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Preferential binding of *E. coli* with type 1 fimbria to D-mannobiose with the Man α 1 \rightarrow 2Man structure

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Manα1→2Man, Manα1→3Man, Manα1→4Man, and Man α 1 \rightarrow 6Man were converted to the glycosylamine derivatives. Then, they were mixed with monobenzyl succinic acid to obtain their amide derivatives. After removing the benzyl group by hydrogenation, the succinylamide derivatives were coupled with the hydrazino groups on BlotGlycoTM beads in the presence of water-soluble carbodiimide. **D-Mannobiose-linked beads were incubated with** fluorescence-labeled Escherichia coli with type 1 fimbria, and the number of the fluorescent dots associated with the beads was counted in order to determine the binding preference among p-mannobiose isomers. The results showed that the bacteria bind strongly to Mana1 \rightarrow 2Man1 \rightarrow beads, Man- $\alpha 1 \rightarrow 3Man1 \rightarrow beads$, $Man\alpha 1 \rightarrow 4Man1 \rightarrow beads$, and Man α 1 \rightarrow 6Man1 \rightarrow beads, in order. In the presence of 0.1 M methyl a-d-mannopyranoside, most of the bacteria failed to bind to these beads. These results indicate that E. coli with type 1 fimbria binds to all types of *D*-mannobiose isomers but preferentially to Man α 1 \rightarrow 2Man disaccharide.

Key words: *E. coli*; carbohydrates; D-mannobiose; binding preference

Many microorganisms recognize cell surface glycans as receptors for cell attachment and tissue colonization. This is because viruses, bacteria, and protozoa express a large number of glycan-binding proteins so-called lectins (reviewed in Ref.¹). For example, human influenza viruses recognize the Sia $\alpha 2 \rightarrow 6$ Gal structure by their hemagglutinins expressed on virus surface, and *Helicobacter pylori* binds Fuc $\alpha 1 \rightarrow 2$ Gal $\beta 1 \rightarrow 3$ (Fuc $\alpha 1 \rightarrow 4$) GlcNAc1 $\beta \rightarrow R$ structure (Le^b antigen) by its adhesin BabA expressed on flagella.^{2,3)} In fact, such glycans are expressed on cell surfaces of human respiratory tract and stomach tissues, respectively. Furthermore, *Escherichia. coli* binds high mannosetype glycans rich in intestinal mucosa and glycans with the Gal α 1 \rightarrow 4Gal group in urinary tract depending on the strains, and the pathogenic strains often cause enterohemorrhagy with verotoxin produced, hemolytic uremic syndrome, acute encephalopathy, pyelonephritis, cystitis, and urethritis for humans.^{4–8)} *E. coli* is a gramnegative bacterium and covered with fimbria. At the top of the fimbria, there are glycan-binding proteins called adhesins. The most well characterized bacterial adhesin is the type 1 fimbrial FimH adhesin which recognizes α -D-mannose.⁹⁾ FimH is displayed on the bacterial surface as a component of the type 1 fimbrial organelle.

Using beads immobilized with a-D-mannose-containing glycans, we found E. coli binds strongly to high mannose-type glycans and wondered how many α -Dmannose residues are required for the recognition by the FimH adhesin. Then, we prepared beads immobilized with D-mannose or D-mannobiose with Man α 1 \rightarrow 2-Man structure, examined the binding activity of E. coli (strain W3110) to these beads, and found that the bacteria bind stronger to D-mannobiose-linked beads than α -D-mannose-linked beads. However, in the case of D-mannobiose, there are four isomers with different glycosidic linkages. In this study, we examined the binding preference of E. coli among D-mannobiose isomers with Man α 1 \rightarrow 2Man. Man α 1 \rightarrow 3Man. Man α 1 \rightarrow 4Man, and Man α 1 \rightarrow 6Man structures.

Materials and methods

Reagents. All the chemical reagents used for the preparation of sugar-linked beads were from Wako Pure Chemicals Co. (Osaka). D-Mannobiose isomers, Man α 1 \rightarrow 2Man, Man α 1 \rightarrow 3Man, and Man α 1 \rightarrow 6Man were synthesized by HCl-catalyzed condensation reaction according to the method described previously.¹⁰ Man α 1 \rightarrow 4Man was obtained from Carbosynth Ltd.

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Abbreviations: E. coli, Escherichia coli; HPLC, high-performance liquid chromatography; HRMS, high-resolution mass spectrometry; PBS, 10 mM phosphate-buffered saline (pH 7.4); WSC, water-soluble carbodiimide.

(Berkshire, UK). A PKH67 green fluorescent cell linker kit for general cell membrane labeling was purchased from Sigma-Aldrich Corp. (St. Louis, MO).

Synthesis of *D*-mannobiosyl succinylamide deriva-Since the method for preparing D-mannobiosetives. linked beads is the same among the isomers, the preparation of Man α 1 \rightarrow 2-Man1 \rightarrow beads was described here as a representative. The amination of the D-mannobiose was performed in a saturated (NH₄)₂CO₃ solution by the method with a slight modification¹¹) as described in Fig. 1. A disaccharide, Man α 1 \rightarrow 2Man, (1, 51.5 mg, 0.15 mmol) was dissolved in a saturated $(NH_4)_2CO_3$ solution (5 mL), and the mixture was stirred at 40 °C for 4 h. After confirming the disappearance of a starting material spot (Rf: 0.43) by thin layer chromatography (CH₃CN/H₂O = 3:1 (v/v) containing 0.1% triethylamine), the solution was subjected to evaporation initially at 45 mm Hg and then at 12 mm Hg.

Monobenzyl succinic acid was synthesized by mixing benzyl alcohol (1.0 mL, 9.6 mmol) and succinic anhydride (1.2 g, 1.2 mmol) in pyridine (2.0 mL) at room temperature for 1 h. The concentrated reaction mixture was dissolved in ethyl acetate, washed with a 1 M HCl solution, brine (a saturated aqueous NaCl solution), a saturated aqueous NaHCO₃ solution and brine, dried in the presence of Na₂SO₄, and then concentrated in vacuo to obtain monobenzyl succinic acid as a pure compound. All the concentrated D-mannobiosyl amine syrup (1) was dissolved in dimethyl formamide (5 mL), to which monobenzyl succinic acid (46.8 mg, 0.225 mmol) and 4-(4,6-dimethoxy-1,3,5-triazin-2-yl)-4-methylmorpholinium chloride n-hydrate¹²⁾ (61.2 mg, 0.225 mmol) were added. The mixture was stirred at 40 °C for 22 h and then concentrated in vacuo to obtain syrup (34.5 mg). The syrup was purified by chromatography using a C-18 Sep-Pak^T column (Nihon Waters KK, Japan) and a Mightysil RP-18 GP^{TM} column (1 × 25 cm) (Kanto Chemical Co., Inc., Tokyo). Fractions eluted at the retention time from 12 to 14 min were collected (the retention times were different among D-mannobiose isomers), and the collected fractions were concentrated to obtain a pure D-mannobiosyl benzylsuccinylamide derivative (**3**, 9.8 mg, 18 µmol).

The compound **3** (7.1 mg, 13 µmol) was dissolved in distilled water (2 mL), and PD-C (10 mg) was added to the solution. It was then placed into a flask attached with a H₂ balloon and stirred at room temperature for 3 h. After completing the reaction, the catalyst was separated by centrifugation at $2500 \times g$ for 3 min. By concentrating the supernatant, D-mannobiosyl succinylamide derivative (4) was obtained quantitatively. The ¹H-NHR and ¹³C-NMR spectra and high-resolution mass spectrometry (HRMS) spectra of D-mannobiosyl succinylamide derivatives were obtained with a Jeol JNM-EC600 spectrometer (JEOL Ltd., Tokyo) and a micrOTOF II spectrometer (Bruker Daltonics Inc., Billerica, MA), respectively.

Analytical data of $Man \alpha 1 \rightarrow 2Man$ succinylamide derivative. Rf: 0.28 (CH₃CN/H₂O = 3:1). ¹H-NMR (600 MHz, D₂O): $\delta_{\rm H}$ 5.57 (0.18H, d, J=1.91 Hz, H-1α), 5.15 (0.82H, bs, H-1β), 5.13 (0.16H, s, H-1α'), 4.97 (0.84H, d, J = 1.81 Hz, H-1 β), 4.00 (1H, dd, J =1.92, 3.05 Hz, H-2), 3.93 (1H, bd, J=3.04 Hz, H-2'), 3.81 (1H, dd, J = 1.21, 11.98 Hz, H-6), 3.78 (1H, dd, J= 3.08, 9.51 Hz, H-3', 3.89 (1H, bd, J = 2.81,11.83 Hz, H-6), 3.70 (1H, dd, J=3.08, 9.66 Hz, H-3), 3.62 (1H, dd, J=6.81, 12.21, H-6'), 3.59 (1H, dd, J= 5.85, 12.01 Hz, H-6') 3.53 (1H, t, J=9.461 Hz, H-4), 3.51 (1H, t, J=9.59 Hz, H-4), 3.34 (1H, ddd, J=2.02, 5.97, 9.67 Hz, H-5), 2.57-2.49 (4H, m, -CH₂-CH₂-), ¹³C-NMR (150 MHz, D₂O): δ_C 177.53, 175.49 (C=O), 103.10 (C-1'), 81.25, 78.91 (C-1a), 78.58 (C-1b), 78.33 (C-3), 77.75, 74.44, 73.89 (C-5), 70.62 (C-2), 67,39, 67.02, 61.45 (C-6), 61.40, 30.43, 29.51 (-CH₂-CH₂-).



Fig. 1. Synthetic pathways of BlotGlycoTM beads linked with D-mannobiose derivative (one of R_1-R_4 is D-mannose, and the others are H's).

HRMS m/z calcd. for $C_{16}H_{26}NO_{13}$ [M-H⁺] 440.1399, found 440.1390.

Analytical data of $Man\alpha 1 \rightarrow 3Man$ succinylamide Rf: 0.27 (CH₃CN/H₂O = 3:1). ¹H-NMR derivative. (600 MHz, D_2O): δ_H 5.28 (0.09H, d, J = 1.95 Hz, H-1 α), 5.13 (0.91H, s, H-1 β), 5.10 (0.09H, s, H-1 α '), 4.97 $(0.91H, d, J=1.38 \text{ Hz}, H-1\beta'), 3.95 (1H bd, J=$ 3.15 Hz, H-2'), 3.93 (1H, dd, J=1.48, 3.06 Hz, H-2), 3.77-3.72 (3H, m, H-6, H-6, H-3), 3.70 (1H, dd, J= 3.32, 9.47 Hz, H-3), 3.67-3.56 (4H, m, H-6', H-6', H-5', H-4), 3.51 (1H, t, J=9.66 Hz, H4), 3.36 (1H, ddd, J=2.00, 5.53, 9.56 Hz, H-5), 2.48 (4H, bs, -CH2-CH2–), ¹³C-NMR (150 MHz, D₂O): δ_C 178.78, 176.14 (C=O), 102.56 (C-1'), 80.86, 78.19, 78.23, 78.23, 74.44, 73.92, 70.91, 70.57, 70.36, 67,36, 66.52, 61.55 (C-6), 61.30 (C-6'), 31.12, 30.43 (-CH₂-CH₂-). HRMS m/z calcd. for C₁₆H₂₆NO₁₃ [M-H⁺] 440.1399, found 440.1393.

Analytical data of Manα1→4Man succinylamide derivative. Rf: 0.25 (CH₃CN/H₂O = 3:1). ¹H-NMR (600 MHz, D₂O): $\delta_{\rm H}$ 5.31 (0.12H, d, J=1.70 Hz, H-1α), 5.14 (0.88H, d, J=1.46 Hz, H-1β), 5.10 (1H, d, J=0.85 Hz, H-1β'), 3.94 (1H, dd, J=1.70, 2.97 Hz, H-2), 3.79 (1H, dd, J=0.81, 3.09 Hz, H-2'), 3.78–3.74 (2H, m, H-6, H-6), 3.71–3.60 (4H, m), 3.59–3.52 (3H, m), 3.43 (1H, ddd, J=1.95, 5.10, 9.40 Hz, H-5), 2.59– 2.50 (4H, m, CH₂–CH₂–), ¹³C-NMR (150 MHz, D₂O) $\delta_{\rm C}$ 177.17, 175.42 (C=O), 101.57 (C-1'), 80.86, 78.37, 76.83, 74.41, 74.27, 71.23, 70.87, 67,12, 61.54 (C-6), 61.48 (C-6'), 30.04, 28.91 (–CH₂–CH₂–). HRMS *m*/*z* calcd. for C₁₆H₂₆NO₁₃ [M-H⁺] 440.1399, found 440.1400.

Analytical data of $Man\alpha 1 \rightarrow 6Man$ succinvlamide Rf: 0.25 (CH₃CN/H₂O = 3:1). ¹H-NMR derivative. (600 MHz, D_2O): δ_H 5.31 (0.11H, d, J=1.70 Hz, H-1 α), 5.11 (0.89H, d, J=0.97 Hz, H-1 β), 4.77 (0.88H, d, J=1.59 Hz, H-1β'), 4.75 (012H, d, J=1.64 Hz, H- $1\alpha'$), 3.86 (1H, dd, J=1.62, 3.98 Hz, H-2), 3.83 (1H, dd, J=1.02, 2.98 Hz, H-2'), 3.81 (1H, dd, J=5.13, 11.46 Hz, H-6), 3.75 (1H, dd, J=2.04, J=12.17 Hz, H-6), 3.71 (1H, dd, J=3.18, 8.93 Hz, H-3'), 3.63 (1H, dd, J=5.56, 12.04, H-6'), 3.59 (1H, dd, J=5.47, 11.64, H-6'), 3.58 (1H, dd, J=3.92, 9.58, H-3), 3.58-3.51 (3H, m, H-4, H-4', H-5'), 3.47 (1H, ddd, J=1.95, 5.10, 9.40 Hz, H-5), 2.59–2.52 (4H, m, -CH₂– CH₂--),¹³C-NMR (150 MHz, D₂O) δ_{C} 177.66, 175.79 (C=O), 100.10 (C-1'), 78.27, 76.34, 74.01, 73.27, 71.04, 74.01, 70.04, 67,23, 66.89, 66.28 (C-6), 61.44 (C-6'), 30.55, 29.40 (-CH2-CH2-). HRMS m/z calcd. for C₁₆H₂₆NO₁₃ [M-H⁺] 440.1399, found 440.1394.

Immobilization of *D*-mannobiose isomers on beads.

To D-mannobiosyl succinylamide derivative (**4**, 2 mg, 4.5 μ mol) dissolved in distilled water (300 μ L), 8 mg of BlotGlycoTM beads (Sumitomo Bakelite Co. Ltd., Tokyo)¹³⁾ and water-soluble carbodiimide (WSC) (86.3 mg, 450 μ mol) was added (Fig. 1). The immobi-

lization was conducted by rotation of the mixture at 40 °C for 48 h, and then, the suspension was centrifuged at $2500 \times g$ for 30 s. The beads were washed with water (300 µL × 3 times) and methanol (500 µL × 3 times) and finally dried *in vacuo* to obtain p-mannobiose-linked BlotGlycoTM beads (5) with keeping its cyclic conformation of p-mannose residue at the reducing end (Fig. 1).

Determination of amounts of *D*-mannose linked to beads. The amount of D-mannobiose linked to BlotGlycoTM beads was determined as follows. The disaccharide-linked beads (5, 0.9-2.3 mg) were placed into 4 M CF₃COOH (300 µL) in an Eppendorf tube. After incubating the tube at 100 °C for 3 h, N₂ gas was flashed into the tube for evaporation of CF₃COOH and water. D-Mannose residues released were dissolved in distilled water (200 μ L), and the sample was filtered through a Millex filter (0.45 µm) (Merk Millipore Corp., Billerica, MA) at $700 \times g$ for 30 s. The amount of D-mannose was determined by high-performance liquid chromatography (HPLC) equipped with an evaporative light scattering detector using an Asahi-Pak NH₂-P50[™] column (Showa Denko K.K., Tokyo).

Culture and labeling with a fluorescent dye of E. coli (strain W3110)¹⁴⁾ donated by Dr. hacteria. M. Fukuda of Nagaoka University of Technology was grown in a Luria-Bertani medium at 37 °C for 14-20 h by shaking. After centrifugation, the bacteria were suspended in Diluent C (1 mL) from the cell membrane labeling kit, mixed with 10 µL of PKH67 solution (4 µL of PKH67 mixed with 996 µL of Diluent C), and incubated at 37 °C for 10 min to label the bacteria with the fluorescent dye according to the manufacturer's protocol. In this study, more than 90% of E. coli was labeled with the dye and alive during the period of experiments, which was confirmed by determining the growth rate of the fluorescence-labeled E coli in culture. Higher percentages of cells labeled with this dye together with a longer survival time of the dye-labeled cells were reported.¹⁵

Assay for binding of bacteria to *D*-mannobiose-linked beads. Fluorescence-labeled E. coli suspended in a Luria-Bertani medium (OD600 = 0.1, 200 μ L) was mixed with 1 mg of D-mannobiose-linked beads suspended in 40 µL of 10 mM phosphate-buffered saline (pH 7.4) (PBS) in a glass-bottom dish and incubated at 37 °C for 1–2 h. Then, beads were washed with PBS (200 μ L) three times and suspended in PBS (200 μ L) for observation under a confocal laser microscope. In some experiments, the bacteria and the beads were incubated in the presence of 0.1 M methyl α-D-mannopyranoside.

Results and discussion

Preparation of *D*-mannobiose isomer-linked beads In order to determine the binding preference of *E. coli* with type 1 fimbria among *D*-mannobiose

Man α 1 \rightarrow 2Man, Man α 1 \rightarrow 3Man, isomers, and Man α 1 \rightarrow 6-Man were synthesized by HCl-catalyzed condensation reaction,¹⁰⁾ and Man α 1 \rightarrow 4-Man were commercially obtained. Individual D-mannobiose isomers were mixed with monobenzyl succinic acid in the presence of 4-(4,6-dimethoxy-1,3,5-triazin-2-yl)-4methylmorpholinium chloride n-hydrate, and then, the benzyl group was removed by hydrogenolysis to obtain D-mannobiosyl succinylamide derivatives. The ¹H-NMR spectra of D-mannobiosyl succinylamide derivatives (Fig. 2(a-d)) showed two sets of signals derived from α -anomer (1 α) and β -anomer (1 β) at the reducing end and at the non-reducing end, the latter of which is indicated with an asterisk shown in Fig. 2. The assignment of H-1 signals was performed by nuclear Overhauser effect techniques in NMR spectroscopy. The H-1 signal giving a cross-peak with H-5 proton signal was assigned to be axial, which indicates that the succinylamide group was conjugated in a β -linkage to the D-mannose residue. Contrary to the empirical rule in the glycosylation of D-mannose, the amount of the β -linked isomer was more than that of the α -linked isomer, and furthermore, the coupling constant between H-1 α and H-2 was larger than that between H-1 β and H-2 in all the derivatives. For all the succinylamide derivatives of D-mannobiose isomers, the percentages of the β -isomers were 82–93% by calculating the peak areas of the proton signals at individual reducing ends. The α -/ β -isomer ratios obtained by ¹H-NMR and HPLC analyses are shown in Table 1.

D-Mannobiosyl succinylamide derivatives were mixed with BlotGlycoTM beads. The immobilization of them on the beads was conducted according to the reaction described in Fig. 1. Although the protocol of



Fig. 2. 600 MHz ¹H-NMR spectra of D-mannobiosyl succinyl-amide derivatives. Panels (a), (b), (c), and (d) indicate Man α 1 \rightarrow 2Man1-, Man α 1 \rightarrow 3Man1-, Man α 1 \rightarrow 4-Man-1-, and Man α 1 \rightarrow 6Man1-succinyl-amide derivatives, respectively. The 1 α and 1 β indicate H-1 α and H-1 β proton signals at the reducing end, respectively. *, a signal from the H-1 proton at the non-reducing end.

Table 1. The α/β percentages of D-mannobiosyl succinylamide derivative 4's.

Sugar part of compound 4	NMR ^a		HPLC ^b	
	α	β	α	β
Manα1→2Man	18	82	11	89
Manα1→3Man	9	91	10	90
Manα1→4Man	12	88	13	87
Manα1→6Man	11	89	10	90

^aDetermined from integral intensity of each H-1 signal in NMR spectra.

^bDetermined from height of each peak in HPLC chart.

BlotGlyco[™] beads shows the coupling between a hydrazino group on the bead and an aldehyde group of the reducing end of glycan,¹³⁾ the present method was to couple the carboxyl group of D-mannobiosyl succinylamide derivatives to a hydrazino group on the bead with an aid of WSC to establish a hydrazide linkage. As the reaction was to hydrazino groups on the bead, a 100-fold higher amount of WSC than that of bead was used. The amount of D-mannobiose isomers linked to BlotGlyco[™] beads was determined, which is described in detail in the materials and methods section, and 0.11-0.16 µmol of D-mannose was linked to 1 mg of the beads (Table 2). As the amount of hydrazino group expressed on the bead is shown to be 1-2 µmol/mg bead according to the manufacturer's protocol,¹³⁾ the obtained values in Table 2 appeared reasonable.

Binding of E. coli to *D*-mannobiose isomer-linked beads

E. coli with type 1 fimbria is shown to bind to α -Dmannose-containing glycans.4,5,9) Our preliminary study showed that the bacteria bind to D-mannobiose-linked beads stronger than D-mannose-linked beads. In order to examine whether E. coli shows binding preference to a D-mannobiose isomer with a certain structure, the fluorescence-labeled bacteria were incubated with Man α 1 \rightarrow 2Man-1 \rightarrow beads, Man α 1 \rightarrow 3-Man1 \rightarrow beads, Man α 1 \rightarrow 4Man1 \rightarrow beads, or Man α 1 \rightarrow 6Man-1 \rightarrow beads, and the reaction mixtures were observed under a confocal laser microscope for counting the numbers of the dots associated with the circumferences of the beads. The bacteria bound to all types of D-mannobiose isomer-linked beads (Fig. 3). In the case of Mana- $1 \rightarrow 2$ -Man $1 \rightarrow$ beads, the average number of the dots associated with the beads was 52 dots/bead, and more dots whose numbers were unable to be counted were associated with the two beads shown in Fig. 3(a).

When the numbers of the fluorescent dots associated with other types of the beads were counted, the average numbers of them were 26 dots/bead in the case of Man α 1 \rightarrow 3-Man-1 \rightarrow bead (Fig. 3(b)), 16 dots/bead in the case of Man α 1 \rightarrow 4-Man1 \rightarrow bead (Fig. 3(c)), and 10 dots/bead in the case of $Man\alpha 1 \rightarrow 6Man 1 \rightarrow bead$ (Fig. 3(d)), respectively. The binding assay was conducted three times, and similar binding preference to those shown in Fig. 3 was obtained. When the bacteria were incubated with Man α 1 \rightarrow 2Man-1 \rightarrow beads in the presence of 0.1 M methyl a-D-mannopyranoside, most of them failed to bind to the beads (Fig. 3(e)). The same results as that shown in Fig. 3(e) were obtained for E. coli using Man α 1 \rightarrow 3Man1 \rightarrow beads, Man α 1 \rightarrow 4-Man-1 \rightarrow beads, and Man α 1 \rightarrow 6Man1 \rightarrow beads, respectively, in the presence of the haptenic sugar (data not shown). These results indicate that E. coli with type 1 fimbria binds to D-mannobiose-linked beads through its type 1 adhesion and that the bacteria have a preference to bind to Man α 1 \rightarrow 2Man disaccharide among the isomers. There are many fluorescent dots not bound to the beads in Fig. 3. They can be removed from the reaction mixtures by harsh washings with PBS. However, such washings also take the beads out and reduce the number of the beads to be observed. Another approach to reduce the background fluorescent dots is to decrease the number of E. coli mixed with the beads. However, in this case, it also decreases the number of the fluorescent dots associated with the beads. Therefore, the present assay condition is suitable for determining the binding preference of E. coli among D-mannobiose isomers. Furthermore, once the bacteria bind to these beads, they are not released from the beads under the present condition unless the haptenic sugar is included in the reaction mixture. Therefore, the fluorescent dots in the background are not from those released from the beads

High mannose-type glycans attached to proteins contain Man α 1 \rightarrow 2Man disaccharides at the non-reducing termini,¹⁶) and *E. coli* may bind to these high mannose-type glycans expressed in the large intestine where they reside. *E. coli* is quite often contaminated in water for domestic use to cause us serious enterohemorrhagic diseases.⁸) If some filtration systems immobilized with the D-mannobiose or high mannosetype glycans are developed and applied to a water supply system to clean, they can trap specifically *E. coli* with type 1 fimbria in water and provide us with safe water. In some underdeveloped countries, local residents are still facing to serious infectious diseases with *E. coli* contaminated in water for domestic use. It has been reported that around 100 bacteria are contaminated in 1 mL of water.¹⁷⁾ Our preliminary study

Table 2. Amounts of D-mannose liberated from sugar-linked BlotGlyco[™] beads.

	Beads ^a (mg)	D-mannose ^b (μg/200 μL)	D-mannose on the beads	
			(µg/mg beads)	(µmol/mg beads)
Manα1→2Man	0.577	32.2	56	0.16
Manα1→3Man	0.576	20.2	37	0.11
Manα1→4Man	0.569	44.0	42	0.12
Manα1→6Man	0.570	28.6	50	0.15

^aWeights of sugar-linked BlotGlyco[™] beads used for the analysis.

^bAmounts of D-mannose from BlotGlyco[™] beads determined by HPLC analysis.



Fig. 3. Incubation of fluorescence-labeled *E. coli* with sugar-linked beads. Panels (a), (b), (c), and (d) indicate the bacteria with Man α 1 \rightarrow 2-Man1 \rightarrow beads, Man α 1 \rightarrow 3Man-1 \rightarrow beads, Man α 1 \rightarrow 4Man-1 \rightarrow beads, and Man α 1 \rightarrow 6Man1 \rightarrow beads, respectively, and panel (e) indicates the bacteria with Man α 1 \rightarrow 2Man1 \rightarrow beads in the presence of 0.1 M methyl α -D-mannopyranoside.

showed that 1 mg of beads immobilized with high mannose-type glycans can trap at least a million of *E. coli*. If such a tool immobilized with Man α 1 \rightarrow 2Man disaccharide or high mannose-type glycans is used in these countries, it can trap the bacteria in water for domestic use and provide the local residents with safe water.

Author contributions

KA and KF designed the research and wrote the paper. KY, KS, NI, IM, RK, and TM performed the experiments.

Disclosure statement

No potential conflict of interest was reported by the authors.

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