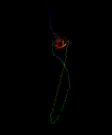
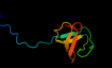
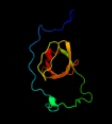
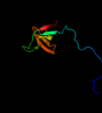


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P15891
Date	Mon Jul 2 19:13:22 BST 2012
Unique Job ID	a839774b4009a372

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlhqz1_	 Alignment		100.0	100	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
2	dludma_	 Alignment		100.0	16	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
3	dlwnja_	 Alignment		100.0	17	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
4	c1x67A_	 Alignment		100.0	20	PDB header: protein binding Chain: A: PDB Molecule: drebrin-like protein; PDBTitle: solution structure of the cofilin homology domain of hip-552 (drebrin-like protein)
5	dlt3ya1	 Alignment		99.9	14	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
6	c1mv3A_	 Alignment		99.9	15	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
7	dlcfya_	 Alignment		99.8	21	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
8	clgriA_	 Alignment		99.8	32	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
9	dlcnua_	 Alignment		99.8	24	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
10	c2kd5A_	 Alignment		99.8	24	PDB header: hormone Chain: A: PDB Molecule: actin severing and dynamics regulatory protein; PDBTitle: solution structure of adf/cofilin (Idcof) from Leishmania donovani
11	c2rqrA_	 Alignment		99.8	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

12	c2kymA_	Alignment		99.7	23	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
13	c2lqnA_	Alignment		99.7	17	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
14	d1ng2a2	Alignment		99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
15	c2oi3A_	Alignment		99.7	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)
16	c2yupA_	Alignment		99.7	26	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
17	c3nmzD_	Alignment		99.7	29	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
18	c2dx1A_	Alignment		99.7	29	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
19	c2l3sA_	Alignment		99.7	14	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
20	d1jo8a_	Alignment		99.7	100	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c2eyzA_	Alignment	not modelled	99.7	16	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
22	c2ebpA_	Alignment	not modelled	99.7	22	PDB header: cell adhesion Chain: A: PDB Molecule: sam and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
23	c2yt6A_	Alignment	not modelled	99.7	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
24	c2rqvA_	Alignment	not modelled	99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
25	c3rnjA_	Alignment	not modelled	99.7	27	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)
26	c2i2qA_	Alignment	not modelled	99.7	22	PDB header: actin-binding protein Chain: A: PDB Molecule: cofilin; PDBTitle: fission yeast cofilin
27	c2xfaA_	Alignment	not modelled	99.7	18	PDB header: protein binding Chain: A: PDB Molecule: actin depolymerization factor 2; PDBTitle: crystal structure of plasmodium berghei actin2 depolymerization factor 2
28	c2ysqA_	Alignment	not modelled	99.7	31	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 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
29	c2e5kA	Alignment	not modelled	99.7	22	
30	d1phta	Alignment	not modelled	99.7	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
31	c2djgA	Alignment	not modelled	99.7	27	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
32	c2ed1A	Alignment	not modelled	99.7	31	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
33	d1f7sa	Alignment	not modelled	99.7	16	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
34	c1x2qA	Alignment	not modelled	99.7	36	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
35	c2cubA	Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
36	c2dbkA	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
37	c3qwxX	Alignment	not modelled	99.7	17	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
38	d1wiea	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
39	c1x43A	Alignment	not modelled	99.7	28	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)
40	c2jxbA	Alignment	not modelled	99.7	21	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	d1spka	Alignment	not modelled	99.7	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2cudA	Alignment	not modelled	99.7	15	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)
43	c1x6gA	Alignment	not modelled	99.7	14	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.
44	c3qwyA	Alignment	not modelled	99.7	21	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
45	c2d1xD	Alignment	not modelled	99.7	52	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
46	c1x6bA	Alignment	not modelled	99.7	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
47	c2kt1A	Alignment	not modelled	99.7	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
48	c2ke9A	Alignment	not modelled	99.7	25	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
49	c2ct3A	Alignment	not modelled	99.7	27	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
50	d1v6fa	Alignment	not modelled	99.7	12	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
51	c2fpeB	Alignment	not modelled	99.7	18	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
52	c1zukA	Alignment	not modelled	99.7	26	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti 1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17

53	c2enmA	Alignment	not modelled	99.7	29	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
54	c2ed0A	Alignment	not modelled	99.7	32	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)
55	c2bzxA	Alignment	not modelled	99.7	28	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
56	c1yn8E	Alignment	not modelled	99.7	31	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
57	c2eyxA	Alignment	not modelled	99.7	22	PDB header: signaling protein Chain: A: PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
58	c2nwmA	Alignment	not modelled	99.7	35	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
59	c2dl3A	Alignment	not modelled	99.7	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
60	d1udla	Alignment	not modelled	99.7	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	c2dl8A	Alignment	not modelled	99.7	30	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
62	c2yunA	Alignment	not modelled	99.7	32	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
63	c2yuoA	Alignment	not modelled	99.7	34	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
64	c2dbmA	Alignment	not modelled	99.7	34	PDB header: transferase, signaling protein Chain: A: PDB Molecule: sh3-containing grb2-like protein 2; PDBTitle: solution structures of the sh3 domain of human sh3-2 containing grb2-like protein 2
65	d1ov3a2	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2j06B	Alignment	not modelled	99.7	21	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
67	c2ct4A	Alignment	not modelled	99.7	18	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
68	c2xmfA	Alignment	not modelled	99.7	44	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
69	c2i0nA	Alignment	not modelled	99.7	26	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
70	c2ekhA	Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structures of the sh3 domain of human kiaa0418
71	c2eyyA	Alignment	not modelled	99.6	18	PDB header: signaling protein Chain: A: PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
72	d1ue9a	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	c2eczA	Alignment	not modelled	99.6	22	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
74	c2epdA	Alignment	not modelled	99.6	31	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
75	c2eqiA	Alignment	not modelled	99.6	32	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
76	c2lcsA	Alignment	not modelled	99.6	32	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
77	c1w70A	Alignment	not modelled	99.6	28	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
78	c2bz8B	Alignment	not modelled	99.6	41	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

79	d2v1ra1	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	d1uffa_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c4esrB_	Alignment	not modelled	99.6	36	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
82	c2d8hA_	Alignment	not modelled	99.6	23	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
83	d1uuea_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	d1q8ga_	Alignment	not modelled	99.6	21	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
85	c1wi7A_	Alignment	not modelled	99.6	27	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
86	c2df6A_	Alignment	not modelled	99.6	29	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
87	c2csqA_	Alignment	not modelled	99.6	26	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
88	c2l0aA_	Alignment	not modelled	99.6	44	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
89	c2dl4A_	Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
90	c1x69A_	Alignment	not modelled	99.6	44	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
91	d1oota_	Alignment	not modelled	99.6	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2dmoA_	Alignment	not modelled	99.6	18	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)
93	d1bb9a_	Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c1bb9A_	Alignment	not modelled	99.6	28	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
95	c2drmB_	Alignment	not modelled	99.6	43	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
96	d1utia_	Alignment	not modelled	99.6	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	d1pwta_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
98	c2l72A_	Alignment	not modelled	99.6	21	PDB header: protein binding Chain: A: PDB Molecule: actin depolymerizing factor, putative; PDBTitle: solution structure and dynamics of adf from toxoplasma gondii (tgadf)
99	d1gria2	Alignment	not modelled	99.6	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	d1k4us_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2egeA_	Alignment	not modelled	99.6	21	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
102	c2dlpA_	Alignment	not modelled	99.6	20	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
103	d1jqqa_	Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c2dl5A_	Alignment	not modelled	99.6	23	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein

105	dlgcqa_	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	clwyxA_	Alignment	not modelled	99.6	36	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
107	dlujya_	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	dluj0a_	Alignment	not modelled	99.6	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	dlwfsa_	Alignment	not modelled	99.6	14	Fold: Gelsolin-like Superfamily: Actin depolymerizing proteins Family: Cofilin-like
110	clz9zA_	Alignment	not modelled	99.6	29	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
111	dlarka_	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	dlsema_	Alignment	not modelled	99.6	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c2azsA_	Alignment	not modelled	99.6	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)
114	c2o2oA_	Alignment	not modelled	99.6	27	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
115	dluhfa_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	dlilja_	Alignment	not modelled	99.6	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	dlj3ta_	Alignment	not modelled	99.6	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	dlwlpb2	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2js0A_	Alignment	not modelled	99.6	37	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
120	clwx6A_	Alignment	not modelled	99.6	29	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