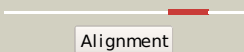

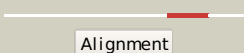

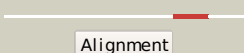

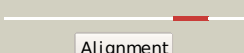

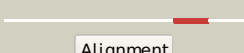

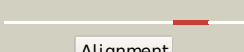

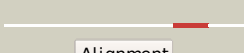

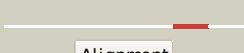







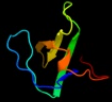
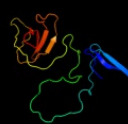










Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q96KQ4
Date	Wed Jun 6 09:44:33 BST 2012
Unique Job ID	0ca6f4d581c1b1e7

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1ycsB_	 Alignment		100.0	81	PDB header: complex (anti-oncogene/ankyrin repeats) Chain: B: PDB Molecule: 53bp2; PDBTitle: p53-53bp2 complex
2	c2vgeA_	 Alignment		99.9	55	PDB header: transcription Chain: A: PDB Molecule: rela-associated inhibitor; PDBTitle: crystal structure of the c-terminal region of human iaspp
3	c1mv3A_	 Alignment		99.7	29	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
4	d1ycsb2	 Alignment		99.7	73	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
5	c1griA_	 Alignment		99.6	22	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
6	c2dx1A_	 Alignment		99.6	40	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
7	c3nmzD_	 Alignment		99.6	36	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
8	c2rqrA_	 Alignment		99.6	14	PDB header: protein binding Chain: A: PDB Molecule: engulfment and cell motility protein 1, linker, dedicator PDBTitle: the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
9	d1l1ja_	 Alignment		99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
10	c2lqnA_	 Alignment		99.5	12	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
11	d1ng2a2	 Alignment		99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

12	c1wyxA	Alignment		99.5	37	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
13	c2eyzA	Alignment		99.5	9	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
14	d1ug1a	Alignment		99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
15	c2dlpA	Alignment		99.5	30	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
16	c1x43A	Alignment		99.5	29	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)
17	c2kymA	Alignment		99.5	32	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
18	c2l3sA	Alignment		99.5	9	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
19	c2dl7A	Alignment		99.5	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
20	c2e5kA	Alignment		99.5	25	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
21	c1x2qA	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: A: PDB Molecule: signal transducing adaptor molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
22	c2dnuA	Alignment	not modelled	99.5	25	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
23	c2nwmA	Alignment	not modelled	99.5	28	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
24	c2ekhA	Alignment	not modelled	99.5	25	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	c2d8hA	Alignment	not modelled	99.5	27	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
26	c2jxbA	Alignment	not modelled	99.5	27	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
27	c2yunA	Alignment	not modelled	99.5	29	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
28	c2creA	Alignment	not modelled	99.5	38	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2 human cdna

29	d1uffa_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
30	c2ysqA_	Alignment	not modelled	99.5	34	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
31	c2yupA_	Alignment	not modelled	99.5	25	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
32	c2xmfA_	Alignment	not modelled	99.5	35	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
33	c2cubA_	Alignment	not modelled	99.5	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
34	c2yt6A_	Alignment	not modelled	99.5	24	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
35	d1ov3a2	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
36	c1w70A_	Alignment	not modelled	99.5	24	PDB header: sh3 domain Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: sh3 domain of p40phox complexed with c-terminal polypoline2 region of p47phox
37	d1j3ta_	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c2uwqA_	Alignment		99.5	67	PDB header: apoptosis Chain: A: PDB Molecule: apoptosis-stimulating of p53 protein 2; PDBTitle: solution structure of aspp2 n-terminus
39	d1gbra_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	c2yuoA_	Alignment	not modelled	99.5	29	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	c2l0aA_	Alignment	not modelled	99.5	30	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
42	c2dl8A_	Alignment	not modelled	99.5	28	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
43	d1uj0a_	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c1ng2A_	Alignment	not modelled	99.5	21	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
45	c2djgA_	Alignment	not modelled	99.4	24	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
46	c2oi3A_	Alignment	not modelled	99.4	37	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)
47	c1x2pA_	Alignment	not modelled	99.4	30	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
48	c2ebpA_	Alignment	not modelled	99.4	20	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
49	c2df6A_	Alignment	not modelled	99.4	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
50	c2csqA_	Alignment	not modelled	99.4	29	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
51	c2dila_	Alignment	not modelled	99.4	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
52	d1jqqa_	Alignment	not modelled	99.4	31	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
53	c2i0nA	Alignment	not modelled	99.4	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
54	c1wxuA	Alignment	not modelled	99.4	26	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
55	c2k6dA	Alignment	not modelled	99.4	37	PDB header: sh3 domain/ubiquitin Chain: A: PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
56	c2eyyA	Alignment	not modelled	99.4	25	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
57	c1zlmA	Alignment	not modelled	99.4	35	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
58	c1x6gA	Alignment	not modelled	99.4	21	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.
59	d1uhca	Alignment	not modelled	99.4	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2ed0A	Alignment	not modelled	99.4	24	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)
61	c2d1xD	Alignment	not modelled	99.4	36	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
62	d1utia	Alignment	not modelled	99.4	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c2dl3A	Alignment	not modelled	99.4	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
64	c2dbmA	Alignment	not modelled	99.4	29	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
65	c2dm1A	Alignment	not modelled	99.4	28	PDB header: signaling protein Chain: A: PDB Molecule: protein vav-2; PDBTitle: solution structure of the second sh3 domain of human2 protein vav-2
66	c1ov3A	Alignment	not modelled	99.4	24	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
67	d1ujya	Alignment	not modelled	99.4	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	d1udla	Alignment	not modelled	99.4	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c1wx6A	Alignment	not modelled	99.4	20	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
70	c2epdA	Alignment	not modelled	99.4	26	PDB header: protein binding Chain: A: PDB Molecule: rho gtpase-activating protein 4; PDBTitle: solution structure of sh3 domain in rho-gtpase-activating2 protein 4
71	c3qwyA	Alignment	not modelled	99.4	25	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
72	d1oeba	Alignment	not modelled	99.4	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2rqvA	Alignment	not modelled	99.4	32	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
74	d1k4us	Alignment	not modelled	99.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c2azsA	Alignment	not modelled	99.4	37	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)
76	c2a28D	Alignment	not modelled	99.4	33	PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein; PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
77	c2egaA	Alignment	not modelled	99.4	20	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

78	dluuea_	Alignment	not modelled	99.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	c3qwxX_	Alignment	not modelled	99.4	20	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
80	c3rnjA_	Alignment	not modelled	99.4	28	PDB header: protein binding Chain: A: PDB Molecule: brain-specific angiogenesis inhibitor 1-associated protein PDBTitle: crystal structure of the sh3 domain from irsp53 (baiap2)
81	dlgcqa_	Alignment	not modelled	99.4	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2enmA_	Alignment	not modelled	99.4	18	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
83	c2egeA_	Alignment	not modelled	99.4	23	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
84	clzx6A_	Alignment	not modelled	99.4	27	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
85	c2dl4A_	Alignment	not modelled	99.4	25	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
86	c2j06B_	Alignment	not modelled	99.4	26	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
87	dlu5sa1	Alignment	not modelled	99.4	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	dlk9aa1	Alignment	not modelled	99.4	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c2k2mA_	Alignment	not modelled	99.4	34	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
90	c2csiA_	Alignment	not modelled	99.4	30	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
91	dlnl1a_	Alignment	not modelled	99.4	35	Fold: beta-hairpin-alpha-hairpin repeat Superfamily: Ankyrin repeat Family: Ankyrin repeat
92	clnl1A_	Alignment	not modelled	99.4	35	PDB header: structural protein Chain: A: PDB Molecule: ankyrin; PDBTitle: d34 region of human ankyrin-r and linker
93	c2pqhA_	Alignment	not modelled	99.4	32	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
94	dlsema_	Alignment	not modelled	99.4	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	clnegA_	Alignment	not modelled	99.4	32	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
96	dlnega_	Alignment	not modelled	99.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2fpeB_	Alignment	not modelled	99.4	17	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
98	c2cudA_	Alignment	not modelled	99.4	28	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adopter protein (slap)
99	c2jteA_	Alignment	not modelled	99.4	29	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
100	c2dmoA_	Alignment	not modelled	99.4	19	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)
101	clu3oA_	Alignment	not modelled	99.4	35	PDB header: signaling protein Chain: A: PDB Molecule: huntingtin-associated protein-interacting PDBTitle: solution structure of rat kalirin n-terminal sh3 domain
102	c2ed1A_	Alignment	not modelled	99.4	31	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
103	clx69A_	Alignment	not modelled	99.4	32	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin

104	c2eyxA	Alignment	not modelled	99.4	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
105	c1wi7A	Alignment	not modelled	99.4	26	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
106	d1pwta	Alignment	not modelled	99.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
107	d1uhfa	Alignment	not modelled	99.4	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2drmB	Alignment	not modelled	99.4	39	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin iB; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
109	d1loota	Alignment	not modelled	99.4	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2kbtA	Alignment	not modelled	99.4	36	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
111	c2ct4A	Alignment	not modelled	99.4	23	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
112	d2v1ra1	Alignment	not modelled	99.4	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1e6ga	Alignment	not modelled	99.4	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d1wlpb2	Alignment	not modelled	99.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2dbkA	Alignment	not modelled	99.4	14	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
116	c2lcsA	Alignment	not modelled	99.4	30	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
117	c1x6bA	Alignment	not modelled	99.4	18	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine exchange factor (gef) 16; PDBTitle: solution structures of the sh3 domain of human rho guanine2 exchange factor (gef) 16
118	d1wiea	Alignment	not modelled	99.4	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c1wxtA	Alignment	not modelled	99.4	29	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
120	c2js0A	Alignment	not modelled	99.4	26	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck