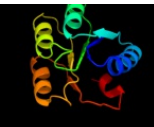
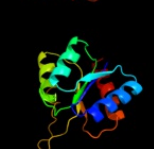


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1ys7B_	 Alignment		100.0	35	PDB header: transcription regulator Chain: B: PDB Molecule: transcriptional regulatory protein prra; PDBTitle: crystal structure of the response regulator protein prra complexed with2 mg2 +
2	c2oqrA_	 Alignment		100.0	37	PDB header: transcription,signaling protein Chain: A: PDB Molecule: sensory transduction protein regx3; PDBTitle: the structure of the response regulator regx3 from mycobacterium2 tuberculosis
3	c3r0jA_	 Alignment		100.0	39	PDB header: dna binding protein Chain: A: PDB Molecule: possible two component system response transcriptional PDBTitle: structure of phop from mycobacterium tuberculosis
4	c2gwrA_	 Alignment		100.0	35	PDB header: signaling protein Chain: A: PDB Molecule: dna-binding response regulator mtra; PDBTitle: crystal structure of the response regulator protein mtra from2 mycobacterium tuberculosis
5	c1kgsA_	 Alignment		100.0	41	PDB header: dna binding protein Chain: A: PDB Molecule: dna binding response regulator d; PDBTitle: crystal structure at 1.50 a of an ompr/phob homolog from thermotoga2 maritima
6	c2hqrA_	 Alignment		100.0	28	PDB header: signaling protein Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: structure of a atypical orphan response regulator protein revealed a2 new phosphorylation-independent regulatory mechanism
7	c1p2fA_	 Alignment		100.0	27	PDB header: transcription Chain: A: PDB Molecule: response regulator; PDBTitle: crystal structure analysis of response regulator drrb, a2 thermotoga maritima ompr/phob homolog
8	c3q9sA_	 Alignment		100.0	43	PDB header: dna binding protein Chain: A: PDB Molecule: dna-binding response regulator; PDBTitle: crystal structure of rra(1-215) from deinococcus radiodurans
9	c1zn2A_	 Alignment		100.0	24	PDB header: transcription regulator Chain: A: PDB Molecule: response regulatory protein; PDBTitle: low resolution structure of response regulator styr
10	c3c3wB_	 Alignment		99.9	21	PDB header: transcription Chain: B: PDB Molecule: two component transcriptional regulatory protein devr; PDBTitle: crystal structure of the mycobacterium tuberculosis hypoxic response2 regulator dosr
11	c2ayxA_	 Alignment		99.9	25	PDB header: transferase Chain: A: PDB Molecule: sensor kinase protein rcsc; PDBTitle: solution structure of the e.coli rcsc c-terminus (residues2 700-949) containing linker region and phosphoreceiver3 domain

12	c3eq2A	Alignment		99.9	33	PDB header: signaling protein Chain: A: PDB Molecule: probable two-component response regulator; PDBTitle: structure of hexagonal crystal form of pseudomonas2 aeruginosa rssb
13	c1rn1A	Alignment		99.9	30	PDB header: signal transduction protein Chain: A: PDB Molecule: nitrate/nitrite response regulator protein narl; PDBTitle: the nitrate/nitrite response regulator protein narl from narl
14	c3breA	Alignment		99.9	26	PDB header: signaling protein Chain: A: PDB Molecule: probable two-component response regulator; PDBTitle: crystal structure of p.aeruginosa pa3702
15	c1ny5A	Alignment		99.9	28	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator (ntrc family); PDBTitle: crystal structure of sigm54 activator (aaa+ atpase) in the inactive2 state
16	c1w25B	Alignment		99.9	29	PDB header: signaling protein Chain: B: PDB Molecule: stalked-cell differentiation controlling protein; PDBTitle: response regulator pled in complex with c-digmp
17	c3rq1A	Alignment		99.9	16	PDB header: transcription Chain: A: PDB Molecule: response regulator protein; PDBTitle: crystal structure of a response regulator protein from burkholderia2 pseudomallei with a phosphorylated aspartic acid, calcium ion and3 citrate
18	d1s8na	Alignment		99.9	23	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
19	d1ys7a2	Alignment		99.9	37	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
20	d1ny5a1	Alignment		99.9	28	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
21	d1kgsa2	Alignment	not modelled	99.9	46	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
22	c3dzdA	Alignment	not modelled	99.9	19	PDB header: transcription regulator Chain: A: PDB Molecule: transcriptional regulator (ntrc family); PDBTitle: crystal structure of sigma54 activator ntrc4 in the inactive2 state
23	d2a9pa1	Alignment	not modelled	99.9	41	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
24	c3i5aA	Alignment	not modelled	99.9	26	PDB header: signaling protein Chain: A: PDB Molecule: response regulator/ggdef domain protein; PDBTitle: crystal structure of full-length wpsr from pseudomonas syringae
25	d1mvoa	Alignment	not modelled	99.9	35	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
26	d1jbea	Alignment	not modelled	99.9	25	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
27	d2pl1a1	Alignment	not modelled	99.9	39	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
28	d2ayxa1	Alignment	not modelled	99.9	26	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
29	d1peya	Alignment	not modelled	99.9	27	Fold: Flavodoxin-like Superfamily: CheY-like

					Family: CheY-related
30	c2jrlA	Alignment	not modelled	99.9	19 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulator (ntrc family); PDBTitle: solution structure of the beryllofluoride-activated ntrc4 receiver2 domain dimer
31	d1krwa	Alignment	not modelled	99.9	30 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
32	c3hdgE	Alignment	not modelled	99.9	22 PDB header: structural genomics, unknown function Chain: E: PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of the n-terminal domain of an2 uncharacterized protein (ws1339) from wolinetlla3 succinogenes
33	c2zwmA	Alignment	not modelled	99.9	38 PDB header: transcription Chain: A: PDB Molecule: transcriptional regulatory protein yycf; PDBTitle: crystal structure of yycf receiver domain from bacillus2 subtilis
34	d1dbwa	Alignment	not modelled	99.9	19 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
35	c3cfyA	Alignment	not modelled	99.9	25 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative luxo repressor protein; PDBTitle: crystal structure of signal receiver domain of putative luxo2 repressor protein from vibrio parahaemolyticus
36	d1qkka	Alignment	not modelled	99.9	24 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
37	d1w25a1	Alignment	not modelled	99.9	29 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
38	c2rjnA	Alignment	not modelled	99.9	19 PDB header: hydrolase Chain: A: PDB Molecule: response regulator receiver:metal-dependent PDBTitle: crystal structure of an uncharacterized protein q2bku2 from2 neptuniibacter caesariensis
39	c3cu5B	Alignment	not modelled	99.9	18 PDB header: transcription regulator Chain: B: PDB Molecule: two component transcriptional regulator, arac family; PDBTitle: crystal structure of a two component transcriptional regulator arac2 from clostridium phytofermentans isdg
40	c3nhzA	Alignment	not modelled	99.9	37 PDB header: dna binding protein Chain: A: PDB Molecule: two component system transcriptional regulator mtra; PDBTitle: structure of n-terminal domain of mtra
41	c2qr3A	Alignment	not modelled	99.9	20 PDB header: transcription Chain: A: PDB Molecule: two-component system response regulator; PDBTitle: crystal structure of the n-terminal signal receiver domain of two-2 component system response regulator from bacteroides fragilis
42	c3gt7A	Alignment	not modelled	99.9	27 PDB header: hydrolase Chain: A: PDB Molecule: sensor protein; PDBTitle: crystal structure of signal receiver domain of signal2 transduction histidine kinase from syntrophus3 aciditrophicus
43	c3eulB	Alignment	not modelled	99.9	25 PDB header: transcription Chain: B: PDB Molecule: possible nitrate/nitrite response transcriptional PDBTitle: structure of the signal receiver domain of the putative2 response regulator narl from mycobacterium tuberculosis
44	d1u0sy	Alignment	not modelled	99.9	27 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
45	c3b2nA	Alignment	not modelled	99.9	21 PDB header: transcription Chain: A: PDB Molecule: uncharacterized protein q99uf4; PDBTitle: crystal structure of dna-binding response regulator, luxr family, from2 staphylococcus aureus
46	d1yioa2	Alignment	not modelled	99.9	23 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
47	d1zh2a1	Alignment	not modelled	99.9	35 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
48	d1dz3a	Alignment	not modelled	99.9	19 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
49	d1zesal	Alignment	not modelled	99.8	36 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
50	c2qzjC	Alignment	not modelled	99.8	28 PDB header: transcription Chain: C: PDB Molecule: two-component response regulator; PDBTitle: crystal structure of a two-component response regulator from2 clostridium difficile
51	c3hv2B	Alignment	not modelled	99.8	23 PDB header: signaling protein Chain: B: PDB Molecule: response regulator/hd domain protein; PDBTitle: crystal structure of signal receiver domain of hd domain-2 containing protein from pseudomonas fluorescens pf-5
52	d1p6qa	Alignment	not modelled	99.8	28 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
53	d1zgza1	Alignment	not modelled	99.8	28 Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
54	c3jteA	Alignment	not modelled	99.8	29 PDB header: protein binding Chain: A: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of response regulator receiver domain2 protein from clostridium thermocellum

55	c3h1gA	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: A: PDB Molecule: chemotaxis protein chey homolog; PDBTitle: crystal structure of chey mutant t84a of helicobacter pylori
56	c2qv0A	Alignment	not modelled	99.8	23	PDB header: transcription Chain: A: PDB Molecule: protein mrke; PDBTitle: crystal structure of the response regulatory domain of2 protein mrke from klebsiella pneumoniae
57	c3t6kB	Alignment	not modelled	99.8	30	PDB header: signaling protein Chain: B: PDB Molecule: response regulator receiver; PDBTitle: crystal structure of a hypothetical response regulator (caur_3799)2 from chloroflexus aurantiacus j-10-fl at 1.86 a resolution
58	d1a04a2	Alignment	not modelled	99.8	28	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
59	c3cz5B	Alignment	not modelled	99.8	23	PDB header: transcription regulator Chain: B: PDB Molecule: two-component response regulator, luxr family; PDBTitle: crystal structure of two-component response regulator, luxr family,2 from aurantiomonas sp. si85-9a1
60	c3cnbC	Alignment	not modelled	99.8	23	PDB header: dna binding protein Chain: C: PDB Molecule: dna-binding response regulator, merr family; PDBTitle: crystal structure of signal receiver domain of dna binding response2 regulator protein (merr) from colwellia psychrerythraea 34h
61	c2zayA	Alignment	not modelled	99.8	21	PDB header: signaling protein Chain: A: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of response regulator from desulfuromonas2 acetoxidans
62	d1xhfa1	Alignment	not modelled	99.8	34	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
63	c3crnA	Alignment	not modelled	99.8	23	PDB header: signaling protein Chain: A: PDB Molecule: response regulator receiver domain protein, chey-like; PDBTitle: crystal structure of response regulator receiver domain protein (chey-2 like) from methanospirillum hungatei jf-1
64	c1a2oB	Alignment	not modelled	99.8	23	PDB header: bacterial chemotaxis Chain: B: PDB Molecule: cheb methyltransferase; PDBTitle: structural basis for methyltransferase cheb regulation by a2 phosphorylation-activated domain
65	d1heya	Alignment	not modelled	99.8	25	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
66	c3lteH	Alignment	not modelled	99.8	28	PDB header: transcription Chain: H: PDB Molecule: response regulator; PDBTitle: crystal structure of response regulator (signal receiver domain) from2 bermanella marisrubri
67	c3c3mA	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: A: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of the n-terminal domain of response regulator2 receiver protein from methanoculleus marisnigri jr1
68	d2r25b1	Alignment	not modelled	99.8	24	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
69	c3a0rB	Alignment	not modelled	99.8	29	PDB header: transferase Chain: B: PDB Molecule: response regulator; PDBTitle: crystal structure of histidine kinase thka (tm1359) in complex with2 response regulator protein trra (tm1360)
70	c3cg4A	Alignment	not modelled	99.8	27	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: response regulator receiver domain protein (chey-like); PDBTitle: crystal structure of response regulator receiver domain protein (chey-2 like) from methanospirillum hungatei jf-1
71	c3t8yA	Alignment	not modelled	99.8	21	PDB header: hydrolase Chain: A: PDB Molecule: chemotaxis response regulator protein-glutamate PDBTitle: crystal structure of the response regulator domain of thermotoga2 maritima cheb
72	c2hqoA	Alignment	not modelled	99.8	30	PDB header: signaling protein Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: structure of a atypical orphan response regulator protein revealed a2 new phosphorylation-independent regulatory mechanism
73	d1k66a	Alignment	not modelled	99.8	24	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
74	c3f6cB	Alignment	not modelled	99.8	20	PDB header: dna binding protein Chain: B: PDB Molecule: positive transcription regulator evga; PDBTitle: crystal structure of n-terminal domain of positive transcription2 regulator evga from escherichia coli
75	c2qxyB	Alignment	not modelled	99.8	23	PDB header: transcription Chain: B: PDB Molecule: response regulator; PDBTitle: crystal structure of a response regulator from thermotoga2 maritima
76	c2jk1A	Alignment	not modelled	99.8	23	PDB header: dna-binding Chain: A: PDB Molecule: hydrogenase transcriptional regulatory protein hupr1; PDBTitle: crystal structure of the wild-type hupr receiver domain
77	c3khtA	Alignment	not modelled	99.8	23	PDB header: signaling protein Chain: A: PDB Molecule: response regulator; PDBTitle: crystal structure of response regulator from hahella chejuensis
78	c3hebB	Alignment	not modelled	99.8	30	PDB header: transcription regulator Chain: B: PDB Molecule: response regulator receiver domain protein (chey); PDBTitle: crystal structure of response regulator receiver domain from2 rhodospirillum rubrum

79	c2qvgA_	Alignment	not modelled	99.8	23	PDB header: transferase Chain: A: PDB Molecule: two component response regulator; PDBTitle: the crystal structure of a two-component response regulator2 from legionella pneumophila
80	c3kcnA_	Alignment	not modelled	99.8	24	PDB header: lyase Chain: A: PDB Molecule: adenylate cyclase homolog; PDBTitle: the crystal structure of adenylate cyclase from2 rhodopirellula baltica
81	d1p2fa2	Alignment	not modelled	99.8	28	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
82	c3hzhA_	Alignment	not modelled	99.8	19	PDB header: signaling protein Chain: A: PDB Molecule: chemotaxis response regulator (cheY-3); PDBTitle: crystal structure of the cheX-cheY-bef3-mg+2 complex from2 borrelia burgdorferi
83	d1a2oa1	Alignment	not modelled	99.8	21	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
84	d1mb3a_	Alignment	not modelled	99.8	18	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
85	c3klnC_	Alignment	not modelled	99.8	12	PDB header: transcription Chain: C: PDB Molecule: transcriptional regulator, luxr family; PDBTitle: vibrio cholerae vpst
86	c3hdvB_	Alignment	not modelled	99.8	19	PDB header: transcription regulator Chain: B: PDB Molecule: response regulator; PDBTitle: crystal structure of response regulator receiver protein from2 pseudomonas putida
87	c3gl9B_	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: B: PDB Molecule: response regulator; PDBTitle: the structure of a histidine kinase-response regulator2 complex sheds light into two-component signaling and3 reveals a novel cis autophosphorylation mechanism
88	d2b4aa1	Alignment	not modelled	99.8	25	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
89	c3grcD_	Alignment	not modelled	99.8	26	PDB header: transferase Chain: D: PDB Molecule: sensor protein, kinase; PDBTitle: crystal structure of a sensor protein from polaromonas sp.2 js666
90	c3m6mF_	Alignment	not modelled	99.8	30	PDB header: lyase/transferase Chain: F: PDB Molecule: sensory/regulatory protein rpfc; PDBTitle: crystal structure of rpff complexed with rec domain of rpfc
91	d1i3ca_	Alignment	not modelled	99.8	22	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
92	c2rdmB_	Alignment	not modelled	99.8	25	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of response regulator receiver protein from2 sinorhizobium medicae wsm419
93	c3cg0A_	Alignment	not modelled	99.8	20	PDB header: lyase Chain: A: PDB Molecule: response regulator receiver modulated diguanylate cyclase PDBTitle: crystal structure of signal receiver domain of modulated diguanylate2 cyclase from desulfovibrio desulfuricans g20, an example of alternate3 folding
94	c3i42A_	Alignment	not modelled	99.8	23	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: response regulator receiver domain protein (cheY-like)2 from methylobacillus flagellatus
95	c2j48A_	Alignment	not modelled	99.8	20	PDB header: transferase Chain: A: PDB Molecule: two-component sensor kinase; PDBTitle: nmr structure of the pseudo-receiver domain of the cika2 protein.
96	c3c97A_	Alignment	not modelled	99.8	21	PDB header: signaling protein, transferase Chain: A: PDB Molecule: signal transduction histidine kinase; PDBTitle: crystal structure of the response regulator receiver domain2 of a signal transduction histidine kinase from aspergillus3 oryzae
97	c3snkA_	Alignment	not modelled	99.8	20	PDB header: signaling protein Chain: A: PDB Molecule: response regulator chey-like protein; PDBTitle: crystal structure of a response regulator chey-like protein (ml16475)2 from mesorhizobium loti at 2.02 a resolution
98	c3mmnA_	Alignment	not modelled	99.8	27	PDB header: transferase Chain: A: PDB Molecule: histidine kinase homolog; PDBTitle: crystal structure of the receiver domain of the histidine kinase cki12 from arabidopsis thaliana complexed with mg2+
99	d1dcfa_	Alignment	not modelled	99.8	18	Fold: Flavodoxin-like Superfamily: CheY-like Family: Receiver domain of the ethylene receptor
100	c3nhmA_	Alignment	not modelled	99.8	25	PDB header: signaling protein Chain: A: PDB Molecule: response regulator; PDBTitle: crystal structure of a response regulator from myxococcus xanthus
101	c3h5iA_	Alignment	not modelled	99.8	25	PDB header: transcription Chain: A: PDB Molecule: response regulator/sensory box protein/ggdef PDBTitle: crystal structure of the n-terminal domain of a response2 regulator/sensory box/ggdef 3-domain protein from3 carboxydothermus hydrogenoformans
102	c3n0rA_	Alignment	not modelled	99.8	25	PDB header: signaling protein Chain: A: PDB Molecule: response regulator; PDBTitle: structure of the phyr stress response regulator at 1.25 angstrom2 resolution
103	c3eodA_	Alianment	not modelled	99.8	28	PDB header: signaling protein Chain: A: PDB Molecule: protein hnr;

						PDBTitle: crystal structure of n-terminal domain of e. coli rssb
104	c3eqzB_	Alignment	not modelled	99.8	23	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: response regulator; PDBTitle: crystal structure of a response regulator from colwellia2 psychrerythraea
105	c3luaA_	Alignment	not modelled	99.8	19	PDB header: transcription regulator Chain: A: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of a signal receiver domain of two component signal2 transduction (histidine kinase) from clostridium thermocellum
106	d1w25a2	Alignment	not modelled	99.8	22	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
107	c2zxb_	Alignment	not modelled	99.7	27	PDB header: transcription Chain: B: PDB Molecule: transcriptional regulatory protein walr; PDBTitle: crystal structure of yycf dna-binding domain from staphylococcus2 aureus
108	c3ilhA_	Alignment	not modelled	99.7	17	PDB header: transcription regulator Chain: A: PDB Molecule: two component response regulator; PDBTitle: crystal structure of two component response regulator from cytophaga2 hutchinsonii
109	d1gxqa_	Alignment	not modelled	99.7	29	Fold: DNA/RNA-binding 3-helical bundle Superfamily: C-terminal effector domain of the bipartite response regulators Family: PhoB-like
110	c2nt3A_	Alignment	not modelled	99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: response regulator homolog; PDBTitle: receiver domain from myxococcus xanthus social motility protein frzs2 (y102a mutant)
111	c2hqnA_	Alignment	not modelled	99.7	27	PDB header: signaling protein Chain: A: PDB Molecule: putative transcriptional regulator; PDBTitle: structure of a atypical orphan response regulator protein revealed a2 new phosphorylation-independent regulatory mechanism
112	c3lufB_	Alignment	not modelled	99.7	25	PDB header: signaling protein Chain: B: PDB Molecule: two-component system response regulator/ggdef PDBTitle: structure of probable two-component system response2 regulator/ggdef domain protein
113	d1k68a_	Alignment	not modelled	99.7	25	Fold: Flavodoxin-like Superfamily: CheY-like Family: CheY-related
114	c3q9vB_	Alignment	not modelled	99.7	48	PDB header: dna binding protein Chain: B: PDB Molecule: dna-binding response regulator; PDBTitle: crystal structure of rra c-terminal domain(123-221) from deinococcus2 radiodurans
115	c2jzvA_	Alignment	not modelled	99.7	33	PDB header: transcription Chain: A: PDB Molecule: transcriptional regulatory protein pcor; PDBTitle: solution structure of c-terminal effector domain of2 putative two-component-system response regulator involved3 in copper resistance from klebsiella pneumoniae
116	d1ys7a1	Alignment	not modelled	99.7	30	Fold: DNA/RNA-binding 3-helical bundle Superfamily: C-terminal effector domain of the bipartite response regulators Family: PhoB-like
117	d1opca_	Alignment	not modelled	99.7	27	Fold: DNA/RNA-binding 3-helical bundle Superfamily: C-terminal effector domain of the bipartite response regulators Family: PhoB-like
118	c3ktoA_	Alignment	not modelled	99.7	16	PDB header: transcription regulator Chain: A: PDB Molecule: response regulator receiver protein; PDBTitle: crystal structure of response regulator receiver protein2 from pseudoalteromonas atlantica
119	d1qo0d_	Alignment	not modelled	99.7	15	Fold: Flavodoxin-like Superfamily: CheY-like Family: Positive regulator of the amidase operon AmiR
120	c2hvvA_	Alignment	not modelled	99.7	26	PDB header: transcription Chain: A: PDB Molecule: dna-binding response regulator vicr; PDBTitle: crystal structure of an essential response regulator dna binding2 domain, vicrc in enterococcus faecalis, a member of the yycf3 subfamily.