





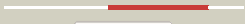










































#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3olmA_</a>	 Alignment		100.0	53	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase rsp5; <b>PDBTitle:</b> structure and function of a ubiquitin binding site within the2 catalytic domain of a hect ubiquitin ligase
2	<a href="#">c1zvdA_</a>	 Alignment		100.0	97	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> smad ubiquitination regulatory factor 2; <b>PDBTitle:</b> regulation of smurf2 ubiquitin ligase activity by anchoring the e2 to2 the hect domain
3	<a href="#">d1nd7a_</a>	 Alignment		100.0	51	<b>Fold:</b> Hect, E3 ligase catalytic domain <b>Superfamily:</b> Hect, E3 ligase catalytic domain <b>Family:</b> Hect, E3 ligase catalytic domain
4	<a href="#">c3g1nA_</a>	 Alignment		100.0	47	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase huwe1; <b>PDBTitle:</b> catalytic domain of the human e3 ubiquitin-protein ligase2 huwe1
5	<a href="#">c2oniA_</a>	 Alignment		100.0	54	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase nedd4-like protein; <b>PDBTitle:</b> catalytic domain of the human nedd4-like e3 ligase
6	<a href="#">d1c4za_</a>	 Alignment		100.0	35	<b>Fold:</b> Hect, E3 ligase catalytic domain <b>Superfamily:</b> Hect, E3 ligase catalytic domain <b>Family:</b> Hect, E3 ligase catalytic domain
7	<a href="#">c2jqzA_</a>	 Alignment		99.9	100	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase smurf2; <b>PDBTitle:</b> solution structure of the c2 domain of human smurf2
8	<a href="#">c3pt3A_</a>	 Alignment		99.9	33	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase ubr5; <b>PDBTitle:</b> crystal structure of the c-terminal lobe of the human ubr5 hect domain
9	<a href="#">c2kxqA_</a>	 Alignment		99.9	98	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase smurf2; <b>PDBTitle:</b> solution structure of smurf2 ww2 and ww3 bound to smad7 py motif2 containing peptide
10	<a href="#">d2nq3a1</a>	 Alignment		99.9	26	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
11	<a href="#">c2nq3A_</a>	 Alignment		99.9	26	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> itchy homolog e3 ubiquitin protein ligase; <b>PDBTitle:</b> crystal structure of the c2 domain of human itchy homolog2 e3 ubiquitin protein ligase

12	<a href="#">c3nsjA_</a>	Alignment		99.8	22	<b>PDB header:</b> immune system <b>Chain:</b> A: <b>PDB Molecule:</b> perforin-1; <b>PDBTitle:</b> the x-ray crystal structure of lymphocyte perforin
13	<a href="#">c1tk7A_</a>	Alignment		99.8	41	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> cg4244-pb; <b>PDBTitle:</b> nmr structure of ww domains (ww3-4) from suppressor of2 deltex
14	<a href="#">d1gmia_</a>	Alignment		99.8	21	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
15	<a href="#">c3m7fB_</a>	Alignment		99.8	18	<b>PDB header:</b> signaling protein/ligase <b>Chain:</b> B: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase nedd4; <b>PDBTitle:</b> crystal structure of the nedd4 c2/grb10 sh2 complex
16	<a href="#">c2dmhA_</a>	Alignment		99.8	21	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> myoferlin; <b>PDBTitle:</b> solution structure of the first c2 domain of human myoferlin
17	<a href="#">c2fk9A_</a>	Alignment		99.8	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein kinase c, eta type; <b>PDBTitle:</b> human protein kinase c, eta
18	<a href="#">d1bdya_</a>	Alignment		99.8	16	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
19	<a href="#">d1rlwa_</a>	Alignment		99.8	16	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
20	<a href="#">c3l9bA_</a>	Alignment		99.8	17	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> otoferlin; <b>PDBTitle:</b> crystal structure of rat otoferlin c2a
21	<a href="#">d1qasa2</a>	Alignment	not modelled	99.8	23	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
22	<a href="#">d2ep6a1</a>	Alignment	not modelled	99.8	23	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
23	<a href="#">c2z0uB_</a>	Alignment	not modelled	99.8	11	<b>PDB header:</b> lipid binding protein <b>Chain:</b> B: <b>PDB Molecule:</b> ww domain-containing protein 1; <b>PDBTitle:</b> crystal structure of c2 domain of kibra protein
24	<a href="#">c2nsgA_</a>	Alignment	not modelled	99.8	21	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase nedd4-like protein; <b>PDBTitle:</b> crystal structure of the c2 domain of the human e3 ubiquitin-protein2 ligase nedd4-like protein
25	<a href="#">c1dqvA_</a>	Alignment	not modelled	99.8	21	<b>PDB header:</b> endocytosis/exocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> synaptotagmin iii; <b>PDBTitle:</b> crystal structure of synaptotagmin iii c2a/c2b
26	<a href="#">d1wfja_</a>	Alignment	not modelled	99.7	15	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
27	<a href="#">c3kwtA_</a>	Alignment	not modelled	99.7	26	<b>PDB header:</b> metal binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> munc13-1; <b>PDBTitle:</b> munc13-1 c2b-domain, calcium-free
28	<a href="#">c3jzyA_</a>	Alignment	not modelled	99.7	20	<b>PDB header:</b> endocytosis <b>Chain:</b> A: <b>PDB Molecule:</b> intersectin 2; <b>PDBTitle:</b> crystal structure of human intersectin 2 c2 domain
						<b>Fold:</b> C2 domain-like

29	<a href="#">d1dsya_</a>	Alignment	not modelled	99.7	28	<b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
30	<a href="#">d1ugka_</a>	Alignment	not modelled	99.7	14	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
31	<a href="#">c2q3xA_</a>	Alignment	not modelled	99.7	21	<b>PDB header:</b> transport protein <b>Chain:</b> A; <b>PDB Molecule:</b> regulating synaptic membrane exocytosis protein 1; <b>PDBTitle:</b> the rim1alpha c2b domain
32	<a href="#">c2chdA_</a>	Alignment	not modelled	99.7	25	<b>PDB header:</b> protein transport <b>Chain:</b> A; <b>PDB Molecule:</b> rabphilin-3a; <b>PDBTitle:</b> crystal structure of the c2a domain of rabphilin-3a
33	<a href="#">c2d8kA_</a>	Alignment	not modelled	99.7	21	<b>PDB header:</b> endocytosis/exocytosis <b>Chain:</b> A; <b>PDB Molecule:</b> synaptotagmin vii; <b>PDBTitle:</b> solution structure of the first c2 domain of synaptotagmin2 vii
34	<a href="#">c2enjA_</a>	Alignment	not modelled	99.7	16	<b>PDB header:</b> transferase <b>Chain:</b> A; <b>PDB Molecule:</b> protein kinase c theta type; <b>PDBTitle:</b> solution strcture of the c2 domain from human protein2 kinase c theta
35	<a href="#">d2cjsa1</a>	Alignment	not modelled	99.7	21	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
36	<a href="#">c2r83B_</a>	Alignment	not modelled	99.7	16	<b>PDB header:</b> endocytosis, exocytosis <b>Chain:</b> B; <b>PDB Molecule:</b> synaptotagmin-1; <b>PDBTitle:</b> crystal structure analysis of human synaptotagmin 1 c2a-c2b
37	<a href="#">d2bwqa1</a>	Alignment	not modelled	99.7	26	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
38	<a href="#">d2r83a1</a>	Alignment	not modelled	99.7	31	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
39	<a href="#">d1wfma_</a>	Alignment	not modelled	99.7	17	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
40	<a href="#">d2cjta1</a>	Alignment	not modelled	99.7	21	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
41	<a href="#">d1rsya_</a>	Alignment	not modelled	99.7	18	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
42	<a href="#">d1v27a_</a>	Alignment	not modelled	99.7	25	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
43	<a href="#">c2enpA_</a>	Alignment	not modelled	99.7	18	<b>PDB header:</b> transport protein <b>Chain:</b> A; <b>PDB Molecule:</b> b/k protein; <b>PDBTitle:</b> solution structure of the first c2 domain from human b/k2 protein
44	<a href="#">c1cjbB_</a>	Alignment	not modelled	99.7	18	<b>PDB header:</b> hydrolase <b>Chain:</b> B; <b>PDB Molecule:</b> protein (cytosolic phospholipase a2); <b>PDBTitle:</b> human cytosolic phospholipase a2
45	<a href="#">d1a25a_</a>	Alignment	not modelled	99.7	27	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
46	<a href="#">c3fbkB_</a>	Alignment	not modelled	99.6	23	<b>PDB header:</b> signaling protein <b>Chain:</b> B; <b>PDB Molecule:</b> regulator of g-protein signaling 3; <b>PDBTitle:</b> crystal structure of the c2 domain of the human regulator2 of g-protein signaling 3 isoform 6 (rgp3), northeast3 structural genomics consortium target hr5550a
47	<a href="#">d2zkmx2</a>	Alignment	not modelled	99.6	20	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
48	<a href="#">c3l4hA_</a>	Alignment	not modelled	99.6	26	<b>PDB header:</b> protein binding <b>Chain:</b> A; <b>PDB Molecule:</b> e3 ubiquitin-protein ligase hecw1; <b>PDBTitle:</b> helical box domain and second ww domain of the human e3 ubiquitin-2 protein ligase hecw1
49	<a href="#">c1djyB_</a>	Alignment	not modelled	99.6	22	<b>PDB header:</b> lipid degradation <b>Chain:</b> B; <b>PDB Molecule:</b> phosphoinositide-specific phospholipase c, <b>PDBTitle:</b> phosphoinositide-specific phospholipase c-delta1 from rat2 complexed with inositol-2,4,5-trisphosphate
50	<a href="#">c2l5fA_</a>	Alignment	not modelled	99.6	18	<b>PDB header:</b> protein binding <b>Chain:</b> A; <b>PDB Molecule:</b> pre-mrna-processing factor 40 homolog a; <b>PDBTitle:</b> solution structure of the tandem ww domains from hypa/fbp11
51	<a href="#">c1o6wA_</a>	Alignment	not modelled	99.6	33	<b>PDB header:</b> nuclear protein <b>Chain:</b> A; <b>PDB Molecule:</b> pre-mrna processing protein prp40; <b>PDBTitle:</b> solution structure of the prp40 ww domain pair of the yeast2 splicing factor prp40
52	<a href="#">d2cm5a1</a>	Alignment	not modelled	99.6	27	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
53	<a href="#">c2cm6A_</a>	Alignment	not modelled	99.6	25	<b>PDB header:</b> protein transport <b>Chain:</b> A; <b>PDB Molecule:</b> rabphilin-3a; <b>PDBTitle:</b> crystal structure of the c2b domain of rabphilin3a
54	<a href="#">c2jxwA_</a>	Alignment	not modelled	99.6	15	<b>PDB header:</b> formin binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> ww domain-binding protein 4; <b>PDBTitle:</b> solution structure of the tandem ww domains of fbp21
55	<a href="#">c2dmgA_</a>	Alignment	not modelled	99.6	23	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A; <b>PDB Molecule:</b> kiaa1228 protein; <b>PDBTitle:</b> solution structure of the third c2 domain of kiaa12282 protein

56	<a href="#">c2b3rA_</a>	 Alignment	not modelled	99.6	21	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> phosphatidylinositol-4-phosphate 3-kinase c2 domain- <b>PDBTitle:</b> crystal structure of the c2 domain of class ii phosphatidylinositide2 3-kinase c2
57	<a href="#">d1rh8a_</a>	 Alignment	not modelled	99.6	25	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
58	<a href="#">d1dqva1</a>	 Alignment	not modelled	99.6	19	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
59	<a href="#">c2zajA_</a>	 Alignment	not modelled	99.5	38	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> membrane-associated guanylate kinase, ww and pdz <b>PDBTitle:</b> solution structure of the short-isoform of the second ww2 domain from the human membrane-associated guanylate kinase,3 ww and pdz domain-containing protein 1 (magi-1)
60	<a href="#">d1byna_</a>	 Alignment	not modelled	99.5	17	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
61	<a href="#">c2l4ja_</a>	 Alignment	not modelled	99.5	44	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> yes-associated protein 2 (yap2); <b>PDBTitle:</b> yap ww2
62	<a href="#">d1dqva2</a>	 Alignment	not modelled	99.5	24	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
63	<a href="#">c2lb0A_</a>	 Alignment	not modelled	99.5	39	<b>PDB header:</b> signaling protein/transcription <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase smurf1; <b>PDBTitle:</b> structure of the first ww domain of human smurf1 in complex with a di-2 phosphorylated human smad1 derived peptide
64	<a href="#">c2lazA_</a>	 Alignment	not modelled	99.5	39	<b>PDB header:</b> signaling protein/transcription <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase smurf1; <b>PDBTitle:</b> structure of the first ww domain of human smurf1 in complex with a2 mono-phosphorylated human smad1 derived peptide
65	<a href="#">c2ysdA_</a>	 Alignment	not modelled	99.5	41	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> membrane-associated guanylate kinase, ww and pdz <b>PDBTitle:</b> solution structure of the first ww domain from the human2 membrane-associated guanylate kinase, ww and pdz domain-3 containing protein 1. magi-1
66	<a href="#">d1tk7a1</a>	 Alignment	not modelled	99.5	52	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
67	<a href="#">d1uowa_</a>	 Alignment	not modelled	99.4	31	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
68	<a href="#">c3ohmB_</a>	 Alignment	not modelled	99.4	19	<b>PDB header:</b> signaling protein / hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> 1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase <b>PDBTitle:</b> crystal structure of activated g alpha q bound to its effector2 phospholipase c beta 3
69	<a href="#">c2djvA_</a>	 Alignment	not modelled	99.4	88	<b>PDB header:</b> ligase/signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> smad ubiquitination regulatory factor 2; <b>PDBTitle:</b> solution structure of smurf2 ww3 domain-smad7 py peptide2 complex
70	<a href="#">c3n5aA_</a>	 Alignment	not modelled	99.4	23	<b>PDB header:</b> protein transport <b>Chain:</b> A: <b>PDB Molecule:</b> synaptotagmin-7; <b>PDBTitle:</b> synaptotagmin-7, c2b-domain, calcium bound
71	<a href="#">d1i5hw_</a>	 Alignment	not modelled	99.4	46	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
72	<a href="#">c3fdwA_</a>	 Alignment	not modelled	99.4	20	<b>PDB header:</b> lipid binding protein <b>Chain:</b> A: <b>PDB Molecule:</b> synaptotagmin-like protein 4; <b>PDBTitle:</b> crystal structure of a c2 domain from human synaptotagmin-2 like protein 4
73	<a href="#">c3qr0A_</a>	 Alignment	not modelled	99.4	16	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> phospholipase c-beta (plc-beta); <b>PDBTitle:</b> crystal structure of s. officinalis plc21
74	<a href="#">c2jmfA_</a>	 Alignment	not modelled	99.4	58	<b>PDB header:</b> ligase/signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase suppressor of deltex; <b>PDBTitle:</b> solution structure of the su(dx) ww4- notch py peptide2 complex
75	<a href="#">d2jmfa1</a>	 Alignment	not modelled	99.4	73	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
76	<a href="#">c2lawA_</a>	 Alignment	not modelled	99.4	53	<b>PDB header:</b> signaling protein/transcription <b>Chain:</b> A: <b>PDB Molecule:</b> yorkie homolog; <b>PDBTitle:</b> structure of the second ww domain from human yap in complex with a2 human smad1 derived peptide
77	<a href="#">c1wmvA_</a>	 Alignment	not modelled	99.4	39	<b>PDB header:</b> oxidoreductase, apoptosis <b>Chain:</b> A: <b>PDB Molecule:</b> ww domain containing oxidoreductase; <b>PDBTitle:</b> solution structure of the second ww domain of wwox
78	<a href="#">c1e0mA_</a>	 Alignment	not modelled	99.4	43	<b>PDB header:</b> de novo protein <b>Chain:</b> A: <b>PDB Molecule:</b> wwprototype; <b>PDBTitle:</b> prototype ww domain
79	<a href="#">c2kq0A_</a>	 Alignment	not modelled	99.4	53	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase nedd4; <b>PDBTitle:</b> human nedd4 3rd ww domain complex with ebola zaire virus matrix2 protein vp40 derived peptide ilptappeyme
80	<a href="#">c2ysgA_</a>	 Alignment	not modelled	99.4	38	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> syntaxin-binding protein 4; <b>PDBTitle:</b> solution structure of the ww domain from the human

						syntaxin-2 binding protein 4
81	<a href="#">d1tk7a2</a>	Alignment	not modelled	99.4	59	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
82	<a href="#">c2kykA</a>	Alignment	not modelled	99.3	54	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase itchy homolog; <b>PDBTitle:</b> the sandwich region between two Imp2a py motif regulates the2 interaction between aip4ww2domain and py motif
83	<a href="#">c1ymzA</a>	Alignment	not modelled	99.3	39	<b>PDB header:</b> unknown function <b>Chain:</b> A: <b>PDB Molecule:</b> cc45; <b>PDBTitle:</b> cc45, an artificial ww domain designed using statistical2 coupling analysis
84	<a href="#">c2ez5W</a>	Alignment	not modelled	99.3	36	<b>PDB header:</b> signalling protein,ligase <b>Chain:</b> W: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase neddd4; <b>PDBTitle:</b> solution structure of the dneddd4 ww3* domain- comm lpsy2 peptide complex
85	<a href="#">c1wr7A</a>	Alignment	not modelled	99.3	49	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> neddd4-2; <b>PDBTitle:</b> solution structure of the third ww domain of neddd4-2
86	<a href="#">c2ysfA</a>	Alignment	not modelled	99.3	50	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> e3 ubiquitin-protein ligase itchy homolog; <b>PDBTitle:</b> solution structure of the fourth ww domain from the human2 e3 ubiquitin-protein ligase itchy homolog, itch
87	<a href="#">c2dmvA</a>	Alignment	not modelled	99.3	38	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> itchy homolog e3 ubiquitin protein ligase; <b>PDBTitle:</b> solution structure of the second ww domain of itchy homolog2 e3 ubiquitin protein ligase (itch)
88	<a href="#">c1yiuA</a>	Alignment	not modelled	99.3	66	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> itchy e3 ubiquitin protein ligase; <b>PDBTitle:</b> itch e3 ubiquitin ligase ww3 domain
89	<a href="#">c2yshA</a>	Alignment	not modelled	99.3	30	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> growth-arrest-specific protein 7; <b>PDBTitle:</b> solution structure of the ww domain from the human growth-2 arrest-specific protein 7, gas-7
90	<a href="#">d1w15a</a>	Alignment	not modelled	99.3	24	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> Synaptotagmin-like (S variant)
91	<a href="#">c2ysbA</a>	Alignment	not modelled	99.3	43	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> salvador homolog 1 protein; <b>PDBTitle:</b> solution structure of the first ww domain from the mouse2 salvador homolog 1 protein (sav1)
92	<a href="#">c2fjuB</a>	Alignment	not modelled	99.3	22	<b>PDB header:</b> signaling protein,apoptosis/hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> 1-phosphatidylinositol-4,5-bisphosphate <b>PDBTitle:</b> activated rac1 bound to its effector phospholipase c beta 2
93	<a href="#">c2dwvB</a>	Alignment	not modelled	99.3	29	<b>PDB header:</b> protein binding <b>Chain:</b> B: <b>PDB Molecule:</b> salvador homolog 1 protein; <b>PDBTitle:</b> solution structure of the second ww domain from mouse2 salvador homolog 1 protein (mww45)
94	<a href="#">c1wr4A</a>	Alignment	not modelled	99.3	56	<b>PDB header:</b> ligase <b>Chain:</b> A: <b>PDB Molecule:</b> ubiquitin-protein ligase neddd4-2; <b>PDBTitle:</b> solution structure of the second ww domain of neddd4-2
95	<a href="#">d1k9ra</a>	Alignment	not modelled	99.3	43	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
96	<a href="#">c2yscA</a>	Alignment	not modelled	99.3	28	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> amyloid beta a4 precursor protein-binding family <b>PDBTitle:</b> solution structure of the ww domain from the human amyloid2 beta a4 precursor protein-binding family b member 3, apbb3
97	<a href="#">d2f21a1</a>	Alignment	not modelled	99.2	50	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
98	<a href="#">d1nmva1</a>	Alignment	not modelled	99.1	33	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
99	<a href="#">d1f8ab1</a>	Alignment	not modelled	99.0	38	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
100	<a href="#">d1i8gb</a>	Alignment	not modelled	99.0	40	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
101	<a href="#">d2itka1</a>	Alignment	not modelled	99.0	43	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
102	<a href="#">c2ysiA</a>	Alignment	not modelled	99.0	21	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> transcription elongation regulator 1; <b>PDBTitle:</b> solution structure of the first ww domain from the mouse2 transcription elongation regulator 1, transcription factor3 ca150
103	<a href="#">d1pina1</a>	Alignment	not modelled	99.0	45	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
104	<a href="#">c1yw5A</a>	Alignment	not modelled	98.2	24	<b>PDB header:</b> isomerase <b>Chain:</b> A: <b>PDB Molecule:</b> peptidyl prolyl cis/trans isomerase; <b>PDBTitle:</b> peptidyl-prolyl isomerase ess1 from candida albicans
105	<a href="#">d1o6wa2</a>	Alignment	not modelled	98.2	39	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
106	<a href="#">d1o6wa1</a>	Alignment	not modelled	98.1	21	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain

107	<a href="#">c2engA_</a>	 Alignment	not modelled	98.0	15	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> phosphatidylinositol-4,5-bisphosphate 3-kinase <b>PDBTitle:</b> solution structure of the c2 domain from human pi3-kinase2 p110 subunit alpha
108	<a href="#">c2dk7A_</a>	 Alignment	not modelled	98.0	28	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcription elongation regulator 1; <b>PDBTitle:</b> solution structure of ww domain in transcription elongation2 regulator 1
109	<a href="#">d2rm0w1</a>	 Alignment	not modelled	97.9	22	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
110	<a href="#">d2e45a1</a>	 Alignment	not modelled	97.7	27	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
111	<a href="#">c3pfqA_</a>	 Alignment	not modelled	97.7	27	<b>PDB header:</b> transferase <b>Chain:</b> A: <b>PDB Molecule:</b> protein kinase c beta type; <b>PDBTitle:</b> crystal structure and allosteric activation of protein kinase c beta2 ii
112	<a href="#">d1zr7a1</a>	 Alignment	not modelled	97.6	19	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
113	<a href="#">d1ywia1</a>	 Alignment	not modelled	97.6	19	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
114	<a href="#">c1zr7A_</a>	 Alignment	not modelled	97.5	19	<b>PDB header:</b> signaling protein <b>Chain:</b> A: <b>PDB Molecule:</b> huntingtin-interacting protein hypa/fbp11; <b>PDBTitle:</b> solution structure of the first ww domain of fbp11
115	<a href="#">c2dk1A_</a>	 Alignment	not modelled	97.5	24	<b>PDB header:</b> gene regulation <b>Chain:</b> A: <b>PDB Molecule:</b> ww domain-binding protein 4; <b>PDBTitle:</b> solution structure of ww domain in ww domain binding2 protein 4 (wbp-4)
116	<a href="#">c1ywjA_</a>	 Alignment	not modelled	97.5	19	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> formin-binding protein 3; <b>PDBTitle:</b> structure of the fbp11ww1 domain
117	<a href="#">d1e7ua2</a>	 Alignment	not modelled	97.4	17	<b>Fold:</b> C2 domain-like <b>Superfamily:</b> C2 domain (Calcium/lipid-binding domain, CaLB) <b>Family:</b> PLC-like (P variant)
118	<a href="#">c1eg4A_</a>	 Alignment	not modelled	97.4	21	<b>PDB header:</b> structural protein <b>Chain:</b> A: <b>PDB Molecule:</b> dystrophin; <b>PDBTitle:</b> structure of a dystrophin ww domain fragment in complex2 with a beta-dystroglycan peptide
119	<a href="#">d2dk1a1</a>	 Alignment	not modelled	97.4	24	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain
120	<a href="#">d2ho2a1</a>	 Alignment	not modelled	97.3	34	<b>Fold:</b> WW domain-like <b>Superfamily:</b> WW domain <b>Family:</b> WW domain