

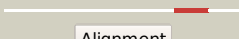

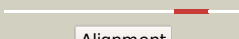











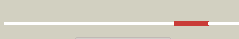
















Phyre2

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Description	A1A535
Date	Wed Jul 10 14:03:03 BST 2013
Unique Job ID	671d1baf5f3b7c66

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2luIA_	 Alignment		99.3	18	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase tec; PDBTitle: solution nmr structure of ph domain of tyrosine-protein kinase tec2 from homo sapiens, northeast structural genomics consortium (nesg)3 target hr3504c
2	c2dtcB_	 Alignment		99.3	23	PDB header: protein binding Chain: B: PDB Molecule: ral guanine nucleotide exchange factor ralgps1a; PDBTitle: crystal structure of ms0666
3	d1baka_	 Alignment		99.2	13	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
4	d2bcja2	 Alignment		99.2	19	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
5	d1btka_	 Alignment		99.2	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
6	d1x1fa1	 Alignment		99.2	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
7	c2q13A_	 Alignment		99.1	22	PDB header: protein transport Chain: A: PDB Molecule: dcc-interacting protein 13 alpha; PDBTitle: crystal structure of bar-ph domain of appl1
8	c2dhkA_	 Alignment		99.1	20	PDB header: immune system Chain: A: PDB Molecule: tbc1 domain family member 2; PDBTitle: solution structure of the ph domain of tbc1 domain family2 member 2 protein from human
9	c2d9xA_	 Alignment		99.1	14	PDB header: lipid transport Chain: A: PDB Molecule: oxysterol binding protein-related protein 11; PDBTitle: solution structure of the ph domain of oxysterol binding2 protein-related protein 11 from human
10	d1u5fa1	 Alignment		99.0	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
11	c3cxB_	 Alignment		99.0	20	PDB header: signaling protein Chain: B: PDB Molecule: pleckstrin homology domain-containing family m PDBTitle: crystal structure of sifa and skip

12	c2d9wA_	Alignment		98.9	23	PDB header: signaling protein Chain: A: PDB Molecule: docking protein 2; PDBTitle: solution structure of the ph domain of docking protein 22 from human
13	d1u5da1	Alignment		98.9	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
14	c3a8qB_	Alignment		98.9	28	PDB header: signaling protein Chain: B: PDB Molecule: t-lymphoma invasion and metastasis-inducing PDBTitle: low-resolution crystal structure of the tiam2 phcxc domain
15	c2kcaA_	Alignment		98.9	18	PDB header: membrane protein Chain: A: PDB Molecule: pleckstrin homology domain-containing family a PDBTitle: solution structure of fapp1 ph domain
16	c2vryA_	Alignment		98.9	23	PDB header: signaling protein Chain: A: PDB Molecule: pleckstrin homology domain-containing family a PDBTitle: solution structure of the ph domain of pleckstrin homology2 domain-containing family a member 6 from human
17	c4h8sA_	Alignment		98.9	24	PDB header: signaling protein Chain: A: PDB Molecule: dccc-interacting protein 13-beta; PDBTitle: crystal structure of human appl2barph domain
18	c3nsuA_	Alignment		98.9	20	PDB header: signaling protein Chain: A: PDB Molecule: phosphatidylinositol 4,5-bisphosphate-binding protein slm1; PDBTitle: a systematic screen for protein-lipid interactions in saccharomyces2 cerevisiae
19	c2d9zA_	Alignment		98.8	22	PDB header: signaling protein Chain: A: PDB Molecule: protein kinase c, nu type; PDBTitle: solution structure of the ph domain of protein kinase c, nu2 type from human
20	c1u5fa_	Alignment		98.8	22	PDB header: signaling protein Chain: A: PDB Molecule: src-associated adaptor protein; PDBTitle: crystal structure of the ph domain of skap-hom with 8 vector-derived2 n-terminal residues
21	d1wgqa_	Alignment	not modelled	98.8	16	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
22	c2dn6A_	Alignment	not modelled	98.8	24	PDB header: signaling protein Chain: A: PDB Molecule: kiaa0640 protein; PDBTitle: solution structure of the ph domain of kiaa0640 protein2 from human
23	d1wi1a_	Alignment	not modelled	98.8	20	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
24	d1v5ua_	Alignment	not modelled	98.8	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
25	c2rloA_	Alignment	not modelled	98.8	19	PDB header: signaling protein Chain: A: PDB Molecule: centaurin-gamma 1; PDBTitle: split ph domain of pi3-kinase enhancer
26	d1v5pa_	Alignment	not modelled	98.8	25	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
27	d1w1ha_	Alignment	not modelled	98.8	20	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
28	d1unqa_	Alignment	not modelled	98.8	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
29	c3tfmA_	Alignment	not modelled	98.7	17	PDB header: motor protein Chain: A: PDB Molecule: myosin x;

						PDBTitle: myosin x ph1n-ph2-ph1c tandem
30	c2da0A_	Alignment	not modelled	98.7	19	PDB header: signaling protein Chain: A: PDB Molecule: 130-kda phosphatidylinositol 4,5-biphosphate- PDBTitle: solution structure of the ph domain of pip2-dependent arf12 gtpase-activating protein from human
31	c2rsgA_	Alignment	not modelled	98.7	19	PDB header: lipid transport Chain: A: PDB Molecule: collagen type iv alpha-3-binding protein; PDBTitle: solution structure of the cert ph domain
32	c2dkpA_	Alignment	not modelled	98.7	21	PDB header: signaling protein Chain: A: PDB Molecule: pleckstrin homology domain-containing family a PDBTitle: solution structure of the ph domain of pleckstrin homology2 domain-containing protein family a member 5 from human
33	c2lkoA_	Alignment	not modelled	98.7	14	PDB header: cell adhesion Chain: A: PDB Molecule: fermitin family homolog 2; PDBTitle: structural basis of phosphoinositide binding to kindlin-2 pleckstrin2 homology domain in regulating integrin activation
34	c2r09A_	Alignment	not modelled	98.7	21	PDB header: signaling protein Chain: A: PDB Molecule: cytohesin-3; PDBTitle: crystal structure of autoinhibited form of grp1 arf gtpase exchange2 factor
35	d1wjma_	Alignment	not modelled	98.7	29	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
36	d1v88a_	Alignment	not modelled	98.7	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
37	d2coaa1	Alignment	not modelled	98.7	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
38	c3a8nA_	Alignment	not modelled	98.7	27	PDB header: signaling protein Chain: A: PDB Molecule: t-lymphoma invasion and metastasis-inducing PDBTitle: crystal structure of the tiam1 phccex domain
39	c2j59O_	Alignment	not modelled	98.7	22	PDB header: hydrolase Chain: O: PDB Molecule: rho-gtpase activating protein 10; PDBTitle: crystal structure of the arf1:arhgap21-arfbd complex
40	d1eaza_	Alignment	not modelled	98.7	25	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
41	d1plsA_	Alignment	not modelled	98.7	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
42	d2elba2	Alignment	not modelled	98.7	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
43	d2i5fa1	Alignment	not modelled	98.6	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
44	c3pp2A_	Alignment	not modelled	98.6	24	PDB header: hydrolase activator Chain: A: PDB Molecule: rho gtpase-activating protein 27; PDBTitle: crystal structure of the pleckstrin homology domain of arhgap27
45	d1u5ea1	Alignment	not modelled	98.6	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
46	d1wg7a_	Alignment	not modelled	98.6	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
47	d1p6sa_	Alignment	not modelled	98.6	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
48	d2j59m1	Alignment	not modelled	98.6	27	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
49	c2ys3A_	Alignment	not modelled	98.6	16	PDB header: signaling protein Chain: A: PDB Molecule: unc-112-related protein 2; PDBTitle: solution structure of the ph domain of kindlin-3 from human
50	d2fjla1	Alignment	not modelled	98.6	15	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
51	c2rovA_	Alignment	not modelled	98.6	19	PDB header: transferase Chain: A: PDB Molecule: rho-associated protein kinase 2; PDBTitle: the split ph domain of rock ii
52	d1upqa_	Alignment	not modelled	98.5	22	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
53	d2coca1	Alignment	not modelled	98.5	16	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
54	d2coda1	Alignment	not modelled	98.5	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
55	c4f7hA_	Alignment	not modelled	98.5	17	PDB header: cell adhesion Chain: A: PDB Molecule: fermitin family homolog 2; PDBTitle: the crystal structure of kindlin-2 pleckstrin homology domain in free2 form

56	c1faoA	Alignment	not modelled	98.5	26	PDB header: signaling protein Chain: A: PDB Molecule: dual adaptor of phosphotyrosine and 3- PDBTitle: structure of the pleckstrin homology domain from 2 dapp1/phish in complex with inositol 1,3,4,5-3 tetrakisphosphate
57	d1faoa	Alignment	not modelled	98.5	26	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
58	c4bbkA	Alignment	not modelled	98.5	17	PDB header: cell adhesion Chain: A: PDB Molecule: fermitin family homolog 1; PDBTitle: structural and functional characterisation of the kindlin-12 pleckstrin homology domain
59	c4gzub	Alignment	not modelled	98.5	23	PDB header: signaling protein Chain: B: PDB Molecule: ferm, rhogef and pleckstrin domain- containing protein 2; PDBTitle: crystal structure of the dh-ph-ph domain of farp2
60	c3fehA	Alignment	not modelled	98.4	19	PDB header: hydrolase activator Chain: A: PDB Molecule: centaurin-alpha-1; PDBTitle: crystal structure of full length centaurin alpha-1
61	c2p0fA	Alignment	not modelled	98.4	20	PDB header: ligand binding protein Chain: A: PDB Molecule: rho gtpase-activating protein 9; PDBTitle: arhgap9 ph domain in complex with ins(1,3,5)p3
62	d1fgya	Alignment	not modelled	98.4	22	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
63	d1v89a	Alignment	not modelled	98.4	25	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
64	c2d9vA	Alignment	not modelled	98.4	14	PDB header: membrane protein Chain: A: PDB Molecule: pleckstrin homology domain-containing protein PDBTitle: solution structure of the ph domain of pleckstrin homology2 domain-containing protein family b member 1 from mouse
65	d1qqga1	Alignment	not modelled	98.4	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Phosphotyrosine-binding domain (PTB)
66	c2dhiA	Alignment	not modelled	98.4	17	PDB header: signaling protein Chain: A: PDB Molecule: pleckstrin homology domain-containing family b PDBTitle: solution structure of the ph domain of evectin-2 from mouse
67	d1u29a1	Alignment	not modelled	98.4	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
68	d1btna	Alignment	not modelled	98.3	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
69	d1x1ga1	Alignment	not modelled	98.3	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
70	c2y7bA	Alignment	not modelled	98.3	19	PDB header: cell cycle Chain: A: PDB Molecule: actin-binding protein anillin; PDBTitle: crystal structure of the ph domain of human actin- binding2 protein anillin anln
71	c1qqgA	Alignment	not modelled	98.2	20	PDB header: signal transduction Chain: A: PDB Molecule: insulin receptor substrate 1; PDBTitle: crystal structure of the ph-ptb targeting region of irs-1
72	d1droa	Alignment	not modelled	98.2	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
73	c3hk0B	Alignment	not modelled	98.2	21	PDB header: signaling protein Chain: B: PDB Molecule: growth factor receptor-bound protein 10; PDBTitle: crystal structure of the ra and ph domains of grb10
74	d2cofa1	Alignment	not modelled	98.2	19	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
75	c4gmvB	Alignment	not modelled	98.2	23	PDB header: cell adhesion Chain: B: PDB Molecule: ras-associated and pleckstrin homology domains-containing PDBTitle: crystal structure of the coiled-coil, ra and ph domains of 2 lamellipodin
76	d1ki1b2	Alignment	not modelled	98.2	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
77	d2dyna	Alignment	not modelled	98.1	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
78	c2k2jA	Alignment	not modelled	98.1	21	PDB header: hydrolase, signaling protein Chain: A: PDB Molecule: 1-phosphatidylinositol-4,5-bisphosphate PDBTitle: nmr solution structure of the split ph domain from 2 phospholipase c gamma 2
79	c3o96A	Alignment	not modelled	98.1	23	PDB header: transferase Chain: A: PDB Molecule: rac-alpha serine/threonine-protein kinase; PDBTitle: crystal structure of human akt1 with an allosteric inhibitor
80	d1xx0a1	Alignment	not modelled	98.1	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
81	c3tcaA	Alignment	not modelled	98.0	19	PDB header: signaling protein Chain: A: PDB Molecule: amyloid beta a4 precursor protein-binding family b member PDBTitle: crystal structure of the ras-associating and pleckstrin-

					homology2 domains of riam PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal structure of the dh-ph-ph domain of farp1	
82	c4h6yA_	Alignment	not modelled	97.9	27	
83	c1w63A_	Alignment		97.8	11	PDB header: endocytosis Chain: A: PDB Molecule: adapter-related protein complex 1 gamma 1 PDBTitle: ap1 clathrin adaptor core
84	d1gw5b_	Alignment		97.6	14	Fold: alpha-alpha superhelix Superfamily: ARM repeat Family: Clathrin adaptor core protein
85	c3ml4D_	Alignment	not modelled	97.5	19	PDB header: signaling protein Chain: D: PDB Molecule: protein dok-7; PDBTitle: crystal structure of a complex between dok7 ph-ptb and the musk2 juxtamembrane region
86	d2dfka2	Alignment	not modelled	97.4	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
87	c2db0B_	Alignment		97.4	12	PDB header: protein binding Chain: B: PDB Molecule: 253aa long hypothetical protein; PDBTitle: crystal structure of ph0542
88	d1rj2a2	Alignment	not modelled	97.3	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
89	d1maia_	Alignment	not modelled	97.3	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
90	d1v61a_	Alignment	not modelled	97.2	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
91	c2jkrL_	Alignment		97.0	12	PDB header: endocytosis Chain: L: PDB Molecule: ap-2 complex subunit alpha-2; PDBTitle: ap2 clathrin adaptor core with dileucine peptide rm(2 phosphos)qikrllse
92	c2lg1A_	Alignment		97.0	16	PDB header: metal binding protein Chain: A: PDB Molecule: a-kinase anchor protein 13; PDBTitle: solution structure of the human akap13 ph domain and stabilizing dh2 helix
93	d1nty2	Alignment	not modelled	96.5	16	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
94	d1zc3b1	Alignment	not modelled	96.4	16	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
95	d1xcga2	Alignment	not modelled	96.4	23	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
96	c3odwB_	Alignment	not modelled	96.3	13	PDB header: signaling protein Chain: B: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the linker-dh/ph domains of p115-rhogef
97	c3mpxA_	Alignment	not modelled	96.3	21	PDB header: lipid binding protein Chain: A: PDB Molecule: fyve, rhogef and ph domain-containing protein 5; PDBTitle: crystal structure of the dh and ph-1 domains of human fgd5
98	c3bjia_	Alignment	not modelled	96.2	19	PDB header: signaling protein Chain: A: PDB Molecule: proto-oncogene vav; PDBTitle: structural basis of promiscuous guanine nucleotide exchange2 by the t-cell essential vav1
99	c3ky9B_	Alignment	not modelled	96.1	23	PDB header: apoptosis Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
100	c2vszA_	Alignment	not modelled	96.0	23	PDB header: apoptosis Chain: A: PDB Molecule: engulfment and cell motility protein 1; PDBTitle: crystal structure of the elmo1 ph domain
101	c3qwmA_	Alignment	not modelled	95.9	15	PDB header: signaling protein Chain: A: PDB Molecule: iq motif and sec7 domain-containing protein 1; PDBTitle: crystal structure of gep100, the plextrin homology domain of iq motif2 and sec7 domain-containing protein 1 isoform a
102	c2vrwB_	Alignment	not modelled	95.9	23	PDB header: signaling protein Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: critical structural role for the ph and c1 domains of the2

						vav1 exchange factor
103	d1dbha2	Alignment	not modelled	95.6	27	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
104	d1kz7a2	Alignment	not modelled	95.5	20	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
105	c1x86A_	Alignment	not modelled	95.1	18	PDB header: signaling protein/membrane protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 12; PDBTitle: crystal structure of the dh/ph domains of leukemia-associated rhogef2 in complex with rhoa
106	c2dfkA_	Alignment	not modelled	95.0	19	PDB header: cell cycle Chain: A; PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
107	c3w3zA_	Alignment	not modelled	95.0	13	PDB header: protein transport/nuclear protein Chain: A; PDB Molecule: importin subunit beta-3; PDBTitle: crystal structure of kap121p bound to rangtp
108	c3odxA_	Alignment	not modelled	95.0	14	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of an n-terminally truncated linker-dh/ph domains of p115-rhogef
109	c3a98B_	Alignment	not modelled	94.7	23	PDB header: signaling protein Chain: B; PDB Molecule: engulfment and cell motility protein 1; PDBTitle: crystal structure of the complex of the interacting regions of dock22 and elmo1
110	c1ki1B_	Alignment	not modelled	94.3	19	PDB header: signaling protein Chain: B; PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
111	c1xcgA_	Alignment	not modelled	94.2	19	PDB header: signaling protein activator/signaling pr Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of human rhoa in complex with dh/ph2 fragment of pdzrhogef
112	d1txda2	Alignment	not modelled	93.5	22	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
113	c3v6aA_	Alignment	not modelled	92.0	20	PDB header: apoptosis inhibitor Chain: A; PDB Molecule: apoptosis inhibitor 5; PDBTitle: helical repeat structure of apoptosis inhibitor 5 reveals protein-2 protein interaction modules
114	c3ohmB_	Alignment	not modelled	91.5	15	PDB header: signaling protein / hydrolase Chain: B; PDB Molecule: 1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase PDBTitle: crystal structure of activated g alpha q bound to its effector2 phospholipase c beta 3
115	d1omwa2	Alignment	not modelled	91.3	20	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
116	c2nppE_	Alignment	not modelled	89.6	18	PDB header: hydrolase/hydrolase inhibitor Chain: E; PDB Molecule: serine/threonine-protein phosphatase 2a 56 kda regulatory PDBTitle: structure of the protein phosphatase 2a holoenzyme
117	d2nppb1	Alignment	not modelled	89.6	18	Fold: alpha-alpha superhelix Superfamily: ARM repeat Family: B56-like
118	d2bpta1	Alignment	not modelled	89.5	7	Fold: alpha-alpha superhelix Superfamily: ARM repeat Family: Armadillo repeat
119	c2rgnB_	Alignment	not modelled	88.8	16	PDB header: signaling protein complex Chain: B; PDB Molecule: rhoa/rac/cdc42 exchange factor; PDBTitle: crystal structure of p63rhogef complex with galpha-q and rhoa
120	c3qr0A_	Alignment	not modelled	88.7	16	PDB header: hydrolase Chain: A; PDB Molecule: phospholipase c-beta (plc-beta); PDBTitle: crystal structure of s. officinalis plc21