



Transferase-catalyzed synthesis of non-natural oligosaccharide-libraries (SLe^a- and SLe^x-analogues)

Reinhold Öhrlein *, Gabi Baisch, Andreas Katopodis, Markus Streiff, Frank Kolbinger

NOVARTIS PHARMA, Postfach, CH-4002 Basle, Switzerland Received 26 September 1997; accepted 19 November 1997

Abstract

A number of non-natural glucosamine derivatives with various N-acyl and aglycon residues are prepared chemically. The N-acyl residues comprise aliphatic, polar, charged, aromatic and heteroaromatic replacements of the natural N-acetyl group. Both $\beta(1-4)$ - and cloned $\beta(1-3)$ -galactosyl-transferase tolerate a wide range of these replacements and yield the corresponding type II and type I disaccharides when incubated together with UDP-galactose. These disaccharides are subsequently sialylated with cloned $\alpha(2-3)$ -sialyl-transferase and CMP-sialic acid to give trisaccharides, which are sialylated at the 3-OH group of the terminal galactose. The sialylated type I compounds are finally incubated with cloned fucosyl-transferase III and type II compounds with cloned fucosyl-transferase VI, respectively. Thus, sialyl-Lewis^a and sialyl-Lewis^x libraries are generated. Additionally, both fucosyl-transferases accept donor substrates which have either the natural fucose-moiety or the guanosine-unit of the natural GDP-fucose donor replaced by non-natural congeners. © 1998 Elsevier Science B.V. All rights reserved.

Keywords: Glycosyl-transferases; Guanosine-diphosphofucose analogues; Oligosaccharide libraries; Sialyl-Lewis*; Sialyl-Lewis*; Enzymatic synthesis

Glycoconjugates play key roles in various adhesion and cell-cell recognition phenomena [1,2]. In order to probe carbohydrates as potential drug candidates [3], a proper understanding of carbohydrate-protein interaction is mandatory. This necessitates a rapid access to a wide range of natural and non-natural carbohydrate epitopes [4]. An alternative to the tedious chemical procedures for oligosaccharide synthesis is given by protocols based on the use of glycosyl-transferases for carbohydrate assem-

Recently, improved protocols make available these activated donor substrates and non-natural congeners thereof in large quantities, e.g., UDP-gal, ¹ CMP-sia [6] and GDP-fuc [7]. In addition, a number of glycosyl-transferases have been

blage [5]. Glycosyl-transferases are a class of enzymes, which transfer a monosaccharide unit highly regio- and stereospecifically onto a growing oligosaccharide chain in vivo. The monosaccharide sources are nucleotide monoor diphosphate sugars.

^{*} Corresponding author. E-mail: reinhold.oehrlein@pharma.novartis.com

¹ For example, UDP-galactose is offered and produced by Yamasa Jpn. in kilogram amounts.

produced by cloning and microbiol overproduction for preparative use [8]. We have recently shown that commercial $\beta(1-4)$ -galactosyltransferase [9], cloned $\alpha(2-3)$ -sialyl-transferase [10] and cloned fucosyl-transferase VI [11] can be used successfully to prepare SLe^x -derivatives, which have the natural N-acetyl-group of the glucosamine-moiety replaced by a broad array of aliphatic, aromatic and charged groups. The versatility of this approach has been extended further. We incubated non-natural acceptor substrates with non-natural 'fucose'-donors [12] (confer Scheme 1 and Table 1). Surpris-

ingly, also these 'fucoses' are transferred to the 4-OH group of the N-acyl glucosamine units3 in the expected α -mode.

Accordingly, a sialyl-Lewis^a-library has been produced. First, a series of type I disaccharides has been synthesized chemically or via enzymatic $\beta(1-3)$ -galactosylation [13]. These disaccharides are subsequently treated with cloned $\alpha(2-3)$ -sialyl-transferase and CMP-sia to α -sialylate, the terminal galactose at the 3-OH group. In a final incubation with cloned fucosyl-transferase III and various GDP-'fucoses' (see Scheme 2 and Table 2), the 'fucose'-

73

68

81

Table 1 Enzymatic preparation of SLe^x derivatives

fuc

L-gal ara

86

Table 2 Enzymatic preparation of SLe^a derivatives

moieties are linked to the 4-OH group of the N-acyl-glucosamine units in an α -mode.

References

- [1] M. Fukuda, Bioorg. Med. Chem. 3 (3) (1995) 207.
- [2] J. Hodgson, Biotechnol. 8 (1990) 108, 441.
- [3] M. Fukuda, O. Hindsgaul, Molecular Glycobiology, Oxford Univ. Press, Oxford, 1994.
- [4] M.J. Sofia, DDT 1 (1) (1996) 27.
- [5] H.J.M. Gijsen, L. Qiao, W. Fitz, C.-H. Wong, Chem. Rev. 96 (1996) 443.

- [6] M. Kittelmann, T. Klein, U. Kragl, C. Wandrey, O. Ghisalba, Appl. Microbiol. Biotechnol. 44 (1995) 59.
- [7] G. Baisch, R. Öhrlein, Bioorg. Med. Chem. 5 (2) (1997) 383.
- [8] M.L. Field, L.J. Wainwright, Glycobiology 5 (1995) 463.
- [9] G. Baisch, R. Öhrlein, B. Ernst, Bioorg. Med. Chem. Lett. 6 (7) (1996) 749.
- [10] G. Baisch, R. Öhrlein, M. Streiff, B. Ernst, Bioorg. Med. Chem. Lett. 6 (7) (1996) 755.
- [11] G. Baisch, R. Öhrlein, A. Katopodis, B. Ernst, Bioorg. Med. Chem. Lett. 6 (7) (1996) 759.
- [12] G. Baisch, R. Öhrlein, A. Katopodis, Enzymatic fucosylations of non-natural acceptors with non-natural donors, Bioorg. Med. Chem. Lett. 7 (18) (1997) 2431.
- [13] G. Baisch, R. Öhrlein, F. Kolbinger, in press.