Selective Treg reconstitution during lymphopenia normalizes DC costimulation and prevents graft-versus-host disease

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Regulatory T cells (Tregs) have been shown to enhance immune reconstitution and prevent graft-versus-host disease (GVHD) after hematopoietic stem cell transplantation; however, it is unclear how Tregs mediate these effects. Here, we developed a model to examine the mechanism of Treg-dependent regulation of immune reconstitution. Lymphopenic mice were selectively reconstituted with Tregs prior to transfer of conventional CD4+ T cells. Full Treg reconstitution prevented the rapid oligoclonal proliferation that gives rise to pathogenic CD4 effector T cells, while preserving the slow homeostatic form of lymphopenia-induced peripheral expansion that repopulates a diverse peripheral T cell pool. Treg-mediated CTLA-4–dependent downregulation of CD80/CD86 on DCs was critical for inhibition of rapid proliferation and was a function of the Treg/DC ratio achieved by reconstitution. In an allogeneic BM transplant model, selective Treg reconstitution before T cell transfer also normalized DC costimulation and provided complete protection against GVHD. In contrast, cotransfer of Tregs was not protective. Our results indicate that achieving optimal recovery from lymphopenia should aim to improve early Treg reconstitution in order to increase the relative number of Tregs to DCs and thereby inhibit spontaneous oligoclonal T cell proliferation.

Introduction

Achieving satisfactory immune reconstitution in lymphopenic subjects remains a major problem in many clinical settings, including following autologous or allogeneic hematopoietic stem–cell transplantation (HSCT) and recovery from cancer chemotherapy. Studies in both mouse and man have indicated that the lymphopenic state is associated with a spectrum of T cell abnormalities, including spontaneous proliferation, conversion to activated/memory phenotype, and tissue infiltration and damage (1–4). In addition, spontaneous proliferation is usually oligoclonal, leading to constriction of the T cell repertoire (5–8). Development of protocols that allow full reconstitution of the peripheral T cell compartment without inflammatory sequelae is an important aspect of ensuring good outcomes after lymphopenia-inducing therapeutic regimens.

Regulatory T cell infusion (Treg infusion) has been shown to promote immune reconstitution and reduce the incidence of graft-versus-host disease (GVHD) after allogeneic HSCT (9–17). However, the mechanistic basis of this effect remains unclear. In mouse models in which the kinetics of CD4+ T cell lymphopenia reconstitution have been studied in detail, 2 phases of lymphopenia-induced proliferation (LIP) have been identified (18, 19). The first is rapid, requires T cell receptor interactions (TCR interactions) with MHC-peptide ligands, and generates a differentiated effector cell population. Only 4%–6% of CD4+ T cells are subject to fast-phase LIP in syngeneic hosts (20). These spontaneously proliferating cells are believed to exhibit low affinity cross-reactivities with endogenous antigens, including self-antigens and gut microflora (21, 22), and their TCRs lie closest to the thymic cut-off for negative selection of self-reactive specificities. Although they represent a minor subpopulation, they rapidly generate a large oligoclonal population that dominates the reconstituted immune system and may induce tissue inflammation and autoimmunity. In contrast, slow-phase LIP is TCR independent, generates cells with a naive phenotype, and can be regarded as truly homeostatic, reconstituting a highly diverse polyclonal immune compartment. Previous studies have indicated that Tregs can partially suppress fast-phase proliferation (23). Importantly, a detailed quantitative study of the suppressive effect of selective Treg reconstitution on fast-phase proliferation has not previously been reported.

We have developed a mouse model to study the mechanism by which Tregs suppress LIP, based on reconstitution of syngeneic immunodeficient mice with pure populations of Tregs. Since Tregs are strictly IL-2 dependent but do not themselves make...
Figure 1. Effect on rapid-phase LIP of reconstitution of the Treg compartment prior to T cell transfer. Lymphopenic Rag–/– mice were reconstituted with purified Tregs by adoptively transferring either 0.25 x 10⁶, 0.5 x 10⁶, 1.25 x 10⁶, or 2.5 x 10⁶ bead-selected CD4⁺CD25⁺ Tregs. Mice were treated with IL-2/JES6-1 complexes second daily to support Treg expansion. (A) Donor cells were distinguished from host cells by a CD45 allelic difference. Shown is expression of CD25 and Foxp3 on donor CD4⁺ T cells in pLN (n = 4 per group). (B) The total number of CD4⁺CD25⁺Foxp3⁺ Tregs was calculated for each Treg dose in pLN, mLN, and spleen of Treg-reconstituted mice, and for WT controls (n = 4 per group). Data in A and B are from a single experiment. The comparison of WT, Rag–/–, and Rag–/– mice reconstituted with 2.5 x 10⁶ CD4⁺CD25⁺ Tregs is representative of more than 10 independent experiments. (C) At day 7 after Treg transfer, 1 x 10⁶ CFSE-labeled CD4⁺CD25– conventional T cells were adoptively transferred into unreconstituted or Treg-reconstituted Rag–/– mice, as described for A. All mice were treated with second daily IL-2/JES6-1 complexes throughout the experiment. Representative CFSE division profiles of conventional CD4⁺ T cells are shown at 7 days after transfer (left panels). (D) The ratio of CFSE⁻ cells (>7 rounds of division) to CFSE⁺ T cells (0–3 rounds of division) was calculated. Data in C and D are from one experiment with n = 3–4 per group. (E) Comparison of the ratio of divided to undivided cells in the pLN, mLN, and spleen of unreconstituted or Treg-reconstituted Rag–/– mice. Data are pooled from 6 independent experiments (n = 9–19 per group). Data in B and D were analyzed using one-way ANOVA with a Newman-Keuls post-test, while data in E were analyzed using a 2-tailed unpaired t-test. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, and ****P < 0.0001.
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Results

Treg reconstitution inhibits spontaneous T cell proliferation in lymphopenic animals

To study the mechanism of Treg-dependent suppression of LIP, we developed a model in which pure populations of Tregs were adoptively transferred into immunodeficient Rag–/– mice and provided with an exogenous source of IL-2 in the form of IL-2/anti–IL-2 mAb complexes (24). After 7 days, reconstituting Tregs were identified in lymph nodes (LN) and spleen by flow cytometry (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI76031DS1). Results are from a single experiment and are representative of more than 20 independent experiments.

CD80, CD86, and CD86 expression by DCs from Treg-deficient mice

Figure 2. Expression of CD80 and CD86 by DCs from Treg-deficient mice. (A) MFI of CD80 (left panel) and CD86 expression (right panel) by DCs from immunodeficient Rag–/– mice and their immunosufficient Rag+/– littermates (n = 6 per group). Migratory DCs and blood-borne DCs in pLN/mLN were gated as MHCII+CD11c+ and MHCII+CD11c−, respectively. Blood-borne DCs in the spleen were further subdivided based on expression of CD8 and CD11b. Results are from a single experiment and are representative of more than 10 independent experiments. (B) Comparison of DC phenotype in mice housed under SPF versus GF conditions. MFI of CD80 expression by CD11c+ B220+ pLN and spleen DCs from immunosufficient BALB/c and immunodeficient C.B-17/lcr-scid mice housed under either SPF or GF conditions. Data are pooled from 2 independent experiments with 2 mice per group in each experiment. (C) Expression of CD80 by migratory DCs in the pLN of Rag–/– mice that received either no cells or were reconstituted with 0.25 × 10⁶, 0.5 × 10⁶, 1.25 × 10⁶, or 2.5 × 10⁶ CD4+CD25+ Tregs (open histograms) compared with expression in untreated WT mice (shaded histograms). (D) Expression of CD80 (left panels) and CD86 (right panels) by migratory DCs in the pLN, mLN, and splenic DCs at day 7 after Treg reconstitution compared with expression in untreated WT and Rag–/– mice. Data in A, C, and D are from a single experiment with n = 4 per group. The comparison of WT, Rag–/– mice, and Rag–/– mice reconstituted with 2.5 × 10⁶ CD4+CD25+ Tregs is representative of more than 10 independent experiments. Data in A were analyzed using a 2-tailed t test, while data in B and D were analyzed using one-way ANOVA with a Newman-Keuls post-test. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.

IL-2, we used IL-2 complexes (24) to support reconstitution with- out the potentially confounding effects of cotransferred conventional T cells as an endogenous source of IL-2. Here, we show that Tregs prevent fast-phase LIP by downregulating the expression of costimulatory molecules by DCs, thereby allowing slow-phase LIP to proceed. In contrast, reconstitution with conventional CD4+ T cells further upregulates costimulation, enhances fast-phase LIP, and inhibits slow-phase LIP. The effect of Tregs is strictly dependent on the numerical ratio of Tregs to DCs in individual secondary lymphoid organs and requires expression of CTLA-4 by Tregs. Using a mouse model of allogeneic BM transplantation (BMT), we show that DC costimulation is also elevated following irradiation and can be reduced by means of reconstitution with either syngeneic or allogeneic Tregs. In addition, Treg reconstitution after BMT completely protected against development of GVHD, whereas cotransfer of Tregs and conventional T cells did not. Our results explain why clinical protocols favoring reconstitution of Tregs before conventional T cells may lead to superior long-term outcomes and suggest that ensuring adequate early Treg recon-stitution is a crucial aspect of the management of lymphopenic patients. In addition, these findings may also explain the mecha-nistic basis behind the association of lymphopenia and autoimmu-nity, which has been noted in both mouse and man.

Results

Treg reconstitution inhibits spontaneous T cell proliferation in lymphopenic animals. To study the mechanism of Treg-dependent suppression of LIP, we developed a model in which pure populations of Tregs were adoptively transferred into immunodeficient Rag–/– mice and provided with an exogenous source of IL-2 in the form of IL-2/anti–IL-2 mAb complexes (24). After 7 days, reconstituting Tregs were identified in lymph nodes (LN) and spleen by flow cytometry (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI76031DS1). CFSE-labeling indicated that the vast majority of Tregs had proliferated after transfer (Supplemental Figure 2) and over 90% retained expression of FoxP3 in the peripheral LN (pLN) (Figure 1A), mesen-teric LN (mLN), and spleen (Supplemental Figure 2).

Figure 2. Expression of CD80 and CD86 by DCs from Treg-deficient mice. (A) MFI of CD80 (left panel) and CD86 expression (right panel) by DCs from immunodeficient Rag–/– mice and their immunosufficient Rag+/– littermates (n = 6 per group). Migratory DCs and blood-borne DCs in pLN/mLN were gated as MHCII+CD11c+ and MHCII+CD11c−, respectively. Blood-borne DCs in the spleen were further subdivided based on expression of CD8 and CD11b. Results are from a single experiment and are representative of more than 10 independent experiments. (B) Comparison of DC phenotype in mice housed under SPF versus GF conditions. MFI of CD80 expression by CD11c+ B220+ pLN and spleen DCs from immunosufficient BALB/c and immunodeficient C.B-17/lcr-scid mice housed under either SPF or GF conditions. Data are pooled from 2 independent experiments with 2 mice per group in each experiment. (C) Expression of CD80 by migratory DCs in the pLN of Rag–/– mice that received either no cells or were reconstituted with 0.25 × 10⁶, 0.5 × 10⁶, 1.25 × 10⁶, or 2.5 × 10⁶ CD4+CD25+ Tregs (open histograms) compared with expression in untreated WT mice (shaded histograms). (D) Expression of CD80 (left panels) and CD86 (right panels) by migratory DCs in the pLN, mLN, and splenic DCs at day 7 after Treg reconstitution compared with expression in untreated WT and Rag–/– mice. Data in A, C, and D are from a single experiment with n = 4 per group. The comparison of WT, Rag–/– mice, and Rag–/– mice reconstituted with 2.5 × 10⁶ CD4+CD25+ Tregs is representative of more than 10 independent experiments. Data in A were analyzed using a 2-tailed t test, while data in B and D were analyzed using one-way ANOVA with a Newman-Keuls post-test. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.
of reconstitution varied according to transferred cell dose (0.25, 0.5, 1.25, or 2.5 × 10^6 Tregs) and between different secondary lymphoid organs, being most efficient in spleen and least efficient in mLN (Figure 1B). To determine the effect of Treg reconstitution on fast-phase LIP, 1 × 10^6 CFSE-labeled CD4^+CD25^− conventional T cells were adoptively transferred into reconstituted hosts on day 7, and the extent of proliferation was assessed after another 7 days. Proliferation was significantly inhibited in reconstituted hosts, with a trend toward increased inhibition at higher doses of Tregs (Figure 1, C and D). As with Treg reconstitution, the degree of inhibition also varied between the different lymphoid organs.

