



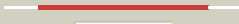



















#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1qoxa_	 Alignment		100.0	35	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
2	c3pn8A_	 Alignment		100.0	50	PDB header: hydrolase Chain: A: PDB Molecule: putative phospho-beta-glucosidase; PDBTitle: the crystal structure of 6-phospho-beta-glucosidase from streptococcus2 mutans ua159
3	d2j78a1	 Alignment		100.0	34	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
4	c3qomA_	 Alignment		100.0	51	PDB header: hydrolase Chain: A: PDB Molecule: 6-phospho-beta-glucosidase; PDBTitle: crystal structure of 6-phospho-beta-glucosidase from lactobacillus2 plantarum
5	c2xhyD_	 Alignment		100.0	54	PDB header: hydrolase Chain: D: PDB Molecule: 6-phospho-beta-glucosidase bgla; PDBTitle: crystal structure of e.coli bgla
6	c3fiyA_	 Alignment		100.0	33	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of bglb
7	d1e4ia_	 Alignment		100.0	33	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
8	c3ahxC_	 Alignment		100.0	34	PDB header: hydrolase Chain: C: PDB Molecule: beta-glucosidase a; PDBTitle: crystal structure of beta-glucosidase a from bacterium clostridium2 cellulovorans
9	c2j75A_	 Alignment		100.0	34	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase a; PDBTitle: beta-glucosidase from thermotoga maritima in complex with2 noeuromycin
10	c2dgaA_	 Alignment		100.0	31	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of hexameric beta-glucosidase in wheat
11	d1gnxa_	 Alignment		100.0	35	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase

12	dlv08a_	Alignment		100.0	29	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
13	c2rgmA_	Alignment		100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: rice bglu1 beta-glucosidase, a plant exoglucanase/beta-glucosidase
14	c3gnoA_	Alignment		100.0	32	PDB header: hydrolase Chain: A: PDB Molecule: os03g0212800 protein; PDBTitle: crystal structure of a rice os3bglu6 beta-glucosidase
15	c3ai0A_	Alignment		100.0	30	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
16	c2z1sA_	Alignment		100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase b; PDBTitle: beta-glucosidase b from paenibacillus polymyxa complexed2 with cellotetraose
17	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
18	dle4mm_	Alignment		100.0	27	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
19	c3u57A_	Alignment		100.0	28	PDB header: hydrolase Chain: A: PDB Molecule: raucaffricine-o-beta-d-glucosidase; PDBTitle: structures of alkaloid biosynthetic glucosidases decode substrate2 specificity
20	c1v02F_	Alignment		100.0	33	PDB header: hydrolase Chain: F: PDB Molecule: dhurrinase; PDBTitle: crystal structure of the sorghum bicolor dhurrinase 1
21	dlv02a_	Alignment	not modelled	100.0	33	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
22	c2jf7B_	Alignment	not modelled	100.0	30	PDB header: hydrolase Chain: B: PDB Molecule: strictosidine-o-beta-d-glucosidase; PDBTitle: structure of strictosidine glucosidase
23	dlcbga_	Alignment	not modelled	100.0	31	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
24	dlpbga_	Alignment	not modelled	100.0	35	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
25	dlwcga1	Alignment	not modelled	100.0	31	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
26	c2e3zB_	Alignment	not modelled	100.0	30	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
27	c3ahyD_	Alignment	not modelled	100.0	31	PDB header: hydrolase Chain: D: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of beta-glucosidase 2 from fungus trichoderma reesei2 in complex with tris
28	c2zoxA_	Alignment	not modelled	100.0	29	PDB header: hydrolase Chain: A: PDB Molecule: cytosolic beta-glucosidase; PDBTitle: crystal structure of the covalent intermediate of human cytosolic2 beta-glucosidase

29	dlqvba_	Alignment	not modelled	100.0	23	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
30	dluwsa_	Alignment	not modelled	100.0	25	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
31	dlug6a_	Alignment	not modelled	100.0	33	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
32	dlvffa1	Alignment	not modelled	100.0	28	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
33	c2cncA_	Alignment	not modelled	100.0	16	PDB header: hydrolase Chain: A: PDB Molecule: endoxylanase; PDBTitle: family 10 xylanase
34	dlxyza_	Alignment	not modelled	100.0	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
35	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
36	dlv6ya_	Alignment	not modelled	100.0	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
37	dlur1a_	Alignment	not modelled	100.0	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
38	dlkwga2	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
39	dlfh9a_	Alignment	not modelled	100.0	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
40	dlmq6a_	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
41	c3emzA_	Alignment	not modelled	100.0	14	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	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
43	c2jepB_	Alignment	not modelled	100.0	15	PDB header: hydrolase Chain: B: PDB Molecule: xyloglucanase; PDBTitle: native family 5 xyloglucanase from paenibacillus pabuli
44	dlv0la_	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
45	dlv6wa2	Alignment	not modelled	100.0	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
46	dlm82a_	Alignment	not modelled	100.0	15	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	c2depA_	Alignment	not modelled	100.0	16	PDB header: hydrolase Chain: A: PDB Molecule: thermostable cellobiohydrolase; PDBTitle: crystal structure of xylanase b from clostridium2 stercorarium f9
49	c3ndyA_	Alignment	not modelled	100.0	16	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
50	dlilwa_	Alignment	not modelled	100.0	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
51	dlrh9a1	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
52	dledga_	Alignment	not modelled	100.0	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
53	dlw91a2	Alignment	not modelled	100.0	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
54	dlbg4a_	Alignment	not modelled	100.0	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
55	dlhjsa_	Alignment	not modelled	100.0	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
56	dlhjga_	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases

					Family: beta-glycanases
57	c3aysA	Alignment	not modelled	99.9	17 PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase; PDBTitle: gh5 endoglucanase from a ruminal fungus in complex with cellotriose
58	d1vjza	Alignment	not modelled	99.9	19 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
59	d1ta3b	Alignment	not modelled	99.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
60	c3ncoA	Alignment	not modelled	99.9	17 PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase fncel5a; PDBTitle: crystal structure of fncel5a from f. nodosum rt17-b1
61	d1ceoa	Alignment	not modelled	99.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
62	c3mmwB	Alignment	not modelled	99.9	16 PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: crystal structure of endoglucanase cel5a from the hyperthermophilic2 thermotoga maritima
63	d1tuxa	Alignment	not modelled	99.9	21 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
64	d1us3a2	Alignment	not modelled	99.9	12 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
65	c3pzaA	Alignment	not modelled	99.9	18 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	c1uz4A	Alignment	not modelled	99.9	17 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
67	d1uuqa	Alignment	not modelled	99.9	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
68	c2fglA	Alignment	not modelled	99.9	16 PDB header: hydrolase Chain: A: PDB Molecule: alkaline thermostable endoxylanase; PDBTitle: an alkali thermostable f/10 xylanase from alkalophilic2 bacillus sp. ng-27
69	d1ur4a	Alignment	not modelled	99.9	19 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
70	d1uhva2	Alignment	not modelled	99.9	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
71	c2bs9B	Alignment	not modelled	99.9	18 PDB header: hydrolase Chain: B: PDB Molecule: beta-xylosidase; PDBTitle: native crystal structure of a gh39 beta-xylosidase xynb12 from geobacillus stearothermophilus
72	d1foba	Alignment	not modelled	99.9	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
73	d1r85a	Alignment	not modelled	99.9	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
74	c1kwgA	Alignment	not modelled	99.9	20 PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of thermus thermophilus a4 beta-galactosidase
75	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
76	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)
77	c2oylB	Alignment	not modelled	99.8	16 PDB header: hydrolase Chain: B: PDB Molecule: endoglycoceramidase ii; PDBTitle: endo-glycoceramidase ii from rhodococcus sp.: cellobiose-like2 imidazole complex
78	d1h1na	Alignment	not modelled	99.8	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
79	c1uhvD	Alignment	not modelled	99.8	18 PDB header: hydrolase Chain: D: PDB Molecule: beta-xylosidase; PDBTitle: crystal structure of beta-d-xylosidase from2 thermoanaerobacterium saccharolyticum, a family 393 glycoside hydrolase
80	c3l55B	Alignment	not modelled	99.8	15 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
81	d1bhga3	Alignment	not modelled	99.8	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases

82	c1bhgB	Alignment	not modelled	99.8	19	PDB header: glycosidase Chain: B: PDB Molecule: beta-glucuronidase; PDBTitle: human beta-glucuronidase at 2.6 a resolution
83	d1ecea	Alignment	not modelled	99.8	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
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	d2pb1a1	Alignment	not modelled	99.7	11	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
86	d2c0ha1	Alignment	not modelled	99.7	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
87	c3lpgA	Alignment	not modelled	99.7	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
88	d1qnra	Alignment	not modelled	99.7	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
89	d1egza	Alignment	not modelled	99.7	12	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
90	c3jugA	Alignment	not modelled	99.6	19	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
91	c2w5fB	Alignment	not modelled	99.6	14	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
92	c3civA	Alignment	not modelled	99.6	15	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
93	c1iszA	Alignment	not modelled	99.6	17	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
94	c2cksB	Alignment	not modelled	99.6	19	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)
95	d1wkya2	Alignment	not modelled	99.6	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
96	d1yq2a5	Alignment	not modelled	99.6	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
97	d1tvna1	Alignment	not modelled	99.6	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
98	c3pzvB	Alignment	not modelled	99.5	13	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: c2 crystal form of the endo-1,4-beta-glucanase from bacillus subtilis2 168
99	d1h4pa	Alignment	not modelled	99.5	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
100	d7a3ha	Alignment	not modelled	99.5	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
101	d1bqca	Alignment	not modelled	99.5	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
102	c3cmgA	Alignment	not modelled	99.5	19	PDB header: hydrolase Chain: A: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
103	d1jz8a5	Alignment	not modelled	99.5	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
104	c3fn9B	Alignment	not modelled	99.5	16	PDB header: hydrolase Chain: B: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
105	d1tg7a5	Alignment	not modelled	99.5	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Glycosyl hydrolases family 35 catalytic domain
106	d1g01a	Alignment	not modelled	99.5	11	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
107	c1wkya	Alignment	not modelled	99.3	18	PDB header: hydrolase 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
108	c1j0yD_	Alignment	not modelled	99.2	12 PDB header: hydrolase Chain: D: PDB Molecule: beta-amylase; PDBTitle: beta-amylase from bacillus cereus var. mycoides in complex2 with glucose
109	c1us2A_	Alignment	not modelled	99.2	12 PDB header: hydrolase Chain: A: PDB Molecule: endo-beta-1,4-xylanase; PDBTitle: xylanase10c (mutant e385a) from cellvibrio japonicus in2 complex with xylopentaose
110	c2w62A_	Alignment	not modelled	99.2	18 PDB header: transferase Chain: A: PDB Molecule: glycolipid-anchored surface protein 2; PDBTitle: saccharomyces cerevisiae gas2p in complex with2 laminaripentaose
111	c3u7vA_	Alignment	not modelled	99.1	16 PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: the structure of a putative beta-galactosidase from caulobacter2 crescentus cb15.
112	c2y8kA_	Alignment	not modelled	99.0	14 PDB header: hydrolase Chain: A: PDB Molecule: carbohydrate binding family 6; PDBTitle: structure of ctgh5-cbm6, an arabinoxylan-specific xylanase.
113	d1vema2	Alignment	not modelled	99.0	15 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
114	d2cyga1	Alignment	not modelled	98.9	12 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
115	c3gm8A_	Alignment	not modelled	98.8	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
116	c3thdD_	Alignment	not modelled	98.8	12 PDB header: hydrolase Chain: D: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of human beta-galactosidase in complex with 1-2 deoxygalactonojirimycin
117	c3clwF_	Alignment	not modelled	98.7	12 PDB header: structural genomics, unknown function Chain: F: PDB Molecule: conserved exported protein; PDBTitle: crystal structure of conserved exported protein from bacteroides2 fragilis
118	d1aq0a_	Alignment	not modelled	98.7	15 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
119	c3f55A_	Alignment	not modelled	98.7	15 PDB header: hydrolase, allergen Chain: A: PDB Molecule: beta-1,3-glucanase; PDBTitle: crystal structure of the native endo beta-1,3-glucanase (hev b 2), a2 major allergen from hevea brasiliensis (space group p41)
120	c1nofA_	Alignment	not modelled	98.7	22 PDB header: hydrolase Chain: A: PDB Molecule: xylanase; PDBTitle: the first crystallographic structure of a xylanase from2 glycosyl hydrolase family 5: implications for catalysis