

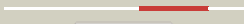


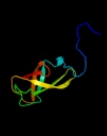






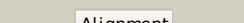






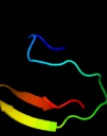





#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2k14A_	 Alignment		98.7	14	PDB header: unknown function Chain: A: PDB Molecule: yuaf protein; PDBTitle: solution structure of the soluble domain of the nfd2 protein yuaf from bacillus subtilis
2	c3cp0A_	 Alignment		97.6	15	PDB header: membrane protein Chain: A: PDB Molecule: membrane protein implicated in regulation of membrane PDBTitle: crystal structure of the soluble domain of membrane protein implicated2 in regulation of membrane protease activity from corynebacterium3 glutamicum
3	c2k5hA_	 Alignment		97.6	16	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: conserved protein; PDBTitle: solution nmr structure of protein encoded by mth693 from2 methanobacterium thermoautotrophicum: northeast structural3 genomics consortium target tt824a
4	d2exda1	 Alignment		96.5	15	Fold: OB-fold Superfamily: NfeD domain-like Family: NfeD domain-like
5	c1l2fA_	 Alignment		69.4	15	PDB header: transcription Chain: A: PDB Molecule: n utilization substance protein a; PDBTitle: crystal structure of nusa from thermotoga maritima: a2 structure-based role of the n-terminal domain
6	c2ebpA_	 Alignment		69.3	19	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
7	c2d8jA_	 Alignment		61.5	20	PDB header: transferase Chain: A: PDB Molecule: fyn-related kinase; PDBTitle: solution structure of the sh3 domain of fyn-related kinase
8	d1wfwA_	 Alignment		60.9	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
9	c1hh2P_	 Alignment		59.6	15	PDB header: transcription regulation Chain: P: PDB Molecule: n utilization substance protein a; PDBTitle: crystal structure of nusa from thermotoga maritima
10	c2ekhA_	 Alignment		58.7	9	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
11	d1wlpb1	 Alignment		57.2	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

12	c3dclC_	Alignment		56.7	16	PDB header: structural genomics, unknown function Chain: C: PDB Molecule: tm1086; PDBTitle: crystal structure of tm1086
13	d1e6ha_	Alignment		51.9	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
14	c2dmoA_	Alignment		51.5	16	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)
15	c1x2pA_	Alignment		50.9	25	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
16	d1utia_	Alignment		49.5	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
17	c2djgA_	Alignment		47.8	29	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
18	c1griA_	Alignment		47.6	9	PDB header: signal transduction adaptor Chain: A: PDB Molecule: growth factor bound protein 2; PDBTitle: grb2
19	d2fo0a1	Alignment		47.5	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
20	c2xmfA_	Alignment		46.3	13	PDB header: motor protein Chain: A: PDB Molecule: myosin 1e sh3; PDBTitle: myosin 1e sh3
21	d1u06a1	Alignment	not modelled	45.7	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
22	d1k4us_	Alignment	not modelled	45.3	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
23	c3nmzD_	Alignment	not modelled	44.2	9	PDB header: cell adhesion/cell cycle Chain: D: PDB Molecule: rho guanine nucleotide exchange factor 4; PDBTitle: crytal structure of apc complexed with asef
24	d1efna_	Alignment	not modelled	43.9	23	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
25	c2drmB_	Alignment	not modelled	43.4	19	PDB header: contractile protein Chain: B: PDB Molecule: acanthamoeba myosin 1b; PDBTitle: acanthamoeba myosin i sh3 domain bound to acan125
26	c2egcA_	Alignment	not modelled	42.9	17	PDB header: signaling protein Chain: A: PDB Molecule: sh3 and px domain-containing protein 2a; PDBTitle: solution structure of the fifth sh3 domain from human2 kiaa0418 protein
27	c2dl4A_	Alignment	not modelled	42.7	9	PDB header: signaling protein Chain: A: PDB Molecule: protein stac; PDBTitle: solution structure of the first sh3 domain of stac protein
28	c2cubA_	Alignment	not modelled	42.5	25	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of the sh3 domain of the human2 cytoplasmic protein nck1

29	d1opka1	Alignment	not modelled	42.5	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
30	c2ysqA	Alignment	not modelled	42.3	9	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
31	d1jo8a	Alignment	not modelled	41.1	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
32	c1x69A	Alignment	not modelled	40.5	9	PDB header: signaling protein Chain: A: PDB Molecule: cortactin isoform a; PDBTitle: solution structures of the sh3 domain of human src2 substrate cortactin
33	c2ed0A	Alignment	not modelled	40.5	19	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)
34	d1oeba	Alignment	not modelled	40.3	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
35	c2kymA	Alignment	not modelled	40.3	13	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
36	c2ct4A	Alignment	not modelled	40.0	6	PDB header: signaling protein Chain: A: PDB Molecule: cdc42-interacting protein 4; PDBTitle: solution strcture of the sh3 domain of the cdc42-2 interacting protein 4
37	d1hh2p1	Alignment	not modelled	39.8	15	Fold: OB-fold Superfamily: Nucleic acid-binding proteins Family: Cold shock DNA-binding domain-like
38	c2dl8A	Alignment	not modelled	39.7	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
39	c3m3hA	Alignment	not modelled	39.2	18	PDB header: transferase Chain: A: PDB Molecule: orotate phosphoribosyltransferase; PDBTitle: 1.75 angstrom resolution crystal structure of an orotate2 phosphoribosyltransferase from bacillus anthracis str. 'ames3 ancestor'
40	c2jmcA	Alignment	not modelled	39.1	16	PDB header: signaling protein Chain: A: PDB Molecule: spectrin alpha chain, brain and p41 peptide PDBTitle: chimera between spc-sh3 and p41
41	d1ov3a1	Alignment	not modelled	38.6	9	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
42	d1ujya	Alignment	not modelled	38.3	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
43	c1x2qA	Alignment	not modelled	38.1	16	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
44	d2pyta1	Alignment	not modelled	37.4	17	Fold: Double-stranded beta-helix Superfamily: RmlC-like cupins Family: EutQ-like
45	c2epdA	Alignment	not modelled	37.3	22	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
46	d1ov3a2	Alignment	not modelled	37.2	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
47	c2yuoA	Alignment	not modelled	37.0	23	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
48	c2dbmA	Alignment	not modelled	36.9	19	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
49	d1ue9a	Alignment	not modelled	36.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
50	c2yunA	Alignment	not modelled	36.5	19	PDB header: protein transport Chain: A: PDB Molecule: nostrin; PDBTitle: solution structure of the sh3 domain of human nostrin
51	d1h8ka	Alignment	not modelled	36.3	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
52	c2ed1A	Alignment	not modelled	36.3	8	PDB header: signaling protein Chain: A: PDB Molecule: 130 kda phosphatidylinositol 4,5-biphosphate- PDBTitle: solution structure of the sh3 domain of 130 kda2 phosphatidylinositol 4,5-biphosphate-dependent arf1 gtpase-3 activating protein
53	c2ct3A	Alignment	not modelled	36.2	13	PDB header: signaling protein Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the sh3 domain of the vinexin protein
54	c1u3oA	Alignment	not modelled	36.0	19	PDB header: signaling protein Chain: A: PDB Molecule: huntingtin-associated protein-interacting

						PDBTitle: solution structure of rat kalirin n-terminal sh3 domain
55	c2dnuA	Alignment	not modelled	36.0	16	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
56	c3zvK_G	Alignment	not modelled	35.8	19	PDB header: antitoxin/toxin/dna Chain: G: PDB Molecule: antitoxin of toxin-antitoxin system vapb; PDBTitle: crystal structure of vapbc2 from rickettsia felis bound to2 a dna fragment from their promoter
57	c2fpeB	Alignment	not modelled	35.5	19	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
58	c1h9sA	Alignment	not modelled	35.1	19	PDB header: transcription regulator Chain: A: PDB Molecule: molybdenum transport protein mode; PDBTitle: molybdate bound complex of dimop domain of mode from e.coli
59	d1nega	Alignment	not modelled	34.9	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
60	c1negA	Alignment	not modelled	34.9	16	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
61	d1qlva	Alignment	not modelled	34.8	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
62	d1gria2	Alignment	not modelled	34.8	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
63	c1x6gA	Alignment	not modelled	34.7	23	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.
64	c2yupA	Alignment	not modelled	34.6	19	PDB header: cell adhesion Chain: A: PDB Molecule: vinexin; PDBTitle: solution structure of the second sh3 domain of human vinexin
65	d1wiea	Alignment	not modelled	34.5	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
66	d1pwta	Alignment	not modelled	34.2	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
67	d1udla	Alignment	not modelled	34.1	19	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
68	c2rqra	Alignment	not modelled	34.1	13	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
69	d1sema	Alignment	not modelled	34.0	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
70	d1ug1a	Alignment	not modelled	34.0	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
71	c3zv0D	Alignment	not modelled	33.9	10	PDB header: cell cycle Chain: D: PDB Molecule: h/aca ribonucleoprotein complex subunit 4; PDBTitle: structure of the shq1p-cbf5p complex
72	d2q07a1	Alignment	not modelled	33.7	14	Fold: PUA domain-like Superfamily: PUA domain-like Family: PUA domain
73	d1uj0a	Alignment	not modelled	33.6	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
74	c1zuyB	Alignment	not modelled	33.5	18	PDB header: contractile protein Chain: B: PDB Molecule: myosin-5 isoform; PDBTitle: high-resolution structure of yeast myo5 sh3 domain
75	c1zx6A	Alignment	not modelled	33.1	25	PDB header: protein binding Chain: A: PDB Molecule: ypr154wp; PDBTitle: high-resolution crystal structure of yeast pin3 sh3 domain
76	c2dl5A	Alignment	not modelled	33.0	6	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: kiaa0769 protein; PDBTitle: solution structure of the first sh3 domain of human2 kiaa0769 protein
77	c1w70A	Alignment	not modelled	32.7	13	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
78	d1uuea	Alignment	not modelled	32.7	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
79	c1zlmA	Alignment	not modelled	32.5	13	PDB header: signaling protein Chain: A: PDB Molecule: osteoclast stimulating factor 1; PDBTitle: crystal structure of the sh3 domain of human osteoclast2 stimulating factor
80	d1e6ga	Alignment	not modelled	32.3	16	Fold: SH3-like barrel Superfamily: SH3-domain

						Family: SH3-domain
81	c2pqhA	Alignment	not modelled	32.3	16	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
82	d1ng2a1	Alignment	not modelled	32.3	9	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
83	d1ng2a2	Alignment	not modelled	32.1	13	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
84	c2eczA	Alignment	not modelled	31.8	13	PDB header: signaling protein Chain: A: PDB Molecule: sorbin and sh3 domain-containing protein 1; PDBTitle: solution structure of the sh3 domain of sorbin and sh32 domain-containing protein 1
85	c1wxba	Alignment	not modelled	31.7	6	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: epidermal growth factor receptor pathway PDBTitle: solution structure of the sh3 domain from human epidermal2 growth factor receptor pathway substrate 8-like protein
86	d1h9ra2	Alignment	not modelled	31.5	17	Fold: OB-fold Superfamily: MOP-like Family: BiMOP, duplicated molybdate-binding domain
87	d1gcqa	Alignment	not modelled	31.1	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
88	c2df6A	Alignment	not modelled	30.9	19	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
89	c2krnA	Alignment	not modelled	30.6	16	PDB header: signaling protein Chain: A: PDB Molecule: cd2-associated protein; PDBTitle: high resolution structure of the second sh3 domain of cd2ap
90	c2l0aA	Alignment	not modelled	30.0	9	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
91	c2js0A	Alignment	not modelled	29.9	26	PDB header: signaling protein Chain: A: PDB Molecule: cytoplasmic protein nck1; PDBTitle: solution structure of second sh3 domain of adaptor nck
92	c1iq8B	Alignment	not modelled	29.8	11	PDB header: transferase Chain: B: PDB Molecule: archaeosine trna-guanine transglycosylase; PDBTitle: crystal structure of archaeosine trna-guanine2 transglycosylase from pyrococcus horikoshii
93	d1wlpb2	Alignment	not modelled	29.8	9	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
94	c1wi7A	Alignment	not modelled	29.1	10	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
95	c2bz8B	Alignment	not modelled	28.9	10	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
96	c1yn8E	Alignment	not modelled	28.7	27	PDB header: unknown function Chain: E: PDB Molecule: nap1-binding protein 2; PDBTitle: sh3 domain of yeast nbp2
97	c2yt6A	Alignment	not modelled	28.6	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
98	d1j6qa	Alignment	not modelled	28.4	11	Fold: OB-fold Superfamily: Heme chaperone CcmE Family: Heme chaperone CcmE
99	c1j6qA	Alignment	not modelled	28.4	11	PDB header: chaperone Chain: A: PDB Molecule: cytochrome c maturation protein e; PDBTitle: solution structure and characterization of the heme2 chaperone ccme
100	c1z9qA	Alignment	not modelled	28.1	13	PDB header: oxidoreductase activator Chain: A: PDB Molecule: neutrophil cytosol factor 4; PDBTitle: solution structure of sh3 domain of p40phox
101	c2eyxA	Alignment	not modelled	27.7	14	PDB header: signaling protein Chain: A: PDB Molecule: v-crk sarcoma virus ct10 oncogene homolog PDBTitle: c-terminal sh3 domain of ct10-regulated kinase
102	c1s1nA	Alignment	not modelled	27.7	20	PDB header: cell adhesion Chain: A: PDB Molecule: nephrocystin 1; PDBTitle: sh3 domain of human nephrocystin
103	c2kgtA	Alignment	not modelled	27.7	13	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase 6; PDBTitle: solution structure of sh3 domain of ptk6
104	c3cqtA	Alignment	not modelled	27.5	22	PDB header: transferase Chain: A: PDB Molecule: proto-oncogene tyrosine-protein kinase fyn; PDBTitle: n53i v55l mutant of fyn sh3 domain
105	d1g2ba	Alignment	not modelled	27.5	16	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain

106	c2ckkA	 Alignment	not modelled	27.0	16	PDB header: nuclear protein Chain: A: PDB Molecule: kin17; PDBTitle: high resolution crystal structure of the human kin172 c-terminal domain containing a kow motif3 kin17.
107	d1ckaa	 Alignment	not modelled	26.7	28	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
108	c2k6dA	 Alignment	not modelled	26.6	17	PDB header: sh3 domain/ubiquitin Chain: A: PDB Molecule: sh3 domain-containing kinase-binding protein 1; PDBTitle: cin85 sh3-c domain in complex with ubiquitin
109	c2nwmA	 Alignment	not modelled	26.3	19	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
110	c2eqiA	 Alignment	not modelled	25.7	6	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
111	d1fmka1	 Alignment	not modelled	25.5	22	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
112	d1tuca	 Alignment	not modelled	25.4	18	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
113	c3es4B	 Alignment	not modelled	25.1	13	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: uncharacterized protein duf861 with a rmlc-like cupin fold; PDBTitle: crystal structure of protein of unknown function (duf861) with a rmlc-2 like cupin fold (17741406) from agrobacterium tumefaciens str. c583 (dupont) at 1.64 a resolution
114	c2dila	 Alignment	not modelled	24.8	16	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
115	c2dl3A	 Alignment	not modelled	24.7	25	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
116	d1lcka1	 Alignment	not modelled	24.6	17	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
117	c2k2mA	 Alignment	not modelled	24.6	3	PDB header: signaling protein Chain: A: PDB Molecule: eps8-like protein 1; PDBTitle: structural basis of pxxdy motif recognition in sh3 binding
118	c2d1xD	 Alignment	not modelled	24.5	10	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
119	d1uffa	 Alignment	not modelled	24.4	10	Fold: SH3-like barrel Superfamily: SH3-domain Family: SH3-domain
120	c2x3wD	 Alignment	not modelled	24.1	10	PDB header: endocytosis Chain: D: PDB Molecule: protein kinase c and casein kinase substrate in neurons PDBTitle: structure of mouse syndapin i (crystal form 2)