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STAT4 and STAT6 are transcription factors that play crucial roles in responding to IL-12 and IL-4, respectively. STAT4 gene knockout (STAT4−/−) mice have markedly reduced Th1 responses and enhanced Th2 responses. STAT6−/− mice show the inverse phenotype. We compared the ability of bone marrow transplantation (BMT) with the inclusion of spleen cells from STAT6−/−, STAT4−/−, and wild-type (WT) mice to produce graft-versus-host disease (GVHD) in lethally irradiated MHC-mismatched recipients. Acute GVHD mortality was more rapid when induced by cells from STAT6−/− mice than when induced by STAT4−/− cells. However, cells from STAT4−/− and STAT6−/− donors both induced delayed GVHD mortality compared with WT controls, or compared with combined STAT4−/− and STAT6−/− cells, indicating a contribution of both Th1 cells and Th2 cells to acute GVHD. Recipients of STAT6−/− BMT showed evidence of acute GVHD with severe diarrhea and marked weight loss. Recipients of STAT4−/− BMT showed signs of GVHD with only initial transient weight loss and later development of severe skin GVHD. Histopathology showed that Th2 responses were required for the induction of both hepatic and severe skin GVHD. In contrast, both Th1 cells and Th2 cells were capable of causing intestinal pathology of GVHD. Our studies demonstrate an additive role for Th1 and Th2 cells in producing acute GVHD, and suggest a cytokine-directed approach to treating end-organ manifestations of GVHD.


Introduction

Despite major advances, the full therapeutic potential of bone marrow transplantation (BMT) has not yet been achieved. Graft-versus-host disease (GVHD), arising from an attack of donor T cells against alloantigens of the recipient, is a major complication of allogeneic BMT, and prohibits the successful application of BMT across extensive human leukocyte antigen barriers. Despite the morbidity and death it causes, GVHD has been associated with 2 clinical benefits. Donor T cells exert graft-versus-leukemia effects by destroying residual leukemic cells, and also enhance the engraftment process, probably by destroying residual host T cells that survive intensive conditioning regimens (1). Therefore, separation of the beneficial engraftment-promoting and anti-leukemic effects of donor T cells from their ability to induce GVHD is a major goal of research in BMT.

Two distinct functional cytokine secretion patterns, termed Th1 and Th2, have been defined for murine as well as human T helper cells. The Th1 response is characterized by the production of IFN-γ, IL-2, and TNF-β, by the stimulation of delayed-type hypersensitivity reactions, and by the production of high-affinity Fcγ receptor–binding Abs (IgG2a in the mouse) (2). Characteristic of the Th2 response in the mouse is the production of IL-4 and IL-5, and the stimulation of the production of IgE and IgG1 Abs (2). These cell types are cross-regulatory in vitro, and the balance of these cells in vivo determines the character of cell-mediated immune and inflammatory responses (2). Recently, it has been suggested that the Th1/Th2 polarization of T helper cell subsets plays an important role in the development of GVHD. In both mice and humans, a “cytokine storm” related to the Th1 phenotype has been found to correlate with the development of GVHD (3, 4). Studies have shown that in vivo injection of IL-2 plus IL-4 into donor mice resulted in the generation of CD4+ IL-4+ and IL-10-producing Th2 cells that, upon transfer into nonirradiated F1 or sublethally irradiated mice, inhibited GVHD lethality (5, 6). Furthermore, production of Th2 cytokines induced by G-CSF administration has been associated with inhibitory effects on the development of GVHD (7). In another study, transplantation of in vitro–polarized murine Th2 cells resulted in a significant increase of survival after BMT across minor histocompatibility antigenic barriers (8).
However, some other studies of the therapeutic approach of using a Th1 to Th2 switch for the prevention or treatment of GVHD have suggested that the roles of Th1 and Th2 cells in GVHD may be far more complicated than was previously believed. Altering the development of GVHD or treating ongoing GVHD by direct administration of Th2 cytokines in vivo has not been successful. In 1 study, recipients of IL-4 showed increased mortality and equivalent, histopathological changes in the skin and liver compared with mice receiving no cytokine. It has also been shown that the exogenous administration of IL-10 was not toxic per se, but did accelerate GVHD lethality (9, 10). Furthermore, recent studies using IFN-γ or IL-4 gene knockout mice as donors showed that the absence of IFN-γ is still associated with severe acute GVHD (11, 12), whereas the inability of donors to produce IL-4 can be associated with less severe GVHD (12).

To elucidate the contribution of T helper cell subsets, we initiated a study using gene knockout mice that were capable of mounting only Th1 or Th2 responses as donors. Interactions between cytokines and their receptors leads to the activation of multiple signaling molecules, including the family of signal transducer and activator of transcription (STAT) proteins. STAT4 is a member of this family that is essential for responsiveness to IL-12. Mice with a disrupted STAT4 gene (STAT4–/– mice) have impaired IL-12 responsiveness of NK and T cells, lack Th1 responsiveness, and have enhanced Th2 cell function (13, 14). STAT6 is activated in response to IL-4 and IL-13. Therefore, T lymphocytes from STAT6–/– mice have impaired IL-4 and IL-13 responsiveness, and almost completely lack Th2 responses, but show enhanced Th1 responses (15–17). We assessed the role of Th1 cells and Th2 cells in GVHD by transferring bone marrow cells (BMC) and spleen cells from normal B6 and normal BALB/c mice, irradiated with 30 Gy, and resuspended to a concentration of 10^6 cells/mL as responders. Stimulators were prepared from normal B6 or BALB/c mice, irradiated with 30 Gy, and resuspended to a concentration of 4 × 10^6 cells/mL. Responder and stimulator cell suspensions of 100 µL each were plated in 96-well U-bottom tissue culture plates. After incubation for 24 hours, culture supernatants were collected for cytokine analyses, and were immediately aliquoted and frozen.

**Methods**

**Mice.** Specific pathogen-free female C57BL/6 (B6, H-2^b^, K^d^D^d^) and BALB/c (H-2^d^, K^d^D^d^) mice were purchased from the Frederick Cancer Research Facility (NCI Laboratories, Frederick, Maryland, USA). STAT4 and STAT6 gene knockout mice were generated on the 129 background and were backcrossed to the BALB/c background 5–10 times. All donor lines were H-2d homozygotes. Mice were housed in sterilized microisolator cages and received autoclaved feed and autoclaved, acidified drinking water.

**Preparation of BMC, spleen cells, and T-cell depleted BMC.** BMC and splenocyte suspensions were prepared as described (18). Host-type (B6) BMC were depleted of T cells with mAbs against CD4 (GK1.5 in ascites, 1:1,000 dilution) (19) and CD8 (2.43 in ascites, 1:1,500 dilution) (20), plus low-toxicity rabbit complement (1:16 dilution), as described previously (21).

**Total body irradiation and BMT.** Recipient mice were lethally irradiated (9.75 Gy, 137Cs source) and were reconstituted within 4–8 hours with a single (1 mL) intravenous inoculum. For each experiment we compared the number of CD4+ and CD8+ T cells in the spleens of wild-type (WT), STAT4–/–, and STAT6–/– mice, and we did not observe significant differences in the percentages of each cell type. To avoid bias from cage-related effects, animals were randomized as described (22).

**Flow cytometry and mAbs.** Spleen cells were analyzed by 2-color flow cytometry on a FACSscan (Becton Dickinson, Mountain View, California, USA). To block nonspecific Fcγ receptor binding of labeled Abs, 2.4G2 (rat anti-mouse Fcγ receptor mAb) (23) was added to the first incubation. To determine the percentage of donor- and host-type T cells in spleen, cells were stained with FITC-conjugated rat anti-mouse CD4 mAb, rat anti-mouse CD8 mAb (PharMingen, San Diego, California, USA), and biotinylated anti-H2D^d^ mAb 34-12-12 (24). The percentage of donor- and host-type T cells was determined by subtracting the percentage of HOPC1-staining cells from the percentage of CD4+ and CD8+ cells in the same quadrant, as described (11).

**Mixed lymphocyte culture (MLC) supernatants.** Spleen cells from randomly chosen GVHD control, STAT4–/–, and STAT6–/– transplant recipients, and from normal B6 and normal BALB/c mice were resuspended to a concentration of 10^6 cells/mL as responders. Stimulators were prepared from normal B6 or BALB/c mice, irradiated with 30 Gy, and resuspended to a concentration of 4 × 10^6 cells/mL. Responder and stimulator cell suspensions of 100 µL each were plated in 96-well U-bottom tissue culture plates. After incubation for 24 hours, culture supernatants were collected for cytokine analyses, and were immediately aliquoted and frozen.

**ELISA for cytokine levels in MLC supernatants and for immunoglobulin levels in sera.** Cytokines in MLC supernatants were measured by ELISA using mouse cytokine–specific rat mAbs (IFN-γ capture mAb R4-6A2, IFN-γ detecting mAb XM1 1:2; IL-2 capture mAb JES6-1A12, IL-2 detecting mAb JES6-5H4; IL-4 capture mAb R4-6A2BVD4-1D11, IL-4 detecting mAb BVD6-24G2; IL-10 capture mAb JES5-2A5, IL-10 detecting mAb SXC-1), and were compared with known concentrations of mouse recombinant cytokines (PharMingen) as described (25). IgE, IgG1, and total IgG levels in sera were measured by ELISA using purified mouse immunoglobulin standards and mouse immunoglobulin-specific Abs (IgE capture mAb R35-72, IgE detecting mAb R35-92; IgG1 capture mAb A85-3, IgG1 detecting mAb A85-1 [PharMingen]; total IgG capture and detecting goat anti-mouse IgG [Southern Biotechnology Associates, Birmingham, Alabama, USA]), as described (26). The sensitivity of each immunoglobulin assay was 2 ng/mL for IgE, 0.16 ng/mL for IgG1, and 0.25 ng/mL for total IgG. The sensitivity of each cytokine assay was 0.31 ng/mL for IFN-γ, 0.25 ng/mL for IL-2, 0.25 ng for IL-10, and 8 pg/mL for IL-4.
Assessment of GVHD. Transplanted animals were followed for more than 100 days after BMT. Body weights were measured on the day of transplantation, and then twice each week during the first month and once a week after that. Animals were also scored for clinical evidence of GVHD by assessment of changes in skin (i.e., alopecia, inflamed or scaly skin), generalized signs (fur texture, posture, and inflammation of the eyes), and diarrhea. Each parameter was quantified by scoring as follows: 0 = normal, 1 = mild, 2 = moderate, 3 = severe (for scoring of eye inflammation, 4 = eyes severely inflamed and closed).

Histopathology. For histopathological analysis of GVHD target tissues, samples were collected from large intestine, kidney, liver, and skin (from the head), and were fixed in 10% formalin. Formalin-preserved tissue samples were embedded in paraffin, cut into 5-μm-thick sections, and stained with hematoxylin and eosin for histological examination. Slides were coded without reference to mouse type or prior treatment status, and were systematically examined by a pathologist.

Statistical analysis. For comparison between groups of cytokine levels in supernatants, the Student’s 2-tailed t test for comparison of means was used. A P value less than 0.05 was considered to be statistically significant. Survival data were analyzed using the Kaplan-Meier method of life table analysis, and statistical analysis was performed by the Mantel-Haenszel test. Unless otherwise noted, data are presented as mean ± SEM.

Results

Both Th1 cells and Th2 cells contribute to GVHD. To evaluate the separate contributions of Th1 and Th2 cells to the induction of GVHD in a fully MHC-mismatched and multiple minor histocompatibility antigen–mismatched strain combination, we compared the ability of BMC plus spleen cells from BALB/c→backcrossed H-2b STAT4–/– mice, STAT6–/– mice, or WT control BALB/c mice to induce GVHD in lethally irradiated C57BL/6 recipients. The number of donor CD4+ and CD8+ T cells administered in WT, STAT4–/–, and STAT6–/– inocula was similar in all experiments, as measured by flow cytometry analysis of each inoculum. As shown in Figure 1, the most severe mortality occurred in the group receiving WT splenocytes, which are capable of producing both Th1 and Th2 responses (median survival time [MST] = 22 days). STAT6–/– BMC, which is capable of producing high Th1 responses and no Th2 responses, induced mortality (MST = 31.5 days) that was less rapid in onset than that induced by WT BMC (P < 0.01). Mortality caused by the injection of STAT4–/– cells, which lack Th1 and produce enhanced Th2 responses, was consistently less rapid (MST = 79 days) than that caused by both WT (P < 0.0001) and STAT6–/– cells (P < 0.001). Figure 1 shows the mortality of 117 animals from 6 separate experiments. We observed that the rapidity of onset of GVHD varied significantly from experiment to experiment, but in each experiment the GVHD mortality was most rapid when induced by WT cells, followed by STAT6–/– cells and finally STAT4–/– cells. Furthermore, when GVHD was induced in the reverse strain combination, using C57BL/6→backcrossed H-2b STAT4–/–, STAT6–/–, or WT control C57BL/6 mice to induce GVHD in lethally irradiated BALB/c recipients, the same hierarchy of rapidity of GVHD mortality was observed: mortality induced by WT cells was the most rapid, followed by STAT6–/– cells and then STAT4–/– cells, the slowest (data not shown).

To further demonstrate that both Th1 cells and Th2 cells contribute to acute GVHD, we compared the severity and mortality of GVHD induced by STAT4–/– cells, STAT6–/– cells, and combined STAT4–/– plus STAT6–/– cells. Mice received either a full dose (13 × 10^6) or a half dose (6.5 × 10^6) of both STAT4–/– and STAT6–/– spleen cells plus BMC, or 13 × 10^6 WT, STAT4–/–, or STAT6–/– cells alone. Whereas recipients of the full dose of combined STAT4–/– and STAT6–/– spleen cells developed accelerated GVHD (MST = 6 days) compared with recipients of WT inocula (MST = 16 days), there was no difference in GVHD induced by 13 × 10^6 WT spleen cells compared with the combination of 6.5 × 10^6 STAT4–/– cells and 6.5 × 10^6 STAT6–/– cells (MST = 15.5 days). Recipients of 13 × 10^6 STAT6–/– or STAT4–/– spleen cells alone showed delayed GVHD compared with recipients of either 13 × 10^6 WT cells or 6.5 × 10^6 STAT4–/– cells and 6.5 × 10^6 STAT6–/– cells, confirming that both Th1 cells and Th2 cells contribute to acute GVHD.

The addition of STAT4–/– cells to WT cells does not protect against GVHD. It has previously been shown that transplantation of polarized Th2 cells, after 7 days in culture with IL-2 and IL-4, reduced the GVHD mortality caused by nonpolarized cell populations (27). We therefore attempted to determine whether naive Th2 cells could ameliorate GVHD induced by Th1 cells by adding STAT4–/– BMC and spleen cells to WT cells. The addition of STAT4–/– (Th2) cells to WT cells not only failed
The addition of STAT4–/– cells to WT BMC plus spleen cell inocula does not protect from GVHD. Lethally irradiated B6 mice received TCD B6 BMC and either syngeneic spleen cells (n = 3), BALB/c WT BMC and spleen cells (WT) (n = 6), STAT4–/– (Th2) BMC and spleen cells (n = 3), or WT BMC and spleen cells plus STAT4–/– BMC and spleen cells (STAT4–/– + WT) (n = 3). Each inoculum contained 13 \times 10^6 BALB/c spleen cells plus 10 \times 10^6 BALB/c BMC (WT-induced control GVHD); 13 \times 10^6 BALB/c STAT4–/– spleen cells plus 10 \times 10^6 BALB/c STAT4–/– BMC (STAT4–/–-induced GVHD); or 6.5 \times 10^6 BALB/c WT spleen cells plus 6.5 \times 10^6 BALB/c STAT4–/– spleen cells plus 5 \times 10^6 BALB/c WT BMC plus 5 \times 10^6 STAT4–/– BMC (STAT4–/– + WT) cells. Results of one of two similar experiments are shown.

to prevent or reduce the severity of acute GVHD (MST = 18 days), but it accelerated GVHD mortality compared with that induced by WT cells alone (MST = 31 days) (Figure 2 shows one of two similar experiments). This result further emphasizes that Th2 cells play a pathogenic role in GVHD, and shows that they do not effectively inhibit GVHD mediated by naive cells that can differentiate into Th1 cells.

Unique clinical and pathological syndromes associated with Th2 and Th1 GVHD. We monitored transplanted animals for clinical signs of GVHD. Figure 3 shows the mortality curve and weight curve from a representative experiment. GVHD scores for generalized signs (hunching, ruffled fur, and periorbital edema) of WT recipients were the most severe in intensity and were the earliest in onset: on day 8, mean GVHD score was 9.8 ± 0.6 out of a possible maximum score of 10. The recipients of STAT6–/– BMT and STAT4–/– BMT had GVHD scores that were lower at this time point (on day 8, mean score for recipients of STAT4–/– BMT was 2.3 ± 1.2, and for recipients of STAT6–/– BMT it was 5.4 ± 0.5) and were similar to each other. However, the time required to achieve the maximal score for generalized signs of GVHD was earlier in recipients of STAT6–/– BMT (on day 12, mean score was 6.7 ± 0.6) than in STAT4–/– BMT recipients (on day 16, mean score was 7 ± 1). A weight curve, which provides a sensitive indicator of GVHD, is shown in the lower panel of Figure 3. Animals receiving WT cells rapidly lost weight before their deaths by day 13. Animals receiving Th1 (STAT6–/–) and Th2 (STAT4–/–) BMT showed initial weight loss during the first 10 days after BMT. However, the clinical signs subsequently diverged markedly in the two groups. Whereas recipients of STAT6–/– BMT developed severe diarrhea and began to show mortality around day 25, recipients of STAT4–/– BMT did not have diarrhea, and had only mild weight loss. As shown in the upper panel of Figure 3, recipients of STAT4–/– BMT began to show mortality around day 30. These mice developed extremely severe skin GVHD, which began around the eyes, mouth, and ears and progressed to involve the whole body, resulting in hair loss and profound scaling. This striking clinical picture was observed in all six experiments, with time of onset ranging from 15 to 50 days after BMT. Whereas these clinical changes were not observed in any experiments in the animals that received BMT from WT or STAT6–/– mice and survived until day 100, they were present in the majority (11 of 14) of STAT4–/– BMT recipients that survived to 10 weeks after BMT.

Different localization of pathological GVHD-associated changes in recipients of WT, STAT4–/–, and STAT6–/– BMT. After death, carcasses were fixed in formaldehyde and the tissues were processed and scored in a blinded fashion for lesions associated with GVHD. All analyzed organs were harvested from animals that died at similar times after transplantation from two experiments where the severity of GVHD was similar (MSTs in days for experiment 1 and 2: WT = 12 and 19, STAT6–/– = 35 and 29, STAT4–/– = 79 and 155). The liver, spleen, lungs, kidney, and large intestine were examined. There were...
no pathological changes in the appearance of liver (Figure 4a), skin (Figure 4e), or large intestine (Figure 4i) in recipients of syngeneic BMT. Moderate to severe GVHD-induced liver changes were observed in 5 of 5 recipients of WT BMT, which died on days 19, 24, 27, and 54 after BMT. Figure 4b shows typical liver lesions found in these mice: portal areas were diffusely infiltrated by a mixed population of leukocytes including neutrophils, lymphocytes, macrophages, and a few plasma cells. There was associated multifocal, single-cell hepatocellular necrosis, consistent with severe hepatic GVHD. Similar changes were observed in 4 of 4 recipients of STAT6−/− BMT, which died on days 20, 39, 47, and 106 after BMT. In striking contrast, liver sections of STAT6−/− BMT recipients that died at similar times (22, 23, 25, 26, 27, 29, and 38 days after BMT) appeared normal, without any infiltrate or evidence of cholestasis (Figure 4c).

All recipients of WT BMT showed signs of slight skin hyperkeratosis. In some mice (3 of 7) that died on days 19 and 27 after BMT, there were very mild focal mononuclear inflammatory cell infiltrates located mostly in the dermis (Figure 4f). In accordance with the clinical picture, recipients of STAT4−/− BMT (3 of 4) that died on days 20, 39, and 106 after BMT showed marked skin abnormalities. In some animals, pustules and crusting with neutrophil infiltration was observed on the skin’s surface (Figure 4h). In all animals, the dermis was infiltrated by a mixed population of leukocytes, including lymphocytes, macrophages, and neutrophils (Figure 4h). In several places the cells formed granulomas. There was also a suggestion of lysis along the epidermal-dermal junction. Consistent with the clinical picture, inflammatory changes were absent in all 12 recipients of STAT6−/− BMT that died on days 5, 22, 23, 25, 26, 27, 29, 38, and 44 after BMT (Figure 4g). Furthermore, these severe skin changes were not observed in mice with GVHD induced by STAT6−/− that had long-term survival (4 animals that died at 74, 122, 149, and 150 days after BMT). Their skin showed signs of mild hyperkeratosis, but without inflammatory changes. Recipients of WT and STAT6−/− BMT showed clinical signs of severe colitis manifested by profuse diarrhea and weight loss. The pathological examination revealed that in 7 of 7 recipients of WT inocula, which died on days 19, 24, 27, 39, and 54 after BMT, there was a thick-
IgG1 in sera are expressed as mean ± SEM. B6 mice received 5·10^6 BMC. Recipients of cells, BALB/c WT BMC and spleen cells (WT), were 26 for recipients of WT cells, 54 for recipients of group, 2–4 animals were analyzed. MSTs in days for this experiment were filled with exudate and severely ulcerated mucosa had extremely severe chronic colitis, with lumens that were undetectable levels of serum IgG 1 (Figure 5b). Total IgG levels were similar in the recipients of STAT6−/− BMT (312.9 ± 152.0 μg/mL) sacrificed at 5 months after BMT and in recipients of STAT4−/− cells (470.3 ± 38.0 μg/mL) sacrificed at 2 weeks after BMT, despite the difference in IgG1 levels cited above.

**Donor engraftment and Thelper phenotype in recipients of WT, STAT4−/−, and STAT6−/− BMT.** Recipients of STAT4−/− cells were sacrificed at 2 weeks, 3 months (87 days), and 5 months (136 days) after BMT. Recipients of syngeneic and WT cells were sacrificed at 2 weeks and at 3 months (87 days) after BMT. For STAT6−/−-induced GVHD, animals were sacrificed only at 5 months (136 days) after BMT. MLCs were prepared with host-type (B6) irradiated stimulator cells. FACS® analysis of peripheral blood lymphocytes and bone marrow showed that these animals were full donor-type hematopoietic chimeras at all 3 time points (data not shown). The splenocytes from these animals showed mixed lymphocyte response reactivity both toward

![Figure 5](image-url)
host C57BL/6 cells and third-party strain A/J, while being nonreactive toward donor antigens (data not shown). Thus, there was constant, ongoing activation and proliferation of donor T cells toward host antigens. By quantifying the levels of IFN-γ, IL-2, IL-4, and IL-10 in MLC supernatants from WT and STAT4–/– BMT recipients, distinct patterns emerged. In supernatants obtained from spleen cells of mice receiving STAT4-deficient cells, the amounts of IL-4 and IL-10 (i.e., the Th2-specific cytokines) were increased compared with those in supernatants of spleen cells from WT recipients (see Figure 6). Similar results were obtained at all 3 time points after BMT. IFN-γ and IL-2, which are both induced during the Th1 response, were detected at higher levels in supernatants of spleen cells from recipients of WT BMT than in recipients of STAT4–/– BMT (see Figure 6). Although produced at lower levels than in the recipients of WT BMT, there was measurable production of IFN-γ in the recipients of STAT4–/– BMT (Figure 6). Due to profound atrophy of the spleen at 5 months after BMT, only 1 recipient of STAT6–/– BMT was available for analysis. This mouse showed the expected Th1 phenotype, with large amounts of IFN-γ (781 pg/mL) and IL-2 (591 pg/mL), low amounts of IL-10 (228 pg/mL), and undetectable IL-4 in the collected supernatants.

Discussion

Our study indicates that the role of Th1 cells and Th2 cells in GVHD is more complicated than previously thought. We have shown, contrary to other studies (6, 8, 27, 30), that both Th1 cells and Th2 cells contribute to acute GVHD. However, each subset causes injury of specific tissues, resulting in a unique clinical picture when Th1 or Th2 responses separate induces GVHD. Most strikingly, Th2 cells appear to be absolutely required for the induction of hepatic GVHD and severe skin GVHD. If confirmed in humans, these observations could have important implications for the therapy of GVHD, because they suggest that Th2-targeted approaches might be used to most effectively and specifically target liver and skin manifestations of acute GVHD, which are frequently seen without intestinal GVHD in clinical BMT.

We consistently observed that WT cells induced the most severe GVHD when compared with that induced by STAT6–/– (Th1) or STAT4–/– (Th2) cells. In other studies, we have observed that GVHD induced by IL-4–/– cells (wherein the disease results mostly from Th1 cells, but with some Th2 component due to residual signaling through IL-13) was of a severity intermediate between that induced by WT cells and STAT6–/– cells (B. Nikolic and M. Sykes, unpublished data). In further support of the hypothesis that both Th1 cells and Th2 cells contribute to GVHD, our group and others have shown that acute GVHD can occur in the absence of IFN-γ (11, 12), and that GVHD induced by donors that cannot produce IL-4 is milder than that induced by WT cells (12). Furthermore, the administration of anti–IL-4 mAb can ameliorate acute GVHD (31). Previous studies have shown that systemic administration of the Th1-inducing or Th1-associated cytokines IL-2 (18), IL-12 (32), or IFN-γ (33) can inhibit GVHD. Paradoxically, in other models, the administration of anti–IL-2 or anti–IL-12 mAbs (31, 34) has also been shown to prevent acute GVHD. Apparently, cytokines play complex roles in the induction and inhibition of GVHD, and the timing, the route of administration, and the strain combination may be critical in determining their effect (11, 35–38).

Although the experiments described herein were performed under similar conditions, with careful matching of the ages of donors and recipients and of the number of injected cells and dose of irradiation, we have always observed wide variability in the severity of GVHD from experiment to experiment, particularly over long time periods. There seem to be periods of months or years in which a given T-cell dose produces GVHD of a certain tempo in a particular model, and then this tempo later changes for reasons that cannot be identified. To minimize the effect of this variability and enable interpretation of data, we included every control group in every

Figure 6

Cytokine profile of splenocytes from mice with WT- or STAT4–/–-induced GVHD. Lethally irradiated C57BL/6 host mice transplanted with 5 × 10^6 TCD B6 BMC and either 13 × 10^6 WT spleen cells plus 10 × 10^6 WT BMC (n = 2) or 13 × 10^6 STAT4–/– spleen cells plus 10 × 10^6 STAT4–/– BMC (n = 4) were sacrificed at 2 weeks after BMT, and splenocyte suspensions were prepared. Their donor-derived phenotype was demonstrated by FACS®. The cells were cultured with irradiated splenocytes from C57BL/6 mice. Supernatants were harvested 72 hours after initiation of MLC, and cytokine levels were determined by ELISA. MSTs for this experiment were 26 days for recipients of WT cells and 54 days for recipients of STAT4–/– cells. Concentrations of IFN-γ, IL-2, IL-4, and IL-10 in cell culture supernatants are expressed as mean ± SEM (pg/mL). *P < 0.01 vs. WT group.
experiment. GVHD induced by WT cells serves as a standard for the severity of GVHD, and syngeneic BMT acts as a negative control. In this study, we consistently observed that GVHD mortality was more rapid when induced by cells from STAT6–/– mice than that induced by STAT4–/– cells. Nevertheless, the observed variability in the severity of GVHD and differences between human and murine GVHD mandate additional studies before direct translation of these results to human transplantation should be considered.

Although our results are at variance with previous studies suggesting that Th2 cells do not cause, and even prevent acute GVHD, these studies used spontaneous or LPS-induced mortality as an endpoint, and did not show an absence of GVHD pathology in recipients of Th2 cells (8, 27). Another possible reason for the inconsistency of our results with others that implicated Th1 cells but not Th2 cells in GVHD is that some of the previous studies relied on polarization of T helper cells by incubation with cytokines for several days in vitro (5, 8, 27). Polarized T cells were then transferred to irradiated recipients. This procedure might result in incomplete polarization, partial anergy, or a failure of subsequent proliferation, and could allow secondary conversion of T cells to a mixed phenotype upon transfer to BMT recipients. Indeed, examination of T-cell subsets in these BMT models revealed that T-cell populations were not polarized into “classical” Th1 or Th2 cytokine phenotypes 2 weeks after BMT (8), and that intermediate cytokine profiles of both CD8+ and CD4+ T-cell subsets developed, as has been described in other systems (30). STAT4– and STAT6-deficient mice have the advantage over these experimental models of allowing more pure Th1 or Th2 responses to be examined, and of not requiring a priming step before T cells are transferred for the induction of GVHD (15, 17, 39, 40). Although STAT6–/– lymphocytes are completely impaired in their ability to generate IL-4–secreting cells, we were able to detect some IFN-γ in supernatants of MLC from STAT4–/–-induced GVHD (Figure 6). A recent study showed that under conditions that promote Th1 cell differentiation, it is possible to detect low levels of production of IFN-γ via a STAT4-independent pathway (41). However, whereas mice that are both STAT4+/– and STAT6–/– are able to produce sufficient amounts of IFN-γ and mount Th1 cell–mediated delayed-type hypersensitivity, STAT4–/– mice retain their predominantly Th2 phenotype, presumably through dominant IL-4 activation of STAT6 or Th2-mediated inhibition of Th1 cytokine production, or both (41).

Perhaps in keeping with our data showing that Th1 cells are associated with severe intestinal GVHD, a requirement for Th1 responses in inducing pathology in 2 different models of experimental colitis was recently demonstrated (42). Treatment with anti-IL-12 Abs or bone marrow reconstitution of T-cell– and NK cell–deficient Tg26 mice from STAT4-deficient mice resulted in modulation of the severity of disease (42). Thus, the reduced intestinal pathology in Th2 recipients may have been an important determinant in the observed prolonged survival of recipients of STAT4–/– BMT. This may also explain data from other experiments in which mortality of recipients of in vitro–polarized Th2 cells was diminished or even absent (5, 7, 8, 27). In some of these studies, models were used that tend to induce milder GVHD, perhaps accounting for the failure to observe the contribution of Th2 cells to acute GVHD. Consistent with the failure of Th2 cells to induce acute GVHD across minor histocompatibility barriers without MHC differences (8), a recent study demonstrated that Th2 cells were unable to reject MHC-matched, minor histocompatibility antigen–mismatched cells, but were able to rapidly reject fully MHC-mismatched grafts (43). Thus, the presence of full MHC disparities in our model may partially explain the apparent discrepancy with the results of Krenger et al. (8, 27).

Consistent with our data showing a requirement for Th2 cells in hepatic and skin GVHD, a recent study has shown that although mortality from GVHD is substantially diminished after infusion with donor in vitro–polarized Th2 cells, clinical signs of GVHD are still present in surviving mice, and target organ pathology persists. Significant cutaneous and hepatic histopathological damage was evident 7 weeks after BMT in mice receiving polarized donor Th2 cells, and was equivalent to that in animals receiving naive donor T cells (8). Furthermore, after induction of GVHD with IFN-γ–/– cells, extensive mononuclear cell infiltrates were seen in livers (12); treatment with anti–IL-4 mAb markedly reduced IgE and IgG1 levels, and liver disease was ameliorated (31). Increased levels of IgE correlated strongly with the histological grades and stages of liver disease in parental mice receiving F1 spleen cell transfer, and treatment with anti–IL-4 mAb prevented both phenomena (44). In humans, predominant activation of Th2 cells, indirectly measured by elevated serum IgE levels, was associated with liver disease in GVHD (28).

T1 (Tc1) and T2 (Tc2) subsets of CD8+ T cells secrete polarized patterns of cytokines similar to the Th1 and Th2 CD4 subsets. Although both T1 and T2 CD8+ T cells exhibit comparable levels of perforin-mediated cytolyis (45), T1 CD8+ T cells exhibit significantly higher Fas-mediated cytosis (45). Fas-mediated anti-repipient cytotoxicity has been implicated in the development of hepatic and cutaneous GVHD against minor histocompatibility antigens (46), but this was not reproduced in a full haplotype-mismatched strain combination (47). In another study, administration of an anti–Fas ligand inhibitory mAb resulted in a reduction of the weight loss and mortality associated with GVHD directed against a full haplotype mismatch, but without improvement in skin or liver lesions (48). Therefore, Tc1-mediated cytosis may not play a major role in hepatic and skin GVHD caused by a full MHC disparity. In accordance with these data, we have observed that only Th2 and Th2 cells induced liver and skin GVHD in a fully MHC-mismatched strain combination.
The preferential injury of gut-associated tissue in Th1-induced GVHD, and of skin and liver in GVHD caused by Th2 cells, may be the result of different homing properties of the different subsets. Indeed, there is growing evidence that in addition to effector function, activated Th1 and Th2 lymphocytes acquire different migratory capacities (49–52). Chemokines and chemokine receptors are probably important for the selective migration of T-cell subsets (50). Distinct profiles of chemokine receptors are acquired by T cells after polarization; these are modulated by cytokines (50–52). Indeed, in vivo studies have confirmed this concept. Mouse Th1 cells express the functional ligand for P-selectin and E-selectin, PSGL-1, which promotes their migration into inflamed tissue (53). It was recently shown that Th1 cells could efficiently enter inflamed sites, such as sensitized skin or artritic joints (53). In our model, preferential migration of Th2 cells, but not Th1 cells, into the liver and skin was observed, suggesting that other receptors or chemokines may play a decisive role in this migratory process.

There are potential dangers associated with therapies based on immune deviation from Th1 responses to Th2 responses. Th2 responses are responsible for the pathology of allergic diseases and at least 1 autoimmune condition (54–57). Our data indicate that manipulation of the Th1/Th2 balance as an approach to the prevention of GVHD may not be ideal. Possible severe side effects of such treatment should be considered before proceeding with human trials. However, our results suggest that the potential to treat hepatic and skin GVHD with anti–Th2 cytokine therapy deserves exploration.

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