







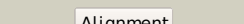

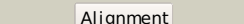

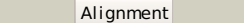

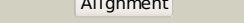

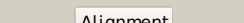

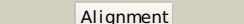





#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3lpgA_	 Alignment		100.0	98	PDB header: hydrolase/hydrolase inhibitor Chain: A: PDB Molecule: beta-glucuronidase; PDBTitle: structure of e. coli beta-glucuronidase bound with a novel, potent2 inhibitor 3-(2-fluorophenyl)-1-(2-hydroxyethyl)-1-((6-methyl-2-oxo-1,3,2-dihydroquinolin-3-yl)methyl)urea
2	c1bhgB_	 Alignment		100.0	48	PDB header: glycosidase Chain: B: PDB Molecule: beta-glucuronidase; PDBTitle: human beta-glucuronidase at 2.6 a resolution
3	c3fn9B_	 Alignment		100.0	24	PDB header: hydrolase Chain: B: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
4	c3cmgA_	 Alignment		100.0	28	PDB header: hydrolase Chain: A: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
5	c3mv14_	 Alignment		100.0	26	PDB header: hydrolase Chain: 4: PDB Molecule: beta-galactosidase; PDBTitle: e.coli (lacZ) beta-galactosidase (r599a) in complex with guanidinium
6	c1jz6C_	 Alignment		100.0	26	PDB header: hydrolase Chain: C: PDB Molecule: beta-galactosidase; PDBTitle: e. coli (lacZ) beta-galactosidase in complex with galacto-2 tetrazole
7	c3obaA_	 Alignment		100.0	23	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: structure of the beta-galactosidase from kluyveromyces fragilis
8	c3gm8A_	 Alignment		100.0	23	PDB header: hydrolase Chain: A: PDB Molecule: glycoside hydrolase family 2, candidate beta-glycosidase; PDBTitle: crystal structure of a beta-glycosidase from bacteroides vulgatus
9	c3bgaB_	 Alignment		100.0	23	PDB header: hydrolase Chain: B: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of beta-galactosidase from bacteroides2 thetaiotaomicron vpi-5482
10	c1yq2C_	 Alignment		100.0	26	PDB header: hydrolase Chain: C: PDB Molecule: beta-galactosidase; PDBTitle: beta-galactosidase from arthrobacter sp. c2-2 (isoenzyme c2-2 2-1)
11	c2je8B_	 Alignment		100.0	18	PDB header: hydrolase Chain: B: PDB Molecule: beta-mannosidase; PDBTitle: structure of a beta-mannosidase from bacteroides2 thetaiotaomicron

12	c2vzvB_	Alignment		100.0	17	PDB header: hydrolase Chain: B: PDB Molecule: exo-beta-d-glucosaminidase; PDBTitle: substrate complex of amycolatopsis orientalis exo-2 chitosanase csxa e541a with chitosan
13	d1bhga3	Alignment		100.0	51	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
14	d1yq2a5	Alignment		100.0	29	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
15	d1jz8a5	Alignment		100.0	25	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
16	d2je8a5	Alignment		100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
17	d2vzsa5	Alignment		100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
18	d1bhga2	Alignment		100.0	46	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
19	d1uuqa_	Alignment		100.0	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
20	c1uz4A_	Alignment		100.0	15	PDB header: hydrolase Chain: A: PDB Molecule: man5a; PDBTitle: common inhibition of beta-glucosidase and beta-mannosidase2 by isofagomine lactam reflects different conformational3 itinieraries for glucoside and mannoside hydrolysis
21	c2w62A_	Alignment	not modelled	100.0	13	PDB header: transferase Chain: A: PDB Molecule: glycolipid-anchored surface protein 2; PDBTitle: saccharomyces cerevisiae gas2p in complex with2 laminariipentaose
22	d1jz8a3	Alignment	not modelled	100.0	24	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
23	d1tg7a5	Alignment	not modelled	100.0	12	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Glycosyl hydrolases family 35 catalytic domain
24	d1yq2a3	Alignment	not modelled	100.0	25	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
25	d1rh9a1	Alignment	not modelled	100.0	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
26	d2c0ha1	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
27	d2je8a4	Alignment	not modelled	100.0	19	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
28	d1qnra_	Alignment	not modelled	100.0	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
29	c3eivA_	Alignment	not modelled	99.9	12	PDB header: hydrolase Chain: A: PDB Molecule: endo-beta-1,4-mannanase;

29	c3civA	Alignment	not modelled	99.9	12	PDBTitle: crystal structure of the endo-beta-1,4-mannanase from2 alicyclobacillus acidocaldarius
30	d1wkya2	Alignment	not modelled	99.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
31	d7a3ha	Alignment	not modelled	99.9	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
32	c3jugA	Alignment	not modelled	99.9	15	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
33	d1egza	Alignment	not modelled	99.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
34	d2vzsa4	Alignment	not modelled	99.9	18	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
35	c3pzvB	Alignment	not modelled	99.9	21	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: c2 crystal form of the endo-1,4-beta-glucanase from bacillus subtilis2 168
36	c3thdD	Alignment	not modelled	99.9	15	PDB header: hydrolase Chain: D: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of human beta-galactosidase in complex with 1-2 deoxygalactonojirimycin
37	d1tvna1	Alignment	not modelled	99.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
38	c2cksB	Alignment	not modelled	99.9	18	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)
39	d1bqca	Alignment	not modelled	99.9	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
40	d1ecea	Alignment	not modelled	99.9	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
41	c1wkya	Alignment	not modelled	99.9	16	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
42	d1kwga2	Alignment	not modelled	99.9	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
43	c2zunB	Alignment	not modelled	99.9	19	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
44	d1tg7a2	Alignment	not modelled	99.9	17	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
45	c3d3aA	Alignment	not modelled	99.9	12	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of a beta-galactosidase from bacteroides2 thetaiotaomicron
46	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
47	c2oylB	Alignment	not modelled	99.9	17	PDB header: hydrolase Chain: B: PDB Molecule: endoglycoceramidase ii; PDBTitle: endo-glycoceramidase ii from rhodococcus sp.: cellobiose-like2 imidazole complex
48	d1higa	Alignment	not modelled	99.9	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
49	d1hjsa	Alignment	not modelled	99.8	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
50	d1vbua1	Alignment	not modelled	99.8	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
51	d1g01a	Alignment	not modelled	99.8	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
52	d1v6ya	Alignment	not modelled	99.8	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
53	c1kwgA	Alignment	not modelled	99.8	14	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of thermus thermophilus a4 beta-galactosidase
54	d1fh9a	Alignment	not modelled	99.8	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
						Fold: TIM beta/alpha-barrel

55	d1nq6a_	Alignment	not modelled	99.8	17	Superfamily: (Trans)glycosidases Family: beta-glycanases
56	d1w91a2	Alignment	not modelled	99.8	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
57	c3ogrA_	Alignment	not modelled	99.8	11	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: complex structure of beta-galactosidase from trichoderma reesei with2 galactose
58	d1uhva2	Alignment	not modelled	99.8	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
59	d1n82a_	Alignment	not modelled	99.8	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
60	d1vjza_	Alignment	not modelled	99.8	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
61	c1xc6A_	Alignment	not modelled	99.8	12	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: native structure of beta-galactosidase from penicillium sp. in complex2 with galactose
62	c3emzA_	Alignment	not modelled	99.8	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
63	d1v6wa2	Alignment	not modelled	99.8	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
64	d1ur4a_	Alignment	not modelled	99.8	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
65	d1us3a2	Alignment	not modelled	99.8	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
66	d1w32a_	Alignment	not modelled	99.8	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
67	c1xyzA_	Alignment	not modelled	99.8	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
68	d1xyza_	Alignment	not modelled	99.8	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
69	c2depA_	Alignment	not modelled	99.8	17	PDB header: hydrolase Chain: A: PDB Molecule: thermostable cellobiohydrolase; PDBTitle: crystal structure of xylanase b from clostridium2 stercorarium f9
70	d1v0la_	Alignment	not modelled	99.8	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
71	c2jepB_	Alignment	not modelled	99.7	13	PDB header: hydrolase Chain: B: PDB Molecule: xyloglucanase; PDBTitle: native family 5 xyloglucanase from paenibacillus pabuli
72	c2cncA_	Alignment	not modelled	99.7	15	PDB header: hydrolase Chain: A: PDB Molecule: endoxylanase; PDBTitle: family 10 xylanase
73	c2bs9B_	Alignment	not modelled	99.7	16	PDB header: hydrolase Chain: B: PDB Molecule: beta-xylosidase; PDBTitle: native crystal structure of a gh39 beta-xylosidase xynb12 from geobacillus stearothermophilus
74	d1ceoa_	Alignment	not modelled	99.7	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
75	d1ta3b_	Alignment	not modelled	99.7	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
76	d1h1na_	Alignment	not modelled	99.7	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
77	c3ncoA_	Alignment	not modelled	99.7	19	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase fncel5a; PDBTitle: crystal structure of fncel5a from f. nodosum rt17-b1
78	d1ur1a_	Alignment	not modelled	99.7	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
79	d1ilwa_	Alignment	not modelled	99.7	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
80	d1foba_	Alignment	not modelled	99.7	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
81	c3aysA_	Alignment	not modelled	99.7	18	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase; PDBTitle: gh5 endoglucanase from a ruminal fungus in complex with cellotriose
82	d2pb1a1	Alignment	not modelled	99.7	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases

					Family: beta-glycanases
83	c3ndyA	Alignment	not modelled	99.7	17 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
84	d1bg4a	Alignment	not modelled	99.7	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
85	c1uhvD	Alignment	not modelled	99.7	14 PDB header: hydrolase Chain: D: PDB Molecule: beta-xylosidase; PDBTitle: crystal structure of beta-d-xylosidase from2 thermoanaerobacterium saccharolyticum, a family 393 glycoside hydrolase
86	c2y8kA	Alignment	not modelled	99.7	21 PDB header: hydrolase Chain: A: PDB Molecule: carbohydrate binding family 6; PDBTitle: structure of ctgh5-cbm6, an arabinoxylan-specific xylanase.
87	c3qr3B	Alignment	not modelled	99.7	15 PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase eg-ii; PDBTitle: crystal structure of cel 5a (eg2) from hypocrea jecorina (trichoderma2 reesei)
88	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
89	c3mmwB	Alignment	not modelled	99.7	19 PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: crystal structure of endoglucanase cel5a from the hyperthermophilic2 thermotoga maritima
90	c3icgD	Alignment	not modelled	99.6	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
91	d1tuxa	Alignment	not modelled	99.6	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
92	d1edga	Alignment	not modelled	99.6	14 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
93	c2fglA	Alignment	not modelled	99.6	14 PDB header: hydrolase Chain: A: PDB Molecule: alkaline thermostable endoxylanase; PDBTitle: an alkali thermostable f/10 xylanase from alkalophilic2 bacillus sp. ng-27
94	d1ug6a	Alignment	not modelled	99.6	21 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
95	c3ahxC	Alignment	not modelled	99.6	18 PDB header: hydrolase Chain: C: PDB Molecule: beta-glucosidase a; PDBTitle: crystal structure of beta-glucosidase a from bacterium clostridium2 cellulovorans
96	d1e4ia	Alignment	not modelled	99.6	19 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
97	c2j75A	Alignment	not modelled	99.6	20 PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase a; PDBTitle: beta-glucosidase from thermotoga maritima in complex with2 noeuromycin
98	d1r85a	Alignment	not modelled	99.6	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
99	d1h4pa	Alignment	not modelled	99.6	14 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
100	c2z1sA	Alignment	not modelled	99.6	23 PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase b; PDBTitle: beta-glucosidase b from paenibacillus polymyxa complexed2 with cellotetraose
101	d1jz8a1	Alignment	not modelled	99.5	29 Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
102	d1qoxa	Alignment	not modelled	99.5	21 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
103	c3qomA	Alignment	not modelled	99.5	19 PDB header: hydrolase Chain: A: PDB Molecule: 6-phospho-beta-glucosidase; PDBTitle: crystal structure of 6-phospho-beta-glucosidase from lactobacillus2 plantarum
104	d1vffa1	Alignment	not modelled	99.5	22 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
105	c3fiyA	Alignment	not modelled	99.5	20 PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of bg1b
106	c2w5fB	Alignment	not modelled	99.5	17 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
107	c3u7vA	Alignment	not modelled	99.5	15 PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: the structure of a putative beta-galactosidase from caulobacter2 crescentus cb15.
					Fold: TIM beta/alpha-barrel

108	d1pbga_	Alignment	not modelled	99.4	18	Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
109	c3pn8A_	Alignment	not modelled	99.4	20	PDB header: hydrolase Chain: A: PDB Molecule: putative phospho-beta-glucosidase; PDBTitle: the crystal structure of 6-phospho-beta-glucosidase from streptococcus2 mutans ua159
110	d1tg7a3	Alignment	not modelled	99.4	11	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
111	d2j78a1	Alignment	not modelled	99.4	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
112	d1gnxa_	Alignment	not modelled	99.4	23	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
113	d1qvba_	Alignment	not modelled	99.4	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
114	d1aq0a_	Alignment	not modelled	99.4	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
115	c2xhyD_	Alignment	not modelled	99.4	20	PDB header: hydrolase Chain: D: PDB Molecule: 6-phospho-beta-glucosidase bgla; PDBTitle: crystal structure of e.coli bgla
116	c3kzsD_	Alignment	not modelled	99.4	18	PDB header: hydrolase Chain: D: PDB Molecule: glycosyl hydrolase family 5; PDBTitle: crystal structure of glycosyl hydrolase family 5 (np_809925.1) from2 bacteroides thetaiotaomicron vpi-5482 at 2.10 a resolution
117	c2dgaA_	Alignment	not modelled	99.4	22	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of hexameric beta-glucosidase in wheat
118	d1v02a_	Alignment	not modelled	99.4	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
119	c1v02F_	Alignment	not modelled	99.4	21	PDB header: hydrolase Chain: F: PDB Molecule: dhurrinase; PDBTitle: crystal structure of the sorghum bicolor dhurrinase 1
120	d1uwsa_	Alignment	not modelled	99.4	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase