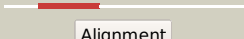

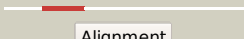
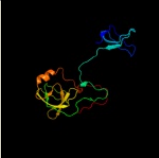
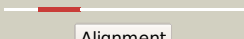


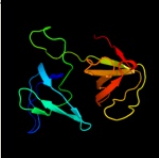

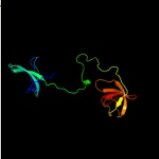



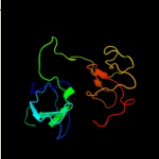







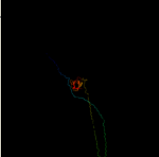







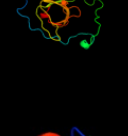



Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	A2AAY5
Date	Tue Jul 30 13:00:07 BST 2013
Unique Job ID	f8c6a4defb6bed86

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dybA	 Alignment		100.0	21	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
2	c1ng2A	 Alignment		100.0	34	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
3	c2lqnA	 Alignment		99.9	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
4	c1griA	 Alignment		99.9	23	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
5	c1ov3A	 Alignment		99.9	42	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
6	c2eyzA	 Alignment		99.9	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
7	c2l3sA	 Alignment		99.9	20	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
8	c2l73A	 Alignment		99.9	21	PDB header: oxidoreductase regulator Chain: A: PDB Molecule: nadph oxidase organizer 1; PDBTitle: structure of the noxo1b px domain
9	d1kq6a	 Alignment		99.9	36	Fold: PX domain Superfamily: PX domain Family: PX domain
10	d1h6ha	 Alignment		99.9	26	Fold: PX domain Superfamily: PX domain Family: PX domain
11	c1mv3A	 Alignment		99.9	12	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

12	d1o7ka_	Alignment		99.9	36	Fold: PX domain Superfamily: PX domain Family: PX domain
13	c2reaA_	Alignment		99.8	32	PDB header: transferase Chain: A: PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: crystal structures of c2alpha-pi3 kinase px-domain domain2 indicate conformational change associated with ligand3 binding.
14	c2iwlX_	Alignment		99.7	28	PDB header: transferase Chain: X: PDB Molecule: phosphatidylinositol-4-phosphate 3-kinase c2 PDBTitle: structure of the px domain of phosphoinositide 3-kinase-2 c2alpha
15	c2egcA_	Alignment		99.7	51	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
16	c2czoA_	Alignment		99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the px domain of bem1p
17	d1udla_	Alignment		99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
18	d1ng2a2	Alignment		99.7	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
19	c2rqrA_	Alignment		99.7	16	PDB header: protein binding Chain: A: PDB Molecule: engulfment and cell motility protein 1, linker, dedicator PDBTitle: the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
20	c2dx1A_	Alignment		99.6	28	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
21	c2wweA_	Alignment	not modelled	99.6	23	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
22	c2yt6A_	Alignment	not modelled	99.6	21	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
23	c2jxbA_	Alignment	not modelled	99.6	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
24	d1ue9a_	Alignment	not modelled	99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2d8jA_	Alignment	not modelled	99.6	26	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
26	d1ov3a2	Alignment	not modelled	99.6	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2dbkA_	Alignment	not modelled	99.6	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
28	c2e5kA_	Alignment	not modelled	99.6	18	PDB header: protein binding Chain: A: PDB Molecule: suppressor of t-cell receptor signaling 1; PDBTitle: solution structure of sh3 domain in suppressor of t-cell2 receptor signaling 1
						Fold: SH3-like barrel

29	d1phta_	Alignment	not modelled	99.5	16	Superfamily: SH3-domain Family: SH3-domain
30	c1zukA_	Alignment	not modelled	99.5	19	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
31	c2ct3A_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
32	c2jmcA_	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
33	c2fpeB_	Alignment	not modelled	99.5	14	PDB header: signaling protein Chain: B: PDB Molecule: c-jun-amino-terminal kinase interacting protein PDBTitle: conserved dimerization of the ib1 src-homology 3 domain
34	c3rniA_	Alignment	not modelled	99.5	24	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)
35	c3nmzD_	Alignment	not modelled	99.5	25	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asef
36	c1zlmA_	Alignment	not modelled	99.5	29	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
37	c2ke9A_	Alignment	not modelled	99.5	19	PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
38	c2kymA_	Alignment	not modelled	99.5	17	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
39	c2cucA_	Alignment	not modelled	99.5	23	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
40	c4afzC_	Alignment	not modelled	99.5	30	PDB header: hydrolase/de novo protein Chain: C: PDB Molecule: fynomer; PDBTitle: human chymase - fynomer complex
41	c2lj0A_	Alignment	not modelled	99.5	25	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
42	c2yupA_	Alignment	not modelled	99.5	23	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
43	d1wlpb2	Alignment	not modelled	99.5	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c2oi3A_	Alignment	not modelled	99.5	28	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)
45	c1x6gA_	Alignment	not modelled	99.5	16	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.
46	d1efna_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	c2rf0D_	Alignment	not modelled	99.5	31	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
48	c2cudA_	Alignment	not modelled	99.5	19	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)
49	c2ebpA_	Alignment	not modelled	99.5	19	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
50	c2kt1A_	Alignment	not modelled	99.5	22	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
51	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
52	c2j06B_	Alignment	not modelled	99.5	23	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
53	c2ed1A_	Alignment	not modelled	99.5	25	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 PDB header: signaling protein

54	c1x2qA	Alignment	not modelled	99.5	22	Chain: A: PDB Molecule: signal transducing adapter molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
55	c2eyxA	Alignment	not modelled	99.5	17	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
56	c1zuyB	Alignment	not modelled	99.5	30	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
57	c2eyyA	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
58	c3cqtA	Alignment	not modelled	99.5	31	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
59	c1x2pA	Alignment	not modelled	99.5	24	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
60	d1gcqc	Alignment	not modelled	99.5	15	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
61	d1shfa	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	c4glmD	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: D: PDB Molecule: dynamamin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
63	c2xmfA	Alignment	not modelled	99.5	24	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
64	c2lcsA	Alignment	not modelled	99.5	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
65	d1spka	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	d1wfwA	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	c1x6bA	Alignment	not modelled	99.5	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
68	c3qwyA	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
69	c2df6A	Alignment	not modelled	99.5	39	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
70	c1yn8E	Alignment	not modelled	99.5	28	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
71	c2dmoA	Alignment	not modelled	99.5	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)
72	d1fmka1	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d1pwta	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2kgtA	Alignment	not modelled	99.5	27	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
75	c2dl4A	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
76	d1uuea	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	d1u06a1	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	d1e6ga	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	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
80	d1uffa	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

81	c2js0A	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
82	c1negA	Alignment	not modelled	99.5	18	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
83	d1nega	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	d1ql5a	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
85	d1k1za	Alignment	not modelled	99.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c2cubA	Alignment	not modelled	99.5	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
87	c2bzxA	Alignment	not modelled	99.5	19	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
88	c2nwmA	Alignment	not modelled	99.5	16	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
89	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
90	c1x27F	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: F: PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of2 p130cas
91	c1bb9A	Alignment	not modelled	99.5	20	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
92	d1bb9a	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1e6ha	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2ed0A	Alignment	not modelled	99.5	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)
95	c1x69A	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
96	d1uj0a	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2ysqA	Alignment	not modelled	99.5	25	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
98	c2egeA	Alignment	not modelled	99.5	16	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
99	d1uhca	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2dl3A	Alignment	not modelled	99.4	15	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
101	d1jo8a	Alignment	not modelled	99.4	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c4esrB	Alignment	not modelled	99.4	23	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
103	d1wiea	Alignment	not modelled	99.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	c3qwxX	Alignment	not modelled	99.4	24	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
105	c2jteA	Alignment	not modelled	99.4	17	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
106	d1gcqa	Alignment	not modelled	99.4	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

107	c2i0nA_	Alignment	not modelled	99.4	13	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
108	d1k4us_	Alignment	not modelled	99.4	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	d1arka_	Alignment	not modelled	99.4	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2d8hA_	Alignment	not modelled	99.4	25	PDB header: unknown function Chain: A: PDB Molecule: sh3y11 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3y11
111	c2pqhA_	Alignment	not modelled	99.4	18	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
112	c2dnuA_	Alignment	not modelled	99.4	41	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
113	d1wlpb1	Alignment	not modelled	99.4	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d1ov3a1	Alignment	not modelled	99.4	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2epdA_	Alignment	not modelled	99.4	28	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
116	d1ujya_	Alignment	not modelled	99.4	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c1x43A_	Alignment	not modelled	99.4	14	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)
118	d1k9aa1	Alignment	not modelled	99.4	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	d1h8ka_	Alignment	not modelled	99.4	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	d2iima1	Alignment	not modelled	99.4	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain