




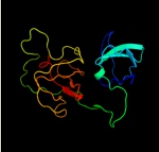

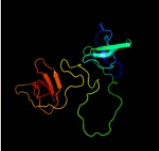
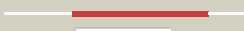




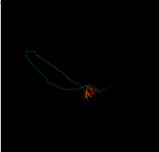

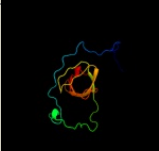



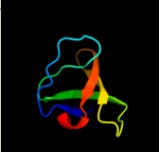

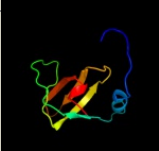


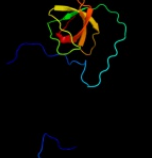
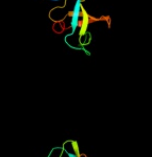
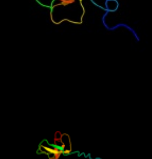
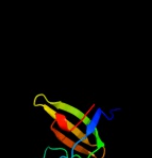
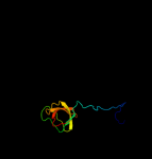
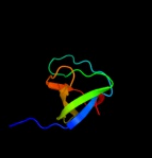
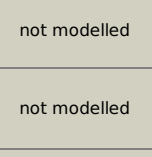


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	O62482
Date	Fri May 3 21:35:38 BST 2013
Unique Job ID	907f535300166207

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	30	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2lqnA_	 Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c2l3sA_	 Alignment		99.9	19	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
4	c2eyzA_	 Alignment		99.9	17	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1ov3A_	 Alignment		99.9	31	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
6	c1ng2A_	 Alignment		99.9	18	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
7	c1mv3A_	 Alignment		99.9	14	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
8	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
9	d1ng2a2	 Alignment		99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
10	c2dx1A_	 Alignment		99.7	23	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
11	c3nmzD_	 Alignment		99.7	19	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef

12	c2kymA	Alignment		99.7	24	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from <i>I.elongisporus</i> in2 complex with ste20p peptide
13	c2yt6A	Alignment		99.7	17	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
14	c2jxbA	Alignment		99.7	18	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
15	c1x6gA	Alignment		99.7	11	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.
16	c2e5kA	Alignment		99.7	36	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
17	c2eyyA	Alignment		99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
18	c2fpeB	Alignment		99.7	32	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
19	c2oi3A	Alignment		99.7	23	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)
20	c2ct3A	Alignment		99.7	53	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
21	d1ov3a2	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2cudA	Alignment	not modelled	99.7	18	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adopter protein (slap)
23	c1zukA	Alignment	not modelled	99.7	31	PDB header: contractile protein Chain: A: PDB Molecule: myosin tail region-interacting protein mti1; PDBTitle: yeast bbc1 sh3 domain complexed with a peptide from las17
24	c3rniA	Alignment	not modelled	99.7	25	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)
25	c2lj0A	Alignment	not modelled	99.7	57	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
26	c2yupA	Alignment	not modelled	99.7	24	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
27	d1phta	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
28	d1ue9a	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: signaling protein

29	c1x2qA	Alignment	not modelled	99.6	26	Chain: A: PDB Molecule: signal transducing adapter molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
30	c3qwyA	Alignment	not modelled	99.6	19	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
31	c2j06B	Alignment	not modelled	99.6	30	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
32	d1bb9a	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
33	c1bb9A	Alignment	not modelled	99.6	32	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
34	c2ysqA	Alignment	not modelled	99.6	21	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
35	c3qwxX	Alignment	not modelled	99.6	18	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
36	c2rqvA	Alignment	not modelled	99.6	30	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
37	c1x6bA	Alignment	not modelled	99.6	23	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
38	c1zlmA	Alignment	not modelled	99.6	36	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
39	c2cubA	Alignment	not modelled	99.6	30	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
40	c2l0aA	Alignment	not modelled	99.6	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
41	c2xmfA	Alignment	not modelled	99.6	42	PDB header: :motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
42	d1jo8a	Alignment	not modelled	99.6	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	d1k4us	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c2kt1A	Alignment	not modelled	99.6	33	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
45	c2yunA	Alignment	not modelled	99.6	36	PDB header: :protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
46	c4afzC	Alignment	not modelled	99.6	27	PDB header: :hydrolase/de novo protein Chain: C: PDB Molecule: fynomer; PDBTitle: human chymase - fynomer complex
47	c2cucA	Alignment	not modelled	99.6	37	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
48	c2lcsA	Alignment	not modelled	99.6	25	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
49	c1yn8E	Alignment	not modelled	99.6	27	PDB header: :unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
50	d1wiea	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	c2ed0A	Alignment	not modelled	99.6	38	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)
52	d1gl5a	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	c2df6A	Alignment	not modelled	99.6	37	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
54	c2dlpA	Alignment	not modelled	99.6	21	PDB header: :structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein

55	c2ekhA	Alignment	not modelled	99.6	23	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
56	d1gcqa	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	c2d8jA	Alignment	not modelled	99.6	23	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
58	c1zuyB	Alignment	not modelled	99.6	30	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
59	d1uffa	Alignment	not modelled	99.6	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c2ed1A	Alignment	not modelled	99.6	23	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
61	c2egcA	Alignment	not modelled	99.6	19	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
62	c2yuoA	Alignment	not modelled	99.6	27	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
63	d1oeba	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
64	c2djqa	Alignment	not modelled	99.6	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
65	d1sema	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2nwmA	Alignment	not modelled	99.6	36	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
67	c2epdA	Alignment	not modelled	99.6	21	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
68	d1spka	Alignment	not modelled	99.6	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	c2i0nA	Alignment	not modelled	99.6	24	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	c4glmD	Alignment	not modelled	99.6	36	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
71	c2dmoA	Alignment	not modelled	99.6	29	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)
72	c2dl4A	Alignment	not modelled	99.6	29	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
73	c1x43A	Alignment	not modelled	99.6	17	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)
74	c3cqtA	Alignment	not modelled	99.6	26	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
75	d1uj0a	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	c2ebpA	Alignment	not modelled	99.6	27	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
77	d1uuea	Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c4esrB	Alignment	not modelled	99.6	36	PDB header: protein binding Chain: B: PDB Molecule: jouberein; PDBTitle: molecular and structural characterization of the sh3 domain of ahi-12 in regulation of cellular resistance of bcr-abl+ chronic myeloid3 leukemia cells to tyrosine kinase inhibitors
79	c2rf0D	Alignment	not modelled	99.6	36	PDB header: transferase Chain: D: PDB Molecule: mitogen-activated protein kinase kinase kinase 10; PDBTitle: crystal structure of human mixed lineage kinase map3k10 sh3 domain
80	d1pwta	Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
81	d1efna_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c1zx6A_	Alignment	not modelled	99.6	40	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
83	c2eyxA_	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
84	c2enmA_	Alignment	not modelled	99.6	22	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
85	c1z9zA_	Alignment	not modelled	99.6	21	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
86	c2dnuA_	Alignment	not modelled	99.6	31	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
87	c2egeA_	Alignment	not modelled	99.6	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein kiaa1666; PDBTitle: solution structure of the third sh3 domain from kiaa1666 protein
88	c2bz8B_	Alignment	not modelled	99.6	37	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
89	d1utia_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	c2d1xD_	Alignment	not modelled	99.6	41	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
91	d1gria2	Alignment	not modelled	99.6	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
92	c2csqA_	Alignment	not modelled	99.6	21	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
93	c1x2pA_	Alignment	not modelled	99.6	25	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
94	d1wlpb2	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c2d8hA_	Alignment	not modelled	99.6	24	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
96	d1fmka1	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2drmb_	Alignment	not modelled	99.6	38	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
98	d1arka_	Alignment	not modelled	99.6	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	d1u06a1	Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	d1ujya_	Alignment	not modelled	99.6	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	d1shfa_	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2dilA_	Alignment	not modelled	99.6	39	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
103	c2dl8A_	Alignment	not modelled	99.6	19	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
104	d2v1ra1	Alignment	not modelled	99.6	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	d2iima1	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c2eqiA_	Alignment	not modelled	99.6	31	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 PDB header: structural protein

107	c2pqhA_	Alignment	not modelled	99.6	26	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
108	c2ke9A_	Alignment	not modelled	99.6	30	PDB header: protein binding Chain: A: PDB Molecule: casikin-2; PDBTitle: nmr solution structure of the casikin sh3 domain
109	d1e6ga_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2dl3A_	Alignment	not modelled	99.6	29	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
111	c2jteA_	Alignment	not modelled	99.6	27	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
112	c2dbmA_	Alignment	not modelled	99.6	36	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
113	c2ct4A_	Alignment	not modelled	99.6	13	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strutcure of the sh3 domain of the cdc42-2 interacting protein 4
114	d1udla_	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	d1ckaa_	Alignment	not modelled	99.6	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
116	c2kgtA_	Alignment	not modelled	99.6	17	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
117	d1y57a1	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	c4iqzA_	Alignment	not modelled	99.6	35	PDB header: protein binding Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 2; PDBTitle: crystal structure of the sh3 domain of human sorbin and sh3 domain-2 containing protein 2
119	c2dl5A_	Alignment	not modelled	99.6	19	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
120	c1negA_	Alignment	not modelled	99.6	26	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