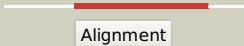

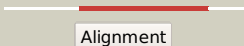

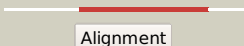







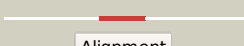











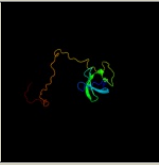
Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	Q814E2
Date	Sat Apr 13 13:13:52 BST 2013
Unique Job ID	3b15295aa727664b

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2rajA_	 Alignment		100.0	34	PDB header: structural protein Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: so4 bound px-bar membrane remodeling unit of sorting nexin 9
2	c3dyuB_	 Alignment		100.0	34	PDB header: transport protein Chain: B: PDB Molecule: sorting nexin-9; PDBTitle: crystal structure of snx9px-bar (230-595), h32
3	c4akvB_	 Alignment		100.0	37	PDB header: transport protein Chain: B: PDB Molecule: sorting nexin-33; PDBTitle: crystal structure of human sorting nexin 33 (snx33)
4	c3hpbA_	 Alignment		99.9	23	PDB header: protein transport Chain: A: PDB Molecule: snx5 protein; PDBTitle: crystal structure of snx5-px domain in p212121 space group
5	c3iq2B_	 Alignment		99.8	31	PDB header: protein transport Chain: B: PDB Molecule: sorting nexin-7; PDBTitle: human sorting nexin 7, phox homology (px) domain
6	c2cskA_	 Alignment		99.8	27	PDB header: protein transport Chain: A: PDB Molecule: sorting nexin 12; PDBTitle: solution structure of px domain from human snx12
7	c2i4kA_	 Alignment		99.8	30	PDB header: protein transport Chain: A: PDB Molecule: sorting nexin-1; PDBTitle: solution structure of the px domain of sorting nexin 1
8	c1ocuA_	 Alignment		99.8	24	PDB header: sorting protein Chain: A: PDB Molecule: sorting nexin; PDBTitle: crystal structure of the yeast px-domain protein grd19p2 (sorting nexin 3) complexed to 3 phosphatidylinositol-3-phospahte.
9	c1xteA_	 Alignment		99.7	18	PDB header: transferase Chain: A: PDB Molecule: serine/threonine-protein kinase sgk3; PDBTitle: crystal structure of cisk-px domain
10	d1xtea_	 Alignment		99.7	18	Fold: PX domain Superfamily: PX domain Family: PX domain
11	d1ocsa_	 Alignment		99.7	24	Fold: PX domain Superfamily: PX domain Family: PX domain

12	c1mv3A_	 Alignment		99.7	36	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
13	c3fogA_	 Alignment		99.7	23	PDB header: protein transport Chain: A; PDB Molecule: sorting nexin-17; PDBTitle: crystal structure of the px domain of sorting nexin-172 (snx17)
14	c2v14A_	 Alignment		99.6	18	PDB header: transport protein Chain: A; PDB Molecule: kinesin-like motor protein c20orf23; PDBTitle: kinesin 16b phox-homology domain (kif16b)
15	c2dybA_	 Alignment		99.6	17	PDB header: oxidoreductase Chain: A; PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
16	c2eyyA_	 Alignment		99.6	21	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
17	c1griA_	 Alignment		99.6	36	PDB header: signal transduction adaptor Chain: A; PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
18	c2enmA_	 Alignment		99.6	35	PDB header: endocytosis Chain: A; PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
19	d1uffa_	 Alignment		99.6	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	d1kmda_	 Alignment		99.6	19	Fold: PX domain Superfamily: PX domain Family: PX domain
21	c3p0cA_	 Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A; PDB Molecule: nischarin; PDBTitle: nischarin px-domain
22	c3jv3A_	 Alignment		99.6	35	PDB header: protein binding Chain: A; PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1
23	c3rniA_	 Alignment	not modelled	99.6	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)
24	c2dipA_	 Alignment	not modelled	99.6	25	PDB header: structural protein Chain: A; PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
25	c2dx1A_	 Alignment	not modelled	99.6	29	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
26	d1spka_	 Alignment	not modelled	99.6	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2e5kA_	 Alignment	not modelled	99.6	25	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
						PDB header: structural genomics, unknown function

28	c1wx6A	Alignment	not modelled	99.6	27	Chain: A; PDB Molecule: cytoplasmic protein nck2; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck2
29	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
30	c2yunA	Alignment	not modelled	99.6	26	PDB header: protein transport Chain: A; PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
31	d1u5sa1	Alignment	not modelled	99.6	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	c2kymA	Alignment	not modelled	99.6	27	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
33	c2dvjA	Alignment	not modelled	99.6	17	PDB header: signaling protein Chain: A; PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
34	c1x2qA	Alignment	not modelled	99.6	28	PDB header: signaling protein Chain: A; PDB Molecule: signal transducing adapter molecule 2; PDBTitle: solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
35	d1ng2a2	Alignment	not modelled	99.6	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
36	c3qwxX	Alignment	not modelled	99.5	30	PDB header: signaling protein Chain: X; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
37	c2lqnA	Alignment		99.5	23	PDB header: signaling protein Chain: A; PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
38	d1h6ha	Alignment	not modelled	99.5	14	Fold: PX domain Superfamily: PX domain Family: PX domain
39	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
40	c1x6qA	Alignment	not modelled	99.5	19	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.
41	c2reaA	Alignment	not modelled	99.5	20	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.
42	d1wiea	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	d1phta	Alignment	not modelled	99.5	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c2csqA	Alignment	not modelled	99.5	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
45	c3qwyA	Alignment	not modelled	99.5	30	PDB header: signaling protein Chain: A; PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
46	d1jqqa	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	c3nmzD	Alignment	not modelled	99.5	28	PDB header: cell adhesion/cell cycle Chain: D; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
48	d2v1ra1	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	c2ct4A	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: A; PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strcture of the sh3 domain of the cdc42-2 interacting protein 4
50	d1kq6a	Alignment	not modelled	99.5	14	Fold: PX domain Superfamily: PX domain Family: PX domain
51	c2l3sA	Alignment	not modelled	99.5	18	PDB header: structural protein Chain: A; PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
52	c2i0nA	Alignment	not modelled	99.5	20	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 PDB header: structural protein

53	c2pqhA	Alignment	not modelled	99.5	27	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
54	c2ke9A	Alignment	not modelled	99.5	28	PDB header: protein binding Chain: A; PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
55	d1qlya	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
56	c2x3wD	Alignment	not modelled	99.5	43	PDB header: endocytosis Chain: D; PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: structure of mouse syndapin i (crystal form 2)
57	c2jxbA	Alignment	not modelled	99.5	29	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
58	c2nwmA	Alignment	not modelled	99.5	33	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	c1z9zA	Alignment	not modelled	99.5	26	PDB header: structural protein Chain: A; PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
60	c2cubA	Alignment	not modelled	99.5	25	PDB header: signaling protein Chain: A; PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
61	c4esrB	Alignment	not modelled	99.5	27	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
62	c2ed0A	Alignment	not modelled	99.5	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)
63	c2rf0D	Alignment	not modelled	99.5	32	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
64	d1gbra	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c2kbtA	Alignment	not modelled	99.5	29	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
66	c2dl5A	Alignment	not modelled	99.5	33	PDB header: structural genomics, unknown function Chain: A; PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
67	c2ekhA	Alignment	not modelled	99.5	21	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
68	c4iqzA	Alignment	not modelled	99.5	38	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
69	d1ugva	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	c2bz8B	Alignment	not modelled	99.5	45	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
71	d1ql5a	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c2ed1A	Alignment	not modelled	99.5	36	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
73	d1ue9a	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	d1gcqa	Alignment	not modelled	99.5	38	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c2xmfA	Alignment	not modelled	99.5	46	PDB header: motor protein Chain: A; PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
76	c1x43A	Alignment	not modelled	99.5	23	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)
77	c2lj0A	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: A; PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
78	d1l1ja	Alignment	not modelled	99.5	23	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
79	c2oi3A	Alignment	not modelled	99.5	24	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)
80	d1cska	Alignment	not modelled	99.5	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	c1ng2A	Alignment		99.5	13	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
82	c2ebpA	Alignment	not modelled	99.5	22	PDB header: cell adhesion Chain: A: PDB Molecule: sam and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
83	d1gria2	Alignment	not modelled	99.5	36	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c1w70A	Alignment	not modelled	99.5	36	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
85	c4az9B	Alignment	not modelled	99.5	16	PDB header: protein transport Chain: B: PDB Molecule: sorting nexin-24; PDBTitle: crystal structure of phox homology domain of human sorting2 nexin 24
86	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
87	c2fpeB	Alignment	not modelled	99.5	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
88	c2csiA	Alignment	not modelled	99.5	25	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
89	c2kt1A	Alignment	not modelled	99.5	32	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
90	c4afzC	Alignment	not modelled	99.5	29	PDB header: hydrolase/de novo protein Chain: C: PDB Molecule: fynomer; PDBTitle: human chymase - fynomer complex
91	c1x6bA	Alignment	not modelled	99.5	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
92	d1oota	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
93	c2j6kE	Alignment	not modelled	99.5	30	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
94	c2egeA	Alignment	not modelled	99.5	25	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
95	c2d1xD	Alignment	not modelled	99.5	38	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
96	c2azsA	Alignment	not modelled	99.5	24	PDB header: signaling protein Chain: A: PDB Molecule: sh2-sh3 adapter protein drk; PDBTitle: nmr structure of the n-terminal sh3 domain of drk2 (calculated without noe restraints)
97	c1wyxA	Alignment	not modelled	99.5	24	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
98	d1k4us	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
99	c2df6A	Alignment	not modelled	99.5	44	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
100	d1nega	Alignment	not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c1negA	Alignment	not modelled	99.5	28	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
102	c2dnuA	Alignment	not modelled	99.5	30	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
103	c2vunA	Alignment	not modelled	99.5	25	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin;

103	c2yupA	Alignment	not modelled	99.5	25	PDBTitle: solution structure of the second sh3 domain of human vinexin PDB header: signal transduction
104	c2j06B	Alignment	not modelled	99.5	26	Chain: B: PDB Molecule: ras gtpase-activating protein 1; PDBTitle: crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
105	c2dl7A	Alignment	not modelled	99.5	32	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
106	c2eqiA	Alignment	not modelled	99.5	27	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
107	c2cucA	Alignment	not modelled	99.5	26	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	c1s1nA	Alignment	not modelled	99.5	35	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
109	d1bb9a	Alignment	not modelled	99.5	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c1bb9A	Alignment	not modelled	99.5	37	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
111	c2ct3A	Alignment	not modelled	99.5	32	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
112	d1k9aa1	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	d1fmka1	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2rqrA	Alignment	not modelled	99.5	21	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
115	c1wxtA	Alignment	not modelled	99.5	25	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: hypothetical protein flj21522; PDBTitle: solution structure of the sh3 domain of human hypothetical2 protein flj21522
116	c1zlmA	Alignment	not modelled	99.5	27	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
117	c2epdA	Alignment	not modelled	99.5	25	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
118	d1sema	Alignment	not modelled	99.5	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
119	c2dmoA	Alignment	not modelled	99.5	20	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)
120	c1zukA	Alignment	not modelled	99.5	25	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