

Separation of graft-versus-host disease from graft-versus-leukemia responses by targeting CC-chemokine receptor 7 on donor T cells

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CC-chemokine receptor 7 (CCR7) is expressed on the surface of naive T cells, and plays a critical role in their movement into secondary lymphoid tissue. Here, we show that murine T cells lacking CCR7 (CCR7^{-/-}) generate attenuated graft-versus-host disease (GVHD) responses compared with wild-type (WT) cells, with the difference varying inversely with the degree of major histocompatibility complex (MHC) disparity between the donor and recipient. CCR7^{-/-} T cells exhibited

an impaired ability to traffic to recipient lymph nodes, with an increased capacity to home to the spleen. CCR7^{-/-} T cells, however, demonstrated a reduced ability to undergo in vivo expansion in the spleen due to impaired interactions with splenic antigen-presenting cells. On a cellular level, CCR7^{-/-} T cells were functionally competent, demonstrating a normal in vitro proliferative capacity and a preserved ability to produce inflammatory cytokines. Importantly, CCR7^{-/-} T cells were capable of generating robust graftversus-leukemia (GVL) responses in vivo, as well as complete donor T-cell reconstitution. CCR7^{-/-} regulatory T cells were able to protect against lethal GVHD when administered before WT conventional T cells. Our data suggest that CCR7 inhibition in the early posttransplantation period may represent a feasible new therapeutic approach for acute GVHD attenuation without compromising GVL responses. (*Blood.* 2010;115(23):4914-4922)

Introduction

Graft-versus-host disease (GVHD) is the greatest complication limiting the clinical utility of allogeneic hematopoietic stem cell transplantation (HSCT). Mechanistically, GVHD involves the early trafficking of donor naive T cells (T_n cells) to recipient secondary lymphoid tissue (SLT) where they undergo activation and expansion, and their subsequent migration to peripheral target organs where they elicit injury.1-3 Some uncertainty exists, however, as to the relative contributions of the various recipient SLTs to GVHD pathogenesis. Several studies have shown that donor T_n cells are imprinted with a particular adhesion molecule profile within a specific lymphatic site that serves to direct the cell to a corresponding area of peripheral inflammation. For example, recipient Peyer patches (PPs) and mesenteric lymph nodes (MLNs) may be important for the induction of gastrointestinal GVHD.^{4,5} Other studies, however, have suggested a more significant redundancy among the various recipient lymphoid tissues, without a direct link between a given lymphatic organ and a specific GVHD manifestation. Notably, irradiated B6 lymphotoxin-a receptordeficient mice, which lack PPs and MLNs but possess an intact spleen, appear to develop intestinal acute GVHD that is similar to that of wild-type (WT) B6 recipients when given transplants across a complete major histocompatibility complex (MHC) mismatch.⁶

Regardless, the movement of donor T cells into recipient lymphoid organs is critical for maximal GVHD induction, as animals lacking all SLTs have consistently been shown to generate absent or attenuated inflammatory responses.^{6,7} The trafficking of donor T cells into SLT is in turn dependent on the exact array of

adhesion molecules expressed by the cell. In general, naive and central memory T cells express a receptor profile that allows for their efficient migration into SLT, where they sample antigen on resident antigen-presenting cells (APCs). Upon activation, T cells down-regulate these homeostatic trafficking receptors and express other adhesion molecules that direct them to peripheral sites of inflammation.³

CC-chemokine receptor 7 (CCR7) is a G protein-coupled receptor expressed on naive T cells, B cells, and activated dendritic cells (DCs), and plays an important role in their trafficking into SLT.8 In the case of T cells, the binding of CCR7 to either CCL19 or CCL21 allows for lymphocyte firm arrest on lymph node (LN) high endothelial venules by stabilizing the binding of T-cell LFA-1 to vascular ICAM-1.9,10 In addition, CCR7 functions to direct T cells to appropriate T cell-rich zones within the LN paracortex after egress from the circulation, and is critical for the movement of T cells to their proper anatomical location within the spleen. Not surprisingly, mice knocked out at the CCR7 locus demonstrate numerous lymphoid abnormalities, including greatly reduced T cell numbers within peripheral LNs and PPs and a near-total absence of T cells within the splenic white pulp.8 Given its multiplicity of effects on the organization of SLTs, we hypothesized that the absence of CCR7 on donor T cells might overcome any lymphatic redundancy present in transplant recipients, and allow for a reduction in GVHD severity.

Here, we show that $CCR7^{-/-}$ T cells generate greatly attenuated GVHD responses due to both impaired migration of donor T_n cells

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into recipient LNs and PPs, and abnormal T_n -cell proliferation in the spleen. Importantly, however, CCR7^{-/-} T_n cells retain the ability to mount aggressive graft-versus-leukemia (GVL) effects and to generate complete donor T-cell reconstitution. In addition, CCR7^{-/-} regulatory T cells (T_{reg} cells) remain capable of protecting against lethal GVHD when administered before conventional WT T_n cells. Taken together, these data suggest that CCR7 inhibition may represent a viable treatment option for the prevention of acute GVHD in the clinical setting.

Methods

Mice

C57BL/6, BALB/c. BALB.B, B6.PL-Thy1^a/CyJ, and B6xDBA/2 F1 mice were purchased from The Jackson Laboratory. Enhanced green fluorescent protein (eGFP)–expressing C57BL/6 mice were generated as described.² CCR7^{-/-} mice backcrossed 4 times onto a C57BL/6 background (B6.129P2-CCR7^{tm1Dgen}) were obtained from The Jackson Laboratory and further backcrossed 7 to 8 generations. eGFP-expressing CCR7^{-/-} mice were generated by crossing CCR7^{-/-} C57BL/6 mice with eGFP-expressing C57BL/6 mice. All experiments were performed in accordance with protocols approved by The University of North Carolina Institutional Animal Care and Use Committee.

Transplantation systems

Donor T cell–depleted (TCD) bone marrow (BM) cells were prepared as previously described.¹¹ Whole CD25-depleted splenic effector T cells were prepared using a total T-cell isolation column kit (Cedarlane Laboratories) followed by negative selection of CD25- and B220-expressing cells using antibodies coupled to ferromagnetic beads and magnetic separation of these cells using a magnetic-activated cell sorter (MACS) column (Miltenyi Biotec).¹² T_n cell purity was measured by flow cytometry before transplantation. For whole splenic T_n cell transplants, the donor CD4⁺ to CD8⁺ ratio approximated 1:1 following our standard purification procedure. Donor T_{reg} cells were prepared as previously described using a column purification method.¹³

P815 tumor model

P815 cells (ATCC; TIB-64) are murine mastocytoma cells derived from DBA/2 (H-2^d) mice.¹⁴ P815 cells were cultured in RPMI-1640 media (Invitrogen) supplemented with 10% fetal bovine serum (Invitrogen) and 100 U/mL of penicillin and streptomycin for 7 to 10 days before all transplantations. Initial tumor dose experiments were conducted to determine an optimal P815 inoculum that would consistently result in 100% recipient mortality when irradiated B6D2 mice were given transplants of TCD B6 BM cells without T_n cells (25 000 tumor cells per recipient).

Organ cytokine analysis

Recipient animals were anesthetized and perfused with phosphate-buffered saline (PBS). Whole organs (or in the case of the liver, the hepatic middle lobe immediately adjacent to the gallbladder) were removed. Cytokine levels were measured in the homogenates using enzyme-linked immunosorbent assay (ELISA) kits directed against IFN- γ and TNF- α (Biolegend).

Flow cytometry and immunofluorescence microscopy

The following antibodies were purchased from eBioscience: anti-CD4 (RM 4.5), anti-CD8 (53-6-7), anti-B220 (RA3-6B2), anti-CD25 (PC61), anti-CD62L (Mel-14), anti-CD44 (IM7), anti-K^b (AF6-88.5.5.3), anti–IFN- γ (XMG1.2), anti–IL-13 (eBio13A), anti–IL-10 (JES5-16E3), anti–TNF- α (MP6-XT22), anti-CD107a (eBioABL-93), anti-CD107b (eBio1D4B), anti-Thy1.1 (HIS51), and anti-CD11c (N418). Antibody to mouse K^d (SF1-1.1) was purchased from Biolegend. Cy3 goat anti–Armenian hamster was purchased from Jackson ImmunoResearch Laboratories. Flow cytometric

acquisition was performed on a Cyan flow cytometer using Summit software (Dako).

Intracellular cytokine staining

Cells from recipient spleens and MLNs were stimulated with 50 ng/mL phorbol 12-myristate 13-acetate (PMA), 500 ng/mL ionomycin, and 3 µg/mL brefeldin A (Sigma-Alrich) for 4 hours, with or without anti-CD107a and anti-CD107b antibody. Cells were then washed and stained for K^d, CD4, or CD8, and intracellular cytokine staining was subsequently performed using a Cytofix/Cytoperm kit (BD Biosciences).

Mixed lymphocyte reactions

A total of 2 × 10⁵ highly pure CD4⁺CD25⁻ or CD8⁺CD25⁻ were obtained from donor spleens using fluorescence-activated cell sorting and cocultured with an equal number of irradiated, TCD B6D2 stimulator cells for 96 hours. During the last 16 to 20 hours of incubation, 0.037 MBq (1 μ Ci) of [³H]thymidine (Amersham Pharmacia Biotech) was added per well. Cells were harvested onto filters, and [³H]thymidine incorporation was measured by scintillation counting.

Stereomicroscopy

Animals were anesthetized, and the organs were imaged with a Zeiss Stereo Lumar V12 microscope with eGFP bandpass filter (Carl Zeiss) at room temperature (RT). eGFP intensities were determined with AxioVision (Carl Zeiss) software. For all time points and organs, WT and $CCR7^{-/-}$ T_n cell recipients were assessed using identical camera exposure times (exp) and similar magnifications (mag). Specific values were as follows: Spleen day +15, exp = 824 milliseconds, mag = 15 × to 18 ×; MLN day +15, exp = 263 milliseconds, mag = 13 × to 15 ×; PP day +15, exp = 318 milliseconds, mag = 22 × to 27 ×; lung day +15, exp = 1.3 seconds, mag = 17 ×; liver day +15, exp = 3.1 seconds, mag = 34 × to 40 ×; colon day +15, exp = 2.1 seconds, mag = 12 ×; spleen day +3, exp = 3.5 seconds, mag = 18 × to 32 ×; MLN day +3, exp = 2.5 seconds, mag = 15 ×; and PP day +3, exp = 976 milliseconds, mag = 32 × to 33 ×.

Organ eGFP quantification

Recipient organs were homogenized, and absolute eGFP levels were measured using an ELISA kit directed against GFP/eGFP (Cell Biolabs). To ensure that there were no intrinsic differences in eGFP expression between B6 eGFP⁺ and CCR7^{-/-} eGFP⁺ cells, 1×10^7 splenocytes were isolated from 2 of each mouse strain, the cells were pelleted by centrifugation and homogenized, and absolute eGFP levels within the homogenates were measured. The eGFP levels from equal numbers of cells from both strains were found to be nearly identical (data not shown).

Fluorescence microscopy

Recipient spleens were immersed in 4% formalin, 10% sucrose PBS solution for 2 hours at 4°C before being flash-frozen over liquid nitrogen in optimal cutting temperature solution.¹⁵ Subsequently 8-µm sections were cut on a cryostat at -20° C and allowed to dry overnight at 25°C. On the day of immunostaining, slides were fixed in cold acetone for 10 minutes, then washed for 5 minutes in PBS. Slides were blocked using 10% goat serum, 2% bovine serum albumin, and 5% FC-block solution, and stained with an Armenian hamster anti–mouse CD11c/Cy3-conjugated goat anti–hamster antibody combination. Slides were visualized on an Olympus BX61 fluorescence microscope using Improvision Velocity Software and a Hammatsu ORCA RC camera at RT. Images were obtained at 4× and 10× magnifications with the following exposures: 4×, 1.5 to 1.8 seconds for Cy3, 700 to 760 milliseconds for eGFP, 10×, 700 to 800 milliseconds for Cy3, 500 to 600 milliseconds for eGFP.

Statistical analysis

Survival curves were constructed using the method of Kaplan and Meier. Overall survival was compared using Fisher exact test. Where indicated, median survival times were compared using the log-rank test. To compare



Figure 1. Survival and GVHD scores of animals receiving TCD BM alone or BM plus T cells from WT B6 or CCR7^{-/-} B6 donors. (A-B) B6D2 recipients were lethally irradiated to 9.5 Gy (950 rads) on day -1, and then administered $3\times10^6~TCD$ BM cells from WT B6 donors with or without $4\times 10^6~\text{CD25-depleted}$ whole (CD4+ and CD8+) splenic T_n cells from either WT B6 or CCR7^{-/-} B6 donors on day 0 by tail vein injection. Recipient animals were followed for survival and scored for GVHD twice weekly using a validated clinical scoring system.²⁰ Animals were assigned a score from 0 to 2 for each of 5 GVHD parameters: weight loss, activity, fur-ruffling, kyphosis, and skin lesions. Scores ranged from 0 (minimum) to 10 (maximum). Error bars depict SEM. Data are combined from 3 separate transplantation experiments. n = 12 animals per group. (A) P < .001 for comparison of overall survival proportions at the end of the study observation period between BM/WT Tn and BM/CCR7^{-/-} T_{n} groups by Fisher exact test. (B) *P < .001 for comparison of GVHD scores between BM/WT T_n and BM/CCR7^{-/-} T_n groups by the Mann-Whitney test. (C-D) BALB.B recipients were lethally irradiated to 9.5 Gy (950 rads) on day -1, and then administered 3 \times 10 6 TCD BM cells from WT B6 donors with or without 5×10^6 CD25-depleted whole T_n cells from either WT B6 or CCR7^{-/-} B6 donors on day 0. Data are combined from 2 separate transplantation experiments. n = 6 BM, 10 BM/WT T_n, 8 BM/CCR7^{-/-} T_n. (C) P < .001 for overall survival comparison between BM/WT T_n and BM/CCR7^{-/-} T_n groups. (D) *P < .001for comparison of GVHD scores. (E-F) BALB/c recipients were lethally irradiated to 8 Gy (800 rads) in a single fraction on day -1, and then administered 5×10^{6} TCD BM cells from WT B6 donors with or without 6×10^5 CD25-depleted whole T_n cells from either WT B6 or CCR7 $^{-/-}$ B6 donors on day 0. n = 2 BM, 5 BM/WT T_n, 6 BM/CCR7^{-/-} T_n. (E) P = .182 for overall survival comparison using the Fisher exact test, and P = .002 for median survival comparison using the log-rank test between BM/WT T_n and BM/CCR7^{-/-} T_n groups.

mean scores that fell within a predetermined range (ie, GVHD scores ranging from 0-10, and pathology scores ranging from 0-4), the nonparametric Mann-Whitney test was used. All other continuous variables, including cytokine levels, eGFP levels, and cell ratios, were compared using the 2-tailed Student *t* test in accordance with previously published reports.¹⁶⁻¹⁹ *P* values less than .05 were considered significant. Unless otherwise indicated, error bars represent SEM.

Results

In the absence of CCR7, T cells generate attenuated GVHD responses

Initially, we set out to determine whether the absence of CCR7 on donor T_n cells would affect on their ability to induce lethal GVHD. Using a well-described C57BL/6 (H-2^b, termed B6) into B6xDBA/2 F1 (H-2^{bxd}; termed B6D2) haplotype-matched murine transplantation model, we found that WT and CCR7^{-/-} T_n cells generated nearly identical GVHD responses during the first 7 to 10 days after transplantation. Thereafter, those mice receiving CCR7^{-/-} T_n cells demonstrated an attenuation of their GVHD, with 83% surviving until the completion of the experiment (Figure 1A-B). In contrast, those animals receiving WT T_n cells exhibited relentlessly progressive disease with a 92% mortality rate by day +70 (P < .001).

To confirm and extend these findings, a minor MHCmismatched B6 into BALB.B (H-2^b) system was used, which more closely resembles the degree of histocompatibility matching typically found in the clinical HSCT setting (Figure 1C-D). Once again, we found a striking difference in outcomes, with recipients of CCR7^{-/-} T_n cells demonstrating a profound survival advantage and much lower GVHD scores (P < .01, Figure 1C; P < .001, Figure 1D).

We then examined how the absence of CCR7 would affect the ability of T_n cells to mediate lethal GVHD in a completely mismatched B6 into BALB/c (H-2^d) model (Figure 1E-F). In this setting, we found the influence of CCR7 on GVHD induction to be somewhat less striking. Although median survival was clearly improved in recipients of CCR7^{-/-} T_n cells compared with those getting WT cells (P = .002; Figure 1E), we did not find a statistical difference in overall survival (P = .182). Further, whereas the onset of severe GVHD was substantially delayed in CCR7^{-/-} T_n cell recipients, these mice eventually went on to develop skin inflammation and weight loss. In total, these findings suggest that the absence of CCR7 on donor T cells affects on their ability to induce GVHD, with the extent of this effect varying with the degree of MHC mismatch between the donor and recipient strains.

CCR7^{-/-} T cells retain a significant GVL effect

We next set out to determine whether the absence of CCR7 would affect on the ability of donor T cells to mount antitumor responses Figure 2. Survival and GVHD scores of B6D2 mice receiving TCD BM cells with or without T_n cells from WT B6 or CCR7^{-/-} B6 donors with P815 murine mastocytoma cells. B6D2 recipients were lethally irradiated and then administered 3 \times 10⁶ TCD BM cells from WT B6 donors with or without 4 \times 10⁶ CD25-depleted T_n cells from either WT B6 or CCR7^{-/-} B6 donors on day 0. In addition, all animals simultaneously received 25 000 P815 murine mastocytoma cells in the same inoculum. Data are combined from 2 separate transplantation experiments. n = 8 animals per group. (A) *P* = .001 for overall survival comparison between WT T_n cell/tumor and CCR7^{-/-} T_n cell/tumor groups. (B) **P* < .001 for comparison of GVHD scores.

in vivo. For these experiments, we returned to our haplotypematched B6 into B6D2 HSCT model, but challenged recipient animals at the time of transplantation with 25 000 P815 murine mastocytoma cells (H-2^d), a tumor line with a propensity to metastasize to the spleen, liver, and spinal cord. As seen in Figure 2A, those mice receiving TCD BM plus tumor cells without T_n cells all died by transplantation day +20. At the time of death, these animals demonstrated massive splenomegaly with diffuse tumor infiltration of the liver. In contrast, animals receiving WT T_n cells at the time of P815 challenge all died of GVHD, with no tumor identified at the time of death and all animals exhibiting high GVHD scores. Importantly, 75% of the mice administered P815 cells and CCR7^{-/-} T_n cells survived long term with no sign of malignancy and only minimal evidence of GVHD clinically (P = .001).

Recipients of CCR7^{-/-} T_n cells demonstrate less hepatic/gastrointestinal inflammation and less colonic inflammatory cytokine production

To evaluate the mechanism by which the absence of CCR7 on donor T_n cells attenuated GVHD severity, lethally irradiated B6D2 mice received transplants of TCD B6 BM with or without WT or CCR7^{-/-} B6 T_n cells, and their organs were harvested between transplantation days 18 and 21 for histopathology (Figure 3A). A separate cohort of recipients underwent transplantation similarly, but their organs were homogenized for cytokine analysis on day +15 (Figure 3B-C). B6D2 recipients that underwent transplantation with CCR7^{-/-} T cells had a significant reduction in pathology scores within the liver (P = .017) and to an even greater extent in the colon (P < .001) compared with animals given WT cells. In addition, less interferon- γ and TNF- α were observed in the colons of CCR7^{-/-} T_n recipients (P = .017 and P = .052, respectively).

CCR7 $^{-/-}$ T cells possess an intact ability to undergo activation and cytokine production

Previous reports have suggested that CCR7^{-/-} T cells possess an intrinsic inability to undergo normal cellular activation and cell cycling.²¹ To determine whether such functional abnormalities would provide an explanation for our findings, we examined the surface expression of various cellular activation markers on both CD4⁺ and CD8⁺ donor T cells within B6D2 spleens and MLNs 7 and 14 days after transplantation, and evaluated the cytokine expression profile of these cells after ex vivo stimulation with PMA and ionomycin (representative data of donor cells extracted from recipient spleens on transplantation day +14 are depicted in supplemental Figure 1A-D, available on the *Blood* Web site; see the Supplemental Materials link at the top of the online article). We also performed in vitro mixed lymphocyte reactions (MLRs) with















Figure 4. Accumulation of WT versus CCR7-/-Tn cells within B6D2 recipients. B6D2 recipients were lethally irradiated and then administered 3×10^6 TCD BM cells from WT B6 donors with or without 4×10^6 CD25-depleted eGFP+ Tn cells from either WT B6 or CCR7^{-/-} B6 donors on day 0. (A-F) Recipient mice were anesthetized on transplantation day +15, and donor T-cell trafficking was studied using stereofluorescence microscopy. Tissues from 1 of 3 representative WT Tn cell recipients are depicted in the top panels, and tissues from 1 of 3 representative CCR7^{-/-} T_n cell recipients are depicted in the bottom panels. Images on the left of each panel depict actual eGFP fluorescence, while those on the right indicate the intensity of the eGFP signal (white > red > yellow > green > blue > black). (G-I) Recipient mice were anesthetized on transplantation day +3, and donor T-cell trafficking to SLTs was evaluated using stereofluorescence microscopy. Images are taken from 1 of 3 representative WT Tn cell recipients and from 1 of 3 CCR7 $^{-/-}$ T_n recipients. (J) Recipient animals were euthanized on transplantation day +15, and their organs were homogenized in a PBS/protease-inhibitor solution. After appropriate dilutions, eGFP levels were measured in each site using an anti-GFP ELISA kit. n = 4 per group. *P = .044; **P = .011; ***P = .002. Organs were pooled from a B6D2 mouse that received a transplant of TCD BM cells plus non-eGFP-expressing B6 T_{n} cells for control purposes. (K) Recipient animals were killed on transplantation day 3, and their spleens were homogenized for analysis by anti-GFP ELISA. n = 3 per group. *P = .002.

freshly purified WT and CCR7^{-/-} CD4⁺ CD25⁻ and CD8⁺ CD25⁻ lymphocytes, plus irradiated, TCD B6D2 splenocytes as stimulator cells (supplemental Figure 1E). WT and CCR7^{-/-} T cells demonstrated a similar ability to undergo activation and proliferation in vitro and a similar phenotype characterized by high levels of surface CD44 and low expression of L-selectin after transplantation. Both CD4⁺ cell populations showed a T helper cell 1 (Th1)-polarized intracellular cytokine expression profile characterized by abundant IFN- γ and TNF- α and little IL-13 or IL-10, and CD8⁺ T cells demonstrated nearly identical levels of surface CD107, a surrogate for cytotoxic T lymphocyte (CTL) lytic activity²² (supplemental Figure 1A-D; data not shown). Further, $CCR7^{-/-}$ T_n cells were found to produce complete donor reconstitution of the CD4⁺ and CD8⁺ compartments by transplantation day +40, although the rate of reconstitution was delayed compared with WT cells (supplemental Figure 1F). Taken together, these findings suggested that in the absence of CCR7, T cells are functional, with an immunophenotype and in vitro cytokine production profile that are similar to those of WT cells.

CCR7^{-/-} T cells demonstrate an impaired ability to accumulate within recipient tissues after HSCT

We next focused on any differences in T-cell in vivo trafficking that might exist in the absence of CCR7. For these studies, we made use of B6 and CCR7^{-/-} mice that were transgenic for eGFP and imaged specific organs by fluorescence stereomicroscopy as previously described.² Upon examining B6D2 recipient mice on transplantation day +15, a time point just before those when our histopathology studies were performed (Figure 3A), we observed a higher fluorescence intensity in the spleens, MLNs, PPs, livers, lungs, and especially colons of those animals receiving WT cells versus those administered CCR7 $^{-\prime-}$ T_n cells (Figure 4A-F). In addition, we quantified the levels of eGFP expressed within several of these organs using an anti-eGFP ELISA (Figure 4J). In accordance with our microscopy findings, we detected significantly less eGFP expression within the spleens, lungs, and colons of those mice receiving CCR7^{-/-} cells, with a strong trend toward less eGFP observed in the liver (P = .044, P = .011, P = .002, and P = .06, respectively).

Given that higher numbers of WT cells were present in recipient MLNs, PPs, and spleens 2 weeks after transplantation, we wondered if a similar difference in accumulation in SLT could be detected at a time point closer to transplantation when donor T cells would have had less time to undergo in vivo expansion. When we examined B6D2 recipients on day +3, we again observed higher numbers of WT cells within the MLN and PPs (Figure 4H-I). Using anti-eGFP ELISA, we also detected more eGFP in the spleens of those mice receiving WT cells (P = .002; Figure 4K). Thus, we

confirmed the important role of CCR7 in the accumulation of T_n cells within LNs, and surprisingly found that CCR7 was important for the early collection of donor T_n cells in the spleen.

CCR7^{-/-} T_n cells show increased trafficking to the spleen and reduced homing to LNs, with less in vivo expansion in both sites

Although the greater number of WT cells within recipient MLNs and spleens as early as transplantation day +3 suggested a homing advantage to both of these sites, we could not rule out the possibility that differential in vivo expansion was also contributing to the difference. To definitively examine the issue of T-cell homing, we made use of a competitive lymphocyte trafficking approach.¹⁶ With this method, Thy1.2⁺ eGFP⁺ WT or eGFP⁺ CCR7^{-/-} T_n cells are coinjected with an equal number of Thy1.1⁺ T_n cells, and the ratio of eGFP⁺ to Thy1.1⁺ cells within recipient tissues are determined by flow cytometry within 24 hours of adoptive transfer, before any substantial T-cell expansion would be expected. As a result, absolute numbers of $CCR7^{-/-}$ and WT eGFP+ cells need not be directly compared, thereby avoiding confounding issues such as variable recipient animal size, circulatory system anatomy, and differences in the number and size of retrieved LNs.

As anticipated, eGFP⁺ WT B6 and Thy1.1⁺ T_n cells were found to home equally well to both lymphoid sites, with the ratio of these 2 populations approaching 1:1 in the spleen and MLNs. In contrast, the eGFP⁺ CCR7^{-/-}/Thy1.1⁺ ratio was found to be reduced in the MLN, but augmented in the spleen, demonstrating an impaired ability for CCR7^{-/-} cells to traffic into LNs, with an increase in their splenic homing (Figure 5A). This finding would be in keeping with previous reports showing higher overall numbers of T cells within the spleens of CCR7^{-/-} mice, with reduced numbers in the peripheral LNs.⁸ It also strongly suggested that our finding decreased numbers of CCR7^{-/-} T cells in the spleen at later time points was not due to impaired trafficking.

To elucidate a mechanism for the lower numbers of CCR7^{-/-} T cells in the spleen on days +3 and +15, we evaluated the proliferation of CCR7^{-/-} and WT T cells within SLTs early after transplantation. For this work, we labeled both B6 WT and CCR7^{-/-} T_n cells with carboxyfluorescein succinimidyl ester (CFSE) immediately before transplantation, and assessed the extent of CFSE dilution within B6D2 recipient spleens and MLNs on day +3. As shown (Figure 5B-D), WT cells had lower levels of residual CFSE within both sites compared with CCR7^{-/-} T cells, demonstrating that donor T-cell proliferation is impaired in the spleen and MLNs in the absence of CCR7.

CCR7^{-/-} donor T cells interact less efficiently with splenic DCs than WT cells

Next, we set out to identify a potential mechanism for the impaired donor T-cell proliferation observed in the absence of CCR7 within the recipient spleen, despite a normal capacity for cellular division in vitro (supplemental Figure 1E). Based on previous reports showing an important role for CCR7 in the normal organization of SLTs,⁸ we hypothesized that CCR7^{-/-} donor T cells might home to recipient spleens after HSCT, but localize poorly to DC-rich zones within the white pulp. To address this possibility, B6D2 recipients received transplants of T_n cells from eGFP⁺ WT or eGFP⁺ CCR7^{-/-} donors, and their spleens were extracted for immunofluorescence microscopy on transplantation days +4 and +7. DCs were stained with an anti-CD11c antibody, and donor T cells were

identified by their eGFP signal. As seen in Figure 6A and 6C, by transplantation day +4, WT donor T cells had efficiently localized into the splenic white pulp, forming discrete, circular T-cell aggregates overlapping with resident DCs. In contrast, $CCR7^{-/-}$ donor T cells were found to colocalize much less frequently with splenic DCs, demonstrating a more diffuse infiltration pattern (Figure 6B,D). These early differences became somewhat less striking 7 days after transplantation (supplemental Figure 2), demonstrating that although $CCR7^{-/-}$ T-cell activation is both diminished and delayed compared with WT cells, it is not entirely impaired.

T_{reg} cells can protect against lethal GVHD in irradiated HSC transplant recipients in the absence of CCR7

Up to this point, our experiments had focused on conventional T cells and the impact of CCR7 on the ability of donor T_n cells to induce GVHD and GVL responses. While these data suggested that blocking CCR7 function in the immediate pre- and posttransplantation periods might prove to be a feasible approach for improving transplantation outcomes, the effect that such an intervention would have on the ability of Foxp3-expressing T_{reg} cells to suppress GVHD responses remained uncertain. To address the contribution of CCR7 to the in vivo function of T_{reg} cells in the HSCT setting, we administered WT B6 T_n cells with WT B6 or CCR7^{-/-} T_{reg} cells to lethally irradiated B6D2 mice. As expected, WT B6 T_{reg} cells afforded 100% survival of B6D2 recipients (Figure 7A). Importantly, however, recipients given CCR7^{-/-} donor T_{reg} cells also demonstrated 100% survival, indicating that naive Treg cells lacking CCR7 remain capable of substantial in vivo immune suppression in lethally irradiated hosts.

Discussion

For the past 20 years, investigators have sought methods to separate the deleterious effects of GVHD from beneficial GVL responses. Here, we demonstrate that the absence of CCR7 on T_n cells leads to impaired T-cell migration into recipient MLNs and PPs and infrequent T-cell/DC interactions within the spleen, which result in impaired donor T-cell expansion. These effects appear to be sufficient for a general attenuation of GVHD severity while still allowing for the elimination of tumor cells.

Several of our results were somewhat unexpected. Previous work had suggested that donor T-cell homing to recipient LNs and/or PPs is not required for lethal GVHD in the presence of an intact spleen, with the latter lymphoid site being sufficient for aggressive inflammatory responses in most if not all peripheral GVHD target organs.⁶ Because CCR7 had previously been shown to be critical for T_n cell homing to LNs and PPs but not for overall splenic trafficking, we hypothesized that CCR7^{-/-} T_n cells might generate GVHD effects that were identical to those of WT cells. Our GVHD survival data showing improved outcomes with $CCR7^{-/-}$ T_n cells were initially surprising, particularly after we confirmed that their splenic homing remained intact after transplantation. Our subsequent studies, however, appear to have reconciled this paradox by demonstrating that early T-cell proliferation within SLTs is also critical for maximal GVHD, and that approaches to diminish this expansion can substantially alter the incidence and severity of acute GVHD independent of effects on T-cell migration.

Also unexpected was the finding that T cells lacking CCR7 were still capable of substantial antitumor activity in vivo. This was



A Lymphocyte Extraction at 24 Hours After Transplant

Figure 5. Trafficking of donor T cells to recipient SLTs 24 hours after transplantation. and their subsequent in vivo expansion. (A) B6D2 recipients were lethally irradiated and then administered 4×10^6 non-eGFP-expressing B6.PL-Thy1a/CyJ Thy1.1+ T_n plus either 4×10^6 eGFP⁺ WT B6 T_n or 4×10^6 eGFP⁺ CCR7^{-/-} B6 T_n cells. Recipient animals were killed after 24 hours, and lymphocytes were extracted from the spleen and MLNs. These cells were then stained for flow cytometry using PerCp-Cy5.5-conjugated anti-Thy1.1 and APC-conjugated anti-CD3 antibodies. The ratios of CD3+eGFP+/ CD3+Thv1.1+ events were then determined for all of the animals in each treatment group. For spleens, n = 6 per group. For MLNs, n = 3 LN pairs per group. *P < .001. **P = .005. Error bars indicate SEM. (B-C) B6D2 recipients were lethally irradiated and then administered 3×10^6 unlabeled TCD BM cells from WT B6 donors plus 3×10^6 CFSE-labeled Tn cells from either WT or CCR7-/- B6 donors. Recipient animals were killed on transplantation day +3, and lymphocytes were extracted from the spleen for flow cytometry. (B) Histograms of cells isolated from the spleen from 1 of 3 representative WT T_n cell recipients and from 1 of 3 representative CCR7^{-/-} T_n cell recipients are depicted, using a CD3⁺ K^d-negative (donor cell) live lymphocyte gate. (C) The mean percentage of proliferating donor T cells within the spleens of WT (left) and CCR7-/- Tn cell (right) recipients are also shown. Error bars indicate SEM. *P = .014. (D) Recipient animals were killed on transplantation day +3, and the MLNs from each of 3 animals per treatment group were pooled for flow cytometry. MLN plots are depicted using a CD3⁺ K^d-negative (donor cell) live lymphocyte gate.

most likely because (1) CCR7^{-/-} T_n cells remained capable of migrating to the liver and spleen, important sites of P815 tumor growth; and that (2) higher T-cell numbers are required for lethal GVHD compared with GVL effects.²⁴ Although CCR7^{-/-} T-cell numbers were reduced relative to WT cells within all SLTs and GVHD target organs 2 weeks after transplantation, CCR7 T_n cells still demonstrated a residual ability to accumulate in those sites and to generate inflammation (Figures 3-4). It is likely that these immune responses were sufficient to eliminate small numbers of tumor cells, but inadequate to produce enough end-organ damage to consistently result in GVHD mortality.

Interestingly, separate anatomic sites appeared to be differentially affected by the absence of CCR7 on T_n cells. Significant differences in pathology scores were noted within the colons and livers of those animals given $CCR7^{-/-}$ T_n cells versus WT cells, whereas similar degrees of histologic injury were observed in recipient spleens and lungs (Figure 3A). Somewhat to our surprise, these variable pathologic differences were observed even though CCR7^{-/-} T_n cell accumulation was generally less than that of WT cells within every SLT and GVHD target organ examined (Figure 4). Although the mechanism underlying this disparity is not entirely clear, several different hypotheses may explain this finding. First, GVHD pathology may be dependent on a threshold number of donor T cells within a specific organ. In such a scenario, any effector T cells accumulating beyond that threshold would be expected to have little effect on the ultimate pathologic score observed in that location, even though the kinetics of the GVHD inflammatory response might be accelerated. Alternatively, it is possible that GVHD pathology correlates better with inflammatory cytokine production than donor T-cell accumulation in certain organs. In our experiments, the greatest differences in both donor T-cell accumulation and IFN- γ /TNF- α production were observed in the colon. We believe that the improved survival outcomes seen in those animals receiving CCR7^{-/-} T_n cells were primarily due to the reduced histopathology observed in that site, since gastrointestinal injury is an important cause of recipient mortality in acute GVHD.

Finally, we were surprised by the finding that CCR7^{-/-} T_{reg} cells were able to prevent lethal GVHD in murine HSC transplant recipients. Previous reports had demonstrated that although CCR7 is not required for the thymic development of Treg cells or for their ability to suppress T-cell expansion in vitro, CCR7^{-/-} T_{reg} cells are impaired in their capacity to suppress immune responses in nonirradiated animals secondary to an inability to efficiently enter LNs and to traffic to appropriate T-cell zones within SLTs.²⁵ Our data would suggest that in the lymphopenic and inflammatory environment that follows ablative conditioning, Treg cells may be capable of trafficking into lymphoid sites using adhesion/chemokine receptors other than CCR7. Alternatively, donor T_{reg} cells might directly suppress developing alloimmune responses within peripheral GVHD target organs²⁶ and rely less on early trafficking into SLTs. Future experiments will focus in greater depth on the influence of CCR7 on the regulatory T-cell arm in the transplantation setting, with an emphasis on where these cells function to suppress GVHD.

While we were pursuing these studies, Anderson et al published work specifically designed to evaluate whether trafficking receptor differences between T_n cells and memory T cells were responsible for the disparate GVHD outcomes produced by the 2 cell types.⁷ For a portion of these experiments, the investigators used CCR7^{-/-}

Figure 6. Donor T cell/DC interactions within recipient spleens on transplantation day +4. B6D2 recipients were lethally irradiated and administered 3×10^{6} TCD BM cells from WT B6 donors with 4 \times 10 6 CD25depleted eGFP⁺ T_n cells from either WT B6 or CCR7^{-/-} B6 donors on day 0. Recipients were killed on transplantation day +4, and their spleens removed for immunofluorescence microscopy. Images are taken from 1 of 3 representative WT T_n cell recipients and from 1 of 3 $CCR7^{-/-}$ T_n cell recipients. Donor cells are identified by their eGFP positivity (green). An anti-CD11c primary/Cy3-conjugated secondary antibody combination was used to visualize CD11c-expressing cells, which are predominantly DCs (red). (A,C) Recipients of WT Tn cells. (B,D) Recipients of CCR7^{-/-} cells. Original magnifications, $\times 4$ (A-B) and ×10 (C-D).



BALB/c mice as donors and B6 mice knocked-out at the glucosyltransferase 2-3 loci (which lack peripheral LN addressin, the ligand for L-selectin) as recipients (GST $2-3^{-/-}$). There, they found CD4⁺ CCR7^{-/-} T_n cells to be capable of mediating lethal GVHD in GST $2-3^{-/-}$ animals. Although our own experiments with haplotype and minor MHC-mismatched transplantation systems clearly yielded different outcomes, our findings using a completely MHCmismatched model (B6 into BALB/c) would be consistent with their data, as CCR7^{-/-} T_n cells were able to elicit substantial GVHD in that setting (Figure 1E-F). The unfavorable outcomes observed in these 2 model systems were likely the result of the extreme histocompatibility differences between the donor and recipient strains, which appear to have allowed for enough T-cell expansion to overcome the beneficial effects of the absence of CCR7. Our results are somewhat divergent, however, in that we found CCR7^{-/-} T_n cells to generate GVHD responses that differed qualitatively from those of WT cells, even in the setting of a complete MHC mismatch. Namely, the onset of severe GVHD was delayed in those animals receiving CCR7^{-/-} cells, and their median survival times were significantly prolonged. The reason for these disparities is not entirely clear, and is most likely related to differences between the recipient mice in the 2 models used.

In summary, we have demonstrated superior HSCT outcomes with CCR7^{-/-} donor T cells in clinically relevant, minor and haplotype MHC-mismatched murine model systems. Importantly, the absence of CCR7 appears to have little impact on the ability of T_n cells to mount robust GVL responses, or on the in vivo suppressive ability of donor T_{reg} cells in irradiated hosts. While long-term inhibition of CCR7 would most likely have deleterious consequences due to the critical role of CCR7 in the homeostatic



Figure 7. Survival and GVHD scores of B6D2 mice receiving WT or CCR7^{-/-} T_{reg} cells before WT T_n cell dosing. (A-B) B6D2 recipients were lethally irradiated on day -1 and then administered 3 \times 10⁶ TCD BM cells from WT B6 donors with or without 1 \times 10⁶ column-purified CD4⁺CD25⁺ T_{reg} cells from WT B6 or CCR7^{-/-} B6 mice on day 0. A total of 4 \times 10⁶ CD25-depleted T_n cells from WT B6 donors were then administered on day $+2.2^3$ n = 4 BM, 7 BM/ T_n , 9 BM/ T_n /WT T_{reg} , 9BM/ T_n /CCR7^{-/-} T_{reg} . (B) *P = .094 for comparison of GVHD scores between WT T_{reg} and CCR7^{-/-} T_{reg} groups on day +70. Error bars indicate SEM.

migration of T- and B-lymphocytes, short-term targeting of this chemokine receptor for approximately 4 to 6 weeks after stem cell transplantation could significantly affect the occurrence of acute GVHD with minimal long-term effects.

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Authorship

Contribution: J.M.C. performed experiments and wrote the manuscript; M.J.C. performed experiments; A.P.-M. analyzed the histopathology specimens; M.L.W. performed experiments and derived the CCR7^{-/-}eGFP⁺ mouse line; J.E.B. performed experiments; B.R.B. reviewed and edited the manuscript; and J.S.S. conceived and supervised the project and edited the manuscript.

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