



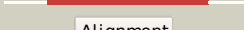

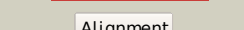

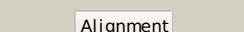







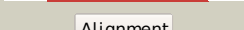

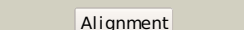

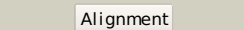



Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	Q46834
Date	Thu Jan 5 12:34:59 GMT 2012
Unique Job ID	ed051d7dd15b0254

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2i4sa1	 Alignment		99.8	38	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: EpsC C-terminal domain-like
2	d2i6va1	 Alignment		99.7	41	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: EpsC C-terminal domain-like
3	c3stjC_	 Alignment		99.7	15	PDB header: hydrolase Chain: C: PDB Molecule: protease degg; PDBTitle: crystal structure of the protease + pdz1 domain of degg from2 escherichia coli
4	c3qo6B_	 Alignment		99.6	12	PDB header: photosynthesis Chain: B: PDB Molecule: protease do-like 1, chloroplastic; PDBTitle: crystal structure analysis of the plant protease deg1
5	c3pv4A_	 Alignment		99.6	14	PDB header: hydrolase Chain: A: PDB Molecule: degq; PDBTitle: structure of legionella fallonii degq (delta-pdz2 variant)
6	c3pv5B_	 Alignment		99.5	16	PDB header: hydrolase Chain: B: PDB Molecule: degq; PDBTitle: structure of legionella fallonii degq (n189g/p190g variant)
7	c1lcyA_	 Alignment		99.5	11	PDB header: hydrolase Chain: A: PDB Molecule: htra2 serine protease; PDBTitle: crystal structure of the mitochondrial serine protease htra2
8	d2z9ia1	 Alignment		99.5	12	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: HtraA-like serine proteases
9	c3gdsA_	 Alignment		99.5	11	PDB header: hydrolase/hydrolase activator Chain: A: PDB Molecule: protease degs; PDBTitle: crystal structure of degs h198p/d320a mutant modified by dfp in2 complex with dnrdgnvyvf peptide
10	c1ky9A_	 Alignment		99.4	17	PDB header: hydrolase Chain: A: PDB Molecule: protease do; PDBTitle: crystal structure of degp (htra)
11	c2kl1A_	 Alignment		99.4	16	PDB header: protein binding Chain: A: PDB Molecule: ylbl protein; PDBTitle: solution structure of gtr34c from geobacillus thermodenitrificans.2 northeast structural genomics consortium target gtr34c

12	c2p3wB_	Alignment		99.4	21	PDB header: protein binding Chain: B: PDB Molecule: probable serine protease htra3; PDBTitle: crystal structure of the htra3 pdz domain bound to a phage-derived2 ligand (fgrwv)
13	d1lcyal	Alignment		99.4	11	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: HtrA-like serine proteases
14	c3i18A_	Alignment		99.4	13	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: lmo2051 protein; PDBTitle: crystal structure of the pdz domain of the sdrc-like protein2 (lmo2051) from listeria monocytogenes, northeast structural3 genomics consortium target lmr166b
15	c2kjpA_	Alignment		99.3	14	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: uncharacterized protein ylbl; PDBTitle: solution structure of protein ylbl (bsu15050) from bacillus2 subtilis, northeast structural genomics consortium target3 sr713a
16	c2joaA_	Alignment		99.3	18	PDB header: protein binding Chain: A: PDB Molecule: serine protease htra1; PDBTitle: htra1 bound to an optimized peptide: nmr assignment of pdz2 domain and ligand resonances
17	c3ossC_	Alignment		99.3	97	PDB header: protein transport Chain: C: PDB Molecule: type 2 secretion system, gspc; PDBTitle: the crystal structure of enterotoxigenic escherichia coli gspc-gspd2 complex from the type ii secretion system
18	d1ky9a1	Alignment		99.3	22	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: HtrA-like serine proteases
19	d1fc6a3	Alignment		99.3	12	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: Tail specific protease PDZ domain
20	d2hgaa1	Alignment		99.2	10	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: MTH1368 C-terminal domain-like
21	c2z9iB_	Alignment	not modelled	99.2	11	PDB header: hydrolase Chain: B: PDB Molecule: probable serine protease pepd; PDBTitle: crystal structure of rv0983 from mycobacterium tuberculosis-2 proteolytically active form
22	d1ky9b2	Alignment	not modelled	99.2	17	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: HtrA-like serine proteases
23	d1sota1	Alignment	not modelled	99.2	9	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: HtrA-like serine proteases
24	c3rleA_	Alignment	not modelled	99.1	10	PDB header: membrane protein Chain: A: PDB Molecule: golgi reassembly-stacking protein 2; PDBTitle: crystal structure of grasp55 grasp domain (residues 7-208)
25	c2zplA_	Alignment	not modelled	99.1	17	PDB header: hydrolase Chain: A: PDB Molecule: regulator of sigma e protease; PDBTitle: crystal structure analysis of pdz domain a
26	c2r3yC_	Alignment	not modelled	99.1	15	PDB header: hydrolase/hydrolase activator Chain: C: PDB Molecule: protease degs; PDBTitle: crystal structure of the degs protease in complex with the2 ywf activating peptide
27	c4a8al_	Alignment	not modelled	99.1	15	PDB header: hydrolase/hydrolase Chain: I: PDB Molecule: periplasmic ph-dependent serine endoprotease degg; PDBTitle: asymmetric cryo-em reconstruction of e. coli degg 12-mer in complex2 with lysozyme
28	c2zpmA_	Alignment	not modelled	99.0	19	PDB header: hydrolase Chain: A: PDB Molecule: regulator of sigma e protease; PDBTitle: crystal structure analysis of pdz domain b

29	c2eaqA	Alignment	not modelled	98.9	21	PDB header: metal binding protein Chain: A: PDB Molecule: lim domain only protein 7; PDBTitle: crystal structure of pdz domain of kiaa0858 (lim), ms07932 from homo sapiens
30	c2krqA	Alignment	not modelled	98.8	16	PDB header: signaling protein Chain: A: PDB Molecule: na(+)/h(+) exchange regulatory cofactor nhe-rf1; PDBTitle: solution structure of human sodium/ hydrogen exchange2 regulatory factor 1(150-358)
31	d1wfga	Alignment	not modelled	98.8	19	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
32	d1m5za	Alignment	not modelled	98.8	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
33	d1uf1a	Alignment	not modelled	98.8	27	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
34	c2eehA	Alignment	not modelled	98.8	20	PDB header: metal binding protein Chain: A: PDB Molecule: pdz domain-containing protein 7; PDBTitle: solution structure of first pdz domain of pdz domain2 containing protein 7
35	c2vwrA	Alignment	not modelled	98.8	21	PDB header: protein-binding Chain: A: PDB Molecule: ligand of numb protein x 2; PDBTitle: crystal structure of the second pdz domain of numb-binding2 protein 2
36	d1wifa	Alignment	not modelled	98.8	13	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
37	c3shuB	Alignment	not modelled	98.8	22	PDB header: cell adhesion Chain: B: PDB Molecule: tight junction protein zo-1; PDBTitle: crystal structure of zo-1 pdz3
38	c3l4fD	Alignment	not modelled	98.7	17	PDB header: signaling protein/protein binding Chain: D: PDB Molecule: sh3 and multiple ankyrin repeat domains protein PDBTitle: crystal structure of betapix coiled-coil domain and shank2 pdz complex
39	c2kjdA	Alignment	not modelled	98.7	14	PDB header: signaling protein Chain: A: PDB Molecule: sodium/hydrogen exchange regulatory cofactor nhe- PDBTitle: solution structure of extended pdz2 domain from nherf1 (150-2 270)
40	c3diwB	Alignment	not modelled	98.7	17	PDB header: signaling protein/cell adhesion Chain: B: PDB Molecule: tax1-binding protein 3; PDBTitle: c-terminal beta-catenin bound tip-1 structure
41	c3shwA	Alignment	not modelled	98.7	22	PDB header: cell adhesion Chain: A: PDB Molecule: tight junction protein zo-1; PDBTitle: crystal structure of zo-1 pdz3-sh3-guk supramodule complex with2 connexin-45 peptide
42	c2egkC	Alignment	not modelled	98.7	13	PDB header: protein binding Chain: C: PDB Molecule: general receptor for phosphoinositides 1- PDBTitle: crystal structure of tamalin pdz-intrinsic ligand fusion2 protein
43	c2komA	Alignment	not modelled	98.7	23	PDB header: signaling protein Chain: A: PDB Molecule: partitioning defective 3 homolog; PDBTitle: solution structure of humar par-3b pdz2 (residues 451-549)
44	d1ueqa	Alignment	not modelled	98.7	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
45	d1x5qa1	Alignment	not modelled	98.7	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
46	c2qt5A	Alignment	not modelled	98.7	12	PDB header: peptide binding protein Chain: A: PDB Molecule: glutamate receptor-interacting protein 1; PDBTitle: crystal structure of grip1 pdz12 in complex with the fras12 peptide
47	d1q3oa	Alignment	not modelled	98.7	18	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
48	d1rgwa	Alignment	not modelled	98.7	13	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
49	d1p1da2	Alignment	not modelled	98.7	14	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
50	d2cssa1	Alignment	not modelled	98.7	19	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
51	c2q3gA	Alignment	not modelled	98.7	11	PDB header: structural genomics Chain: A: PDB Molecule: pdz and lim domain protein 7; PDBTitle: structure of the pdz domain of human pdlim7 bound to a c-2 terminal extension from human beta-tropomyosin
52	d1qaua	Alignment	not modelled	98.6	14	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
53	d1t2ma1	Alignment	not modelled	98.6	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
54	c3khfA	Alignment	not modelled	98.6	18	PDB header: transferase Chain: A: PDB Molecule: microtubule-associated serine/threonine-protein

					PDBTitle: the crystal structure of the pdz domain of human microtubule2 associated serine/threonine kinase 3 (mast3)
55	d2f5ya1	Alignment	not modelled	98.6	14 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
56	d1wf7a_	Alignment	not modelled	98.6	12 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
57	d2fe5a1	Alignment	not modelled	98.6	17 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
58	c2he4A_	Alignment	not modelled	98.6	18 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: na(+)/h(+) exchange regulatory cofactor nhe-rf2; PDBTitle: the crystal structure of the second pdz domain of human2 nherf-2 (slc9a3r2) interacting with a mode 1 pdz binding3 motif
59	d1g9oa_	Alignment	not modelled	98.6	14 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
60	d1w9ea1	Alignment	not modelled	98.6	15 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
61	c2v1wB_	Alignment	not modelled	98.6	18 PDB header: structural protein Chain: B: PDB Molecule: pdz and lim domain protein 4; PDBTitle: crystal structure of human lim protein ril (pdlim4) pdz2 domain bound to the c-terminal peptide of human alpha-3 actinin-1
62	d1ozia_	Alignment	not modelled	98.6	16 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
63	d1p1da1	Alignment	not modelled	98.6	14 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
64	c1p1dA_	Alignment	not modelled	98.6	14 PDB header: protein binding Chain: A: PDB Molecule: glutamate receptor interacting protein; PDBTitle: structural insights into the inter-domain chaperoning of2 tandem pdz domains in glutamate receptor interacting3 proteins
65	d1ry4a_	Alignment	not modelled	98.6	17 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
66	c2d90A_	Alignment	not modelled	98.6	22 PDB header: protein binding Chain: A: PDB Molecule: pdz domain containing protein 1; PDBTitle: solution structure of the third pdz domain of pdz domain2 containing protein 1
67	c3qglD_	Alignment	not modelled	98.6	20 PDB header: protein binding Chain: D: PDB Molecule: sorting nexin-27; PDBTitle: crystal structure of pdz domain of sorting nexin 27 (snx27) in complex2 with the eseskv peptide corresponding to the c-terminal tail of girk3
68	c2gzvA_	Alignment	not modelled	98.6	9 PDB header: signaling protein Chain: A: PDB Molecule: prkca-binding protein; PDBTitle: the cystal structure of the pdz domain of human pick1 (casp target)
69	c2omjA_	Alignment	not modelled	98.6	17 PDB header: cell adhesion Chain: A: PDB Molecule: rho guanine nucleotide exchange factor 12; PDBTitle: solution structure of larg pdz domain
70	c1u38A_	Alignment	not modelled	98.6	12 PDB header: protein transport Chain: A: PDB Molecule: amyloid beta a4 precursor protein-binding, PDBTitle: auto-inhibition mechanism of x11s/mints family scaffold2 proteins revealed by the closed conformation of the tandem3 pdz domains
71	d1b8qa_	Alignment	not modelled	98.6	15 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
72	c3eggC_	Alignment	not modelled	98.6	11 PDB header: hydrolase Chain: C: PDB Molecule: spinophilin; PDBTitle: crystal structure of a complex between protein phosphatase 1 alpha2 (pp1) and the pp1 binding and pdz domains of spinophilin
73	c3k1rA_	Alignment	not modelled	98.6	13 PDB header: structural protein Chain: A: PDB Molecule: harmonin; PDBTitle: structure of harmonin npdz1 in complex with the sam-pbm of2 sans
74	d1pdra_	Alignment	not modelled	98.6	17 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
75	d1be9a_	Alignment	not modelled	98.6	15 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
76	c2v90E_	Alignment	not modelled	98.6	15 PDB header: protein-binding Chain: E: PDB Molecule: pdz domain-containing protein 3; PDBTitle: crystal structure of the 3rd pdz domain of intestine- and2 kidney-enriched pdz domain ikepp (pdzd3)
77	c2jxoA_	Alignment	not modelled	98.6	16 PDB header: protein binding Chain: A: PDB Molecule: ezrin-radixin-moesin-binding phosphoprotein 50; PDBTitle: structure of the second pdz domain of nherf-1
78	d1ihja_	Alignment	not modelled	98.6	27 Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
79	c2iwnA_	Alignment	not modelled	98.6	21 PDB header: signaling protein Chain: A: PDB Molecule: multiple pdz domain protein; PDBTitle: 3rd pdz domain of multiple pdz domain protein mpdz (casp2 target)

80	c2vsvB	Alignment	not modelled	98.6	16	PDB header: protein-binding Chain: B: PDB Molecule: rhophilin-2; PDBTitle: crystal structure of the pdz domain of human rhophilin-2
81	d1d5ga	Alignment	not modelled	98.6	14	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
82	c1u37A	Alignment	not modelled	98.6	12	PDB header: protein transport Chain: A: PDB Molecule: amyloid beta a4 precursor protein-binding, PDBTitle: auto-inhibition mechanism of x11s/mints family scaffold2 proteins revealed by the closed conformation of the tandem3 pdz domains
83	c2i04B	Alignment	not modelled	98.6	16	PDB header: peptide binding protein Chain: B: PDB Molecule: membrane-associated guanylate kinase, ww and pdz PDBTitle: x-ray crystal structure of magi-1 pdz1 bound to the c-2 terminal peptide of hpv18 e6
84	c2ogpA	Alignment	not modelled	98.5	23	PDB header: signaling protein Chain: A: PDB Molecule: partitioning-defective 3 homolog; PDBTitle: solution structure of the second pdz domain of par-3
85	c2o2tB	Alignment	not modelled	98.5	26	PDB header: structural protein Chain: B: PDB Molecule: multiple pdz domain protein; PDBTitle: the crystal structure of the 1st pdz domain of mpdz
86	d2byga1	Alignment	not modelled	98.5	14	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
87	c2ka9A	Alignment	not modelled	98.5	16	PDB header: cell adhesion Chain: A: PDB Molecule: disks large homolog 4; PDBTitle: solution structure of psd-95 pdz12 complexed with cypin2 peptide
88	d1q7xa	Alignment	not modelled	98.5	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
89	c2q9vA	Alignment	not modelled	98.5	16	PDB header: transferase Chain: A: PDB Molecule: membrane-associated guanylate kinase, ww and pdz domain- PDBTitle: crystal structure of the c890s mutant of the 4th pdz domain of human2 membrane associated guanylate kinase
90	d1uepa	Alignment	not modelled	98.5	17	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
91	c2z17A	Alignment	not modelled	98.5	15	PDB header: protein binding Chain: A: PDB Molecule: pleckstrin homology sec7 and coiled-coil domains- PDBTitle: crystal sturcture of pdz domain from human pleckstrin2 homology, sec7
92	d1wh1a	Alignment	not modelled	98.5	21	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
93	c2iwoA	Alignment	not modelled	98.5	14	PDB header: signaling protein Chain: A: PDB Molecule: multiple pdz domain protein; PDBTitle: 12th pdz domain of multiple pdz domain protein mpdz (casp2 target)
94	c2qg1A	Alignment	not modelled	98.5	27	PDB header: signaling protein Chain: A: PDB Molecule: multiple pdz domain protein; PDBTitle: crystal structure of the 11th pdz domain of mpdz (mupp1)
95	d1uhpa	Alignment	not modelled	98.5	20	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
96	c2k1zA	Alignment	not modelled	98.5	23	PDB header: signaling protein Chain: A: PDB Molecule: partitioning-defective 3 homolog; PDBTitle: solution structure of par-3 pdz3
97	d1um1a	Alignment	not modelled	98.5	17	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
98	c2dm8A	Alignment	not modelled	98.5	16	PDB header: protein binding Chain: A: PDB Molecule: inad-like protein; PDBTitle: solution structure of the eighth pdz domain of human inad-2 like protein
99	d1wf8a1	Alignment	not modelled	98.5	11	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
100	c3bpuA	Alignment	not modelled	98.5	25	PDB header: transferase Chain: A: PDB Molecule: membrane-associated guanylate kinase, ww and pdz domain- PDBTitle: crystal structure of the 3rd pdz domain of human membrane associated2 guanylate kinase, c677s and c709s double mutant
101	c3cyyA	Alignment	not modelled	98.5	22	PDB header: peptide binding protein Chain: A: PDB Molecule: tight junction protein zo-1; PDBTitle: the crystal structure of zo-1 pdz2 in complex with the cx43 peptide
102	c2ji1A	Alignment	not modelled	98.5	11	PDB header: membrane protein Chain: A: PDB Molecule: glutamate receptor interacting protein-1; PDBTitle: crystal structure of 2nd pdz domain of glutamate receptor2 interacting protein-1 (grip1)
103	d2f0aa1	Alignment	not modelled	98.5	21	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
104	d1tp5a1	Alignment	not modelled	98.5	17	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain

105	d1ueza_	Alignment	not modelled	98.5	20	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
106	c2jikB_	Alignment	not modelled	98.5	21	PDB header: membrane protein Chain: B: PDB Molecule: synaptojanin-2 binding protein; PDBTitle: crystal structure of pdz domain of synaptojanin-2 binding2 protein
107	c2osgB_	Alignment	not modelled	98.5	26	PDB header: cell adhesion Chain: B: PDB Molecule: tight junction protein zo-2; PDBTitle: solution structure and binding property of the domain-2 swapped dimer of zo2pdz2
108	c3r0hA_	Alignment	not modelled	98.5	12	PDB header: peptide binding protein Chain: A: PDB Molecule: inactivation-no-after-potential d protein; PDBTitle: structure of inad pdz45 in complex with ng2 peptide
109	d1rzxa_	Alignment	not modelled	98.5	16	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
110	c3ggeA_	Alignment	not modelled	98.5	13	PDB header: protein binding Chain: A: PDB Molecule: pdz domain-containing protein gipc2; PDBTitle: crystal structure of the pdz domain of pdz domain-containing protein2 gipc2
111	c2edzA_	Alignment	not modelled	98.5	19	PDB header: signaling protein Chain: A: PDB Molecule: pdz domain-containing protein 1; PDBTitle: solution structures of the pdz domain of mus musculus pdz2 domain-containing protein 1
112	d1nf3c_	Alignment	not modelled	98.5	15	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
113	d1whaa_	Alignment	not modelled	98.5	22	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
114	c2dkrA_	Alignment	not modelled	98.5	16	PDB header: protein transport Chain: A: PDB Molecule: lin-7 homolog b; PDBTitle: solution structure of the pdz domain from human lin-72 homolog b
115	c3hpmA_	Alignment	not modelled	98.5	10	PDB header: protein binding Chain: A: PDB Molecule: protein interacting with prkca 1; PDBTitle: oxidized dimeric pick1 pdz c46g mutant in complex with the carboxyl2 tail peptide of glur2
116	d1kwaa_	Alignment	not modelled	98.5	13	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
117	c2db5A_	Alignment	not modelled	98.5	29	PDB header: protein binding Chain: A: PDB Molecule: inad-like protein; PDBTitle: solution structure of the first pdz domain of inad-like2 protein
118	d1x5ra1	Alignment	not modelled	98.5	15	Fold: PDZ domain-like Superfamily: PDZ domain-like Family: PDZ domain
119	c2djtA_	Alignment	not modelled	98.5	14	PDB header: signaling protein Chain: A: PDB Molecule: unnamed protein product; PDBTitle: solution structures of the pdz domain of human unnamed2 protein product
120	c2yuyA_	Alignment	not modelled	98.5	22	PDB header: signaling protein Chain: A: PDB Molecule: rho gtpase activating protein 21; PDBTitle: solution structure of pdz domain of rho gtpase activating2 protein 21