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The Absence of Donor-Derived IL-13 Exacerbates the Severity of Acute Graft-Versus-Host Disease Following Allogeneic Bone Marrow Transplantation

Gerhard C. Hildebrandt, MD,^{1,2}* Sung W. Choi, MD,² Gunnar Mueller,¹ Krystyna M. Olkiewicz, MS,² Bethany B. Moore, PhD,³ and Kenneth R. Cooke, MD²

Acute graft versus host disease (aGVHD) after allogeneic bone marrow transplantation (allo-BMT) predominantly involves a Th1type cytokine response. Interestingly, the Th2-cytokine, Interleukin-13 (IL-13), produced by alloreactive donor T cells in vitro was recently shown to correlate with clinical aGVHD severity. Using an established mouse model, we show that the systemic cytokine milieu following allo-BMT with IL-13–/– donors is characterized by decreases in serum Th α , and ultimately correlates with higher aGVHD mortality compared to allogeneic controls. In vitro studies further demonstrate that both exogenous and T cell-derived IL-13 can regulate TNF α production by macrophages following lipopolysaccharide stimulation. Thus, donor-derived IL-13 may have a role in modulating inflammatory cytokine release that is associated with aGVHD. Pediatr Blood Cancer 2008;50:911–914. © 2007 Wiley-Liss, Inc.

Key words: tumor necrosis factor α ; acute GVHD; T cell; cytokines; IL-13

INTRODUCTION

Allogeneic (allo-) bone marrow transplantation (BMT) is the only curative option for a number of malignant and non-malignant diseases. The utility of allo-BMT is limited by the development of acute graft versus host disease (aGVHD) [1]. Experimental and clinical data suggest that aGVHD is predominantly mediated by a Th1-type cytokine response [2,3], although an additive role for Th1 and Th2 cells has been described [4]. Interestingly, the production of Interleukin-13 (IL-13) by donor T cells responding to host antigens in vitro was recently shown to correlate with the severity of clinical aGVHD [5]. IL-13 is a typical Th2-type cytokine that is produced by a wide variety of cells including T cells [6,7]. IL-13 binds to its functional receptor (IL-13R) that consists of the IL-4Ra chain and the IL-13R α 1 chain [8] or to a decoy receptor, called IL-13R α 2 [9]. IL-13 exhibits a range of biologic activities, some of which overlap with those observed with IL-4 [10-12]. Although T lymphocytes are a significant producer of IL-13, these cells do not express the IL-13R on their surface and therefore do not respond to this cytokine [13]. We used a well-established murine BMT model (BALB/cB6) and mice deficient in IL-13 as BMT donors to more clearly determine the role of donor T cell derived IL-13 in the pathogenesis of aGVHD.

MATERIAL AND METHODS

Mice and Bone Marrow Transplantation

Mice received BMT as described previously [16]. C57BL/6 mice received 13 Gy total body irradiation (TBI; ¹³⁷Cs source) split into two doses followed by intravenous injection of 5×10^6 bone marrow (BM) and 2×10^6 CD4+ and CD8+ magnetic-bead-purified (Miltenyi Biotec GmbH, Bergisch Gladbach, Germany) splenic T cells from either syngeneic B6, allogeneic BALB/c IL-13^{+/+} or allogeneic BALB/c IL-13^{-/-} donors. Survival was monitored daily until day 180.

Fluorescence-activated Cell Sorting (FACS) Analysis

Cells were analyzed by FACS using fluorescein isothiocyanateconjugated monoclonal antibodies to CD8 and phycoerythrinconjugated antibodies to CD4 (BD Pharmingen, San Diego, CA) as described [15].

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Female BALB/c IL- $13^{+/+}$ (H- 2^{d}) and C57BL/6 (H- 2^{b}) mice were purchased from Jackson Laboratories (Bar Harbor, ME). BALB/c IL- $13^{-/-}$ (H- 2^{d}) mice were provided by Andrew McKenzie [14]. Animals were housed and fed as previously described [15].

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¹Department of Hematology and Oncology, University of Regensburg Medical School, D-93053 Regensburg, Germany; ²Department of Pediatrics, Division of Hematology and Oncology, Bone Marrow Transplantation Program, University of Michigan, Ann Arbor, 48109, Michigan; ³Department of Internal Medicine, Division of Pulmonary and Critical Care Medicine, University of Michigan, Ann Arbor, 48109, Michigan

^{*}Correspondence to: Gerhard C. Hildebrandt, Department of Hematology and Oncology, University of Regensburg Medical School, Franz-Josef-Strauß Allee 11, D-93053 Regensburg, Germany. E-mail: gerhard.hildebrandt@klinik.uni-regensburg.de

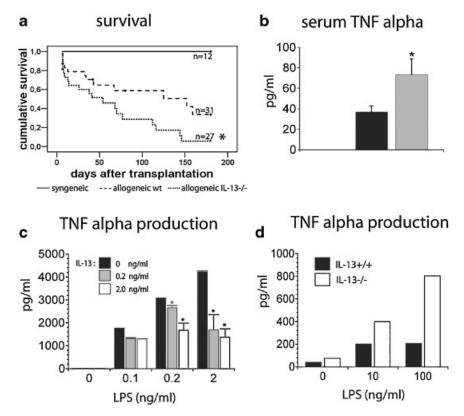


Fig. 1. IL-13 production regulates mortality after allogeneic BMT and TNF α production in vivo and in vitro. Lethally irradiated C57BL/6 (\Box ;—), allogeneic BALB/c IL-13+/+ (\blacksquare ;--) or allogeneic BALB/c IL-13-/- (\blacksquare ;[…]) donors as described in Materials and Methods. **a**: Allogeneic BMT from IL-13-/- donors resulted in significantly increased GVHD-related mortality compared to animals receiving allogeneic IL-13+/+ donor cells. Data presented are combined from four experiments; n = 12-31 per group; each experiment concurrently included animals of all groups; P < 0.02. **b**: Compared to animals receiving allogeneic IL-13+/+ BMT, serum TNF α levels in recipients of allogeneic IL-13-/- cells were significantly elevated on day 7 (and on day 14; data not shown) after BMT. Data are presented as mean \pm standard error and are combined from three experiments; n = 9-15 per group; P < 0.05. **c**: Pre-incubation of peritoneal cells with recombinant IL-13 reduces TNF α production following LPS stimulation. Data are presented as mean \pm standard error. **d**: TNF α production of peritoneal cells to LPS stimulation is reduced following pre-incubation with supernatant obtained from MLRs using IL-13+/+ T cells + allogeneic APCs compared to those obtained when peritoneal cells were pre-incubated with supernatants collected following MLRs with IL-13+/+ T cells or IL-13-/- T cells + media alone. Data are from one of two similar experiments.

T Cell Proliferation

Donor T cell expansion in the spleen was assessed in vivo 7 and 14 days following BMT as described [15]. Antigen-specific T cell proliferation in vitro was assessed in mixed lymphocyte reactions (MLR) [15]. C57BL/6, BALB/c IL-13^{+/+} or BALB/c IL-13^{-/-} T cells were plated at a concentration of 2×10^5 cells/well with 2×10^4 C57BL/6 splenic DCs isolated as described [15]. At 96 hr, supernatants were collected for cytokine analysis, responder cells were pulsed with ³[H] thymidine and proliferation in response to Concavalin A was also determined [15]. In some experiments, C57BL/6 peritoneal cells were pre-incubated with recombinant IL-13 (6 hr) or MLR supernatants (4 hr) prior to stimulation with lipopolysaccharide (LPS) to produce TNF α .

Measurement of Cytokine Protein Levels by ELISA

Concentrations of IFN γ , IL-13 in serum (day +7 after BMT) and/or cell culture supernatants were measured in duplicate by

ELISA (IFNγ: OptEIA; BD Pharmingen. IL-13: Biosource; Camarillo, CA). Serum levels of TNFα, IL-2,-4,-5 were determined by Cytokine Bead Array (Th1/Th2 cytokine kit, BD Pharmingen). Assay methods and limits of detection were followed as per the specific protocol.

Statistical Analysis

Values are expressed as the mean \pm S.E.M. Statistical comparisons between groups were completed using the parametric independent sample *t*-test. Survival curves were plotted using Kaplan–Meier estimates. The log-rank test was used to analyze survival data.

RESULTS AND DISCUSSION

We investigated the role of IL-13 produced by donor cells in the pathogenesis of aGVHD. First, we determined whether the absence of IL-13 in T lymphocytes alters alloantigen-specific responses in vitro. C57BL/6, BALB/c IL-13^{+/+} and BALB/c IL-13^{-/-} T cells

were stimulated with B6 splenic DCs. BALB/c IL-13^{+/+} and IL-13^{-/-} T cells proliferated comparably well in response to B6 alloantigens compared to syngeneic controls (Supplemental Figure 1a) and only modest changes in IFN γ (P < 0.05) and IL-2 (n.s.) production were seen in vitro but not in vivo between allogeneic groups (Supplemental Figure 1b and c). While allogeneic wild-type T cells produced significant amounts of IL-13 in culture, IL-13 levels in MLRs using allogeneic IL-13^{-/-} T cells were low and did not differ from syngeneic controls (Supplemental Figure 1d). Further, IL-5 (Supplemental Figure 1e) and IL-4 (not shown) production by IL-13^{-/-} T cells were also significantly reduced, indicating a general decrease in Th2 cytokine production. Consistent with previous reports [14], similar results were obtained after non-specific T cell stimulation with Con A (Supplemental Figure 1a-e).

We next determined the effects of IL-13 deficiency in donor cells on splenic T cell expansion and the resultant proinflammatory cytokine milieu characteristically observed following allo-BMT. Lethally irradiated B6 mice received BMT from either syngeneic (B6) or allogeneic (Balb/c IL-13^{+/+} or Balb/c IL-13^{-/-}) donors. Compared to syngeneic controls, T cells from allogeneic IL-13^{+/+} and IL-13^{-/-} expanded significantly and equally in the spleen following BMT (Supplemental Figure 2a). Serum levels of IFN γ , IL-2, IL-5 and IL-13 were all elevated in mice 7 days after allo-BMT from IL-13^{+/+} donors compared to syngeneic controls. Concentrations of IL-13 and IL-5, but not IFN γ or IL-2 were significantly reduced following IL-13^{-/-} BMT compared to levels measured in allogeneic IL-13^{+/+} controls (Supplemental Figure 2b–e). Serum IL-4 levels were not detectable in any group.

Th2 cytokine responses have been shown to abrogate the intensity of aGVHD in vivo [17]. We hypothesized that the reduction in Th2 cytokine production following IL-13^{-/-} BMT would correlate with increased aGVHD severity. As shown in Figure 1a, 90% of recipients of allo-BMT from IL-13^{-/-} donors died of aGVHD by day 150. By contrast, animals receiving allo-BMT from IL-13^{+/+} donors showed a significant reduction in aGVHD mortality (P < 0.02). Mortality was associated with increased serum TNFa levels after allo-BMT, and TNFa levels were significantly higher after allo-BMT with IL-13-/- cells compared to recipients of allogeneic IL-13^{+/+} BMT (Fig. 1b). We hypothesized that the enhanced mortality seen following allo-BMT with IL- $13^{-/-}$ donors were secondary to the loss of the immunomodulatory effects of IL-13 on macrophage function and TNFa secretion [18]. To test this, we pre-incubated B6 peritoneal cells prior to LPS stimulation with either recombinant IL-13, or supernatants collected from MLRs using either IL-13^{+/+} or IL- $13^{-/-}$ T cells. The latter design was chosen to reproduce in vitro, aspects of immune dysregulation that were occurring in vivo following allo-BMT. As shown in Figure 1c, the addition of recombinant IL-13 to cell culture prior to LPS stimulation resulted in a significant reduction in TNF α production by peritoneal cells in a dose-dependent fashion. Similar reductions were observed when peritoneal cells were pre-incubated with supernatants collected from MLRs using T cells capable of producing IL-13 (IL- $13^{+/+}$) compared to the conditioned media collected from T cells that could not (Fig. 1d). Collectively, these data suggest that donor derived IL-13 down-regulates TNFa production following allo-BMT. This effect in combination with the ability of IL-13 to augment the secretion of Th2 cytokines such as IL-4 or IL-5 or other immunomodulatory proteins like

TGF β [19–21] may contribute to a protective role of IL-13 in aGVHD pathophysiology. The previously described correlation between the in vitro production of IL-13 by donor T cells and clinical aGVHD [5] offers an interesting approach to predict GVHD pre-transplant. However, since alloreactive T cells are a significant source of IL-13 (Supplemental Figure 1d), this correlation may simply reflect the overall allo-activation status of these cells rather that a direct link between IL-13 and GVHD severity. While our data support this hypothesis, it is possible that IL-13 could directly enhance target organ injury in a Th2mediated manner [4]. Whether proinflammatory effects of IL-13, such as those seen with airway hyperresponsiveness, eosinophilia and fibrosis [19,22,23], play a functional role in chronic GVHD remains to be determined.

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Identification of A Novel p53 In-Frame Deletion in a Li-Fraumeni-Like Family

Joshua D. Schiffman, MD,¹* Nicolette Chun, MS,² Paul G. Fisher, MD,¹ Gary V. Dahl, MD,¹ James M. Ford, MD,² and Faye A. Eggerding, MD, PhD³

We describe a 2-year-old female with a completely resected cerebral pilocytic astrocytoma who subsequently developed Bprogenitor acute lymphoblastic leukemia (ALL). Her father and paternal uncle were previously diagnosed with glioblastoma multiforme. Sequence analysis of the patient's p53 gene revealed a novel germline three base-pair deletion (339_341delCTT) in exon 4, resulting in removal of an evolutionarily conserved phenylalanine amino acid residue at codon 113. The same mutation was found in the patient's two clinically unaffected siblings. The in-frame deletion we describe has not previously been reported and adds to our understanding of the biologic effects of p53 gene mutation in Li–Fraumeni syndrome (LFS). Pediatr Blood Cancer 2008;50: 914–916. © 2007 Wiley-Liss, Inc.

Key words: brain tumors; leukemia; Li-Fraumeni-like syndrome p53 gene

INTRODUCTION

Li–Fraumeni syndrome (LFS) is a rare familial cancer syndrome originally identified through analysis of rhabdomyosarcoma patients and their families [1]. Classic LFS is characterized by bone or soft tissue sarcoma before age 45 years in an individual with a first degree relative diagnosed with cancer before age 45, and one additional first or second degree relative diagnosed with cancer at an early age or sarcoma at any age [2–4]. Families that exhibit a clustering of LFS cancers but do not conform to all the features of classic LFS are labeled Li–Fraumeni-like (LFL) families. LFL syndrome is characterized by occurrence of a LFS-related cancer before age 45 years in an individual with two first or second degree relative with LFS-related malignancies [5].

Germline mutations in the p53 tumor suppressor gene located on chromosome 17p13 have been identified in 70–80% of LFS families and in 20–40% of LFL families [2,4,6,7]. Germline and somatic p53 mutations are similar, with the majority missense substitutions resulting in defective transcriptional regulation. Inactivating germline mutations in the p53 gene are highly penetrant and individuals harboring p53 mutations have an elevated risk for multiple primary cancers [8]. Here we report the identification of a novel p53 germline mutation in a child with pilocytic astrocytoma and leukemia, in a LFL family with a clustering of brain tumors.

CLINICAL REPORT

A previously healthy 2 1/2-year-old presented with a seizure. Brain MRI revealed a left parietal tumor. The tumor was completely resected and found to be a pilocytic astrocytoma (WHO grade I). No adjuvant therapy was necessary.

At age 4 years, she presented with pallor, hepatosplenomegaly, and fever. Peripheral blood white cell count was 21,800/mm³ with 90% lymphoblasts. Bone marrow aspirate confirmed the diagnosis of B-progenitor cell acute lymphoblastic leukemia (ALL). Cytogenetic analysis identified the TEL/AML-1 fusion gene in lymphoblasts. There was no central nervous system disease. After two years of treatment per Pediatric Oncology Group protocols 9900 and 9904 regimen D, she is now on maintenance chemotherapy.

A family history revealed the death of the patient's father secondary to glioblastoma multiforme at age 46. The patient's paternal uncle died secondary to glioblastoma at age 48. The patient's mother is healthy. The maternal grandmother developed breast cancer at age 60, and died at age 61 from heart failure. A family pedigree is in Figure 1. A diagnosis of LFL syndrome was considered. The mother consented to p53 genotyping for the patient

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¹Division of Pediatric Hematology/Oncology, Stanford University School of Medicine, Stanford, California 94304; ²Division of Oncology, Stanford University School of Medicine, Stanford, California 94304; ³Huntington Medical Research Institutes, Pasadena, California 91101

^{*}Correspondence to: Joshua D. Schiffman, Division of Pediatric Hematology/Oncology, Stanford University School of Medicine, 1000 Welch Road, Suite 300, Stanford, CA 94304. E-mail: jschiffman@lpch.org