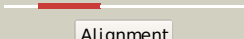

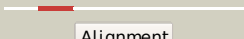

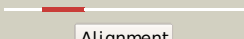








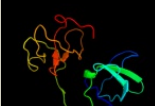

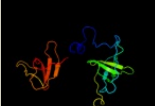

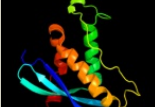




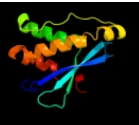
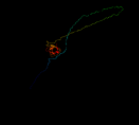





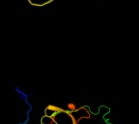



Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	A1X283
Date	Thu Apr 26 09:27:51 BST 2012
Unique Job ID	5ef591bb9fd08c0a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dybA_	 Alignment		100.0	21	PDB header: oxidoreductase Chain: A; PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
2	c2I73A_	 Alignment		100.0	21	PDB header: oxidoreductase regulator Chain: A; PDB Molecule: nadph oxidase organizer 1; PDBTitle: structure of the noxo1b px domain
3	c1ng2A_	 Alignment		100.0	34	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
4	c1griA_	 Alignment		100.0	17	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
5	c1ov3A_	 Alignment		100.0	42	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
6	d1kq6a_	 Alignment		100.0	37	Fold: PX domain Superfamily: PX domain Family: PX domain
7	c2I3sA_	 Alignment		100.0	20	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
8	c2eyzA_	 Alignment		100.0	20	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
9	d1h6ha_	 Alignment		99.9	27	Fold: PX domain Superfamily: PX domain Family: PX domain
10	d1o7ka_	 Alignment		99.9	37	Fold: PX domain Superfamily: PX domain Family: PX domain
11	c2reaA_	 Alignment		99.9	31	PDB header: transferase Chain: A; PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: crystal structures of c2alpha-pi3 kinase px-domain domain2 indicate conformational change associated with ligand3 binding.

12	c2iw1X_	Alignment		99.9	28	PDB header: transferase Chain: X: PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: structure of the px domain of phosphoinositide 3-kinase-2 c2alpha
13	c1mv3A_	Alignment		99.9	11	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
14	c2czoA_	Alignment		99.9	29	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the px domain of bem1p
15	c2wweA_	Alignment		99.8	22	PDB header: transferase Chain: A: PDB Molecule: phosphoinositide-3-kinase, class 2, gamma PDBTitle: crystal structure of the phox homology domain of human2 phosphoinositide-3-kinase-c2-gamma
16	c2egcA_	Alignment		99.7	54	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
17	d1ng2a2	Alignment		99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
18	d1wlpb2	Alignment		99.7	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
19	c2rqrA_	Alignment		99.7	15	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
20	d1udla_	Alignment		99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c2jmcA_	Alignment	not modelled	99.7	20	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
22	d1ov3a1	Alignment	not modelled	99.6	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	c2eyyA_	Alignment	not modelled	99.6	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
24	c2ekhA_	Alignment	not modelled	99.6	36	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
25	d1ng2a1	Alignment	not modelled	99.6	47	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
26	c2yt6A_	Alignment	not modelled	99.6	22	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
27	c2v14A_	Alignment	not modelled	99.6	16	PDB header: transport protein Chain: A: PDB Molecule: kinesin-like motor protein c20orf23; PDBTitle: kinesin 16b phox-homology domain (kif16b)
28	c2dx1A_	Alignment	not modelled	99.6	17	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
						Fold: SH3-like barrel

29	d1ue9a_	Alignment	not modelled	99.6	21	Superfamily: SH3-domain Family: SH3-domain
30	c2egaA_	Alignment	not modelled	99.6	73	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
31	c1x2qA_	Alignment	not modelled	99.6	14	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
32	c3nmzD_	Alignment	not modelled	99.6	14	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
33	c2dila_	Alignment	not modelled	99.6	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
34	c2kymA_	Alignment	not modelled	99.6	17	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
35	c2cskA_	Alignment	not modelled	99.6	15	PDB header: protein transport Chain: A: PDB Molecule: sorting nexin 12; PDBTitle: solution structure of px domain from human snx12
36	c2xmfA_	Alignment	not modelled	99.6	27	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
37	c2epdA_	Alignment	not modelled	99.6	18	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
38	c2yunA_	Alignment	not modelled	99.6	27	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
39	c3iq2B_	Alignment	not modelled	99.6	21	PDB header: protein transport Chain: B: PDB Molecule: sorting nexin-7; PDBTitle: human sorting nexin 7, phox homology (px) domain
40	c2yuoA_	Alignment	not modelled	99.6	17	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
41	c1xteA_	Alignment	not modelled	99.6	23	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase sgk3; PDBTitle: crystal structure of cisk-px domain
42	d1xtea_	Alignment	not modelled	99.6	23	Fold: PX domain Superfamily: PX domain Family: PX domain
43	c2dmoA_	Alignment	not modelled	99.6	23	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)
44	c2d8jA_	Alignment	not modelled	99.6	30	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
45	d1ug1a_	Alignment	not modelled	99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	c2dlpA_	Alignment	not modelled	99.6	19	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
47	c2dnuA_	Alignment	not modelled	99.6	32	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
48	c2i4kA_	Alignment	not modelled	99.6	22	PDB header: protein transport Chain: A: PDB Molecule: sorting nexin-1; PDBTitle: solution structure of the px domain of sorting nexin 1
49	c2ed0A_	Alignment	not modelled	99.6	27	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)
50	c2e5kA_	Alignment	not modelled	99.6	18	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-cell12 receptor signaling 1
51	c2bz8B_	Alignment	not modelled	99.6	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
52	c2eyxA_	Alignment	not modelled	99.6	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
53	c1zukA_	Alignment	not modelled	99.6	21	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mt1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
54	c2fpeB_	Alignment	not modelled	99.6	14	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
55	c2dl4A_	Alignment	not modelled	99.6	20	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
56	d1ov3a2	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	d1ujya_	Alignment	not modelled	99.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
58	c2ysqA_	Alignment	not modelled	99.6	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
59	d1nega_	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c1negA_	Alignment	not modelled	99.5	21	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
61	d1wlpb1	Alignment	not modelled	99.5	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c2ke9A_	Alignment	not modelled	99.5	22	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
63	d1e6ga_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c1x6gA_	Alignment	not modelled	99.5	15	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.
65	d1pwta_	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	d1phta_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c1zlmA_	Alignment	not modelled	99.5	13	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
68	c2djgA_	Alignment	not modelled	99.5	17	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
69	c3rniA_	Alignment	not modelled	99.5	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)
70	c2yupA_	Alignment	not modelled	99.5	20	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
71	c2dbkA_	Alignment	not modelled	99.5	16	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
72	d1e6ha_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2cucA_	Alignment	not modelled	99.5	24	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	c2pqhA_	Alignment	not modelled	99.5	23	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
75	c2cudA_	Alignment	not modelled	99.5	23	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)
76	c2ct3A_	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
77	c2ebpA_	Alignment	not modelled	99.5	17	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
78	c1s1nA_	Alignment	not modelled	99.5	30	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
79	d1k4us_	Alignment	not modelled	99.5	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c2dviA_	Alignment	not modelled	99.5	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog,

						PDBTitle: phosphorylated crk-ii PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
81	c2cubA_	Alignment	not modelled	99.5	26	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
82	c2ed1A_	Alignment	not modelled	99.5	13	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)
83	c2oi3A_	Alignment	not modelled	99.5	30	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
84	c2dbmA_	Alignment	not modelled	99.5	10	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2
85	c2df6A_	Alignment	not modelled	99.5	25	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
86	c2l0aA_	Alignment	not modelled	99.5	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	d1uhca_	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	d1uuea_	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c2jxbA_	Alignment	not modelled	99.5	16	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
90	d1gcqc_	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	d1uj0a_	Alignment	not modelled	99.5	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c1x69A_	Alignment	not modelled	99.5	24	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
93	d1uhfa_	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2dl8A_	Alignment	not modelled	99.5	18	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
95	c2csqA_	Alignment	not modelled	99.5	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
96	c1wi7A_	Alignment	not modelled	99.5	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
97	c2dl3A_	Alignment	not modelled	99.5	20	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
98	c2d1xD_	Alignment	not modelled	99.5	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
99	c1x2pA_	Alignment	not modelled	99.5	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
100	c2nwmA_	Alignment	not modelled	99.5	21	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
101	c1x43A_	Alignment	not modelled	99.5	22	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)
102	d1u06a1	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2i0nA_	Alignment	not modelled	99.5	14	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
104	c2kbtA_	Alignment	not modelled	99.5	22	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
						Fold: SH3-like barrel

105	d1wfwA_	Alignment	not modelled	99.5	27	Superfamily: SH3-domain Family: SH3-domain
106	c1zuyB_	Alignment	not modelled	99.5	32	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
107	d1gria2	Alignment	not modelled	99.5	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2o2oA_	Alignment	not modelled	99.5	24	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
109	c1zx6A_	Alignment	not modelled	99.5	22	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
110	d1oeba_	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c1x6bA_	Alignment	not modelled	99.5	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
112	c3qwxX_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
113	d1efna_	Alignment	not modelled	99.5	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2bzxA_	Alignment	not modelled	99.5	20	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
115	c3qwyA_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
116	c2krnA_	Alignment	not modelled	99.5	32	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
117	d1h8ka_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c2eqiA_	Alignment	not modelled	99.5	20	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
119	c1z9qA_	Alignment	not modelled	99.5	14	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
120	d1ql5a_	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain