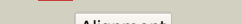
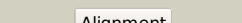
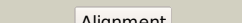
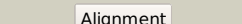
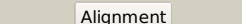
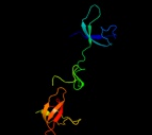
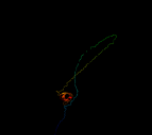
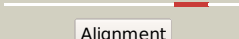
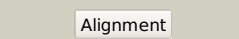
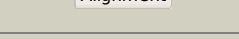
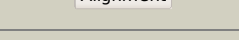
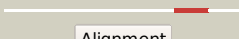
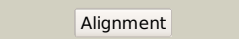
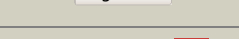
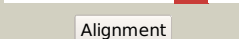
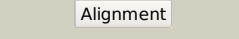
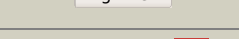
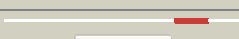


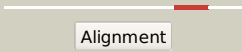
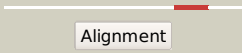
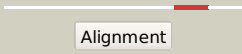
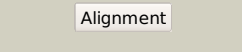
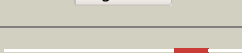
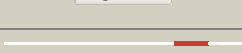
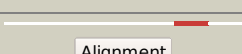
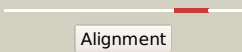
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2khnA_	 Alignment		100.0	37	PDB header: signaling protein Chain: A: PDB Molecule: intersectin-1; PDBTitle: nmr solution structure of the eh 1 domain from human2 intersectin-1 protein. northeast structural genomics3 consortium target hr3646e.
2	c1griA_	 Alignment		100.0	25	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
3	c2jq6A_	 Alignment		99.9	32	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	38	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	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
6	c2kfhA_	 Alignment		99.9	32	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	d2jxca1	 Alignment		99.9	36	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
8	c2eyzA_	 Alignment		99.9	19	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
9	c2l3sA_	 Alignment		99.9	22	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
10	c2qptA_	 Alignment		99.9	31	PDB header: endocytosis Chain: A: PDB Molecule: eh domain-containing protein-2; PDBTitle: crystal structure of an ehd atpase involved in membrane remodelling
11	c1ov3A_	 Alignment		99.9	26	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex

12	d1iq3a_	Alignment		99.9	32	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
13	c1ng2A_	Alignment		99.9	27	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
14	c2kgrA_	Alignment		99.9	34	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
15	d1fi6a_	Alignment		99.9	33	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
16	d1c07a_	Alignment		99.8	28	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
17	c1mv3A_	Alignment		99.8	20	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	25	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
19	c2dx1A_	Alignment		99.7	42	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
20	d1udla_	Alignment		99.7	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
21	d1ng2a2	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2rqra_	Alignment	not modelled	99.6	15	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
23	c2kymA_	Alignment	not modelled	99.6	25	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of the bem1p sh3-ci domain from I.elongisporus in2 complex with ste20p peptide
24	c2jxbA_	Alignment	not modelled	99.6	27	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
25	c3nmzD_	Alignment	not modelled	99.6	37	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
26	c1x6gA_	Alignment	not modelled	99.6	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.
27	c2yt6A_	Alignment	not modelled	99.6	30	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
28	d1ov3a2	Alignment	not modelled	99.6	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

29	c2ed1A		Alignment	not modelled	99.6	40	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
30	c1zukA		Alignment	not modelled	99.6	28	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
31	c2rqvA		Alignment	not modelled	99.6	23	PDB header: signaling protein Chain: A: PDB Molecule: bud emergence protein 1; PDBTitle: solution structure of sh3ci domain of bem1p
32	c1zlmA		Alignment	not modelled	99.6	44	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
33	c2eyyA		Alignment	not modelled	99.6	26	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
34	c2fpeB		Alignment	not modelled	99.6	21	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
35	d1ue9a		Alignment	not modelled	99.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
36	c2yupA		Alignment	not modelled	99.5	29	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
37	c2j06B		Alignment	not modelled	99.5	27	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
38	c2xmFA		Alignment	not modelled	99.5	36	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
39	d1l1ja		Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
40	c2e5kA		Alignment	not modelled	99.5	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
41	d1k4us		Alignment	not modelled	99.5	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	c2lj0A		Alignment	not modelled	99.5	29	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: the third sh3 domain of r85fl
43	c2d8jA		Alignment	not modelled	99.5	32	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
44	c2cudA		Alignment	not modelled	99.5	16	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)
45	c3qwY_A		Alignment	not modelled	99.5	23	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
46	c1x2qA		Alignment	not modelled	99.5	41	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
47	c2egcA		Alignment	not modelled	99.5	25	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
48	c2ysqA		Alignment	not modelled	99.5	40	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
49	c2df6A		Alignment	not modelled	99.5	39	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 7; PDBTitle: crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2
50	c3qwxX		Alignment	not modelled	99.5	22	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174
51	d1uj0a		Alignment	not modelled	99.5	48	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2ct3A		Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
53	c2yunA		Alignment	not modelled	99.5	38	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
54	c2m0yA		Alignment	not modelled	99.5	26	PDB header: apoptosis Chain: A: PDB Molecule: dedicator of cytokinesis protein 1; PDBTitle: solution structure of the sh3 domain of dock180

55	c1x6bA	Alignment	not modelled	99.5	20	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
56	d1phta	Alignment	not modelled	99.5	27	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
57	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
58	c2ekhA	Alignment	not modelled	99.5	28	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
59	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
60	c2ebpA	Alignment	not modelled	99.5	28	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
61	c4esrB	Alignment	not modelled	99.5	39	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
62	c2cubA	Alignment	not modelled	99.5	32	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	c2nwmA	Alignment	not modelled	99.5	30	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
64	c2ke9A	Alignment	not modelled	99.5	29	PDB header: protein binding Chain: A; PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
65	c2dlpA	Alignment	not modelled	99.5	30	PDB header: structural protein Chain: A; PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
66	c2csqA	Alignment	not modelled	99.5	31	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
67	c2djqa	Alignment	not modelled	99.5	33	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
68	c2yuoA	Alignment	not modelled	99.5	33	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
69	c1zuyB	Alignment	not modelled	99.5	38	PDB header: contractile protein Chain: B; PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
70	c4glmD	Alignment	not modelled	99.5	33	PDB header: signaling protein Chain: D; PDB Molecule: dynammin-binding protein; PDBTitle: crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
71	c2lcsA	Alignment	not modelled	99.5	31	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
72	c2ed0A	Alignment	not modelled	99.5	34	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)
73	d1gcqa	Alignment	not modelled	99.5	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c2kgtA	Alignment	not modelled	99.5	27	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
75	d1oeba	Alignment	not modelled	99.5	47	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
76	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
77	c2dilA	Alignment	not modelled	99.5	34	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
78	c1yn8E	Alignment	not modelled	99.5	32	PDB header: unknown function Chain: E; PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
79	c2oi3A	Alignment	not modelled	99.5	32	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	26	Fold: SH3-like barrel Superfamily: SH3-domain

						Family:SH3-domain
81	c2dl4A_	Alignment	not modelled	99.5	31	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
82	c2dmoA_	Alignment	not modelled	99.5	21	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)
83	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
84	c2cucA_	Alignment	not modelled	99.5	31	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
85	c2bz8B_	Alignment	not modelled	99.5	37	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
86	d1utia_	Alignment	not modelled	99.5	50	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
87	d1wlpb2	Alignment	not modelled	99.5	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	c2i0nA_	Alignment	not modelled	99.5	31	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
89	c2d1xD_	Alignment	not modelled	99.5	39	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
90	d1bb9a_	Alignment	not modelled	99.5	32	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c1bb9A_	Alignment	not modelled	99.5	32	PDB header: transferase Chain: A: PDB Molecule: amphiphysin 2; PDBTitle: crystal structure of the sh3 domain from rat amphiphysin 2
92	c2enmA_	Alignment	not modelled	99.5	29	PDB header: endocytosis Chain: A: PDB Molecule: sorting nexin-9; PDBTitle: solution structure of the sh3 domain from mouse sorting2 nexin-9
93	d1jo8a_	Alignment	not modelled	99.5	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c2kt1A_	Alignment	not modelled	99.5	30	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
95	d1sema_	Alignment	not modelled	99.5	40	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2js0A_	Alignment	not modelled	99.5	38	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
97	c2d8hA_	Alignment	not modelled	99.5	28	PDB header: unknown function Chain: A: PDB Molecule: sh3yl1 protein; PDBTitle: solution structure of the sh3 domain of hypothetical2 protein sh3yl1
98	c1w70A_	Alignment	not modelled	99.5	35	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
99	c2dbkA_	Alignment	not modelled	99.5	21	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
100	c2dl8A_	Alignment	not modelled	99.5	27	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
101	c2dnuA_	Alignment	not modelled	99.5	26	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
102	c4afzC_	Alignment	not modelled	99.5	32	PDB header: hydrolase/de novo protein Chain: C: PDB Molecule: fyonomer; PDBTitle: human chymase - fynomer complex
103	c2drnB_	Alignment	not modelled	99.5	51	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
104	d1wiea_	Alignment	not modelled	99.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
105	d1gria2	Alignment	not modelled	99.5	41	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
106	d1uffa_	Alignment	not modelled	99.5	33	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

107	c2rf0D_		not modelled	99.5	40	PDB header: transferase Chain: D: PDB Molecule: mitogen-activated protein kinase kinase 10; PDBTitle: crystal structure of human mixed lineage kinase map3k10 sh3 domain
108	d1uuea_		not modelled	99.5	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	c2dl3A_		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
110	c2eyxA_		not modelled	99.5	25	PDB header: signaling protein Chain: A: PDB Molecule: v-crck sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
111	c1x2pA_		not modelled	99.5	31	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
112	c3rnjA_		not modelled	99.5	24	PDB header: protein binding Chain: A: PDB Molecule: brain-specific angiogenesis inhibitor 1-associated protein PDBTitle: crystal structure of the sh3 domain from irsp53 (baiap2)
113	c1zx6A_		not modelled	99.5	35	PDB header: protein binding Chain: A: PDB Molecule: yp154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
114	d1fmka1		not modelled	99.5	35	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
115	c4igzA_		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
116	c2eqiA_		not modelled	99.5	37	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
117	c2kbtA_		not modelled	99.5	37	PDB header: signaling protein Chain: A: PDB Molecule: chimera of proto-oncogene vav, linker, tag2 (inset) using a sortase-mediated protein ligation method
118	c1x43A_		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)
119	d2iima1		not modelled	99.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2krnA_		not modelled	99.5	42	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap