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Activation, Immune Polarization, and Graft-versus-Leukemia Activity of Donor T-cells are Regulated by Specific Subsets of Donor Bone Marrow Antigen-Presenting Cells in Allogeneic Hematopoietic Stem Cell Transplantation¹

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Abstract

We investigated the roles of specific subsets of donor APCs purified from bone marrow in donor T cell activation and graft-vs-leukemia (GvL) activity in murine models of hemopoietic stem cell transplantation. Lineage⁻CD11c⁺ APC precursors were separated from donor bone marrow based on expression of CD11b. Transplanting lineage⁻CD11c⁺CD11b⁻ APC (CD11b⁻ APC) in combination with c-kit+Sca-1+lineage- hemopoietic stem cells (HSC) and congenic donor T cells led to increased donor CD4⁺ and CD8⁺ T cell proliferation and higher donor T cell chimerism than with transplanting grafts containing HSC, T cells, and lineage⁻CD11c⁺CD11b⁺ APCs (CD11b⁺ APC), or grafts containing only HSC and T cells. Transplanting CD11b⁻ APCs induced Th1/type 1 cytotoxic T lymphocyte donor T cell immune polarization and enhanced GvL activity of donor T cells without increased graft-vs-host disease in both MHC- and minor histocompatibility Ag-mismatched murine hemopoietic stem cell transplantation models, whereas CD11b⁺ APCs led to Th2/type 2 cytotoxic T lymphocyte donor T cell immune polarization. Donor CD11b⁻ APCs were plasmacytoid dendritic cell progenitors (>90% CD317; PDCA-1⁺) and upregulated CD80, CD86, and IL-12 during alloantigen presentation, whereas CD11b⁺ APCs expressed Gr-1 and up-regulated expression of programmed death ligands-1 and 2 after activation. These results are the first to show that manipulation of the content of donor APCs in allogeneic HSC grafts can regulate donor T cell immunity and enhance GvL without increasing graft-vs-host disease activity.

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Disclosures

E.K.W. has nonlicensed patents in China and Australia that describe the potential clinical utility of manipulating the DC content of hemopoietic progenitor cell allografts. The other authors declare no competing financial interests.

Keywords

Antigen presenting cells; Dendritic Cells; T-cells; Graft versus Host Disease; Cell Activity; Tumor Immunity

Introduction

Graft-vs-host disease $(GvHD)^3$ and relapsed leukemia are the primary complications of hemopoietic stem cell transplantation (HSCT) in patients with leukemia. Both GvHD and graft-vs-leukemia (GvL) reactions require APCs to activate T cell effectors. Host APCs persist after high-dose chemotherapy (1), HSCT, and HSCT (2, 3) and initiate GvHD in mouse models of HSCT (4, 5, 6).

In contrast to the requirement for host-type APC in the initiation of GvHD, the role for donor APC on transplant outcomes is less clear (7). Experiments using MHC class II (MHC-II)-deficient radiation chimeras have demonstrated a requirement for host APCs on the initiation of CD4⁺ T cell-mediated GvHD (5) and a role for donor APC in increasing the severity of CD8⁺ T cell-dependent acute GvHD (5) and intestinal chronic GvHD (8). In allogeneic bone marrow (BM) transplant (BMT) from HLA-matched siblings, larger numbers of donor plasmacytoid dendritic cells (pDC) in allogeneic BM grafts were associated with more relapse and worse survival (9, 10). Additionally, higher numbers of donor dendritic cells (DC) in the blood of transplant recipients and higher serum IL-12 levels are associated with less relapse and less GvHD after allogeneic HSCT (11). Although these clinical studies suggest a role for donor APCs and DC in transplant outcomes, they do not formally evaluate the relative contribution of one donor cell subset over another. In this study, we tested the effects of transplanting phenotypically defined subsets of murine donor BM APCs on immune reconstitution following allogeneic HSCT.

Mouse APC and DC (the most potent type of APC) progenitors are contained within the population of CD11c⁺ cells that lack lineage (Lin) markers (12, 13, 14, 15). Specific DC subpopulations can be further defined by their expression of B220, CD11b, CD4, and CD8 (12). We have previously demonstrated that donor BM depleted of CD11b⁺ cells by MACS has an enhanced ability to polarize donor T cells to Th1 immunity and that tumor-bearing transplant recipients of allogeneic CD11b⁻ depleted BM cells had prolonged survival compared with recipients of unmanipulated BM (16). These data suggested that phenotypically defined donor populations could modulate alloreactivity and GvL activity of donor T cells in allogeneic HSCT (17) and raise the interesting question of whether specific subsets of donor APCs could affect immune reconstitution in allogeneic HSCT.

To test this hypothesis, we used defined populations of donor APCs purified by FACS from donor BM to clarify their role in donor immunity following allogeneic HSCT. The expression of CD11b was used to divide Lin⁻CD11c⁺ BM APC into two distinct populations: a fairly homogeneous population of Lin⁻ CD11c⁺ CD11b⁻ APC precursors that are predominantly B220⁺CD11b⁻. and CD317⁺ (plasmacytoid DC Ag (PDCA)-1⁺) pDC precursors (18), and a more heterogeneous Lin⁻CD11c⁺CD11b⁺ population that includes myeloid DC progenitors as well as myeloid suppressor cells (16, 19, 20). We predicted that the murine CD11b⁻ APC precursors would augment posttransplant cellular immune

³Abbreviations used in this paper: GvHD, graft-vs-host disease; HSCT, hemopoietic stem cell transplantation; MHC-II, MHC class II; BM, bone marrow; DC, dendritic cell; GvL, graft-vs-leukemia; B6, C57BL/6; B10, B10.BR; B/c, BALB/c; Lin, lineage; MiHA, minor histocompatibility Ag; Syn, syngenic; pDC, plasmacytoid DC precursor; PD-L, programmed death ligand; PD-L2, programmed death ligand-2; Tc, cytotoxic T lymphocyte donor T cell; PDCA, plasmacytoid DC Ag; Tc, cytotoxic T lymphocyte donor T cell; HSC, hematopoietic stem cell.

responses based on their ability to polarize cognate T cells toward Th1 immune responses (21, 22, 23, 24, 25) and that CD11b⁺ APC precursors would polarize T cells toward Th2 or mixed Th1/Th2 responses (23, 26). We herein demonstrate a striking ability of FACS-purified donor APC to regulate posttransplant immunity including the proliferation, cytokine synthesis, and antitumor cytotoxic activity of donor T cells.

Materials and Methods

Mice

B10.BR (B10, H-2K^k), PL/J (H-2K^s), BALB/c (B/c, H-2K^d), C3H.SW (H-2K^b), C57BL/6 (B6, H-2K^b) and congenic B6.SJL (H-2K^b, CD45.1, CD90.2) mice were purchased from The Jackson Laboratory. Congenic strains expressing CD90.1 and CD45.1 on a B6 (H-2K^b) background and CD90.1 and CD45.2 on a B10 (H-2K^k) background were bred at Emory University (Atlanta, GA). GFP-expressing B6 mice were a gift from Dr. Robert Taylor (Emory University).

Tumor cells

MMB3.19, a retrovirus-transformed myeloid leukemia line from B6 (27), was provided by Dr. R. Korngold (Jefferson Medical University, Philadelphia, PA). LBRM 33-5A4, a B10 T cell lymphoma cell line (28), was purchased from American Type Culture Collection (ATCC). Tumor cell lines were cultured according to ATCC recommendations, and per tests were free of lymphocytic choriomeningitis virus, mouse hepatitis virus, minute virus of mice, and mouse parvovirus by the University of Missouri Research Animal Diagnostic Laboratory (Columbia, MO).

Cell preparations

Donor mice were killed in a humane manner, and femurs, tibias, and spleens were removed aseptically. BM cells and splenocytes were harvested with sterile RPMI 1640 containing 1% heat-inactivated FCS. For purification of APC subsets, BM cells were incubated with anti-CD11c microbeads (Miltenyi Biotec) followed by anti-CD11c-allophycocyanin, CD11ballophycocyanin-Cy7, and a mixture of PE-conjugated anti-CD3, IgM, CD19, DX5, and TER119 Abs (BD Biosciences). CD11c⁺ cells were then selected using LS magnetic columns (MACS; Miltenyi Biotec), and Lin⁻CD11c⁺CD11b⁺ and Lin⁻CD11c⁺CD11b⁻ populations were sorted using the FACSAria (BD Biosciences). For hematopoietic stem cell (HSC) selection, BM cells were stained with biotinylated Abs to Lin markers CD11b, Gr-1, CD3, CD4, CD8, DX5, B220, I-A^b, and TER119 (BD Biosciences) and antibiotin microbeads (Miltenyi Biotech). Lin⁻ cells were collected after MACS magnetic separation and stained with Abs to the stem cell factor receptor c-kit and to stem cell Ag (Sca)-1 (BD Biosciences); then Lin⁻c-kit⁺Sca-1⁺ HSCs were sorted by FACS. BM from one donor mouse provided sufficient sorted stem cells for three recipients and APC for five recipients. T cells were purified by incubating splenocytes with biotinylated anti-CD11b, B220, DX5, and TER119 Abs, followed by antibiotin microbeads, and negative MACS selection using an LS column. CD8 T cells were obtained by the addition of biotinylated anti-CD4 Ab to the T cell-negative selection mixture.

Characterization of BM APCs

MHC-II, CD40, CD80, CD86, ICOS ligand (ICOS-L), programmed death ligand-1 (PD-L1), and PD-L2 costimulatory molecule expression, and CD317, Ly-6C/G (Gr-1), F4/80, CD115, CD135, CD90, and NK1.1 lineage-related molecule expression on BM APCs were analyzed by flow cytometry using directly conjugated Abs as described by the manufacturer (BD Biosciences). For in vitro experiments, FACS-purified BM APCs (2×10^{5} /ml) were cultured

with CD40L (1 µg/ml; PeproTech); with 1×10^{6} /ml irradiated (11 Gy) allogeneic splenocytes; with 1×10^{6} /ml syngeneic splenic T cells; with 1×10^{6} /ml irradiated allogeneic splenocytes plus 1×10^{6} /ml syngeneic splenic T cells; or with medium alone for 72 h in RPMI 1640 with 10% FCS. Flow cytometric analyses of costimulatory molecule expression were performed before and after the culture period. To analyze the maturation of BM APCs in vivo, mice were transplanted with 1×10^{6} FACS-sorted APCs from GFP⁺ transgenic donors plus HSCs and donor T cells and then sacrificed 10 days posttransplant. Mononuclear cells were harvested from the BM and spleen and analyzed for Ag expression by flow cytometry, after electronically gating on GFP⁺ donor APC cells in list mode files of at least 1,000,000 events.

Transplantation

Recipient B10, PL/J, or B6 mice were irradiated with two doses of 5.5 Gy separated by 3 h on day -2 (29). On day 0, recipient mice were transplanted with combinations of 3×10^3 HSCs, with varying numbers of APCs (5×10^4 or 1×10^6), and T cells (3×10^5 or 3×10^6) using B6 and C3H.SW donors in the B6 \rightarrow B6, B6 \rightarrow B10, B6 \rightarrow PL/J, and C3H.SW \rightarrow B6 models. For tumor experiments, B10 mice received an i.v. dose of 1×10^5 cells of viable (Ficolled, from log-phase culture) LBRM (28) on day -1 (16, 30), whereas B6 recipients received an i.p. dose of 5×10^4 MMB3.19 cells on day -1 (27). Mice were weighed twice weekly and examined daily for signs of GvHD as described (31). Moribund animals losing >25% of initial body weight and mice surviving until the end of the experiment were euthanized, and tissues were processed for histopathological analysis of tumor-trophic sites including brain, lung, liver, spleen, and kidney. Flow cytometric chimerism analyses were performed on blood leukocytes on (mean \pm SD) days 10, 30 ± 1 , 60 ± 2 , and 100 ± 5 posttransplant.

T cell proliferation, cytokine expression, and donor cell localization

The proliferation of donor T cells in recipient spleen was analyzed by CFSE dilution as described (32). Briefly, donor T cells were stained with CFSE before transplant and recipient spleens were removed 3 days later, and cell suspensions were prepared. Proliferation of donor T cells was determined by flow cytometric analysis of CFSE dilution profiles. In secondary MLR, responder T cells were recovered 15 days posttransplant, stained with CFSE before setting up the MLR culture, and analyzed 3 days later. Intracellular cytokine expression (IL-4, IL-10, IL-17, and IFN- γ ; BD Biosciences) by CD4⁺ and CD8⁺ T cells was analyzed as described (30). In transplant experiments using GFP⁺ donors, the presence of GFP⁺ donor DCs in frozen sections of spleens, mesenteric lymph nodes, and ileum tissues was visualized by confocal microscopy (Zeiss LSM 510; Thornwood) using Zeiss LSM 5 Image Browser software with images captured using ×40 and $\times 100$ objectives. Donor T cells were visualized using rat anti-mouse CD90.2 (BD Biosciences) and goat anti-rat IgG Alexa Fluor 555 (Invitrogen) with 4',6'-diamidino-2phenylindole to stain nuclei. The number of APCs per microliter of spleen tissue was determined by averaging the APCs found in 6-µm-thick sections of tissue (volumes calculated as area × thickness) from three mice per group in each of three separate transplant experiments.

Measurement of serum and secreted cytokine levels

Recipient mice were anesthetized by isofluorane inhalation, and peripheral blood was collected by tail vein bleeding into Microtainer brand serum separation tubes (BD Biosciences). After centrifugation, the serum was stored at -20° C, and cytokines were assayed in duplicate wells using OptEIA ELISA sets (IL-2, IL-10, IFN- γ , IL-12 p70, and TNF- α ; BD Biosciences) and Ready-SET-Go ELISA kits (IL-4, IL-5, IL-23; eBioscience) and analyzed using a SpectraMax 340PC spectrophotometer (Molecular Devices).

CTL activity

Recipients of C3H.SW \rightarrow B6 or B6 \rightarrow B10 transplants were euthanized on day 34 or 81, respectively, and donor T cells were isolated by MACS-negative selection of splenocytes using a mixture of Abs to CD11b, DX5, B220, and TER119 (BD Biosciences). To isolate spleen-derived donor T cells from B6 \rightarrow B10 transplant recipients, anti-CD45.2 (BD Biosciences) was added to the Ab cocktail to deplete host and HSC-derived T cells. In the C3H.SW \rightarrow B6 transplant, S90% of splenic T cells were derived from the T cells in the graft on day 34 posttransplant. T cells (2 × 10⁶/ml) were cultured with 2 × 10⁶ cells/ml irradiated (30 Gy) allogeneic host-type splenocytes or tumor cells in 24-well plates in RPMI 1640 supplemented with 10% FCS and antibiotics. After 5 days, viable T cells were collected and assayed for cytotoxic activity against LBRM and MMB3.19 tumor cells by a flow cytometry assay using the CyToxiLux PLUS kit (OncoImmunin). Briefly, 2 × 10⁶ effector cells were mixed with 2 × 10⁵ surface-labeled target cells and incubated with caspase substrate for 30 min, washed twice with PBS, and the percentage of apoptotic target cells was calculated after flow cytometry analysis.

Statistical analyses

Analyses of data were performed using SPSS (version 17 Mac, SPSS Inc. Chicago, IL). Data are presented as mean \pm SD. Survival differences between groups were calculated with the Kaplan-Meier log rank test in a pairwise manner. Differences in GvHD outcome between groups were compared using the Kruskal-Wallis test. Differences in the levels of donor T cells and other parametric tests were compared using the one-way ANOVA.

Results

Addition of BM CD11b⁻ APCs to HSC grafts enhanced donor CD4 and CD8 T cell proliferation

To test the hypothesis that the activation status of donor T cells can be modulated by the presence of donor APC in the graft, we used two MHC-mismatched allogeneic HSCT models (B6 \rightarrow B10 and B6 \rightarrow PL/J), as well as MHC-matched syngenic controls (B6 \rightarrow B6). Using high-speed FACS, we enriched APC from BM by selecting cells lacking lineage markers (TER119, CD3, DX5, CD19, and IgM) and expressing CD11c, and then sorted Lin⁻CD11c⁺CD11b⁻ APCs and APC precursors (CD11b⁻ APC) and Lin⁻CD11c⁺CD11b⁺ APCs and APC precursors (CD11b⁺ APCs) to average purities (\pm SD) of 97 \pm 1% and 93 ±2%, respectively. Using separate congenic strains as donors of HSC and T cells, Lin⁻ckit⁺Sca-1⁺ HSC were purified by FACS from donor BM and T cells were purified by MACS from splenocytes with average purities of $90 \pm 1\%$ and $95 \pm 2\%$, respectively (Fig. 1, A-F). The CD11b⁻ and CD11b⁺ APC subpopulations had the scatter properties of lymphocytes (Fig. 1, B and C). The majority of BM CD11b⁻ APCs expressed B220 (72%), CD90 (51%), and CD317 (93%), had low levels of MHC-II, partial expression of CD4, and lacked expression of other markers including CD24, CD80, and CD86 as well as markers associated with lymphoid cell precursors and NK cell (CD135, NK1.1) or myeloid cell differentiation (F4/80, Gr-1, and CD115; Fig. 1G). In contrast, the majority of BM CD11b⁺ APCs lacked B220, CD4, CD8, CD24, CD80, CD86, MHC-II, and CD317 but expressed high levels of F4/80 and low levels Gr-1, consistent with recently described BM myeloid suppressor cells (Refs. 16, 19, and 20; Fig. 1G). Applying the same phenotypic analysis to Lin⁻CD11c^{low/+} cells in the spleen revealed three distinct populations: the same two populations of CD11clowCD11b⁺ APCs and CD11clowCD11b⁻ APCs described above and a third population of Lin⁻CD11c^{high} cells that expressed high levels of CD4, CD80, CD86, and MHC-II which has been previously described as classical splenic DCs (Refs. 33 and 34 and supplemental Fig. 1).⁴

Next we measured the in vivo proliferation of CFSE-labeled donor T cells 3 days posttransplant in mice that received either CD11b⁻ APCs or CD11b⁺ APCs. We transplanted 3×10^3 HSCs combined with 3×10^5 CFSE-labeled congenic CD90.1⁺ T cells and 5×10^4 FACS-sorted CD11b⁻ APCs or CD11b⁺ APCs. The proliferation of donor T cells in syngeneic recipients were equivalent regardless of the presence or absence of donor APCs (Fig. 2, *A* and *B*; only the HSC plus T cells data are shown for clarity). In B6 \rightarrow B10 allogeneic recipients, there was greater initial proliferation of donor CD8⁺ T cells than of donor CD4⁺ T cells, and cotransplantation of CD11b⁻ APCs led to higher proliferation rates of both donor CD4⁺ and CD8⁺ T cells compared with the proliferation of corresponding donor T cell subsets cotransplanted with CD11b⁺ APCs, or T cells transplanted into allogeneic mice that received no donor APCs (Fig. 2, *A* and *B*). After 3 days, CD4⁺ and CD8⁺ donor T cells recovered from recipients of CD11b⁻ APCs expressed higher levels of CD25 and CD69 (data not shown) and Ki-67 (Fig. 2*C*) by FACS compared with T cells cotransplanted with HSCs alone or the combination of T cells, HSCs, and CD11b⁺ APCs.

The kinetics of donor T cell engraftment was analyzed by sampling blood on days 10, 30, 60, and 100 posttransplant and using congenic markers to distinguish T cells derived from donor HSCs (CD90.2⁺CD45.1⁺), donor BM APC subsets (CD90.1⁺CD45.2⁺), MACS-purified mature T cells in the graft (CD90.1⁺CD45.1⁺), and residual host T cells (CD90.2⁺CD45.2⁺). There was significantly greater expansion of donor T cells derived from the T cells in the graft among mice that received CD11b⁻ APCs, compared with those that received CD11b⁺ APCs or grafts containing donor HSC and T cells without APC (Fig. 2*D*). In contrast, expansion of T cells derived from donor HSC was not significantly different among the groups (Fig. 2*E*). Persistence of host-type T cells was seen only among recipients of HSCs alone, consistent with mixed chimerism in the absence of donor T cells (Fig. 2*F*). Among recipients of CD11b⁻ APCs, the graft-derived donor T cells were predominantly CD8⁺ (CD4:CD8 0.67:1) with an effector memory phenotype, compared with donor T cells from recipients of HSCs plus T cells (CD4:CD8 1.16:1), whereas the comparable donor T cells among recipients of CD11b⁺ APCs were predominantly CD4⁺ (CD4:CD8 1.28:1; *p* < 0.02) suggesting preferential activation of CD8⁺ donor T cells by CD11b⁻ donor APCs.

Transplantation of purified CD11b⁻ APCs promotes Th1 and cytotoxic T lymphocyte donor T cell (Tc) type 1 immune polarization of donor T cells

To characterize the effects of donor APC on the immune polarization of donor T cells, we measured intracellular synthesis of cytokines (IL-4, IL-10, IL-17, and IFN- γ) in donor (CD45.1⁺CD90.1⁺) CD4⁺ and CD8⁺ T cells on day 10 posttransplant and serum levels of IL-2, IL-4, IL-5, IL-10, IL-12, IL-23, IFN- γ and TNF- α . Recipients of CD11b⁻ APC had higher frequencies and total numbers of Th1/Tc1 donor CD4⁺ and CD8⁺ T cells, lower numbers of Th2/Tc2 donor T cells, and significantly higher serum levels of Th1 cytokines compared with recipients of CD11b⁺ APC in which Th2/Tc2-polarized immune responses predominated (Fig. 3). There were no differences in frequencies of IL-17⁺ T cells comparing recipients of different donor APC preparations (data not shown).

To test whether $B6 \rightarrow B10$ T cell transfer resulted in activation of donor T cells specifically responding directly to host APCs (H2^k, host restricted), we assessed the proliferative capacity of B6 donor T cells harvested 15 days posttransplant from B10 recipients in oneway MLR cultures. Donor T cells were harvested from HSCT recipients, stained with CFSE, and cultured for 3 days with irradiated syngeneic B6 splenocytes, recipient-type B10 splenocytes, or third-party BALB/c splenocytes. For all groups, donor T cells had minimal

⁴The online version of this article contains supplemental material.

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proliferation when cultured with syngeneic B6 or third-party BALB/c irradiated splenocytes (data not shown). In MLRs with recipient-type B10 irradiated splenocytes, donor CD4 and CD8 T cells from recipients of CD11b⁻ APCs proliferated more than T cells from recipients of HSCs and T cells or recipients of HSCs, T cells, and CD11b⁺ APCs (supplemental Fig. 2*A* and *B*). Donor T cells from recipients of CD11b⁻ APCs were Th1/Tc1 polarized compared with Th2/Tc2-polarization of T cells from recipients of CD11b⁺ APCs (supplemental Fig. 2*C* and *D*).

Donor APC and donor T cells colocalize in peripheral lymphoid tissues

FACS-purified CD11b⁺ APC or CD11b⁻ APCs from GFP-transgenic B6 donor mice were transplanted along with non-GFP B6 CD90.1⁺ HSCs and CD90.2⁺ T cells into lethally irradiated CD90.1⁺ B10 recipients. No GFP signal was observed in spleens of control recipients of HSCs or HSCs and T cells (Fig. 4, *A* and *B*). Equal numbers of GFP⁺ donor CD11b⁺, and CD11b⁻ APCs with visible dendrites were seen in close physical proximity to donor T cells (and unstained host mononuclear cells) in sections of recipient spleen on days +3 and +10 posttransplant (Fig. 4, *C*–*E*), and the numbers of both APC subsets increased in the recipient spleen from day +3 to day +10 posttransplant over this time (Fig. 4*F*), consistent with in situ proliferation and/or continued migration of donor GFP⁺ APCs from other sites (35). GFP⁺CD11b⁻ APCs and CD11b⁺ APCs were also observed in proximity to donor T cells in mesenteric lymph nodes and Peyer's patches of transplant recipients (data not shown). These observations suggest equivalent abilities of both donor APC subsets to home to secondary lymphoid organs and argue that increased proliferation and Th1 polarization of donor T cells cotransplanted with donor CD11b⁻ APCs is not due to differences in microanatomic localization.

Differential development of APC precursors and expression of costimulatory molecules on CD11b⁺ APCs vs CD11b⁻ APCs

To explore the impact of CD11b⁻ and CD11b⁺ APC precursor development and differentiation on donor immune polarization, we tracked expression of lineage markers on GFP⁺ donor cells in recipient BM and spleen 10 days after transplantation of GFP⁺CD11c⁺CD11b⁻ or CD11b⁺ APCs. The frequencies of GFP⁺CD11c⁺CD11b⁻ and CD11b⁺ APCs were similar in BM (2.2 ± 0.2 and $1.8 \pm 0.3\%$, respectively; Fig. 5A) and spleen (data not shown). Based on scatter plots, GFP⁺CD11b⁺ APCs had higher side scatter, consistent with myelomonocytic differentiation compared with GFP⁺CD11b⁻ APC (Fig. 5A). Analysis of donor GFP⁺CD11b⁻ APCs in recipient BM showed similar lineage expression patterns compared with cells taken directly from BM, whereas donor GFP+CD11b⁻ APCs in recipient spleen showed moderately decreased frequencies of cells expressing B220, Gr-1, and CD317. In contrast, the frequency of GFP+CD11b+ APCexpressing F4/80 and Gr-1 were slightly increased in the spleen compared with GFP⁺CD11b⁻ cells recovered from BM (Figs. 1G and 5B). Expression of CD3 and NK1.1 on GFP⁺ donor cells isolated from recipients of either GFP⁺CD11b⁻ APCs or GFP⁺CD11b⁺ APCs was absent (data not shown). GFP+CD11b- APCs recovered from the spleens and BM of allogeneic transplant recipients had higher levels of MHC-II, CD80, and CD86, as well as lower levels of PD-L1 and PD-L2, compared with the GFP⁺ progeny of CD11b⁺ APCs (Fig. 5C). Low levels of expression of CD40 were seen only on CD11b⁺ APCs, and ICOS-L was largely absent on both CD11b⁻ and CD11b⁺ APCs (data not shown). There were higher levels of costimulatory and coinhibitory molecule expression on GFP⁺ donor APCs recovered from recipient spleen vs recipient BM, indicating that factors in the tissue microenvironment may also have a significant role in the maturation of transplanted donor APCs. In parallel, FACS-purified CD11b⁺ and CD11b⁻ APCs were analyzed for MHC-II, CD80, CD86, PD-L1, and PD-L2 expression before and after stimulation with CD40L, irradiated allogeneic splenocytes, and the combination of syngeneic T cells and allogeneic

Ags. In vitro stimulation with CD40L (36, 37) or alloantigen with syngeneic T cells led to marked up-regulation of MHC-II, CD80, and CD86 expression on CD11b⁻ APC, and up-regulation of PD-L1 and PD-L2 on CD11b⁺ APC (Fig. 5*D*). The level of expression of activation and maturation markers on donor APC in vivo were lower than the levels of the same markers expressed on APC populations cultured in vitro with CD40L (Fig. 5, *C* and *D*).

Donor CD11b⁻ APCs augmented GvL activity of donor T cells without increasing GvHD

We next tested the effect of varying numbers of donor T cells transplanted with CD11b⁺ APCs or CD11b⁻ APCs on GvHD in the MHC-mismatched ($B6 \rightarrow B10$) murine allogeneic HSCT model. As expected, recipients of a larger dose of donor T cells (3×10^6) developed lethal GvHD compared with minimal GvHD among recipients of a lower donor T cell dose $(3 \times 10^5;$ Fig. 6, A and B). Clinical GvHD scores and body weight loss for the recipients of 3 $\times 10^5$ T cells plus CD11b⁻ APCs were low and not significantly different from other treatment groups with the same T cell dose and different donor APCs (Fig. 6, B and C); mean clinical GvHD scores for recipients of 3×10^6 T cells were much higher but did not vary significantly among groups with different donor APCs (Fig. 6B). In the MHC-matched, clinical sores, and body weight loss between recipients of different grafts in the absence of tumor (Fig. 6, D–F). To test whether donor CD11b⁻ APC could augment the GvL activity of low-dose T cells, we transplanted 3×10^5 donor T cells and 3000 HSC with 5×10^4 amounts of either population of donor BM APCs, using two distinct allogeneic HSCT models. Hosttype tumor cells were injected one day before HSCT and following total body irradiation. In the MHC-mismatched $B6 \rightarrow B10.BR$ model, recipients of CD11b⁻ APCs had 45% durable survival (Fig. 6G) compared with <5% survival among tumor-bearing recipients of HSCs alone; HSCs plus T cells; or HSCs, T cells, and CD11b⁺ APCs (p < 0.001). Necropsy of moribund B10.BR recipient mice that were euthanized showed a lower incidence of detectable leukemia among recipients of CD11b⁻ APCs compared with other groups (Fig. 6H). In the C3H.SW \rightarrow B6 MiHA-mismatched model, recipients of CD11b⁻ APCs had 35% long-term survival compared with uniform early mortality among other treatment groups (Fig. 6*I*). The GvL activities in the C3H.SW \rightarrow B6 model observed using mixed CD4⁺ and CD8⁺ donor T cells or purified CD8 T cells were identical (data not shown), indicating that our transplant model is comparable with others that use only CD8 T cells (4, 17).

Donor CD11b⁻ APCs enhanced T cell proliferation and CTL activity ex vivo

To further address mechanisms whereby CD11b⁻ APC enhanced the GvL activity of lowdose donor T cells, we assessed the cytotoxic activity of donor T cells recovered from recipients of CD11b⁻ APCs compared with recipients of other graft combinations. Wellappearing recipients of MHC-mismatched and MiHA-mismatched transplants were euthanized on days 81 and 34 posttransplant, respectively, and donor T cells derived from the mature donor T cells in the original transplant graft were selected by MACS using the CD45.1 congenic marker. Donor T cells recovered from recipients of CD11b⁻ APCs had increased proliferation in a one-way MLR following culture with irradiated host-type allogeneic splenocytes compared with donor T cells recovered from recipients of CD11b⁺ APCs or T cells from recipients without donor APCs (Fig. 7, *A* and *B*). In addition, antileukemia cytotoxic activity was significantly greater among T cells recovered from recipients of CD11b⁻ APCs than among T cells from other treatment groups (Fig. 7, *C* and *D*).

IFN-y does not directly kill leukemia cells

To rule out the possibility that the observed GvL effect of CD11b⁻ APCs was due to direct antitumor effects of IFN- γ , we cocultured LBRM or MMB3.19 cells (2 × 10/ml) with various concentrations of IFN- γ . In vitro exposure of leukemia cells to IFN- γ , at doses of 10–300 pg/ml, similar to concentrations observed in vivo, demonstrated neither cytotoxicity nor long-term growth-inhibitory effects on either leukemia cell line over 5 days of culture (data not shown).

Discussion

Our findings indicate that distinct subsets of donor APC are associated with quite different immunological effects posttransplant. Using MHC-mismatched and MiHA-mismatched HSCT model systems, leukemia-bearing recipients of CD11b⁻ APC had substantially enhanced survival compared with recipients of HSCs alone, HSCs plus T cells, or HSCs plus CD11b⁺ APCs. Significantly, the addition of CD11b⁻ APCs to grafts containing a low number of donor T cells did not significantly affect clinical GvHD mortality compared with mice transplanted with HSC and T cells (without donor APC). These findings represent, to our knowledge, the first clear demonstration that a purified subset of donor APCs can increase GvL activity without a concomitant increase in GvHD when added to grafts containing purified HSCs and T cells.

Previous studies have shown that donor T cells and DCs isolated from donors pretreated with hemopoietic cytokines or cytokine receptor agonists led to decreased GvL (38) and increased acute and chronic GvHD (38, 39). In contrast to using donor DCs mobilized by cytokines (23, 40), or DCs derived from cultured BM (41) or spleens (17), or bulk BM cells as donor APCs (4), we chose to use unstimulated populations of phenotypically defined APC subsets isolated by FACS from the BM in an attempt to recapitulate our previous clinical observations that more BM pDCs were associated with reduced GvL activity after allogeneic HSCT (42). The BM CD11b⁻ APC used in this study were nearly all pDC progenitors as evidenced by their expression of B220 and PDCA-1 (43, 44). In contrast, the CD11b⁺ APCs are heterogeneous and contained cells with the phenotypes of myelomonocytic precursors (16, 19, 20).

Although the current findings support clinical observations that donor DCs have a role in transplant outcomes, there is no direct correspondence between the immunological functions of murine vs human pDC. Our previous report on human BMT indicated that larger numbers of donor pDC suppressed GvL (42), whereas the current study, using murine BMT models, indicates that donor CD11b⁻ APCs (the majority of which are pDC) augment the GvL activity of donor T cells. Although both human and murine pDCs are important in innate immunity and synthesize large amounts of IFN- α in response to viral infection or to the binding of CpG sequences to TLR9 (24, 45), their effects on T cell immune polarization are quite different. Mature murine BM-derived pDCs (but not blood-derived pDCs) produce significant amounts of IL-12 and polarize T cells toward Th1 immune responses in vitro (21, 22, 23, 24). In contrast, CD11c⁻ human pDCs do not make significant amounts of IL-12, and they generate Th2 immunity in cognate T cells responding to direct or indirect Ag presentation (21, 46). In humans, the CD11c⁺ myeloid DCs make IL-12 and polarize responding T cells toward Th1 immune responses (46, 47). Thus, from the standpoint of IL-12 production and Th1 polarization of donor T cells, we believe that the function of the murine CD11b⁻ APCs is analogous to the CD11c⁺ human myeloid DC subset.

To delineate the mechanisms by which the CD11b⁻ subset of donor APCs enhanced GvL activity compared with CD11b⁺ APCs, we evaluated the homing and proliferation of APCs to secondary lymphoid organs on transplant recipients and equivalent numbers of both APC

subsets were seen in the spleens and lymph nodes during the first 10 days posttransplant. Both types of APC subsets freshly isolated from BM lacked activation and maturation markers MHC-II, CD80, CD86, and ICOS-L, indicating an immature level of differentiation that was not affected by FACS isolation or exposure to medium containing FBS (48, 49, 50). Donor CD11b⁻ APCs that homed to the BM retained an immature phenotype, whereas donor CD11b⁻ APCs that homed to and proliferated in the spleen up-regulated the same pattern of activation and maturation markers after in vitro exposure to CD40L, or following exposure to alloantigen and homologous T cells in one-way MLR, consistent with pDC differentiation. The levels of activation and maturation marker expression on the donor APCs recovered from allogeneic transplant recipients were lower compared with the expression of those markers in vitro, suggesting that GvHD may impair the maturation of pDC (51). The lack of increased GvHD seen among recipients of CD11b⁻ APCs are consistent with the conclusions of Banovic et al. (51) that bone marrow pDCs are immunomodulatory and may decrease GvHD. The Th1 polarization of donor T cells seen in our study was not described by Banovic, possibly due to differences in the model systems and the absence of more potent APCs (donor conventional DCs) and other accessory cells in grafts composed of purified cell subsets that were used in our studies.

Transplanting CD11b⁻ APCs resulted in a global effect of early Th1 polarization with higher serum levels of IL-12, IFN- γ , IL-2, and Th1-polarized donor T cells. The increased numbers of donor T cells with direct antitumor cytotoxicity seen with CD11b⁻ APCs in the MHC-mismatched model are consistent with cross-presentation (52) of alloantigen by CD11b⁻ APCs to donor T cells, leading to activation and subsequent activation and Th1 polarization of additional donor T cells with direct cytotoxic activity against host hemopoietic cells (including leukemia targets). Furthermore, IFN- γ did not have direct tumoricidal activity, indicating that the observed increase in GvL activity of donor T cells cotransplanted with CD11b⁻ APCs was due to their enhanced antitumor cytotoxicity. We have also shown, using IFN- γ knockout mice as T cell donors in the B6 \rightarrow B10 transplant model, that IFN- γ synthesis by donor T cells is critical to the survival of tumor-bearing recipients of CD11b⁻ APCs (Y. Lu et al., manuscript in preparation), consistent with models in which IFN- γ 53, 54, 55).

These results are in contrast with reports by Shlomchik and others in which persistent host APCs are integral to the initiation of GvHD, and donor APCs have only a weak effect on GvHD and no effect on GvL (4, 5, 17). These models used much higher doses of T cells that caused lethal GvHD. Other differences between our results and previous reports include the use of FACS-purified populations of donor BM APC in our study, whereas other studies used unfractionated BM or splenic donor APCs that may have obscured the effect of donor APC subpopulations on donor T cell immune function. A recent report by Koyama et al. (25) also show that host-type pDCs harvested from Flt2/Flk3 ligand-treated mice primed Th1 immune responses in donor T cells and induced lethal GvHD when transplanted into lethally irradiated allogeneic recipients. The higher level of GvHD observed in this study likely results from the use of larger numbers of donor T cells (2×10^6) and adoptive transfer of recipient-type Flt2/Flk3 ligand-activated pDC rather than resting BM donor pDCs in our study. The radiation doses used in the study by Koyama et al. (56) ablate endogenous hosttype pDCs, raising the question of the physiological relevance of host pDCs in the initiation of GvHD following myeloablative conditioning. Nevertheless, these results support our findings that donor pDCs can initiate Th1 immune responses in responding T cells. A recent report by Markey et al. (57) explored the role for different donor DC populations in activating donor T cells in allogeneic BMT. In this study, transgenic donor T cells were administered on day 10 posttransplant, and different subpopulations of donor DCs deleted through the administration of toxins or mAbs. The results indicate that classical CD11c⁺

DCs were most efficient in stimulating donor T cell proliferation by indirect presentation of donor alloantigens and that the role of donor pDC was minimal. Although these two model systems are quite different, both results support the importance of donor APCs in activating donor T cells. Ongoing experiments will test whether the salutorious antitumor effect of donor pDC in the experiments presented herein are due either to 1) an ability of pDCs to present alloantigen indirectly to donor T cells (in contrast to the data presented by Markey et al.) or 2) the ability of donor pDCs to modulate T cell function in response to host APCs through local production of cytokines such as IL-12 or IFN- α . Additional reasons for the improved survival seen among recipients of pDCs in our study may be that donor pDCs more efficiently activate donor T cells administered with the stem cell graft vs donor T cells administered at later time points posttransplant; that donor pDCs may facilitate donor HSC engraftment (33, 58); or that donor pDCs may induce IFN- γ production by donor T cells and thereby limit the development of GvHD (7, 53, 54, 55).

In contrast to the Th1 polarization seen with donor pDCs, transplanting CD11b⁺ APCs yielded donor T cells with Th2 polarization and higher serum IL-4, IL-5, and IL-10. Whereas the CD11b⁻ APC are nearly all pDC precursors, the CD11b⁺ APCs are more heterogeneous, and express a pattern of markers that are expressed on precursors for monocytes as well as myeloid suppressor cells. The expression of MHC-II and CD80CD86 on GFP⁺CD11b⁺ APCs recovered from transplant recipients was minimal, indicating limited differentiation to DCs. In contrast, the progeny of CD11b⁺ APC expressed high levels of immunosuppressing costimulatory molecules PD-L1 and PD-L2 (36, 59) and increased levels of Gr-1 and F4/80 expression, consistent with differentiation of some CD11b⁺ APCs into myeloid suppressor cells (16, 19, 20) or immunosuppressive monocytes (60). The expression of PD-L1 and PD-L2 on CD11b⁺ APCs is consistent with previous reports showing that CD11b⁺ donor splenocytes and CD11b⁺ donor BM cells suppress host-vs-graft alloreactivity in myeloablative and nonmyeloablative transplants, respectively (16, 32), and that murine CD11b⁺CD8a⁻ DCs result in Th2 or mixed responses (23, 26).

Some of the limitations of this study are that we did not attempt to isolate the subpopulations of pDCs with varying abilities to polarize Th1 and Th2 immunity depending on the level of the Rag1 promoter (61, 62), nor could we vary the level of alloantigen presented by donor DCs, an additional factor that has been shown to affect the ability of DC subsets to direct Th1 vs Th2 T cell polarization (63). The population of CD11b⁺ APC was clearly heterogeneous, and we cannot be certain which subpopulation is responsible for the observed Th2 polarization of donor T cells. Nevertheless, our data support a model in which different APC subsets result in distinctly different types of T cell polarization (14).

The current findings from murine HSCT models may have potential for clinical translation, as selective depletion of human pDCs from mononuclear cells isolated from blood, using a mAb to BDCA2 and immunomagnetic columns, resulted in marked enhancement of the ability of the remaining cells to activate homologous T cells to alloantigen via cross presentation (46). Taken together, these data suggest that selective manipulation of the content of donor APCs from BM grafts could be a novel method to enhance the GvL activity of allogeneic HSC transplantation.

In summary, using two distinct HSCT model systems in which donors and recipients are either MHC or MiHA mismatched, we observed highly significant Th1 polarization of donor T cells and improvements in the survival of tumor-bearing recipients of FACS-purified CD11b⁻ pDCs compared with other transplant groups. These observations indicate a role for donor APC in regulating GvL activities.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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FIGURE 1.

Isolation of purified populations of donor APC subsets, HSCs, and donor T cells using FACS and MACS. *A*, Gating strategies for sorting CD11b⁻ APC subsets and CD11b⁺ APC subsets from CD11c-enriched B6 BM. *Insets*, Histogram of lineage expression in BM before (*top*) and after (*bottom*) MACS enrichment of CD11c⁺ cells. *B*, Purity of sorted CD11b⁻ APCs. *Inset*, Purified CD11c⁺CD11b⁻ population back-gated in lymphocyte area in forward scatter (FSC) vs sideward scatter (SSC) flow cytometry plot. *C*, Purity of sorted CD11b⁺ APC. Insert: purified CD11c⁺CD11b⁺ population back-gated in lymphocyte area in a forward scatter vs sideward scatter flow cytometry plot. *D*, Gates (RUQ) for sorting Lin⁻Sca-1⁺c-kit⁺ HSCs from B6 BM. *Inset*, Histogram of lineage expression in BM after MACS enrichment of Lin⁻ cells. *E*, Purity of sorted HSCs. *F*, Analysis of T cells purified by negative MACS selection. Insert shows the percentage of CD3⁺ T cells before immunomagnetic depletion. *G*, Phenotypes of CD11b⁻ and CD11b⁺ APCs from BM. FACS plots are from a single representative experiment. Mean percentages (±SD) for each subset are shown from five replicate experiments.



FIGURE 2.

The addition of CD11b⁻ donor APCs accelerated in vivo donor T cell proliferation in allogeneic BMT. *A* and *B*, Percentages of: donor CD4⁺ T cells (*A*) and CD8⁺ T cells (*B*) according to numbers of cell divisions (CFSE dilution). For B6 \rightarrow B6 syngeneic (Syn) transplants only data from HSCs plus T cells transplant; transplants adding CD11b⁺ or CD11b⁻ APCs yielded similar results (data not shown). *C*, Numbers of total and Ki67⁺ donor CD4⁺ and CD8⁺ donor T cells at day 3 in the spleen of B6 \rightarrow B10 transplant recipients. Data are means ± SD from four replicate experiments with three mice per treatment group. *D*–*F*, T cell chimerism of blood T cells derived from mature T cells in the graft (*D*), donor HSC-derived T cells (*E*), and residual host T cells (*F*). Data are means (±SD) from four replicate experiments, five mice per group. *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001 comparing T cell chimerism in groups receiving donor APCs with HSC and T cell groups.

Li et al.

Page 19



FIGURE 3.

Transplanting CD11b⁻ donor APCs lead to increased IFN- γ production by donor T cells. *A*–*C*, Mean numbers (±SD) of cytokine producing CD4 (*top*) and CD8 (*bottom*) T cells were determined per spleen. *A*, IFN- γ^+ donor T cells; *B*, IL-4⁺ donor T cells; *C*, IL-10⁺ donor T cells. *D*, Th1 (*top*) and Th2 (*bottom*) cytokine concentrations determined in serum of recipients 3 and 10 days posttransplant. Data are from three experiments with a total of nine mice at each time point per group. *p* values are based on comparisons of cytokine levels from mice that received donor APCs with those that received HSCs and T cells. *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001.

Li et al.



FIGURE 4.

CD11b⁺ and CD11b⁻ donor APCs home equivalently to lymphoid organs after transplantation. CD90.1⁺ B10 mice were transplanted with 3×10^3 B6 CD90.1⁺ HSC, 3×10^5 CD90.2⁺ T cells, and 5×10^4 GFP⁺ APC subsets. Spleen sections were prepared from mice sacrificed on days 3 and 10 posttransplant and were examined by confocal microscopy for donor APC (green) and donor T cells (red). Day 10 spleen sections were photographed using $\times 100$ (*A*–*D*) and $\times 40$ (*E*) objectives. *A*, Mice transplanted with HSCs alone. *B*, Mice transplanted with HSCs T cells. *C*, Mice transplanted with HSCs, T cells, and GFP⁺CD11b⁺ APCs. *D*–*E*, Mice transplanted with HSCs, T cells, and GFP⁺CD11b⁺ APCs. *F*, The average number of APCs per microliter of splenic tissue was determined by microscopic analysis of histological sections. The experiment was repeated four times with three mice per group. Three slides per organ were screened, and five high-powered fields per slide were examined.



FIGURE 5.

Costimulatory molecule expression is increased following activation of CD11b⁻ APCs. *A*, Initial gating on donor-derived GFP⁺ cells in recipient BM and spleen, and subsequent identification of GFP⁺CD11c⁺CD11b⁺ APC populations and GFP⁺CD11c⁺CD11b⁻ APC populations. *B–D*, Isotype controls are shown as dotted line histograms. *B*, Differentiation of CD11b⁻ APC precursors and CD11b⁺ APC precursors tracked by expression of lineage markers B220, F4/80, Gr-1, and CD317 on GFP⁺ donor cells 10 days posttransplant. *C*, Expression of costimulatory molecules on APC subsets from B6 GFP⁺ BM 10 days following transplantation in B10 recipients. Recipient BM and splenocytes (SP) were analyzed by flow cytometry with electronic gates on GFP⁺ cells and concatenation of list mode files of at least 1,000,000 events from each of 3 mice per treatment group. Results are from one of five independent experiments with similar results. *D*, Costimulatory molecule expression on BM CD11b⁻ and CD11b⁺ APC subpopulations after stimulation with either CD40L or syngeneic T cells with allogeneic Ag. Data represent analysis of 6×10^5 cultured APCs and are representative of three replicate experiments, each with thee parallel wells per experimental condition. FSC, Forward scatter.

Li et al.





FIGURE 6.

Addition of CD11b⁻ APCs to an allogeneic graft of HSC and T cells increased leukemiafree survival without increasing GvHD. A, Survival of mice that received various doses (0, 3 $\times 10^5$, or 3×10^6) of donor splenic T cells and CD11b⁻ or CD11b⁺ APCs in B6 \rightarrow B10 transplant pair with no leukemia cells (see legend in B). B, Mean GvHD clinical scores in B6 \rightarrow B10 recipients with no leukemia. C, Body weight loss in B6 \rightarrow B10 recipients with no leukemia. D–I, HSCs and 3×10^5 T cell dose groups only. D, Survival of mice in C3H \rightarrow B6 transplants with no leukemia cells. E, Mean GvHD clinical scores in C3H \rightarrow B6 recipients with no leukemia cells. F, Body weight loss in C3H \rightarrow B6 recipients with no leukemia cells. G, Survival of B10 mice that received B6 transplants with 1×10^5 viable LBRM leukemia cells in 5 separate experiments with 10 mice per experimental group. H, The fraction of mice from G with pathological evidence for tumor at necropsy. I, Survival of B6 mice that received C3H.SW transplants with 5×10^4 viable MMB3.19 leukemia cells, in 2 experiments, with 10 mice per experimental group. Arrows, i transplant groups that received CD11b⁻ APC. #, Mice euthanized due to body weight loss were 25%. p values (***, p <0.001) represent log-rank comparison of survival of recipients of CD11b⁻ APCs with recipients of HSCs and T cells.

Li et al.



FIGURE 7.

Transplanting donor CD11b⁻ APC increased long-term alloreactivity, Th1 polarization and antitumor cytotoxicity of donor T cells. Donor T cells were recovered from the spleens of B6→B10 or C3H.SW→B6 recipients on days 81 or 34 posttransplant, respectively, and CFSE-labeled as described in *Materials and Methods. A* and *B*, Mean percentages (±SD) of nondivided T cells and T cells that had undergone five to six cell divisions after 3 days of culture with irradiated host-type splenocytes are shown (three mice from each transplant group). *C* and *D*, Killing of allogeneic leukemia cells by donor T cells. Mean percentages (±SD) of caspase⁺ LBRM targets (*C*) or MMB3.19 targets (*D*) following a 3-h incubation with T cells at an E:T ratio of 10:1 are shown (three to five mice from each transplant group). *, *p* < 0.05; **, *p* < 0.01, comparing T cells from recipients of HSCs and T cells plus CD11b⁻ APCs with T cells from recipients of HSCs and T cells without DC.