







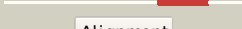
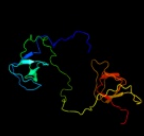
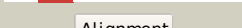
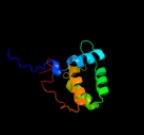
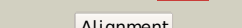

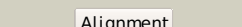

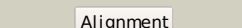

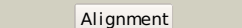

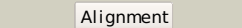



Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q81NU2
Date	Tue Jan 22 11:33:37 GMT 2013
Unique Job ID	be4083f9631c26b4

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		99.9	24	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2khnA_	 Alignment		99.9	50	PDB header: signaling protein Chain: A; PDB Molecule: intersectin-1; PDBTitle: nmr solution structure of the eh 1 domain from human2 intersectin-1 protein. northeast structural genomics3 consortium target hr3646e.
3	c2jq6A_	 Alignment		99.9	39	PDB header: metal binding protein Chain: A; PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of eh-domain of ehd1
4	c3fiaA_	 Alignment		99.9	51	PDB header: protein binding Chain: A; PDB Molecule: intersectin-1; PDBTitle: crystal structure of the eh 1 domain from human intersectin-2 1 protein. northeast structural genomics consortium target3 hr3646e.
5	c2lqnA_	 Alignment		99.9	22	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
6	c2kfhA_	 Alignment		99.9	40	PDB header: protein binding Chain: A; PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of the c-terminal domain of ehd1 with fnyestgpftak
7	c2eyzA_	 Alignment		99.9	23	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
8	c2l3sA_	 Alignment		99.9	23	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
9	d2jxca1	 Alignment		99.9	39	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
10	c1ng2A_	 Alignment		99.9	25	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
11	c2qptA_	 Alignment		99.9	37	PDB header: endocytosis Chain: A; PDB Molecule: eh domain-containing protein-2; PDBTitle: crystal structure of an ehd atpase involved in membrane remodelling

12	c1ov3A_	Alignment		99.8	24	PDB header: oxidoreductase activator Chain: A; PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
13	c2kgrA_	Alignment		99.8	36	PDB header: protein binding Chain: A; PDB Molecule: intersectin-1; PDBTitle: solution structure of protein itsn1 from homo sapiens.2 northeast structural genomics consortium target hr5524a
14	d1iq3a_	Alignment		99.8	39	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
15	d1c07a_	Alignment		99.8	39	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
16	d1fi6a_	Alignment		99.8	42	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
17	c1mv3A_	Alignment		99.8	19	PDB header: endocytosis/exocytosis Chain: A; PDB Molecule: myc box dependent interacting protein 1; PDBTitle: nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
18	d1qjta_	Alignment		99.7	30	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
19	d1ludla_	Alignment		99.7	45	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	d1ng2a2	Alignment		99.6	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	c2dx1A_	Alignment	not modelled	99.6	30	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
22	c3jv3A_	Alignment	not modelled	99.6	45	PDB header: protein binding Chain: A; PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-11
23	c2cubA_	Alignment	not modelled	99.6	29	PDB header: signaling protein Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
24	c2eyvA_	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
25	c1x2qA_	Alignment	not modelled	99.6	37	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
26	c2yunA_	Alignment	not modelled	99.6	32	PDB header: protein transport Chain: A; PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
27	c2ed0A_	Alignment	not modelled	99.5	36	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)
28	d1ov3a2	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: cell adhesion/cell cycle

29	c3nmzD_	Alignment	not modelled	99.5	24	Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
30	c2rqrA_	Alignment	not modelled	99.5	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
31	c2yuoA_	Alignment	not modelled	99.5	31	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
32	c2epdA_	Alignment	not modelled	99.5	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
33	c2dlpA_	Alignment	not modelled	99.5	32	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
34	c2yt6A_	Alignment	not modelled	99.5	29	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	c2df6A_	Alignment	not modelled	99.5	42	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
36	c2ekhA_	Alignment	not modelled	99.5	20	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
37	c2dmoA_	Alignment	not modelled	99.5	23	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)
38	c2dbmA_	Alignment	not modelled	99.5	32	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
39	c4glmD_	Alignment	not modelled	99.5	35	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
40	d1ue9a_	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
41	c2dbkA_	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
42	c2csqA_	Alignment	not modelled	99.5	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
43	c2djgA_	Alignment	not modelled	99.5	29	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: sh3 domain containing ring finger 2; PDBTitle: the solution structure of the first sh3 domain of mouse sh32 domain containing ring finger 2
44	c2ysqA_	Alignment	not modelled	99.5	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
45	c2nwmA_	Alignment	not modelled	99.5	29	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
46	c2kymA_	Alignment	not modelled	99.5	28	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
47	d1oeba_	Alignment	not modelled	99.5	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
48	d1sema_	Alignment	not modelled	99.5	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	d1utia_	Alignment	not modelled	99.5	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c1x69A_	Alignment	not modelled	99.5	38	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
51	d1k4us_	Alignment	not modelled	99.5	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2i0nA_	Alignment	not modelled	99.5	23	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
53	d1ujya_	Alignment	not modelled	99.5	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: motor protein

54	c2xmfA_	Alignment	not modelled	99.5	45	Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
55	c2cudA_	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpoter protein (slap)
56	c2dl3A_	Alignment	not modelled	99.5	26	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
57	c2dilA_	Alignment	not modelled	99.5	33	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
58	c2l0aA_	Alignment	not modelled	99.5	42	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
59	c2egaA_	Alignment	not modelled	99.5	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
60	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
61	c2pqhA_	Alignment	not modelled	99.5	31	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
62	c2ebpA_	Alignment	not modelled	99.5	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
63	c2dl4A_	Alignment	not modelled	99.5	26	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
64	c2yupA_	Alignment	not modelled	99.5	24	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
65	c2jxbA_	Alignment	not modelled	99.5	23	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
66	d1gcqa_	Alignment	not modelled	99.5	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	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
68	c1zx6A_	Alignment	not modelled	99.5	41	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
69	c2e5kA_	Alignment	not modelled	99.5	27	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
70	c2egcA_	Alignment	not modelled	99.5	20	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
71	d1nega_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c1negA_	Alignment	not modelled	99.5	29	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
73	c2oi3A_	Alignment	not modelled	99.5	29	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)
74	c2lj0A_	Alignment	not modelled	99.5	41	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
75	c2dl7A_	Alignment	not modelled	99.5	38	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
76	c2fpeB_	Alignment	not modelled	99.5	24	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
77	c1x6gA_	Alignment	not modelled	99.5	18	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.

78	d1pwta_	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	d1wlpb2	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
80	c1x43A	Alignment	not modelled	99.5	26	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)
81	d1uuea_	Alignment	not modelled	99.5	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2ed1A	Alignment	not modelled	99.5	30	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
83	d1gria2	Alignment	not modelled	99.5	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c3qwyA	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
85	c1x2pA	Alignment	not modelled	99.5	28	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
86	c2eyxA	Alignment	not modelled	99.5	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
87	c1x6bA	Alignment	not modelled	99.5	24	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
88	c1w70A	Alignment	not modelled	99.5	30	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
89	c2ct3A	Alignment	not modelled	99.5	29	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
90	c2bz8B	Alignment	not modelled	99.5	39	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
91	c2drmB	Alignment	not modelled	99.5	47	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
92	d1uhfa_	Alignment	not modelled	99.5	34	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1uj0a_	Alignment	not modelled	99.5	42	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2kbtA	Alignment	not modelled	99.5	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
95	c1zlmA	Alignment	not modelled	99.5	38	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
96	c1zukA	Alignment	not modelled	99.5	37	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
97	c3qwxX	Alignment	not modelled	99.5	20	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
98	c4esrB	Alignment	not modelled	99.5	43	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
99	c2rf0D	Alignment	not modelled	99.5	40	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
100	c2egeA	Alignment	not modelled	99.5	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
101	c1wi7A	Alignment	not modelled	99.5	37	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
102	d1uhca_	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
						PDB header: cell invasion

103	c2d1xD_	Alignment	not modelled	99.5	47	Chain: D; PDB Molecule: cortactin isoform a; PDBTitle: the crystal structure of the cortactin-sh3 domain and amap1-2 peptide complex
104	d1e6ga_	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	d1ug1a_	Alignment	not modelled	99.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	d1u06a1	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
107	c2cucA_	Alignment	not modelled	99.5	35	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
108	c3rnjA_	Alignment	not modelled	99.5	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)
109	c2eqiA_	Alignment	not modelled	99.5	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
110	d1wfwA_	Alignment	not modelled	99.5	14	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c2o2oA_	Alignment	not modelled	99.5	36	PDB header: protein binding Chain: A; PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
112	c1wx6A_	Alignment	not modelled	99.4	30	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
113	d1ov3a1	Alignment	not modelled	99.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2j06B_	Alignment	not modelled	99.4	32	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
115	c2krnA_	Alignment	not modelled	99.4	39	PDB header: signaling protein Chain: A; PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
116	c2k2mA_	Alignment	not modelled	99.4	25	PDB header: signaling protein Chain: A; PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
117	d1ycsb2	Alignment	not modelled	99.4	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
118	d1e6ha_	Alignment	not modelled	99.4	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	d1ng2a1	Alignment	not modelled	99.4	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c1zuyB_	Alignment	not modelled	99.4	28	PDB header: contractile protein Chain: B; PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain