
















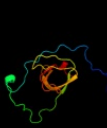









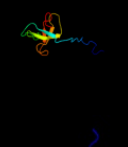
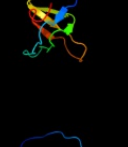

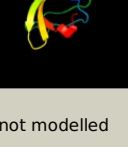


# Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	E6N0W1
Date	Fri May 3 21:47:45 BST 2013
Unique Job ID	d228f0f148128221

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c1griA_</a>	 Alignment		99.9	26	<b>PDB header:</b> signal transduction adaptor <b>Chain:</b> A: <b>PDB Molecule:</b> growth factor bound protein 2; <b>PDBTitle:</b> grb2
2	<a href="#">c2lqnA_</a>	 Alignment		99.9	24	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> crk-like protein; <b>PDBTitle:</b> solution structure of crkl
3	<a href="#">c2eyzA_</a>	 Alignment		99.9	28	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> v-crk sarcoma virus ct10 oncogene homolog <b>PDBTitle:</b> ct10-regulated kinase isoform ii
4	<a href="#">c2l3sA_</a>	 Alignment		99.9	23	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> autoinhibited crk protein; <b>PDBTitle:</b> structure of the autoinhibited crk
5	<a href="#">c1ov3A_</a>	 Alignment		99.8	26	<b>PDB header:</b> oxidoreductase activator <b>Chain:</b> A: <b>PDB Molecule:</b> neutrophil cytosol factor 1; <b>PDBTitle:</b> structure of the p22phox-p47phox complex
6	<a href="#">c1ng2A_</a>	 Alignment		99.8	27	<b>PDB header:</b> oxidoreductase activator <b>Chain:</b> A: <b>PDB Molecule:</b> neutrophil cytosolic factor 1; <b>PDBTitle:</b> structure of autoinhibited p47phox
7	<a href="#">c1mv3A_</a>	 Alignment		99.8	20	<b>PDB header:</b> endocytosis/exocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> myc box dependent interacting protein 1; <b>PDBTitle:</b> nmr structure of the tumor suppressor bin1: alternative2 splicing in melanoma and interaction with c-myc
8	<a href="#">c2dx1A_</a>	 Alignment		99.7	42	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> rho guanine nucleotide exchange factor 4; <b>PDBTitle:</b> crystal structure of rhogef protein asef
9	<a href="#">c2rqrA_</a>	 Alignment		99.7	15	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> engulfment and cell motility protein 1, linker, dedicator <b>PDBTitle:</b> the solution structure of human dock2 sh3 domain - elmo1 peptide2 chimera complex
10	<a href="#">d1ng2a2</a>	 Alignment		99.7	30	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
11	<a href="#">c2kymA_</a>	 Alignment		99.7	24	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> bud emergence protein 1; <b>PDBTitle:</b> solution structure of the bem1p sh3-ci domain from l.elongisporus in2 complex with ste20p peptide

12	<a href="#">d1udla_</a>	Alignment		99.6	44	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
13	<a href="#">c3nmzD_</a>	Alignment		99.6	29	<b>PDB header:</b> cell adhesion/cell cycle <b>Chain:</b> D: <b>PDB Molecule:</b> rho guanine nucleotide exchange factor 4; <b>PDBTitle:</b> crytal structure of apc complexed with asef
14	<a href="#">c1zukA_</a>	Alignment		99.6	28	<b>PDB header:</b> contractile protein <b>Chain:</b> A: <b>PDB Molecule:</b> myosin tail region-interacting protein mt1; <b>PDBTitle:</b> yeast bbc1 sh3 domain complexed with a peptide from las17
15	<a href="#">c1zlmA_</a>	Alignment		99.6	43	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> osteoclast stimulating factor 1; <b>PDBTitle:</b> crystal structure of the sh3 domain of human osteoclast2 stimulating factor
16	<a href="#">c2ed1A_</a>	Alignment		99.6	38	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> 130 kda phosphatidylinositol 4,5-biphosphate- <b>PDBTitle:</b> solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein
17	<a href="#">c2eyvA_</a>	Alignment		99.6	25	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> v-crk sarcoma virus ct10 oncogene homolog <b>PDBTitle:</b> ct10-regulated kinase isoform i
18	<a href="#">c2rqvA_</a>	Alignment		99.6	23	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> bud emergence protein 1; <b>PDBTitle:</b> solution structure of sh3ci domain of bem1p
19	<a href="#">c2yt6A_</a>	Alignment		99.6	29	<b>PDB header:</b> cell cycle <b>Chain:</b> A: <b>PDB Molecule:</b> adult male urinary bladder cdna, riken full- <b>PDBTitle:</b> solution structure of the sh3_1 domain of yamaguchi sarcoma2 viral (v-yes) oncogene homolog 1
20	<a href="#">d1ue9a_</a>	Alignment		99.6	26	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
21	<a href="#">c3qwvA_</a>	Alignment	not modelled	99.6	22	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cell death abnormality protein 2; <b>PDBTitle:</b> ced-2
22	<a href="#">d1ov3a2</a>	Alignment	not modelled	99.6	30	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
23	<a href="#">c2jxbA_</a>	Alignment	not modelled	99.6	28	<b>PDB header:</b> signaling protein complex <b>Chain:</b> A: <b>PDB Molecule:</b> t-cell surface glycoprotein cd3 epsilon chain, <b>PDBTitle:</b> structure of cd3epsilon-nck2 first sh3 domain complex
24	<a href="#">c1x2qA_</a>	Alignment	not modelled	99.6	40	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> signal transducing adapter molecule 2; <b>PDBTitle:</b> solution structure of the sh3 domain of the signal2 transducing adaptor molecule 2
25	<a href="#">c2yupA_</a>	Alignment	not modelled	99.6	27	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> vinexin; <b>PDBTitle:</b> solution structure of the second sh3 domain of human vinexin
26	<a href="#">d1uj0a_</a>	Alignment	not modelled	99.6	48	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
27	<a href="#">c2df6A_</a>	Alignment	not modelled	99.6	39	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> rho guanine nucleotide exchange factor 7; <b>PDBTitle:</b> crystal structure of the sh3 domain of betapix in complex2 with a high affinity peptide from pak2
28	<a href="#">c1bb9A_</a>	Alignment	not modelled	99.6	32	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> amphiphysin 2; <b>PDBTitle:</b> crystal structure of the sh3 domain from rat amphiphysin 2
						<b>Fold:</b> SH3-like barrel

29	<a href="#">d1bb9a_</a>	Alignment	not modelled	99.6	32	<b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
30	<a href="#">c3qwxX_</a>	Alignment	not modelled	99.6	21	<b>PDB header:</b> signaling protein <b>Chain:</b> X: <b>PDB Molecule:</b> cell death abnormality protein 2; <b>PDBTitle:</b> ced-2 1-174
31	<a href="#">d1l1ja_</a>	Alignment	not modelled	99.6	33	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
32	<a href="#">c2nwmA_</a>	Alignment	not modelled	99.6	30	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> vinexin; <b>PDBTitle:</b> solution structure of the first sh3 domain of human vinexin2 and its interaction with the peptides from vinculin
33	<a href="#">c2d8jA_</a>	Alignment	not modelled	99.6	32	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> fyn-related kinase; <b>PDBTitle:</b> solution structure of the sh3 domain of fyn-related kinase
34	<a href="#">c2lj0A_</a>	Alignment	not modelled	99.6	29	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sorbin and sh3 domain-containing protein 1; <b>PDBTitle:</b> the third sh3 domain of r85fl
35	<a href="#">c2ysqA_</a>	Alignment	not modelled	99.6	37	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> rho guanine nucleotide exchange factor 9; <b>PDBTitle:</b> solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
36	<a href="#">d1k4us_</a>	Alignment	not modelled	99.6	38	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
37	<a href="#">c2fpeB_</a>	Alignment	not modelled	99.6	20	<b>PDB header:</b> signaling protein <b>Chain:</b> B: <b>PDB Molecule:</b> cc-jun-amino-terminal kinase interacting protein <b>PDBTitle:</b> conserved dimerization of the ib1 src-homology 3 domain
38	<a href="#">c3rnjA_</a>	Alignment	not modelled	99.6	28	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> brain-specific angiogenesis inhibitor 1-associated protein <b>PDBTitle:</b> crystal structure of the sh3 domain from irsp53 (baiap2)
39	<a href="#">d1oeba_</a>	Alignment	not modelled	99.6	47	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
40	<a href="#">c2l0aA_</a>	Alignment	not modelled	99.6	42	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> signal transducing adapter molecule 1; <b>PDBTitle:</b> solution nmr structure of signal transducing adapter molecule 1 stam-12 from homo sapiens, northeast structural genomics consortium target3 hr4479e
41	<a href="#">c2e5kA_</a>	Alignment	not modelled	99.6	23	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> suppressor of t-cell receptor signaling 1; <b>PDBTitle:</b> solution structure of sh3 domain in suppressor of t-cell2 receptor signaling 1
42	<a href="#">c4glmD_</a>	Alignment	not modelled	99.6	33	<b>PDB header:</b> signaling protein <b>Chain:</b> D: <b>PDB Molecule:</b> dynamamin-binding protein; <b>PDBTitle:</b> crystal structure of the sh3 domain of dnmbp protein [homo sapiens]
43	<a href="#">c2ed0A_</a>	Alignment	not modelled	99.6	34	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> abl interactor 2; <b>PDBTitle:</b> solution structure of the sh3 domain of abl interactor 22 (abelson interactor 2)
44	<a href="#">c2yunA_</a>	Alignment	not modelled	99.6	38	<b>PDB header:</b> protein transport <b>Chain:</b> A: <b>PDB Molecule:</b> nostrin; <b>PDBTitle:</b> solution structure of the sh3 domain of human nostrin
45	<a href="#">d1gcqa_</a>	Alignment	not modelled	99.6	40	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
46	<a href="#">c2ebpA_</a>	Alignment	not modelled	99.6	28	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> sam and sh3 domain-containing protein 1; <b>PDBTitle:</b> solution structure of the sh3 domain from human sam and sh32 domain containing protein 1
47	<a href="#">c2yuoA_</a>	Alignment	not modelled	99.6	32	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> run and tbc1 domain containing 3; <b>PDBTitle:</b> solution structure of the sh3 domain of mouse run and tbc12 domain containing 3
48	<a href="#">c2cubA_</a>	Alignment	not modelled	99.6	32	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cytoplasmic protein nck1; <b>PDBTitle:</b> solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
49	<a href="#">c4esrB_</a>	Alignment	not modelled	99.6	39	<b>PDB header:</b> protein binding <b>Chain:</b> B: <b>PDB Molecule:</b> jouberein; <b>PDBTitle:</b> 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
50	<a href="#">c2dlpA_</a>	Alignment	not modelled	99.6	30	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> kiaa1783 protein; <b>PDBTitle:</b> solution structure of the sh3 domain of human kiaa17832 protein
51	<a href="#">c1x6gA_</a>	Alignment	not modelled	99.6	19	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> megakaryocyte-associated tyrosine-protein kinase; <b>PDBTitle:</b> solution structures of the sh3 domain of human2 megakaryocyte-associated tyrosine-protein kinase.
52	<a href="#">c2ekhA_</a>	Alignment	not modelled	99.6	28	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 and px domain-containing protein 2a; <b>PDBTitle:</b> solution structures of the sh3 domain of human kiaa0418
53	<a href="#">c2j06B_</a>	Alignment	not modelled	99.6	27	<b>PDB header:</b> signal transduction <b>Chain:</b> B: <b>PDB Molecule:</b> ras gtpase-activating protein 1; <b>PDBTitle:</b> crystal structure of the rasgap sh3 domain at 1.8 angstrom2 resolution
54	<a href="#">c2xmfA_</a>	Alignment	not modelled	99.6	36	<b>PDB header:</b> motor protein <b>Chain:</b> A: <b>PDB Molecule:</b> myosin 1e sh3; <b>PDBTitle:</b> myosin 1e sh3

55	<a href="#">c2ke9A_</a>	Alignment	not modelled	99.6	31	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> caskin-2; <b>PDBTitle:</b> nmr solution structure of the caskin sh3 domain
56	<a href="#">c2epdA_</a>	Alignment	not modelled	99.6	28	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> rho gtpase-activating protein 4; <b>PDBTitle:</b> solution structure of sh3 domain in rho-gtpase-activating2 protein 4
57	<a href="#">c1wx6A_</a>	Alignment	not modelled	99.6	23	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> cytoplasmic protein nck2; <b>PDBTitle:</b> solution structure of the sh3 domain of the human2 cytoplasmic protein nck2
58	<a href="#">d1utia_</a>	Alignment	not modelled	99.6	50	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
59	<a href="#">c2ct3A_</a>	Alignment	not modelled	99.5	33	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> vinexin; <b>PDBTitle:</b> solution structure of the sh3 domain of the vinexin protein
60	<a href="#">c2djqa_</a>	Alignment	not modelled	99.5	33	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 domain containing ring finger 2; <b>PDBTitle:</b> the solution structure of the first sh3 domain of mouse sh32 domain containing ring finger 2
61	<a href="#">c1x6bA_</a>	Alignment	not modelled	99.5	21	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> rho guanine exchange factor (gef) 16; <b>PDBTitle:</b> solution structures of the sh3 domain of human rho guanine2 exchange factor (gef) 16
62	<a href="#">d1gria2</a>	Alignment	not modelled	99.5	40	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
63	<a href="#">d1uuea_</a>	Alignment	not modelled	99.5	28	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
64	<a href="#">c2eyxA_</a>	Alignment	not modelled	99.5	23	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> v-crk sarcoma virus ct10 oncogene homolog <b>PDBTitle:</b> c-terminal sh3 domain of ct10-regulated kinase
65	<a href="#">d1sema_</a>	Alignment	not modelled	99.5	40	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
66	<a href="#">d1wlpb2</a>	Alignment	not modelled	99.5	30	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
67	<a href="#">c2egcA_</a>	Alignment	not modelled	99.5	24	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 and px domain-containing protein 2a; <b>PDBTitle:</b> solution structure of the fifth sh3 domain from human2 kiaa0418 protein
68	<a href="#">c2cudA_</a>	Alignment	not modelled	99.5	16	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> src-like-adaptor; <b>PDBTitle:</b> solution structure of the sh3 domain of the human src-like2 adppter protein (slap)
69	<a href="#">c2m0yA_</a>	Alignment	not modelled	99.5	27	<b>PDB header:</b> apoptosis <b>Chain:</b> A: <b>PDB Molecule:</b> dedicator of cytokinesis protein 1; <b>PDBTitle:</b> solution structure of the sh3 domain of dock180
70	<a href="#">d1ckaa_</a>	Alignment	not modelled	99.5	31	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
71	<a href="#">c2enmA_</a>	Alignment	not modelled	99.5	29	<b>PDB header:</b> endocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> sorting nexin-9; <b>PDBTitle:</b> solution structure of the sh3 domain from mouse sorting2 nexin-9
72	<a href="#">c2egeA_</a>	Alignment	not modelled	99.5	23	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> uncharacterized protein kiaa1666; <b>PDBTitle:</b> solution structure of the third sh3 domain from human2 kiaa1666 protein
73	<a href="#">c2dl4A_</a>	Alignment	not modelled	99.5	30	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> protein stac; <b>PDBTitle:</b> solution structure of the first sh3 domain of stac protein
74	<a href="#">c1zx6A_</a>	Alignment	not modelled	99.5	34	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> ypr154wp; <b>PDBTitle:</b> high-resolution crystal structure of yeast pin3 sh3 domain
75	<a href="#">c2dmoA_</a>	Alignment	not modelled	99.5	21	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> neutrophil cytosol factor 2; <b>PDBTitle:</b> refined solution structure of the 1st sh3 domain from human2 neutrophil cytosol factor 2 (ncf-2)
76	<a href="#">c2kgtA_</a>	Alignment	not modelled	99.5	27	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase 6; <b>PDBTitle:</b> solution structure of sh3 domain of ptk6
77	<a href="#">d1cska_</a>	Alignment	not modelled	99.5	26	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
78	<a href="#">c2d1xD</a>	Alignment	not modelled	99.5	38	<b>PDB header:</b> cell invasion <b>Chain:</b> D: <b>PDB Molecule:</b> cortactin isoform a; <b>PDBTitle:</b> the crystal structure of the cortactin-sh3 domain and amap1-2 peptide complex
79	<a href="#">d1uffa_</a>	Alignment	not modelled	99.5	33	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
80	<a href="#">c2kbtA_</a>	Alignment	not modelled	99.5	37	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> chimera of proto-oncogene vav, linker, <b>PDBTitle:</b> attachment of an nmr-invisible solubility enhancement tag2 (inset) using a sortase-mediated protein ligation method
81	<a href="#">d1phta_</a>	Alignment	not modelled	99.5	29	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain

82	<a href="#">c2i0nA_</a>	Alignment	not modelled	99.5	32	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> class vii unconventional myosin; <b>PDBTitle:</b> structure of dictyostelium discoideum myosin vii sh3 domain2 with adjacent proline rich region
83	<a href="#">c2dl8A_</a>	Alignment	not modelled	99.5	26	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> slit-robo rho gtpase-activating protein 2; <b>PDBTitle:</b> solution structure of the sh3 domain of human slit-robo rho2 gtpase-activating protein 2
84	<a href="#">c2csqA_</a>	Alignment	not modelled	99.5	28	<b>PDB header:</b> endocytosis/exocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> rim binding protein 2; <b>PDBTitle:</b> solution structure of the second sh3 domain of human rim-2 binding protein 2
85	<a href="#">d1pwta_</a>	Alignment	not modelled	99.5	26	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
86	<a href="#">d2v1ra1</a>	Alignment	not modelled	99.5	29	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
87	<a href="#">d1gl5a_</a>	Alignment	not modelled	99.5	28	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
88	<a href="#">c2lcsA_</a>	Alignment	not modelled	99.5	33	<b>PDB header:</b> transferase, signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> nap1-binding protein 2; <b>PDBTitle:</b> yeast nbp2p sh3 domain in complex with a peptide from ste20p
89	<a href="#">c2dbkA_</a>	Alignment	not modelled	99.5	21	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> crk-like protein; <b>PDBTitle:</b> solution structures of the sh3 domain of human crk-like2 protein
90	<a href="#">c1zuyB_</a>	Alignment	not modelled	99.5	36	<b>PDB header:</b> contractile protein <b>Chain:</b> B: <b>PDB Molecule:</b> myosin-5 isoform; <b>PDBTitle:</b> high-resolution structure of yeast myo5 sh3 domain
91	<a href="#">c2rf0D_</a>	Alignment	not modelled	99.5	40	<b>PDB header:</b> transferase <b>Chain:</b> D: <b>PDB Molecule:</b> mitogen-activated protein kinase kinase kinase 10; <b>PDBTitle:</b> crystal structure of human mixed lineage kinase map3k10 sh3 domain
92	<a href="#">c2dnuA_</a>	Alignment	not modelled	99.5	26	<b>PDB header:</b> structural genomics, structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 multiple domains 1; <b>PDBTitle:</b> solution structure of rsgi ruh-061, a sh3 domain from human
93	<a href="#">d1u06a1</a>	Alignment	not modelled	99.5	28	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
94	<a href="#">c2dilA_</a>	Alignment	not modelled	99.5	34	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> proline-serine-threonine phosphatase-interacting <b>PDBTitle:</b> solution structure of the sh3 domain of the human proline-2 serine-threonine phosphatase-interacting protein 1
95	<a href="#">d1u5sa1</a>	Alignment	not modelled	99.5	28	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
96	<a href="#">c2bz8B_</a>	Alignment	not modelled	99.5	37	<b>PDB header:</b> sh3 domain <b>Chain:</b> B: <b>PDB Molecule:</b> sh3-domain kinase binding protein 1; <b>PDBTitle:</b> n-terminal sh3 domain of cin85 bound to cbl-b peptide
97	<a href="#">d1wiea_</a>	Alignment	not modelled	99.5	22	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
98	<a href="#">c2oi3A_</a>	Alignment	not modelled	99.5	32	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> tyrosine-protein kinase hck; <b>PDBTitle:</b> nmr structure analysis of the hematopoietic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
99	<a href="#">c1w70A_</a>	Alignment	not modelled	99.5	35	<b>PDB header:</b> sh3 domain <b>Chain:</b> A: <b>PDB Molecule:</b> neutrophil cytosol factor 4; <b>PDBTitle:</b> sh3 domain of p40phox complexed with c-terminal polyproline2 region of p47phox
100	<a href="#">c1x2pA_</a>	Alignment	not modelled	99.5	31	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein arginine n-methyltransferase 2; <b>PDBTitle:</b> solution structure of the sh3 domain of the protein2 arginine n-methyltransferase 2
101	<a href="#">c2d8hA_</a>	Alignment	not modelled	99.5	28	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> sh3yl1 protein; <b>PDBTitle:</b> solution structure of the sh3 domain of hypothetical2 protein sh3yl1
102	<a href="#">c2dl7A_</a>	Alignment	not modelled	99.5	33	<b>PDB header:</b> structural genomics, unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> kiaa0769 protein; <b>PDBTitle:</b> solution structure of the second sh3 domain of human2 kiaa0769 protein
103	<a href="#">d1jo8a_</a>	Alignment	not modelled	99.5	48	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
104	<a href="#">c2drmB_</a>	Alignment	not modelled	99.5	50	<b>PDB header:</b> contractile protein <b>Chain:</b> B: <b>PDB Molecule:</b> acanthamoeba myosin ib; <b>PDBTitle:</b> acanthamoeba myosin i sh3 domain bound to acan125
105	<a href="#">c2dl3A_</a>	Alignment	not modelled	99.5	25	<b>PDB header:</b> cell adhesion, signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sorbin and sh3 domain-containing protein 1; <b>PDBTitle:</b> solution structure of the first sh3 domain of human sorbin2 and sh3 domain-containing protein 1
106	<a href="#">c4igzA_</a>	Alignment	not modelled	99.5	38	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> sorbin and sh3 domain-containing protein 2; <b>PDBTitle:</b> crystal structure of the sh3 domain of human sorbin and sh3 domain-2 containing protein 2
107	<a href="#">c2dhnA_</a>	Alignment	not modelled	99.5	30	<b>PDB header:</b> transferase, signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3-containing grb2-like protein 2;

107	<a href="#">c2g0m1A_</a>	Alignment	not modelled	99.5	39	<b>PDBTitle:</b> solution structures of the sh3 domain of human sh3-2 containing grb2-like protein 2
108	<a href="#">c2js0A_</a>	Alignment	not modelled	99.5	37	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cytoplasmic protein nck1; <b>PDBTitle:</b> solution structure of second sh3 domain of adaptor nck
109	<a href="#">c1x43A_</a>	Alignment	not modelled	99.5	25	<b>PDB header:</b> endocytosis/exocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 domain grb2-like protein b1; <b>PDBTitle:</b> solution structure of the sh3 domain of endophilin b12 (sh3g1b1)
110	<a href="#">c2pqhA_</a>	Alignment	not modelled	99.5	28	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> spectrin alpha chain, brain; <b>PDBTitle:</b> structure of sh3 chimera with a type ii ligand linked to the chain c-2 terminal
111	<a href="#">c2eqiA_</a>	Alignment	not modelled	99.5	37	<b>PDB header:</b> immune system, hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> phospholipase c, gamma 2; <b>PDBTitle:</b> solution structure of the sh3 domain from phospholipase c,2 gamma 2
112	<a href="#">d1t0ha_</a>	Alignment	not modelled	99.5	23	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
113	<a href="#">d1qlya_</a>	Alignment	not modelled	99.5	29	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
114	<a href="#">c2krnA_</a>	Alignment	not modelled	99.5	42	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cd2-associated protein; <b>PDBTitle:</b> high resolution structure of the second sh3 domain of cd2ap
115	<a href="#">c4afzC_</a>	Alignment	not modelled	99.5	30	<b>PDB header:</b> hydrolase/de novo protein <b>Chain:</b> C: <b>PDB Molecule:</b> fynomer; <b>PDBTitle:</b> human chymase - fynomer complex
116	<a href="#">c2cucA_</a>	Alignment	not modelled	99.5	31	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 domain containing ring finger 2; <b>PDBTitle:</b> solution structure of the sh3 domain of the mouse2 hypothetical protein sh3rf2
117	<a href="#">d1e6ga_</a>	Alignment	not modelled	99.5	25	<b>Fold:</b> SH3-like barrel <b>Superfamily:</b> SH3-domain <b>Family:</b> SH3-domain
118	<a href="#">c1yn8E_</a>	Alignment	not modelled	99.5	34	<b>PDB header:</b> unknown function <b>Chain:</b> E: <b>PDB Molecule:</b> nap1-binding protein 2; <b>PDBTitle:</b> sh3 domain of yeast nbp2
119	<a href="#">c2egaA_</a>	Alignment	not modelled	99.5	25	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> sh3 and px domain-containing protein 2a; <b>PDBTitle:</b> solution structure of the first sh3 domain from human2 kiaa0418 protein
120	<a href="#">c1x69A_</a>	Alignment	not modelled	99.5	33	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cortactin isoform a; <b>PDBTitle:</b> solution structures of the sh3 domain of human src2 substrate cortactin