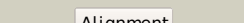
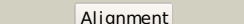
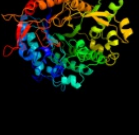
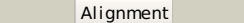
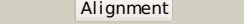
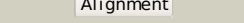
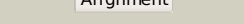
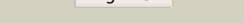


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3pn8A_	 Alignment		100.0	49	PDB header: hydrolase Chain: A: PDB Molecule: putative phospho-beta-glucosidase; PDBTitle: the crystal structure of 6-phospho-beta-glucosidase from streptococcus2 mutans ua159
2	dlqoxa_	 Alignment		100.0	34	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
3	c2xhyD_	 Alignment		100.0	54	PDB header: hydrolase Chain: D: PDB Molecule: 6-phospho-beta-glucosidase bgla; PDBTitle: crystal structure of e.coli bgla
4	c3qomA_	 Alignment		100.0	49	PDB header: hydrolase Chain: A: PDB Molecule: 6-phospho-beta-glucosidase; PDBTitle: crystal structure of 6-phospho-beta-glucosidase from lactobacillus2 plantarum
5	d2j78a1	 Alignment		100.0	33	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
6	c3fiyA_	 Alignment		100.0	33	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of bglb
7	dle4ia_	 Alignment		100.0	33	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
8	c3ahxC_	 Alignment		100.0	33	PDB header: hydrolase Chain: C: PDB Molecule: beta-glucosidase a; PDBTitle: crystal structure of beta-glucosidase a from bacterium clostridium2 cellulovorans
9	dlgnxa_	 Alignment		100.0	32	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
10	c2dgaA_	 Alignment		100.0	30	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of hexameric beta-glucosidase in wheat
11	c2j75A_	 Alignment		100.0	32	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase a; PDBTitle: beta-glucosidase from thermotoga maritima in complex with2 noeuromycin

12	c2rgmA_	Alignment		100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: rice bglu1 beta-glucosidase, a plant exoglucanase/beta-glucosidase
13	c3gnoA_	Alignment		100.0	32	PDB header: hydrolase Chain: A: PDB Molecule: os03g0212800 protein; PDBTitle: crystal structure of a rice os3bglu6 beta-glucosidase
14	d1v08a_	Alignment		100.0	30	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
15	c2z1sA_	Alignment		100.0	30	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase b; PDBTitle: beta-glucosidase b from paenibacillus polymyxa complexed2 with cellotetraose
16	d1e4mm_	Alignment		100.0	28	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
17	c3ai0A_	Alignment		100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of beta-glucosidase from termite neotermes2 koshunensis in complex with para-nitrophenyl-beta-d-glucopyranoside
18	c3ptkB_	Alignment		100.0	30	PDB header: hydrolase Chain: B: PDB Molecule: beta-glucosidase os4bglu12; PDBTitle: the crystal structure of rice (oryza sativa l.) os4bglu12
19	c3u57A_	Alignment		100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: raucaffricine-o-beta-d-glucosidase; PDBTitle: structures of alkaloid biosynthetic glucosidases decode substrate2 specificity
20	d1pbga_	Alignment		100.0	36	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
21	c1v02F_	Alignment	not modelled	100.0	30	PDB header: hydrolase Chain: F: PDB Molecule: dhurrinase; PDBTitle: crystal structure of the sorghum bicolor dhurrinase 1
22	d1v02a_	Alignment	not modelled	100.0	30	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
23	c2jf7B_	Alignment	not modelled	100.0	29	PDB header: hydrolase Chain: B: PDB Molecule: strictosidine-o-beta-d-glucosidase; PDBTitle: structure of strictosidine glucosidase
24	c2e3zB_	Alignment	not modelled	100.0	29	PDB header: hydrolase Chain: B: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of intracellular family 1 beta-2 glucosidase bgl1a from the basidiomycete phanerochaete3 chrysosporium in substrate-free form
25	d1cbga_	Alignment	not modelled	100.0	31	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
26	c3ahyD_	Alignment	not modelled	100.0	30	PDB header: hydrolase Chain: D: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of beta-glucosidase 2 from fungus trichoderma reesei2 in complex with tris
27	c2zoxA_	Alignment	not modelled	100.0	28	PDB header: hydrolase Chain: A: PDB Molecule: cytosolic beta-glucosidase; PDBTitle: crystal structure of the covalent intermediate of human cytosolic2 beta-glucosidase
28	d1wcga1	Alignment	not modelled	100.0	30	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase

29	dluwsa_	Alignment	not modelled	100.0	26	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
30	dlqvba_	Alignment	not modelled	100.0	23	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
31	dlug6a_	Alignment	not modelled	100.0	31	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
32	dlvffa1	Alignment	not modelled	100.0	29	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
33	c2cncA_	Alignment	not modelled	100.0	12	PDB header: hydrolase Chain: A: PDB Molecule: endoxylanase; PDBTitle: family 10 xylanase
34	clxyzA_	Alignment	not modelled	100.0	14	PDB header: glycosyltransferase Chain: A: PDB Molecule: 1,4-beta-d-xylan-xylanohydrolase; PDBTitle: a common protein fold and similar active site in two2 distinct families of beta-glycanases
35	dlxyza_	Alignment	not modelled	100.0	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
36	dlv6ya_	Alignment	not modelled	100.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
37	dlur1a_	Alignment	not modelled	100.0	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
38	dlfh9a_	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
39	dlmq6a_	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
40	dlkwga2	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
41	c3emzA_	Alignment	not modelled	100.0	15	PDB header: hydrolase Chain: A: PDB Molecule: endo-1,4-beta-xylanase; PDBTitle: crystal structure of xylanase xynb from paenibacillus2 barcinonensis complexed with a conduramine derivative
42	dlvbua1	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
43	dlv0la_	Alignment	not modelled	100.0	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
44	dlv6wa2	Alignment	not modelled	100.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
45	c2jepB_	Alignment	not modelled	100.0	15	PDB header: hydrolase Chain: B: PDB Molecule: xyloglucanase; PDBTitle: native family 5 xyloglucanase from paenibacillus pabuli
46	dlm82a_	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
47	dlw32a_	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
48	c3ndyA_	Alignment	not modelled	100.0	19	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase d; PDBTitle: the structure of the catalytic and carbohydrate binding domain of2 endoglucanase d from clostridium cellulovorans
49	c2depA_	Alignment	not modelled	100.0	15	PDB header: hydrolase Chain: A: PDB Molecule: thermostable celloxylanase; PDBTitle: crystal structure of xylanase b from clostridium2 stercorarium f9
50	dlilwa_	Alignment	not modelled	100.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
51	dlwedga_	Alignment	not modelled	100.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
52	dlrh9a1	Alignment	not modelled	99.9	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
53	dlbg4a_	Alignment	not modelled	99.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
54	dlw91a2	Alignment	not modelled	99.9	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
55	c3aysA_	Alignment	not modelled	99.9	20	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase; PDBTitle: gh5 endoglucanase from a ruminal fungus in complex with cellotriose
56	dlvjza_	Alignment	not modelled	99.9	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases

					Family: beta-glycanases
57	dlhjsa_	Alignment	not modelled	99.9	13 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
58	dlhjqa_	Alignment	not modelled	99.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
59	dlta3b_	Alignment	not modelled	99.9	13 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
60	c3ncoA_	Alignment	not modelled	99.9	19 PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase fncel5a; PDBTitle: crystal structure of fncel5a from f. nodosum rt17-b1
61	dltxa_	Alignment	not modelled	99.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
62	c3mmwB_	Alignment	not modelled	99.9	15 PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: crystal structure of endoglucanase cel5a from the hyperthermophilic2 thermotoga maritima
63	dlceoa_	Alignment	not modelled	99.9	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
64	dlus3a2	Alignment	not modelled	99.9	14 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
65	c3pzqA_	Alignment	not modelled	99.9	16 PDB header: hydrolase Chain: A: PDB Molecule: mannan endo-1,4-beta-mannosidase. glycosyl hydrolase family PDBTitle: structure of the hyperthermostable endo-1,4-beta-d-mannanase from2 thermotoga petrophila rku-1 with maltose and glycerol
66	c2fglA_	Alignment	not modelled	99.9	17 PDB header: hydrolase Chain: A: PDB Molecule: alkaline thermostable endoxylanase; PDBTitle: an alkali thermostable f/10 xylanase from alkalophilic2 bacillus sp. ng-27
67	cluz4A_	Alignment	not modelled	99.9	16 PDB header: hydrolase Chain: A: PDB Molecule: man5a; PDBTitle: common inhibition of beta-glucosidase and beta-mannosidase2 by isofagomine lactam reflects different conformational3 itineraries for glucoside and mannoside hydrolysis
68	dluuqa_	Alignment	not modelled	99.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
69	dlr85a_	Alignment	not modelled	99.8	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
70	dlur4a_	Alignment	not modelled	99.8	21 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
71	c3qr3B_	Alignment	not modelled	99.8	13 PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase eg-ii; PDBTitle: crystal structure of cel5a (eg2) from hypocrea jecorina (trichoderma2 reesei)
72	c3icgD_	Alignment	not modelled	99.8	16 PDB header: hydrolase Chain: D: PDB Molecule: endoglucanase d; PDBTitle: crystal structure of the catalytic and carbohydrate binding domain of2 endoglucanase d from clostridium cellulovorans
73	dlfoba_	Alignment	not modelled	99.8	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
74	dlh1na_	Alignment	not modelled	99.8	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
75	dluhva2	Alignment	not modelled	99.8	20 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
76	c2oylB_	Alignment	not modelled	99.8	17 PDB header: hydrolase Chain: B: PDB Molecule: endoglycoceramidase ii; PDBTitle: endo-glycoceramidase ii from rhodococcus sp.: cellobiose-like2 imidazole complex
77	c2bs9B_	Alignment	not modelled	99.8	18 PDB header: hydrolase Chain: B: PDB Molecule: beta-xylosidase; PDBTitle: native crystal structure of a gh39 beta-xylosidase xynb12 from geobacillus stearothermophilus
78	clkwgA_	Alignment	not modelled	99.8	15 PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of thermus thermophilus a4 beta-galactosidase
79	c3l55B_	Alignment	not modelled	99.7	14 PDB header: structural genomics, unknown function Chain: B: PDB Molecule: b-1,4-endoglucanase/cellulase; PDBTitle: crystal structure of a putative beta-1,4-endoglucanase /2 cellulase from prevotella bryantii
80	cluhvD_	Alignment	not modelled	99.7	21 PDB header: hydrolase Chain: D: PDB Molecule: beta-xylosidase; PDBTitle: crystal structure of beta-d-xylosidase from2 thermoanaerobacterium saccharolyticum, a family 393 glycoside hydrolase
81	dlecea_	Alignment	not modelled	99.7	13 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases

82	d1bhga3	Alignment	not modelled	99.7	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
83	c1bhgB_	Alignment	not modelled	99.7	20	PDB header: glycosidase Chain: B: PDB Molecule: beta-glucuronidase; PDBTitle: human beta-glucuronidase at 2.6 a resolution
84	c2zunB_	Alignment	not modelled	99.7	14	PDB header: hydrolase Chain: B: PDB Molecule: 458aa long hypothetical endo-1,4-beta-glucanase; PDBTitle: functional analysis of hyperthermophilic endocellulase from2 the archaeon pyrococcus horikoshii
85	cliszA_	Alignment	not modelled	99.7	18	PDB header: hydrolase Chain: A: PDB Molecule: endo-1,4-beta-d-xylanase; PDBTitle: crystal structure of xylanase from streptomyces2 olivaceoviridis e-86 complexed with galactose
86	d2c0ha1	Alignment	not modelled	99.7	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
87	d2pb1a1	Alignment	not modelled	99.6	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
88	d1egza_	Alignment	not modelled	99.6	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
89	c3lpgA_	Alignment	not modelled	99.6	16	PDB header: hydrolase/hydrolase inhibitor Chain: A: PDB Molecule: beta-glucuronidase; PDBTitle: structure of e. coli beta-glucuronidase bound with a novel, potent2 inhibitor 3-(2-fluorophenyl)-1-(2-hydroxyethyl)-1-((6-methyl-2-oxo-1,3,2-dihydroquinolin-3-yl)methyl)urea
90	c2w5fB_	Alignment	not modelled	99.6	12	PDB header: hydrolase Chain: B: PDB Molecule: endo-1,4-beta-xylanase y; PDBTitle: high resolution crystallographic structure of the2 clostridium thermocellum n-terminal endo-1,4-beta-d-3 xylanase 10b (xyn10b) cbm22-1-gh10 modules complexed with4 xylohexaose
91	c3civA_	Alignment	not modelled	99.6	18	PDB header: hydrolase Chain: A: PDB Molecule: endo-beta-1,4-mannanase; PDBTitle: crystal structure of the endo-beta-1,4-mannanase from2 alicyclobacillus acidocaldarius
92	d1qnra_	Alignment	not modelled	99.6	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
93	c3jugA_	Alignment	not modelled	99.5	22	PDB header: hydrolase Chain: A: PDB Molecule: beta-mannanase; PDBTitle: crystal structure of endo-beta-1,4-mannanase from the alkaliphilic2 bacillus sp. n16-5
94	d1yq2a5	Alignment	not modelled	99.5	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
95	d1wkyA2	Alignment	not modelled	99.5	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
96	c2cksB_	Alignment	not modelled	99.5	14	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase e-5; PDBTitle: x-ray crystal structure of the catalytic domain of2 thermobifida fusca endoglucanase cel5a (e5)
97	d1tvna1	Alignment	not modelled	99.5	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
98	d7a3ha_	Alignment	not modelled	99.5	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
99	d1tg7a5	Alignment	not modelled	99.5	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Glycosyl hydrolases family 35 catalytic domain
100	c3cmgA_	Alignment	not modelled	99.4	19	PDB header: hydrolase Chain: A: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
101	c3pzvB_	Alignment	not modelled	99.4	18	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: c2 crystal form of the endo-1,4-beta-glucanase from bacillus subtilis2 168
102	d1h4pa_	Alignment	not modelled	99.4	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
103	d1g01a_	Alignment	not modelled	99.4	9	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
104	d1bqca_	Alignment	not modelled	99.4	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
105	d1jz8a5	Alignment	not modelled	99.4	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
106	c3fn9B_	Alignment	not modelled	99.4	16	PDB header: hydrolase Chain: B: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
107	c1us2A_	Alignment	not modelled	99.4	13	PDB header: hydrolase Chain: A: PDB Molecule: endo-beta-1,4-xylanase; PDBTitle: xylanase10c (mutant e385a) from cellvibrio japonicus in2 complex with xylopentaose PDB header: hydrolase

108	c1wkyA_	Alignment	not modelled	99.2	13	Chain: A: PDB Molecule: endo-beta-1,4-mannanase; PDBTitle: crystal structure of alkaline mannanase from bacillus sp. strain jamb-2 602: catalytic domain and its carbohydrate binding module
109	c2y8kA_	Alignment	not modelled	99.1	15	PDB header: hydrolase Chain: A: PDB Molecule: carbohydrate binding family 6; PDBTitle: structure of ctgh5-cbm6, an arabinoxylan-specific xylanase.
110	c2w62A_	Alignment	not modelled	99.0	16	PDB header: transferase Chain: A: PDB Molecule: glycolipid-anchored surface protein 2; PDBTitle: saccharomyces cerevisiae gas2p in complex with2 laminaripentaose
111	c1j0yD_	Alignment	not modelled	98.9	12	PDB header: hydrolase Chain: D: PDB Molecule: beta-amylase; PDBTitle: beta-amylase from bacillus cereus var. mycoides in complex2 with glucose
112	c3gm8A_	Alignment	not modelled	98.7	22	PDB header: hydrolase Chain: A: PDB Molecule: glycoside hydrolase family 2, candidate beta-glycosidase; PDBTitle: crystal structure of a beta-glycosidase from bacteroides vulgatus
113	c3thdD_	Alignment	not modelled	98.7	12	PDB header: hydrolase Chain: D: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of human beta-galactosidase in complex with 1-2 deoxygalactonojirimycin
114	d1vema2	Alignment	not modelled	98.7	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
115	c3u7vA_	Alignment	not modelled	98.6	11	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: the structure of a putative beta-galactosidase from caulobacter2 crescentus cb15.
116	c3clwF_	Alignment	not modelled	98.5	14	PDB header: structural genomics, unknown function Chain: F: PDB Molecule: conserved exported protein; PDBTitle: crystal structure of conserved exported protein from bacteroides2 fragilis
117	c2wnwB_	Alignment	not modelled	98.4	12	PDB header: hydrolase Chain: B: PDB Molecule: activated by transcription factor ssrb; PDBTitle: the crystal structure of srfj from salmonella typhimurium
118	c1nofA_	Alignment	not modelled	98.4	19	PDB header: hydrolase Chain: A: PDB Molecule: xylanase; PDBTitle: the first crystallographic structure of a xylanase from2 glycosyl hydrolase family 5: implications for catalysis
119	d2je8a5	Alignment	not modelled	98.3	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
120	c2y2wC_	Alignment	not modelled	98.3	19	PDB header: hydrolase Chain: C: PDB Molecule: arabinofuranosidase; PDBTitle: elucidation of the substrate specificity and protein2 structure of abfb, a family 51 alpha-l-arabinofuranosidase3 from bifidobacterium longum.