6.0	8	PDB header: transcription regulator Chain: C: PDB Molecule: transcription regulator; PDBTitle: crystal structure of a tetr transcription regulator from haloarcula2 marismortui atcc 43049
93	c2zcmB	Alignment	not modelled	5.9	6	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
94	c3mvpA	Alignment	not modelled	5.9	21	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
95	c2dcia	Alignment	not modelled	5.8	45	PDB header: viral protein Chain: A: PDB Molecule: hemagglutinin; PDBTitle: nmr structure of influenza ha fusion peptide mutant w14a in2 dpc in ph5
96	c2qwtA	Alignment	not modelled	5.8	16	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator, tetr family; PDBTitle: crystal structure of the tetr transcription regulatory2 protein from mycobacterium vanbaalenii
97	c2qtqB	Alignment	not modelled	5.7	16	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
98	d1v7ba1	Alignment	not modelled	5.5	10	Fold: DNA/RNA-binding 3-helical bundle Superfamily: Homeodomain-like Family: Tetracyclin repressor-like, N-terminal domain
99	c2i10A	Alignment	not modelled	5.5	14	PDB header: transcription Chain: A: PDB Molecule: putative tetr transcriptional regulator; PDBTitle: putative tetr transcriptional regulator from rhodococcus sp. rha1