
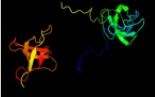


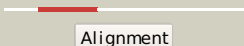



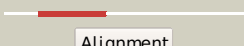
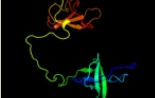
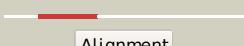

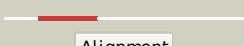




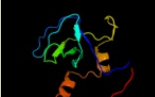






Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P29366
Date	Mon Jul 2 19:15:38 BST 2012
Unique Job ID	ea28636af6198f4b

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2lqnA_	 Alignment		100.0	21	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
2	c2dybA_	 Alignment		100.0	19	PDB header: oxidoreductase Chain: A; PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
3	c1griA_	 Alignment		100.0	27	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
4	c2l3sA_	 Alignment		100.0	21	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
5	c2eyzA_	 Alignment		100.0	18	PDB header: signaling protein Chain: A; PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
6	c1ng2A_	 Alignment		99.9	18	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
7	c1ov3A_	 Alignment		99.9	20	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
8	c2czoA_	 Alignment		99.9	100	PDB header: signaling protein Chain: A; PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the px domain of bem1p
9	c2rqvA_	 Alignment		99.9	97	PDB header: signaling protein Chain: A; PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
10	c2kymA_	 Alignment		99.9	54	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
11	c2l73A_	 Alignment		99.9	18	PDB header: oxidoreductase regulator Chain: A; PDB Molecule: nadh oxidase organizer 1; PDBTitle: structure of the noxo1b px domain

12	d1kq6a_	Alignment		99.8	20	Fold: PX domain Superfamily: PX domain Family: PX domain
13	d1h6ha_	Alignment		99.8	18	Fold: PX domain Superfamily: PX domain Family: PX domain
14	d1o7ka_	Alignment		99.8	22	Fold: PX domain Superfamily: PX domain Family: PX domain
15	d1ip9a_	Alignment		99.8	96	Fold: beta-Grasp (ubiquitin-like) Superfamily: CAD & PB1 domains Family: PB1 domain
16	d1udla_	Alignment		99.8	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
17	c1mv3A_	Alignment		99.7	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
18	c2rqrA_	Alignment		99.7	20	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
19	c3nmzD_	Alignment		99.6	20	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asf
20	c2ed1A_	Alignment		99.6	27	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
21	c2yt6A_	Alignment	not modelled	99.6	23	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
22	d1ue9a_	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	c2yupA_	Alignment	not modelled	99.6	20	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
24	d1jo8a_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2ysqA_	Alignment	not modelled	99.6	17	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
26	c2wweA_	Alignment	not modelled	99.6	25	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
27	c2cubA_	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
						PDB header: transferase Chain: A: PDB Molecule: phosphatidylinositol-4-phosphate 3-

28	c2reaA	Alignment	not modelled	99.6	23	kinase c2 PDBTitle: crystal structures of c2alpha-pi3 kinase px-domain domain2 indicate conformational change associated with ligand3 binding.
29	d1ng2a2	Alignment	not modelled	99.6	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
30	c3qwxX	Alignment	not modelled	99.6	24	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
31	c2enmA	Alignment	not modelled	99.6	25	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
32	c2yuoA	Alignment	not modelled	99.5	21	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
33	c2ed0A	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: abl interactor 2; PDBTitle: solution structure of the sh3 domain of abl interactor 2 (abelson interactor 2)
34	c1zukA	Alignment	not modelled	99.5	15	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
35	c2oi3A	Alignment	not modelled	99.5	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)
36	c3qwyA	Alignment	not modelled	99.5	26	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
37	c2eyyA	Alignment	not modelled	99.5	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
38	c1x2qA	Alignment	not modelled	99.5	27	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
39	c2d1xD	Alignment	not modelled	99.5	22	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
40	c2d8jA	Alignment	not modelled	99.5	20	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
41	c2jxbA	Alignment	not modelled	99.5	15	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
42	c2epdA	Alignment	not modelled	99.5	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
43	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
44	c2yunA	Alignment	not modelled	99.5	23	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
45	c1wx6A	Alignment	not modelled	99.5	21	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
46	c2i0nA	Alignment	not modelled	99.5	15	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
47	c1x43A	Alignment	not modelled	99.5	21	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)
48	c2ct3A	Alignment	not modelled	99.5	17	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
49	c2iwlX	Alignment	not modelled	99.5	24	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
50	c2df6A	Alignment	not modelled	99.5	19	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
51	c2dl8A	Alignment	not modelled	99.5	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
52	d1pwta	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

53	c2eyxA	Alignment	not modelled	99.5	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
54	c1yn8E	Alignment	not modelled	99.5	39	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
55	d1gria2	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	c2ekhA	Alignment	not modelled	99.5	11	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
57	c2nwmA	Alignment	not modelled	99.5	24	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
58	d1uuea	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	c2dlpA	Alignment	not modelled	99.5	21	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
60	c2dmoA	Alignment	not modelled	99.5	24	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)
61	c2dbmA	Alignment	not modelled	99.5	21	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
62	d1j3ta	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c2ebpA	Alignment	not modelled	99.5	14	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
64	c2dl4A	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
65	c3a98A	Alignment	not modelled	99.5	18	PDB header: signaling protein Chain: A: PDB Molecule: dedicator of cytokinesis protein 2; PDBTitle: crystal structure of the complex of the interacting regions of dock22 and elmo1
66	d1uhfa	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c2jmcA	Alignment	not modelled	99.5	24	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
68	d2v1ra1	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c2cudA	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
70	c2dx1A	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
71	d1utiA	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c2ke9A	Alignment	not modelled	99.5	21	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
73	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.
74	c2lcsA	Alignment	not modelled	99.5	38	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
75	c2dl3A	Alignment	not modelled	99.5	21	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
76	c2bz8B	Alignment	not modelled	99.5	36	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
77	d1wlpb2	Alignment	not modelled	99.5	12	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c2eczA	Alignment	not modelled	99.5	17	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
79	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
80	d1e6ga	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c2ct4A	Alignment	not modelled	99.5	16	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
82	c1wi7A	Alignment	not modelled	99.5	18	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
83	c2csqA	Alignment	not modelled	99.5	18	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
84	c2krnA	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A; PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
85	c1x69A	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A; PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
86	c2l0aA	Alignment	not modelled	99.5	23	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
87	c2e5kA	Alignment	not modelled	99.5	22	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
88	c2drmB	Alignment	not modelled	99.5	28	PDB header: contractile protein Chain: B; PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
89	c2dbkA	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
90	c2xmfA	Alignment	not modelled	99.5	27	PDB header: motor protein Chain: A; PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
91	c2o2oA	Alignment	not modelled	99.5	17	PDB header: protein binding Chain: A; PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
92	d1gcqa	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1e6ha	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	d1ov3a2	Alignment	not modelled	99.5	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c1wxuA	Alignment	not modelled	99.5	25	PDB header: protein transport Chain: A; PDB Molecule: peroxisomal biogenesis factor 13; PDBTitle: solution structure of the sh3 domain of mouse peroxisomal2 biogenesis factor 13
96	c2d8hA	Alignment	not modelled	99.5	14	PDB header: unknown function Chain: A; PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
97	c2dnuA	Alignment	not modelled	99.5	15	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
98	d1spka	Alignment	not modelled	99.5	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	c1x2pA	Alignment	not modelled	99.5	19	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	c2dl7A	Alignment	not modelled	99.5	22	PDB header: structural genomics, unknown function Chain: A; PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
101	c2fpeB	Alignment	not modelled	99.5	20	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
102	c1x6bA	Alignment	not modelled	99.5	13	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
103	c1w70A	Alignment	not modelled	99.5	29	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
						Fold: SH3-like barrel

104	d1nega_	Alignment	not modelled	99.5	23	Superfamily: SH3-domain Family: SH3-domain
105	c1negA_	Alignment	not modelled	99.5	23	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
106	c1z9qA_	Alignment	not modelled	99.5	26	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
107	c2dlfA_	Alignment	not modelled	99.5	24	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
108	c2egaA_	Alignment	not modelled	99.5	17	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
109	c3rnjA_	Alignment	not modelled	99.5	12	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)
110	c2cucA_	Alignment	not modelled	99.5	17	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
111	d1sema_	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	d1l1ja_	Alignment	not modelled	99.5	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1ujya_	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c4esrB_	Alignment	not modelled	99.5	25	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
115	c1z9zA_	Alignment	not modelled	99.5	33	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
116	d1k4us_	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2eqiA_	Alignment	not modelled	99.5	22	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
118	d1ql5a_	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	d1uhca_	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2dilA_	Alignment	not modelled	99.5	25	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