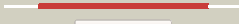


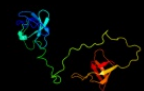
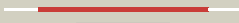














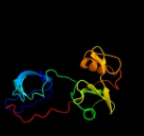



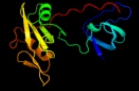

















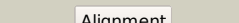

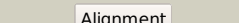

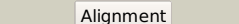
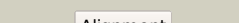
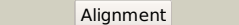



Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	O89100
Date	Wed Jul 10 14:33:29 BST 2013
Unique Job ID	51ca4f3ac066ec0d

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	58	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2lqnA_	 Alignment		100.0	26	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c2l3sA_	 Alignment		100.0	26	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
4	c2eyzA_	 Alignment		100.0	34	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1ng2A_	 Alignment		100.0	33	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
6	c1ov3A_	 Alignment		100.0	34	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
7	c2eyvA_	 Alignment		99.9	27	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
8	c1mv3A_	 Alignment		99.9	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
9	c1oplA_	 Alignment		99.9	31	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase; PDBTitle: structural basis for the auto-inhibition of c-abl tyrosine2 kinase
10	c2fo0A_	 Alignment		99.9	31	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase abl1 (1b isoform); PDBTitle: organization of the sh3-sh2 unit in active and inactive forms of the2 c-abl tyrosine kinase
11	c1k9aB_	 Alignment		99.9	33	PDB header: transferase Chain: B: PDB Molecule: carboxyl-terminal src kinase; PDBTitle: crystal structure analysis of full-length carboxyl-terminal2 src kinase at 2.5 a resolution

12	c2c0iA_	 Alignment		99.9	32	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
13	c2dvjA_	 Alignment		99.9	35	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
14	c1x27F_	 Alignment		99.9	39	PDB header: signaling protein Chain: F: PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of2 p130cas
15	c1y57A_	 Alignment		99.9	29	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
16	c2h8hA_	 Alignment		99.9	32	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
17	c2ablA_	 Alignment		99.8	35	PDB header: transferase Chain: A: PDB Molecule: abl tyrosine kinase; PDBTitle: sh3-sh2 domain fragment of human bcr-abl tyrosine kinase
18	c3nhnA_	 Alignment		99.8	38	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: crystal structure of the src-family kinase hck sh3-sh2-linker2 regulatory region
19	c3qwxX_	 Alignment		99.8	31	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
20	c3qwyA_	 Alignment		99.8	38	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
21	d1ng2a2	 Alignment	not modelled	99.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2dx1A_	 Alignment	not modelled	99.8	42	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
23	c1g83A_	 Alignment	not modelled	99.8	35	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: crystal structure of fyn sh3-sh2
24	c2dybA_	 Alignment	not modelled	99.8	30	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
25	c3nmzD_	 Alignment	not modelled	99.8	33	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
26	c2cubA_	 Alignment	not modelled	99.8	24	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
27	d1ov3a2	 Alignment	not modelled	99.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
28	c2df6A_	 Alignment	not modelled	99.8	34	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
						Fold: SH3-like barrel

29	d1utia_	Alignment	not modelled	99.8	100	Superfamily: SH3-domain Family: SH3-domain
30	c1zx6A_	Alignment	not modelled	99.8	36	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
31	c2ed0A_	Alignment	not modelled	99.8	26	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)
32	c2xmfA_	Alignment	not modelled	99.8	37	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
33	c2dmoA_	Alignment	not modelled	99.8	29	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)
34	c2fpeB_	Alignment	not modelled	99.8	24	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
35	c2yt6A_	Alignment	not modelled	99.8	28	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
36	c2yunA_	Alignment	not modelled	99.8	35	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
37	d1sema_	Alignment	not modelled	99.8	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c3jv3A_	Alignment	not modelled	99.8	28	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
39	c1x2qA_	Alignment	not modelled	99.8	39	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
40	c4glmD_	Alignment	not modelled	99.8	34	PDB header: signaling protein Chain: D: PDB Molecule: dynamamin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
41	c2dlpA_	Alignment	not modelled	99.8	25	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
42	c2yuoA_	Alignment	not modelled	99.8	33	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
43	d1uuea_	Alignment	not modelled	99.8	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	d1udla_	Alignment	not modelled	99.8	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
45	d1oeba_	Alignment	not modelled	99.8	95	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2nwmA_	Alignment	not modelled	99.8	36	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
47	d1wlpb2	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	c2dl4A_	Alignment	not modelled	99.8	32	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
49	c2rqrA_	Alignment	not modelled	99.8	12	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
50	c2kymA_	Alignment	not modelled	99.8	29	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from I.elongisporus in2 complex with ste20p peptide
51	c2dilA_	Alignment	not modelled	99.8	28	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
52	d1gcqa_	Alignment	not modelled	99.8	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	d1gria2	Alignment	not modelled	99.8	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
54	c2bz8B_	Alignment	not modelled	99.8	30	PDB header: sh3 domain Chain: B: PDB Molecule: sh3-domain kinase binding protein 1; PDBTitle: n-terminal sh3 domain of cin85 bound to cbl-b peptide
55	d1pwta_	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
56	d1j3ta_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	c2cudA_	Alignment	not modelled	99.7	19	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)
58	c2csqA_	Alignment	not modelled	99.7	25	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
59	c2dl3A_	Alignment	not modelled	99.7	29	PDB header: cell adhesion, signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the first sh3 domain of human sorbin2 and sh3 domain-containing protein 1
60	c2djqa_	Alignment	not modelled	99.7	23	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: the solution structure of the first sh3 domain of mouse sh32 domain containing ring finger 2
61	c2jxbA_	Alignment	not modelled	99.7	22	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
62	c2pqhA_	Alignment	not modelled	99.7	29	PDB header: structural protein Chain: A: PDB Molecule: spectrin alpha chain, brain; PDBTitle: structure of sh3 chimera with a type ii ligand linked to the chain c-2 terminal
63	d1u06a1	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2oi3A_	Alignment	not modelled	99.7	29	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoetic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
65	d1k4us_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2ct3A_	Alignment	not modelled	99.7	35	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
67	c2lj0A_	Alignment	not modelled	99.7	33	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
68	c1x69A_	Alignment	not modelled	99.7	28	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
69	c2drmB_	Alignment	not modelled	99.7	39	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
70	c2dnuA_	Alignment	not modelled	99.7	25	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
71	d1e6ga_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c2l0aA_	Alignment	not modelled	99.7	43	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
73	c2ekha_	Alignment	not modelled	99.7	19	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
74	c2ysqa_	Alignment	not modelled	99.7	34	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
75	c4igzA_	Alignment	not modelled	99.7	36	PDB header: protein binding Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 2; PDBTitle: crystal structure of the sh3 domain of human sorbin and sh3 domain-2 containing protein 2
76	c2egaA_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structure of the first sh3 domain from human2 kiaa0418 protein
77	c1wi7A_	Alignment	not modelled	99.7	25	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase binding protein 1; PDBTitle: solution structure of the sh3 domain of sh3-domain kinase2 binding protein 1
78	c2epdA_	Alignment	not modelled	99.7	26	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
79	c2eyxA_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
80	c2d1xD_	Alignment	not modelled	99.7	27	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

81	d1e6ha_	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c1x43A_	Alignment	not modelled	99.7	26	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)
83	c2azsA_	Alignment	not modelled	99.7	49	PDB header: signaling protein Chain: A: PDB Molecule: sh2-sh3 adapter protein drk; PDBTitle: nmr structure of the n-terminal sh3 domain of drk2 (calculated without noe restraints)
84	c1negA_	Alignment	not modelled	99.7	28	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
85	d1nega_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c1x6bA_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine exchange factor (gef) 16; PDBTitle: solution structures of the sh3 domain of human rho guanine2 exchange factor (gef) 16
87	c2yupA_	Alignment	not modelled	99.7	29	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
88	c2ebpA_	Alignment	not modelled	99.7	26	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
89	d1h8ka_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2dbmA_	Alignment	not modelled	99.7	37	PDB header: transferase, signaling protein Chain: A: PDB Molecule: sh3-containing grb2-like protein 2; PDBTitle: solution structures of the sh3 domain of human sh3-2 containing grb2-like protein 2
91	d1ue9a_	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c1x6gA_	Alignment	not modelled	99.7	19	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.
93	c1w70A_	Alignment	not modelled	99.7	33	PDB header: sh3 domain Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: sh3 domain of p40phox complexed with c-terminal polyproline2 region of p47phox
94	d1ujya_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c2dbkA_	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
96	c2ed1A_	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: 130 kda phosphatidylinositol 4,5-biphosphate- PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein
97	d1ov3a1	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c1zlmA_	Alignment	not modelled	99.7	43	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
99	c1x2pA_	Alignment	not modelled	99.7	27	PDB header: transferase Chain: A: PDB Molecule: protein arginine n-methyltransferase 2; PDBTitle: solution structure of the sh3 domain of the protein2 arginine n-methyltransferase 2
100	c3rniA_	Alignment	not modelled	99.7	20	PDB header: protein binding Chain: A: PDB Molecule: brain-specific angiogenesis inhibitor 1-associated protein PDBTitle: crystal structure of the sh3 domain from irsp53 (baiap2)
101	c2egeA_	Alignment	not modelled	99.7	24	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
102	c4esrB_	Alignment	not modelled	99.7	38	PDB header: protein binding Chain: B: PDB Molecule: jouberin; PDBTitle: molecular and structural characterization of the sh3 domain of ahi-12 in regulation of cellular resistance of bcr-abl+ chronic myeloid3 leukemia cells to tyrosine kinase inhibitors
103	d1uj0a_	Alignment	not modelled	99.7	54	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	d1uhfa_	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	c2dl8A_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: slit-robo rho gtpase-activating protein 2; PDBTitle: solution structure of the sh3 domain of human slit-robo rho2 gtpase-activating protein 2
						PDB header: signaling protein

106	c2js0A_	Alignment	not modelled	99.7	32	Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
107	c2eqiA_	Alignment	not modelled	99.7	30	PDB header: immune system, hydrolase Chain: A; PDB Molecule: phospholipase c, gamma 2; PDBTitle: solution structure of the sh3 domain from phospholipase c,2 gamma 2
108	c2j06B_	Alignment	not modelled	99.7	23	PDB header: signal transduction Chain: B; PDB Molecule: ras gtpase-activating protein 1; PDBTitle: crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
109	d1wfwA_	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2krnA_	Alignment	not modelled	99.7	40	PDB header: signaling protein Chain: A; PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
111	c2cucA_	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A; PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: solution structure of the sh3 domain of the mouse2 hypothetical protein sh3rf2
112	c3cqtA_	Alignment	not modelled	99.7	45	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
113	c2j6kE_	Alignment	not modelled	99.7	33	PDB header: protein binding Chain: E; PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
114	c2o2oA_	Alignment	not modelled	99.7	27	PDB header: protein binding Chain: A; PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
115	d1phta_	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2kbtA_	Alignment	not modelled	99.7	33	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
117	c2e5kA_	Alignment	not modelled	99.7	19	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
118	c1wvxA_	Alignment	not modelled	99.7	38	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
119	c2i0nA_	Alignment	not modelled	99.7	23	PDB header: structural protein Chain: A; PDB Molecule: class vii unconventional myosin; PDBTitle: structure of dictyostelium discoideum myosin vii sh3 domain2 with adjacent proline rich region
120	c2dl7A_	Alignment	not modelled	99.7	29	PDB header: structural genomics, unknown function Chain: A; PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein