

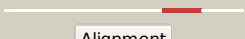

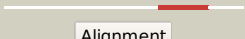

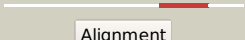

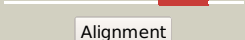

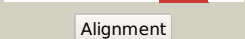

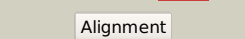

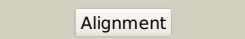

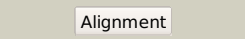

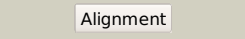

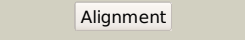



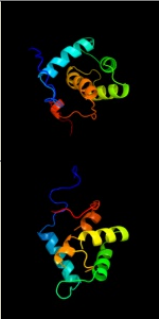
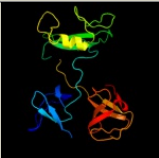
Phyre2

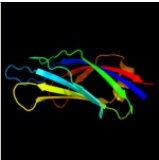
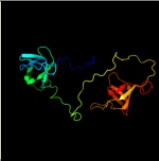
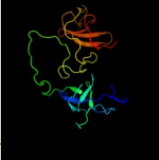
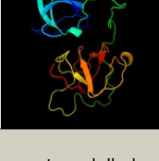
Email	l.a.kelley@imperial.ac.uk
Description	Q9Z0R6
Date	Wed Jul 10 14:14:37 BST 2013
Unique Job ID	b50ef405e1b95f7f

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dx1A_	 Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
2	c3jv3A_	 Alignment		100.0	76	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
3	c3odoA_	 Alignment		100.0	26	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the dh/ph domains of p115-rhogef
4	c3ky9B_	 Alignment		100.0	22	PDB header: apoptosis Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
5	c1xcgA_	 Alignment		100.0	26	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
6	c1ki1B_	 Alignment		100.0	77	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
7	c1x86A_	 Alignment		100.0	26	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
8	c3ksyA_	 Alignment		100.0	16	PDB header: signaling protein Chain: A: PDB Molecule: son of sevenless homolog 1; PDBTitle: crystal structure of the histone domain, dh-ph unit, and2 catalytic unit of the ras activator son of sevenless (sos)
9	c3t06E_	 Alignment		100.0	26	PDB header: signaling protein Chain: E: PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of the dh/ph fragment of pdzrhogef with n-terminal2 regulatory elements in complex with human rhoa
10	c1xdvB_	 Alignment		100.0	15	PDB header: signaling protein Chain: B: PDB Molecule: son of sevenless protein homolog 1; PDBTitle: experimentally phased structure of human the son of2 sevenless protein at 4.1 ang.
11	c2dfkA_	 Alignment		100.0	20	PDB header: cell cycle Chain: A: PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex

12	c2vrwB_	Alignment		100.0	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
13	c1foeA_	Alignment		100.0	22	PDB header: signaling protein, immune system/signali Chain: A: PDB Molecule: t-lymphoma invasion and metastasis inducing PDBTitle: crystal structure of rac1 in complex with the guanine2 nucleotide exchange region of tiam1
14	c4gouA_	Alignment		100.0	13	PDB header: signaling protein Chain: A: PDB Molecule: ehrgs-rhogef; PDBTitle: crystal structure of an rgs-rhogef from entamoeba histolytica
15	c2z0gA_	Alignment		100.0	27	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
16	c3odxA_	Alignment		100.0	27	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 of2 p115-rhogef
17	c3bjjA_	Alignment		100.0	23	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
18	c3odwB_	Alignment		100.0	27	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
19	c4gzuB_	Alignment		100.0	24	PDB header: signaling protein Chain: B: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal strucure of the dh-ph-ph domain of farp2
20	c1dbhA_	Alignment		100.0	15	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hso1
21	c4h6yA_	Alignment	not modelled	100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal strucure of the dh-ph-ph domain of farp1
22	c2rgnB_	Alignment	not modelled	100.0	20	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
23	c3mpxA_	Alignment	not modelled	100.0	20	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
24	c1lb1C_	Alignment	not modelled	100.0	20	PDB header: signaling protein Chain: C: PDB Molecule: guanine nucleotide exchange factor dbs; PDBTitle: crystal structure of the dbl and pleckstrin homology2 domains of dbs in complex with rhoa
25	c1ntyA_	Alignment	not modelled	100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: triple functional domain protein; PDBTitle: crystal structure of the first dh/ph domain of trio to 1.7 a
26	d1xcga1	Alignment	not modelled	100.0	28	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
27	d1txda1	Alignment	not modelled	100.0	28	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
28	d1ki1b1	Alignment	not modelled	100.0	77	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)

29	c4gyvA	Alignment	not modelled	100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal structure of the dh domain of farp2
30	d1dbha1	Alignment	not modelled	100.0	16	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
31	d2dfka1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
32	c3eo2A	Alignment	not modelled	100.0	30	PDB header: splicing Chain: A: PDB Molecule: neuroepithelial cell-transforming gene 1 protein; PDBTitle: crystal structure of the rhogef domain of human neuroepithelial cell-2 transforming gene 1 protein
33	d1foea1	Alignment	not modelled	100.0	26	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
34	c3gf9A	Alignment	not modelled	100.0	94	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
35	d1f5xa	Alignment	not modelled	99.9	27	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
36	d1by1a	Alignment	not modelled	99.9	20	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
37	d1kz7a1	Alignment	not modelled	99.9	23	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
38	d1ntya1	Alignment	not modelled	99.9	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
39	c2khnA	Alignment		99.9	69	PDB header: signaling protein Chain: A: PDB Molecule: intersectin-1; PDBTitle: nmr solution structure of the eh 1 domain from human2 intersectin-1 protein. northeast structural genomics3 consortium target hr3646e.
40	c3fiaA	Alignment		99.9	68	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: crystal structure of the eh 1 domain from human intersectin-2 1 protein. northeast structural genomics consortium target3 hr3646e.
41	c2jq6A	Alignment	not modelled	99.9	38	PDB header: metal binding protein Chain: A: PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of eh-domain of ehd1
42	c2kfhA	Alignment	not modelled	99.8	39	PDB header: protein binding Chain: A: PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of the c-terminal domain of ehd1 with fnyestgpftak
43	d2jxca1	Alignment	not modelled	99.8	38	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
44	c2kgrA	Alignment	not modelled	99.8	38	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: solution structure of protein itsn1 from homo sapiens.2 northeast structural genomics consortium target hr5524a
45	d1iq3a	Alignment	not modelled	99.7	35	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
46	d1fi6a	Alignment	not modelled	99.7	39	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
47	d1c07a	Alignment	not modelled	99.7	37	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
48	c2qptA	Alignment	not modelled	99.7	38	PDB header: endocytosis Chain: A: PDB Molecule: eh domain-containing protein-2; PDBTitle: crystal structure of an ehd atpase involved in membrane remodelling
49	c1gria	Alignment		99.7	23	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
50	c2lg1A	Alignment		not modelled	99.7	19

51	c3jzyA_	Alignment		99.6	96	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 c2 domain
52	d1qjta_	Alignment	not modelled	99.6	26	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
53	d1ki1b2	Alignment	not modelled	99.6	73	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
54	c2lqnA_	Alignment		99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
55	c2eyzA_	Alignment		99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: zv-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
56	c2l3sA_	Alignment		99.5	21	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
57	c2fk9A_	Alignment	not modelled	99.5	25	PDB header: transferase Chain: A: PDB Molecule: protein kinase c, eta type; PDBTitle: human protein kinase c, eta
58	c1ng2A_	Alignment	not modelled	99.4	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
59	c1ov3A_	Alignment	not modelled	99.4	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
60	d2nq3a1	Alignment	not modelled	99.4	23	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
61	c2nq3A_	Alignment	not modelled	99.4	23	PDB header: ligase Chain: A: PDB Molecule: itchy homolog e3 ubiquitin protein ligase; PDBTitle: crystal structure of the c2 domain of human itchy homolog2 e3 ubiquitin protein ligase
62	c1mv3A_	Alignment	not modelled	99.4	18	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
63	d1xcga2	Alignment	not modelled	99.3	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
64	d1gmia_	Alignment	not modelled	99.3	23	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
65	d2dfka2	Alignment	not modelled	99.3	20	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
66	d1zc3b1	Alignment	not modelled	99.2	18	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
67	c3nsjA_	Alignment	not modelled	99.2	23	PDB header: immune system Chain: A: PDB Molecule: perforin-1; PDBTitle: the x-ray crystal structure of lymphocyte perforin
68	d1udla_	Alignment	not modelled	99.2	85	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c3kwtA_	Alignment	not modelled	99.2	25	PDB header: metal binding protein Chain: A: PDB Molecule: munc13-1; PDBTitle: munc13-1 c2b-domain, calcium-free
70	d1bdya_	Alignment	not modelled	99.2	18	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
71	d1dbha2	Alignment	not modelled	99.2	16	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
72	c2dmhA_	Alignment	not modelled	99.2	22	PDB header: lipid binding protein Chain: A: PDB Molecule: myoferlin; PDBTitle: solution structure of the first c2 domain of human myoferlin
73	d2ep6a1	Alignment	not modelled	99.2	36	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)

74	d1wfja_	Alignment	not modelled	99.2	21	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
75	d1txda2	Alignment	not modelled	99.1	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
76	c3l9bA_	Alignment	not modelled	99.1	17	PDB header: membrane protein Chain: A: PDB Molecule: otoferlin; PDBTitle: crystal structure of rat otoferlin c2a
77	d2cjsa1	Alignment	not modelled	99.1	22	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
78	c3m7fB_	Alignment	not modelled	99.1	31	PDB header: signaling protein/ligase Chain: B: PDB Molecule: e3 ubiquitin-protein ligase nedd4; PDBTitle: crystal structure of the nedd4 c2/grb10 sh2 complex
79	d1rlwa_	Alignment	not modelled	99.1	20	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
80	d1ng2a2	Alignment	not modelled	99.1	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c2jqzA_	Alignment	not modelled	99.0	25	PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase smurf2; PDBTitle: solution structure of the c2 domain of human smurf2
82	c3qwmA_	Alignment	not modelled	99.0	18	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
83	c1dqvA_	Alignment	not modelled	99.0	24	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: synaptotagmin iii; PDBTitle: crystal structure of synaptotagmin iii c2a/c2b
84	c2enjA_	Alignment	not modelled	99.0	18	PDB header: transferase Chain: A: PDB Molecule: protein kinase c theta type; PDBTitle: solution structure of the c2 domain from human protein2 kinase c theta
85	d1qasa2	Alignment	not modelled	99.0	25	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
86	d1dsya_	Alignment	not modelled	99.0	30	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
87	c2rqrA_	Alignment	not modelled	99.0	13	PDB header: protein binding Chain: A: PDB Molecule: engulfment and cell motility protein 1, linker, dedicator PDBTitle: the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
88	c3nmzD_	Alignment	not modelled	99.0	29	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asef
89	c1x2gA_	Alignment	not modelled	99.0	30	PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adapter molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
90	c2ysqA_	Alignment	not modelled	98.9	20	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
91	c2cubA_	Alignment	not modelled	98.9	33	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
92	c2jxbA_	Alignment	not modelled	98.9	25	PDB header: signaling protein complex Chain: A: PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain, PDBTitle: structure of cd3epsilon-nck2 first sh3 domain complex
93	c2eyyA_	Alignment	not modelled	98.9	21	PDB header: signaling protein Chain: A: PDB Molecule: sv-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
94	c2ed0A_	Alignment	not modelled	98.9	39	PDB header: signaling protein Chain: A: PDB Molecule: abl interactor 2; PDBTitle: solution structure of the sh3 domain of abl interactor 22 (abelson interactor 2)
95	c2yuoA_	Alignment	not modelled	98.9	25	PDB header: signaling protein Chain: A: PDB Molecule: run and tbc1 domain containing 3; PDBTitle: solution structure of the sh3 domain of mouse run and tbc12 domain containing 3
96	d1ov3a2	Alignment	not modelled	98.9	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2ekhA_	Alignment	not modelled	98.9	20	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structures of the sh3 domain of human kiaa0418
98	c2yunA_	Alignment	not modelled	98.9	27	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
99	c2r83B_	Alignment	not modelled	98.9	26	PDB header: endocytosis, exocytosis Chain: B: PDB Molecule: synaptotagmin-1; PDBTitle: crystal structure analysis of human synaptotagmin 1 c2a-c2b
100	c2dlnA_	Alignment	not modelled	98.9	28	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein;

100	c2qipA	Alignment	not modelled	98.9	28	PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein PDB header: cell adhesion Chain: A; PDB Molecule: proline-serine-threonine phosphatase-interacting
101	c2dilA	Alignment	not modelled	98.9	29	PDBTitle: solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1 PDB header: protein binding Chain: A; PDB Molecule: rho gtpase-activating protein 4;
102	c2epdA	Alignment	not modelled	98.9	26	PDBTitle: solution structure of sh3 domain in rho-gtpase-activating2 protein 4 PDB header: signaling protein Chain: A; PDB Molecule: signal transducing adapter molecule 1;
103	c2l0aA	Alignment	not modelled	98.9	44	PDBTitle: solution nmr structure of signal transducing adapter molecule 1 stam-12 from homo sapiens, northeast structural genomics consortium target3 hr4479e PDB header: ligase Chain: A; PDB Molecule: e3 ubiquitin-protein ligase nedd4-like protein;
104	c2nsgA	Alignment	not modelled	98.9	28	PDBTitle: crystal structure of the c2 domain of the human e3 ubiquitin-protein2 ligase nedd4-like protein PDB header: endocytosis/exocytosis Chain: A; PDB Molecule: rim binding protein 2;
105	c2csqA	Alignment	not modelled	98.9	30	PDBTitle: solution structure of the second sh3 domain of human rim-2 binding protein 2 Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
107	c2d8kA	Alignment	not modelled	98.9	28	PDB header: endocytosis/exocytosis Chain: A; PDB Molecule: synaptotagmin vii; PDBTitle: solution structure of the first c2 domain of synaptotagmin2 vii PDB header: cell adhesion Chain: A; PDB Molecule: vinexin;
108	c2nwmA	Alignment	not modelled	98.9	29	PDBTitle: solution structure of the first sh3 domain of human vinexin2 and its interaction with the peptides from vinculin PDB header: signaling protein Chain: A; PDB Molecule: slit-robo rho gtpase-activating protein 2;
109	c2dl8A	Alignment	not modelled	98.9	31	PDBTitle: solution structure of the sh3 domain of human slit-robo rho2 gtpase-activating protein 2 PDB header: structural genomics, structural protein Chain: A; PDB Molecule: sh3 multiple domains 1;
110	c2dnuA	Alignment	not modelled	98.9	28	PDBTitle: solution structure of rsgi ruh-061, a sh3 domain from human PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2;
111	c3qwyA	Alignment	not modelled	98.8	26	PDBTitle: ced-2 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	d1k4us	Alignment	not modelled	98.8	33	PDB header: cell adhesion Chain: A; PDB Molecule: sam and sh3 domain-containing protein 1;
113	c2ebpA	Alignment	not modelled	98.8	22	PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1 PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2;
114	c3qwxX	Alignment	not modelled	98.8	26	PDBTitle: ced-2 1-174 PDB header: signaling protein Chain: A; PDB Molecule: bud emergence protein 1;
115	c2kymA	Alignment	not modelled	98.8	27	PDBTitle: solution structure of the bem1p sh3-ci domain from l.elongisporus in2 complex with ste20p peptide PDB header: signaling protein Chain: A; PDB Molecule: cortactin isoform a;
116	c1x69A	Alignment	not modelled	98.8	32	PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin PDB header: signaling protein Chain: A; PDB Molecule: 130 kda phosphatidylinositol 4,5-biphosphate-
117	c2ed1A	Alignment	not modelled	98.8	35	PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein PDB header: signaling protein Chain: D; PDB Molecule: dynamin-binding protein;
118	c4glmD	Alignment	not modelled	98.8	30	PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens] PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein;
119	c2dbkA	Alignment	not modelled	98.8	20	PDBTitle: solution structures of the sh3 domain of human crk-like2 protein PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 7;
120	c2df6A	Alignment	not modelled	98.8	45	PDBTitle: crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2