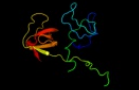
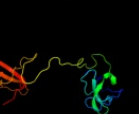
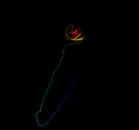
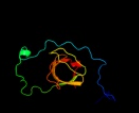
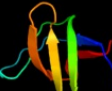


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q9U559
Date	Sun Jul 8 12:02:30 BST 2012
Unique Job ID	c127a9c2b84eb636

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		99.9	20	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2l3sA_	 Alignment		99.9	15	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
3	c2lqnA_	 Alignment		99.9	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
4	c1ov3A_	 Alignment		99.9	14	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
5	c1mv3A_	 Alignment		99.9	17	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
6	c1ng2A_	 Alignment		99.9	14	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
7	c2eyzA_	 Alignment		99.9	16	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
8	c3nmzD_	 Alignment		99.8	25	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asef
9	c2rqrA_	 Alignment		99.8	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
10	c2dx1A_	 Alignment		99.8	31	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
11	c3qwyA_	 Alignment		99.8	27	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2

12	d1ng2a2	Alignment		99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
13	c3qwxX	Alignment		99.8	27	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
14	c1zuka	Alignment		99.8	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
15	c2kymA	Alignment		99.8	36	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
16	c1x6gA	Alignment		99.8	18	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.
17	c2yupA	Alignment		99.8	24	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
18	c2yt6A	Alignment		99.8	26	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
19	c2ed0A	Alignment		99.8	28	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)
20	c2ysqA	Alignment		99.8	26	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
21	c2oi3A	Alignment	not modelled	99.8	25	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)
22	c2lcsA	Alignment	not modelled	99.8	45	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
23	c2cudA	Alignment	not modelled	99.8	21	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpoter protein (slap)
24	d1jo8a	Alignment	not modelled	99.8	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	d1uj0a	Alignment	not modelled	99.8	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
26	c2ct3A	Alignment	not modelled	99.8	23	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
27	c1x2qA	Alignment	not modelled	99.8	31	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
28	d1phta	Alignment	not modelled	99.8	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

29	clyn8E_	Alignment	not modelled	99.8	45	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
30	c2d1xD_	Alignment	not modelled	99.8	33	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
31	c3rnjA_	Alignment	not modelled	99.8	18	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)
32	d1ov3a2	Alignment	not modelled	99.8	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
33	c2fpeB_	Alignment	not modelled	99.8	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
34	c2eyyA_	Alignment	not modelled	99.8	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
35	c2ed1A_	Alignment	not modelled	99.8	26	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
36	c2ktiA_	Alignment	not modelled	99.8	23	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
37	d1udla_	Alignment	not modelled	99.8	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c1x6bA_	Alignment	not modelled	99.8	21	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
39	c2ebpA_	Alignment	not modelled	99.8	21	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
40	c2jxbA_	Alignment	not modelled	99.8	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
41	c2yunA_	Alignment	not modelled	99.8	26	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
42	c2nwmA_	Alignment	not modelled	99.8	27	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
43	c2cubA_	Alignment	not modelled	99.8	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
44	c2yuoA_	Alignment	not modelled	99.8	31	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
45	d1ue9a_	Alignment	not modelled	99.8	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2d8hA_	Alignment	not modelled	99.8	22	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
47	d1utia_	Alignment	not modelled	99.8	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	c2e5kA_	Alignment	not modelled	99.8	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
49	d1ilja_	Alignment	not modelled	99.8	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c2epdA_	Alignment	not modelled	99.8	25	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
51	c4esrB_	Alignment	not modelled	99.8	24	PDB header: protein binding Chain: B: PDB Molecule: jouberein; 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
52	c2l0aA_	Alignment	not modelled	99.8	28	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 PDB header: signaling protein

53	c1x69A	Alignment	not modelled	99.8	30	Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
54	c2df6A	Alignment	not modelled	99.8	23	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
55	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
56	c1x43A	Alignment	not modelled	99.8	18	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)
57	c2enmA	Alignment	not modelled	99.8	28	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
58	c2ekhA	Alignment	not modelled	99.8	18	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
59	c2dl8A	Alignment	not modelled	99.8	27	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
60	c2dbkA	Alignment	not modelled	99.8	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
61	c1zlmA	Alignment	not modelled	99.8	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
62	dlwiea	Alignment	not modelled	99.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c2djgA	Alignment	not modelled	99.8	22	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
64	c2dl3A	Alignment	not modelled	99.8	27	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
65	c2dl4A	Alignment	not modelled	99.8	28	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
66	dlspka	Alignment	not modelled	99.8	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c2dmoA	Alignment	not modelled	99.7	25	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)
68	c2j06B	Alignment	not modelled	99.7	22	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
69	dl0eba	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c1zuyB	Alignment	not modelled	99.7	27	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
71	c2eczA	Alignment	not modelled	99.7	24	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
72	c2dlpA	Alignment	not modelled	99.7	23	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
73	c2cucA	Alignment	not modelled	99.7	27	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
74	c1wi7A	Alignment	not modelled	99.7	23	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
75	dlwlpb2	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c2xmfA	Alignment	not modelled	99.7	25	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
77	c2dbmA	Alignment	not modelled	99.7	32	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
78	dlfmka1	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

79	dlgrla2	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2i0nA	Alignment	not modelled	99.7	20	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
81	c1w70A	Alignment	not modelled	99.7	34	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
82	d1k4us	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	d1jqqa	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	d2v1ra1	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	c2rqvA	Alignment	not modelled	99.7	36	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
86	c2drmB	Alignment	not modelled	99.7	27	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
87	c1wx6A	Alignment	not modelled	99.7	25	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
88	c1z9zA	Alignment	not modelled	99.7	32	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
89	c2dl7A	Alignment	not modelled	99.7	32	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
90	c2ct4A	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strcture of the sh3 domain of the cdc42-2 interacting protein 4
91	d1gcqa	Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	d1sema	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1uffa	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c3cqtA	Alignment	not modelled	99.7	30	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
95	d1pwta	Alignment	not modelled	99.7	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2eqiA	Alignment	not modelled	99.7	23	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
97	d1efna	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c2eyxA	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
99	d1uuea	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2rf0D	Alignment	not modelled	99.7	33	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
101	c2dnuA	Alignment	not modelled	99.7	20	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
102	d1arka	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2azsA	Alignment	not modelled	99.7	30	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)
104	d1k9aa1	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein

105	c2egaA	Alignment	not modelled	99.7	21	2a; PDBTitle: solution structure of the first sh3 domain from human2 kiaa0418 protein
106	c2js0A	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
107	d1u06a1	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2krnA	Alignment	not modelled	99.7	26	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
109	c2dl5A	Alignment	not modelled	99.7	16	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
110	c1zx6A	Alignment	not modelled	99.7	34	PDB header: protein binding Chain: A: PDB Molecule: yp154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
111	d1zuua1	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c1x2pA	Alignment	not modelled	99.7	23	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
113	d1bb9a	Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c1bb9A	Alignment	not modelled	99.7	19	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
115	c2pqhA	Alignment	not modelled	99.7	28	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
116	d1qlya	Alignment	not modelled	99.7	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2d8jA	Alignment	not modelled	99.7	27	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
118	d1u5sa1	Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2ke9A	Alignment	not modelled	99.7	27	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
120	d1y57a1	Alignment	not modelled	99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain