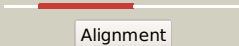


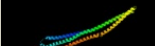

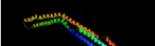

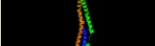

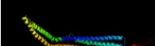


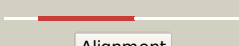


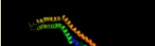





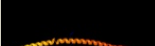
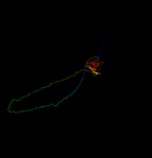

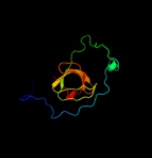
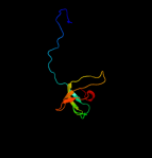
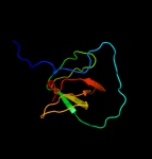

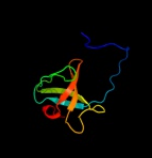
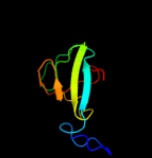



Phyre2

Email	I.a.kelley@imperial.ac.uk
Description	Q9XUS7
Date	Fri May 3 21:43:12 BST 2013
Unique Job ID	ae632157046dfe9d

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2efka1	 Alignment		100.0	21	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
2	d2efla1	 Alignment		100.0	26	Fold: BAR/IMD domain-like Superfamily: BAR/IMD domain-like Family: FCH domain
3	c3lllA_	 Alignment		100.0	13	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: crystal structure of mouse pacsin2 f-bar domain
4	c3i2wB_	 Alignment		100.0	14	PDB header: endocytosis Chain: B: PDB Molecule: syndapin; PDBTitle: crystal structure of efc/f-bar domain of drosophila2 syndapin/pacsin
5	c3hajA_	 Alignment		100.0	12	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin2 f-bar; PDBTitle: crystal structure of human pacsin2 f-bar domain (p2121212 lattice)
6	c3m3wA_	 Alignment		100.0	17	PDB header: endocytosis Chain: A: PDB Molecule: protein kinase c and casein kinase ii substrate protein 3; PDBTitle: crystal structure of mouse pacsin3 bar domain mutant
7	c2v0oB_	 Alignment		100.0	15	PDB header: lipid-binding protein Chain: B: PDB Molecule: fch domain only protein 2; PDBTitle: fcho2 f-bar domain
8	c3acoB_	 Alignment		100.0	12	PDB header: endocytosis Chain: B: PDB Molecule: protein kinase c and casein kinase substrate in PDBTitle: crystal structure of the efc/f-bar domain of human2 pacsin2/syndapin ii (2.7 a)
9	c3hahA_	 Alignment		99.9	14	PDB header: endocytosis Chain: A: PDB Molecule: human pacsin1 f-bar; PDBTitle: crystal structure of human pacsin1 f-bar domain (c2 lattice)
10	c3syvB_	 Alignment		99.9	17	PDB header: endocytosis Chain: B: PDB Molecule: protein kinase c and casein kinase ii substrate protein 3; PDBTitle: crystal structure of mpacsin 3 f-bar domain mutant
11	c3qweA_	 Alignment		99.9	13	PDB header: protein binding Chain: A: PDB Molecule: gem-interacting protein; PDBTitle: crystal structure of the n-terminal domain of the gem interacting2 protein

12	c1mv3A_	Alignment		99.8	8	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	c3g9gA_	Alignment		99.8	11	PDB header: endocytosis Chain: A: PDB Molecule: suppressor of yeast profilin deletion; PDBTitle: crystal structure of the n-terminal efc/f-bar domain of syp1
14	c2rqra_	Alignment		99.7	18	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
15	c1gria_	Alignment		99.6	15	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
16	c2jxbA_	Alignment		99.6	19	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
17	c1x6gA_	Alignment		99.6	12	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.
18	c1x43A_	Alignment		99.6	16	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)
19	d1wiea_	Alignment		99.6	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2dl5A_	Alignment		99.6	29	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
21	c3nmzD_	Alignment	not modelled	99.6	16	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
22	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
23	c2d8hA_	Alignment	not modelled	99.6	16	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
24	c2l3sA_	Alignment	not modelled	99.6	11	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
25	c2oi3A_	Alignment	not modelled	99.6	18	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)
26	c2lqnA_	Alignment	not modelled	99.6	13	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
27	c3qwxX_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
28	c2csqA_	Alignment	not modelled	99.5	23	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
29	c2dlbA_	Alignment	not modelled	99.5	17	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein;

29	c2ulpA	Alignment	not modelled	99.5	17	PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein PDB header: cell cycle
30	c2yt6A	Alignment	not modelled	99.5	20	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
31	c2eyyA	Alignment	not modelled	99.5	15	PDB header: signaling protein Chain: A: PDB Molecule: v-crck sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
32	c1wx6A	Alignment	not modelled	99.5	22	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
33	c3qwyA	Alignment	not modelled	99.5	25	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
34	d1ng2a2	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
35	c2yupA	Alignment	not modelled	99.5	18	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
36	c2kymA	Alignment	not modelled	99.5	23	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
37	d1l1ja	Alignment	not modelled	99.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
38	c1zukA	Alignment	not modelled	99.5	14	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
39	c3rniA	Alignment	not modelled	99.5	25	PDB header: protein binding Chain: A: PDB Molecule: brain-specific angiogenesis inhibitor 1-associated protein PDBTitle: crystal structure of the sh3 domain from irsp53 (baiap2)
40	c2ct4A	Alignment	not modelled	99.5	40	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
41	c2enmA	Alignment	not modelled	99.5	28	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
42	c2kbtA	Alignment	not modelled	99.5	23	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
43	d1spka	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
44	c2eyzA	Alignment	not modelled	99.5	12	PDB header: signaling protein Chain: A: PDB Molecule: v-crck sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
45	c1x2qA	Alignment	not modelled	99.5	20	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
46	c2ed0A	Alignment	not modelled	99.5	18	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)
47	c2ysqA	Alignment	not modelled	99.5	17	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
48	d1u5sa1	Alignment	not modelled	99.5	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
49	c2azsA	Alignment	not modelled	99.5	20	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)
50	d1udla	Alignment	not modelled	99.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
51	c2dl7A	Alignment	not modelled	99.5	21	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
52	d1jqqa	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
53	c2cudA	Alignment	not modelled	99.5	13	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adapter; PDBTitle: solution structure of the sh3 domain of the human src-like2 adopter protein (slap)
54	d1zuua1	Alignment	not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
55	c2epdA	Alignment	not modelled	99.5	27	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
56	c2egeA_	Alignment	not modelled	99.5	13	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
57	c2a28D_	Alignment	not modelled	99.5	31	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
58	d1gbra_	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	d1uffa_	Alignment	not modelled	99.5	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c1x6bA_	Alignment	not modelled	99.5	18	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
61	c2i0nA_	Alignment	not modelled	99.5	17	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
62	c2cubA_	Alignment	not modelled	99.5	17	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
63	c1z9zA_	Alignment	not modelled	99.5	17	PDB header: structural protein Chain: A: PDB Molecule: cytoskeleton assembly control protein sla1; PDBTitle: crystal structure of yeast sla1 sh3 domain 3
64	d2v1ra1	Alignment	not modelled	99.5	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	d1ugva_	Alignment	not modelled	99.5	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	c2x3wD_	Alignment	not modelled	99.5	24	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)
67	d1ov3a2	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c2rqvA_	Alignment	not modelled	99.5	19	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
69	c2csiA_	Alignment	not modelled	99.5	21	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
70	c1u3oA_	Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: A: PDB Molecule: huntingtin-associated protein-interacting PDBTitle: solution structure of rat kalirin n-terminal sh3 domain
71	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
72	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
73	c2e5kA_	Alignment	not modelled	99.5	13	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
74	c2lcsA_	Alignment	not modelled	99.5	13	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
75	c2m0yA_	Alignment	not modelled	99.5	17	PDB header: apoptosis Chain: A: PDB Molecule: dedicator of cytokinesis protein 1; PDBTitle: solution structure of the sh3 domain of dock180
76	c2lj0A_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
77	c2ed1A_	Alignment	not modelled	99.5	14	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
78	c2ekbA_	Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structures of the sh3 domain of human kiaa0418
79	c2yunA_	Alignment	not modelled	99.5	20	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
80	c2ct3A_	Alignment	not modelled	99.5	16	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
81	c2ke9A_	Alignment	not modelled	99.5	20	PDB header: protein binding Chain: A: PDB Molecule: caskin-2;

						PDBTitle: nmr solution structure of the caskin sh3 domain
82	c2d1xD	Alignment	not modelled	99.5	19	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
83	d1cska	Alignment	not modelled	99.5	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2df6A	Alignment	not modelled	99.4	24	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
85	d1oota	Alignment	not modelled	99.4	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
86	d1k9aa1	Alignment	not modelled	99.4	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	c2j06B	Alignment	not modelled	99.4	15	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
88	d1ue9a	Alignment	not modelled	99.4	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	c1bb9A	Alignment	not modelled	99.4	13	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
90	d1bb9a	Alignment	not modelled	99.4	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2l0aA	Alignment	not modelled	99.4	19	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
92	c2jteA	Alignment	not modelled	99.4	15	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: third sh3 domain of cd2ap
93	c1x69A	Alignment	not modelled	99.4	19	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
94	d1jo8a	Alignment	not modelled	99.4	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	d1phta	Alignment	not modelled	99.4	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	d1uj0a	Alignment	not modelled	99.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
97	c1k9aB	Alignment	not modelled	99.4	16	PDB header: transferase Chain: B: PDB Molecule: carboxyl-terminal src kinase; PDBTitle: crystal structure analysis of full-length carboxyl-terminal2 src kinase at 2.5 a resolution
98	c2creA	Alignment	not modelled	99.4	17	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
99	c2yuoA	Alignment	not modelled	99.4	13	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
100	c2ebpA	Alignment	not modelled	99.4	13	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
101	c1wyxA	Alignment	not modelled	99.4	33	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
102	d1k4us	Alignment	not modelled	99.4	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c2djQA	Alignment	not modelled	99.4	15	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
104	d1utia	Alignment	not modelled	99.4	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	c4glmD	Alignment	not modelled	99.4	15	PDB header: signaling protein Chain: D: PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
106	c2lx7A	Alignment	not modelled	99.4	23	PDB header: protein binding Chain: A: PDB Molecule: growth arrest-specific protein 7; PDBTitle: solution nmr structure of sh3 domain of growth arrest-specific protein2 7 (gas7) (fragment 1-60) from homo sapiens, northeast structural3 genomics consortium (nesg) target hr8574a
107	d1oeba	Alignment	not modelled	99.4	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

108	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
109	d1ycsb2	Alignment	not modelled	99.4	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c2kt1A_	Alignment	not modelled	99.4	24	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
111	c2dbkA_	Alignment	not modelled	99.4	15	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
112	c2j6kE_	Alignment	not modelled	99.4	17	PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
113	c2o2oA_	Alignment	not modelled	99.4	13	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
114	c1yn8E_	Alignment	not modelled	99.4	22	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
115	c2xmfA_	Alignment	not modelled	99.4	15	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
116	c4esrB_	Alignment	not modelled	99.4	24	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
117	c2dl8A_	Alignment	not modelled	99.4	23	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
118	c1zlmA_	Alignment	not modelled	99.4	19	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
119	c1w70A_	Alignment	not modelled	99.4	21	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
120	d1uhca_	Alignment	not modelled	99.4	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain