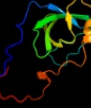
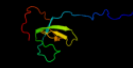
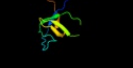
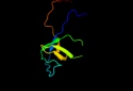


Phyre2

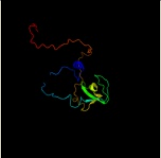
Email	I.a.kelley@imperial.ac.uk
Description	Q06449
Date	Mon Jul 2 19:31:53 BST 2012
Unique Job ID	3f84c9c62279e9a8

Detailed template information

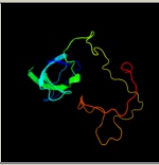
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1mv3A_	 Alignment		99.9	22	PDB header: endocytosis/exocytosis Chain: A; PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
2	c1griA_	 Alignment		99.9	36	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
3	d1ng2a2	 Alignment		99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
4	c2dx1A_	 Alignment		99.8	29	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein aseq
5	c2ysqA_	 Alignment		99.8	22	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
6	c3nmzD_	 Alignment		99.8	31	PDB header: cell adhesion/cell cycle Chain: D; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with aseq
7	c2ed0A_	 Alignment		99.8	31	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)
8	c1x2qA_	 Alignment		99.8	22	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
9	c2cubA_	 Alignment		99.8	26	PDB header: signaling protein Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
10	c2yunA_	 Alignment		99.8	36	PDB header: protein transport Chain: A; PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
11	c2yuoA_	 Alignment		99.8	27	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

12	c2eyyA	Alignment		99.8	26	PDB header: signaling protein Chain: A: PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
13	c2kymA	Alignment		99.8	20	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
14	c2djgA	Alignment		99.8	36	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
15	c2dlpA	Alignment		99.8	28	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
16	c2rqrA	Alignment		99.8	19	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
17	c2nwmA	Alignment		99.8	33	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
18	c2yupA	Alignment		99.8	25	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
19	c2oi3A	Alignment		99.8	23	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)
20	c2dmoA	Alignment		99.8	33	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)
21	d1ov3a2	Alignment	not modelled	99.8	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2ebpA	Alignment	not modelled	99.8	25	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
23	c2ekhA	Alignment	not modelled	99.8	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
24	c2xmfA	Alignment	not modelled	99.8	29	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
25	c2epdA	Alignment	not modelled	99.8	21	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
26	c2df6A	Alignment	not modelled	99.8	31	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
27	c2dbmA	Alignment	not modelled	99.8	34	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
28	c2dl3A	Alignment	not modelled	99.8	33	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
29	d1udla_	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
30	d1oeba_	Alignment	not modelled	99.8	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
31	c2i0nA_	Alignment	not modelled	99.8	25	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
32	c2eyxA_	Alignment	not modelled	99.8	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
33	c2eyzA_	Alignment	not modelled	99.8	17	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
34	c2dbkA_	Alignment	not modelled	99.8	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
35	c2dl4A_	Alignment	not modelled	99.8	42	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
36	d1ujya_	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
37	c2pqhA_	Alignment	not modelled	99.8	33	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
38	d1utiA_	Alignment	not modelled	99.8	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	c2jxbA_	Alignment	not modelled	99.8	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
40	c1wi7A_	Alignment	not modelled	99.8	33	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
41	d1k4us_	Alignment	not modelled	99.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	d1gria2_	Alignment	not modelled	99.8	49	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	d1sema_	Alignment	not modelled	99.8	46	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	d1gcqa_	Alignment	not modelled	99.8	52	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
45	d1pwta_	Alignment	not modelled	99.8	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2l0aA_	Alignment	not modelled	99.8	26	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
47	c1zx6A_	Alignment	not modelled	99.8	100	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
48	c2yt6A_	Alignment	not modelled	99.8	25	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
49	d1wlpb2_	Alignment	not modelled	99.8	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c2dl8A_	Alignment	not modelled	99.8	21	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
51	c2dilA_	Alignment	not modelled	99.8	24	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	c1x43A_	Alignment	not modelled	99.7	16	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)
53	d1uuea_	Alignment	not modelled	99.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: structural protein

54	c1negA_	Alignment	not modelled	99.7	33	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
55	d1nega_	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	d1j3ta_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	c3qwxX_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
58	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
59	c2ed1A_	Alignment	not modelled	99.7	21	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
60	c1zlmA_	Alignment	not modelled	99.7	33	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
61	d1uj0a_	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	d1ue9a_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c1w70A_	Alignment	not modelled	99.7	36	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
64	c2bz8B_	Alignment	not modelled	99.7	27	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
65	c2e5kA_	Alignment	not modelled	99.7	28	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
66	c2csqA_	Alignment	not modelled	99.7	19	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
67	c2d1xD_	Alignment	not modelled	99.7	34	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
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	d1uffa_	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c2ct3A_	Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
72	c2lqnA_	Alignment		99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
73	d1u06a1	Alignment	not modelled	99.7	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c4esrB_	Alignment	not modelled	99.7	31	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
75	c1wyxA_	Alignment	not modelled	99.7	27	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
76	c1wx6A_	Alignment	not modelled	99.7	23	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
77	c2dnuA_	Alignment	not modelled	99.7	21	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
78	c2dviA_	Alignment	not modelled	99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii

79	c2cudA	Alignment	not modelled	99.7	14	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adopter protein (slap)
80	c3qwyA	Alignment	not modelled	99.7	25	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
81	d1e6ga	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2o2oA	Alignment	not modelled	99.7	25	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
83	c1ng2A	Alignment	not modelled	99.7	22	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
84	c1zukA	Alignment	not modelled	99.7	25	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti 1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
85	c2fpeB	Alignment	not modelled	99.7	25	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
86	c2egaA	Alignment	not modelled	99.7	25	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
87	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.
88	d1uhfa	Alignment	not modelled	99.7	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c2bzxA	Alignment	not modelled	99.7	27	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
90	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
91	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
92	c2krnA	Alignment	not modelled	99.7	31	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
93	c1x6bA	Alignment	not modelled	99.7	25	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
94	d1oota	Alignment	not modelled	99.7	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c1z9qA	Alignment	not modelled	99.7	33	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
96	d1e6ha	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2dl7A	Alignment	not modelled	99.7	30	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
98	c2js0A	Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
99	d2v1ra1	Alignment	not modelled	99.7	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	d1ov3a1	Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	d1jo8a	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2csiA	Alignment	not modelled	99.7	27	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the third sh3 domain of human rim-2 binding protein 2
103	d1uhca	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c2d8hA	Alignment	not modelled	99.7	35	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2

					protein sh3yl1
105	c2cucA_	Alignment	not modelled	99.7	36 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
106	c2l3sA_	Alignment		99.7	22 PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
107	d1h8ka_	Alignment	not modelled	99.7	30 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c3rnjA_	Alignment	not modelled	99.7	23 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)
109	c2rqvA_	Alignment	not modelled	99.7	21 PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
110	c1wxtA_	Alignment	not modelled	99.7	27 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein flj21522; PDBTitle: solution structure of the sh3 domain of human hypothetical2 protein flj21522
111	d1wiea_	Alignment	not modelled	99.7	15 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	d1ng2a1	Alignment	not modelled	99.7	23 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2d8jA_	Alignment	not modelled	99.7	30 PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
114	c2eczA_	Alignment	not modelled	99.7	28 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
115	d1ugva_	Alignment	not modelled	99.7	29 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	d1ug1a_	Alignment	not modelled	99.7	21 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2egeA_	Alignment	not modelled	99.7	29 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
118	d1spka_	Alignment	not modelled	99.7	25 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c1wxbA_	Alignment	not modelled	99.7	28 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: epidermal growth factor receptor pathway PDBTitle: solution structure of the sh3 domain from human epidermal2 growth factor receptor pathway substrate 8-like protein
120	c2k2mA_	Alignment	not modelled	99.7	21 PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding