
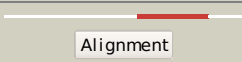
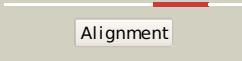
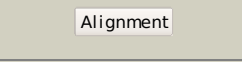

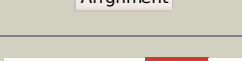
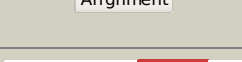
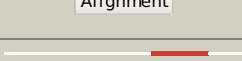
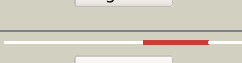
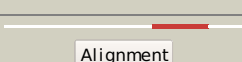
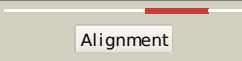
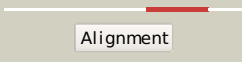

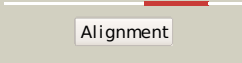
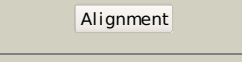
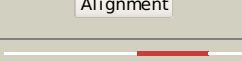


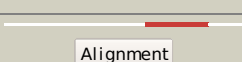
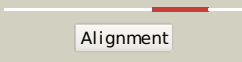
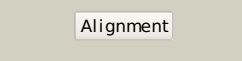

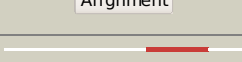




#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1griA_	 Alignment		100.0	100	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
2	c2lqnA_	 Alignment		100.0	20	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
3	c2l3sA_	 Alignment		100.0	23	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
4	c2eyzA_	 Alignment		100.0	25	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
5	c1ng2A_	 Alignment		100.0	24	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
6	c1ov3A_	 Alignment		100.0	25	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
7	c2eyyA_	 Alignment		99.9	27	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
8	c1k9aB_	 Alignment		99.9	29	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
9	c1oplA_	 Alignment		99.9	30	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase; PDBTitle: structural basis for the auto-inhibition of c-abl tyrosine2 kinase
10	c2fo0A_	 Alignment		99.9	30	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase abl1 (1b isoform); PDBTitle: organization of the sh3-sh2 unit in active and inactive forms of the2 c-abl tyrosine kinase
11	c3qwxX_	 Alignment		99.9	22	PDB header: signaling protein Chain: X: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2 1-174

12	c2dvjA_	Alignment		99.9	28	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog, PDBTitle: phosphorylated crk-ii
13	c3qwyA_	Alignment		99.9	24	PDB header: signaling protein Chain: A: PDB Molecule: cell death abnormality protein 2; PDBTitle: ced-2
14	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
15	c2dybA_	Alignment		99.8	23	PDB header: oxidoreductase Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: the crystal structure of human p40(phox)
16	c1y57A_	Alignment		99.8	33	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: structure of unphosphorylated c-src in complex with an inhibitor
17	c2c0iA_	Alignment		99.8	37	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: src family kinase hck with bound inhibitor a-420983
18	c2h8hA_	Alignment		99.8	32	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase src; PDBTitle: src kinase in complex with a quinazoline inhibitor
19	d1ng2a2	Alignment		99.8	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2ablA_	Alignment		99.8	30	PDB header: transferase Chain: A: PDB Molecule: abl tyrosine kinase; PDBTitle: sh3-sh2 domain fragment of human bcr-abl tyrosine kinase
21	d1gria2	Alignment	not modelled	99.8	100	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	c2dx1A_	Alignment	not modelled	99.8	33	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
23	c1x27F_	Alignment	not modelled	99.8	35	PDB header: signaling protein Chain: F: PDB Molecule: proto-oncogene tyrosine-protein kinase lck; PDBTitle: crystal structure of lck sh2-sh3 with sh2 binding site of2 p130cas
24	c3nmzD_	Alignment	not modelled	99.8	27	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
25	c2ed0A_	Alignment	not modelled	99.8	27	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)
26	d1sema_	Alignment	not modelled	99.8	55	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
27	c2epdA_	Alignment	not modelled	99.8	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
28	c2xmfA_	Alignment	not modelled	99.8	36	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3

29	c2cubA	Alignment	not modelled	99.8	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
30	d1pwta	Alignment	not modelled	99.8	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
31	c2eyxA	Alignment	not modelled	99.8	20	PDB header: signaling protein Chain: A: PDB Molecule: v-crK sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
32	c2bz8B	Alignment	not modelled	99.8	32	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
33	d1wlpb2	Alignment	not modelled	99.8	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
34	c2yuoA	Alignment	not modelled	99.8	32	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
35	c1x2qA	Alignment	not modelled	99.8	40	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
36	d1uuea	Alignment	not modelled	99.8	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
37	c2dmoA	Alignment	not modelled	99.8	29	PDB header: signaling protein Chain: A: PDB Molecule: neutrophil cytosol factor 2; PDBTitle: refined solution structure of the 1st sh3 domain from human2 neutrophil cytosol factor 2 (ncf-2)
38	c2yunA	Alignment	not modelled	99.8	28	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
39	c2djgA	Alignment	not modelled	99.8	27	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
40	c2df6A	Alignment	not modelled	99.7	40	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
41	c3nhnA	Alignment	not modelled	99.7	37	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase hck; PDBTitle: crystal structure of the src-family kinase hck sh3-sh2-linker2 regulatory region
42	c2dl8A	Alignment	not modelled	99.7	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
43	c2kymA	Alignment	not modelled	99.7	25	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
44	c2dbkA	Alignment	not modelled	99.7	16	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein
45	d1udla	Alignment	not modelled	99.7	30	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
46	d1lutia	Alignment	not modelled	99.7	47	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	c2dl3A	Alignment	not modelled	99.7	32	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
48	c1x69A	Alignment	not modelled	99.7	30	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
49	c2oi3A	Alignment	not modelled	99.7	22	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)
50	c2d1xD	Alignment	not modelled	99.7	36	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
51	c2ebpA	Alignment	not modelled	99.7	20	PDB header: cell adhesion Chain: A: PDB Molecule: sam and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
52	c2yt6A	Alignment	not modelled	99.7	23	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
53	d1gcqa	Alignment	not modelled	99.7	98	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

54	c2nwmA_	Alignment	not modelled	99.7	34	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
55	c2ekhA_	Alignment	not modelled	99.7	17	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
56	c2dl4A_	Alignment	not modelled	99.7	33	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
57	c2ysqA_	Alignment	not modelled	99.7	26	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
58	d1e6ga_	Alignment	not modelled	99.7	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
59	d1ov3a2_	Alignment	not modelled	99.7	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c1g83A_	Alignment	not modelled	99.7	31	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: crystal structure of fyn sh3-sh2
61	c2dlpA_	Alignment	not modelled	99.7	29	PDB header: structural protein Chain: A: PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
62	c2fpeB_	Alignment	not modelled	99.7	26	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
63	c2pqhA_	Alignment	not modelled	99.7	24	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
64	d1ue9a_	Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
65	c1zx6A_	Alignment	not modelled	99.7	51	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
66	c1wi7A_	Alignment	not modelled	99.7	33	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	c2dilA_	Alignment	not modelled	99.7	25	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
68	c2drmB_	Alignment	not modelled	99.7	39	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin ib; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
69	c1negA_	Alignment	not modelled	99.7	24	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
70	d1nega_	Alignment	not modelled	99.7	24	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	d1u06a1_	Alignment	not modelled	99.7	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
72	c2dnuA_	Alignment	not modelled	99.7	18	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
73	c2ed1A_	Alignment	not modelled	99.7	25	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
74	d1h8ka_	Alignment	not modelled	99.7	21	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
75	c1w70A_	Alignment	not modelled	99.7	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
76	d1k4us_	Alignment	not modelled	99.7	29	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
77	c2rqrA_	Alignment	not modelled	99.7	11	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
78	c2azsA_	Alignment	not modelled	99.7	72	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)

79	c1zukA		Alignment	not modelled	99.7	27	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
80	c2yupA		Alignment	not modelled	99.7	23	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
81	d1jo8a		Alignment	not modelled	99.7	39	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
82	c2ct3A		Alignment	not modelled	99.7	34	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
83	d1e6ha		Alignment	not modelled	99.7	20	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2dbmA		Alignment	not modelled	99.7	40	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
85	c2l0aA		Alignment	not modelled	99.7	44	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
86	c2csqA		Alignment	not modelled	99.7	26	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
87	c2krnA		Alignment	not modelled	99.7	32	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
88	d1ujya		Alignment	not modelled	99.7	31	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
89	d1loota		Alignment	not modelled	99.7	43	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
90	d1ov3a1		Alignment	not modelled	99.7	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
91	c2egaA		Alignment	not modelled	99.7	18	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
92	c1x43A		Alignment	not modelled	99.7	20	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)
93	c2i0nA		Alignment	not modelled	99.7	27	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
94	d1oeba		Alignment	not modelled	99.7	44	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
95	d1uj0a		Alignment	not modelled	99.7	51	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
96	c2cudA		Alignment	not modelled	99.7	19	PDB header: signaling protein Chain: A: PDB Molecule: src-like-adaptor; PDBTitle: solution structure of the sh3 domain of the human src-like2 adpoter protein (slap)
97	c3rniA		Alignment	not modelled	99.7	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)
98	c2eqiA		Alignment	not modelled	99.7	31	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
99	c2o2oA		Alignment	not modelled	99.7	32	PDB header: protein binding Chain: A: PDB Molecule: sh3-domain kinase-binding protein 1; PDBTitle: solution structure of domain b from human cin85 protein
100	c2bzxA		Alignment	not modelled	99.7	23	PDB header: sh3 domain Chain: A: PDB Molecule: crk-like protein; PDBTitle: atomic model of crkl-sh3c monomer
101	c2dl7A		Alignment	not modelled	99.7	29	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the second sh3 domain of human2 kiaa0769 protein
102	d1jqqa		Alignment	not modelled	99.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
103	c1z9qA		Alignment	not modelled	99.7	31	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
104	c1x2pA		Alignment	not modelled	99.7	23	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
105	c2j6kE	Alignment	not modelled	99.7	32 PDB header: protein binding Chain: E: PDB Molecule: cd2-associated protein; PDBTitle: n-terminal sh3 domain of cms (cd2ap human homolog)
106	c1x6gA	Alignment	not modelled	99.7	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.
107	c2e5kA	Alignment	not modelled	99.7	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
108	d1uhfa	Alignment	not modelled	99.7	31 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
109	d1uhca	Alignment	not modelled	99.7	17 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
110	c1zlmA	Alignment	not modelled	99.7	46 PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
111	c1x6bA	Alignment	not modelled	99.7	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
112	c1wyxA	Alignment	not modelled	99.7	30 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
113	d1ng2a1	Alignment	not modelled	99.7	21 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
114	c2egeA	Alignment	not modelled	99.7	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
115	c2cucA	Alignment	not modelled	99.7	30 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
116	c2ke9A	Alignment	not modelled	99.7	33 PDB header: protein binding Chain: A: PDB Molecule: caskin-2; PDBTitle: nmr solution structure of the caskin sh3 domain
117	c2kbtA	Alignment	not modelled	99.7	31 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
118	c2lcsA	Alignment	not modelled	99.7	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
119	d2v1ra1	Alignment	not modelled	99.7	32 Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2js0A	Alignment	not modelled	99.7	33 PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck