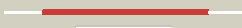








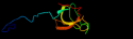


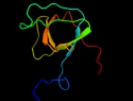

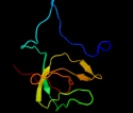





Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	A1ZAY1
Date	Wed Feb 13 11:18:13 GMT 2013
Unique Job ID	c8a3442ff5a43ab9

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	22	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c21qnA_	 Alignment		100.0	16	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c213sA_	 Alignment		100.0	15	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
4	c2eyzA_	 Alignment		100.0	13	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1ng2A_	 Alignment		99.9	23	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
6	c1ov3A_	 Alignment		99.9	24	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
7	c1mv3A_	 Alignment		99.9	14	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
8	c2rqrA_	 Alignment		99.9	11	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
9	c2yt6A_	 Alignment		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
10	d1ng2a2	 Alignment		99.8	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
11	c2fpeB_	 Alignment		99.8	21	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

12	c2oi3A_	Alignment		99.8	26	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
13	c2dlpA_	Alignment		99.8	22	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
14	c2cudA_	Alignment		99.8	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)
15	c2e5kA_	Alignment		99.8	23	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
16	c2j06B_	Alignment		99.8	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
17	d1udla_	Alignment		99.8	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
18	c3rnjA_	Alignment		99.8	28	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)
19	c1x43A_	Alignment		99.8	20	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)
20	c2kymA_	Alignment		99.8	28	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
21	c2d8hA_	Alignment	not modelled	99.8	23	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
22	c1x6gA_	Alignment	not modelled	99.8	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.
23	d1uffa_	Alignment	not modelled	99.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	c2yupA_	Alignment	not modelled	99.8	22	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
25	c2jteA_	Alignment	not modelled	99.8	21	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
26	c2dx1A_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
27	c2dm1A_	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
28	d1spka_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain of hypothetical protein

29	c2dl5A_	Alignment	not modelled	99.7	19	Chain: A: PDB Molecule: klaa0/69 protein; PDBTitle: solution structure of the first sh3 domain of human2 klaa0769 protein
30	c1zuyB_	Alignment	not modelled	99.7	31	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
31	c2csgA_	Alignment	not modelled	99.7	17	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
32	c2lcsA_	Alignment	not modelled	99.7	24	PDB header: transferase, signaling protein Chain: A: PDB Molecule: nap1-binding protein 2; PDBTitle: yeast nbp2p sh3 domain in complex with a peptide from ste20p
33	c1x6bA_	Alignment	not modelled	99.7	18	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
34	d1k4us_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
35	c2ct3A_	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
36	c1wx6A_	Alignment	not modelled	99.7	15	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: cytoplasmic protein nck2; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck2
37	c3nmzD_	Alignment	not modelled	99.7	21	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asf
38	c2jxbA_	Alignment	not modelled	99.7	14	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
39	d1ue9a_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	c2lj0A_	Alignment	not modelled	99.7	29	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
41	d1wiea_	Alignment	not modelled	99.7	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	d1phta_	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	c2rf0D_	Alignment	not modelled	99.7	29	PDB header: transferase Chain: D: PDB Molecule: mitogen-activated protein kinase kinase kinase 10; PDBTitle: crystal structure of human mixed lineage kinase map3k10 sh3 domain
44	c2egeA_	Alignment	not modelled	99.7	17	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein klaa1666; PDBTitle: solution structure of the third sh3 domain from human2 klaa1666 protein
45	c1bb9A_	Alignment	not modelled	99.7	21	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
46	d1bb9a_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	d1ov3a2	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	c1z9zA_	Alignment	not modelled	99.7	23	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
49	c2cucA_	Alignment	not modelled	99.7	28	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
50	c2dmoA_	Alignment	not modelled	99.7	18	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)
51	c3jv3A_	Alignment	not modelled	99.7	23	PDB header: protein binding Chain: A: PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
52	c1x2qA_	Alignment	not modelled	99.7	16	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
53	c1zukA_	Alignment	not modelled	99.7	22	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
54	d1arka_	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: endocytosis/exocytosis

55	c2csiA	Alignment	not modelled	99.7	21	Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the third sh3 domain of human rim-2 binding protein 2
56	c3cqtA	Alignment	not modelled	99.7	35	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
57	c1yn8E	Alignment	not modelled	99.7	27	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
58	c4afzC	Alignment	not modelled	99.7	32	PDB header: hydrolase/de novo protein Chain: C: PDB Molecule: fynomer; PDBTitle: human chymase - fynomer complex
59	c2rqvA	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
60	d2iima1	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c2cubA	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
62	d1k9aa1	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c2eyyA	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
64	c2kt1A	Alignment	not modelled	99.7	28	PDB header: protein binding Chain: A: PDB Molecule: phosphatidylinositol 3-kinase regulatory subunit PDBTitle: solution nmr structure of the sh3 domain from the p85beta2 subunit of phosphatidylinositol 3-kinase from h.sapiens,3 northeast structural genomics consortium target hr5531e
65	d1fmka1	Alignment	not modelled	99.7	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2yunA	Alignment	not modelled	99.7	24	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
67	c2m0yA	Alignment	not modelled	99.7	22	PDB header: apoptosis Chain: A: PDB Molecule: dedicator of cytokinesis protein 1; PDBTitle: solution structure of the sh3 domain of dock180
68	c1zx6A	Alignment	not modelled	99.7	24	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
69	c2ct4A	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution structure of the sh3 domain of the cdc42-2 interacting protein 4
70	d1y57a1	Alignment	not modelled	99.7	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	d1gcqa	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	d1u5sa1	Alignment	not modelled	99.7	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d1oeba	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2enmA	Alignment	not modelled	99.7	25	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
75	c2df6A	Alignment	not modelled	99.7	24	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
76	c1w1fA	Alignment	not modelled	99.7	33	PDB header: sh3-domain Chain: A: PDB Molecule: tyrosine-protein kinase lyn; PDBTitle: sh3 domain of human lyn tyrosine kinase
77	c1x27F	Alignment	not modelled	99.7	23	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
78	d1efna	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	c1s1nA	Alignment	not modelled	99.7	28	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
80	c2ed1A	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: 130 kda phosphatidylinositol 4,5-bisphosphate- PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-bisphosphate-dependent arf1 gtpase-3

						activating protein
81	c2yuoA_	Alignment	not modelled	99.7	18	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
82	d1sema_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	c2ysqA_	Alignment	not modelled	99.7	18	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
84	c2i0nA_	Alignment	not modelled	99.7	19	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
85	c2dl4A_	Alignment	not modelled	99.7	26	PDB header: signaling protein Chain: A; PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
86	d1uhca_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2k6dA_	Alignment	not modelled	99.7	23	PDB header: sh3 domain/ubiquitin Chain: A; PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
88	d1lutia_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	d1ug1a_	Alignment	not modelled	99.7	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2kgtA_	Alignment	not modelled	99.7	23	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
91	c4esrB_	Alignment	not modelled	99.7	28	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
92	d1shfa_	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1wfwA_	Alignment	not modelled	99.7	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2dilA_	Alignment	not modelled	99.7	30	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
95	c2xmfA_	Alignment	not modelled	99.7	28	PDB header: motor protein Chain: A; PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
96	c2eczA_	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A; PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain of sorbin and sh32 domain-containing protein 1
97	c2lx7A_	Alignment	not modelled	99.7	21	PDB header: protein binding Chain: A; PDB Molecule: growth arrest-specific protein 7; PDBTitle: solution nmr structure of sh3 domain of growth arrest-specific protein2 7 (gas7) (fragment 1-60) from homo sapiens, northeast structural3 genomics consortium (nesg) target hr8574a
98	d1gcqc_	Alignment	not modelled	99.7	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	d1jo8a_	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2eqiA_	Alignment	not modelled	99.7	28	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
101	d1cska_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	d1pwta_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2dnuA_	Alignment	not modelled	99.7	17	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
104	c4glmD_	Alignment	not modelled	99.7	29	PDB header: signaling protein Chain: D; PDB Molecule: dynamidin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
105	c3qwyA_	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
						PDB header: structural genomics, unknown function

106	c2creA	Alignment	not modelled	99.7	21	Chain: A; PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2 human cdna
107	d1wlpb2	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2egcA	Alignment	not modelled	99.7	14	PDB header: signaling protein Chain: A; PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structure of the fifth sh3 domain from human2 kiaa0418 protein
109	c1zlmA	Alignment	not modelled	99.7	26	PDB header: signaling protein Chain: A; PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
110	c1y57A	Alignment	not modelled	99.7	34	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
111	d1ujya	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c3qwxX	Alignment	not modelled	99.7	15	PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
113	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
114	d1uuea	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2k2mA	Alignment	not modelled	99.7	23	PDB header: signaling protein Chain: A; PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
116	c2ed0A	Alignment	not modelled	99.7	25	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)
117	d2fo0a1	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c2djgA	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
119	c2h8hA	Alignment	not modelled	99.7	33	PDB header: transferase Chain: A; PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
120	d1u06a1	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain