# Chemoselection of Allogeneic HSC After Murine Neonatal Transplantation Without Myeloablation or Post-transplant Immunosuppression

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The feasibility of allogeneic transplantation, without myeloablation or post-transplant immunosuppression, was tested using in vivo chemoselection of allogeneic hematopoietic stem cells (HSCs) after transduction with a novel tricistronic lentiviral vector (MGMT<sup>P140K</sup>-2A-GFP-IRES-TK (MAGIT)). This vector contains P140K-O<sup>6</sup>-methylguanine-methyltransferase (MGMT<sup>P140K</sup>), HSV-thymidine kinase (TK<sup>HSV</sup>), and enhanced green fluorescent protein (eGFP) enabling (i) in vivo chemoselection of HSC by conferring resistance to benzylguanine (BG), an inhibitor of endogenous MGMT, and to chloroethylating agents such as 1,3-bis(2-chloroethyl)nitrosourea (BCNU) and, (ii) depletion of proliferating cells such as malignant clones or transduced donor T cells mediating graft versus host disease (GVHD), by expression of the suicide gene TK<sup>HSV</sup> and Ganciclovir (GCV) administration. Nonmyeloablative transplantation of transduced, syngeneic, lineage-depleted (Lin<sup>-</sup>) BM in neonates resulted in 0.67% GFP<sup>+</sup> mononuclear cells in peripheral blood. BG/ BCNU chemoselection, 4 and 8 weeks post-transplant, produced 50-fold donor cell enrichment. Transplantation and chemoselection of major histocompatibility complex (MHC)-mismatched MAGIT-transduced Lin-BM also produced similar expansion for >40 weeks. The efficacy of this allotransplant approach was validated in Hbb<sup>th3</sup> heterozygous mice by correction of β-thalassemia intermedia, without toxicity or GVHD. Negative selection, by administration of GCV resulted in donor cell depletion without graft ablation, as re-expansion of donor cells was achieved with BG/BCNU treatment. These studies show promise for developing non-ablative allotransplant approaches using in vivo positive/negative selection.

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### **INTRODUCTION**

Sibling or matched unrelated allogeneic hematopoietic stem cell (HSC) transplantation remain the only curative therapy for

many hereditary disorders.<sup>1</sup> However, the toxicity of myeloablative preparative regimens, risks of graft versus host disease (GVHD), infectious complications of immunosuppression, and limited availability of suitable donors, restrict application of this approach. While early transplantation could reduce or abrogate the pathogenic consequences of many genetic disorders, myeloablative transplantation approaches, especially very early in life, have been associated not only with morbidity and mortality, but also with subsequent abnormal development and growth.<sup>2,3</sup> Thus, approaches to diminish these risks have focused on reducing the intensity and toxicity of preparatory and immunosuppressive regimens, without compromising engraftment or increasing the incidence of GVHD.

One approach to addressing the risks of transplantation has been to genetically modify donor cell populations to enable positive selection and expansion of donor HSC, or negative selection of donor T cells causing GVHD. Treatment of GVHD caused by transduced donor T cells using HSV thymidine kinase suicide gene (TK<sup>HSV</sup>)/Ganciclovir (GCV)-mediated negative selection has been validated in a number of recent clinical studies.<sup>4</sup> Introduction of a drug resistance gene in HSC followed by *in vivo* chemoselection has increased donor chimerism and may enable reduction of the intensity, and therefore toxicity, of conditioning regimens. In early studies, limitations of positive selection of HSC by *in vivo* chemotherapy included the requirement for ongoing drug administration with associated cumulative toxicity, and the unanticipated transformation and autonomous proliferation of selected cells.<sup>5-7</sup>

The observation that transfer of the O<sup>6</sup>-alkylguanine-DNA alkyltransferase gene into mammalian cells decreased sensitivity to 1,3-bis(2-chloroethyl)nitrosourea (BCNU), a well-established HSC toxin, suggested that alkyltransferases such as MGMT could be used for *in vivo* chemoselection.<sup>8,9</sup> Identification of O<sup>6</sup>-Benzylguanine (BG)<sup>10</sup> as an inhibitor of endogenous MGMT, and the derivation of BG-resistant forms of MGMT (P140K and G156A)<sup>11,12</sup> provided dramatic improvements in the durability of this *in vivo* chemoselection strategy. The MGMT<sup>P140K</sup> mutant repair enzyme exhibits 1,000-fold resistance to alkylators and nitrosoureas compared with the wild-type MGMT enzyme following treatment with BG.<sup>13-15</sup> When BG is administered to inhibit endogenous MGMT activity followed by delivery of the

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nitrosourea BCNU, or alkylating agents such as temozolomide, cells in the bone marrow not expressing MGMT<sup>P140K</sup> are eliminated. Unlike previous *in vivo* selection systems, genetic modification of HSC by transduction with the MGMT<sup>P140K</sup> gene results in *in vivo* chemoselection at the HSC level and stable donor chimerism, even after discontinuation of drug treatment.<sup>11-20</sup>

Using this approach, enrichment of MGMT<sup>P140K</sup>-expressing HSC has been successfully demonstrated in congenic, myeloablated adult mice,<sup>15</sup> human nonobese diabetic/severe combined immunodeficiency repopulating cells,<sup>14,21</sup> and with allogeneic transplantation requiring post-transplant immunosuppression in a canine model.<sup>18-20</sup> Recently, this approach has also been evaluated in nonhuman primates.<sup>22,23</sup> In these studies transplantation was predicated on myeloablation, the use of immuno-incompetent hosts, or on administration of standard immunosuppressive regimens, respectively.

Despite the promise of this in vivo HSC selection approach, concerns remain regarding the proliferative stress placed on small numbers of MGMTP140K modified repopulating HSC clones following successive cycles of chemoselection.<sup>19,20,24</sup> In addition, there are well-established risks of genotoxicity associated with integrating gene transfer vectors and subsequent autonomous clonal proliferation.<sup>25-27</sup> Thus, incorporation of a negative selection strategy to enable control or elimination of malignant clones, should they emerge, would be desirable. TK<sup>HSV</sup>-expressing cells metabolize GCVinto its active triphosphate form resulting in cell death.<sup>4</sup> GCV is more toxic to proliferating cells<sup>28</sup> than to quiescent populations such as HSC, suggesting that administration of GCV could preferentially eliminate autonomously replicating clones arising from insertional mutagenesis and also proliferative alloreactive T cells causing GVHD. In the clinical setting, infusion of donor T cells modified to express the TKHSV suicide gene to improve immune reconstitution and subsequent control of GVHD by GCV has been validated in several studies (reviewed in ref. 4).

To initially test the lentivirally based MGMT<sup>P140K</sup>/TK<sup>HSV</sup> "positive/negative" selection strategy, a neonatal transplant model was established. Transplantation at days 1–2 of life, before maturation of alloreactive T cells, enables studies of engraftment and mechanisms of tolerance induction during immune ontogeny while reducing confounding variables of host-mediated immune responses and graft rejection.<sup>29,30</sup>

Donor HSC engraftment and in vivo chemoselection was assessed after HSC transduction with a tricistronic lentiviral vector (MGMT<sup>P140K</sup>-2A-GFP-IRES-TK (MAGIT)) that incorporates MGMT<sup>P140K</sup> for positive selection, enhanced green fluorescent protein (eGFP) for flow cytometric analysis, and the sr39TK<sup>HSV</sup> suicide gene for enhanced negative selection. Transduction and transplantation of either syngeneic or allogeneic HSC into neonates preconditioned with a busulfan (BU)-based non-toxic, non-ablative regimen, produced successful engraftment with up to 50-fold expansion of donor cells after in vivo chemoselection. Neither GVHD nor autonomous proliferation of transduced clones was observed. Importantly, negative selection by administration of GCV enabled depletion of committed progenitors and circulating cells, without ablation of engrafted, transduced HSC. The promise of this approach was validated by correction of anemia and RBC morphology after transplantation and chemoselection in a murine model of  $\beta$ -thalassemia intermedia.

### RESULTS

# Efficient *in vitro* chemoselection of 293T cells and transduction of lineage-depleted BM with MAGIT lentivirus

The expression efficiency of the tricistronic MAGIT lentivirus (**Figure 1a**) was first tested by transduction of 293T cells. After transduction and culture, >90% of 293T cells co-expressed eGFP and intracellular MGMT<sup>P140K</sup> (**Figure 1b**) and were viable, with similar efficiency to previously reported MGMT<sup>P140K</sup>-mediated selection,<sup>31</sup> after one cycle of BG/BCNU treatment (**Figure 1c**). Control, untransduced cells were largely eliminated after BG/BCNU treatment (**Figure 1c**). Negative selection of 293T cells by co-culture with GCV confirmed the functionality of the sr39TK<sup>HSV</sup> cassette (**Figure 1d**). Efficient transduction of lineage-depleted primary bone marrow (Lin<sup>-</sup> BM) with MAGIT and expression of eGFP was also verified in c-Kit<sup>+</sup>Sca-1<sup>+</sup> HSC (**Figure 1e**).

### Sustained multilineage hematopoietic repopulation by MAGIT-expressing cells after non-myeloablative syngeneic neonatal transplantation

To test the feasibility of non-myeloablative transplantation and in vivo chemoselection in the absence of immune barriers to engraftment, transplantation of syngeneic MAGIT-transduced Lin<sup>-</sup> BM was performed at days 1–2 of life after a conditioning regimen that included BU and 200cGy radiation. Importantly, neonates treated with this regimen showed normal growth, and comparable survival to untreated, age-matched cohorts at 8 weeks, regardless of whether mice were subsequently transplanted or not (data not shown). Peripheral blood counts including total WBC (5.98  $\pm$  0.63  $\times$  10<sup>4</sup>/µl), lymphocytes (2.37  $\pm$  0.56  $\times$  $10^4/\mu$ l), and absolute neutrophil counts ( $3.16 \pm 0.45 \times 10^4/\mu$ l) in BU/200 treated neonates at day 2 of life, before transplantation, were not statistically different from those in untreated neonates (n = 3). After two cycles of BG/BCNU administered at 4 and 8 weeks of life, robust enrichment of transduced populations from low levels (0.67%) to  $\sim$ 50% of peripheral blood mononuclear cell (PBMC) was observed (Figure 2a, dot plots). The mean fluorescence of the GFP<sup>+</sup> population did not differ significantly following each cycle of BG/BCNU suggesting that clones expressing higher levels of MGMT<sup>P140K</sup> were not being preferentially selected. The frequency of GFP expression in PBMC was proportional to the total number of transduced cells in the thymus and spleen, but under-represented the levels of transduced cells in BM and Lin<sup>-</sup> BM fractions from chemoselected animals (Figure 2a, histograms). Mice transplanted with MAGIT-transduced Lin<sup>-</sup> BM, without subsequent BG/BCNU treatment, had gradually decreasing numbers of GFP<sup>+</sup> PBMC persisting for more than 10 months (data not shown). Initiation of chemoselection 10 months after transplantation of MAGIT-transduced cells was also successful in expanding transduced populations, suggesting targeting of longlived HSC and persistence of MGMTP140K expression even in the absence of selection (data not shown). Co-expression of intracellular MGMT<sup>P140K</sup> and GFP was confirmed in these chemoselected populations by flow cytometry (data not shown).

To further evaluate potential cumulative effects of successive cycles of chemoselection, a toxicity study was undertaken in which different doses of BG/BCNU were administered to

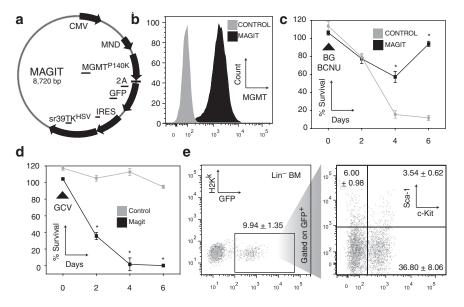


Figure 1 *In vitro* validation of transduction, expression, and function of MGMT<sup>P140K</sup>-2A-GFP-IRES-TK (MAGIT) lentivirus. (**a**) Map of MAGIT tricistronic plasmid co-expressing MGMT<sup>P140K</sup>, enhanced green fluorescent protein (eGFP) linked by a 2A sequence, and followed by TK<sup>HSV</sup> separated by an internal ribosome entry site (IRES). (**b**) 293T cell line was transduced with MAGIT lentivirus and intracellular MGMT<sup>P140K</sup> and GFP expression analyzed by flow cytometry. The proportion of cells shown expressing intracellular MGMT<sup>P140K</sup> and GFP expression was >90% positive cells as compared with untransduced cells. In all histograms the y-axis represents cell counts. (**c**) The same cell population in **b** was exposed to BG followed by BCNU, separated by 1 hour. Survival as a percentage of untreated cells was determined in triplicates at 2-day intervals. (**d**) Simultaneous to **c**, MAGIT transduced cells were exposed to Garciclovir (GCV) and monitored similarly. (**e**) Lineage-depleted bone marrow from C57Bl/6 mice was transduced with MAGIT lentivirus and the levels of GFP, Sca-1, and c-Kit expression determined by flow cytometry 48 hours following transduction. The dot plot showing Sca-1 and c-Kit expression is gated to include only GFP<sup>+</sup> cells. These data are representative of at least three independent experiments and any differences with *P* < 0.05 are indicated with an asterisk (\*). Gray: control-untransduced cells; black: MAGIT-transduced cells. BCNU, 1,3-bis(2-chloroethyl)nitrosourea; BG, benzylguanine; CMV, cytomegalovirus.

adult mice, weekly four times, and total lymphocyte counts and survival monitored (**Supplementary Figure S1**). Cumulative lymphopenia was observed in all treated groups at day 28, after four doses of BG/BCNU (**Supplementary Figure S1a**). There was no impact on survival in groups receiving 10 mg/kg BG/5 mg/kg BCNU or 20 mg/kg BG/5 mg/kg BCNU (**Supplementary Figure S1b**). Decreased survival was observed in the groups receiving 30 mg/kg BG/5mg/kg BCNU or 10 mg/kg BG/10 mg/kg BCNU (**Supplementary Figure S1b**). Thereafter, recovery of counts was observed over the next 2 months in all treated mice (**Supplementary Figure S1a**).

Multilineage repopulation by MAGIT-transduced, chemoselected Lin<sup>-</sup> BM was assessed before and after secondary transplantation into lethally irradiated syngeneic hosts. Flow cytometric analysis after secondary transplantation of Lin<sup>-</sup> BM from mice receiving BG/BCNU *in vivo* chemoselection showed that GFP<sup>+</sup> splenocytes contained lymphoid and myeloid progenitor populations including CD3<sup>+</sup>, CD3<sup>+</sup>CD49b<sup>+</sup>, CD4<sup>+</sup>, CD4<sup>+</sup>CD25<sup>+</sup>, CD19<sup>+</sup>, Gr-1<sup>+</sup>, and CD11c<sup>+</sup> populations, respectively (Figure 2b). Similar frequencies of GFP-expressing cells, and multilineage reconstitution were observed in the primary and secondary recipients.

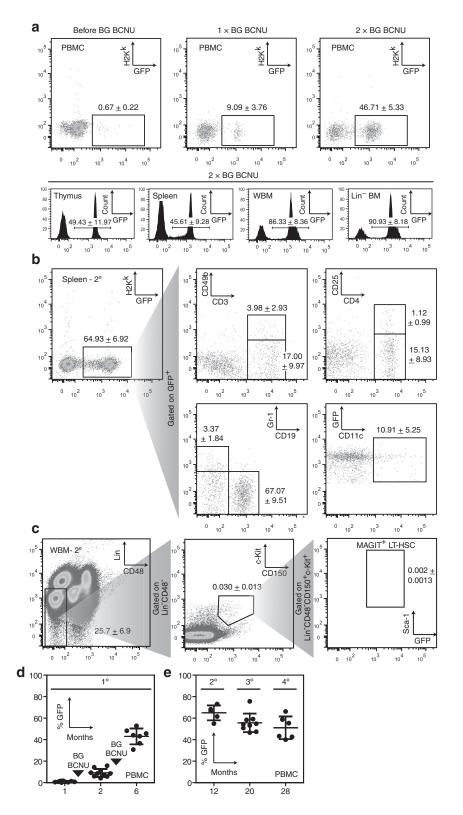
The presence, chemoselection, and persistence of long-term repopulating HSC was demonstrated by additional flow cytometric studies. A population of Lin<sup>-</sup>CD48<sup>-</sup>c-Kit<sup>+</sup>CD150<sup>+</sup>Sca-1<sup>+</sup> cells<sup>32</sup> expressing GFP were detected at comparable percentage to that of GFP<sup>+</sup> cells within the bone marrow of serially transplanted recipients (**Figure 2c**). After discontinuing BG/BCNU in primary transplants at 2 months, the frequency of GFP<sup>+</sup> PBMC populations persisted at 6 months (Figure 2d) and remained stable for 28 months in secondary, tertiary, and quaternary transplants (Figure 2e). These data are consistent with transduction and chemoselection of pluripotent HSC.

### Sustained multilineage hematopoietic repopulation and chemoselection of MAGIT-expressing cells after non-myeloablative allogeneic neonatal transplantation

The efficacy of this non-myeloablative preparative regimen and post-transplant chemoselection protocol was then evaluated in a fully major histocompatibility complex (MHC) mismatched allogeneic setting by transplantation of MAGIT-transduced B10.Br (H-2<sup>k</sup>) Lin<sup>-</sup> BM into C57Bl/6 (H-2<sup>b</sup>) neonatal recipients. Engraftment of allogeneic Lin<sup>-</sup> BM and the presence of MAGITtransduced GFP<sup>+</sup> PBMC were detected before chemoselection (Figure 3a, left dot plot). Efficient expansion of transduced donor allogeneic GFP+ cells was also achieved with BG/BCNU treatment in this MHC-mismatched allogeneic transplant model (Figure 3a, middle and right dot plots). Recipients of MHCmismatched cells had no signs of GVHD and showed comparable growth and appearance to naive littermates. Secondary transplantation of chemoselected B10.Br Lin- BM from primary C57Bl/6 recipients into lethally irradiated B10.Br adults produced multilineage reconstitution of lymphoid and myeloid progenitor compartments (Figure 3b). Humoral responses to GFP neoantigens described by others<sup>33</sup> were not detected in neonatally transplanted mice (Figure 3c).

# Negative selection of MAGIT-expressing engrafted cells by GCV administration

Several potential risk factors were considered in developing this non-ablative allogeneic transplant model based on lentiviral HSC transduction. The first potential limitation was evolution of GVHD after allogeneic transplantation in the absence of any post-transplant immunosuppression. The second concern was the potential transformation/autonomous proliferation of lentivirally transduced cells due to insertional mutagenesis.<sup>25–27</sup> We therefore incorporated a negative selection strategy into the MAGIT vector that would enable elimination of both alloreactive TK<sup>HSV</sup>-expressing T cell populations contributing to GVHD, and



autonomously proliferating clonal populations. Since GVHD was not detected either by examining the appearance of animals, or by testing liver function, it was not feasible to directly evaluate the efficacy of TK<sup>HSV</sup> expression and GCV treatment in this regard. However, proof of concept for GCV-mediated negative selection in vivo was established by administration of GCV (50 mg/kg) to a cohort of lethally irradiated adult mice after secondary transplantation of MAGIT-transduced and -chemoselected, syngeneic Lin<sup>-</sup> BM. A decrease in the mean frequency of GFP<sup>+</sup> donor cells to 0.47%, compared with 68.8% in untreated group, was demonstrated 12 weeks after a single cycle of GCV (Figure 3d, left graph). GCV-mediated negative selection in vivo was not associated with severe pancytopenia, as changes in blood counts were transient and returned to normal levels (Figure 3d, right graph). Re-expansion of transduced donor cells was achieved after readministration of two cycles of BG/BCNU in animals previously treated with GCV (Figure 3d, left graph). Thus, although GCV mediated the depletion of committed progenitors and circulating transduced cells, these data suggest that quiescent HSC remained viable and that GCV may be administered in this setting without ablation of the graft.

Multiple studies have demonstrated that transplantation and *in vivo* chemoselection of MGMT<sup>P140K</sup>-expressing BM results in polyclonal engraftment and proliferation.<sup>18–20,24,34–36</sup> To verify the multiclonality of engrafted, selected populations after positive and negative section with GCV and further re-expansion with BG/BCNU, linear amplification-mediated PCR (LAM-PCR) analysis of repopulating cells was performed (**Figure 3e,f**). Data from individual tertiary transplants demonstrate polyclonal engraftment (**Figure 3e**). After negative selection with GCV and re-expansion of transduced BM by re-administration of BG/BCNU multiple repopulating clones were also observed. Secondary transplants of Lin<sup>-</sup> BM from GCV-treated mice produced engraftment and persistence of multiple independent clones suggesting HSC remained viable (**Figure 3f**).

### Correction of $\beta$ -thalassemia with neonatal nonmyeloablative allogeneic transplantation and *in vivo* chemoselection

Effective, non-myeloablative allogeneic transplantation protocols with less toxicity would enable broader application of this approach for treatment of hematopoietic and other genetic disorders. Mutations in the human globin loci causing thalassemia are among the most common genetic disorders worldwide. Several studies have demonstrated correction of mouse models of  $\beta$ -thalassemia

or sickle cell disease by *ex vivo* lentiviral-mediated gene transfer of globin genes and transplantation.<sup>34</sup>,<sup>37–41</sup> The feasibility of *in vivo* chemoselection of syngeneic BM transduced with globin/MGMTexpressing vectors has been demonstrated by transplantation of adult murine models of  $\beta$ -thalassemia.<sup>34,39</sup> However, variability in persistence and levels of gene expression were observed.

To assess the potential for correction of  $\beta$ -thalassemia with this non-ablative allogeneic neonatal transplantation approach, Hbbth-3 thalassemic mice carrying deletions of the adult murine  $\beta$ -minor and  $\beta$ -major globin genes<sup>42</sup> were transplanted. Heterozygous Hbbth-3/+ mice have thalassemia intermedia with characteristic red cell changes on peripheral blood smear, and extramedullary hematopoiesis (Figure 4a,b).43 Homozygous Hbb<sup>th-3/th-3</sup> knockout mice do not survive gestation due to severe anemia and hydrops fetalis. MAGIT transduction ex vivo and transplantation of either syngeneic normal C57Bl/6 Lin- BM or allogeneic B10.Br Lin- BM was performed at days 1-2 of life in Hbbth-3/+ mice. After two cycles of BG/BCNU at 4 and 8 weeks post-transplantation, normalization of hematocrit, hemoglobin, RBC counts (Figure 4a), and RBC half-life were achieved (Figure 4b). Representative blood smears from transplanted groups show reduced numbers of target cells and other red cell abnormalities, consistent with the correction of the disease phenotype (Figure 4c). The lack of toxicity of the preconditioning regimen was underscored by the comparable survival of BU-conditioned Hbb<sup>th-3/+</sup> neonates that were not transplanted, to naive Hbbth-3/+ littermates. Also, adult Hbbth-3/+ naive controls tolerated at least two cycles of BG/BCNU without adverse effects on survival. Thus, both the BU/low-dose irradiation of this preparative regimen and in vivo chemoselection were well tolerated by the thalassemic Hbb<sup>th-3/+</sup> mice.

### DISCUSSION

Gene transfer approaches have shown great potential for the treatment of genetic disorders. In animal models, disease correction has required high-level transduction and ablative preconditioning. However, for many other genetic diseases, the corrective transgene cassette may not encompass all the regulatory elements required to reconstitute stable, tissue-specific, and physiologically appropriate levels of expression. Further, the design and validation of specific vectors for each disease constitutes a time-consuming and costly process.

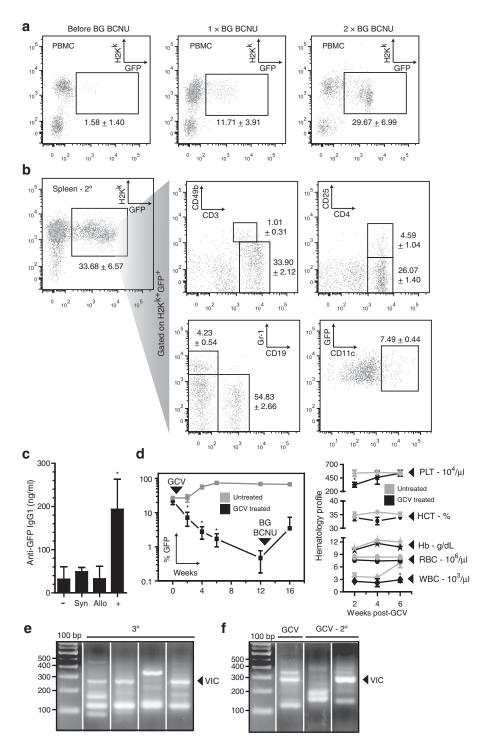
Rather than seeking direct genetic correction of autologous cells, there may be significant advantages to employing gene transfer strategies to improve the safety and outcome of allogeneic

**Figure 2** Non-myeloablative syngeneic neonatal transplantation with efficient chemoselection of MAGIT-expressing cells and multilineage hematopoietic repopulation. (a) Neonatal C57Bl/6 mice were transplanted with MAGIT-transduced C57Bl/6 Lin<sup>-</sup> BM and the percentage of peripheral blood cells expressing GFP and H-2K<sup>k</sup> determined by FACS 1 month post-transplant, and after each successive cycle of chemoselection with BG/BCNU (n = 10). (b) Lin<sup>-</sup> BM from chemoselected mice (similar to a) were harvested and transplanted into lethally irradiated C57Bl/6 mice. One month post-transplantation, the repopulation of indicated lineages in secondary recipients was evaluated. The data shown is gated to include only GFP-positive cells (n = 6). (c) Bone marrow harvested from serially transplanted mice (similar to b) was assessed by flow cytometry for the presence of MAGIT-expressing GFP<sup>+</sup> cells in long-term hematopoietic stem cells (LT-HSC). MAGIT<sup>+</sup> LT-HSC were defined by coexpression of GFP, Sca-1, c-Kit, and CD150 on cells lacking lineage markers (lineage-specific antibodies to CD5, B220, CD11b, Anti-Gr-1, 7-4, and Ter-119) and CD48 were used in these studies (n = 5). These data are representative of at least two independent experiments. (d) Dot plot depicting the percent of GFP<sup>+</sup> peripheral blood cells as determined by FACS at 1, 2, and 6 months after BG/BCNU administration, 1 and 2 months post-transplant as in **a**. (e) The percent GFP-expressing PBMC following serial transplantation of mice from **e**. Each dot represents a single mouse. The data from each panel are representative of at least two independent experiments as fluorescence-activated cell sorting; GFP, green fluorescent protein; PBMC, peripheral blood mononuclear cell.

transplantation, an established curative approach. However, application of allogeneic transplantation also has limitations due to the lack of suitably matched donors, the need for toxic recipient conditioning, risks of morbidity and mortality from GVHD, and infectious complications of immunosuppression and delayed immune reconstitution.

To address some of the current issues associated with allogeneic transplantation, a novel positive/negative selection strategy was pursued to provide both a selective advantage for donor HSC, and a mechanism for depletion of alloreactive T cells and potential autonomously proliferating clones after transplantation. The functionality of a tricistronic lentivirus including an eGFP reporter with linked MGMT<sup>P140K</sup> and TK<sup>HSV</sup> cassettes were demonstrated in both *in vitro* and *in vivo* studies. In these initial transplantation studies, positive/negative chemoselection was combined with transplantation in the neonatal period, when hostmediated immune responses are decreased.

The combination of a minimal, non-myeloablative conditioning regimen with *in vivo* BG/BCNU chemoselection was sufficient to enable engraftment and efficient *in vivo* expansion of lentivirally



transduced HSC. Stable levels of donor chimerism were observed after cessation of drug therapy, and after secondary and tertiary transplantation. Significantly, this minimal preparative regimen also enabled stable engraftment and enrichment of lentivirally transduced allogeneic cells without post-transplant immunosuppression. No evidence of GVHD was detected after discontinuing *in vivo* chemoselection.

The efficacy of this approach was further validated by transplantation and chemoselection of lentivirally transduced allogeneic Lin<sup>-</sup> BM in a murine model of  $\beta$ -thalassemia. Neither the preparative regimen, nor the administration of successive cycles of BG/BCNU produced adverse effects or decreased survival in naive thalassemic controls. Secondary transplantation of lentivirally transduced, chemoselected Lin<sup>-</sup> BM resulted in full multilineage reconstitution in lethally irradiated recipients suggesting that self-renewing HSC were successfully transduced and expanded in thalassemic recipients.

Several transplantation studies in both murine and large animal models, using BM transduced with MGMT<sup>P140K</sup> or another variant, MGMT<sup>G156A</sup>, have shown that successive cycles of chemotherapy with BG/BCNU or temozolomide results in in vivo selection at the stem cell level.<sup>11,12,16-20,30,39</sup> In the canine model, stable donor chimerism was observed for up to 6 years following discontinuation of the selective agent.<sup>18</sup> Multiclonal, stable engraftment, and further enrichment of donor cells with readministration of chemoselection more than a year after initial treatment was observed.<sup>18</sup> LAM-PCR-based genomic studies of retrovirally transduced canine HSC-expressing MGMTP140K showed no evidence for increased transcriptional start sites in proximity to proto-oncogenes, or differences in retroviral integration site patterns after chemoselection with either BCNU or temozolomide.<sup>18</sup> The successful transduction and in vivo selection of human CD34<sup>+</sup> cells transduced with MGMT<sup>P140K</sup> in immunodeficient models<sup>14,21</sup> further supports the promise of clinical application of MGMTP140K-mediated in vivo expansion of modified HSC.

Importantly, to date there are no reports of malignant transformation of BM with viral transduction and *in vivo* expansion of MGMT-expressing donor cells. The incorporation of the HSV<sup>TK</sup> suicide gene described here facilitates depletion of autonomous clones that might arise from insertional mutagenesis. The goal of the GCV-mediated negative selection experiment after BG/ BCNU chemoselection was to test the hypothesis that proliferating clones in the periphery or in the bone marrow would have greater sensitivity to GCV than more quiescent HSC in the bone marrow. Hence, a single dose of GCV was administered to assess the degree of peripheral depletion and to demonstrate that multilineage re-expansion of selected donor HSC was feasible after GCV. Since autonomous clones could persist after single dose GCV further negative selection might be required to eliminate these clones.

In contrast to the successful *in vivo* chemoselection and stable chimerism reported in canine models, recent studies in a rhesus macaque model, showed only transient expansion of autologous cells transduced with MGMT<sup>P140K</sup>, and toxicity with administration of a more dose-intensive BG/BCNU regimen.<sup>22</sup> Use of the MND promoter used in our and other successful chemoselection studies may have advantages in directing higher levels of gene expression and decreased silencing effects.<sup>44</sup>

Although genetic modification of donor cells to enable engraftment and *in vivo* amplification may make further reduction of the intensity and toxicity of preparative regimens feasible, risks of GVHD and the infectious consequences of required immunosuppression remain.<sup>4</sup> T-cell depletion of donor cells reduces the incidence and severity of GVHD, but has been associated with protracted immunodeficiency and increased mortality.<sup>4</sup> The TK<sup>HSV</sup>-based negative selection strategy incorporated in our studies has already been validated in clinical trials where recipients developing GVHD after donor TK<sup>HSV</sup> transduced T-cell lymphocyte infusion post-transplant were uniformly responsive to administration GCV.<sup>4</sup>

In the present study, efficient depletion of transduced donor cell populations was also observed with GCV administration after *in vivo* chemoselection of MAGIT-transduced BM. Importantly, the graft was not ablated after GCV administration enabling re-expansion with further chemoselection. The use of BM transduced with the bifunctional MAGIT lentiviral construct may enable modulation of the donor graft to limit risks associated with transplantation and gene delivery. This model system also enables studies of mechanisms underlying tolerance induction to neoantigens, and allogeneic stem cells during immune ontogeny.

Figure 3 Non-myeloablative allogeneic neonatal transplantation with chemoselection of MAGIT-expressing cells and multilineage hematopoietic repopulation. (a) Neonatal C57BI/6 mice were transplanted with MAGIT-transduced B10.Br Lin<sup>-</sup> BM and the percentage of peripheral blood cells expressing GFP and H-2K<sup>k</sup> determined by flow cytometry beginning 1 month after transplantation, and following two cycles of chemoselection with BG/BCNU at 8 and 12 weeks after transplant (n = 9 mice). (b) Lin<sup>-</sup> BM from mice after BG/BCNU treatment × 2 (similar to a) were harvested and secondarily transplanted into lethally irradiated B10.Br mice. One month after secondary transplantation, the repopulation of indicated lineages was re-evaluated by flow cytometry. The data shown is gated to include only GFP<sup>+</sup> cells (n = 6 mice). These data are representative of at least two independent experiments. (c) The serum of mice after neonatal transplantation of syngeneic (Syn) or allogeneic (Allo) MAGIT-transduced Lin<sup>-</sup> BM and BG/BCNU chemoselection (n = 5 mice per group) was tested for the presence of antibodies specific to GFP. Positive control (+) samples were generated by injection of C57BI/6 mice with alum/GFP. (d) Lin<sup>-</sup> BM from mice treated with BG/BCNU were harvested and transplanted into lethally irradiated C57BI/6 mice (similar to Figure 2b). One month after secondary transplantation, Ganciclovir (GCV) was administered intraperitoneally. The frequency of GFP<sup>+</sup> PBMC was monitored before administration of GCV and for 12 weeks after GCV administration at the indicated time points. One cycle of BG/BCNU was administered 12 weeks after GCV administration and the percent GFP-positive PBMC quantified (left graph). Hematocrit (HCT), hemoglobin (Hb), red blood cell (RBC), white blood cell (WBC), and platelet (PLT) levels were determined 2, 4, and 6 weeks post-GCV treatment (right graph). These data are representative of one experiment (n = 4 mice per group) and any differences with P < 0.05 are indicated with an asterisk (\*). (e) Clonality as indicated by presence of 5' lentiviral LTR adjacent genomic amplicons was imaged following LAM-PCR. Genomic DNA from PBMC following tertiary transplantation of Lin<sup>-</sup> BM as described in Figure 2b (n = 4 mice) was utilized. (f) Clonality of PBMC from mice treated with GCV as described in d and following secondary transplantation of Lin- BM from GCV-treated mice is shown. A 100 bp ladder and corresponding vector internal control (VIC) band are indicated. BCNU, 1,3-bis(2-chloroethyl)nitrosourea; BG, benzylguanine; GFP, green fluorescent protein; LAM-PCR, linear amplication-mediated PCR; LTR, long terminal repeat; PBMC, peripheral blood mononuclear cell.

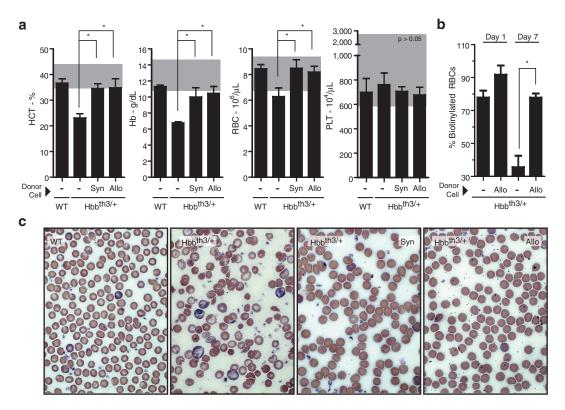


Figure 4 Correction of  $\beta$ -thalassemia following non-myeloablative neonatal transplantation and chemoselection of MAGIT. (a) Heterozygous Hbb<sup>th-3/+</sup> neonatal mice underwent busulfan/200cGy preconditioning, and were transplanted with MAGIT-transduced syngeneic or allogeneic Lin-BM, and treated with two successive BG/BCNU chemoselection cycles. Peripheral blood counts were followed. Gray area indicates the normal range for mice. These data are representative of at least two independent experiments and any differences with *P* < 0.05 are indicated with an asterisk (\*) (*n* = at least four mice per group). (b) RBC half-life, indicated by the decay of biotinylated RBCs in peripheral blood, was monitored by flow cytometry on days 1 and 7 following treatment with Biotin-NHS (*n* = 4 mice per group). (c) Representative Giemsa-stained blood smears of mice from each group from a are shown. BCNU, 1,3-bis(2-chloroethyl)nitrosourea; BG, benzylguanine; Hb, hemoglobin; HCT, hematocrit; PLT, platelet; RBC, red blood cell; WT, wild-type.

### MATERIALS AND METHODS

*Mice.* Wild-type C57Bl/6 (H-2<sup>b</sup>), and B10.Br (H-2<sup>k</sup>) mice were obtained from Charles River (Wilmington, MA), and the Jackson Laboratory (Bar Harbor, ME), respectively. Hbb<sup>th-3</sup> mice carrying a deletion of the murine β-major and β-minor globin genes (gift of Dr Oliver Smithies, University of North Carolina, Chapel Hill, NC)<sup>43</sup> were backcrossed at least 20 generations and maintained on a C57Bl/6 background. The mouse β-globin deletion was detected by PCR screening for a 385 bp product using sense primer 5'-AACAAGAGCAAACTAAGTAAGATGC-3' and antisense primer 5'-AATTCGCCAATGACAAGACG-3'. PCR results were confirmed by analysis of hematocrit and hemoglobin using a Hemavet 850FS system (Drew Scientific, Waterbury, CT). All strains were housed under specific pathogen-free conditions at the University of California, San Francisco, CA. Experiments were conducted under the supervision of the University of California, San Francisco Institutional Animal Care and Use Committee according to an approved protocol.

*Fluorescence-activated cell sorting.* Antibodies to H-2K<sup>k</sup> (36-7-5), MGMT (MT5.1), CD3 (2C11), CD4 (RM4-5), CD25 (PC61), CD49b (HMa2), Ly-6G and Ly-6C (Gr-1), CD19 (1D3), CD11c (HL3), c-Kit (2B8), Sca-1 (D7), and Fc Block (2.4G2) were purchased from BD Biosciences (San Jose, CA). Antibodies to CD150 (TC15-12F12.2) and CD48 (HM48-1) were purchased from BioLegend (San Diego, CA). Flow cytometry was performed on a LSR II and data analyzed on FlowJo software (Treestar, Ashland, OR) using indicated antibodies and Fc Block, with shown gates excluding at least 99.5% of control cells. Intracellular staining specific for human MGMT was performed after fixation with 4% paraformaldhyde, permeabilization with 0.5% saponin, and detection of bound anti-MGMT with a Goat anti-mouse IgG1 PE as described (Invitrogen, Carlsbad, CA).  $^{\rm 45}$ 

Construction of MAGIT plasmid. The tricistronic MAGIT expression plasmid (pMAGIT) coexpressing MGMT<sup>P140K</sup>, eGFP linked by a 2A sequence, and followed by  $\mathrm{TK}^{\scriptscriptstyle\mathrm{HSV}}$  separated by an internal ribosome entry site (IRES) was constructed by inserting an 2,873 bp NdeI/SpeI fragment containing the MGMT<sup>P140K</sup> and GFP sequences from MAG plasmid<sup>31</sup> into NdeI/SpeI digested MGMT<sup>P140K</sup>-IRES-TK (a bicistronic plasmid lacking the GFP gene that was generated in our laboratory). This approach reintroduced  $MGMT^{{}_{P140K}}$  and added a GFP cassette to create a tricistronic plasmid under the control of the MND promoter for stable long-term expression.<sup>44</sup> The plasmid containing the sr39TK<sup>HSV</sup> was a gift from Dr Noriyuki Kasahara (University of California Los Angeles, Los Angeles, CA). This sr39 mutant of TKHSV has enhanced ability to convert acyclovir/GCV into cytotoxic agents and greater in vivo positron emission tomography imaging sensitivity.46 The IRES-TKHSV cassette was cloned from plasmid pACE-TK, which encodes a replication-competent amphotropic MLV vector derived from pACE-GFP47 but with the GFP gene replaced by the TKHSV gene amplified from plasmid pFRA-TK48 by PCR using Pfu polymerase (Stratagene, La Jolla, CA). Plasmid DNA was prepared with PureLink HiPure Plasmid DNA Megaprep and Gigaprep Kits according to the manufacturer's instructions (Invitrogen).

**Production and titering of lentivirus vectors.** MAGIT lentivirus was produced by overnight transfection of 293T cells grown to 80% confluence on poly-lysine coated (Sigma-Aldrich, St Louis, MO) 10 cm dishes with three plasmids (pCMVΔR8.91,<sup>49</sup> pMD.G,<sup>49</sup> and pMAGIT) and Lipofectamine 2000 (Invitrogen). Supernatants were collected 24 and 48 hours after removal of transfection media, and pooled before concentration on 70% ethanol washed Centricon-80 columns (Millipore, Billerica, MA). Aliquots of virus were frozen at -80 °C. The titer of concentrated virus was determined by serial dilution onto 293T cells in the presence of 8µg/ml polybrene. Using this procedure, we observed titers at  $1.2 \pm 0.12 \times 10^9$  IU/ml (data from six viral lots).

In vitro *BG/BCNU* chemoselection and *GCV* killing curves. MAGITtransduced 293T cells (>90% GFP expression by fluorescence-activated cell sorting) were either treated with 10µmol/l BG (Gift from Dr Gary Pauly, National Cancer Institute, Frederick, MD) followed 1 hour later by 200µmol/l BCNU (Bristol-Myers Squibb, Princeton, NJ) or with 1µg/ml GCV (InvivoGen, San Diego, California). Cell survival was assessed every other day by evaluating conversion of MTS (Promega, Madison, WI) into the aqueous soluble formazan product. The quantity of formazan product as measured by the amount of 490 nm absorbance is directly proportional to the number of living cells in culture.

Lentiviral transduction of murine Lin– BM cells. Bone marrow was harvested from femurs and tibias of C57Bl/6 or B10.Br mice. Lineage positive populations were depleted using the Lineage Cell Depletion Kit (Miltenyi Biotec, Auburn, CA) according to the manufacturer's instructions. The resulting Lin<sup>-</sup> BM was >98% depleted of lineage populations with undetectable CD3<sup>+</sup> cells (data not shown). Lin<sup>-</sup> BM cells were transduced with 100 multiplicity of infection of MAGIT lentivirus at a density of 10<sup>6</sup> cells/ml in DMEM (Invitrogen) containing 10% fetal calf serum supplemented with 100 ng/ml rmFlt-3L, 100 ng/ml rmSCF (R&D Systems, Minneapolis, MN), and 8µg/ml polybrene (Sigma-Aldrich). Following overnight incubation, the transduced cells were harvested, washed, and 10<sup>6</sup> cells injected *via* facial vein in 50µl of phosphate-buffered saline. GFP expression, an indicator of transduction efficiency, was evaluated by flow cytometric analysis after culture for an additional 48 hours. This transduction protocol consistently produced GFP expression in 9–12% of Lin<sup>-</sup> BM.

**Preconditioning and transplantation of neonates.** C57Bl/6 females were timed-mated with either wild-type C57Bl/6 or Hbb<sup>th-3/+</sup> heterozygous C57Bl/6 male mice. On day 18 of gestation (E18), 15 mg/kg of Busulfan (Aldrich, Milwaukee, WI) was delivered intraperitoneally into pregnant dams.<sup>3</sup> Neonates from BU-treated dams were exposed to 2 Gy (200 rad) on day 1 or 2 of life and injected with the indicated cell numbers on the same day.

In vivo positive and negative selection procedures for MAGIT-transduced populations. At 1 month of life and before chemoselection, mice were bled via submandibular vein. The fractions of GFP- and H-2K<sup>k</sup>-expressing PBMC were determined via flow cytometry. *In vivo* chemoselection was performed by intraperitoneal administration of BG at 30 mg/kg and then BCNU at 7.5 mg/kg, 1 hour later. The level of donor cell enrichment was evaluated 1 month after each cycle of chemoselection before drug administration was repeated. Negative selection by *in vivo* depletion of MAGIT-expressing cells was performed by a single intraperitoneal administration of GCV at 50 mg/kg.<sup>50</sup>

Secondary transplantation of MAGIT-transduced cells. Bone marrow, spleen, and thymus were harvested after neonatal transplantation and chemoselection. Lineage repopulation of each of these tissues with MAGIT-transduced cells was evaluated *via* flow cytometric analysis after staining with the panel of indicated antibodies. The remainder of the harvested bone marrow was depleted of lineage positive populations, as described above, and injected intravenously into lethally irradiated (10 GY) syngeneic adult recipients. The frequency of GFP-expressing cells and repopulation of different lineages were again evaluated after secondary transplantation.

Anti-GFP ELISA. Plates were coated with eGFP (Promega) overnight. After blocking, the plate was incubated with serum for 1 hour followed by

incubation with anti-mouse IgG HRP (Calbiochem, Gibbstown, NJ) for an additional hour separated by a wash step. The amount of anti-GFP antibody was quantified with reference to a known control anti-mouse GFP IgG (Abcam, Cambridge, MA). Positive control serum was produced by repeated injection of C57Bl/6 mice with a suspension of Alum and eGFP.

**RBC** biotinylation decay assay. Red blood cells were pulse biotinylated with N-hydroxysuccinimide biotin (Biotin-NHS; Calbiochem, San Diego, CA) dissolved in N,-N,-dimethylacetamide at 50 mg/ml. Before tail vein injection, the dissolved Biotin-NHS was diluted in phosphate-buffered saline to 5 mg/ml and delivered at 50 mg/kg per mouse. The percentage of biotinylated RBCs in peripheral blood was quantified by flow cytometry at indicated time points after incubation with strepavidin-PE (BD Biosciences, San Jose, California).

*Lentivirus insertion site analysis by LAM-PCR.* LAM-PCR analyses to identify 5' long terminal repeat lentiviral vector adjacent genomic sequences was performed in collaboration with the National Gene Vector Biorepository (NGVB) and Dr. Kenneth Cornetta (Indiana University, Indianapolis, IN) as described previously.<sup>24</sup> PBMC from indicated mice were collected and 200 ng of DNA used as template for two rounds of 50-cycle linear PCR with biotinylated lentiviral long terminal repeat-specific primer. Following magnetic separation of PCR products, the linear product was exposed to hexanucleotide random priming, restriction digestion with Mse1, ligation with a linker, and exponential PCR. Resulting amplicons adjacent to the 5' long terminal repeat were imaged following a second round of magnetic separation and nested PCR as described.<sup>24</sup>

**Statistics.** All experiments were repeated at least two times unless indicated otherwise. Results are reported as the mean  $\pm$  SD of independent experiments. The significance of differences was determined using Student's paired *t*-test.

### SUPPLEMENTARY MATERIAL

**Figure S1**. Effects of different weekly BG/BCNU regimens on total lymphocyte counts and survival.

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