The Role of Cell-mediated Cytotoxicity in Acute GVHD after MHC-Matched Allogeneic Bone Marrow Transplantation in Mice

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Summary

The role of cell-mediated cytotoxicity in the complex pathophysiology of graft-versus-host disease (GVHD) has remained poorly defined for several decades. We transplanted T cells from Fas-ligand (FasL)-defective and perforin-deficient mutant donor mice into lethally irradiated MHC-matched allogeneic recipient mice to characterize the role of cell-mediated cytotoxicity in GVHD. Although recipients of allogeneic FasL-defective donor T cells underwent severe GVHD-associated cachexia, they exhibited only minimal signs of hepatic and cutaneous GVHD pathology. Recipients of perforin-deficient allogeneic donor T cells developed signs of acute GVHD, but the time of onset was significantly delayed. These findings demonstrate that Fas-mediated anti-recipient cytotoxicity may be critical for the development of hepatic and cutaneous GVHD, but is not required for GVHD-associated cachexia. In addition, perforin-mediated anti-recipient cytotoxicity appears to play an important role in the kinetics of GVHD pathophysiology, but is not required for GVHD-associated tissue damage.

Allogeneic bone marrow transplantation (BMT) has greatly expanded as a clinical treatment modality for several disorders of hematopoiesis and certain hematological malignancies (1). Graft-versus-host disease (GVHD) remains a principal complication following allogeneic BMT occurring in up to 75% of recipients of unmanipulated HLA-matched marrow (2). The immunopathophysiology of GVHD is complex, and is generally considered to involve two phases: an afferent (inductive) phase, and an efferent (effector) phase (3). In the afferent phase, mature T cells present in the donor marrow inoculum recognize antigenic disparities expressed on recipient tissues resulting in alloactivation and proliferation of the allogeneic donor T cells. In the efferent phase, inflammatory reactions may develop in specific host target tissues such as skin, liver, and gastrointestinal tract that are characterized by mononuclear cell infiltration and histopathological damage (4). Studies of experimental models of allogeneic BMT using T cell-depletion have demonstrated that mature T cells must be present in the donor marrow inoculum in order to induce GVHD, and several clinical studies have confirmed this finding (5, 6). However, the precise role of T cell-mediated anti-recipient cytotoxicity in the pathophysiology of GVHD remains controversial (7–13).

Recently, it has been demonstrated that perforin-dependent cytolysis and Fas-mediated apoptosis together constitute the major mechanisms of short-term T cell-mediated cytotoxicity (14–16). T cells from mice which are homozygous for the gld (generalized lymphoproliferative disease) mutation are known to express a functionally defective Fas ligand (FasL) molecule which is not capable of transducing an apoptotic signal to Fas-bearing target cells (17). The perforin-deficient (perforin 0/0) mutant mouse strain was developed by homologous recombination in a B6 embryonic stem cell line, and bred to homozygosity on a C57BL/6 background (18). We have transplanted T cells from FasL-defective (gld) and perforin-deficient (perforin 0/0) donor mice into lethally irradiated MHC-matched allogeneic recipient mice to determine the role of these cytotoxic pathways in acute GVHD.

The present studies demonstrate that Fas-mediated cytotoxicity plays an essential role in the pathophysiology of hepatic and cutaneous GVHD, but is not required for GVH-induced cachexia. Furthermore, the absence of perforin-mediated cytotoxicity significantly delays the onset of GVHD, but does not prevent or diminish GVHD. These results suggest that certain local processes of GVHD such as tissue damage are separable from the systemic process of cachexia.
In addition, the Fas-mediated and perforin-dependent cytotoxic pathways appear to act during different stages in the complex pathophysiology of GVHD. The role of inflammatory cytokines is discussed.

**Materials and Methods**

*Mice.* C57BL/6J (H-2b), B6.SmnC3H-gld (H-2b), LP/J [H-2b], and C3H.SW (H-2b) mice were obtained from Jackson Laboratory (Bar Harbor, ME). Perforin-deficient C57BL/6 mice (B6-perforin 0/0) were obtained from D. Kägi and H. Hengartner (University of Zürich, Switzerland) and B. Ledermann and K. Bürki (Sandoz Pharma Ltd., Basel, Switzerland). B6-perforin 0/0 mice were propagated at the University of Miami School of Medicine Specialized Animal Facility and were maintained in a pathogen-free colony until use.

**Preparation of Cells.** Bone marrow cells were aspirated from the femurs and tibias of donor mice. T cells were depleted from the bone marrow by incubation with anti-Thy1.2 mAb (30- H-12 culture supernatant) at 1:5 dilution and 4°C for 30 min followed by low-Tox M complement (Accurate Chemical Co., Westbury, NY) at 1:20 dilution and 37°C for 45 min. Spleen and lymph node cells were harvested, pooled, and treated with anti-B220 mAb (14/8 culture supernatant) at 1:2.5 dilution and 4°C for 30 min, followed by a secondary mouse anti-rat mAb (18.5 ascites) at 1:50 and 4°C for an additional 30 min. The labeled cells were then treated with rabbit complement at 1:10 dilution and 37°C for 45 min to remove B cells and enrich for T cells. This procedure routinely enriches the T cell population to levels of purity between 75-80% as determined by flow cytometric analysis.

**Assay for GVHD.** Recipient mice were exposed to 900 cGy TBI from a 46Co source at a dose rate of 50 cGy/min 24 h before the BMT. In murine models of GVHD, precise numbers of mature allogeneic donor T cells are routinely added together with bone marrow to induce lethal acute GVHD. The severity of GVHD correlates directly with the number of donor T cells transplanted (5). The number of donor T cells was selected based upon the ability of the wild-type inoculum to induce characteristic signs of severe acute GVHD with an incidence of 100% in either donor/recipient strain combination. We transplanted 1 × 10⁷ donor T cells in the B6→LP combination, and 2 × 10⁷ donor T cells in the B6→C3H.SW combination. To ensure reproducible results, the number of CD3⁺ T cells in the donor inoculum was precisely quantified by flow cytometry for each BMT. Precisely the same numbers of CD3⁺ T cells from wild-type, perforin-deficient, or FasL-defective B6 donors were added to the T cell-depleted bone marrow cells (5-10 × 10⁶) from normal wild-type B6 donor mice and injected together into irradiated recipient mice intravenously via the lateral tail vein in a volume of 0.5 cc. Mice receiving transplants were distributed into groups containing 4-8 mice per group in each experiment. Recipients were maintained on acidified water (pH 3.0) containing antibiotics (100 µg/ml neomycin sulfate, 10 mg/L polymyxin B) from day -3 to day 14 post-BMT. Recipient mice were monitored for clinical signs of GVHD including weight loss, skin lesions, alopecia, diarrhea, hunched posture, and mortality. Representative mice were killed at various times post-BMT to harvest tissues for histopathological analysis.

**Histopathology.** Skin and liver sections were harvested from recipients at various intervals after BMT. Tissues were placed in 10% buffered formalin phosphate (Fisher Scientific, Orlando, FL). The fixed tissues were paraffin embedded, sectioned, and stained with hematoxylin and eosin by the core service of the Department of Comparative Pathology at the University of Miami School of Medicine. Slides were coded and examined in a blinded fashion by NHA, and the tissue histology was graded.

**Immunophenotyping.** Pooled spleen and lymph node cells were stained with FITC-conjugated anti-CD3 mAb (145–2C11) or biotinylated anti-B220 mAb (RA3-6B2) obtained from PharMingen (San Diego, CA). Briefly, 0.5–1 × 10⁷ cells were washed in FACS® buffer (PBS with 1% BSA, 0.02% sodium azide), then incubated with the FITC-conjugated and biotinylated mAbs for 20 min on ice. Cells were again washed in FACS® buffer and incubated with streptavidin-PE (Fisher Scientific, Orlando, FL). The cells were then resuspended in FACS® buffer at 2 × 10⁶/ml and analyzed on a FACScan® flow cytometer (Beckton Dickinson, San Jose, CA). Data was analyzed within a gate established for lymphocytes using forward (180°) and side (90°) angle light scatter.

**Lymphocyte Stimulation Assay.** The enriched donor T cell populations were cultured 2 d in the presence of either soluble anti-CD3 mAb (2C11–145 culture supernatant) at 10% (vol/vol) or 5 µg/ml concanavalin A (Sigma Chemical Co., St. Louis, MO) to induce polyclonal activation. Mixed lymphocyte reactions were performed in which the enriched donor T cells were cultured 5 d at a 1:1 ratio with irradiated whole C3H (H-2k) spleen cells to assess alloseactivity. Proliferative responses were determined by measuring [3H]thymidine incorporation following a 6–8-h pulse label period.

**Results**

**Donor T Cells from B6-gld Mice Are Phenotypically and Functionally Normal.** Mice homozygous for the gld mutation develop lymphadenopathy characterized by progressive accumulation of functionally anergic B220⁺, CD4⁻, CD8⁻ double negative (DN) T cells with significant numbers appearing after 6 wk of age (19, 20). To ensure that we were not transplanting significant numbers of non-functional B220⁺ DN T cells, we used B6-gld mice which were 5–6 wk of age as donors. In addition, all detectable DN T cells and most B cells were removed by treating spleen and lymph node cells with anti-B220 mAb (14.8) and complement that concurrently enriched the phenotypically normal T cells (Fig. 1A). Notably, the RA3-6B2 anti-B220 mAb used for immunophenotyping recognizes an epitope that is distinct from the epitope recognized by the 14.8 anti-B220 mAb, and binding of 14.8 does not block subsequent binding of RA3-6B2 (21). Syngeneic donor mice, wild-type allogeneic donor mice, and perforin-deficient donor mice were 6–8 wk of age. All donor T cell populations were prepared in the same fashion, and exhibited similar cellular subpopulations as determined by phenotypic analysis (data not shown).

To confirm that the enriched T cell population purified from the gld donor mice was functionally competent, these cells were cultured in vitro in the presence of either soluble anti-CD3 mAb (2C11–145), concanavalin A, or irradiated allogeneic (C3H) spleen cells. The proliferative responses of the gld T cells to these polyclonal and alloantigen-specific stimuli was within normal limits compared to wild-type B6 T cells (Fig. 1B and C).

**Marked Reduction of Rash but High Incidence of Cachexia and Mortality after Transplantation of FasL-Defective Allogeneic T Cells.** Fig. 2 shows results from one representative trans-
Figure 1. Phenotypic and functional analysis of B6-gld T cells. (A) Flow cytometric analysis of pooled spleen and lymph node cells from wild-type B6 and B6-gld donor mice before and after treatment with anti-B220 (14.8) mAb and complement. The abnormal CD3+B220+ lymphocytes represent only a small percentage (3.1%) of spleen and lymph node cells from B6-gld donor mice before treatment, and these cells are absent after treatment. The CD3+B220− T cell populations are enriched following this procedure. (B) Polyclonal activation of enriched donor T cell populations. The normal B6 and B6-gld T cells were cultured 2 d in the presence of either soluble anti-CD3 (2Cl1-145) mAb culture supernatant at 10% (vol/vol) or 5 µg/ml concanavalin A. Proliferative responses by both T cell populations were within normal limits. Data are shown from a single representative experiment, and columns represent mean incorporation values among quadruplicate wells. (C) Mixed lymphocyte reaction by the T cell-enriched donor lymphocyte populations. The B6 and B6-gld donor T cells were cultured 5 d at a 1:1 ratio with irradiated C3H (H-2k) spleen cells as a source of allograft. Both T cell populations responded similarly to allogeneic stimulation.
Figure 2. Body weight and mortality during acute GVHD after allogeneic bone marrow transplantation of cytotoxicity-defective T cells. Lethally irradiated LP recipient mice (seven per group) were transplanted with $1 \times 10^7$ T cells from either syngeneic (LP), wild-type allogeneic (B6), or FasL-defective allogeneic (B6-gld) donor mice. After BMT, recipients were monitored for mortality (A) and weight loss (B). Mice receiving either wild-type B6 or B6-gld T cells developed acute GVHD with 100% mortality and severe weight loss, while recipients of syngeneic T cells and marrow survived and rapidly regained normal body weight.

Independent experiments confirm an absence of clinical cutaneous GVHD (0/32) despite the presence of cachexia (32/32) following BMT with FasL-defective donor T cells (Table 1).

Marked Delay of GVHD and Mortality after Transplantation of Perforin-Deficient Allogeneic T Cells. An MHC-matched model of allogeneic BMT was employed to determine whether perforin-deficient T cells could induce severe acute GVHD across minor histocompatibility barriers. Lethally irradiated LP (H-2b) mice were transplanted with $1 \times 10^7$ T cells and bone marrow from wild-type or perforin-deficient B6 (H-2b) donor mice. Recipients of syngeneic (LP) T cells maintained normal body weight and did not develop signs of GVHD (Figs. 4 and 5, A). The mice receiving wild-type B6 T cells developed signs of severe acute GVHD with 28 d after the transplant (Figs. 4 and 5, B). The recipients of T cells from B6-perforin 0/0 donors also developed signs of severe acute GVHD including weight loss (Fig. 4 B), alopecia, hunched posture, diarrhea, and desquamative skin rash with an incidence of 100% (Fig. 5 C). However, the average time of onset of these signs was delayed approximately twofold compared to the recipients of wild-type cells. The delay of onset in the recipients of perforin-deficient T cells was also apparent in the kinetics of mortality. There was a more than twofold increase in the mean survival time (65.3 vs 23.7 d) in the perforin-deficient recipient group compared to the wild-type recipient group (Fig. 4 A). Results from seven independent experiments using either C3H.SW or LP strain mice as recipients confirmed that transplantation of perforin-deficient allogeneic
Recipient mice were examined thrice weekly for clinical signs of acute GVHD including cachexia, alopecia, and desquamative rash. Results are expressed as the number of mice exhibiting signs compared to the total number of mice examined. Notably, the recipients of FasL-defective B6-gld T cells all developed cachexia, but none exhibited any signs of alopecia or rash throughout the study.

T cells uniformly produces acute GVHD with a significant delay in time of onset (Table 1).

Cutaneous GVHD Is Markedly Diminished in the Absence of FasL-mediated but Not Perforin-mediated Anti-Recipient Cytotoxicity. Skin sections were harvested from representative mice at 28 d post-transplant to determine if the absence of perforin or Fas-mediated anti-recipient cytotoxicity would effect the incidence or severity of GVHD-associated pathology. Cutaneous sections from recipients of syngeneic T cells were unremarkable (Fig. 6 A). Skin sections from C3H.SW recipients of 2 × 10^7 wild-type B6 T cells exhibited severe inflammation with mononuclear cell infiltrates, dermal fibrosis, loss of hair follicles, and epidermal hypertrophy consistent with cutaneous GVHD (Fig. 6 B). In marked contrast, skin from C3H.SW recipients of B6-gld T cells exhibited evidence of only minimal inflammation (Fig. 6 C). On day 52, cutaneous sections from recipients of perforin-deficient B6 T cells exhibited alterations that were identical in character and severity to those observed in recipients of wild-type B6 T cells on day 28 (Fig. 6 D).

A second MHC-matched donor/recipient strain combination (B6→LP) was employed to confirm this differential pattern of tissue damage after transplantation of cytotoxically defective T cells. These studies also demonstrated that cutaneous GVHD develops in the absence of perforin-mediated anti-recipient cytotoxicity, but is markedly reduced in the absence of FasL-mediated anti-recipient cytotoxicity (data not shown).

Hepatic GVHD Is Markedly Diminished in the Absence of FasL-mediated but Not Perforin-mediated Anti-Recipient Cytotoxicity. Liver sections were harvested from recipients at various time intervals following transplantation to investigate how the absence of Fas and perforin-mediated cytotoxicity affected the pathophysiology of hepatic GVHD. Sections of liver from C3H.SW recipients of syngeneic T cells 28 d post-transplant were unremarkable (Fig. 7 A). Hepatic sections from recipients of 2 × 10^7 wild-type B6 T cells exhibited a marked infiltrate of predominantly PMNs around bile ducts with associated partial inflammation and fibrosis that was evaluated as severe acute cholangitis consistent with hepatic GVHD (Fig. 7 B). In marked contrast, liver sections from C3H.SW recipients of B6-gld T cells exhibited evidence of only minimal involvement (Fig. 7 C). It is important to note that at no time after transplant of B6-gld T cells did the liver sections of these recipients exhibit significant hepatic pathology. On day 52, hepatic tissues from recipients of perforin-deficient B6 T cells exhibited severe cholangiohepatitis (Fig. 7 D). A second MHC-matched donor/recipient strain combination (B6→LP) was employed to confirm this differential pattern of tissue damage.
after transplantation of the cytotoxically defective T cells. These studies also demonstrated that hepatic GVHD develops in the absence of perforin-mediated anti-recipient cytotoxicity, but is markedly reduced in the absence of FasL-mediated anti-recipient cytotoxicity (data not shown).

There is evidence in the literature that epithelial damage in gastrointestinal GVHD lesions is characterized by apoptosis and DNA fragmentation (22). Therefore, we were interested to examine the role of Fas-mediated cytotoxicity in gastrointestinal GVHD. Clinical evidence of diarrhea was observed in some but not all recipients of wild-type, FasL-defective, and perforin-deficient allogeneic T cells. In addition, histopathological evidence of acute colitis and enteritis was occasionally observed in recipients of wild-type, FasL-defective, and perforin-deficient allogeneic T cells (data not shown). Therefore, defects in individual cytotoxic pathways did not consistently abrogate either clinical or histopathological sequelae associated with gastrointestinal GVHD.

**Recipients of FasL-defective Allogeneic T Cells Do Not Exhibit Lymphoid Atrophy.** Recipient splenic lymphocytes were harvested and analyzed by flow cytometry 28 d after transplant to examine the status of the lymphoid compartment. Table 2 shows the numbers of spleen cells recovered from these recipients. Mice receiving wild-type or perforin-deficient allogeneic T cells exhibited profoundly decreased spleen cell recoveries that are typically observed in murine models of acute GVHD (Table 2). Notably, recipients of FasL-defective T cells exhibited a somewhat increased spleen cell recovery, but did not exhibit evidence of uncontrolled lymphoproliferative disease (Table 2). These findings have been observed in seven independent transplantation experiments.

Mice that received syngeneic (C3H.SW) T cells exhibited normal percentages of B220+ cells (54.0%) and CD3+ cells (23.1%) in the spleen (Fig. 8 E). Light scatter profiles of spleen cells from recipients of wild-type allogeneic T cells (Fig. 8 F) showed a markedly reduced lymphoid cell population (Region 1), which was comprised of 71.3% CD3+ T cells (Fig. 8 F). This pattern is consistent with acute GVHD (23). A very similar pattern of lymphoid hypoplasia and spleen cell phenotype was observed in spleens from recipients of perforin-deficient allogeneic T cells in which 81.2% of the gated cells were CD3+ (Fig. 8 C and G). In contrast, the degree of lymphoid hypoplasia was less severe in the recipients of FasL-defective T cells as determined by light scatter (Fig. 8 D). In addition, the percentage of B220+ cells was increased to 28.0% in the recipients of FasL-defective T cells (Fig. 8 H) compared to 2.6% in the recipients of wild-type T cells. However, as expected the percentage of CD3+ T cells (54.2%) was elevated in the recipients of FasL-defective T cells consistent with a GVHD-associated expansion of alloreactive donor T cells.

**Discussion**

The role of cell-mediated anti-recipient cytotoxicity in the pathophysiology of GVHD has remained poorly defined and controversial for several decades. To begin resolving the long standing controversy regarding the role of anti-recipient specific cell-mediated cytotoxicity in GVHD, we have compared the ability of FasL-defective, perforin-deficient, and wild-type allogeneic T cells to induce severe acute GVHD across non-MHC antigenic barriers in mice. The results of the present study demonstrate that Fas-mediated cytotoxicity plays an important role in the pathophysiology of hepatic and cutaneous GVHD after BMT between MHC-matched allogeneic mice. However, Fas-mediated anti-recipient cytotoxicity is not required for the induction of GVHD-associated cachexia. Furthermore, transplantation of perforin-
deficient T cells results in a marked delay in onset of GVHD
and mortality compared to GVHD induced by wild-type
allogeneic T cells. However, severe acute GVHD ultimately
does develop with all of the classical hallmarks of GVHD
including hunched posture, weight loss, alopecia, rash, cho-
langitis, and dermatitis.

The morphological and ultrastructural features of lesions
in hepatic, cutaneous, and gastrointestinal GVHD have pre-
viously been reported to involve some degree of individual
cell necrosis, pyknotic nuclei, and apoptotic bodies consist-
tent with a role for apoptosis in GVHD-associated tissue
damage (24–26). In addition, both liver and skin tissues are
known to express Fas (27), and mice injected with the anti-
Fas monoclonal antibody Jo2 rapidly develop a fulminant
lethal hepatitis (28). Furthermore, it has been reported re-
cently that expression of Fas is upregulated in epidermal tis-
sues undergoing certain inflammatory conditions (29). Thus,
skin and liver could be potential targets for FasL-bearing
cytotoxic donor T cells after allogeneic BMT. Consistent
with our findings that GVHD-associated cholangitis and
dermatitis is markedly reduced in the absence of Fas-medi-
ated cytotoxicity, we propose that this pathway may be an
essential step in the multi-step process that ultimately leads
to GVHD-associated inflammation and tissue damage in

Figure 6. Histopathological analysis of skin from C3H.SW recipient mice 28 d after BMT with 2 × 10^7 T cells. Representative recipient mice were
sacrificed and skin sections were harvested for histopathological analysis. (A) Skin from recipients of syngeneic (C3H.SW) T cells were unremarkable. (B)
Analysis of skin from recipients of wild-type allogeneic (B6) T cells exhibited severe inflammatory infiltrates, dermal fibrosis, loss of hair follicles, and epider-
dermal hypertrophy consistent with cutaneous GVHD. (C) Cutaneous sections from recipients of FasL-defective allogeneic (B6-gld) T cells exhibited ev-
idence of minimal inflammatory changes or no involvement. (D) Skin from recipients of perforin-deficient allogeneic (B6-perforin 0/0) T cells on day 52
post-transplant exhibited severe inflammatory infiltrates, dermal fibrosis, loss of hair follicles, and epidermal hypertrophy consistent with cutaneous
GVHD.
recipient liver and skin tissues. However, the findings in the present study do not suggest that Fas-mediated cytotoxicity is the only effector mechanism contributing to hepatic and cutaneous GVHD.

While hepatic and cutaneous pathology was markedly reduced in recipients of FasL-defective donor T cells, these recipients exhibited severe cachexia and lost on average 35% of their initial body weight and virtually all body fat by day 28 post-BMT. Tumor necrosis factor (TNF-α, cachectin) produced by alloactivated FasL-defective donor T cells could account for the profound wasting and mortality observed in the recipients of the B6-gld T cells. TNF-α has been identified as a principal mediator of cachexia in rodents (30). In addition, serum levels of TNF have been shown to be increased in patients undergoing GVHD after allogeneic BMT (31). Notably, Piguet et al. demonstrated that administration of anti-TNF-α anti-sera markedly reduces recipient weight loss and mortality in a mouse model of GVHD (32).

Despite significant work, it has remained difficult to reconcile the classical concept of MHC-restricted T cell-mediated cytotoxicity with the observation that highly purified T cell subset populations (CD4+ or CD8+) are equally capable of inducing identical GVHD pathology irrespective of class I or class II expression by the target tissues (33). Consistent with our results, one interpretation of the finding that either subset alone can induce identical

Figure 7. Histopathological analysis of liver from C3H.SW recipient mice 28 d after BMT with 2 × 10^7 T cells. Representative recipient mice were killed and liver sections were harvested for histopathological analysis. (A) Liver sections from recipients of syngeneic (C3H.SW) T cells were unremarkable. (B) Analysis of liver from recipients of wild-type allogeneic (B6) T cells demonstrated severe subacute cholangiohepatitis consistent with hepatic GVHD. (C) Hepatic sections from recipients of FasL-defective allogeneic (B6-gld) T cells exhibited evidence of minimal inflammatory changes or no involvement. (D) Liver sections from recipients of perforin-deficient allogeneic (B6-perforin 0/0) T cells on day 52 post-transplant exhibited severe subacute cholangiohepatitis consistent with hepatic GVHD.

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Figure 8. Light scatter and phenotypic analysis of spleen cells from C3H.SW recipient mice 28 d after transplantation of 2 \times 10^7 T cells. Spleen cells were stained with anti-CD3-FITC (FL1) and anti-B220-biotin + avidin-PE (FL2), and analyzed within a gate established for lymphocytes using forward (FSC) and side (SSC) angle light scatter. (A and E) Spleen cells from recipients of syngeneic (C3H.SW) T cells exhibited a normal light scatter and phenotypic pattern of CD3+ and B220+ cells. (B and F) Analysis of spleen from recipients of wild-type allogeneic (B6) T cells demonstrated severe lymphoid hypoplasia and a predominance of CD3+ T cells. (C and G) Spleen cells from recipients of perforin-deficient allogeneic (B6-perforin 0/0) T cells exhibited a pattern of lymphoid hypoplasia and predominance of CD3+ T cells that was similar to the recipients of wild-type allogeneic T cells. (D and H) Analysis of spleen cells from recipients of FasL-defective allogeneic (B6-gld) T cells demonstrated less severe lymphoid hypoplasia (R1) and increased numbers of B220+ cells compared to recipients of wild-type allogeneic T cells.

GVHD pathology is that cytotoxic T cells (CTL) of either phenotype may effect allogeneic cytotoxicity via FasL. Notably, it has recently been reported that CD4+ CTL may kill primarily through the Fas pathway (34).

It is important to note that we have observed no evidence of lpr-GVH or lymphoproliferative disease in recipients of B6-gld donor T cells. These recipients did not develop lpr-GVH-associated skin or liver inflammation, nor did they develop lpr-GVH-associated lymphoid aplasia (Figs. 6–8). The B6-gld T cells that were transplanted have a defect only in the Fas-ligand molecule resulting in an inability to kill through the Fas pathway. However, the B6-gld T cells express normal levels of functional Fas antigen, and therefore have the potential to be regulated or deleted by cells of host origin such as stromal cells that may be capable of expressing functional FasL (35). Thus, our observations are consistent with the established finding that transplantation of bone marrow from gld/gld mutant donors into wild-type recipients does not induce the lpr-GVH phenomenon (36). Moreover, the transplantation of wild-type B6 marrow in our studies results in the de novo generation of T cells that are able to express normal non-mutant FasL molecules (Baker, M.B. and R.B. Levy, manuscript in preparation). Therefore, wild-type donor marrow-derived T cells that express the non-mutant FasL could also regulate the gld T cells that express a functional Fas molecule.

The finding that recipients of perforin-deficient allogeneic T cells exhibited clinical and histopathological signs of GVHD equivalent in severity to the signs observed in recipients of wild-type allogeneic T cells conclusively demonstrates that perforin-mediated cytotoxicity is not a critical effector function during the effector phase of GVHD.

Table 2. Spleen Cell Recoveries from C3H.SW Mice 28 d after Receiving 2 \times 10^7 Donor T Cells

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<thead>
<tr>
<th>Spleen Cell Recovery</th>
<th>CD3+ B220-</th>
<th>B220+ CD3-</th>
<th>CD3+ B220+</th>
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<tr>
<td>C3H.SW→C3H.SW</td>
<td>41.3 \times 10^6</td>
<td>9.5 \times 10^6</td>
<td>22.3 \times 10^6</td>
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<tr>
<td>B6→C3H.SW</td>
<td>6.9 \times 10^6</td>
<td>4.9 \times 10^6</td>
<td>0.18 \times 10^6</td>
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<tr>
<td>B6-perforin 0/0→C3H.SW</td>
<td>5.2 \times 10^6</td>
<td>4.2 \times 10^6</td>
<td>0.14 \times 10^6</td>
</tr>
<tr>
<td>B6-gld→C3H.SW</td>
<td>12.6 \times 10^6</td>
<td>6.8 \times 10^6</td>
<td>3.5 \times 10^6</td>
</tr>
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Lymphocyte subpopulation numbers are calculated based on percentages derived from phenotypic analysis (see Fig. 8). Significantly increased numbers of phenotypically normal B cells (B220+CD3-) are recovered from recipients of FasL-defective allogeneic T cells compared to recipients of wild-type and perforin-deficient allogeneic T cells. Notably, mice receiving B6-gld T cells did not exhibit an abnormally expanded population of B220+ DN T cells (B220+CD3+).
We conclude that perforin-mediated donor anti-host cytotoxic function is not an absolute requirement for the development of GVHD across minor histocompatibility barriers.

While perforin-deficient allogeneic T cells are clearly capable of inducing severe acute GVHD, the recipients of these cells have consistently exhibited a significant delay in the time of onset of clinical signs associated with GVHD and a prolonged MST. This finding suggests that perforin-mediated cytotoxicity may be playing an important role in the early post-transplant period during the afferent phase of GVHD that could lead to a shift in kinetics without diminishing the ultimate severity of tissue damage and clinical signs of GVHD. Notably, preliminary results in our laboratory suggest that transplantation of perforin-deficient T cells at a dose twofold higher than the wild-type T cells results in onset of GVHD with no delay compared to the recipients of the lower dose of normal cells. In addition, we have found that when very low numbers of bone marrow-derived perforin-deficient T cells are transplanted the onset of GVHD is delayed indefinitely (37).

One potential explanation for the delayed onset of GVHD in recipients of perforin-deficient T cells is that perforin-mediated cytotoxicity may accelerate and amplify the donor anti-host immune reaction. This may occur as a result of the release of inflammatory cytoplasmic contents following cytolysis of host target cells. When perforin-mediated anti-host cytolytic activity is absent, the resulting allogeneic reaction might be slowed. Alternatively, perforin-mediated donor anti-host cytotoxic activity may be critical for overcoming residual host resistance in the recipient. Thus, in the absence of perforin-mediated anti-host cytotoxicity, host resistance would remain stronger and persist longer in the recipient diminishing the relative alloaggressive capacity of the perforin-deficient inoculum.

One advantage of employing genetically modified or naturally occurring mutant mouse strains as a source of donor T cells is that only one cytotoxic pathway has been selectively abrogated leaving the other cytotoxic effector functions intact. Cytotoxic T cells and NK cells from perforin-deficient mice exhibit profoundly diminished in vitro cytolytic function, but retain the ability to effect Fas-mediated and TNF-mediated killing (15). Alternatively, the Fas-ligand molecule expressed by T cells from gld mice is non-functional, but perforin-dependent cytolytic function remains intact. Therefore, these experiments were able to directly examine the role of each cytotoxic pathway by its absence, while simultaneously confirming the function of the other by its presence. Accordingly, consistent with the marked diminishment of hepatic and cutaneous GVHD following transplantation of FasL-defective T cells, GVHD developed in these tissues following transplantation of perforin-deficient T cells because Fas-mediated cytotoxic function remains intact. Another advantage of employing T cells with molecular defects in cytolytic function is that the defect is present in every donor cell of all phenotypes regardless of how these populations may interact and contribute to the development of GVHD.

Previous studies designed to examine effector functions in GVHD have employed transplantation of cell populations that have been negatively selected on the basis of immunophenotype (e.g., CD8+, CD4+, NK1.1+ cells). In studies involving negative selection, all potential cytotoxic and cytokine effector functions of the depleted cell population are removed along with the negatively selected population. In contrast, the present study is designed so that the potential to produce multiple inflammatory cytokines such as IL-1, IFN-γ, and TNF-α by the cytotoxically defective T cells remains intact. These cytokines are known to be produced by alloactivated donor T cells during GVHD, and appear to contribute significantly to both inductive and effector phases of GVHD (38).

The results of the present studies represent the first characterization of the role of Fas and perforin-mediated anti-recipient cytotoxic function in the pathogenesis of GVHD after allogeneic bone marrow transplantation across non-MHC genetic disparities. These results demonstrate that Fas-mediated cytotoxicity is required for the development of hepatic and cutaneous GVHD, but is not required for GVHD-associated cachexia. This finding has shown that local and systemic effects of GVHD are separable. Furthermore, the results also demonstrate that absence of perforin-mediated anti-recipient cytotoxicity does not prevent or diminish GVHD, but significantly delays the time of onset. We interpret this finding to indicate that while perforin-mediated cytotoxic function is not required for the effector phase of GVHD, this cytotoxic pathway appears to play a significant role during the inductive phase of GVHD. We conclude that the two major pathways of cell-mediated cytotoxicity play distinct roles in the complex pathophysiology of GVHD, and that these roles appear to be expressed during different stages of GVHD pathogenesis.

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