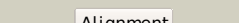
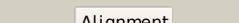
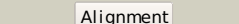
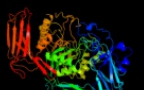


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3mv14_	 Alignment		100.0	100	PDB header: hydrolase Chain: 4: PDB Molecule: beta-galactosidase; PDBTitle: e.coli (lacZ) beta-galactosidase (r599a) in complex with guanidinium
2	c1jz6C_	 Alignment		100.0	100	PDB header: hydrolase Chain: C: PDB Molecule: beta-galactosidase; PDBTitle: e. coli (lacZ) beta-galactosidase in complex with galacto-2 tetrazole
3	c3obaA_	 Alignment		100.0	32	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: structure of the beta-galactosidase from kluyveromyces lactis
4	c3bgaB_	 Alignment		100.0	33	PDB header: hydrolase Chain: B: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of beta-galactosidase from bacteroides2 thetaiotaomicron vpi-5482
5	c1yq2C_	 Alignment		100.0	35	PDB header: hydrolase Chain: C: PDB Molecule: beta-galactosidase; PDBTitle: beta-galactosidase from arthrobacter sp. c2-2 (isoenzyme c2-2 2-1)
6	c3gm8A_	 Alignment		100.0	24	PDB header: hydrolase Chain: A: PDB Molecule: glycoside hydrolase family 2, candidate beta-glycosidase; PDBTitle: crystal structure of a beta-glycosidase from bacteroides vulgatus
7	c3cmgA_	 Alignment		100.0	21	PDB header: hydrolase Chain: A: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
8	c3fn9B_	 Alignment		100.0	21	PDB header: hydrolase Chain: B: PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
9	c3lpgA_	 Alignment		100.0	27	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
10	c2je8B_	 Alignment		100.0	20	PDB header: hydrolase Chain: B: PDB Molecule: beta-mannosidase; PDBTitle: structure of a beta-mannosidase from bacteroides2 thetaiotaomicron
11	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

12	c1bhgB_	Alignment		100.0	24	PDB header: glycosidase Chain: B: PDB Molecule: beta-glucuronidase; PDBTitle: human beta-glucuronidase at 2.6 a resolution
13	d1jz8a4	Alignment		100.0	100	Fold: Supersandwich Superfamily: Galactose mutarotase-like Family: beta-Galactosidase, domain 5
14	d1yq2a4	Alignment		100.0	24	Fold: Supersandwich Superfamily: Galactose mutarotase-like Family: beta-Galactosidase, domain 5
15	d1jz8a5	Alignment		100.0	100	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
16	d1yq2a5	Alignment		100.0	43	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
17	d1jz8a3	Alignment		100.0	100	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
18	d1yq2a3	Alignment		100.0	44	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
19	d2je8a5	Alignment		100.0	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
20	d2vzsa5	Alignment		100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
21	d1bhga3	Alignment	not modelled	100.0	23	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
22	d1uuqa_	Alignment	not modelled	100.0	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
23	c1uz4A_	Alignment	not modelled	100.0	21	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
24	d1tg7a5	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Glycosyl hydrolases family 35 catalytic domain
25	d1bhga2	Alignment	not modelled	100.0	22	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
26	c2w62A_	Alignment	not modelled	100.0	16	PDB header: transferase Chain: A: PDB Molecule: glycolipid-anchored surface protein 2; PDBTitle: saccharomyces cerevisiae gas2p in complex with2 laminaripentaose
27	d2je8a4	Alignment	not modelled	100.0	21	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
28	d2c0ha1	Alignment	not modelled	100.0	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases

29	d1wkya2	Alignment	not modelled	100.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
30	d2vzsa4	Alignment	not modelled	100.0	20	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
31	d1qnra	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
32	d1rh9a1	Alignment	not modelled	100.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
33	c3jugA	Alignment	not modelled	99.9	17	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
34	d1tvna1	Alignment	not modelled	99.9	11	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
35	d1bqca	Alignment	not modelled	99.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
36	d7a3ha	Alignment	not modelled	99.9	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
37	d1egza	Alignment	not modelled	99.9	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
38	c3pzvB	Alignment	not modelled	99.9	15	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: c2 crystal form of the endo-1,4-beta-glucanase from bacillus subtilis2 168
39	c2cksB	Alignment	not modelled	99.9	16	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)
40	d1jz8a2	Alignment	not modelled	99.9	100	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
41	c3civA	Alignment	not modelled	99.9	16	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
42	d1ecea	Alignment	not modelled	99.9	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
43	d1tg7a2	Alignment	not modelled	99.9	14	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
44	c1wkyA	Alignment	not modelled	99.9	19	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
45	c3thdD	Alignment	not modelled	99.9	17	PDB header: hydrolase Chain: D: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of human beta-galactosidase in complex with 1-2 deoxygalactonojirimycin
46	d1yq2a1	Alignment	not modelled	99.8	28	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
47	d1jz8a1	Alignment	not modelled	99.8	100	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
48	c2zunB	Alignment	not modelled	99.8	17	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
49	d1kwga2	Alignment	not modelled	99.7	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
50	c3d3aA	Alignment	not modelled	99.7	13	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of a beta-galactosidase from bacteroides2 thetaiotaomicron
51	c2oylB	Alignment	not modelled	99.6	20	PDB header: hydrolase Chain: B: PDB Molecule: endoglycoceramidase ii; PDBTitle: endo-glycoceramidase ii from rhodococcus sp.: cellobiose-like2 imidazole complex
52	c3ogrA	Alignment	not modelled	99.6	14	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: complex structure of beta-galactosidase from trichoderma reesei with2 galactose
53	d2vzsa1	Alignment	not modelled	99.6	15	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
54	d1g01a	Alignment	not modelled	99.6	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
55	c1xc6A	Alignment	not modelled	99.6	14	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase;

55	c1xcoA	Alignment	not modelled	99.6	14	PDBTitle: native structure of beta-galactosidase from penicillium sp. in complex2 with galactose PDB header: hydrolase
56	c3pzqA	Alignment	not modelled	99.6	16	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
57	c1kwgA	Alignment	not modelled	99.5	20	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of thermus thermophilus a4 beta-galactosidase
58	d1vjza	Alignment	not modelled	99.5	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
59	d1tg7a3	Alignment	not modelled	99.5	21	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
60	d1yq2a2	Alignment	not modelled	99.5	27	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
61	d1bhga1	Alignment	not modelled	99.5	22	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
62	d2je8a1	Alignment	not modelled	99.4	21	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
63	d1hjqa	Alignment	not modelled	99.4	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
64	c3ncoA	Alignment	not modelled	99.3	15	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase fncel5a; PDBTitle: crystal structure of fncel5a from f. nodosum rt17-b1
65	d1ceoa	Alignment	not modelled	99.3	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
66	c2jepB	Alignment	not modelled	99.3	17	PDB header: hydrolase Chain: B: PDB Molecule: xyloglucanase; PDBTitle: native family 5 xyloglucanase from paenibacillus pabuli
67	c3aysA	Alignment	not modelled	99.2	15	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase; PDBTitle: gh5 endoglucanase from a ruminal fungus in complex with cellotriose
68	c3mmwB	Alignment	not modelled	99.2	18	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: crystal structure of endoglucanase cel5a from the hyperthermophilic2 thermotoga maritima
69	c3icgD	Alignment	not modelled	99.2	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
70	d1fh9a	Alignment	not modelled	99.2	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
71	d1v6wa2	Alignment	not modelled	99.2	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
72	d1vbua1	Alignment	not modelled	99.2	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
73	d1w91a2	Alignment	not modelled	99.2	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
74	d1hjsa	Alignment	not modelled	99.2	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
75	d2pb1a1	Alignment	not modelled	99.1	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
76	d1h1na	Alignment	not modelled	99.1	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
77	c3l55B	Alignment	not modelled	99.1	16	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
78	d1uhva2	Alignment	not modelled	99.1	12	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
79	d1nq6a	Alignment	not modelled	99.1	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
80	c3qr3B	Alignment	not modelled	99.1	17	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase eg-ii; PDBTitle: crystal structure of cel5a (eg2) from hypocrea jecorina (trichoderma2 reesei)
81	c3ndyA	Alignment	not modelled	99.1	13	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
82	c3u7vA_	Alignment	not modelled	99.1	18 PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: the structure of a putative beta-galactosidase from caulobacter2 crescentus cb15.
83	d1v6ya_	Alignment	not modelled	99.1	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
84	d1v0la_	Alignment	not modelled	99.0	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
85	d1h4pa_	Alignment	not modelled	98.9	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
86	d1ur4a_	Alignment	not modelled	98.9	15 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
87	c2y8kA_	Alignment	not modelled	98.9	22 PDB header: hydrolase Chain: A: PDB Molecule: carbohydrate binding family 6; PDBTitle: structure of ctgh5-cbm6, an arabinoxylan-specific xylanase.
88	d1ilwa_	Alignment	not modelled	98.9	18 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
89	c3kzsD_	Alignment	not modelled	98.9	17 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
90	d1us3a2	Alignment	not modelled	98.9	16 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
91	d1n82a_	Alignment	not modelled	98.8	15 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
92	d1ta3b_	Alignment	not modelled	98.8	14 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
93	d1w32a_	Alignment	not modelled	98.8	21 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
94	d1edga_	Alignment	not modelled	98.8	19 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
95	c2depA_	Alignment	not modelled	98.7	14 PDB header: hydrolase Chain: A: PDB Molecule: thermostable celloxylanase; PDBTitle: crystal structure of xylanase b from clostridium2 stercorarium f9
96	c3emzA_	Alignment	not modelled	98.7	10 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
97	d1tuxa_	Alignment	not modelled	98.7	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
98	d1bg4a_	Alignment	not modelled	98.7	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
99	c1xyzA_	Alignment	not modelled	98.7	15 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
100	d1xyza_	Alignment	not modelled	98.7	15 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
101	c2okxB_	Alignment	not modelled	98.6	28 PDB header: hydrolase Chain: B: PDB Molecule: rhamnosidase b; PDBTitle: crystal structure of gh78 family rhamnosidase of bacillus sp. gl1 at2 1.9 a
102	d1ur1a_	Alignment	not modelled	98.6	13 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
103	d1foba_	Alignment	not modelled	98.6	17 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
104	c2bs9B_	Alignment	not modelled	98.5	15 PDB header: hydrolase Chain: B: PDB Molecule: beta-xylosidase; PDBTitle: native crystal structure of a gh39 beta-xylosidase xynb12 from geobacillus stearothermophilus
105	c2cncA_	Alignment	not modelled	98.5	14 PDB header: hydrolase Chain: A: PDB Molecule: endoxylanase; PDBTitle: family 10 xylanase
106	c1uhvD_	Alignment	not modelled	98.4	13 PDB header: hydrolase Chain: D: PDB Molecule: beta-xylosidase; PDBTitle: crystal structure of beta-d-xylosidase from2 thermoanaerobacterium saccharolyticum, a family 393 glycoside hydrolase
107	d1ug6a_	Alignment	not modelled	98.2	13 Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
					PDB header: hydrolase

108	c3ahxC_	Alignment	not modelled	98.0	12	Chain: C: PDB Molecule: beta-glucosidase a; PDBTitle: crystal structure of beta-glucosidase a from bacterium clostridium2 cellulovorans
109	d1r85a_	Alignment	not modelled	98.0	12	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
110	c1iszA_	Alignment	not modelled	98.0	16	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
111	c2vrgB_	Alignment	not modelled	98.0	14	PDB header: hydrolase Chain: B: PDB Molecule: alpha-l-arabinofuranosidase; PDBTitle: structure of an inactive mutant of arabinofuranosidase from2 thermobacillus xylanilyticus in complex with a3 pentasaccharide
112	c2j75A_	Alignment	not modelled	97.9	13	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase a; PDBTitle: beta-glucosidase from thermotoga maritima in complex with2 noeuromycin
113	d1aq0a_	Alignment	not modelled	97.8	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
114	c2w5fB_	Alignment	not modelled	97.8	13	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
115	d1e4ia_	Alignment	not modelled	97.7	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
116	c2fglA_	Alignment	not modelled	97.7	11	PDB header: hydrolase Chain: A: PDB Molecule: alkaline thermostable endoxylanase; PDBTitle: an alkali thermostable f/10 xylanase from alkalophilic2 bacillus sp. ng-27
117	c3fiyA_	Alignment	not modelled	97.6	17	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of bglb
118	d1vffa1	Alignment	not modelled	97.6	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
119	d2cyga1	Alignment	not modelled	97.5	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
120	d1vema2	Alignment	not modelled	97.4	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain