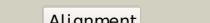
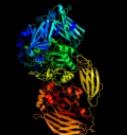
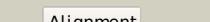
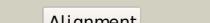
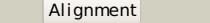
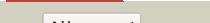
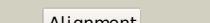
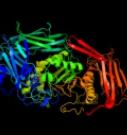


Phyre²

Email	i.a.kelley@imperial.ac.uk
Description	P06864
Date	Thu Jan 5 10:59:27 GMT 2012
Unique Job ID	de0c3862a9060e5a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3mv14_			100.0	33	PDB header: hydrolase Chain: 4; PDB Molecule: beta-galactosidase; PDBTitle: e.coli (lacZ) beta-galactosidase (r599a) in complex with guanidinium
2	c1jz6C_			100.0	34	PDB header: hydrolase Chain: C; PDB Molecule: beta-galactosidase; PDBTitle: e. coli (lacZ) beta-galactosidase in complex with galacto-2 tetrazole
3	c3bgab_			100.0	32	PDB header: hydrolase Chain: B; PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of beta-galactosidase from bacteroides2 thetaiotaomicron vpi-5482
4	c3obaA_			100.0	36	PDB header: hydrolase Chain: A; PDB Molecule: beta-galactosidase; PDBTitle: structure of the beta-galactosidase from kluyveromyces lactis
5	c1yq2C_			100.0	33	PDB header: hydrolase Chain: C; PDB Molecule: beta-galactosidase; PDBTitle: beta-galactosidase from arthrobacter sp. c2-2 (isoenzyme c2-2 2-1)
6	c3gm8A_			100.0	21	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	c3fn9B_			100.0	23	PDB header: hydrolase Chain: B; PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
8	c3cmgA_			100.0	24	PDB header: hydrolase Chain: A; PDB Molecule: putative beta-galactosidase; PDBTitle: crystal structure of putative beta-galactosidase from bacteroides2 fragilis
9	c2je8B_			100.0	19	PDB header: hydrolase Chain: B; PDB Molecule: beta-mannosidase; PDBTitle: structure of a beta-mannosidase from bacteroides2 thetaiotaomicron
10	c3lpgA_			100.0	24	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-di hydroquinolin-3-yl)methyl)urea
11	c2vzvB_			100.0	16	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			100.0	24	PDB header: glycosidase Chain: B: PDB Molecule: beta-glucuronidase; PDBTitle: human beta-glucuronidase at 2.6 a resolution
13	d1jz8a4			100.0	21	Fold: Supersandwich Superfamily: Galactose mutarotase-like Family: beta-Galactosidase, domain 5
14	d1yq2a4			100.0	24	Fold: Supersandwich Superfamily: Galactose mutarotase-like Family: beta-Galactosidase, domain 5
15	d1yq2a5			100.0	43	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
16	d1jz8a5			100.0	43	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
17	d1jz8a3			100.0	42	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
18	d1yq2a3			100.0	39	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
19	d2je8a5			100.0	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
20	d2vzsa5			100.0	13	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
21	d1uuqa		not modelled	100.0	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
22	c1uz4A		not modelled	100.0	20	PDB header: hydrolase Chain: A: PDB Molecule: man5a; PDBTitle: common inhibition of beta-glucosidase and beta-mannosidase2 by isofagomine lactam reflects different conformational3 intineraries for glucoside and mannose hydrolysis
23	d1bhga3		not modelled	100.0	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
24	d1tg7a5		not modelled	100.0	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Glycosyl hydrolases family 35 catalytic domain
25	d1bhga2		not modelled	100.0	21	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
26	d2je8a4		not modelled	100.0	20	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
27	c2w62A		not modelled	100.0	19	PDB header: transferase Chain: A: PDB Molecule: glycolipid-anchored surface protein 2; PDBTitle: saccharomyces cerevisiae gas2p in complex with2 laminarinpentaoe
28	d2c0ha1		not modelled	100.0	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases

29	d2vzsa4	Alignment	not modelled	100.0	20	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: beta-Galactosidase/glucuronidase, N-terminal domain
30	d1wkya2	Alignment	not modelled	99.9	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
31	d1rh9a1	Alignment	not modelled	99.9	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
32	d1qnra	Alignment	not modelled	99.9	18	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	12	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
35	d1bqca	Alignment	not modelled	99.9	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
36	d7a3ha	Alignment	not modelled	99.9	15	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	18	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	14	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase e-5; PDBTitle: x-ray crystal structure of the catalytic domain of thermobifida fusca endoglucanase cel5a (e5)
40	d1tg7a2	Alignment	not modelled	99.9	16	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
41	c3civA	Alignment	not modelled	99.9	19	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	d1ceea	Alignment	not modelled	99.9	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
43	c3thdD	Alignment	not modelled	99.9	18	PDB header: hydrolase Chain: D: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of human beta-galactosidase in complex with 1-2 deoxygalactonojirimycin
44	c1wkyA	Alignment	not modelled	99.8	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
45	d1yq2a1	Alignment	not modelled	99.8	22	Fold: immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
46	d1jz8a2	Alignment	not modelled	99.8	29	Fold: immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
47	d1jz8a1	Alignment	not modelled	99.8	32	Fold: immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
48	c2zunB	Alignment	not modelled	99.7	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	c3d3aA	Alignment	not modelled	99.7	17	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: crystal structure of a beta-galactosidase from bacteroides2 thetaiotaomicron
50	c3ogrA	Alignment	not modelled	99.7	14	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: complex structure of beta-galactosidase from trichoderma reesei with2 galactose
51	c1xc6A	Alignment	not modelled	99.7	12	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: native structure of beta-galactosidase from penicillium sp. in complex2 with galactose
52	d1kwga2	Alignment	not modelled	99.7	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Amylase, catalytic domain
53	d2vzsa1	Alignment	not modelled	99.6	21	Fold: immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
54	c2oylB	Alignment	not modelled	99.6	17	PDB header: hydrolase Chain: B: PDB Molecule: endoglycoceramidase ii; PDBTitle: endo-glycoceramidase ii from rhodococcus sp.: cellobiose-like2 imidazole complex
55	c1wkyA	All	not modelled	99.5	21	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase;

55	c1nwym	Alignment	not modelled	99.5	21	PDBTitle: crystal structure of thermus thermophilus a4 beta-galactosidase
56	d2je8a1	Alignment	not modelled	99.5	16	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
57	dlg01a	Alignment	not modelled	99.5	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
58	c3pzqA	Alignment	not modelled	99.5	18	PDB header: hydrolase Chain: A: PDB Molecule: mannan endo-1,4-beta-mannosidase, glycosyl hydrolase family PDBTitle: structure of the hyperthermstable endo-1,4-beta-d-mannanase from2 thermotoga petrophila rku-1 with maltose and glycerol
59	d1vjza	Alignment	not modelled	99.5	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
60	d1tg7a3	Alignment	not modelled	99.4	18	Fold: Galactose-binding domain-like Superfamily: Galactose-binding domain-like Family: Beta-galactosidase LacA, domains 4 and 5
61	d1yq2a2	Alignment	not modelled	99.4	25	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
62	d1hjqa	Alignment	not modelled	99.4	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
63	d1bhga1	Alignment	not modelled	99.4	25	Fold: Immunoglobulin-like beta-sandwich Superfamily: beta-Galactosidase/glucuronidase domain Family: beta-Galactosidase/glucuronidase domain
64	c3ncoA	Alignment	not modelled	99.4	16	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	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
66	c2jepB	Alignment	not modelled	99.2	14	PDB header: hydrolase Chain: B: PDB Molecule: xyloglucanase; PDBTitle: native family 5 xyloglucanase from paenibacillus pabuli
67	d1vbua1	Alignment	not modelled	99.2	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
68	c3l55B	Alignment	not modelled	99.2	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
69	d1hjsa	Alignment	not modelled	99.2	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
70	c3aysA	Alignment	not modelled	99.2	14	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase; PDBTitle: gh5 endoglucanase from a ruminal fungus in complex with cellotriose
71	d1h1na	Alignment	not modelled	99.2	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
72	c3mmwB	Alignment	not modelled	99.1	12	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase; PDBTitle: crystal structure of endoglucanase cel5a from the hyperthermophilic2 thermotoga maritima
73	d1w91a2	Alignment	not modelled	99.1	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
74	c3icgD	Alignment	not modelled	99.1	15	PDB header: hydrolase Chain: D: PDB Molecule: endoglucanase d; PDBTitle: crystal structure of the catalytic and carbohydrate binding domain2 of endoglucanase d from clostridium cellulovorans
75	c3qr3B	Alignment	not modelled	99.1	13	PDB header: hydrolase Chain: B: PDB Molecule: endoglucanase eg-ii; PDBTitle: crystal structure of cel5a (eg2) from hypocreah jecorina (trichoderma2 reesei)
76	d1nq6a	Alignment	not modelled	99.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
77	d1uhva2	Alignment	not modelled	99.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
78	d1ur4a	Alignment	not modelled	99.0	16	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
79	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.
80	d1v6wa2	Alignment	not modelled	99.0	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
81	d1v6ya	Alignment	not modelled	99.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases

82	c3ndyA	Alignment	not modelled	98.9	13	PDB header: hydrolase Chain: A: PDB Molecule: endoglucanase d; PDBTitle: the structure of the catalytic and carbohydrate binding domain of 2 endoglucanase d from clostridium cellulovorans
83	c3u7vA	Alignment	not modelled	98.9	11	PDB header: hydrolase Chain: A: PDB Molecule: beta-galactosidase; PDBTitle: the structure of a putative beta-galactosidase from caulobacter2 crescentus cb15.
84	d1v0la	Alignment	not modelled	98.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
85	d1edga	Alignment	not modelled	98.9	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
86	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 thetaotaomicron vpi-5482 at 2.10 a resolution
87	d1fh9a	Alignment	not modelled	98.9	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
88	d1us3a2	Alignment	not modelled	98.9	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
89	d1n82a	Alignment	not modelled	98.9	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
90	d1i1wa	Alignment	not modelled	98.9	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
91	c2bs9B	Alignment	not modelled	98.8	15	PDB header: hydrolase Chain: B: PDB Molecule: beta-xylosidase; PDBTitle: native crystal structure of a gh39 beta-xylosidase xynb12 from geobacillus stearothermophilus
92	d1w32a	Alignment	not modelled	98.8	20	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
93	d2pb1a1	Alignment	not modelled	98.8	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
94	d1ta3b	Alignment	not modelled	98.8	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
95	d1h4pa	Alignment	not modelled	98.7	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
96	c3emzA	Alignment	not modelled	98.7	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
97	d1bg4a	Alignment	not modelled	98.7	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
98	c1xyzA	Alignment	not modelled	98.6	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
99	d1xyza	Alignment	not modelled	98.6	15	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
100	d1tuxa	Alignment	not modelled	98.6	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
101	d1urla	Alignment	not modelled	98.6	14	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
102	d1foba	Alignment	not modelled	98.6	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
103	c2depA	Alignment	not modelled	98.6	17	PDB header: hydrolase Chain: A: PDB Molecule: thermostable cellobiohydrolase; PDBTitle: crystal structure of xylanase b from clostridium2 stercorarium f9
104	c2cncA	Alignment	not modelled	98.6	15	PDB header: hydrolase Chain: A: PDB Molecule: endoxylanase; PDBTitle: family 10 xylanase
105	c2okxB	Alignment	not modelled	98.5	17	PDB header: hydrolase Chain: B: PDB Molecule: rhamnosidase b; PDBTitle: crystal structure of gh78 family rhamnosidase of bacillus sp. g1 at2 1.9 a
106	d1ug6a	Alignment	not modelled	98.5	22	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
107	c1uhvD	Alignment	not modelled	98.5	16	PDB header: hydrolase Chain: D: PDB Molecule: beta-xylosidase; PDBTitle: crystal structure of beta-d-xylosidase from2 thermoanaerobacterium saccharolyticum, a family 393 glycoside hydrolase
108	c3ahvC	Alignment	not modelled	98.2	18	PDB header: hydrolase Chain: C: PDB Molecule: beta-glucosidase a;

100	c2qaxA	Alignment	not modelled	98.2	10	PDBTitle: crystal structure of beta-glucosidase a from bacterium clostridium2 cellulovorans PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase b; PDBTitle: beta-glucosidase b from paenibacillus polymyxia complexed2 with cellobetaose
109	c2z1sA	Alignment	not modelled	98.2	25	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
110	d1e4ia	Alignment	not modelled	98.1	21	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase a; PDBTitle: beta-glucosidase from thermotoga maritima in complex with2 noeuromycin
111	c2j75A	Alignment	not modelled	98.1	15	PDB header: hydrolase Chain: A: PDB Molecule: 6-phospho-beta-glucosidase; PDBTitle: crystal structure of 6-phospho-beta-glucosidase from lactobacillus2 plantarum
112	c3qomA	Alignment	not modelled	98.1	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
113	d1r85a	Alignment	not modelled	98.0	18	PDB header: hydrolase Chain: A: PDB Molecule: putative phospho-beta-glucosidase; PDBTitle: the crystal structure of 6-phospho-beta-glucosidase from streptococcus2 mutans ua159
114	d1vffa1	Alignment	not modelled	98.0	17	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
115	c3fiyA	Alignment	not modelled	97.9	25	PDB header: hydrolase Chain: A: PDB Molecule: beta-glucosidase; PDBTitle: crystal structure of bg1b
116	c3pn8A	Alignment	not modelled	97.9	19	PDB header: hydrolase Chain: A: PDB Molecule: putative phospho-beta-glucosidase; PDBTitle: the crystal structure of 6-phospho-beta-glucosidase from streptococcus2 mutans ua159
117	d1aq0a	Alignment	not modelled	97.9	18	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
118	d2cyga1	Alignment	not modelled	97.8	19	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: beta-glycanases
119	d1qoxa	Alignment	not modelled	97.7	21	Fold: TIM beta/alpha-barrel Superfamily: (Trans)glycosidases Family: Family 1 of glycosyl hydrolase
120	cliszA	Alignment	not modelled	97.7	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