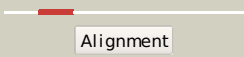
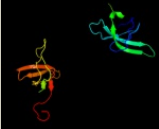
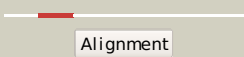
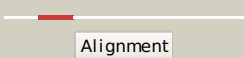
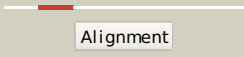
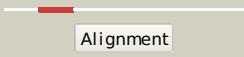
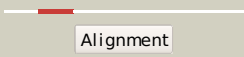
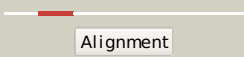
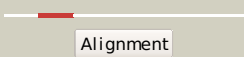
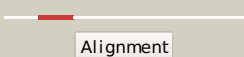
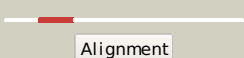
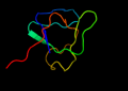


Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	P32790
Date	Mon Jul 2 19:16:44 BST 2012
Unique Job ID	d06943c0b0937d0c

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2lqnA_	 Alignment		100.0	19	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
2	c1griA_	 Alignment		100.0	37	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
3	c1ng2A_	 Alignment		100.0	20	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
4	c2l3sA_	 Alignment		100.0	18	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
5	c2eyzA_	 Alignment		100.0	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
6	c1ov3A_	 Alignment		100.0	23	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
7	c1mv3A_	 Alignment		99.8	24	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	c2dybA_	 Alignment		99.8	19	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
9	d1udla_	 Alignment		99.8	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
10	d1ng2a2	 Alignment		99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
11	c2jmCA_	 Alignment		99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41

12	dlwiea_	Alignment		99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
13	c2rqra_	Alignment		99.7	17	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
14	c2egeA_	Alignment		99.6	20	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
15	c2yt6A_	Alignment		99.6	32	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
16	c2d8jA_	Alignment		99.6	25	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
17	c2csqA_	Alignment		99.6	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
18	c3nmzD_	Alignment		99.6	15	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
19	dlwfa_	Alignment		99.6	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2jxbA_	Alignment		99.6	38	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
21	c3qwyA_	Alignment	not modelled	99.6	16	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
22	dlilja_	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	dluffa_	Alignment	not modelled	99.6	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
24	dlue9a_	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2yunA_	Alignment	not modelled	99.6	24	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
26	c1x6gA_	Alignment	not modelled	99.6	23	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.
27	c1wx6A_	Alignment	not modelled	99.6	25	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
28	c2kymA_	Alignment	not modelled	99.6	22	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 PDB header: signaling protein

29	c2eyyA	Alignment	not modelled	99.6	26	Chain: A: PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
30	c2dmoA	Alignment	not modelled	99.6	30	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)
31	c2oi3A	Alignment	not modelled	99.6	33	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)
32	c2dx1A	Alignment	not modelled	99.6	15	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
33	c1zlmA	Alignment	not modelled	99.6	45	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
34	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
35	d1uhca	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
36	c2egcA	Alignment	not modelled	99.6	24	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
37	c1x2qA	Alignment	not modelled	99.6	29	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
38	c2ekhA	Alignment	not modelled	99.6	24	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
39	c1x43A	Alignment	not modelled	99.6	25	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	c2e5kA	Alignment	not modelled	99.6	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
41	c2ysqA	Alignment	not modelled	99.6	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
42	c2fpeB	Alignment	not modelled	99.6	25	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
43	c2kbtA	Alignment	not modelled	99.6	32	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
44	c2ebpA	Alignment	not modelled	99.6	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
45	d1ug1a	Alignment	not modelled	99.6	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	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
47	c2dl4A	Alignment	not modelled	99.6	32	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
48	c3rnjA	Alignment	not modelled	99.6	23	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)
49	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
50	d1efna	Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	c2ct3A	Alignment	not modelled	99.6	35	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
52	c2ed0A	Alignment	not modelled	99.6	31	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)
53	c3qwxX	Alignment	not modelled	99.6	21	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174

54	c2i0nA	Alignment	not modelled	99.6	18	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
55	c2bzxA	Alignment	not modelled	99.6	18	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
56	c2yupA	Alignment	not modelled	99.6	14	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
57	c2cudA	Alignment	not modelled	99.6	14	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)
58	c2nwmA	Alignment	not modelled	99.6	27	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	c2yuoA	Alignment	not modelled	99.6	22	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
60	c1zukA	Alignment	not modelled	99.6	16	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
61	c2eyxA	Alignment	not modelled	99.6	14	PDB header: signaling protein Chain: A: PDB Molecule: v-crkl sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
62	c2dbmA	Alignment	not modelled	99.6	29	PDB header: transferase, signaling protein Chain: A: PDB Molecule: sh3-containing grb2-like protein 2; PDBTitle: solution structures of the sh3 domain of human sh3-2 containing grb2-like protein 2
63	c2df6A	Alignment	not modelled	99.6	22	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
64	c2l0aA	Alignment	not modelled	99.6	33	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
65	c2a28D	Alignment	not modelled	99.6	30	PDB header: signaling protein Chain: D: PDB Molecule: bzz1 protein; PDBTitle: atomic-resolution crystal structure of the second sh32 domain of yeast bzz1 determined from a pseudomerohedrally3 twinned crystal
66	c1wi7A	Alignment	not modelled	99.6	24	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
67	c3cqtA	Alignment	not modelled	99.6	30	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
68	d1wlpb2	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
69	d1ov3a2	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2cucA	Alignment	not modelled	99.6	28	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
71	d1gcqa	Alignment	not modelled	99.6	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c2cubA	Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
73	c2kt1A	Alignment	not modelled	99.6	28	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
74	d1ujya	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c2djqa	Alignment	not modelled	99.6	20	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
76	c2xmfa	Alignment	not modelled	99.6	28	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
77	d1shfa	Alignment	not modelled	99.6	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
78	c2dbkA	Alignment	not modelled	99.6	13	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2

						protein
79	c1x69A_	Alignment	not modelled	99.6	27	PDB header: signaling protein Chain: A: PDB Molecule: contactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
80	c2egaA_	Alignment	not modelled	99.6	29	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
81	c2bz8B_	Alignment	not modelled	99.6	25	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
82	c2csiA_	Alignment	not modelled	99.6	30	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: rim binding protein 2; PDBTitle: solution structure of the third sh3 domain of human rim-2 binding protein 2
83	d1spka_	Alignment	not modelled	99.6	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1wyxA_	Alignment	not modelled	99.6	22	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
85	d1uuea_	Alignment	not modelled	99.6	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	c2kgtA_	Alignment	not modelled	99.6	24	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
87	c1x6bA_	Alignment	not modelled	99.6	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
88	c2k6dA_	Alignment	not modelled	99.6	28	PDB header: sh3 domain/ubiquitin Chain: A: PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
89	c2dl3A_	Alignment	not modelled	99.5	33	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
90	d1uhfa_	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	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
92	d1pwta_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	d1phta_	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	d1k4us_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	c1x2pA_	Alignment	not modelled	99.5	16	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
96	d1e6ga_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c2ct4A_	Alignment	not modelled	99.5	28	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution structure of the sh3 domain of the cdc42-2 interacting protein 4
98	d1ycsb2_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	d1fmka1_	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
100	c2epdA_	Alignment	not modelled	99.5	34	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
101	d1arka_	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
102	c2rqvA_	Alignment	not modelled	99.5	17	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
103	d1u5sa1_	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
104	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
105	d1gria2	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	c2dila	Alignment	not modelled	99.5	36	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
107	c2dl5a	Alignment	not modelled	99.5	31	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
108	d1opka1	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	c1nega	Alignment	not modelled	99.5	31	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
110	d1nega	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
111	c2j06B	Alignment	not modelled	99.5	29	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
112	d1gl5a	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d2v1ra1	Alignment	not modelled	99.5	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	d2fo0a1	Alignment	not modelled	99.5	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c2j6kE	Alignment	not modelled	99.5	20	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
116	c2creA	Alignment	not modelled	99.5	26	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hef-like protein; PDBTitle: solution structure of rsgi ruh-036, an sh3 domain from2 human cdna
117	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
118	d1bb9a	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2jteA	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
120	c2d1xD	Alignment	not modelled	99.5	30	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