



























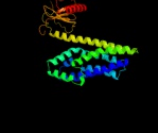
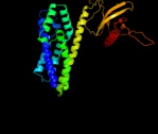



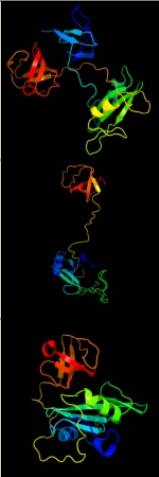

Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P27870
Date	Tue Jul 30 13:02:10 BST 2013
Unique Job ID	08c87f45799f37fa

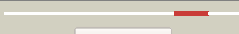


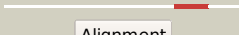
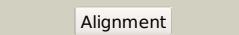
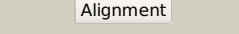
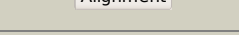





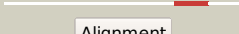
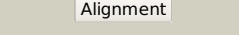
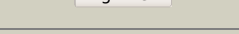




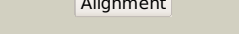
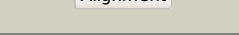



Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3ky9B_	 Alignment		100.0	95	PDB header: apoptosis Chain: B: PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
2	c2vrwB_	 Alignment		100.0	99	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
3	c3bjjA_	 Alignment		100.0	93	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
4	c3ksyA_	 Alignment		100.0	18	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)
5	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
6	c1xdvB_	 Alignment		100.0	19	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.
7	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
8	c2dfkA_	 Alignment		100.0	24	PDB header: cell cycle Chain: A: PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex
9	c1x86A_	 Alignment		100.0	23	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
10	c4gzvB_	 Alignment		100.0	20	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
11	c1xcgA_	 Alignment		100.0	23	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

12	c3t06E_	Alignment		100.0	21	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
13	c4gouA_	Alignment		100.0	15	PDB header: signaling protein Chain: A: PDB Molecule: ehrgs-rhogef; PDBTitle: crystal structure of an rgs-rhogef from entamoeba histolytica
14	c1foeA_	Alignment		100.0	23	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
15	c4h6yA_	Alignment		100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal strcure of the dh-ph-ph domain of farp1
16	c2rgnB_	Alignment		100.0	22	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
17	c3mpxA_	Alignment		100.0	17	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
18	c3odwB_	Alignment		100.0	25	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	c3odxA_	Alignment		100.0	24	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
20	c1ki1B_	Alignment		100.0	21	PDB header: signaling protein Chain: B: PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
21	c1dbhA_	Alignment	not modelled	100.0	20	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hsos1
22	c1lb1C_	Alignment	not modelled	100.0	21	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
23	c2z0qA_	Alignment	not modelled	100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
24	c1ntyA_	Alignment	not modelled	100.0	20	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
25	c3jv3A_	Alignment	not modelled	100.0	21	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-11
26	d1f5xa_	Alignment	not modelled	100.0	99	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
27	d1xcga1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
28	d1dbha1	Alignment	not modelled	100.0	18	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
29	d1txda1	Alignment	not modelled	100.0	22	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)

30	c4gyvA	Alignment	not modelled	100.0	20	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
31	d1ki1b1	Alignment	not modelled	100.0	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
32	d2dfka1	Alignment	not modelled	100.0	27	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
33	c2d86A	Alignment	not modelled	100.0	72	PDB header: signaling protein, protein binding Chain: A; PDB Molecule: vav-3 protein; PDBTitle: solution structure of the ch domain from human vav-3 protein
34	d1foea1	Alignment	not modelled	100.0	29	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
35	c3eo2A	Alignment	not modelled	100.0	23	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
36	d1kz7a1	Alignment	not modelled	100.0	21	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
37	c3gf9A	Alignment	not modelled	100.0	28	PDB header: endocytosis Chain: A; PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
38	d1ntyA1	Alignment	not modelled	100.0	20	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
39	d1by1a	Alignment	not modelled	100.0	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
40	c1gria	Alignment		99.9	24	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
41	c2eyyA	Alignment		99.9	23	PDB header: signaling protein Chain: A; PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
42	c2lqnA	Alignment		99.9	25	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
43	c2l3gA	Alignment	not modelled	99.9	38	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: solution nmr structure of ch domain of rho guanine nucleotide exchange2 factor 7 from homo sapiens, northeast structural genomics consortium3 target hr4495e
44	d1ujoa	Alignment	not modelled	99.9	24	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
45	c1wymA	Alignment	not modelled	99.9	21	PDB header: structural protein Chain: A; PDB Molecule: transgelin-2; PDBTitle: solution structure of the ch domain of human transgelin-2
46	c2dviA	Alignment		99.9	28	PDB header: signaling protein Chain: A; PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
47	c3qwxX	Alignment		99.9	25	PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
48	c2eyzA	Alignment		99.9	28	PDB header: signaling protein Chain: A; PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
49	c3qwyA	Alignment	not modelled	99.9	23	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
						PDB header: structural protein

50	c1wyrA	Alignment	not modelled	99.9	31	Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 6; PDBTitle: solution structure of the ch domain of human rho guanine2 nucleotide exchange factor 6
51	c1mv3A	Alignment	not modelled	99.8	15	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
52	c1wypA	Alignment	not modelled	99.8	25	PDB header: structural protein Chain: A: PDB Molecule: calponin 1; PDBTitle: solution structure of the ch domain of human calponin 1
53	c2rr8A	Alignment	not modelled	99.8	22	PDB header: protein binding Chain: A: PDB Molecule: iqgap1 protein; PDBTitle: solution structure of calponin homology domain of iqgap1
54	c1wynA	Alignment	not modelled	99.8	25	PDB header: structural protein Chain: A: PDB Molecule: calponin-2; PDBTitle: solution structure of the ch domain of human calponin-2
55	c3i6xC	Alignment	not modelled	99.8	23	PDB header: calmodulin-binding, membrane protein Chain: C: PDB Molecule: ras gtpase-activating-like protein iqgap1; PDBTitle: crystal structure of the calponin homology domain of iqgap1
56	c2l3sA	Alignment	not modelled	99.8	14	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
57	d1p2xa	Alignment	not modelled	99.8	26	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
58	c1p5sA	Alignment	not modelled	99.8	26	PDB header: cytokine Chain: A: PDB Molecule: ras gtpase-activating-like protein rng2; PDBTitle: structure and function of the calponin-homology domain of2 an iqgap protein from schizosaccharomyces pombe
59	c2lg1A	Alignment	not modelled	99.8	12	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
60	d1h67a	Alignment	not modelled	99.8	30	Fold: CH domain-like Superfamily: Calponin-homology domain, CH-domain Family: Calponin-homology domain, CH-domain
61	c1ov3A	Alignment	not modelled	99.7	20	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
62	c1ng2A	Alignment	not modelled	99.7	35	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
63	d1ng2a2	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c3nmzD	Alignment	not modelled	99.7	23	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
65	c2rqrA	Alignment	not modelled	99.7	17	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
66	d1kz7a2	Alignment	not modelled	99.7	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
67	c2kbtA	Alignment	not modelled	99.7	94	PDB header: signaling protein Chain: A: PDB Molecule: chimera of proto-oncogene vav, linker, PDBTitle: attachment of an nmr-invisible solubility enhancement tag2 (inset) using a sortase-mediated protein ligation method
68	d1ov3a2	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	d2dfka2	Alignment	not modelled	99.7	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
70	c1zlmA	Alignment	not modelled	99.7	41	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
71	c2jxbA	Alignment	not modelled	99.7	24	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
72	c2ed0A	Alignment	not modelled	99.7	35	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)
73	c2csgA	Alignment	not modelled	99.7	32	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the second sh3 domain of human rim-2 binding protein 2
74	c1x2qA	Alignment	not modelled	99.7	34	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
75	c2nwmA	Alignment	not modelled	99.7	41	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the first sh3 domain of human vinexin2 and its interaction with the peptides from vinculin
76	d1uj0a	Alignment	not modelled	99.7	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

77	c2dlpA	 Alignment	not modelled	99.7	31	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
78	c1x6gA	 Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: megakaryocyte-associated tyrosine-protein kinase; PDBTitle: solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
79	c2e5kA	 Alignment	not modelled	99.7	24	PDB header: protein binding Chain: A: PDB Molecule: suppressor of t-cell receptor signaling 1; PDBTitle: solution structure of sh3 domain in suppressor of t-cell2 receptor signaling 1
80	c2xmfA	 Alignment	not modelled	99.7	39	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
81	c2cubA	 Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
82	c1zx6A	 Alignment	not modelled	99.7	36	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
83	d1l1ja	 Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2df6A	 Alignment	not modelled	99.6	40	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2
85	d1gcqa	 Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c4glmD	 Alignment	not modelled	99.6	36	PDB header: signaling protein Chain: D: PDB Molecule: dynamamin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
87	c2l0aA	 Alignment	not modelled	99.6	34	PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adapter molecule 1; PDBTitle: solution nmr structure of signal transducing adapter molecule 1 stam-12 from homo sapiens, northeast structural genomics consortium target3 hr4479e
88	c1x43A	 Alignment	not modelled	99.6	24	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: sh3 domain grb2-like protein b1; PDBTitle: solution structure of the sh3 domain of endophilin b12 (sh3g1b1)
89	c2yt6A	 Alignment	not modelled	99.6	31	PDB header: cell cycle Chain: A: PDB Molecule: adult male urinary bladder cdna, riken full- PDBTitle: solution structure of the sh3_1 domain of yamaguchi sarcoma2 viral (v-yes) oncogene homolog 1
90	d1utia	 Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2ysqA	 Alignment	not modelled	99.6	33	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
92	d1rj2a2	 Alignment	not modelled	99.6	25	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
93	c2ekhA	 Alignment	not modelled	99.6	25	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
94	d1k4us	 Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	d1sema	 Alignment	not modelled	99.6	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2kymA	 Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from l.elongisporus in2 complex with ste20p peptide
97	d1oeba	 Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	d1uuea	 Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	c2egeA	 Alignment	not modelled	99.6	23	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein kiaa1666; PDBTitle: solution structure of the third sh3 domain from human2 kiaa1666 protein
100	c2epdA	 Alignment	not modelled	99.6	30	PDB header: protein binding Chain: A: PDB Molecule: rho gtpase-activating protein 4; PDBTitle: solution structure of sh3 domain in rho-gtpase-activating2 protein 4
101	c2yuoA	 Alignment	not modelled	99.6	41	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
102	c2yupA	Alignment	not modelled	99.6	34	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin

103	c2ebpA	Alignment	not modelled	99.6	38	PDB header: cell adhesion Chain: A: PDB Molecule: sam and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
104	d1wlpb2	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	c2d1xD	Alignment	not modelled	99.6	44	PDB header: cell invasion Chain: D: PDB Molecule: cortactin isoform a; PDBTitle: the crystal structure of the cortactin-sh3 domain and amap1-2 peptide complex
106	c2cudA	Alignment	not modelled	99.6	17	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
107	d1loota	Alignment	not modelled	99.6	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2a28D	Alignment	not modelled	99.6	35	PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein; PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
109	c2dilA	Alignment	not modelled	99.6	33	PDB header: cell adhesion Chain: A: PDB Molecule: proline-serine-threonine phosphatase-interacting PDBTitle: solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1
110	c2yunA	Alignment	not modelled	99.6	47	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
111	c1wyxA	Alignment	not modelled	99.6	41	PDB header: cell adhesion Chain: A: PDB Molecule: crk-associated substrate; PDBTitle: the crystal structure of the p130cas sh3 domain at 1.1 a2 resolution
112	d1gria2	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2dmoA	Alignment	not modelled	99.6	27	PDB header: signaling protein Chain: A: PDB Molecule: neutrophil cytosol factor 2; PDBTitle: refined solution structure of the 1st sh3 domain from human2 neutrophil cytosol factor 2 (ncf-2)
114	d1u06a1	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2dnuA	Alignment	not modelled	99.6	18	PDB header: structural genomics, structural protein Chain: A: PDB Molecule: sh3 multiple domains 1; PDBTitle: solution structure of rsgi ruh-061, a sh3 domain from human
116	c2fpeB	Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: B: PDB Molecule: c-jun-amino-terminal kinase interacting protein PDBTitle: conserved dimerization of the ib1 src-homology 3 domain
117	d1pwta	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c1zukA	Alignment	not modelled	99.6	19	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mt11; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
119	c1negA	Alignment	not modelled	99.6	33	PDB header: structural protein Chain: A: PDB Molecule: spectrin alpha chain, brain; PDBTitle: crystal structure analysis of n-and c-terminal labeled sh3-2 domain of alpha-chicken spectrin
120	d1nega	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain