
















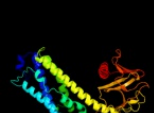





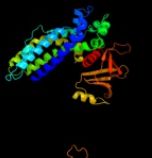
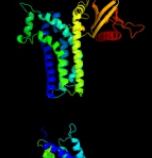

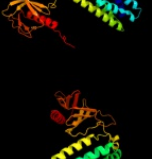
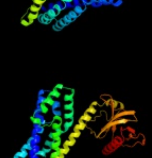


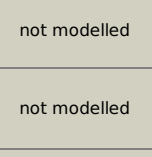






Phyre2

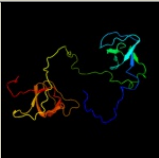
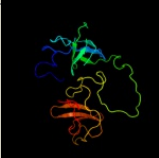
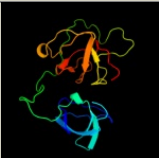
Email	l.a.kelley@imperial.ac.uk
Description	Q9Z0R4
Date	Tue Jul 30 13:06:10 BST 2013
Unique Job ID	298496d1861f1601

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2dx1A_	 Alignment		100.0	22	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of rhogef protein asef
2	c3jv3A_	 Alignment		100.0	99	PDB header: protein binding Chain: A; PDB Molecule: intersectin-1; PDBTitle: structure of sh3e-dh unit of murine intersectin-1l
3	c3odoA_	 Alignment		100.0	26	PDB header: signaling protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the dh/ph domains of p115-rhogef
4	c1x86A_	 Alignment		100.0	27	PDB header: signaling protein/membrane protein Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 12; PDBTitle: crystal structure of the dh/ph domains of leukemia-associated rhogef2 in complex with rhoa
5	c1ki1B_	 Alignment		100.0	94	PDB header: signaling protein Chain: B; PDB Molecule: intersectin long form; PDBTitle: guanine nucleotide exchange region of intersectin in2 complex with cdc42
6	c3ky9B_	 Alignment		100.0	19	PDB header: apoptosis Chain: B; PDB Molecule: proto-oncogene vav; PDBTitle: autoinhibited vav1
7	c1xcqA_	 Alignment		100.0	28	PDB header: signaling protein activator/signaling pr Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of human rhoa in complex with dh/ph2 fragment of pdzrhogef
8	c3ksyA_	 Alignment		100.0	18	PDB header: signaling protein Chain: A; PDB Molecule: son of sevenless homolog 1; PDBTitle: crystal structure of the histone domain, dh-ph unit, and2 catalytic unit of the ras activator son of sevenless (sos)
9	c3t06E_	 Alignment		100.0	28	PDB header: signaling protein Chain: E; PDB Molecule: rho guanine nucleotide exchange factor 11; PDBTitle: crystal structure of the dh/ph fragment of pdzrhogef with n-terminal2 regulatory elements in complex with human rhoa
10	c2vrwB_	 Alignment		100.0	21	PDB header: signaling protein Chain: B; PDB Molecule: proto-oncogene vav; PDBTitle: critical structural role for the ph and c1 domains of the2 vav1 exchange factor
11	c2dfkA_	 Alignment		100.0	20	PDB header: cell cycle Chain: A; PDB Molecule: collybistin ii; PDBTitle: crystal structure of the cdc42-collybistin ii complex

12	c4gouA	Alignment		100.0	15	PDB header: signaling protein Chain: A: PDB Molecule: ehrgs-rhogef; PDBTitle: crystal structure of an rgs-rhogef from entamoeba histolytica
13	c1xdvB	Alignment		100.0	17	PDB header: signaling protein Chain: B: PDB Molecule: son of sevenless protein homolog 1; PDBTitle: experimentally phased structure of human the son of2 sevenless protein at 4.1 ang.
14	c1foeA	Alignment		100.0	21	PDB header: signaling protein, immune system/signali Chain: A: PDB Molecule: t-lymphoma invasion and metastasis inducing PDBTitle: crystal structure of rac1 in complex with the guanine2 nucleotide exchange region of tiam1
15	c2z0qA	Alignment		100.0	28	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 3; PDBTitle: crystal structure of dh-ph domain of rhogef3(xpln)
16	c3bjjA	Alignment		100.0	19	PDB header: signaling protein Chain: A: PDB Molecule: proto-oncogene vav; PDBTitle: structural basis of promiscuous guanine nucleotide exchange2 by the t-cell essential vav1
17	c4gzuB	Alignment		100.0	24	PDB header: signaling protein Chain: B: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal structure of the dh-ph-ph domain of farp2
18	c30dwB	Alignment		100.0	26	PDB header: signaling protein Chain: B: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of the linker-dh/ph domains of p115-rhogef
19	c30dxA	Alignment		100.0	27	PDB header: signaling protein Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 1; PDBTitle: crystal structure of an n-terminally truncated linker-dh/ph domains of2 p115-rhogef
20	c4h6yA	Alignment		100.0	23	PDB header: signaling protein Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 1; PDBTitle: crystal strucure of the dh-ph-ph domain of farp1
21	c1dbhA	Alignment	not modelled	100.0	16	PDB header: gene regulation Chain: A: PDB Molecule: protein (human sos 1); PDBTitle: dbl and pleckstrin homology domains from hsos1
22	c2rgnB	Alignment	not modelled	100.0	17	PDB header: signaling protein complex Chain: B: PDB Molecule: rhoa/rac/cdc42 exchange factor; PDBTitle: crystal structure of p63rhogef complex with galpha-q and rhoa
23	c1lb1C	Alignment	not modelled	100.0	19	PDB header: signaling protein Chain: C: PDB Molecule: guanine nucleotide exchange factor dbs; PDBTitle: crystal structure of the dbl and pleckstrin homology2 domains of dbs in complex with rhoa
24	c3mpxA	Alignment	not modelled	100.0	20	PDB header: lipid binding protein Chain: A: PDB Molecule: fyve, rhogef and ph domain-containing protein 5; PDBTitle: crystal structure of the dh and ph-1 domains of human fgd5
25	c1ntyA	Alignment	not modelled	100.0	21	PDB header: signaling protein Chain: A: PDB Molecule: triple functional domain protein; PDBTitle: crystal structure of the first dh/ph domain of trio to 1.7 a
26	d1xcga1	Alignment	not modelled	100.0	30	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
27	d1txda1	Alignment	not modelled	100.0	28	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
28	d1ki1b1	Alignment	not modelled	100.0	99	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
						PDB header: signaling protein

29	c4gyvA	Alignment	not modelled	99.9	21	Chain: A: PDB Molecule: ferm, rhogef and pleckstrin domain-containing protein 2; PDBTitle: crystal structure of the dh domain of farp2
30	d2dfka1	Alignment	not modelled	99.9	20	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
31	d1dbha1	Alignment	not modelled	99.9	16	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
32	c3eo2A	Alignment	not modelled	99.9	32	PDB header: splicing Chain: A: PDB Molecule: neuroepithelial cell-transforming gene 1 protein; PDBTitle: crystal structure of the rhogef domain of human neuroepithelial cell-2 transforming gene 1 protein
33	d1foea1	Alignment	not modelled	99.9	25	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
34	c3gf9A	Alignment	not modelled	99.9	80	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 rhogef domain
35	d1by1a	Alignment	not modelled	99.9	20	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
36	d1f5xa	Alignment	not modelled	99.9	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
37	d1kz7a1	Alignment	not modelled	99.9	23	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
38	d1nty1	Alignment	not modelled	99.9	24	Fold: DBL homology domain (DH-domain) Superfamily: DBL homology domain (DH-domain) Family: DBL homology domain (DH-domain)
39	c2khnA	Alignment		99.9	96	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.
40	c3fiaA	Alignment		99.8	92	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.
41	c2jq6A	Alignment	not modelled	99.8	36	PDB header: metal binding protein Chain: A: PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of eh-domain of ehd1
42	c2kfhA	Alignment	not modelled	99.7	38	PDB header: protein binding Chain: A: PDB Molecule: eh domain-containing protein 1; PDBTitle: structure of the c-terminal domain of ehd1 with fnyestgpftak
43	d2jxca1	Alignment	not modelled	99.7	36	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
44	c2qptA	Alignment	not modelled	99.7	33	PDB header: endocytosis Chain: A: PDB Molecule: eh domain-containing protein-2; PDBTitle: crystal structure of an ehd atpase involved in membrane remodelling
45	c2kgrA	Alignment	not modelled	99.6	37	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
46	d1iq3a	Alignment	not modelled	99.6	38	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
47	d1fi6a	Alignment	not modelled	99.6	40	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
48	d1c07a	Alignment	not modelled	99.6	36	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
49	c1griA	Alignment		99.5	25	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
50	c3jzyA	Alignment		99.5	73	PDB header: endocytosis Chain: A: PDB Molecule: intersectin 2; PDBTitle: crystal structure of human intersectin 2 c2 domain
51	c2lq1A	Alignment	not modelled	99.5	19	PDB header: metal binding protein Chain: A: PDB Molecule: a-kinase anchor protein 13; PDBTitle: solution structure of the human akap13 ph domain and stabilizing dh2 helix

52	d1qjta_	Alignment	not modelled	99.4	22	Fold: EF Hand-like Superfamily: EF-hand Family: Eps15 homology domain (EH domain)
53	d1ki1b2	Alignment	not modelled	99.4	89	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
54	c2lqnA_	Alignment		99.4	23	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structure of crkl
55	c2eyzA_	Alignment		99.4	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crak sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform ii
56	c2fk9A_	Alignment	not modelled	99.4	24	PDB header: transferase Chain: A: PDB Molecule: protein kinase c, eta type; PDBTitle: human protein kinase c, eta
57	c2l3sA_	Alignment		99.3	25	PDB header: structural protein Chain: A: PDB Molecule: autoinhibited crk protein; PDBTitle: structure of the autoinhibited crk
58	c2nq3A_	Alignment	not modelled	99.2	24	PDB header: ligase Chain: A: PDB Molecule: itchy homolog e3 ubiquitin protein ligase; PDBTitle: crystal structure of the c2 domain of human itchy homolog2 e3 ubiquitin protein ligase
59	d2nq3a1	Alignment	not modelled	99.2	24	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
60	c1mv3A_	Alignment	not modelled	99.2	14	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
61	c1ng2A_	Alignment	not modelled	99.2	27	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosolic factor 1; PDBTitle: structure of autoinhibited p47phox
62	c1ov3A_	Alignment	not modelled	99.2	26	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 1; PDBTitle: structure of the p22phox-p47phox complex
63	d1gmia_	Alignment	not modelled	99.1	25	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
64	c3nsjA_	Alignment	not modelled	99.1	25	PDB header: immune system Chain: A: PDB Molecule: perforin-1; PDBTitle: the x-ray crystal structure of lymphocyte perforin
65	d2ep6a1	Alignment	not modelled	99.0	33	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
66	d1bdya_	Alignment	not modelled	99.0	18	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
67	d1wfla_	Alignment	not modelled	99.0	21	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
68	c2dmhA_	Alignment	not modelled	99.0	23	PDB header: lipid binding protein Chain: A: PDB Molecule: myoferlin; PDBTitle: solution structure of the first c2 domain of human myoferlin
69	c3l9bA_	Alignment	not modelled	99.0	16	PDB header: membrane protein Chain: A: PDB Molecule: otoferlin; PDBTitle: crystal structure of rat otoferlin c2a
70	c2jqzA_	Alignment	not modelled	99.0	27	PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase smurf2; PDBTitle: solution structure of the c2 domain of human smurf2
71	c3kwtA_	Alignment	not modelled	99.0	28	PDB header: metal binding protein Chain: A: PDB Molecule: munc13-1; PDBTitle: munc13-1 c2b-domain, calcium-free
72	d1udla_	Alignment	not modelled	98.9	65	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
73	d2cjsa1	Alignment	not modelled	98.9	21	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
74	c3m7fB_	Alignment	not modelled	98.9	31	PDB header: signaling protein/ligase Chain: B: PDB Molecule: e3 ubiquitin-protein ligase nedd4; PDBTitle: crystal structure of the nedd4 c2/grb10 sh2 complex
75	d1rlwa_	Alignment	not modelled	98.9	18	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)

76	d1dsya_	Alignment	not modelled	98.9	29	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
77	c2enjA_	Alignment	not modelled	98.8	19	PDB header: transferase Chain: A: PDB Molecule: protein kinase c theta type; PDBTitle: solution structure of the c2 domain from human protein2 kinase c theta
78	c1dqva_	Alignment	not modelled	98.8	23	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: synaptotagmin iii; PDBTitle: crystal structure of synaptotagmin iii c2a/c2b
79	d1xcqa2	Alignment	not modelled	98.8	24	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
80	d1ng2a2	Alignment	not modelled	98.8	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
81	d1zc3b1	Alignment	not modelled	98.8	17	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
82	d1qasa2	Alignment	not modelled	98.7	22	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: PLC-like (P variant)
83	d2dfka2	Alignment	not modelled	98.7	21	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
84	c2r83B_	Alignment	not modelled	98.7	23	PDB header: endocytosis, exocytosis Chain: B: PDB Molecule: synaptotagmin-1; PDBTitle: crystal structure analysis of human synaptotagmin 1 c2a-c2b
85	c2d8ka_	Alignment	not modelled	98.7	27	PDB header: endocytosis/exocytosis Chain: A: PDB Molecule: synaptotagmin vii; PDBTitle: solution structure of the first c2 domain of synaptotagmin2 vii
86	c2nsgA_	Alignment	not modelled	98.7	27	PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase nedd4-like protein; PDBTitle: crystal structure of the c2 domain of the human e3 ubiquitin-protein2 ligase nedd4-like protein
87	c3qwmA_	Alignment	not modelled	98.7	21	PDB header: signaling protein Chain: A: PDB Molecule: iq motif and sec7 domain-containing protein 1; PDBTitle: crystal structure of gep100, the plextrin homology domain of iq motif2 and sec7 domain-containing protein 1 isoform a
88	d1txda2	Alignment	not modelled	98.7	25	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
89	d1dbha2	Alignment	not modelled	98.7	19	Fold: PH domain-like barrel Superfamily: PH domain-like Family: Pleckstrin-homology domain (PH domain)
90	c3nmzD_	Alignment	not modelled	98.7	25	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crystal structure of apc complexed with asef
91	c2chdA_	Alignment	not modelled	98.6	30	PDB header: protein transport Chain: A: PDB Molecule: rabphilin-3a; PDBTitle: crystal structure of the c2a domain of rabphilin-3a
92	c1x2qa_	Alignment	not modelled	98.6	31	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
93	d2r83a1	Alignment	not modelled	98.6	24	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
94	d1a25a_	Alignment	not modelled	98.6	28	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
95	c2cubA_	Alignment	not modelled	98.6	29	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1
96	c2rqrA_	Alignment	not modelled	98.6	7	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
97	c2eyyA_	Alignment	not modelled	98.6	23	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: ct10-regulated kinase isoform i
98	c2ed0A_	Alignment	not modelled	98.6	35	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)
99	c2enpA_	Alignment	not modelled	98.6	23	PDB header: transport protein Chain: A: PDB Molecule: b/k protein; PDBTitle: solution structure of the first c2 domain from human b/k2 protein
100	d1ov3a2	Alignment	not modelled	98.6	26	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
101	c2dbkA_	Alignment	not modelled	98.6	28	PDB header: signaling protein Chain: A: PDB Molecule: crk-like protein; PDBTitle: solution structures of the sh3 domain of human crk-like2 protein PDB header: signaling protein

102	c2ysqA_	Alignment	not modelled	98.6	24	Chain: A; PDB Molecule: rho guanine nucleotide exchange factor 9; PDBTitle: solution structure of the sh3 domain from rho guanine2 nucleotide exchange factor 9
103	c2jxbA_	Alignment	not modelled	98.6	29	PDB header: signaling protein complex Chain: A; PDB Molecule: t-cell surface glycoprotein cd3 epsilon chain, PDBTitle: structure of cd3epsilon-nck2 first sh3 domain complex
104	c2oi3A_	Alignment	not modelled	98.6	27	PDB header: transferase Chain: A; PDB Molecule: tyrosine-protein kinase hck; PDBTitle: nmr structure analysis of the hematopoetic cell kinase sh32 domain complexed with an artificial high affinity ligand3 (pd1)
105	c2yunA_	Alignment	not modelled	98.6	25	PDB header: protein transport Chain: A; PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
106	c2dlpA_	Alignment	not modelled	98.6	27	PDB header: structural protein Chain: A; PDB Molecule: kiaa1783 protein; PDBTitle: solution structure of the sh3 domain of human kiaa17832 protein
107	c2ebpA_	Alignment	not modelled	98.6	23	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
108	d1rsya_	Alignment	not modelled	98.5	28	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
109	c2l0aA_	Alignment	not modelled	98.5	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
110	c3fbkB_	Alignment	not modelled	98.5	22	PDB header: signaling protein Chain: B; PDB Molecule: regulator of g-protein signaling 3; PDBTitle: crystal structure of the c2 domain of the human regulator2 of g-protein signaling 3 isoform 6 (rgp3), northeast3 structural genomics consortium target hr5550a
111	d1wiea_	Alignment	not modelled	98.5	25	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	c2yt6A_	Alignment	not modelled	98.5	27	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
113	c2dl8A_	Alignment	not modelled	98.5	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
114	c2yupA_	Alignment	not modelled	98.5	22	PDB header: cell adhesion Chain: A; PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
115	c2kymA_	Alignment	not modelled	98.5	32	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
116	c2yu0A_	Alignment	not modelled	98.5	25	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
117	c2dnuA_	Alignment	not modelled	98.5	29	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
118	d1wfma_	Alignment	not modelled	98.5	20	Fold: C2 domain-like Superfamily: C2 domain (Calcium/lipid-binding domain, CaLB) Family: Synaptotagmin-like (S variant)
119	c2djqa_	Alignment	not modelled	98.5	24	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
120	d1k4us_	Alignment	not modelled	98.5	37	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain