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Molecular Design and Biological Potential of Galacto-Type Trehalose as a Nonnatural Ligand of Shiga Toxins

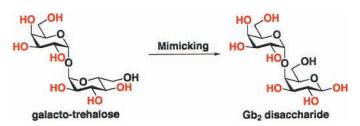
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ABSTRACT



Galacto-type trehalose, a "C-4 epimer of trehalose", possesses a stereochemical structure around the $\alpha(1-1)$ -linkage analogous to that of the globobiosyl $\alpha(1-4)$ -linkage in Gb₂ and Gb₃ ceramides, which are known as the ligands of Shiga toxins produced by pathogenic *E. coli*. This paper presents evidence supporting the new idea of using a trehalosyl $\alpha(1-1)$ -linkage as a substitute for the galactobiosyl $\alpha(1-4)$ -linkage.

Shiga toxins (Stx-I and Stx-II) produced by *Escherichia coli* O-157:H-7 and other pathogenic *E. coli* species are multisubunit proteins comprising one toxic A-subunit and five receptor-binding B-subunits.¹ Both Stx-I and Stx-II recognize a galactobiosyl α(1-4)-linkage in globosyl Gb₂ and Gb₃ ceramides and induce carbohydrate-mediated internalization into host cells.² There have been extensive studies on the molecular design of artificial Stx ligands that possess simpler structures and higher activity for blocking the toxin-host cell adhesion than the natural globosyl ceramides. Most of

the synthetic studies have been focused on the replacement of the ceramide group with simpler lipids³ or on the molecular assembly to construct carbohydrate clusters as glycopolymers, glycodendrimers, and starfish models.⁴ However, an effective mimic of the galactobiosyl $\alpha(1-4)$ -linkage has not yet been found. This is ironic since the synthesis of many mimics has been hampered by the difficulty of constructing the $\alpha(1-4)$ -linkage.⁵ In this communication, we

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propose the use of a trehalose $\alpha(1-1)$ -linkage as the substitute for the galactobiosyl $\alpha(1-4)$ -linkage and demonstrate the potential of this mimic.

Trehalose 1a possesses many biological and medicinal potentials applicable to sugar-based therapeutic reagents.⁶ This is mainly because the glycoside $\alpha(1-1)$ -linkage is highly tolerant to both chemical and enzymatic (mammalian α -glycosidases) degradation compared with glycosyl $\alpha(1-$ 4)-linkages. Here, it can be easily noticed that the trehalose $\alpha(1-1)$ -linkage has a molecular topology analogous to that of the galactobiosyl $\alpha(1-4)$ -linkage 2 (Gb₂) with binding activity to Shiga toxins. A critical difference arises from the configuration at OH-4, which should be axial to be recognized by Shiga toxins. Accordingly, a galacto-trehalose 1b can be designed as a novel Gb₂ mimic and expected to show a binding affinity to Stxs similar to that of the natural Gb₂ ligand as well as the advantageous property mentioned above. Molecular force field (MM-2) calculation supported the similarity of the stereochemical environment around the glycoside linkage between **2** and **1b** (Figure 1).

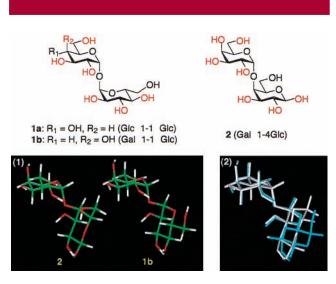


Figure 1. Preferred conformations of galacto $\alpha(1-4)$ -bioside 2 and galacto-type trehalose 1b (1) and superimposed structures (2). The structures were minimized by MM-2 without optimization of all OH bonds in the calculation (Insight II/Discover program and Amber force field); the white and blue lines represent 2 and 1b, respectively, in panel 2.

In this study, we prepared galacto-type trehalose **6**⁷ and its cluster model **7** in order to verify our expectation.⁸ The synthesis was carried out via conventional chemical pathways

starting from 4,6-mono-O-benzylidene trehalose $\mathbf{3}^9$ as summarized in Scheme 1. As a key reference compound to

Scheme 1. Synthesis of Galacto-Trehalose Cluster Model 7^a

a Reaction conditions: (a) (i) TBDPSCI, Et₃N, DMAP, pyridine, room temperature, 12 h, 91%; (ii) BnBr, NaH, DMF, room temperature, 12 h, 95%; (iii) BH₃¬NHMe₂, AlCl₃, THF, 0 °C→room temperature, 2 h, 94%; (b) (i) Tf₂O, pyridine, CH₂Cl₂, 0 °C→room temperature, 45 min, quant; (ii) CsOAc, 18-crown-6, toluene, ultrasound, 40 °C, 2 h, 97%; (c) (i) NaOMe, MeOH, THF, room temperature, 12 h, quant; (ii) TBAF, THF, room temperature, 24 h, 95%; (iii) 6-azido-1-bromohexane, TBAI, NaH, DMF, 60 °C, 12 h, 92%; (iv) Pd(OH)₂/C, H₂, THF, MeOH, room temperature, 1.5 h; (v) acryloyl chloride, Et₃N, CH₂Cl₂, MeOH, 0 °C→room temperature, 2 h, 91% (2 steps); (d) 2,2-azobis (2-amidinopropane) dihydrochloride, H₂O, Me₂SO, 60 °C, 12 h, 72%.
b Sloping line in the structure represents the random copolymer comprised of the two constitutional units.

investigate the role of the OH-4 configuration, a cluster model 10 carrying trehalose at the side chain was also prepared in the same way (6-azidohexylation at O-6, reduction of the azido group, N-acryloylation, and copolymerization with acrylamide in a redox radical manner) (Figure 2). Acrylamido copolymers $\bf 8$ and $\bf 9$ carrying natural Gb₂ and α -D-galactoside clusters, respectively, were prepared in our previous manner. 5,10

Hemagglutination inhibition assay¹¹ was performed for trehalose **1a**, galacto-trehalose **6**, *p-N*-acrylamidophenyl Gb₂,⁵

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⁽⁶⁾ Highly effective use of an α-D-mannosyl(1–1)- β -D-galactosyl linkage as a scaffold constructing a sialyl Lewis X mimic was reported: Hiruma, K.; Kajimoto, T.; Schumidt, G. W.; Ollmann, I.; Wong, C.-H. *J. Am. Chem. Soc.* **1996**, *39*, 9265–9270.

⁽⁷⁾ Compound **6**: ¹H NMR (δ ppm, 500 MHz, D₂O): 5.65–6.24 (dd × 3, 3H, H-olefin), 5.15 (d, 1H, $J_{1,2} = 3.4$ Hz, GalH-1), 5.14 (d, $J_{1,2} = 3.1$ Hz, GlcH-1), 4.15 (d, $J_{3,4} = 3.0$ Hz, GalH-4), 3.36 (dd, 1H, $J_{2,3} = 6.7$ Hz, $J_{3,4} = 9.0$ Hz, GlcH-4), 1.40–3.50 (dd × 2 and m × 4, 12H, H-methylene).

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Gb₂ copolymer **8** $M_{\rm n} = 4.4 \times 10^5$ Mol fraction of the sugar unit = 0.21

% of Gal copolymer 9 $M_{\rm n}{=}~4.4\times10^5$ Mol fraction of the sugar unit =0.21

trehalose copolymer **10** $M_{\rm n} = 3.5 \times 10^5$ Mol fraction of the sugar unit = 0.19

Figure 2. Structures and properties of glycoconjugate copolymers 8–10

and *p*-nitrophenyl α -D-galactoside¹⁰ to show that none of these monomeric glycosides possess notable activity to block the Stx binding to erythrocytes (MIC > 10 mM). The result accords well with the report¹² that a monomeric methyl globotriaoside (Gb₃) has a very low binding affinity to Stx-I ($K_a < 1.0 \times 10^3 \text{ M}^{-1}$). On the other hand, activity was observed clearly for copolymers due to a cluster effect (Table 1).¹³ The galacto-trehalose copolymer **7** showed blocking activity for both Stx-I (MIC = ca. 60 μ M) and Stx-II (ca. 240 μ M). Though the activity was slightly lower than that of the natural Gb₂ copolymer **8** (MIC = 40 μ M for Stx-I

Table 1. Hemagglutination Inhibition Assay and Stx-I Neutralization Activity Assay

	MIC (M) ^a		neutralization activity
copolymer	Stx-I	Stx-II	against Stx-I ^b
7	6.1×10^{-5}	$2.4 imes 10^{-4}$	0.88
8	$4.1 imes 10^{-5}$	$1.6 imes 10^{-4}$	0.95
9	$1.1 imes 10^{-3}$	$> 1.1 \times 10^{-3}$	0.65
10	$>\!5.7\times10^{-2}$	$> 5.7 \times 10^{-2}$	c

 $[^]a$ Minimum inhibition concentration. 13 b Relative activity to that of Gb₃ copolymer. c Not determined.

and 160 μ M for Stx-II), the activity was apparently higher than that of the α -galactoside copolymer 9 (MIC = 1 mM for Stx-I and >1 mM for Stx-II). Moreover, the trehalose copolymer 10 did not show binding activity to Stx-I or Stx-II. These data are suggestive of the structural characteristics of the galacto-type trehalose 1b. The axial 4-OH in the α -galactoside moiety is essential for the Stx binding, and the α -glucoside moiety in 1b, similarly to the β -galactoside moiety in 2, plays an important role as a supplementary binding site.

Stx-I neutralization assay using HeLa cells was performed for the binding-active copolymers 7–9. Under conditions in which the Gb₃/acrylamide copolymer¹⁰ (10 μ M) can keep all cells alive in the presence of Stx-I, galacto-trehalose copolymer 7 could rescue 88% of the cells. The relative activity matches well with the preceding hemagglutination inhibition activity and also supports our expectation that the galacto-type trehalose 1b serves as a nonnatural carbohydrate ligand of Shiga toxins.

In conclusion, we have found a potential utility of the galacto-trehalose $\alpha(1-1)$ -linkage as a promising substitute for the globosyl $\alpha(1-4)$ -linkage. Though the binding of the galacto-trehalose to Stx-I and Stx-II, particularly to the latter, is still weaker than that of the natural Gb_2 and Gb_3 globosides, galacto-trehalose may integrate the binding affinity significantly by additionally introducing a β -glucoside moiety or other related units. Such efforts are in progress in our group and will be reported elsewhere. Moreover, the idea of applying α - (or even β -) (1–1)-linked sugars, instead of natural and labile glycosyl linkages, will open a promising pathway leading to sugar-based therapeutic agents.

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Supporting Information Available: Experimental procedures and full characterization for all new compounds (PDF). This material is available free of charge via the Internet at http://pubs.acs.org.

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