

Communication

## Practical Preparation of Lacto-*N*-biose I, a Candidate for the Bifidus Factor in Human Milk

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**A one-pot enzymatic reaction to produce lacto-*N*-biose I (LNB), which is supposed to represent the bifidus factor in human milk oligosaccharides, was demonstrated. Approximately 500 mM of LNB was generated in 10-liter of reaction mixture initially containing 660 mM of sucrose and 600 mM of GlcNAc by the concurrent actions of four enzymes, sucrose phosphorylase, UDP-glucose—hexose-1-phosphate uridylyltransferase, UDP-glucose 4-epimerase, and lacto-*N*-biose phosphorylase, in the presence of UDP-Glc and phosphate, indicating a reaction yield of 83%. LNB was isolated from the mixture by crystallization after yeast treatment. Finally, 1.4 kg of LNB of 99.6% purity was recovered after recrystallization.**

**Key words:** lacto-*N*-biose I; human milk oligosaccharides; lacto-*N*-biose phosphorylase; bifidus factor; *Bifidobacterium*

Bifidobacteria, gram-positive intestinal anaerobes, have been recognized to be beneficial to human health. Rapid intestinal colonization of bifidobacteria in breast-fed infants has been known since early twentieth century.<sup>1,2)</sup> The resulting intestinal flora is occupied by bifidobacteria and appears to prevent infection of some pathogenic bacteria and diarrhea.<sup>3)</sup> Such rapid colonization does not occur in the intestine of bottle-fed infants.<sup>3)</sup> It is widely accepted that oligosaccharides in human milk other than lactose (human milk oligosaccharides, HMOs) play a key role in the growth of bifidobacteria,<sup>4,5)</sup> but the growth mechanism is a long-term enigma due to the difficulty of research into the complicated contents of HMOs, which include more than 100 kinds of molecules.<sup>4)</sup> Now some prebiotic oligosaccharides such as lactulose<sup>6)</sup> are added to formula milk, which is made from cow's milk, which contains no oligosaccharides other than lactose,<sup>4,5,7)</sup> to grow bifidobacteria in the intestine, but the effect is considerably less than HMOs in forming intestinal flora, consisting

90% of bifidobacteria and 10% of enterobacteria.<sup>8)</sup> It has often been reported that early bifidobacterial colonization may reduce allergy.<sup>9)</sup> Thus, it is still desirable to improve the intestinal flora of bottle-fed infants to resemble the flora of breast-fed infants.

We found that bifidobacteria had a special galactose metabolism pathway specific to lacto-*N*-biose I (Gal $\beta$ 1 $\rightarrow$ 3GlcNAc, LNB) and galacto-*N*-biose (Gal $\beta$ 1 $\rightarrow$ 3GalNAc, GNB).<sup>10)</sup> The metabolism starts with the phosphorylation of disaccharides into  $\alpha$ -galactose 1-phosphate (Gal1P) and corresponding *N*-acetylhexosamines by lacto-*N*-biose phosphorylase (LNBP, EC 2.4.1.211). Then the Gal1P is converted into glucose 1-phosphate (Glc1P) as in the Leloir pathway. Since bifidobacteria possess endo- $\alpha$ -*N*-acetylgalactosaminidase, which liberates O- $\alpha$  linked GNB,<sup>11)</sup> the operon is considered to play a key role in intestinal colonization by utilizing GNB, which exists as a core I sugar chain in mucin.

Lacto-*N*-tetraose (Gal $\beta$ 1 $\rightarrow$ 3GlcNAc $\beta$ 1 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4Glc) and lacto-*N*-fucopentaose I (Fuc $\alpha$ 1 $\rightarrow$ 2 Gal $\beta$ 1 $\rightarrow$ 3GlcNAc $\beta$ 1 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4Glc), which show the LNB structure, are predominant components in HMOs as well as 2'-fucosyllactose.<sup>4)</sup> We noticed another possibility in the operon, that the LNB structure in HMOs can be metabolized in the same pathway if bifidobacteria possess lacto-*N*-biosidase and  $\alpha$ -fucosidase. Though the presence of lacto-*N*-biosidase has not been proved,  $\alpha$ -fucosidase has been found in *Bifidobacterium bifidum*.<sup>12)</sup> Hence we hypothesized that LNB is the real bifidus factor in human milk.<sup>10)</sup> To prove this hypothesis, practical preparation of LNB is required. In this paper we explain kg-scale preparation of LNB, which is scale-up ready.

The scheme of the reaction is summarized in Fig. 1. Sucrose and GlcNAc were chosen as the starting materials. These compounds were industrially prepared and are available at reasonable cost. Four enzymes, sucrose phosphorylase (SP, EC 2.4.1.7), UDP-glucose—

The nucleotide and deduced amino acid sequences reported here are available in the GenBank sequence database under accession nos. AB303573 (GalT), AB303838 (SP), and AB303839 (GalE).

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Abbreviations: GalE, UDP-glucose 4-epimerase; GalT, UDP-glucose—hexose 1-phosphate uridylyltransferase; GNB, galacto-*N*-biose; HMO, human milk oligosaccharide; LNB, lacto-*N*-biose I; LNBP, lacto-*N*-biose phosphorylase; SP, sucrose phosphorylase

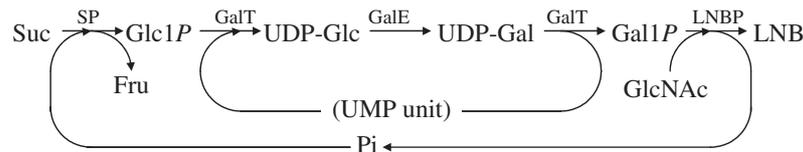


Fig. 1. Reaction Scheme for LNB Production.

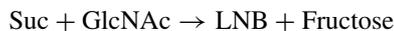
Table 1. Nucleotide Sequences for Cloning Primers, Preparation of Expression Plasmids, and Productivity and Specific Activity of Recombinant Enzymes

Enzyme	Primer sequence (5' → 3')	Vector/Restriction endonuclease	Productivity (mg/l-culture)	Specific activity (U/mg)
SP	gtcccttcgaggtaatcgaac cgggcgcacgtacggtccaacat gatatacatatgaaaaacaagtccaactc ggccaactcgaggtcgatcggcaatcggcgggtggc	pET30/ <i>NdeI</i> and <i>XhoI</i>	34	66
GalT	gatttagtaaagagctcgtgctgcc gaacagcttgtagcttcgaaacgcc caaacataggtcattcgccaactac ttctgcccgcgctgcgcatgctgatgga	pET30/ <i>NdeI</i> and <i>NotI</i>	60	210
GalE	ggcaacacggccgttccaagcagaagc ttcgaatcgaccgagctcaccgcgtggc gatatacatatgactactgttctggttacgggc ctgctcctcgagctcgcgctcggaaccgttggggttc	pET30/ <i>NdeI</i> and <i>XhoI</i>	125	140
LNBP*			32	3.5

\*derived from reference 12).

Primers are listed in the following order: cloning forward, cloning reverse, expression forward, and expression reverse. The restriction endonuclease sites are underlined.

hexose-1-phosphate uridylyltransferase (GalT, EC 2.7.7.12), UDP-glucose 4-epimerase (GalE, EC 5.1.3.2), and LNBP, were involved in the reaction. Sucrose is phosphorylated into  $\alpha$ -glucose 1-phosphate (Glc1P) and fructose by SP. Glc1P is converted into UDP-Glc concomitantly with the conversion of UDP-Gal into Gal1P by GalT. UDP-Glc is converted into UDP-Gal by GalE, and the resulting UDP-Gal is consumed in the reaction of GalT. Finally, LNB is synthesized from Gal1P and GlcNAc by LNBP. Because phosphate and UDP-Glc are recycled during the reaction system, the overall reaction is described as follows:



LNBP was prepared with the expression plasmid of *lnpA2* from *Bifidobacterium bifidum* JCM1254 in *Escherichia coli* as described previously.<sup>13)</sup> The SP, GalT, and GalE genes were isolated by PCR with KOD plus polymerase (Toyobo, Osaka, Japan) from *B. longum* JCM1217 genomic DNA prepared by InstaGene (Bio-Rad, Hercules, CA). Primers were designed based on the DNA sequence of the BL0536, BL1211, and BL1644 genes from *B. longum* NCC2705,<sup>14)</sup> as shown in Table 1. After sequencing of the PCR products, each gene was amplified by expression primers and inserted into pET 30 vector (Novagen, Darmstadt, Germany) with the restriction endonuclease site using Ligation High (Toyobo), as shown in Table 1. Transformation of *E. coli* BL21 (DE3) and purification of recombinant

enzymes from the transformant was performed according to the manufacturer's manual. Purification of recombinant enzymes from cell-free extract was performed using Ni-NTA agarose (Qiagen, Hilden, Germany) column chromatography. The homogeneity of the purified enzymes was confirmed by SDS-PAGE. Protein concentrations were determined using theoretical extinction coefficients calculated from the amino acid sequences of the protein.<sup>15)</sup> The yield of the four enzymes per 1-liter culture was calculated from the concentration of the enzymes after the purification step (Table 1).

The activities of the four enzymes were measured in 100 mM MOPS buffer (pH 7.5) at 30 °C. SP and GalT activities were quantified by determining the increase in the concentration of Glc1P<sup>16)</sup> from 100 mM sucrose and 100 mM phosphate (SP), and 4 mM UDP-Glc and 4 mM Gal1P (GalT), as substrates. GalE activity was determined by measuring the increase in UDP-Glc from 10 mM UDP-Gal as the substrate. LNBP activity was determined by measuring the increase in phosphate<sup>17)</sup> from 10 mM Gal1P and 10 mM GlcNAc as substrates. One unit of enzyme was defined as the amount of enzyme that produced 1  $\mu$ mol of product under above conditions. The specific activities of the four enzymes are shown in Table 1.

The production of LNB was performed in 10-liter of a reaction mixture consisting of 660 mM sucrose, 600 mM GlcNAc, 1 mM UDP-Glc, 30 mM phosphate buffer

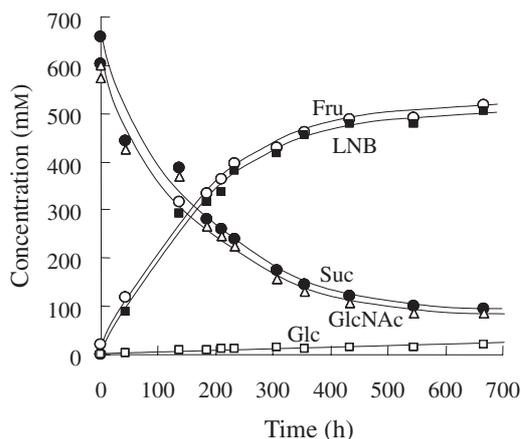


Fig. 2. The Time Course of LNB Production.

The concentrations of LNB, sucrose, fructose, GlcNAc, and glucose are shown by closed square, closed circle, open circle, open triangle, and open square respectively.

(pH 7.0), 10 mM  $MgCl_2$ , 26 mg/l (1716 U/l) SP, 22 mg/l (3080 U/l) GalE, 17 mg/l (3570 U/l) GalT, and 43 mg/l (150 U/l) LNBP at 30 °C. The concentrations of LNB, GlcNAc, sucrose, fructose, and glucose were monitored by high performance liquid chromatography (Shimadzu, Kyoto, Japan) with a Corona Charged Aerosol Detector (ESA Inc., Chelmsford, MA) using a Shodex Asahipak NH2P50-4E column (4.6 mm ID  $\times$  250 mm, Showa Denko, Tokyo, Japan) with acetonitrile- $H_2O$  (75:25 by volume) as the solvent at a flow rate of 1 ml/min. The time course of the reaction is shown in Fig. 2. Production of LNB was observed with decreases in sucrose and GlcNAc at an early stage of the reaction. The LNB concentration was reached 500 mM after 600 h of reaction. The yield was 83% based on the GlcNAc used.

After the reaction, 50 g of DEAE-cellulose (Wako Pure Chemical Industries, Osaka, Japan) pre-equilibrated with 50 mM phosphate buffer (pH 7.0) were added to the reaction mixture to adsorb the enzymes. The mixture was stirred at room temperature for 90 min. After removal of DEAE-cellulose by filtration, 200 g of baker's yeast (Oriental Yeast, Tokyo, Japan) was added to the mixture, which was incubated at 30 °C for 12 h to eliminate fructose (byproduct, 516 mM) and sucrose (remaining substrate, 96 mM).<sup>18)</sup> After centrifugation (18,000  $\times g$ , 30 min) to remove yeast cells, the supernatant was concentrated to 4.8-liter using a rotary evaporator. After crystallization at 4 °C for 12 h, 1.5 kg of crystalline LNB (96.9% purity) was recovered by filtration, followed by vacuum drying. In addition, 0.3 kg of crystalline LNB (95.0% purity) was recovered from the filtrate after concentration to 1.8-liter and crystallization under the same conditions.

The two LNB samples (1.5 kg and 0.3 kg) were mixed and dissolved in 3.6-liter water at 60 °C, followed by the addition of 3.6-liter of ethanol at 4 °C for crystallization with filtration and vacuum drying. Finally, 1.4 kg of

crystalline LNB (99.6% purity) was obtained after filtration and vacuum drying. The  $^1H$ -NMR spectrum of the compound was taken in  $D_2O$  using an Avance 800 spectrometer (Bruker Biospin, Rheinstetten, Germany) at 298 K. It was identical with that of authentic LNB. Anomeric analysis using HPLC<sup>19,20)</sup> revealed that the crystal was a  $\alpha$ -anomer of LNB.

The above results indicate that LNB was produced by a one-pot enzymatic reaction. All the unit processes used in this method were scale-up ready. It should be noted that the reaction time can easily be shortened only by increasing the amounts of the four enzymes at the same ratio. Although this method requires expensive UDP-Glc, the amount of UDP-Glc used was only 1/600 of GlcNAc, indicating that UDP-Glc acted as a catalyst with a turn-over number of 500. Considering these facts, this method can be used in industrial production of LNB by constructing a bio-reactor system using immobilized enzymes. Since LNB has been hypothesized to be a bifidus factor in HMOs, LNB may be applicable as an additive to formula milk to achieve bifidus flora in bottle-fed infants as well as breast-fed infants.

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## References

- 1) Benno, Y., and Mitsuoka, T., The development of gastrointestinal micro-flora in humans and animals. *Bifidobact. Microflora*, **5**, 13–25 (1986).
- 2) Rotimi, V. O., and Duerden, B. I., The development of the bacterial flora in normal neonates. *J. Med. Microbiol.*, **14**, 51–62 (1981).
- 3) Bezkorovainy, A., Ecology of bifidobacteria. In "Biochemistry and Physiology of Bifidobacteria," eds. Bezkorovainny, A., and Miller-Catchpole, R., CRC Press, Cleveland, pp. 29–72 (1989).
- 4) Kunz, C., Rudloff, S., Baier, W., Klein, N., and Strobel, S., Oligosaccharides in human milk: structural, functional, and metabolic aspects. *Annu. Rev. Nutr.*, **20**, 699–722 (2000).
- 5) Newburg, D. S., and Neubauer, S. H., Carbohydrate in milks: analysis, quantities, and significance. In "Handbook of Milk Composition," ed. Jensen, R. G., Academic Press, San Diego, pp. 273–349 (1995).
- 6) Petuely, F., Bifidusflora bei Fräschenkindern durch bifidogene Substanzen (Bifidusfaktor). *Z. Kinderheilkd.*, **79**, 174–179 (1957).
- 7) Nakamura, T., and Urashima, T., The milk oligosaccharides of domestic farm animals. *Trends Glycosci. Glycotech.*, **16**, 135–142 (2004).
- 8) Benno, Y., Sawada, K., and Mitsuoka, T., The intestinal microflora of infants: composition of fecal flora in breast-fed and bottle-fed infants. *Microbiol. Immunol.*, **28**, 975–986 (1984).
- 9) Moro, G., Arslanoglu, S., Stahl, B., Jelinek, J., Wahn, U.,

- and Boehm, G., A mixture of prebiotic oligosaccharides reduces the incidence of atopic dermatitis during the first six months of age. *Arch. Dis. Child.*, **91**, 814–819 (2006).
- 10) Kitaoka, M., Tian, J., and Nishimoto, M., Novel putative galactose operon involving lacto-*N*-biose phosphorylase in *Bifidobacterium longum*. *Appl. Environ. Microbiol.*, **71**, 3158–3162 (2005).
  - 11) Fujita, K., Oura, F., Nagamine, N., Katayama, T., Hiratake, J., Sakata, K., Kumagai, H., and Yamamoto, K., Identification and molecular cloning of a novel glycoside hydrolase family of core 1 type *O*-glycan-specific endo- $\alpha$ -*N*-acetylgalactosaminidase from *Bifidobacterium longum*. *J. Biol. Chem.*, **280**, 37415–37422 (2005).
  - 12) Katayama, T., Sakuma, A., Kimura, T., Makimura, Y., Hiratake, J., Sakata, K., Yamanoi, T., Kumagai, H., and Yamamoto, K., Molecular cloning and characterization of *Bifidobacterium bifidum* 1,2- $\alpha$ -L-fucosidase (AfcA), a novel inverting glycosidase (glycoside hydrolase family 95). *J. Bacteriol.*, **186**, 4885–4893 (2004).
  - 13) Nishimoto, M., and Kitaoka, M., Identification of the putative proton donor residue of lacto-*N*-biose phosphorylase (EC 2.4.1.211). *Biosci. Biotechnol. Biochem.*, **71**, 1587–1591 (2007).
  - 14) Schell, M. A., Karmirantzou, M., Snel, B., Vilanova, D., Berger, B., Pessi, G., Zwahlen, M. C., Desiere, F., Bork, P., Delley, M., Pridmore, R. D., and Arigoni, F., The genome sequence of *Bifidobacterium longum* reflects its adaptation to the human gastrointestinal tract. *Proc. Natl. Acad. Sci. USA*, **99**, 14422–14427 (2002).
  - 15) Gill, S. C., and von Hippel, P. H., Calculation of protein extinction coefficients from amino acid sequence data. *Anal. Biochem.*, **182**, 319–326 (1989).
  - 16) Michael, G., D-Glucose-1-phosphate. In “Methods of Enzymatic Analysis” Vol. 6, 3rd ed., eds. Bergmyer, H. U., Bergmeyer, J., and Grassl, M., Verlag Chemie, Weinheim, pp. 185–191 (1984).
  - 17) Lowry, O. H., and Lopez, J. A., The determination of inorganic phosphate in the presence of labile phosphate esters. *J. Biol. Chem.*, **162**, 421–428 (1946).
  - 18) Yoon, S. H., Mukerjea, R., and Robyt, J. F., Specificity of yeast (*Saccharomyces cerevisiae*) in removing carbohydrates by fermentation. *Carbohydr. Res.*, **338**, 1127–1132 (2003).
  - 19) Honda, Y., and Kitaoka, M., A family 8 glycoside hydrolase from *Bacillus halodurans* C-125 (BH2105) is a reducing end xylose-releasing exo-oligoxylanase. *J. Biol. Chem.*, **279**, 55097–55103 (2004).
  - 20) Honda, Y., Kitaoka, M., Tokuyasu, K., Sasaki, C., Fukamizo, T., and Hayashi, K., Kinetic studies on the hydrolysis of *N*-acetylated and *N*-deacetylated derivatives of 4-methylumbelliferyl chitobioside by the family 18 chitinases ChiA and ChiB from *Serratia marcescens*. *J. Biochem. (Tokyo)*, **133**, 253–258 (2003).