

Unusual Nonreducing Sugar GlcNAc β (1 \leftrightarrow 1)Man β Formation by β -*N*-Acetylhexosaminidase from *Aspergillus oryzae*¹

Vladimír Křen*, Eva Rajnochová, Zdenka Huňková, Jana Dvořáková and Petr Sedmera

*Institute of Microbiology, Academy of Sciences of the Czech Republic, Laboratory of Biotransformation, Videňská 1083,
CZ 142 20 Prague 4, Czech Republic*

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Abstract: β -*N*-Acetylhexosaminidase from *Aspergillus oryzae* catalyzes the transfer of *N*-acetylglucosamine moiety from *p*NP- β -GlcNAc to mannose giving major product GlcNAc β (1 \leftrightarrow 1)Man β (**1a**), and GlcNAc β (1 \rightarrow 3)Man α / β (**2a**) and GlcNAc β (1 \rightarrow 6)Man α / β (**3a**) as two minor products. © 1998 Elsevier Science Ltd. All rights reserved.

β -*N*-Acetylhexosaminidase (E.C. 3.2.1.52) from *Aspergillus oryzae* is a good tool in enzymatic synthesis of oligosaccharides.²⁻⁴ It is able to transfer β -GlcNAc and β -GalNAc moieties in comparable yields to both glycosidic^{2,3b} and non-glycosidic substrates.^{3a} The regioselectivity towards monosaccharides is influenced by their anomeric substitution. This enzyme has been commercially available from Sigma since 1997.

β -GlcNAc bound to Man residues occurs frequently in various glycoproteins especially in multiantennary *N*-glycans. Recently, an enzymatic synthesis of GlcNAc β (1 \rightarrow 2)Man and GlcNAc β (1 \rightarrow 6)Man using reversed glycosylation with β -*N*-acetylhexosaminidase from *Bacillus circulans* was described.⁵ To extend the scope of applications of the *A. oryzae* enzyme and to prepare biologically useful oligosaccharides we have chosen mannose as the acceptor for transglycosylation reactions.

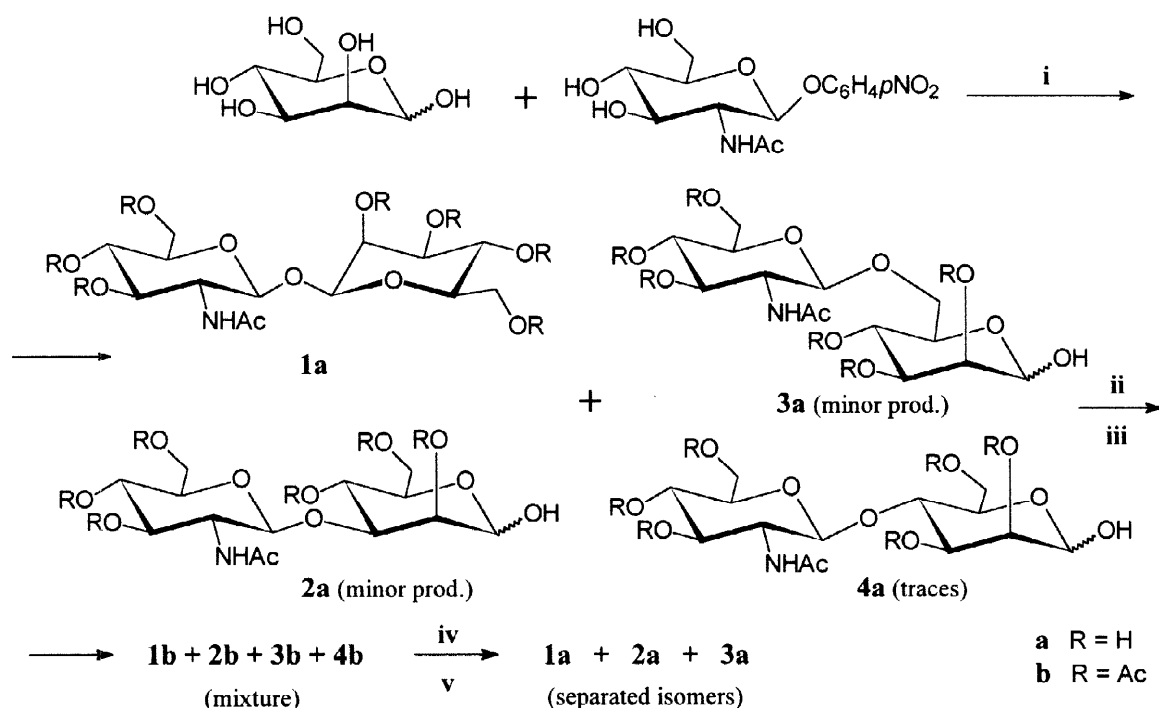
No transglycosidation product of methyl- α -mannoside was detected with various β -*N*-acetylhexosaminidases (from *Aspergillus oryzae*, *A. terreus*, *A. tamarii*,⁴ *Penicillium oxalicum*,⁶ jack beans (Sigma) and bovine kidneys (Sigma)) and *p*-nitrophenyl- β -*O*-*N*-acetylglucosaminide (*p*NP- β -GlcNAc) as a donor. However, the same reactions with mannose yielded two to four disaccharides. Further investigation was continued with enzyme from *Aspergillus oryzae* that gave the best yields of transglycosylation products.⁷

The reaction mixture was subjected to gel filtration and the disaccharide fraction was purified by semipreparative HPLC.⁸ NMR spectroscopy suggested two major components: a non-reducing sugar GlcNAc β (1 \leftrightarrow 1)Man β (**1a**) and a disaccharide(s) GlcNAc β (1 \rightarrow x)Man α / β .

To make the separation and structure elucidation easier, the crude products after the gel filtration step

were peracetylated and the anomeric hydroxyls were deprotected (piperidine/THF). Flash chromatography on silica gel neatly separated **1b** from regioisomeric disaccharides **2b** and **3b**. Large dispersion of chemical shifts and observation of N-H protons made full proton assignment possible.⁹⁻¹² The anomeric configuration of GlcNAc was β in all cases ($J_{1,2}$ 8.0 - 8.6 Hz). Mannose C-1 in **1a** and in its acetate **1b** has also β configuration ($J^{13}\text{C}(1)\text{-H}(1)$ = 165.3 Hz, ref.¹³, NOE between H-1 and H-5). In addition, NOE between both H-1 was also observed. The site of GlcNAc attachment in **2b** and **3b** was inferred from the upfield shift of the involved mannose proton and the downfield shift of the corresponding carbon (bold in footnotes^{11,12}). Some impurity signals¹¹ are consistent with the presence of the (1 \rightarrow 4) isomer **4b**.

Thus, mannose acts as an "ambident" acceptor for β -N-acetylhexosaminidases from *A. oryzae*. The formation of **1a** is probably the first example of 1-OH glycosidation by glycosidases. Two cases of enzymatic anomeric hydroxyl glycosylation have been already described, however, with glycosyltransferases: α -glucosylation of lactose¹⁴ and β -galactosylation of xylose.¹⁵



i β -N-Acetylhexosaminidase from *Aspergillus oryzae*, ii $\text{Ac}_2\text{O/Py}$, iii 5% piperidine/THF, iv flash chromatography/ SiO_2 , v MeONa/MeOH .

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7. Procedure: Mannose (7.2 g, 40 mmol) and *p*NP- β -GlcNAc (130 mg, 0.38 mmol) were dissolved by heating in McIlvain buffer (10 ml, 0.1 M, pH 5.0). After cooling, β -*N*-acetylhexosaminidase from *A. oryzae* (10 U, ammonium sulphate precipitate)^{3b} was added and the mixture was incubated with stirring at 37 °C. After 20 min., another portion of *p*NP- β -GlcNAc (100 mg, 0.3 mmol) was added and the reaction was terminated after 4 hours by boiling (10 min.). The reaction mixture was diluted with H₂O and liberated *p*-nitrophenol was removed by extraction with diethyl ether (2 x 10 ml). The rest of the solvent was removed by short heating and the last traces of *p*-nitrophenol and unreacted *p*NP- β -GlcNAc were removed by passing the mixture through small column of polystyrene resin SM-2 (BioRad, USA) followed by washing with H₂O. The mixture was then fractionated by gel filtration through a BioGel P4 column (90 x 103 cm) being eluted with water. The disaccharidic fraction (56.5 mg, 24.6 %) was acetylated with pyridine/Ac₂O. Mixtures of the acetates were partially deacetylated with piperidine in THF (5 %, 5 ml) at 0 °C for 15 hours. Acetates **1b**, **2b** and **3b** were isolated by flash chromatography on silica gel (Merck 40 - 60 μ m) eluted with CHCl₃ with 2 % (v/v) MeOH. Zemplén deacetylation yielded pure **1a** (4.2 mg, 1.8 %) [α]_D²³ = + 3.5 (*c* = 0.59, H₂O), and minor amounts of **2a** and **3a**.
8. Semipreparatory HPLC was carried out on a Waters high performance liquid chromatograph equipped with solvent delivery system 6000A, universal injector U6K and with refractometric detector ALC 202; GL-PACK LiChrosorb NH₂-5 column (250 x 10 mm) (GL Sciences Inc., USA); the mobile phase was acetonitrile - water (78 : 22, v/v) at a flow rate 5.0 ml/min, temperature 50 °C; retention times were 12.86 and 14.63 min for **1a** and **2a**, respectively.
9. ¹H NMR (400 MHz, D₂O, 25 °C) data of **1a** (GlcNAc moiety denoted by primed numbers in all following cases) δ 2.046 (3 H, s, Ac), 3.369 (1 H, ddd, *J* = 10.0, 6.8, 2.1, H-5'), 3.451 (1 H, dd, *J* = 9.8, 9.8, H-3'), 3.473 (1 H, ddd, *J* = 9.7, 5.5, 2.1, H-5), 3.532 (1 H, dd, *J* = 9.7, 9.6, H-4), 3.577 (1 H, dd, *J* = 10.0, 9.8, H-4'), 3.648 (1 H, dd, *J* = 9.6, 3.3, H-3), 3.713 (1 H, dd, *J* = 12.1, 6.8, H-6'u), 3.740 (1 H, dd, *J* = 12.1, 5.5, H-6u), 3.757 (1 H, dd, *J* = 9.8, 8.6, H-2'), 3.916 (1 H, dd, *J* = 12.1, 2.0, H-6d), 3.933 (1 H, dd, *J* = 12.1, 2.1, H-6'd), 3.984 (1 H, dd, *J* = 3.3, 1.6, H-2), 4.907 (1 H, d, *J* = 8.6, H-1'), 4.986 (1 H, d, *J* = 1.6, H-1). ¹³C NMR (100 MHz, D₂O, 25 °C) data of **1a** δ 25.09 q (CH₃CO), 58.09 d (C-2'), 63.49 t (C-6'),

- 64.06 t (C-6), 69.72 d (C-4), 72.66 d (C-4'), 73.40 d (C-2), 75.57 d (C-3), 76.78 d (C-3'), 78.94 d (C-5), 79.34 d (C-5'), 99.57 d (C-1), 100.64 d (C-1'), 177.92 s (C=O).
10. ¹H NMR (400 MHz, CDCl₃, 30 °C) data of **1b** δ 1.965, 1.994, 2.024, 2.030, 2.062, 2.115, 2.133, 2.143 (all s, each 3 H, 8×Ac), 3.677 (1 H, m, *J* = 9.7, 5.9, 2.7, H-5), 3.683 (1 H, ddd, *J* = 9.6, 4.6, 2.7, H-5'), 4.029 (1 H, ddd, *J* = 10.4, 9.3, 8.4, H-2'), 4.141 (1 H, dd, *J* = 12.3, 2.7, H-6'u), 4.143 (1 H, dd, *J* = 12.2, 2.7, H-6u), 4.239 (1 H, dd, *J* = 12.3, 4.6, H-6'd), 4.369 (1 H, dd, *J* = 12.2, 5.9, H-6d), 4.796 (1 H, d, *J* = 8.4, H-1'), 4.965 (1 H, d, *J* = 1.3, H-1), 5.061 (1 H, dd, *J* = 9.9, 3.3, H-3), 5.070 (1 H, dd, *J* = 9.6, 9.2, H-4'), 5.154 (1 H, dd, *J* = 10.4, 9.2, H-3'), 5.207 (1 H, dd, *J* = 9.9, 9.7, H-4), 5.488 (1 H, dd, *J* = 3.3, 1.3, H-2), 5.535 (1H, d, *J* = 9.3, NH). ¹³C NMR (100 MHz, CDCl₃, 30 °C) data of **1b** δ 20.53 q (2 C), 20.58 q, 20.62 q, 20.67 q, 20.75 q, 20.84 q, 23.15 q, (8×CH₃CO), 53.59 d (C-2'), 61.89 t (C-6'), 62.45 t (C-6), 66.00 d (C-4), 68.11 d (C-2), 68.24 d (C-4'), 70.61 d (C-3), 72.25 d (C-3'), 72.37 d (C-5'), 72.96 d (C-5), 94.76 d (C-1), 97.41 d (C-1'), 169.19 s, 169.61 s, 169.88 s, 169.95 s, 170.12 s, 170.66 s, 170.70 s, 170.86 s (8×CH₃CO).
11. ¹H NMR (400 MHz, CD₃Cl, 30 °C) data of **2b** [α/β 2:1] δ 1.937, 2.008, 2.016, 2.055, 2.072, 2.103, 2.157 (all s, each 3 H, 7×Ac), 3.710 (1 H, ddd, *J* = 10.2, 4.7, 2.4, H-5'α), 3.713 (1 H, dd, *J* = 8.5, 8.0, H-2'β), 3.759 (1 H, dd, *J* = 10.3, 8.3, H-2'α), 4.086 (1 H, dd, *J* = 12.2, 4.7, H-6'uα), 4.170 (1 H, dd, *J* = 12.4, 1.2, H-6uα), 4.181 (1H, 1-OHα), 4.200 (1 H, ddd, *J* = 10.0, 6.5, 1.2, H-5α), 4.217 (1 H, dd, *J* = 12.4, 6.5, H-6dα), **4.262** (1 H, dd, *J* = 9.6, 3.4, H-3α), 4.336 (1 H, dd, *J* = 12.2, 4.7, H-6'dα), 4.739 (1 H, d, *J* = 8.3, H-1'α), 4.779 (1 H, d, *J* = 8.8, H-1'β), 5.037 (1 H, dd, *J* = 10.8, 9.6, H-4'β), 5.075 (1 H, dd, *J* = 10.3, 9.2, H-3'α), 5.161 (1 H, dd, *J* = 10.0, 9.6, H-4α), 5.236 (1 H, d, *J* = 1.8, H-1α), 5.244 (1 H, dd, *J* = 10.2, 9.2, H-4'α), 5.285 (1 H, dd, *J* = 3.4, 1.8, H-2α), 5.325 (1 H, dd, *J* = 10.8, 8.5, H-3'β), 5.616 (1H, d, *J* = 8.4, NH'α), 5.821 (1H, d, *J* = 8.5, NH'β). ¹³C NMR (100 MHz, CDCl₃, 30 °C) data of **2b** δ 20.4 q, 20.6 q, 20.7 q, 20.8 q, 20.9 q (7*×CH₃CO), 54.83 d (C-2'α), 61.95 t (C-6'α), 62.71 t (C-6α), 66.36 d (C-4α), 68.32 d (C-3'α), 68.98 d (C-2α), 69.12 d (C-5α), 71.96 d (C-5'α), 73.05 d (C-4'α), **74.04** d (C-3α), 90.96 d (C-1), 92.57 d (C-1α), 98.66 d (C-1'α). * Signal overlaps. ¹H NMR of **4b** - present as a minor impurity (under 10 %) in the sample of **2b** - δ 3.867 (1H, dd, *J* = 9.5, 9.5, H-4α), 5.449 (1H, dd, *J* = 9.5, 3.3, H-3α).
12. ¹H NMR (400 MHz, CDCl₃, 30 °C) data of **3b** [α/β 3:1] δ 1.996, 1.998, 2.024, 2.041, 2.080, 2.092, 2.169, (all s, each 3 H, 7×Ac), 3.509 (1 H, dd, *J* = 10.2, 8.2, H-2'β), **3.607** (1 H, dd, *J* = 12.3, 2.6, H-6uα), 3.702 (1 H, ddd, *J* = 10.0, 5.5, 1.6, H-5'α), 3.707 (1 H, ddd, *J* = 10.0, 4.4, 2.3, H-5'β), **3.712** (1 H, dd, *J* = 12.5, 4.4, H-6uβ), 3.763 (1 H, ddd, *J* = 10.5, 8.3, 8.2, H-2'α), **3.915** (1 H, dd, *J* = 12.0, 1.6, H-6dα), 3.954 (1H, d, *J* = 3.4, 1-OHα), **4.003** (1 H, dd, *J* = 12.7, 2.3, H-6dβ), 4.119 (1 H, dd, *J* = 12.5, 2.3, H-6'uβ), 4.131 (1 H, ddd, *J* = 10.3, 5.5, 1.6, H-5α), 4.173 (1 H, dd, *J* = 12.3, 2.6, H-6'uα), 4.214 (1 H, ddd, *J* = 9.9, 4.6, 2.3, H-5β), 4.243 (1 H, dd, *J* = 12.0, 1.6, H-6'dα), 4.293 (1 H, dd, *J* = 12.5, 4.6, H-6'dβ), 4.481 (1 H, d, *J* = 8.2, H-1'β), 4.942 (1 H, d, *J* = 8.3, H-1'α), 5.012 (1 H, dd, *J* = 10.3, 9.9, H-4α), 5.051 (1 H, dd, *J* = 10.2, 10.0, H-3'β), 5.071 (1 H, dd, *J* = 10.0, 9.2, H-4'α), 5.071 (1 H, dd, *J* = 10.0, 10.0, H-4'β), 5.190 (1 H, dd, *J* = 3.3, 1.7, H-1β), 5.229 (1 H, d, *J* = 1.8, H-1α), 5.240 (1 H, dd, *J* = 9.9, 9.9, H-4β), 5.257 (1 H, dd, *J* = 3.5, 1.8, H-2α), 5.310 (1 H, dd, *J* = 10.5, 9.2, H-3'α), 5.320 (1 H, dd, *J* = 3.5, 1.7, H-2β), 5.435 (1 H, dd, *J* = 9.9, 3.5, H-3β), 5.444 (1 H, dd, *J* = 9.9, 3.5, H-3α), 5.607 (1H, d, *J* = 10.9, NH'β), 5.616 (1H, d, *J* = 3.3, 1-OHβ), 5.955 (1H, d, *J* = 8.2, NH'α). ¹³C NMR (100 MHz, CDCl₃, 30 °C) data of **3b** δ 19.77 q, 20.35 q, 20.62 q, 21.10 q, 23.21 q. (7*×CH₃CO), 53.8 d (C-2'α), 62.1 t (C-6'α), **69.5** t (C-6α), 67.8 d (C-4α), 68.4 d (C-4'α), 69.7 d (C-2α), 68.9 d (C-3α), 70.7 d (C-5α), 72.2 d (C-5'α), 72.5 d (C-3'α), 92.5 d (C-1α), 100.8 d (C-1'α). * Signal overlaps.
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