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### **Humanized bone-marrow mouse model as a pre-clinical tool to assess therapy-mediated hematotoxicity**

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#### **Abstract**

**Purpose—**Pre-clinical *in vivo* studies can help guide the selection of agents and regimens for clinical testing. However, one of the challenges in screening anti-cancer therapies is the assessment of off-target human toxicity. There is a need for *in vivo* models that can simulate efficacy and toxicities of promising therapeutic regimens. For example, hematopoietic cells of human origin are particularly sensitive to a variety of chemotherapeutic regimens but *in vivo* models to assess potential toxicities have not been developed. In this study, a xenograft model containing humanized bone marrow is utilized as an *in vivo* assay to monitor hematotoxicity.

**Experimental Design—**A proof-of-concept, temozolomide-based regimen was developed that inhibits tumor xenograft growth. This regimen was selected for testing since it has been previously shown to cause myelosuppression in mice and humans. The dose-intensive regimen was administered to NOD/SCID/γchain<sup>null</sup> mice reconstituted with human hematopoietic cells and the impact of treatment on human hematopoiesis was evaluated.

**Results—**The dose-intensive regimen resulted in significant decreases in growth of humanglioblastoma xenografts. When this regimen was administered to mice containing humanized bone marrow, flow cytometric analyses indicated that the human bone-marrow cells were significantly more sensitive to treatment than the murine bone-marrow cells, and that the regimen was highly toxic to human-derived hematopoietic cells of all lineages (progenitor, lymphoid, and myeloid).

**Conclusions—**The humanized bone-marrow xenograft model described has the potential to be used as a platform for monitoring the impact of anti-cancer therapies on human hematopoiesis and could lead to subsequent refinement of therapies prior to clinical evaluation.

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#### **Keywords**

human xenograft model; myelosuppression; stem-cell

#### **INTRODUCTION**

Murine xenograft models are one of the primary tools used for screening of new therapeutic compounds and regimens (1). One major limitation of using murine xenograft studies to determine therapeutic efficacy, however, is that significant inter-species differences in drug sensitivity can exist between mouse- and human-derived cells. It is possible that the levels of a therapeutic compound reached in a mouse xenograft model may not be achievable in humans due to differential profiles of absorption, distribution, metabolism, excretion (ADME), and toxicity. Therefore, while a regimen may exhibit an acceptable toxicity and efficacy profile in mice, this may not necessarily be the case in patients.

#### **Statement of Translational Relevance (150 words max)**

A variety of human tumor xenograft models are currently used to evaluate the efficacy of novel compounds and regimens to kill tumor cells *in vivo*. However, *in vivo* models to screen for potential toxicity to normal human cells in their appropriate microenvironment, early in the drug discovery, need development. Since bone-marrow toxicity can be a major life-threatening side effect of treatment, models to screen for the impact of treatment on human hematopoiesis would improve our ability to select compounds with decreased off-target toxicities. A humanized bone-marrow xenograft model was developed to assess toxicity to human hematopoietic cells *in vivo*. As the first step towards validation of this model on a broader scale, we used a temozolomide-based regimen already known to exhibit dose-limiting toxicities in the clinic. This model holds promise as a new approach for *in vivo* hematotoxicity screening of new compounds and regimens.

Furthermore, *in vitro* clonogenic assays have demonstrated that human and mouse hematopoietic cells show a wide diversity in sensitivity to many anti-cancer drugs; mouse hematopoietic cells in many cases exhibit increased resistance to compound exposure compared to human hematopoietic cells (2-5). Masubuchi et al previously demonstrated differential sensitivities of mice, dog, and human bone-marrow cells exposed to camptothecin derivatives. For example, camptothecin-like compounds-SN-38 and topotecan-exhibited differential inter-species resistance with mouse CFU being the most resistant to treatment (4). However, inter-species differences in sensitivity were not always observed. Three other compounds (DX-8951f, 9-aminocamptothecin, and camptothecin) demonstrated fairly similar sensitivities amongst all three species. Studies by Kurtzberg et al (3) indicated significant differences in sensitivities for tubulin-binding agents; the  $IC_{90}$ values for vincristine and paclitaxel were 30 and 27 nM and for mouse CFU-GM and 3 and 9 nM for human CFU-GM respectively. In addition,  $IC_{90}$  values for tasidotin treatment were > 300 nM for mouse CFU-GM and 65 nM for human CFU-GM.

Evaluation of human hematopoietic toxicity in immunodeficient mice could represent an additional benchmark in the final screening and selection of new therapeutics. This *in vivo* screening approach would take into account the influence of the bone-marrow microenvironment on damaged cells or the cycling kinetics of hematopoietic cells following treatment. An unexplored use of the NOD.Cg-*PrkdcscidIL2rgtm1Wjl*/Sz (NOD/SCID/ γchainnull) mouse strain is to treat mice reconstituted with human hematopoietic cells with a regimen related to clinical practice, and determine the impact of *in vivo* treatment on human

hematopoiesis (6). As a proof-of-concept, we chose to evaluate the impact of a combination therapy consisting of  $O^6$ -benzylguanine and temozolomide ( $O^6$ -BG/TMZ) on human hematopoiesis *in vivo* since it is currently being evaluated in clinical trials and the main dose-limiting toxicity in these patients is myelosuppression (7-10). We first developed a dosing regimen consisting of  $O^6$ -benzylguanine and temozolomide ( $O^6$ -BG/TMZ) followed by stem-cell rescue that would significantly inhibit the growth of TMZ-resistant human xenografts in NOD.Cg-*Prkdcscid IL2rgtm1Wjl*/Sz mice. The underlying strategy of this regimen is to prevent repair of TMZ-mediated DNA damage by inhibiting the DNA repair protein, O<sup>6</sup>-methylguanine DNA- methyltransferase (MGMT) (11-14). Previous studies have demonstrated that numerous cancers can express high levels of MGMT, and therefore can efficiently repair TMZ-mediated DNA damage, decreasing the efficacy of tumor-cell kill (15). In addition, a variety of tumors can be sensitized to alkylating agents both *in vitro* and in xenograft studies with the addition of  $O^6$ -BG, a direct inhibitor of MGMT activity (11,13-14,16). The down side of this approach is that both immature and mature hematopoietic cells can be extremely sensitive to this regimen due to low levels of endogenous DNA repair activity (17).

In this study, the delivery of two cycles of a high-dose regimen of  $O<sup>6</sup>$ -BG/TMZ in combination with stem-cell rescue significantly inhibited the growth of a TMZ-resistant glioma. This course of therapy was then used to test the hypothesis that administration of the regimen would be toxic to human hematopoietic cells *in vivo*. NOD/SCID/γchainnull mice were transplanted with human CD34<sup>+</sup> cells and reconstitution confirmed one month posttransplantation by monitoring the peripheral blood (PB) for human-cell chimerism in the transplanted mice. Treatment with  $O<sup>6</sup>$ -BG/TMZ resulted in significant loss of humanderived hematopoietic cells in the bone-marrow and enumeration of total number of human and mouse cells in vehicle versus treated mice indicated that the human bone-marrow cells were significantly more sensitive to treatment compared to the murine-derived bone-marrow cells. This proof-of-concept study indicates that use of NOD/SCID/γchain<sup>null</sup> mice with humanized bone marrow can be used as an *in vivo* toxicity measure of human hematopoiesis following drug treatment and holds merit as a model for improving screening strategies used to develop and test novel compounds and dosing regimens.

#### **Materials and Methods**

#### **Isolation of umbilical cord blood (UCB) CD34+ cells**

All protocols were approved by Indiana University School of Medicine's Institutional Review Board (IRB) and St. Vincent Hospital's IRB (Indianapolis, IN). Samples of UCB were collected from normal, full-term infants delivered by cesarean section and the CD34<sup>+</sup> cells isolated using the CD34 MicroBead kit and VarioMACS Separator™ (Miltenyi Biotech Inc., Auburn, CA) according to the manufacturer's instructions. Following magnetic bead separation, the viability of the isolated CD34+ cells was routinely 95%. The quality of the isolated CD34+ cells was analyzed by a Colony-forming Unit (CFU) assay as described below on an aliquot of the CD34<sup>+</sup> cells used for transplantation. The frequency of clonogenic cells in UCB products used for transplant in experiments I and II was similar with 60-65 colonies per  $2 \times 10^3$  CD34<sup>+</sup> cells plated.

**Clonogenic survival assays—**Colony forming-Unit (CFU) assays were performed using enriched human CD34+ cells (Methocult GF, H4434, Stem Cell Technologies, Inc.) and unfractionated murine-bone marrow (Methocult GF M3434), or bone marrow from NOD/SCID/γchain<sup>null</sup> transplanted with human CD34<sup>+</sup> cells. The cells were seeded in triplicate dishes at concentrations of  $2 \times 10^3$  for human CD34<sup>+</sup> cells and  $2 \times 10^5$  for murine bone marrow to obtain 60-70 colonies per 35 mm dish. For survival assays, cells were

exposed to drug combinations ( $O^6$ -BG, TMZ or  $O^6$ -BG/TMZ) prior to plating. For analysis of human cells from the bone marrow of NOD/SCID/γchain<sup>null</sup> transplants, chimeric humanmouse bone marrow was plated at  $2 \times 10^5$  bone-marrow cells per dish. After 10 - 14 days of incubation at  $37^{\circ}$ C in 5% CO<sub>2</sub>, colony forming units-granulocyte-macrophage (CFU-GM) and burst-forming units-erythroid (BFU-E) were enumerated using the Axiovert 25 invertedlight microscope. For assays using clonogenic SF767 cells, cells were seeded at 150 cells/ well in triplicate in 6-well plates and allowed to grow overnight in IMDM medium with supplements (10% Cosmic Calf Serum, 1% Pen/Strep, and 1% Glutamine.) The cells were treated the next day and allowed to grow for 13 days at 37°C. Following incubation, the cells were rinsed with PBS, fixed with 0.05% methylene blue, and counted visually.

#### **Animals**

A breeding colony of NOD.Cg-*Prkdcscid IL2rgtm1Wjl*/Sz (NOD/SCID/γchainnull) mice (6) was established at the Laboratory Animal Research Center and maintained by the *In Vivo* Therapeutics Core at the Indiana University Simon Cancer Center (IUSCC) (Indianapolis, IN). BALB/c and C57BL/6N inbred mice were purchased from Harlan Laboratories, Inc (Indianapolis, IN). All protocols and establishment of pre-death endpoints-decreases in body weight, activity, and grooming-were approved by the Institutional Animal Care and Use Committee.

#### **Xenograft study**

The human glioblastoma cell line, SF767, was originally developed by The Brain Tumor Research Center, University of California at San Francisco and has been previously described (18). Dr. Len Erickson (Indiana University School of Medicine, Indianapolis, IN) provided this line; early-passage cells were cryopreserved in 2005 (18). Prior to use in xenograft studies, SF767 cells were tested for confirmation of human MGMT expression and sensitivity to  $O<sup>6</sup>$ -BG/TMZ. Human-cell line authentication was confirmed by the Research Animal Diagnostic Laboratory (Columbia, MO). SF767 cells  $(5 \times 10^6)$  were implanted subcutaneously into the flank of NOD/SCID/γchain<sup>null</sup> mice and tumors allowed to grow to  $100 - 150$  mm<sup>3</sup>. Tumor volume measurements were taken weekly by caliper and calculated according to the formula ( $\alpha^2 \times \beta$ )/2, where α is the shorter and β is the longer of the two dimensions.

#### *Transplantation of* **NOD/SCID/γchainnull** *mice with human CD34+ cells*

NOD/SCID/γchainnull mice were placed on food pellets containing 0.0625% doxycycline for 5-7 days prior to irradiation. Mice were then conditioned with 300-cGy total-body irradiation using a GammaCell 40 (Nordion International Inc., Ontario Canada) equipped with two opposing Cesium-137 sources. UCB CD34<sup>+</sup> cells were resuspended in IMDM containing 0.2 % endotoxin-free bovine serum albumin and injected into the lateral tail vein of each animal.

#### **Chemotherapy administration**

O<sup>6</sup>-BG (Sigma-Aldrich, St Louis, MO) was dissolved in 40% polyethylene glycol-400 (v/v) and 60% saline (v/v). TMZ (LKT laboratories, Inc., St. Paul, MN) was sonicated in phosphate buffered saline (PBS). The myeloablative dosing regimen consisted of 30 mg/kg O<sup>6</sup>-BG followed by 80 mg/kg TMZ one hour later and 15 mg/kg O<sup>6</sup>-BG 7 hours later (O<sup>6</sup>-BG/TMZ). One treatment cycle consisted of 3-consecutive days of treatment with this dosing regimen. Two days after a treatment cycle,  $8 \times 10^6$  fresh murine bone marrow cells were injected intravenously.

#### **Analysis of human cell engraftment**

Mice were sacrificed at 6 and 11 days post-injection and the BM and spleen prepared as previously described (19). Human cell engraftment was measured by human CD45 immunostaining. The proportion of engraftment in various lineages was determined by immunostaining and flow cytometric analysis as previously described using a Becton-Dickinson FACSCalibur and CellQuest software (19). The lack of crossreactivity of humanspecific antibodies with murine cells was confirmed in each experiment by staining BM from a nontransplanted mouse with each antibody combination. Cells were stained with allophycocyanin (APC)-conjugated anti-human-CD45 (anti-HLe-1; Becton Dickinson Immunocytometry, San Jose, CA) alone or in combination with phycoerthyrin (PE) conjugated anti-human CD33 (anti-Leu-M9; Becton Dickinson). Identical aliquots were stained with APC-conjugated anti-human CD34 (clone 581; PharMingen, San Diego, CA) in combination with anti-human CD19-PE (PharMingen).

**Statistical Analyses—**Generalized Linear Mixed Model with mouse as the random effect was used to compare tumor volume between  $O<sup>6</sup>$ -BG /TMZ and vehicle treatments. Twotailed t-tests were performed to determine significance of vehicle-versus drug-treated for tumor weights, clonogenic survival assays, and flow cytometric analyses.

#### **RESULTS**

#### **Development of a proof-of-concept TMZ-based regimen that results in significant decrease in tumor-xenograft growth**

A combination treatment consisting of  $O<sup>6</sup>$ -BG and TMZ was selected to serve as the proofof-concept treatment for development of the *in vivo* human toxicity model since myelosuppression is the major dose-limiting toxicity associated with this treatment in the clinic. (8-9) To gain insight into the drug sensitivity of the cell populations to be investigated (i.e. primary murine hematopoietic cells, primary human hematopoietic cells, and glioma cells), clonogenic survival assays were set up in the absence and presence of 20 μM  $O<sup>6</sup>$ -BG and increasing doses of TMZ (0 - 200 μM) (Fig. 1). Surrogate bone-marrow toxicity studies of new compounds and regimens are commonly evaluated in immunocompetent mice. Since immunodeficient mice would be required for engraftment of human CD34<sup>+</sup> cells, it was important to compare the drug sensitivity of murine clonogenic cells derived from immunocompetent (BALB/c, C57BL/6N) and immunodeficient mice (NOD/SCID/γchain<sup>null</sup>). Murine bone-marrow cells were set up in the absence or presence of drug combinations and sensitivity of clonogenic cells determined by CFU assays. The dose-dependent sensitivities of clonogenic cells derived from BALB/c, C57BL/6N and NOD/SCID/ychain<sup>null</sup> mice were similar to TMZ in the presence and absence of  $O^6$ -BG (Fig. 1A). Therefore, at least for the drug combination studied here, it was reasonable to assume that the sensitivity of the murine bone marrow in NOD/SCID/γchain<sup>null</sup> can serve as an accurate read-out of normal murine hematotoxicity. In the next experiments, human  $CD34<sup>+</sup>$  cells and NOD/SCID/γchain<sup>null</sup> bone-marrow cells were treated with increasing doses of TMZ in the absence or presence of  $O^6$ -BG. The concentrations of TMZ alone that resulted in 50% decrease in the colony number  $(IC_{50})$  varied slightly between mouse and human clonogenic cells (Human IC<sub>50</sub> human = 160 μM; Mouse IC<sub>50</sub>= 190 μM TMZ). However, once  $O^6$ -BG was included, the IC<sub>50</sub> doses varied significantly: a 3-fold difference in sensitivity between mouse and human clonogenic cells was evident  $(IC_{50})$  human = 25  $\mu$ M TMZ; mouse = 75 μM). The IC<sub>90</sub> doses for both mouse and human were similar (IC<sub>90</sub>) human = 150 μM; mouse = 180 μM TMZ), indicating that when cells were exposed to highdoses of O<sup>6</sup> -BG/TMZ, differences in the *in vitro* sensitivity were less pronounced *in vitro*. Clonogenic assays for the human SF767 glioblastoma cell line were also set up under these same conditions. SF767 cells have been shown by us and others to constitutively express

MGMT and are resistant to the methylating agent, TMZ, unless  $O<sup>6</sup>$ -BG is included (18,20). As expected, the SF767 tumor cells were highly resistant to TMZ (Fig. 1) and exhibited significantly increased sensitivity to TMZ in the presence of  $O^6$ -BG (IC<sub>90</sub>= 40 µM TMZ; Fig. 1), demonstrating the dependency of the SF767 cells on MGMT-mediated DNA repair.

We next determined to what extent the *in vitro* sensitivities of the cell populations correlated with *in vivo* anti-tumor efficacy and toxicity to the bone-marrow compartment in NOD/ SCID/ $\gamma$ chain<sup>null</sup> mice. To accomplish this, it was essential to develop an efficacious O<sup>6</sup>-BG/ TMZ regimen that resulted in significant tumor kill *in vivo*. Treatment cycles containing increasing doses of TMZ were first tested for toxicity in NOD/SCID/γchain<sup>null</sup> mice (Fig. 2A&B). Each cycle consisted of 3-consecutive days of treatment with 30 mg/kg  $O^6$ -BG followed one hour later by 40, 60 or 80 mg/kg TMZ and a second bolus of  $O<sup>6</sup>$ -BG delivered seven hours later. The  $O^6$ -BG-double bolus has been shown previously to deplete MGMT for at least 18 hours and prevent repair of TMZ-mediated DNA damage (21). Survival was determined using a pre-death endpoint system (see Material and Methods). Two cycles of treatment initiated a week apart resulted in morbidity in  $>85\%$  of the NOD/SCID/ $\gamma$ chain<sup>null</sup> mice (Fig 2A). Histopathological evaluation of major organs (brain, colon, kidney, liver, lung, heart, spleen, and stomach) were performed on 2 vehicle-treated mice and 4 mice treated with  $O^6$ -BG and followed by TMZ at 80 mg/kg and no gross abnormalities were noted. However, in mice treated with  $O<sup>6</sup>$ -BG and TMZ at 80 mg/kg, bone-marrow cellularity was decreased by at least 70% in the  $O<sup>6</sup>$ -BG/TMZ-treated mice compared to vehicle-treated mice when analyzed at the pre-death endpoint (data not shown). In contrast, when an autologous murine bone-marrow transplant was given 2 days after each cycle of drug administration, all mice maintained normal body weight and morbidity was prevented (Fig 2A). We next tested the efficacy of this regimen using NOD/SCID/γchain<sup>null</sup> mice containing ectopic SF767 glioblastoma xenografts (Figure 2B). In 2 independent experiments, mice received 2 cycles of either vehicle control or  $O<sup>6</sup>$ -BG/TMZ treatment. After each 3-day treatment cycle,  $8 \times 10^6$  autologous murine bone-marrow cells were transplanted via tail vein and tumor growth monitored over time. Following 2 cycles of treatment and stem-cell rescue, a significant delay in tumor progression was observed in vehicle- versus  $O^6$ -BG/TMZ-treated mice (Fig. 2C). At 8 weeks post-treatment, mice were sacrificed for tumor and bone-marrow analysis. Tumors from drug-treated mice weighed significantly less than tumors from vehicle-treated mice (Fig. 2D). Bone-marrow cellularity of vehicle-treated mice versus mice receiving  $O<sup>6</sup>$ -BG/TMZ and stem-cell rescue, was not significantly different (data not shown). These data indicate that significant decreases in growth of alkylator-resistant tumor cells can be obtained with high-dose  $O<sup>6</sup>$ -BG/TMZ when combined with autologous stem-cell rescue and provided us with a relevant proof-of-concept regimen that could be used to monitor impact of the treatment on human hematopoiesis *in vivo*.

#### *In vivo* **bio-toxicity assay to evaluate impact of anti-cancer therapies on normal humanhematopoietic cell function**

Our next objective was to determine to what extent the high-dose  $O<sup>6</sup>$ -BG/TMZ regimen hat effectively kills tumor cells may affect human hematopoiesis *in vivo*. NOD/SCID/γchainnull mice were sublethally irradiated and transplanted with  $1 \times 10^5$  human CD34<sup>+</sup> cells (Fig 3A). At 4 weeks post-transplant, a sample of PB was analyzed for the presence of human CD45<sup>+</sup> cells to confirm human cell engraftment (Fig. 3A and Table 1). Next, cohorts of mice were treated with vehicle or one cycle of  $O<sup>6</sup>$ -BG/TMZ. Bone marrow was harvested at 6 or 11 days post-treatment, and analyzed for the percent and total number of human versus mouse cells (Table 1). Our objective was to determine the maximum impact of the treatment regimen on human hematopoiesis. Therefore, we analyzed the bone marrow at 6 days posttreatment since we have previously found that the nadir in human myeloid progenitor

numbers is typically reached at day 6 following TMZ treatment *in vitro* (data not shown). In addition, the bone marrow was also analyzed at 11 days post-treatment in one experiment to assess marrow function and potential recovery after the nadir period. For both Experiments I and II, the fold decrease in the number of human  $CD45^+$  cells in vehicle versus  $O^6$ -BG/TMZ treatment and for human versus mouse chimerism following treatment, was statistically significant (Table 1,  $p < 0.001$ ). Although different base-line levels of human engraftment in the bone marrow were observed in Experiment I and II (see vehicle treated mice, Table 1), 2  $\times$  10<sup>5</sup> CD34<sup>+</sup> cells were transplanted per animal in both experiments. It has been our experience that the frequency of SCID-repopulating cells can vary widely amongst different cord blood products. In either situation (high or low human-cell engraftment), the drug sensitivity of the human bone-marrow cells was similar. In Experiment I, a 9-fold decrease in the number of human cells in the bone marrow was observed following  $O<sup>6</sup>$ -BG/TMZ treatment. In contrast, only a 1.5-fold decrease in the number of murine bone-marrow cells was observed in these same mice (Table 1). In addition, due to high levels of human engraftment in Experiment I, the spleen was also analyzed for human hematopoietic cells; significant decreases in human cells was observed in  $O<sup>6</sup>$ -BG/TMZ-treated mice compared to vehicle-treated mice (data not shown). In Experiment II, total human engraftment pretreatment was lower than in experiment I, which is not unexpected, since different cord blood products were used in each experiment (Table I, see %huCD45<sup>+</sup> cells in the PB pretreatment). This also enabled us to see if lower levels of human chimerism would result in a different toxicity profile (human versus mouse) from that observed in Experiment I. Differential sensitivity of human and mouse bone-marrow cells was observed at 6 and 11 days post-drug treatment, with the human cells still exhibiting a higher sensitivity to the drug treatment than the mouse cells at both time points. A 13-fold decrease in human cell chimerism and a 1.5 fold-decrease in mouse cell chimerism were observed at 6 days post-O<sup>6</sup>-BG/TMZ treatment. At day 11 post-treatment, a 3-fold decrease and a 1.2 fold-decrease was observed for human and mouse cell chimerism respectively (Table 1). In Experiment II, it is important to note that the level of human cells in the PB pre-treatment differed from the two cohorts used for the day 11 analysis (Table 1, see Experiment II-%huCD45<sup>+</sup> cells in the PB pre-treatment at day 11). Human cells were detected at higher levels in the mouse cohort selected for drug treatment versus the cohort selected for vehicle treatment. Even with higher percentage of human cells in the PB of the cohort selected for  $O<sup>6</sup>$ -BG/TMZ treatment, the fold-decrease in human versus mouse bone-marrow cells was still evident and statistically significant. These data collectively demonstrate that the human hematopoietic cells, even at different levels of overall human cell chimerism in the bone marrow, were significantly more sensitive to  $O<sup>6</sup>$ -BG/TMZ treatment *in vivo* than mouse bone-marrow cells.

The impact of treatment on multi-lineage differentiation typically found in the bone marrow of mice reconstituted with human cells was also evaluated (Figs. 3-5). The number of CD34+ cells significantly decreased following treatment (Fig. 3b). In contrast to vehicletreated mice, CFU assays indicated that human erythroid and myeloid progenitor cells were no longer present in the bone marrow from drug-treated mice (Fig. 3c). The CD34+ cell population was further analyzed for different B-cell CD19+ subsets typically found in humanized bone marrow (Fig. 4a  $\&$  4b). There was a significant decrease in the most primitive cells (CD34<sup>+</sup>, CD19<sup>-</sup>), B-cell progenitors (CD34<sup>+</sup>, CD19<sup>+</sup>), and more mature Blymphoid cells (CD34<sup>-</sup>, CD19<sup>+</sup>). In addition, a significant decrease in human CD33<sup>+</sup> myeloid cells was also evident (Fig. 5a & 5b). These studies indicate that differential sensitivities of mouse and human cells to cytotoxic dosing regimens are apparent both *in vivo* and *in vitro*, and that the regimen sensitivities of human versus mouse may be even more pronounced in the bone marrow.

#### **DISCUSSION**

Development of human xenograft models that focus on treatment efficacy as well as potential hematopoietic toxicities *in vivo* could yield crucial insights into how to best translate basic research findings to the clinic. The main dose-limiting toxicity in >50% of anti-cancer drugs used in the past 15 years is therapy-mediated myelosuppression (22). Therefore, pre-clinical predictive models of this adverse event in humans could help guide the selection of new and safer therapeutic agents for clinical testing. While the use of *in vitro* clonogenic assays and mice for initial testing and regimen development is a logical starting point, incorporation of models that mimic to a larger degree human response to multiple cycles of therapy could aid in screening for safer and effective lead compounds. It is well recognized that doses of a compound that prevent growth of tumor xenografts in mice may not be achievable in humans due to compound-induced myelotoxicity (2-5). Our data collectively demonstrate that the human hematopoietic cells were significantly more sensitive *in vivo* than mouse bone-marrow cells when mice previously transplanted with human CD34<sup>+</sup> cells received a high-dose regimen that can inhibit growth of human glioma xenografts. Although humanizing the bone-marrow compartment of NOD/SCID/ $\gamma$ chain<sup>null</sup> mice cannot necessarily take into account all pharmacological and inter-species differences in human and mouse, it can provide improved correlations of compound levels obtained *in vivo* versus regimen efficacy and human hematotoxicity.

We built our study based on the scientific questions observed from clinical observations in an effort to research improvements in therapy (e.g. from the bedside back to the bench). Middleton *et al* originally determined the pharmacokinetic profile of TMZ in plasma from adult patients diagnosed with malignant melanoma. In patients that received TMZ starting at 200 mg/m<sup>2</sup>/day, the range of peak TMZ levels (C<sub>max</sub>) was 0.1-10.0 μg/ml or approximately 0.5-50 μM in the plasma. (23) Similar TMZ plasma levels were also obtained in an adult glioma clinical study by Osterman *et al.* (24) Additionally, when 267 mg/m<sup>2</sup>/day was administered to pediatric brain tumor patients, levels of TMZ detected in the plasma ranged from 2.7-19.5 μg/ml or 14.0-100 μM which is similar to TMZ peak levels obtained in adults. (25) These TMZ dosing schedules have been incorporated into the standard-of-care protocols for GBM.(26) In the clonogenic assays performed in Figure 1, we used clinically relevant doses of TMZ (25-100 μM) that could be obtained in the plasma of patients, and demonstrated that human hematopoietic progenitor cells were significantly more sensitive than murine hematopoietic progenitors to  $O^6$ -BG in combination with TMZ.

We used a  $O<sup>6</sup>$ -BG/TMZ regimen clinically known to be toxic to the bone marrow, confirmed that it did indeed block tumor growth of ectopic human xenografts; and importantly simulated therapy-mediated toxicity in the bone marrow. An  $O<sup>6</sup>$ -BG/TMZ treatment regimen was recently tested in Phase II clinical trials in the absence of stem-cell rescue (9). In that study, 5 of 32 patients with TMZ-resistant anaplastic glioma responded to O<sup>6</sup>-BG in combination with TMZ. For TMZ-resistant glioblastoma multiforme, only one of 34 patients exhibited a response to treatment. The reason for lack of response in most patients is likely a consequence of the necessity to reduce the levels of TMZ administered due to myelosuppression; a 25% reduction in TMZ dose was necessary in 48% of patients due to therapy-induced grade-4 neutropenia and thrombocytopenia. To increase doses of  $O^6$ -BG/TMZ in these patients, myelosuppression will need to be overcome and this is an area of active investigation. Strategies that protect hematopoietic cells from high-dose  $O<sup>6</sup>$ -BG/TMZ such as viral vector-mediated gene transfer and expression of O<sup>6</sup>-BG-resistant mutant MGMT in autologous hematopoietic stem and progenitor cells (27), or strategies that incorporate stem-cell rescue following administration of  $O<sup>6</sup>$ -BG/TMZ, may be the only treatment strategies to alleviate  $O<sup>6</sup>$ -BG/TMZ-associated bone-marrow toxicity.

When NOD/SCID/γchain<sup>null</sup> mice transplanted with human CD34<sup>+</sup> cells were treated with one cycle of  $O<sup>6</sup>$ -BG/TMZ, significant differential toxicity of human versus mouse hematopoietic cells was observed. The reason for the differential sensitivities between mouse- and human-hematopoietic cells could be due in part to subtle differences in expression or in the amino acid structure of murine and human MGMT in the bone marrow cells. While the active site of murine and human MGMT is identical (Pro-Cys-His-Arg), differences in the MGMT amino acid sequence are evident between the two species. Gerson and colleagues demonstrated that, in contrast to human MGMT, murine MGMT has a leucine at residue 180. This amino acid residue is located just outside the active site of MGMT, and correlated with increased resistance of murine MGMT to  $O<sup>6</sup>$ -BG (28).

A variety of human cancers and diseases are currently being modeled in immunodeficient mouse strains such as NOD/SCID(29), NOD/SCID mice transgenic for human Stem-cell Factor, Granulocyte-macrophage-colony-stimulating factor, and Interleukin-3 (30-31), NOD/SCID/γchain<sup>null</sup> (32-38) and Rag2-/-γc-/- (39). Due to increased and consistent levels of human hematopoietic cell engraftment and an increased life span compared to NOD/ SCID mice, NOD/SCID/γchain<sup>null</sup> mice are beginning to replace NOD/SCID mice in studies that are testing transfer-vector designs and strategies for hematopoietic gene therapy (38). Additionally, development of pre-clinical humanized models that will allow for reconstitution of a mature human immune system will be instrumental in gaining a better understanding of *in vivo* human cell-mediated immune responses and for testing of novel immune therapy or chemotherapy that target the tumor and its microenvironment. Shultz *et al* recently demonstrated that HLA-restricted human immune responses can now be studied *in vivo* using NOD/SCID/γchain<sup>null</sup> mice that have been engineered to constitutively express HLA class I heavy and light chains.(40) The use of immunodeficient mouse models to study cancer progression and testing of anti-cancer therapies is still in its infancy. Integrated modeling approaches that investigate human tumor-cell growth in mice with humanized bone marrow capable of producing mature human immune cells will be instrumental in understanding the interplay between tumor progression, the tumor microenvironment, and the hematopoietic and immune systems. For example, our research team recently demonstrated that human circulating progenitor cells that are phenotypically and functionally distinct from circulating endothelial cells can promote growth of human melanoma xenografts in NOD/SCID and NOD/SCID/γchain<sup>null</sup> mice. (Mund, Case, and Pollok, unpublished observations and (41)

The engrafted human hematopoietic  $CD34<sup>+</sup>$  cells and progeny derived from the initial transplanted CD34+ cells interact at least to some extent with stromal cells and cytokines located in the murine bone-marrow microenvironment since the transplanted human CD34<sup>+</sup> cells differentiate into B-lymphoid and myeloid cells *in vivo*. Murine stromal feeder cells clearly interact with human hematopoietic cells, since numerous studies have used stroma of murine origin for *in vitro* expansion of primitive human hematopoietic cells. (42-45) Additionally, it is important to note that *in vivo* models in which recapitulation of the complete spectrum of human hematopoiesis can be accomplished have been developed. For example, implantation of human fetal thymus and fetal liver into SCID mice provided a human microenvironment which allowed for differentiation of human hematopoietic cells into multiple cell lineages.(46) An expansion of this model was done by Fraser *et al* in which SCID mice were transplanted subcutaneously with human fetal bone, thymus, and spleen fragments; this allowed for differentiation of human hematopoietic cells into multiple lineages and homing of cells to the thymus for further differentiation into mature T-cell subsets.(47) More recently, it has been demonstrated that human marrow mesenchymal stem/stromal cells (MSC) can organize a hematopoietic microenvironment with murine hematopoietic cells after transplantation into NOD/SCID mice. (48) It is possible that the greater drug susceptibility of the engrafted human hematopoietic cells compared to the

mouse hematopoietic cells is that the mouse bone-marrow microenvironment may not completely support human hematopoietic cell survival particularly when under chemotherapy-mediated stress. However, *in vitro* clonogenic data also indicated striking differences in drug sensitivity of murine and human progenitor cells. Interactions between human MSC and hematopoietic stem and progenitor cells are important for survival and differentiation *in vivo* for others have demonstrated that co-transplantation of human MSC and CD34+ cells into NOD/SCID mice enhanced engraftment of human myeloid cells in the bone marrow. (49) In addition, it is clear that the marrow microenvironment and MSC, as well as the hematopoietic stem and progenitor cells, are adversely affected by chemotherapy. (50) At the present time, however, it is not known to what extent the implantation of human MSC into the bone marrow of mice engrafted with human hematopoietic cells would increase or decrease the sensitivity of human hematopoietic cells to myelotoxic therapy. The humanized bone-marrow model described here serves as a starting point for these types of investigations. Our data indicate that differential sensitivities of mouse versus hematopoietic cells to  $O^6$ -BG in combination with TMZ exist *in vitro* (Fig. 1) and that these differences in drug sensitivity could be recapitulated in the *in vivo* environment (Table 1 and Figs 3-5). The humanized bone-marrow model described in our study could be used on a much larger scale for drug and regimen screening since only transplant of reasonable numbers of human CD34+ cells into sublethally irradiated NOD/ SCID/ychain<sup>null</sup> is required, and confirmation of human engraftment in the bone marrow can be done by analyzing a small sample of peripheral blood from the transplanted animals.

The humanized bone-marrow mouse model described here represents a feasible and physiologically relevant tool in which to determine if a balance between treatment efficacy and toxicity is achievable. In contrast to many other immunodeficient mouse strains, previous studies have shown that NOD/SCID/γchain<sup>null</sup> mice can be readily and consistently reconstituted with human hematopoietic cells that have the capacity to differentiate into various blood-cell lineages (6). In addition, we have found that human tumors grow consistently in NOD/SCID/γchain<sup>null</sup> mice, and represent a reliable model for testing anticancer strategies. In future studies, it will be essential to utilize orthotopic xenografts when new therapeutic regimens are tested, since the microenvironment in which the tumor resides could affect tumor sensitivity to treatment. (51) Integration of humanized hematopoietic and tumor models may be a way for anti-cancer therapies to be tested earlier in the drug development process which will ultimately lead to more efficient translation of therapies into clinical practice. While this model has the potential to be used as an *in vivo* screening tool that can aid in testing and refinement of treatment regimens, the validation of this model will require testing of an additional number of compounds and regimens to establish the model's predictive value. We are currently in the process of organizing a consortium of scientists to evaluate a broader set of therapeutics to determine the predictive clinical value of this modeling approach.

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#### **Figure 1. Determination of** *in vitro* **sensitivities to TMZ and combination O<sup>6</sup> -BG/TMZ**

(A) BALB/c, C57BL/6N, and NOD/SCID/γchainnull bone marrow cells were exposed to increasing doses of TMZ in the absence or presence of  $O<sup>6</sup>$ -BG. At days 10-12, hematopoietic colonies (human and mouse) were scored visually by inverted light microscopy. (B) Human CD34<sup>+</sup> cells, NOD/SCID/γchain<sup>null</sup> mice bone marrow cells, and SF767 tumor cells were exposed to increasing doses of TMZ in the absence or presence of O<sup>6</sup>-BG. At days 10-12, hematopoietic colonies (human and mouse) were enumerated. SF767 colonies were scored visually by counting methylene-blue stained colonies. p < 0.0001 for TMZ versus  $O<sup>6</sup>$ -BG/TMZ for all cell types, mouse versus human for  $O<sup>6</sup>$ -BG/TMZ, mouse versus tumor for  $O^6$ -BG/TMZ at all TMZ concentrations, human versus tumor for  $O^6$ -BG/ TMZ at all TMZ concentrations except 200 μM TMZ. HC = hematopoietic cells.





(A) Development of  $O^6$ -BG/TMZ myeloablative dosing-regimen. Mice (n = 4-8 mice per group) were injected ip with vehicle,  $O^6$ -BG (30 mg/kg double bolus), TMZ (80 mg/kg) or O<sup>6</sup>-BG/TMZ (40-60 mg/kg). Kaplan-Meier survival plot of data compiled from 2independent experiments. (B) Schema showing how NOD/SCID/γchainnull mice with SF767 xenografts ( $n = 8-10$  mice per group) were treated with double-bolus  $O<sup>6</sup>$ -BG/TMZ and stemcell rescue. Two cycles of treatment were given 2 weeks apart. (C) Tumor volumes were determined via caliper measurements. (D) Mice were euthanized and harvested tumors

weighed. These data are representative of 2-independent experiments with similar results. \*p < 0.05, vehicle versus drug-treated (tumor volumes and tumor weight).



**Figure 3. Human toxicity model to assess impact of treatment on human bone-marrow cells** (A) Human CD34+ cells were isolated from umbilical cord blood and transplanted into sublethally irradiated NOD/SCID/γchain<sup>null</sup> mice. At 4 weeks post-transplant, a sample of PB was analyzed from each mouse to confirm that all mice were similarly engrafted. Cohorts of mice received vehicle or one cycle of the  $O<sup>6</sup>$ -BG/TMZ regimen. At 6-11 days post-treatment, the bone marrow was harvested and analyzed via flow cytometry for the presence of human cells. (B) At 6 days post-treatment, the bone marrow was harvested, counted, and analyzed via flow cytometry to determine the number of human CD34+ cells in the bone marrow from vehicle and  $O^6$ -BG/TMZ-treated mice. \*p < 0.001 for vehicle versus drug-treated mice. (C) The bone marrow from vehicle and  $O<sup>6</sup>$ -BG/TMZ-treated mice was

analyzed for the presence of human colony-forming unit cells and analyzed 12 days after plating. \*p < 0.001 for vehicle versus drug-treated mice.



**Figure 4. Analysis of human B-lymphoid cells in the bone marrow of NOD/SCID/γchainnull mice following O<sup>6</sup> -BG/TMZ treatment**

(A) Bone marrow from vehicle and  $O<sup>6</sup>$ -BG/TMZ-treated mice was analyzed for the presence of  $CD34<sup>+</sup>$  and  $CD19<sup>+</sup>$  cells via flow cytometry. (B) Based on flow cytometry and bonemarrow cell numbers, the number of human CD34<sup>+</sup>CD19<sup>-</sup> (progenitors), CD34<sup>+</sup>CD19<sup>+</sup> (Blymphoid progenitors), and CD34-CD19+ (more mature B-lymphoid cells) were determined by flow cytometry. These data are representative examples of data from Experiment I. Data are similar from Experiment II except the level of overall human chimerism was lower in vehicle and  $O^6$ -BG/TMZ-treated mice. \*p < 0.0001 for vehicle versus drug-treated mice.



**Figure 5. Analysis of human myeloid cells in the bone marrow of NOD/SCID/γchainnull mice following O<sup>6</sup> -BG/TMZ treatment**

(A) At  $\vec{6}$  days post-treatment, bone marrow from vehicle and  $O^6$ -BG/TMZ-treated mice was analyzed for the presence of human  $CD45<sup>+</sup>$  and  $CD33<sup>+</sup>$  cells via flow cytometry. (B) Based on flow cytometry and bone-marrow cell numbers, the number of human CD33<sup>+</sup> myeloid cells was determined. These data are representative examples of data from Experiment I. Data are similar from Experiment II except the level of overall human chimerism was lower in vehicle and  $O^6$ -BG/TMZ-treated mice. \*p < 0.0001 for vehicle versus drug-treated mice.



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# **Table 1**

Analysis of human versus mouse bone-marrow cells post-treatment. Analysis of human versus mouse bone-marrow cells post-treatment.



Mice were transplanted with human CD34<sup>+</sup> cells and the peripheral blood (PB) analyzed at one month post-transplantation for the presence of human CD45<sup>+</sup> cells. This analysis was done prior to O<sup>6</sup>-BG/  $^+$  cells. This analysis was done prior to  $O^6$ -BG/ + cells and the peripheral blood (PB) analyzed at one month post-transplantation for the presence of human CD45 Mice were transplanted with human CD34 TMZ treatment. TMZ treatment.

 $^{\dagger}$ At 6-11 days post-O<sup>G</sup>-BG/TMZ treatment, the bone marrow was harvested and the percent human-cell chimerism (huCD45<sup>+</sup>) was determined by flow cytometry. +) was determined by flow cytometry. *†*At 6-11 days post-O6-BG/TMZ treatment, the bone marrow was harvested and the percent human-cell chimerism (huCD45

 $t$ The total number of human and mouse bone marrow cells in vehicle- and O<sup>6</sup>-BG/TMZ-treated mice was determined using a coulter counter. *‡*The total number of human and mouse bone marrow cells in vehicle- and O6-BG/TMZ-treated mice was determined using a coulter counter.

 ${}^8$ Total number of human cells = total number bone marrow cells  $\times$ % human cells. Total number of murine cells = total number bone marrow cells - total number human cells. *§*Total number of human cells = total number bone marrow cells × % human cells. Total number of murine cells = total number bone marrow cells – total number human cells.

\*\*<br>The fold decrease between vehicle-treated and drug-treated human or mouse bone marrow cells = total number of human or mouse movement of human or mouse cells The fold decrease between vehicle-treated and drug-treated human or mouse bone marrow cells = total number of human or mouse in vehicle-treated mice ÷ total number of human or mouse cells remaining in O<sup>6</sup>-BG/TMZ-treated mice. remaining in O<sup>6</sup>-BG/TMZ-treated mice.