Galactosyl carbohydrate residues on hematopoietic stem/progenitor cells are essential for homing and engraftment to the bone marrow

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The role of carbohydrate chains in leukocyte migration to inflamed sites during inflammation and trafficking to the lymph nodes under physiological conditions has been well characterized. For example, selectins and carbohydrate ligands such as sialyl Lewis x (sLex) and 6-sulfo sLex are now known to facilitate the leukocyte-endothelium interaction, the first step in these processes¹. Accordingly, mice deficient in both E- and P-selectins, which bind to sLex², show severe impairment in leukocyte migration to inflamed sites³,⁴, while mice deficient in L-selectin, which binds to 6-sulfo sLex⁵, show compromised leukocyte trafficking to the lymph nodes⁶. In addition, insight into the roles of the selectin ligand biosynthesis pathway in leukocyte trafficking has also been obtained using mouse models. For example, mice deficient in both fucosyl transferase-IV and VII, which are responsible for the synthesis of sLex⁷ and 6-sulfo sLex⁸, show impaired selectin-dependent leukocyte recruitment and lymphocyte homing⁹. Similarly, we have shown that mice deficient in β¹,4-galactosyltransferase-1 (β¹4GalT-1) exhibit reduced inflammatory responses⁴ and delayed wound healing⁸ due to impaired leukocyte infiltration following repressed biosynthesis of endothelial selectin ligands. These results indicate that β¹4GalT-1 is also involved in the biosynthesis of selectin ligands. However, β¹4GalT-1 deficiency did not affect leukocyte trafficking to the lymph nodes⁸.

The role of carbohydrate chains in the homing and engraftment of hematopoietic stem/progenitor cells (HSPCs) to the bone marrow (BM) remains largely undefined, although previous studies have suggested the role of galactosyl and mannosyl residues¹⁰. Notably, several cell adhesion molecules, including integrin family

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members, are known to be essential in homing and engraftment\(^{11-13}\). For example, very late antigen-4 (VLA-4, \(\alpha_4\beta_1\) integrin) and VLA-5 (\(\alpha_5\beta_1\) integrin) in HSPCs bind to vascular cell adhesion molecule-1 (VCAM-1) and fibronectin in the BM, respectively\(^{14,15}\). Accordingly, antibodies against these molecules\(^{16,17}\) or genetic deletion of hematopoietic \(\beta_4\) integrin\(^{18}\) severely impair HSPC homing and engraftment. Leukocyte function-associated antigen-1 (LFA-1, \(\alpha_\beta_3\beta_2\) integrin) and lymphocyte Peyer’s patch cell adhesion molecule-1 (LPaM-1, \(\alpha_\beta_3\beta_7\) integrin) in HSPCs interact with intercellular adhesion molecule-1 (ICAM-1) and mucosal adressin cell adhesion molecule-1 (MadCAM-1) in the BM, respectively, which have also been shown to promote homing and engraftment\(^{17,19}\). Several other factors are also known to be important in HSPC homing/engraftment, including hyaluronic acid/CD44\(^{20}\), guanosine triphosphatases rac1 and rac2\(^{21}\), osteopontin\(^{22}\), sphingosine 1-phosphate receptor\(^{23}\), prograssdin E\(^{24}\), and membrane-bound SCF/c-Kit\(^{25}\).

Moreover, stromal-derived factor 1 (SDF-1, also known as CXCL12) acts as a major HSPC chemoattractant through its receptor CXCR4\(^{26}\), and mediates HSPC homing and engraftment in cooperation with LFA-1, VLA-4, and VLA-5\(^{27}\); accordingly, deficiency in SDF-1 or CXCR4 severely compromises these events\(^{27,28}\). Despite this extensive research and identification of the key players involved, the role of carbohydrate chains in the function of these cell adhesion and chemoattractant molecules is unknown. However, there is ample evidence to indicate the involvement of endothelial selectins and their carbohydrate ligands. For example, lethally irradiated mice deficient in both \(\beta_2\)- and \(\beta_3\)-selectin do not survive when transplanted with \(5 \times 10^4\) wild type BM cells, the minimum number that would otherwise be required, whereas approximately 80% of wild-type recipient mice do. Homing in P/E-selectin double-knockout recipient mice is mildly reduced, and further compromised by treatment with antibodies to VCAM-1\(^{29}\). E-selectin ligands and \(\alpha_4\) integrin expressed in HSPCs have also been shown to cooperate in HSPC homing and engraftment\(^{30}\). Overall, these results suggest that LFA-4/CXCL1 and CXCR4/SDF-1 are dominant players in HSPC homing/engraftment, with LFA-1/CXCL1 and selectin ligands/endothelial selectins playing more minor roles.

The aim of the present study was to elucidate the role of carbohydrate chains in HSPC homing/engraftment to the BM using carbohydrate-modified mice. Specifically, we transplanted HSPCs deficient in \(\beta_4\)GalT-1\(^{(\beta_4\text{GalT-1}^-)}\) in lethally irradiated mice, and determined the effect on survival of the mice, and tracked their homing ability to the BM compared to those of BM cells harboring a point mutation in UDP-N-acetylglucosamine 2-epimerase/N-acetylmannosamine kinase (GNE), a key enzyme involved in sialic acid biosynthesis. In addition, homing and engraftment of immature fetal liver HSPCs of \(\beta_4\)GalT-1\(^{−}\) mice to adult BM were examined. Overall, these results provide insights into the relative roles of galactosyl and sialyl residues compared to the more well-known roles of selectin ligands in HSPC homing and engraftment to the BM.

**Results**

**HSPC population and colony-forming activities of BM cells.** Hematopoietic cells, including lymphocytes, neutrophils, monocytes, and red blood cells, are present in the peripheral blood of \(\beta_4\)GalT-1\(^{−}\) mice, although leukocytosis and mild anemia are observed\(^{31}\). The T/B cell ratio and CD4/CD8 cell ratio were normal in the spleen and thymus of these mice, respectively (data not shown). To examine the hematopoietic stem cell (HSC) population in BM cells, we performed flow cytometry analysis using various cell surface markers (Fig. 1A). The number of PHSCs\(^{32}\), defined as lineage (Ter-119, B220, CD3e, CD4, CD8a, Gr-1, and CD11b, IL-7R)-negative, Sca-1\(^−\), c-Kit\(^−\), Flk-2\(^−\), CD150\(^−\), CD34\(^−\)/low was comparable between \(\beta_4\)GalT-1\(^{+}\) and \(\beta_4\)GalT-1\(^{−}\) mice (Fig. 1B). In addition, colony formation by BM cells from \(\beta_4\)GalT-1\(^{−}\) mice was comparable to that of BM cells from \(\beta_4\)GalT-1\(^{+}\) mice (Fig. 1C), consistent with our previous reports showing that colony formation in the presence of granulocyte-colony stimulating factor or IL-3 was comparable between these mice\(^{33}\). These results imply that the hematopoietic population and colony-forming activity of HSPCs are not affected by \(\beta_4\) GalT-1 deficiency.

To further examine HSPC populations in BM cells, as illustrated in Fig. 2A\(^32\), we performed flow cytometry analysis according to the gating scheme (Fig. S1), as described previously\(^{34}\). The numbers of multipotent progenitor (MPP), common myeloid progenitor (CMP), common lymphoid progenitor (CLP), granulocyte-macrophage progenitor (GMP), and megakaryocyte-erythroid progenitor (MEP) are shown in Fig. 2B. The number of MPP increased 3-fold in \(\beta_4\)GalT-1\(^{−}\) BM cells compared to \(\beta_4\)GalT-1\(^{+}\) BM cells. Although the numbers of CMP and GMP were equivalent between the genotypes, the numbers of MEP and CLP decreased by approximately half in \(\beta_4\)GalT-1\(^{−}\) BM cells compared to \(\beta_4\)GalT-1\(^{+}\) BM cells. These results suggest that HSC differentiation was disturbed to some extent downstream of MPP.

**Impaired hematopoietic reconstitution by \(\beta_4\)GalT-1\(^{−}\) BM cells.** We previously reported that \(\beta_4\)GalT-1 is involved in the biosynthesis of selectin ligands, indicated by the impairment of inflammatory responses and wound healing in \(\beta_4\)GalT-1\(^{−}\) mice\(^35,36\). As the carbohydrate chains in HSPCs may also regulate homing and engraftment to the BM, we first examined the extent of hematopoietic reconstitution in lethally irradiated mice (Table 1), which generally die within 10 days unless transplanted with functional BM cells. Remarkably, the lethally irradiated mice intravenously transplanted with \(\beta_4\)GalT-1\(^{−}\) BM cells also died within 10 days, whereas those transplanted with \(\beta_4\)GalT-1\(^{+}\) cells survived for more than 60 days. Conversely, lethally irradiated \(\beta_4\)GalT-1\(^{−}\) mice transplanted with BM cells from wild type C57BL/6 mice also survived for more than 60 days, while lethally irradiated \(\beta_4\)GalT-1\(^{−}\) mice transplanted with \(\beta_4\)GalT-1\(^{+}\) BM cells died within 10 days. Collectively, these results indicated that \(\beta_4\)GalT-1 activity in transplanted BM cells, but not in recipient mice, is critical for hematopoietic reconstitution. It is known that BM cells engraft to the BM more efficiently when they are directly transplanted into the BM cavity using the intra-BM transplantation (IBM-BMT) method\(^{37}\). Similar to the intravenous BM transplantation (IV-BMT), lethally irradiated wild type mice transplanted with BM cells from \(\beta_4\)GalT-1\(^{−}\) mice by IBM-BMT died within 10 days, indicating that \(\beta_4\)GalT-1\(^{−}\) BM cells could not reconstitute hematopoiesis even when the IBM-BMT method was used.
We then transplanted a mixture of β4GalT-1−/− and β4GalT-1+/− BM cells to ensure survival (Table 2), and the former cells were labeled with green fluorescent protein (GFP) to enable tracking. Strikingly, GFP-positive cells were rarely detected in the spleen, thymus, and peripheral blood of recipient mice 9 weeks after transplantation even when 90% of the transplanted cells were β4GalT-1−/− (Table 2, Exp. 2), whereas more than 90% of the 

**Figure 1.** Hematopoietic stem cell (HSC) population and colony-forming activity of bone marrow (BM) cells. (A) Flow cytometry analysis of BM cells using various antibodies against indicated cell surface markers. pHSCs were defined as lineage−, Sca-1+, c-Kit+, Flk-2−, CD150+, CD34−/low. (B) The number of HSCs examined by flow cytometry (A) per 1 × 10⁶ BM cells obtained from β4GalT-1+/− (ht, n = 4) and β4GalT-1−/− mice (mt, n = 4). (C) Colony-forming ratios of BM cells obtained from β4GalT-1+/− (ht, n = 9) and β4GalT-1−/− mice (mt, n = 9). Error bars indicate the S.D.

**Figure 2.** Hematopoietic stem/progenitor cell (HSPC) differentiation. (A) Illustration of HSPC differentiation to terminal blood cells32. LT-HSC; long term-HSC, ST-HSC; short term-HSC, MPP; multipotent progenitor, CMP; common lymphoid progenitor, GMP; granulocyte-macrophage progenitor, MEP; megakaryocyte-erythroid progenitor. (B) The numbers of MPP, CMP, CLP, GMP and MEP examined by flow cytometry per 1 × 10⁶ BM cells obtained from β4GalT-1+/− (ht, n = 6) and β4GalT-1−/− mice (mt, n = 6). Error bars indicate S.D. *p < 0.05. MPP; Lin−, IL-7R−, c-kit+, Sca-1+, CD34+, Flk-2−, CMP; Lin−, IL-7R−, c-kit+, Sca-1−, CD34+, FcgRlow, CLP; Lin−, IL-7R+, Flk-2−, GMP; Lin−, IL-7R−, c-kit+, Sca-1−, CD34+, FcgR+, MEP; Lin−, IL-7R−, c-kit+, Sca-1−, CD34+, FcgRlow, as described in the gating scheme (Fig. S1).
cells in these tissues were GFP-positive when only GFP-positive wild type cells were transplanted (Table 2, Exp. 4). These results suggest that \( \beta^4 \text{GalT-1}^-/^- \) BM cells are impaired in homing and engraftment after transplantation.

**Colony formation by transplanted \( \beta^4 \text{GalT-1}^-/^- \) BM cells.** To examine the homing ability of the transplanted HSPCs, splenocytes and BM cells were collected from the recipient mice at 3 h and 24 h after IV-BMT, and analyzed using a colony formation assay. Colony-forming transplanted \( \beta^4 \text{GalT-1}^-/^- \) cells were approximately 0.5- and 0.3-fold as abundant in recipient splenocytes as in the transplanted \( \beta^4 \text{GalT-1}^+/^- \) cells at 3 h and 24 h after IV-BMT, respectively. Similarly, the colony-forming ratio of transplanted \( \beta^4 \text{GalT-1}^-/^- \) cells increased from 1.5% to 6% between 3 h and 24 h, but that of transplanted \( \beta^4 \text{GalT-1}^-/^- \) cells was less than 0.5% (Fig. 3A,B). These results indicated that \( \beta^4 \text{GalT-1} \) deficiency severely impaired the homing ability of transplanted HSPCs to the BM.

In these experiments, C57BL/6 mice were used as the recipient, although the donor cells were harvested from \( \beta^4 \text{GalT-1}^-/^- \) or \( \beta^4 \text{GalT-1}^+/^- \) mice on a mixed 129/Sv and C57BL/6 genetic background. Nevertheless,

**Table 1. Survival ratio of recipient mice after IV-BMT and IBM-BMT of \( \beta^4 \text{GalT-1}^-/^- \) deficient BMCs.** *Days after BMT.

| BMT methods | Donor BMCs | Recipient mice | Survival ratio (alive/total) |
|-------------|------------|----------------|-----------------------------|
| IV-BMT      | \( \beta^4 \text{GalT-1}^-/^- \) C57BL/6 (wt) | 3/3 | 3/3 | 0/3 | 0/3 | 0/3 |
|             | \( \beta^4 \text{GalT-1}^+/^- \) C57BL/6 (wt) | 4/4 | 4/4 | 4/4 | 4/4 |
|             | \( \beta^4 \text{GalT-1}^-/^- \) C57BL/6 (wt) | 2/2 | 2/2 | 0/2 | 0/2 | 0/2 |
|             | \( \beta^4 \text{GalT-1}^-/^- \) C57BL/6 (wt) | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 |
|             | no C57BL/6 (wt) | 3/3 | 3/3 | 0/3 | 0/3 | 0/3 |
| IBM-BMT     | \( \beta^4 \text{GalT-1}^-/^- \) C57BL/6 (wt) | 4/4 | 4/4 | 0/4 | 0/4 | 0/4 |
|             | \( \beta^4 \text{GalT-1}^+/^- \) C57BL/6 (wt) | 5/5 | 5/5 | 5/5 | 5/5 | 5/5 |
|             | no C57BL/6 (wt) | 4/4 | 3/4 | 0/4 | 0/4 | 0/4 |
| IV-BMT      | \( \beta^4 \text{GalT-1}^-/^- \) NOD/SCID | 4/4 | 4/4 | 0/4 | 0/4 | 0/4 |
|             | \( \beta^4 \text{GalT-1}^+/^- \) NOD/SCID | 4/4 | 4/4 | 0/4 | 0/4 | 0/4 |
|             | no NOD/SCID | 2/2 | 1/2 | 0/2 | 0/2 | 0/2 |

**Table 2. Engraftment of mixed donor bone marrow cells (BMCs).** Exp 1: \( 1.0 \times 10^7 \) mt/GFP BMCs + \( 1.0 \times 10^7 \) ht BMCs. Exp 2: \( 1.8 \times 10^7 \) mt/GFP BMCs + \( 0.2 \times 10^7 \) ht BMCs. Exp 3: \( 0.4 \times 10^7 \) ht BMCs. Exp 4: \( 2.0 \times 10^7 \) wt/GFP BMCs. 9 weeks after transplantation.

| Tissues       | Ratio of GFP-positive cells (%) |
|---------------|--------------------------------|
|               | Exp 1 | Exp 2 | Exp 3 | Exp 4 |
| Peripheral blood | 0.00  | 0.00  | 0.00  | 92.90  |
| Spleen        | 0.04  | 0.46  | 0.00  | 95.05  |
| Thymus        | 0.01  | 0.00  | 0.00  | 98.60  |

**Figure 3. Homing of transplanted \( \beta^4 \text{GalT-1}^-/^- \) BM cells.** (A,B) Colony-forming ratios of splenocytes (A) and BM cells (B) obtained from wild-type recipient mice transplanted with \( \beta^4 \text{GalT-1}^+/^- \) (n = 6–8) and \( \beta^4 \text{GalT-1}^-/^- \) BM cells (n = 6–8), as measured 3 h and 24 h after transplantation. (C) Colony-forming ratios of splenocytes and BM cells obtained from NOD/SCID recipient mice transplanted with \( \beta^4 \text{GalT-1}^+/^- \) (n = 4) and \( \beta^4 \text{GalT-1}^-/^- \) BM cells (n = 4), as measured 24 h after transplantation. Blue bars, \( \beta^4 \text{GalT-1}^+/^- \) cells; red bars, \( \beta^4 \text{GalT-1}^-/^- \) cells. Error bars indicate the S.D. *p < 0.05, **p < 0.01, and ***p < 0.001.
βGFP-labeled BM cells in the recipient femur using fluorescence microscopy (Fig. 4A). Abundant GFP-positive cells specifically recognize Gal residues in HSPCs, lineage-negative BM cells were analyzed by a lectin blot using RCA120 and ECA, which highlight 4GalT-1 development, hematopoiesis mainly proceeds in the fetal liver. We also examined homing and engraftment of immature HSPCs in the fetal liver of embryonic stage 14.5 (E14.5). Lethally irradiated wild-type mice received transplantations of fetal liver cells. Approximately 0.2-fold survival ratio of recipient mice after transfer of β4GalT-1-deficient fetal liver cells. *Days after transplantation of fetal liver cells.

| Donor fetal liver cells | Recipient mice | Survival ratio (alive/total) |
|------------------------|----------------|-----------------------------|
|                        |                | 0*  | 5*  | 10* | 15* |
| β4GalT-1 −/−           | C57BL/6 (wt)   | 5/5 | 5/5 | 0/5 | 0/5 |
| β4GalT-1 +/+           | C57BL/6 (wt)   | 7/7 | 7/7 | 6/7 | 6/7 |
| no                     | C57BL/6 (wt)   | 2/2 | 2/2 | 0/2 | 0/2 |

Figure 4. In situ observation of transplanted donor-derived bone marrow (BM) cells. (A) Frozen sections of the femur of wild-type recipient mice 24 h after BMT of GFP-labeled β4GalT-1+/− (upper, ht) and β4GalT-1−/− (lower, mt) BM cells. Bright field (left), DAPI (middle), GFP (right). Scale bar, 100 μm. (B) The number of GFP-positive cells per mm² in (A) from β4GalT-1+/− (ht, n = 3) and β4GalT-1−/− (mt, n = 3) mice. *p < 0.05.

Impaired hematopoietic reconstitution by β4GalT-1−/− fetal liver cells. During embryonic development, hematopoiesis mainly proceeds in the fetal liver. We also examined homing and engraftment of immature HSPCs in the fetal liver of embryonic stage 14.5 (E14.5). Lethally irradiated wild-type mice transplanted with β4GalT-1−/− fetal liver cells also died within 10 days, whereas those transplanted with β4GalT-1+/− cells survived for more than 15 days (Table 3). Furthermore, the colony-forming ratio of transplanted β4GalT-1−/− fetal liver cells were approximately 0.22-fold as abundant in recipient BM as they were in donor BM cells. These results indicate that the homing efficiency of β4GalT-1−/− BM cells was not due to immunological rejection.

Table 3. Survival ratio of recipient mice after transfer of β4GalT-1-deficient fetal liver cells. *Days after transplantation of fetal liver cells.

Lectin blot of lineage-negative β4GalT-1−/− BM cells. To examine the galactosyl carbohydrate residues in HSPCs, lineage-negative BM cells were analyzed by a lectin blot using RCA120 and ECA, which specifically recognize Galβ1-4GlcNAc. Since most proteins are sialylated at the non-reducing carbohydrate terminus, RCA120- and ECA-reactive bands were rarely detected in both β4GalT-1+/− and β4GalT-1−/− lineage-negative BM cells. However, digestion of cells with sialidase generated strong and smeary RCA120- and ECA-reactive bands between 100 and 200 kDa in β4GalT-1+/− lineage-negative cells, but not in β4GalT-1−/− lineage-negative cells (Fig. 6). This binding specificity was confirmed by the addition of lactose to specifically block RCA120 binding to Galβ1-4GlcNAc. Considering that the galactosyl residues in high-molecular-weight glycoproteins were lost in lineage-negative BM cells from β4GalT-1−/− mice, these results suggested that they significantly promoted homing and engraftment.
Figure 5. Homing of transplanted β4GalT-1−/− fetal liver cells. (A) The number of colony-forming cells of BM cells obtained from wild-type recipient mice transplanted with β4GalT-1+/− (ht, n = 5) and β4GalT-1−/− (mt, n = 6) fetal liver cells, as measured 24 h after transplantation. (B) The number of colony-forming cells of fetal livers obtained from β4GalT-1+/− (ht, n = 6) and β4GalT-1−/− (mt, n = 8) E14.5 embryos. *p < 0.05.

Figure 6. Lectin blot of lineage-negative bone marrow (BM) cells. Lanes 1, 2: lineage-negative BM cells (2.6 × 10⁴ cells) stained with Colloidal Gold. Lanes 3–12: lineage-negative BM cells (2.6 × 10⁴ cells) reacted with RCA-120 (lanes 3–8) or ECA (lanes 9–12). Lanes 5–6 and 11–12 were digested with sialidase, while lanes 7–8 were digested with sialidase and blocked with lactose. Lanes 1, 3, 5, 7, 9, 11 are β4GalT-1+/− cells (ht), while lanes 2, 4, 6, 8, 10, 12 are β4GalT-1−/− cells (mt). Lanes 1–12 were from the same gel. A representative lectin blot of three mice per genotype is shown.
Colony formation by transplanted Gne (V572L) BM cells. Sialic acids are well known to modify non-reducing terminal carbohydrates, including the galactosyl residues in glycoproteins and glycolipids. Notably, the lack of GNE, a key enzyme in sialic acid biosynthesis, is an embryonic lethal mutation37. However, mice with a V572L point mutation in GNE survive for several months, but suffer from a nephrotic-like syndrome because of severe hyposialylation of podocyte glycoproteins38. Colony formation by BM cells obtained from such mice was comparable with that of wild-type cells, suggesting that BM cell proliferation and differentiation were normal (Fig. 7A). However, colony-forming transplanted Gne (V572L) cells were about 60% as abundant in recipient BM cells as transplanted wild type cells (Fig. 7B).

The survival of lethally irradiated wild type mice clearly depended on the number of Gne (V572L) BM cells transplanted (Fig. 7C). For example, survival was lower in mice transplanted with $1 \times 10^5$ Gne (V572L) BM cells (60%) than that in mice transplanted with the same number of wild-type cells (90%). The difference in survival was even larger (0% vs 80%) when $5 \times 10^5$ cells were transplanted. However, all mice died within 10 days.
when 2 × 10^6 cells were transplanted. Collectively, these data indicated that homing was mildly impaired in Gne (V572L) BM cells, suggesting that sialyl residues ligated to Galβ1-4GlcNAc also played a role in homing and engraftment, which was diminished by the Gne (V572L) point mutation.

**Discussion**

Colony-forming activity of HSPCs and the number of pHSC (short-term [ST]-HSC) were not different between β4GalT-1−/− BM cells and β4GalT-1+/− BM cells. Although the number of functional HSC can be quantitatively examined using in vivo limiting dilution assay by BMT of serial diluted BM cells, it was difficult to estimate its number in β4GalT-1−/− BM cells because homing and engraftment to the BM was severely impaired. In the differentiation pathway of HSC, the number of MPP was higher and those of MEP and CLP were lower in β4GalT-1−/− BM cells than in β4GalT-1+/− BM cells. These results suggest that HSC differentiation was disturbed to some extent downstream of MPP. Although the number of MPP increased in β4GalT-1−/− BM cells, colony-forming activity of HSPCs was comparable between β4GalT-1−/− and β4GalT-1+/− BM cells. Therefore, the disturbed differentiation downstream of MPP did not seem to have an influence on our findings that homing and engraftment of β4GalT-1−/− HSPCs after BMT were severely impaired. Colony-forming activity of HSPCs from β4GalT-1−/− fetal liver cells slightly increased. These results indicated that the number, proliferation activity, or both of fetal liver HSPCs in β4GalT-1−/− mice were slightly enhanced, not reduced, which also did not seem to affect the homing and engraftment of transferred β4GalT-1−/− fetal liver HSPCs.

Several cell adhesion systems such as VLA-4/VCAM-1 and CXCR4/SDF-1 play a major role in HSPC homing and engraftment to the BM, while other systems such as LFA-1/ICAM-1 and sLe^x endothelial selectins play a supportive role. However, these systems interact to ensure efficient homing and engraftment. As β4GalT-1−/− mice have compromised biosynthesis of selectin ligands and reduced inflammatory reactions, the contribution of sLe^x/endothelial selectins to HSPC homing and engraftment may also be impaired. However, lethally irradiated mice deficient in both P- and E-selectin survive when transplanted with at least 5 × 10^6 BM cells. Furthermore, homing in these mice deficient in both P- and E-selectin is reduced by approximately 50% compared with that in wild-type recipient mice. These results strongly suggest that the defect in the sLe^x/endothelial selectins system alone cannot explain the loss of homing and engraftment from β4GalT-1−/− BM cells. Furthermore, the present results imply that a novel cell adhesion system based on galactosyl carbohydrates promotes HSPC homing and engraftment following transplantation. On the other hand, the abundance of colony-forming transplanted Gne (V572L) cells in recipient tissues, as well as the survival of lethally irradiated recipient mice transplanted with such cells, was highly similar to that previously observed in P/E-selectin double-knockout mice. These results suggest that the impaired homing and engraftment of Gne (V572L) cells is due to the defect in the sLe^x/endothelial selectins system. Alternatively, the mild phenotype of Gne (V572L) mice may be attributed to reduced, but not abolished, GNE activity.

During the latter half of mouse embryonic development until birth, hematopoiesis mainly occurs in the fetal liver. Neonatal migration of HSPCs from the fetal liver to the adult BM seems to be normal in β4GalT-1−/− mice, because colony-forming activity of HSPCs and ST-HSC populations in the adult BM were comparable between β4GalT-1−/− and β4GalT-1+/− mice. However, homing and engraftment of immature fetal liver HSPCs of β4GalT-1−/− mice to the adult recipient BM was impaired similar to that in the BM HSPCs of β4GalT-1−/− mice. These results suggest that fetal liver HSPCs and adult BM HSPCs use the similar galactosyl residues in homing and engraftment to the BM.

Lectin blots show that galactosyl residues in high-molecular-weight glycoproteins were lost in lineage-negative BM cells from β4GalT-1−/− mice. Accordingly, these glycoproteins are good candidates as critical regulators of HSPC homing and engraftment. We noted that integrins such as α4, α5, β3, and β1 are larger than 100 kDa and contain many possible glycosylation sites, some of which are actually N-glycosylated (Glycoprotein Database, http://gcggdb.jp/rcmg/gpdb/index.action). Therefore, it is possible that the function of VLA-4 (α4β1 integrin), VLA-5 (α5β1 integrin), or LFA-1 (αLβ2 integrin) in HSPC homing and engraftment is compromised by β4GalT-1 and GNE deficiency. Another possibility is that β4GalT-1 and GNE deficiency may disrupt the function of unknown carbohydrate ligands that regulate homing and engraftment. Thus, further studies are necessary to fully elucidate the role of carbohydrate residues in HSPC homing and engraftment.

In clinical applications of BMT, especially in cord blood transplantation, it is essential to enhance the efficiency of HSPC homing and engraftment. The present study suggests the possibility of modifying carbohydrate structures on the surface of HSPCs to improve their homing and engraftment to the BM in clinical application. Indeed, a recent study demonstrated that the ex vivo fucosylation of cord blood cells improved their homing abilities, leading to faster neutrophil and platelet engraftments. Accordingly, it might be possible that enforced galactosylation and sialylation of HSPCs would also improve their homing and engraftment to the BM.

In conclusion, we have demonstrated that β4GalT-1 activity in donor BM cells, but not recipient mice, is critical for hematopoietic reconstitution and homing/engraftment to the BM after transplantation. However, BM cells from Gne (V572L) mice only showed relatively mild impairment. The deficiency of BM cells in sLe^x/endothelial selectins system might explain the defect of BM cells from Gne (V572L) mice, but cannot explain the defect of β4GalT-1−/− BM cells. Collectively, these data suggest that a novel cell adhesion system containing galactosyl or sialyl residues or both may promote homing and engrafting of HSPCs to the BM.

**Materials and Methods**

**Mice.** β4GalT-1−/− mice on a mixed 129/Sv and C57BL/6 genetic background, and mice with a V572L point mutation in Gne [Gne (V572L) mice] on a C57BL/6 background were described previously.[34,38] To produce GFP-labeled BM cells, these mice were crossed with CAGGFP mice on a C57BL/6 background, which were kindly provided by Dr. Okabe at Osaka University[41]. NOD/SCID mice and pseudo-pregnant ICR mice were purchased from Charles River Japan and CLEA Japan, Inc., respectively. Animal experiments were conducted in
Preparation of BM cells. BM cells were harvested by aseptically flushing the femur and tibia using a 22-gauge needle with Dulbecco's modified Eagle's medium (DMEM, Life Technologies, Grand Island, NY, USA) containing 5% fetal calf serum (FCS). The obtained cell suspension was filtered through a 70-μm mesh, treated with 140 mM NH₄Cl in 17 mM Tris-HCl (pH 7.2) buffer for 5 min to lyse red blood cells, washed, and suspended in DMEM with 5% FCS for BMT and the colony formation assay.

Preparation of fetal liver cells. Oocytes from β4GalT-1−/− mice with the CAGGFP gene in homozygotes were fertilized in vitro by sperms from β4GalT-1−/− mice and fertilized two-cell stage eggs were transferred to the oviduct of pseudo-pregnant ICR females. Embryos were collected at E14.5 and fetal livers were prepared by crushing using a plunger on 70-μm strainer in Ca²⁺- and Mg²⁺-free phosphate-buffered saline (PBS) containing 3% FCS. Fetal liver cells were collected after centrifugation, suspended in Hank's solution, and passed through a 40-μm strainer for transplantation and colony formation assay.

Flow cytometry. Flow cytometry was performed using a FACS Aria II and Aria IIIu cell sorter (BD Biosciences, Franklin Lakes, NJ, USA) and analyzed using FlowJo software (Tree Star). BM cells were collected from the bilateral femur and tibia by flushing using a 22-gauge needle in PBS containing 3% FCS. Cells were passed through a 70-μm strainer and treated with 140 mM NH₄Cl, 17 mM Tris-HCl (pH 7.2) buffer for 5 min to lyse red blood cells. After washing in RPMI-1640 (Life Technologies) three times, the cells were passed through a 40-μm strainer to exclude cell aggregates before analysis. For BM cell analysis, the cells were stained with combinations of antibodies against the following surface markers: c-Kit (clone 2B8; Thermo Fisher Scientific, Waltham, MA, USA), Sca-1 (clone D7, BD Biosciences), Flk2 (clone A2F10, Thermo Fisher Scientific), CD150 (clone TC15-12F12.2, BioLegend, San Diego, CA, USA), CD34 (clone RAM34, Thermo Fisher Scientific), and Fc blocker CD16/32 (clone 93, Thermo Fisher Scientific). The lineage markers used for Fig. 1 included Ter-119 (clone TER-119, BioLegend), B220 (clone RA3-6B2, BioLegend), CD3 (clone 145-2C11, BioLegend), and IL-7Rα (clone A7R34, Thermo Fisher Scientific). The lineage markers (BioLegend 133313) used for Fig. 2 included CD3 (clone 17A2), Gr-1 (clone RB6-8C5, BioLegend), CD11b (clone M1/70, BD Biosciences), and CD4 (clone TC15-12F12.2, BioLegend). BM cells were prepared, fixed in 4% paraformaldehyde for 5 h, and equilibrated in 30% sucrose/PBS. Fixed bone samples were embedded in SCEM-L1 (SECTION-LAB, Hiroshima, Japan) and frozen in cooled hexane with liquid nitrogen gas (N₂). Cryostat sections (8 mm thick) were generated using Kawamoto's technique of Japan, and were approved by the Committee on Animal Experimentation at Kanazawa University and Kyoto University, Japan.

Preparation of colony-negative cells. BM cells (1 × 10⁶ cells/mL) were prepared in PBS containing 3% FCS, incubated on ice for 15 min with 2.5 μg anti-mouse CD16/CD32 (Fc Block, BD Biosciences) per 10⁷ cells, and then incubated on ice for 15 min with Biotin Mouse Lineage Depletion Cocktail (BD Biosciences) consisting of biotinylated monoclonal anti-mouse CD3e, CD11b, CD45R/B220, Ly-6G, Ly-6C, and TER-119. Subsequently, the mixture was washed with 10 volumes of PBS containing 0.5% bovine serum albumin and 2 mM ethylenediaminetetraacetic acid, and centrifuged. The resulting cell pellets were mixed with Streptavidin Particles...
Lectin blot. Lineage-negative BM cells (2.6 × 10^5 cells) were suspended in NuPAGE LDS Sample Buffer (Thermo Fisher Scientific), electrophoresed on SuperSep Ace 5–12% precast gels (Wako Pure Chemical Industries, Tokyo, Japan), and transferred to polyvinylidene difluoride membranes by electroblotting (Bio-Rad, Hercules, CA, USA). Lectin blotting was performed as described previously. In brief, the membranes were washed in blocking buffer (10 mM Tris-HCl pH 7.4, 0.15 M NaCl, and 0.05% Tween 20), and immersed for 1 h with gentle shaking in blocking buffer containing 2 μg/mL biotinylated RAC120 (Seikagaku Corporation, Tokyo, Japan) or 5 μg/mL biotinylated ECA (Seikagaku Corporation). The membranes were then washed again, immersed for 30 min with gentle shaking in blocking buffer containing 1:200 avidin conjugated to horseradish peroxidase, washed, incubated with ImmunoStar (Wako Pure Chemical Industries), and visualized with ChemiStage (Kurabo, Osaka, Japan). Total proteins were also stained with Colloidal Gold for comparison.

In the RAC120 blots, certain membranes were treated with 0.05 U/mL sialidase (Roche, Mannheim, Germany) at 37 °C for 30 min before probing with lectins, or with 0.2 M lactose during probing with lectins.

Statistical analysis. Differences with two-sided p < 0.05 were deemed statistically significant, as evaluated using the Student’s t-test. Results are reported as mean ± standard deviation (S.D.).

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**Author Contributions**

S.T., R.Y., N.H., K.S. and M.M.: Collection and/or assembly of data S.O., C.N.: Data analysis and interpretation M.A.: Conception and design, financial support, manuscript writing, final approval of manuscript. All authors reviewed the manuscript.

**Additional Information**

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