

# Expression and Characterization of Human $\beta$ -1,4-Galactosyltransferase 1 ( $\beta$ 4GalT1) Using Silkworm–Baculovirus Expression System

Daisuke Morokuma<sup>1</sup> · Jian Xu<sup>1</sup> · Masato Hino<sup>1</sup> · Hiroaki Mon<sup>1</sup> · Jasmine S. Merzaban<sup>2</sup> · Masateru Takahashi<sup>3</sup> · Takahiro Kusakabe<sup>1</sup> · Jae Man Lee<sup>1</sup>

Published online: 24 March 2017  
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**Abstract** Baculovirus expression vector system (BEVS) is widely known as a mass-production tool to produce functional recombinant glycoproteins except that it may not be always suitable for medical practice due to the differences in the structure of *N*-linked glycans between insects and mammalian. Currently, various approaches have been reported to alter *N*-linked glycan structures of glycoproteins derived from insects into terminally sialylated complex-type *N*-glycans. In the light of those studies, we also proposed in vitro maturation of *N*-glycan with mass-produced and purified glycosyltransferases by silkworm–BEVS.  $\beta$ -1,4-Galactosyltransferase 1 ( $\beta$ 4GalT1) is known as one of type II transmembrane enzymes that transfer galactose in a  $\beta$ -1, 4 linkage to acceptor sugars, and a key enzyme for further sialylation of *N*-glycans. In this study, we developed a large-scale production of recombinant human  $\beta$ 4GalT1 (rh $\beta$ 4GalT1) with N- or C-terminal tags in silkworm–BEVS. We demonstrated that rh $\beta$ 4GalT1 is *N*-glycosylated and without mucin-type glycosylation.

Interestingly, we found that purified rh $\beta$ 4GalT1 from silkworm serum presented higher galactosyltransferase activity than that expressed from cultured mammalian cells. We also validated the UDP-galactose transferase activity of produced rh $\beta$ 4GalT1 proteins by using protein substrates from silkworm silk gland. Taken together, rh $\beta$ 4GalT1 from silkworms can become a valuable tool for producing high-quality recombinant glycoproteins with mammalian-like *N*-glycans.

**Keywords** Baculovirus expression vector system ·  $\beta$ -1,4-Galactosyltransferase 1 · *N*-glycosylation · Silkworm

## Introduction

The migration of cells to tissues/organs is controlled by a specific cell surface sugar decoration, mainly sialyl Lewis<sup>x/a</sup> (sLe<sup>x/a</sup>). This key epitope is created through the activity of specific glycosyltransferases (GTs). When these GTs are accumulated and/or activated in particular cells or are introduced ex vivo, they migrate to specific locations where the counter-receptors (selectins) are expressed. Selectin-dependent tethering and rolling decreases cellular velocity below that of the prevailing hemodynamic stream. *P*- and *E*-selectin are membrane-bound lectins that bind to glycoprotein or glycolipid ligands (e.g., PSGL-1 and CD43) on the cell in flow via their lectin domains. The specialized glycan determinant for selectin ligand is comprised of sialofucosylations containing an  $\alpha$  (2,3)-linked sialic acid substitution on galactose, and an  $\alpha$  (1,3)-linked fucose modification on *N*-acetylglucosamine, prototypically displayed as the terminal tetrasaccharide sLe<sup>x</sup> (or its isomer sLe<sup>a</sup>).

In order for selectin ligands to be formed, GTs responsible for creating the sLe<sup>ax</sup> structures must be

✉ Jae Man Lee  
jaemanle@agr.kyushu-u.ac.jp

<sup>1</sup> Laboratory of Insect Genome Science, Kyushu University Graduate School of Bioresource and Bioenvironmental Sciences, Hakozaki 6-10-1, Higashi-ku, Fukuoka 812-8581, Japan

<sup>2</sup> Laboratory of Cell Migration and Signaling, Division of Biological and Environmental Sciences and Engineering Division, King Abdullah University of Science and Technology, 4700 KAUST Thuwal, Jeddah 23955, Saudi Arabia

<sup>3</sup> Laboratory of DNA Replication and Recombination, Division of Biological and Environmental Sciences and Engineering, King Abdullah University of Science and Technology, 4700 KAUST Thuwal, Jeddah 23955, Saudi Arabia

endogenously expressed and active. The native display of these glycan structures on selectin ligands requires the various active GTs, including the action of  $\alpha$ 1,3- or  $\alpha$ 1,4-fucosyltransferases,  $\alpha$ 2,3-sialyltransferases,  $\beta$ 1,4-galactosyltransferases, and  $\beta$ 1,6-*N*-acetyl-glucosaminyltransferases. The knowledge of which GTs are required to create particular structures allows researchers to manipulate cells in order to create ligands on cells' surface *ex vivo* with the intention of helping to direct the migration of therapeutic cells to specific tissues that express the corresponding selectins [1–3]. Selectin ligands must express sLe<sup>x/a</sup> structures on extended lactosamines of *N*- or *O*-glycans. Adult stem cells are often inadequately  $\alpha$ 1,3-fucosylated, this being the case with mesenchymal stem cells (MSC) [1], hematopoietic stem cells (HSC) [4–7] and neural stem cells [2] and so exhibit migration defects. Indeed, *ex vivo* treatment of stem cells with FTs, particularly FT-VI and FT-VII, increases cell surface sLe<sup>x</sup> determinants, boosts binding to *E*-selectin and *P*-selectin, and enhances homing and engraftment into bone marrow of immunocompromised mice [4–7]. By generating a comprehensive set of tools (i.e.,  $\beta$ 1,4-galactosyltransferases,  $\beta$ 1,6-*N*-acetyl-glucosaminyltransferases,  $\alpha$ 1,3- or  $\alpha$ 1,4-fucosyltransferases and  $\alpha$ 2,3-sialyltransferases) for *ex vivo* design of sLe<sup>x</sup> structures on the surface of a therapeutic cell, strategies to enhance the migration of these cells for the treatment of diseases can be further anticipated.

The  $\beta$ 1,4-galactosyltransferase ( $\beta$ 4GalT1) gene family consists of seven members that transfer galactose in a  $\beta$ -1,4 linkage to acceptor sugars.  $\beta$ 4GalT1 (NCBI accession number NP\_001488) is unique among the seven enzymes because it can be expressed either as membrane associated form or secreted form [8–10]. It is also one of the key enzymes responsible for extending the lactosamine structures on *N*- and *O*-glycans as well as adding the terminal galactose that will be capped with  $\alpha$ (2,3)-linked sialic acid to generate the sLe<sup>x</sup> selectin ligand. Thus, expression and purification of this enzyme would be beneficial for the *ex vivo* creation of selectin ligands on therapeutic cells.

To this end, the mammalian expression system is often applied for the production of glycoproteins with high activity and stability, although it offers suffers from several disadvantages, such relatively poor productivity and higher cost. To date, various approaches have been reported to alter the *N*-linked glycan structure of insect-derived proteins into terminally sialylated complex-type *N*-glycans by transgenic expression of mammalian glycosyltransferases in insect cells [11–13]. However, the efficiency in the formation of matured complex-type *N*-glycans from the insect cells is still low because the activity of introduced mammalian  $\beta$ -1,4-galactosyltransferase 1 ( $\beta$ 4GalT1) is unstable in insect cell [14]. Additionally, the negative effect of exogenously expressed  $\beta$ 4GalT1 in insect cells

must be taken into account especially in transgenic insects. Recently, the baculovirus expression vector system (BEVS) using lepidopteran insects and cells has been reported to be suitable for mass-production of glycoproteins with mammalian-like post-translational modifications and safety profiles [15–19]. Therefore, we propose another approach in which the recombinant glycoproteins produced by silkworm–BEVS are further processed by *in vitro N*-glycosylation using mammalian glycosyltransferases purified from larvae. Because galactosylated *N*-glycans provide substrates for sialylation by sialyltransferases, it is desirable to mass-produce the recombinant  $\beta$ 4GalT1 by silkworm–BEVS. In this study, we constructed the recombinant BmNPV to express the secreted form human  $\beta$ 4GalT1 in cultured silkworm cells and larvae. We also report the expression, purification and characterization of this recombinant human  $\beta$ 4GalT1 secreted from silkworm larvae.

## Materials and Methods

### Cells and Silkworm Larvae

The NIAS-Bm-oyanagi2 (BmO2, kindly provided from Dr. Imanishi) and Bme21 cells [20] were cultured in IPL-41 medium (Sigma, St. Louis, MO) supplemented with 10% fetal bovine serum (Gibco, Grand Island, NY); the silkworm larvae were provided from the Institute of Genetic Resources, Kyushu University Graduate School. The larvae were reared on fresh mulberry leaves at 25–27 °C.

### Construction of Recombinant Human $\beta$ 4GalT1

Matured form of human  $\beta$ 4GalT1 (a.a.77–398) gene was amplified using the specific primers  $\beta$ 4GalT1-matured-5 (5'-CGGACCGGAGGGGCCCGCC-3') and  $\beta$ 4GalT1-3-Xho1 (5'-GTCCCTCGAGCTCGGTGTCCCGATGTCCA C-3'). The amplified PCR product without 1–77 amino acid residues of  $\beta$ 4GalT1 including native cytosolic (C) and transmembrane (T) domains was digested by *Xho*1, and further inserted into the *Eco*RV and *Xho*1 double-digested site of the modified pENTR11 vectors (pENTR11L2130KT EVH8STREP and pENTR11L2130KH8STREPTEV) containing a lobster L21 sequence at the N-terminal, and the tobacco etch virus (TEV) protease cleavage site and the tobacco etch virus (TEV) protease cleavage site for removing the affinity tag from the recombinant protein in case terminal tags affect the protein stability and/or activity, the 8 × histidine tag and the Strep-tag at the N- or C-terminal. The resulting constructs were named pENTR11/30K\_rh $\beta$ 4GalT1 (N-tag) and pENTR11/30K\_rh $\beta$ 4GalT1 (C-tag), respectively.

## Generation of Recombinant BmNPV

The transfer plasmids for generating the recombinant baculoviruses were subsequently constructed by LR Gateway reaction between the pENTR11/30K\_rh $\beta$ 4GalT1 (N-tag) and pENTR11/30K\_rh $\beta$ 4GalT1 (C-tag) and pDEST8 vector (Invitrogen, Carlsbad, California) according to the manufacturer's protocol. The recombinant baculoviruses were generated using BmNPV/T3 bacmid system as described previously [21]. A 200 ng of the bacmid DNA was transfected into  $1 \times 10^5$  BmO2 cells in 24-well plates using FuGENE HD transfection reagent (Promega, Madison, WI) to yield the recombinant virus particles, BmNPV/*polh*-30K\_rh $\beta$ 4GalT1 (N-tag) and BmNPV/*polh*-30K\_rh $\beta$ 4GalT1 (C-tag), respectively. Before transfection, the culture medium was replaced by FBS-free COSMEDIUM 009X medium (Cosmo-Bio, Tokyo, Japan). At 4-day post-infection (dpi), cell culture mediums containing P1 virus particles were collected and stored at 4 °C in the dark. High-titer virus (P3) stocks were prepared by serial infection following the protocols recommended in the manufacturer's manual (Invitrogen).

## Expression of rh $\beta$ 4GalT1 in Cultured Silkworm Cell

$2 \times 10^6$  Bme21 cells in a 12-well plate were infected with the recombinant BmNPV at a multiplicity of infection (MOI) of 10. At 4 dpi, after centrifugation at 1000g for 10 min at 4 °C, the supernatant was subjected to Western blotting analysis. The cell pellet was suspended in ice-cold lysis buffer (20 mM Tris–HCl pH 7.4, 0.5 M NaCl, 1% Triton X-100, 10% glycerol, 10 mM 2-mercaptoethanol, 1 mM PMSF) and homogenized. After centrifugation at 15,000g for 30 min at 4 °C, the supernatant containing soluble proteins was also subjected to Western blotting analysis as cell lysate.

## Purification of rh $\beta$ 4GalT1 from Silkworm Larvae

The recombinant virus was injected into the 5th instar of n17 silkworm larvae. At 4 dpi, 10 ml of the serum from roughly 25 silkworm larvae was collected into a 15-ml tube containing 20 mM 1-phenyl-2-thiourea, followed by centrifugation at 10,000g for 30 min at 4 °C. Then, the supernatant was transferred into a new tube. The resulting serum was diluted by buffer A (20 mM Tris–HCl pH 7.4, 0.5 M NaCl, 1 mM PMSF), and centrifuged at 20,000g for 30 min at 4 °C. After the filtration through a 0.45- $\mu$ m filter (Millipore, USA), the supernatant containing rh $\beta$ 4GalT1 was loaded to a nickel affinity chromatography with 5 ml HisTrap excel column (GE Healthcare Bioscience, USA), and eluted by 500 mM imidazole solution buffers. After concentrated by ultrafiltration using Amicon 10 K filters

(Millipore, USA) in buffer B (100 mM Tris–HCl pH 8.4, 150 mM NaCl, 1 mM EDTA), the rh $\beta$ 4GalT1 was applied to StrepTrap HP column (GE Healthcare Bioscience, USA), and then eluted by the buffer containing 2.5 mM desthiobiotin. The purified proteins were determined of quantity by using DC<sup>TM</sup> Protein Assay (BioRad).

## Glycosylation Analysis with Glycosidase Digestions

The purified rh $\beta$ 4GalT1s and the commercially available rh $\beta$ 4GalT1 (a.a.44–398; R&D systems, Inc., Minneapolis, USA) expressed in mouse myeloma cell lines were incubated in Glycoprotein Denaturing Buffer (NEB, USA) at 100 °C for 10 min. Subsequently, the denatured protein samples were treated by adding PNGase F (NEB, USA), or *O*-glycosidase (NEB, USA) with  $\alpha$ 2-3,6,8 Neuraminidase (NEB, USA) for 1 h at 37 °C according to the manufacturer's protocol.

## SDS-PAGE and Western Blotting

The purified or deglycosylated rh $\beta$ 4GalT1 was subjected to SDS-PAGE (with 10% polyacrylamide gel) and visualized by Coomassie Brilliant Blue R-250 staining or transferred to a polyvinylidene difluoride (PVDF) membrane for Western blotting. As for Western blotting, the membrane was blocked for 1 h in TBST buffer (20 mM Tris–HCl pH 7.6, 500 mM NaCl, 0.1% w/v Tween-20) with 5% w/v skim milk (Wako, Japan) followed by incubation with HisProbe-HRP (1:5000 v/v; Thermo Scientific, USA) for 1 h. After washing, the HRP signals were detected by the Super Signal West Pico Chemiluminescent Substrate (Thermo Scientific, USA).

## Galactosyltransferase Activity Against N-acetylglucosamin

The transfer activity of rh $\beta$ 4GalT1 was determined in vitro using the Glycosyltransferase Activity Kit (R&D Systems, Inc., Minneapolis, USA), following the assay procedure of rh $\beta$ 4GalT1 from R&D systems. Briefly, the dissociative UDP after the galactosyltransferase reaction is further hydrolyzed by the calcium-dependent nucleotidase ENTPD3/CD39L3 giving rise to free inorganic phosphate, which can be detected by the malachite green reagent. The reaction was initiated by incubating 5 nmol of UDP-galactose (UDP-Gal, SIGMA) as a donor substrate, 500 nmol *N*-Acetyl-alpha-D-glucosamine (GlcNAc, SIGMA) as an acceptor substrate, 200 ng of ENTPD3/CD39L3, 150 ng of the recombinant proteins in the assay buffer (25 mM Tris–HCl pH 7.5, with 10 mM CaCl<sub>2</sub>, 10 mM MnCl<sub>2</sub>) using 96-well microplate for 15 min at 37 °C, and terminated by adding the malachite green

development solution. After 20 min incubation with Malachite Green Reagent B, the microplate was read absorbance at 620 nm in endpoint mode of Synergy HTX Plate Reader (BioTek, Japan). As a negative control, recombinant human alpha-1-acid glycoprotein ( $\alpha$ 1AGP) purified in our previous study [22] was used. All reactions were performed in triplicate independently.

### Galactosyltransferase Activity Against Endogenous Glycoproteins Extracted from Silk Glands of Silkworm Larvae

The endogenous glycoproteins in posterior silk glands (PSG) from 5 silkworm larvae (5th instar) were lysed in 5 ml ice-cold extraction buffer (20 mM Tris-HCl pH 7.4, 0.5 M NaCl, 1% Triton X-100, 10% glycerol, 10 mM 2-mercaptoethanol, 1 mM PMSF) for 6 h. Subsequently, 300  $\mu$ L crude extraction of PSG was precipitated in acetone (1:3, v/v) at  $-20$  °C for 2 h. After centrifugation at 12,000g in 4 °C for 5 min, the protein pellet was dried and re-resolved in 100  $\mu$ L the assay buffer (25 mM Tris-HCl pH 7.5, with 10 mM CaCl<sub>2</sub>, 10 mM MnCl<sub>2</sub>). The resulting solution including endogenous glycoprotein in silk gland was named PSG. The in vitro glycosylation assay was initiated by incubating 1  $\mu$ L SG with 100 ng of rh $\beta$ 4GalT1 (C-tag or N-tag) and 10 mM UDP-galactose (UDP-Gal, SIGMA) as a donor in a total volume of 10  $\mu$ L assay buffer for 1 h at 37 °C. After incubation, the resulting solution was subjected to SDS-PAGE and transferred to a PVDF membrane for lectin blotting. Briefly, the membrane was blocked for 30 min in a carbo-free blocking solution (Vector Laboratories, USA), followed by incubation with *Ricinus Communis* Agglutinin I (RCA120, specifically for terminal  $\beta$ -D-galactosyl residues; HRP-conjugated, 2 ppm, J-OILMILLS, Japan) diluted in PBST buffer (PBS pH 7.5, 0.05% Tween-20) for 1 h. After washing twice with PBST buffer, the HRP signals were visualized by the Super Signal West Pico Chemiluminescent Substrate (Thermo Scientific, USA).

## Results

### Construction of Expression Vectors

As illustrated in Fig. 1a, human  $\beta$ 4GalT1 reported to be composed of 398 amino acid residues, contains a cytoplasmic domain (a.a.1–24), a type-2 transmembrane domain (a.a.25–44) and a luminal domain (a.a.45–398). Generally, human  $\beta$ 4GalT1 is localized primarily in the plasma membrane and exists in the *trans* cisternae of the Golgi complex [10]. Previous studies suggest that the mature form of the human  $\beta$ 4GalT1 is secreted into body

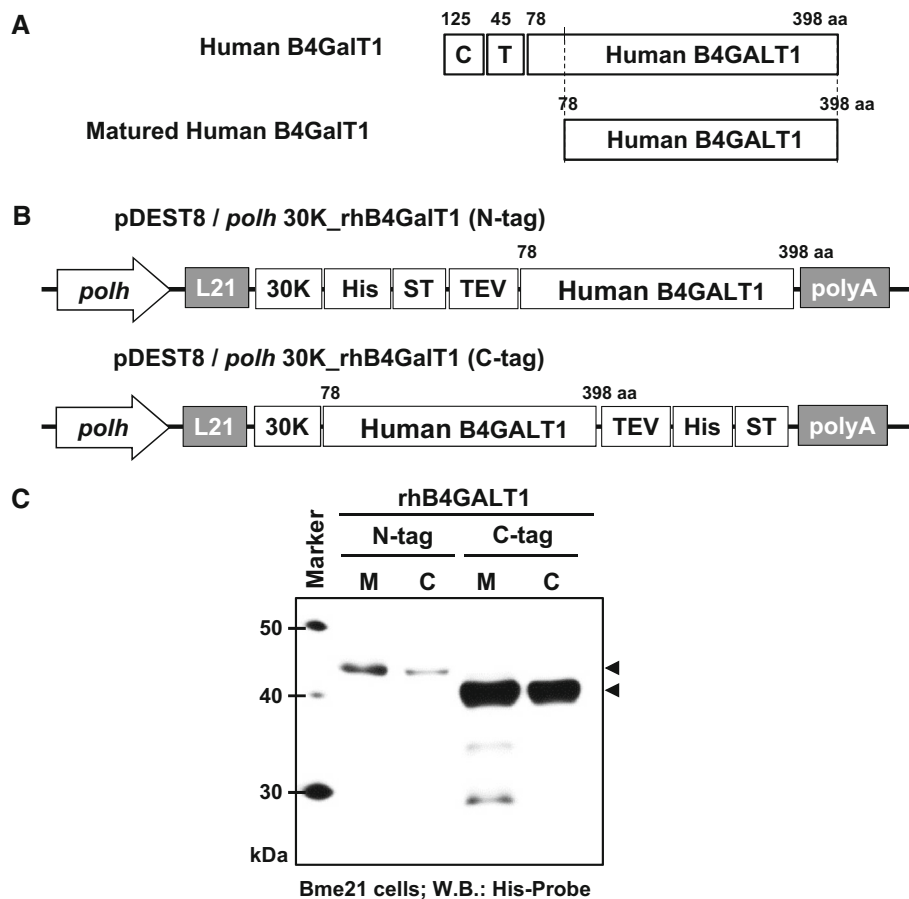
fluids by cleaving its catalytic domain between L77 and R78 [8, 9] (Fig. 1a). Accordingly, it is reasonable to consider that a significant level of the recombinant  $\beta$ 4GalT1 flows out from Golgi apparatus, and the intracellular GT activity becomes relatively lower [14], even when a large amount of protein is generated using the BEVS. Here, we created the expression construct for the catalytic domain of human  $\beta$ 4GalT1 (a.a.78–398) (rh $\beta$ 4GalT1) with a silkworm 30 K signal peptide to induce effective secretion [23]. In addition to aid in the efficient purification of the enzyme, 8  $\times$  His-tag and Strep-tag were added to the N- or C-terminus of the recombinant human  $\beta$ 4GalT1. Therefore, we generated two types of expression vectors for the expression of recombinant human  $\beta$ 4GalT1 (Fig. 1b).

### Expression Verification of rh $\beta$ 4GalT1 in BmNPV-Infected Cultured Silkworm Cells

The expressions of two kinds recombinant baculoviruses of rh $\beta$ 4GalT1 were firstly examined in BmNPV-hypersensitive Bme21 cells [20]. The predicted molecular masses of the rh $\beta$ 4GalT1 s with N-terminal or C-terminal tags were 39.7 and 39.9 kDa, respectively. As demonstrated in Fig. 1c, both rh $\beta$ 4GalT1 s with N-tag or C-tag were expressed and secreted into the culture medium (M). Interestingly, the molecular weight of the N-tagged rh $\beta$ 4GalT1 estimated from the mobility on SDS-PAGE was higher than the predicted size. Besides, the C-tagged rh $\beta$ 4GalT1 secreted into culture medium generated three bands containing 8  $\times$  His-tag, indicating the possibilities that C-tagged rh $\beta$ 4GalT1 was hydrolyzed from its N-terminus or simply because alternative ATG start codons were used for protein translation. Furthermore, the expression level of the C-tagged rh $\beta$ 4GalT1 was significant higher than of the N-tagged, suggesting that the C-terminus is ideal for improving the yields of rh $\beta$ 4GalT1.

### Purification of rh $\beta$ 4GalT1 from Silkworm Larvae

The rh $\beta$ 4GalT1 s secreted into silkworm hemolymph were collected at 4 dpi of the indicated recombinant BmNPV, and then purified using two-step affinity chromatography. As a result, approximately 124  $\mu$ g of the N-tagged (Fig. 2a) and 239  $\mu$ g of the C-tagged (Fig. 2b) highly purified rh $\beta$ 4GalT1 were recovered from 10 ml of hemolymph from silkworm larvae. As far as we are aware, this is the first study reporting the expression and purification of human  $\beta$ 4GalT1 on this scale using this system. Unlike the expression in Bme21 cells illustrated in Fig. 1c, the purified N-tagged and C-tagged rh $\beta$ 4GalT1 s generated one major band at approximately 42 or 40 kDa, and smear multiply bands ranging from 42–45 to 40–45 kDa, respectively (Fig. 2a, b). These results suggest that the



**Fig. 1** Functional domains of full-length human  $\beta$ 4GalT1 are shown, including the cytoplasmic domain (C), transmembrane (T) and luminal domain containing  $\beta$ 4GalT1 catalytic domain. By cleaving between L77 and R78, matured form of human  $\beta$ 4GalT1 was generated (a). Construction of the vectors for expression the rh $\beta$ 4GalT1 (matured form). The plasmid pDEST8/*polh* 30 K\_rh $\beta$ 4GalT1 (N-tag) and pDEST8/*polh* 30K\_rh $\beta$ 4GalT1 (C-tag) were generated from pENTR and pDEST8 vector by Gateway

reaction. These vectors were under the control of polyhedron (*polh*) promoter and followed by SV40 polyadenylation signal (polyA). L21: leader sequence for enhancing translation efficiency; 30K: signal peptide of silkworm 30 kDa protein; TEV: tobacco etch virus protease cleavage site; His: 8 x histidine tag; ST: strep-tag (b). Expression of the rh $\beta$ 4GalT1 in Bme21 cells. At 4 dpi, the rh $\beta$ 4GalT1 s were detected from culture medium (M) and cell lysate (C) by Western blotting using His-Probe (c)

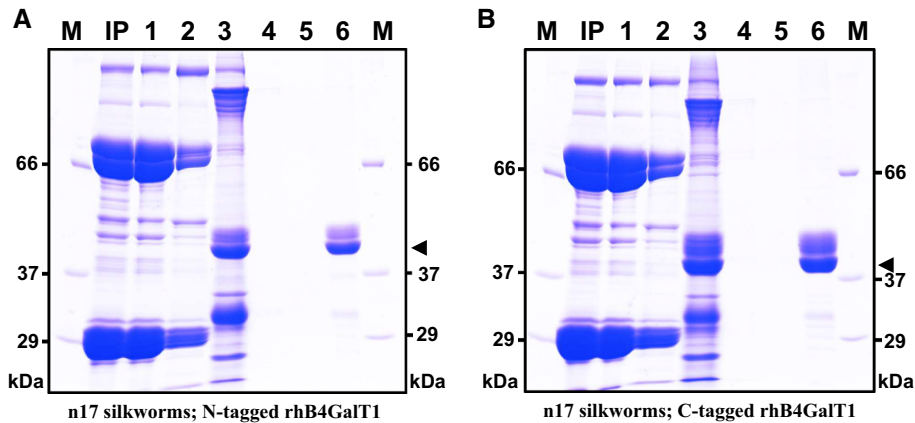
resulting proteins were potentially with certain post-translational modifications (PTMs), e.g., glycosylation [24].

### Molecular Characterization of rh $\beta$ 4GalT1 s from Silkworm Larvae

Human  $\beta$ 4GalT1 is expected to contain 1 *N*-glycosylation site (N113) and the previously study showed its *N*-linked glycan has triantennary or biantennary complex-type structures [24]. In addition, 5 potential *O*-glycosylation sites are predicted in the amino acid sequence of human  $\beta$ 4GalT1, but confirmation of the existence of these *O*-glycans in human  $\beta$ 4GalT1 has not been reported until now. Therefore, we compared the *N*- or *O*-linked glycan structures of rh $\beta$ 4GalT1 s from silkworm larvae with the commercially available rh $\beta$ 4GalT1 (a.a.44–398; R&D

systems, Inc., Mineapolis, USA) expressed in mouse myeloma cell lines as control. As shown in Fig. 3, the rh $\beta$ 4GalT1 s and the commercial  $\beta$ 4GalT1 were treated with PNGase F to cleave potential *N*-linked glycans, or *O*-glycosidase along with  $\alpha$ 2-3,6,8 Neuraminidase to cleave possible *O*-linked glycans. As shown in Fig. 3, all C-tagged, N-tagged and commercial rh $\beta$ 4GalT1s were sensitive and cleavable by PNGase F, which proved that rh $\beta$ 4GalT1s secreted from silkworm larvae were *N*-glycosylated similarly to that expressed by mouse cells. On the other hand, after treatment with *O*-glycosidase, the electrophoretic mobility of C-tagged or N-tagged rh $\beta$ 4GalTs was not changed. Besides, even with treatment of the PNGase F, the rh $\beta$ 4GalT1 s from silkworm larvae did not show a single product, which we considered that the products from smear bands were not generated by glycosylation, possibly an unknown PTMs.

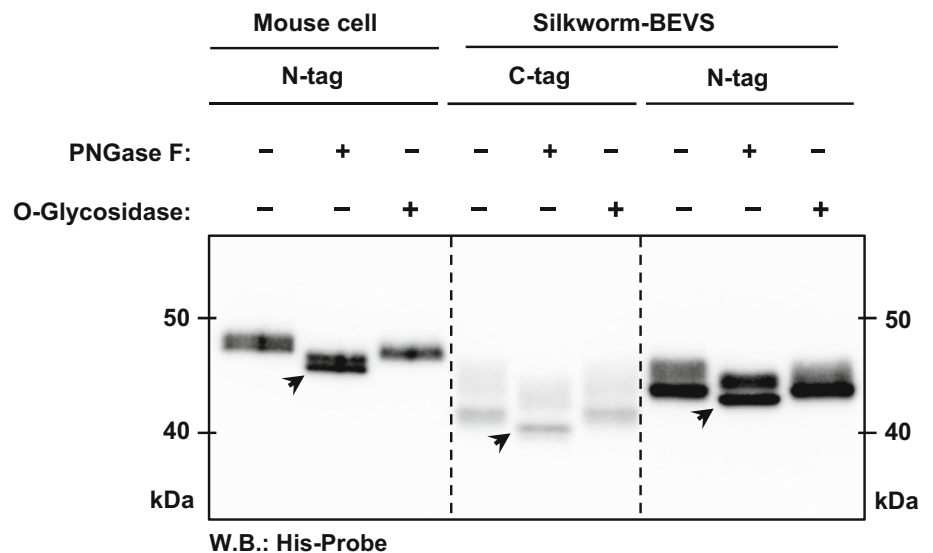




**Fig. 2** Purification of the N- or C-tagged rh $\beta$ 4GalT1 from sera of infected silkworm larvae (**a** C-tagged; **b** N-tagged). The His and Strep double-tagged rh $\beta$ 4GalT1 were purified through nickel affinity chromatography (lanes IP, 1, 2, 3) and Strep-Tactin affinity chromatography (lanes 4, 5, 6) as described in Methods. Each of

the fractions was resolved on 10% SDS-PAGE and visualized by Coomassie Brilliant Blue (CBB) R-250. *M* molecular mass markers; *Lane 1* flow-through fraction; *Lane 2* wash fraction; *Lane 3* eluent fraction (500 mM imidazole); *Lane 4* flow-through fraction; *Lane 5* wash fraction; *Lane 6* eluent fraction (2.5 mM desthiobiotin)

**Fig. 3** Characterization of the C-tagged and N-tagged rh $\beta$ 4GalT1 secreted in silkworm larval serum. The commercial rh $\beta$ 4GalT1 (with N-terminal tag) from mouse cell lines, C-tagged and N-tagged rh $\beta$ 4GalT1 produced by BEVS were incubated with (+) or without (–) PNGase F or O-glycosidase with  $\alpha$  2-3,6,8 Neuraminidase for 1 h at 37 °C. After reaction, each mixture was resolved on 10% SDS-PAGE and visualized by Western blotting using His-Probe

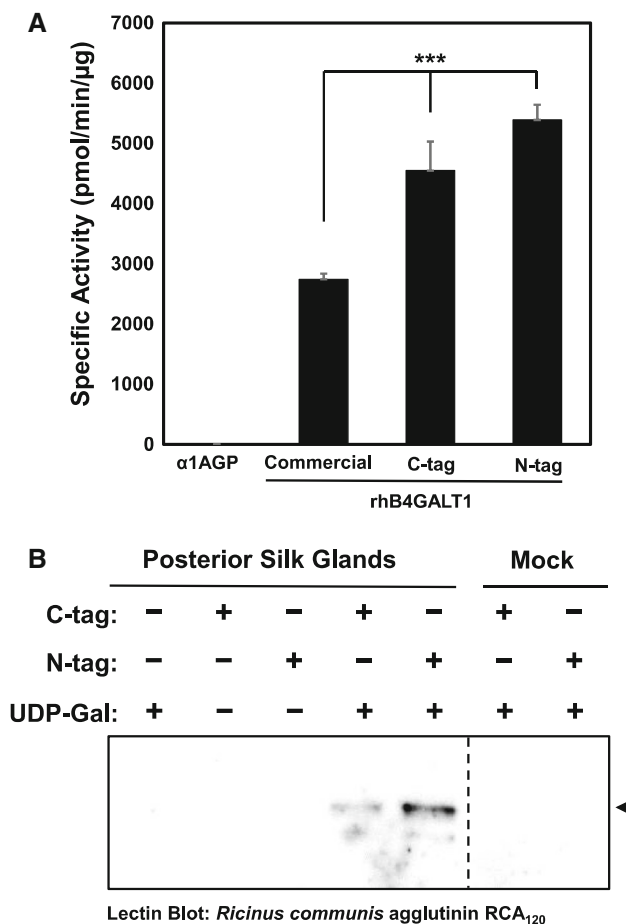


### Galactosyltransferase Activity Assay to Test the Activity

To compare the galactosyltransferase activities of the rh $\beta$ 4GalT1 s from silkworm larvae with that from mouse cells, we performed an in vitro activity assay using a Glycosyltransferase Activity Kit (R&D Systems, Inc., Minneapolis, USA). As a negative control, the secretion protein from silkworm larvae, recombinant human  $\alpha$ 1AGP [22] was utilized in this study. As shown in Fig. 4a, both of the C-tagged and N-tagged rh $\beta$ 4GalT1 exhibited much higher activity of about 1.6–2.0-fold than the rh $\beta$ 4GalT1 expressed in mammalian cell lines did. Furthermore, we also learned that the N-tagged rh $\beta$ 4GalT1 showed higher activity than the C-tagged protein, though the C-tagged construct provided more production, indicating C-terminal

tags may slightly affect the glycol-transferring and thus it is rescannable to consider that N-terminal design is better for functional production of rh $\beta$ 4GalT1. Moreover, from the results in Fig. 1c, the C-tagged rh $\beta$ 4GalT1 appears to have less enzyme stability than that of N-tagged protein although it was expressed at a higher level.

Commonly, *N*-glycans of glycoproteins secreted from insect cells have paucimannosidic structures that lack the terminal GlcNAc that could act as an acceptor for galactose, likely due to the extensive  $\beta$ -*N*-acetylglucosaminidase (FDL) activity causing the removal of the terminal GlcNAc from the hybrid-type structure [25]. However, the previous research showed glycoproteins produced in silk glands in silkworm larvae have complex-type *N*-glycans with terminal GlcNAc [26]. In order to confirm the in vitro galactosyltransferase activity of rh $\beta$ 4GalT1, *N*-glycans of



**Fig. 4** Galactosyltransferase activity assay of the rh $\beta$ 4GalT1. The specific activities of the commercial rh $\beta$ 4GalT1 produced in mammalian cell lines, the C-tagged and N-tagged rh $\beta$ 4GalT1 from silkworm larvae were measured by Glycosyltransferase Activity Kit (a). As a negative control of recombinant glycoprotein secreted silkworm larvae, we used the recombinant human  $\alpha$ 1AGP. Bars and error bars indicate the mean and SD of values, respectively. Significant differences in each rh $\beta$ 4GalT1 activity were evaluated using one-way ANOVA and Turkey's test. \*\*\* $p < 0.05$ . Galactosyltransferase activities against *N*-glycan of endogenous glycoproteins from posterior silk glands in silkworm larvae were detected by lectin blotting using *Ricinus communis* agglutinin I (RCA120) (b). The extraction from posterior silk glands was incubated with (+) or without (–) the rh $\beta$ 4GalT1 (N-tag or C-tag) and UDP-Galactose (UDP-Gal) as a donor for 1 h at 37 °C. Mock: the assay buffer (25 mM Tris–HCl pH 7.5, with 10 mM CaCl<sub>2</sub>, 10 mM MnCl<sub>2</sub>) instead of the extraction from posterior silk glands

glycoproteins from silkworm PSG with terminal GlcNAc and UDP-Galactose were used as a substrate and donor, respectively. We then extracted endogenous glycoproteins from PSG and incubated with UDP-Galactose together with indicated rh $\beta$ 4GalT1. As shown in Fig. 4b, lectin blotting with galactose-specific detecting RCA120, both of the C- and N-tagged rh $\beta$ 4GalT1 s exhibited galactosyltransferase activities against *N*-glycans of endogenous glycoprotein in PSG, verifying that rh $\beta$ 4GalT1 from

silkworm–BEVS provides both high yields and promising activity.

## Discussion

In the current study, the N-tagged and C-tagged rh $\beta$ 4GalT1 was expressed, purified and recovered at approximately 124 and 239  $\mu$ g, respectively, from 10 ml sera of roughly 25 infected silkworm larvae. The rh $\beta$ 4GalT1s were N-glycosylated with galactosyltransferase activities higher than that of rh $\beta$ 4GalT1 expressed and purified from mammalian cell lines. Although the final yield of the C-tagged rh $\beta$ 4GalT1 is higher than that of the N-tagged, it is better to choose N-tag for production of rh $\beta$ 4GalT1 in silkworm–BEVS if taking the purity and overall activity into considerations. Besides, it was shown that the glycoproteins extracted from silk glands of silkworm larvae could be further galactosylated by in vitro processing with the rh $\beta$ 4GalT1s. The production of these recombinant human  $\beta$ 4GalT1 s from silkworm–BEVS will be valuable tools for both therapeutics as well as biotechnology. For therapeutics, rh $\beta$ 4GalT1 s can be used along with  $\alpha$ 1,3- or  $\alpha$ 1,4-fucosyltransferases [1, 2, 6, 7] and  $\alpha$ 2,3-sialyltransferases to help create the sLe<sup>x</sup> for the ex vivo on therapeutic cells such as stem cells in order to treat disease. In biotechnology, rh $\beta$ 4GalT1 s can be used in the production of high-quality recombinant glycoproteins with mammalian-like *N*-glycans. Since there are several differences in *N*-linked glycosylation pathways between mammals and insects, proteins secreted from insect cells are paucimannosidic and generally not terminally galactosylated or sialylated [27] likely due to the extensive FDL activity which removes the terminal GlcNAc of the hybrid-type structure [25]. From this point of view, the recombinant glycoproteins generated using BEVS are presumably somewhat less stable in mammalian blood than those from native animals. Accordingly, it has been, for a long time, a problem to use recombinant glycoproteins produced from BEVS in practical medicine and clinical research. To date, various approaches have been reported to alter the *N*-linked glycan structure of insect-derived proteins into terminally sialylated complex-type *N*-glycans by transgenic expression of mammalian glycosyltransferases in insect cells [11–13]. However, the efficiency of formation of matured complex-type *N*-glycans in the insect cells is still low because the activity of mammalian  $\beta$ 4GalT1 is unstable in insect cells [14]. In addition, the negative effect of exogenously expressed  $\beta$ 4GalT1 on insect cells must be taken into account especially in transgenic insects. Here, we propose another approach in which the recombinant glycoproteins produced by BEVS are processed by in vitro *N*-glycosylation using mammalian glycosyltransferases

mass-produced and purified from insect cells or larvae. Because galactosylated *N*-glycans provide substrates for sialylation by sialyltransferases, it is desirable to mass-produce the recombinant rh $\beta$ 4GalT1 s by BEVS.

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