Tregs (Figure 1, C and D). As with Treg reconstitution, the degree of inhibition also varied between the different lymphoid organs. A comparison of fast-phase LIP in Rag^-/- controls and Rag^-/- mice reconstituted with the highest dose of Tregs showed that inhibition was highest in the pLN (30-fold) and spleen (25-fold), compared with the mLN (9-fold) (Figure 1E), reflecting the differences in Treg-reconstitution efficiency in these organs (Figure 1B). In both untreated Rag^-/- controls and Treg-reconstituted mice, the relative number of divided cells was higher in spleen than in LN. This is likely due to preferential recirculation of divided effector T cells from LN to spleen, rather than reflecting the local rate of proliferation in the spleen itself.

Treg reconstitution reduces expression of CD80 and CD86 in lymphopenic animals. We next tested whether Treg reconstitution inhibited fast-phase LIP by downregulating expression of CD80 and CD86 by DCs, since CD28-CD80/CD86 signaling is known to be an absolute requirement for fast-phase LIP (19, 25). Tonic downregulation of DC costimulation by Tregs has been reported in vitro (26–29) but not in vivo. DCs from pLN, mLN, and spleen of untreated immunodeficient Rag^-/- mice consistently expressed more CD80 and CD86 than those from immunosufficient Rag^-/- littermates (Figure 2A). These statistically significant increases, although relatively small, were seen in both MHCII^+CD11c^+ migratory and MHC^-/+CD11c^- blood-derived DC populations, including both the CD8^-/^- and CD11b^-/^- splenic subsets. We excluded the possibility that elevated costimulation was due to subclinical microbial infection of Rag^-/- mice by showing that costimulatory molecules were equivalently elevated on DCs from immunodeficient mice housed under germ-free (GF) and specific pathogen-free (SPF) conditions (Figure 2B).

We next examined whether Treg reconstitution downregulated DC expression of CD80/CD86 in Rag^-/- mice. Preliminary experiments indicated that expression of CD80/CD86 was unaffected by the IL-2 treatment required to support Treg reconstitution (Supplemental Figure 3). DCs in Treg-reconstituted mice showed a uniform shift in surface expression of CD80/CD86, as illustrated for migratory DCs in the pLN (Figure 2C), indicating that comparing the mean fluorescent intensity (MFI) of CD80/CD86 expression provided a valid measure of the extent of downregulation. Splenic DCs and migratory DCs from pLN and mLN downregulated both CD80 and CD86 in response to Treg reconstitution (Figure 2D). In pLN, the extent of CD80/CD86 downregulation correlated with the number of Tregs transferred, with costimulation returning to WT levels only in mice that received at least 0.5 × 10^6 Tregs. In mLN, CD80 was significantly reduced only at the highest Treg cell dose (Figure 2D). To determine whether this difference could be explained by the comparatively poor Treg reconstitution in mLN (Figure 1B), we measured the correlation between Treg frequency and expression of CD80/CD86 on migratory DCs in individual
pLN and mLN of Treg-reconstituted mice, using the data shown in Figure 2D. Because the steady-state levels of costimulation in pLN and mLN are different, costimulation data from each LN was normalized relative to the appropriate mean in WT mice. This analysis demonstrated a clear inverse correlation between Treg frequency and costimulation in individual LN (Figure 3A).

In the spleen, transfer of as few as 0.25 × 10⁶ Tregs reduced costimulation to WT levels (Figure 2D). This might reflect a more activated Treg phenotype in spleen as compared with LN, consistent with recent studies of BLIMP-reporter mice (30). However our data indicated that almost all Tregs in reconstituted pLN were lower for all groups of Treg-reconstituted mice compared with WT controls (Supplemental Figure 4B). The splenic and mLN are different, costimulation data from each LN was normalized relative to the appropriate mean in WT mice. This analysis was limited to pLN and spleen, and summarized for all LIP experiments performed, totaling 6 independent experiments for each group. Statistical analysis was performed using one-way ANOVA with a Newman-Keuls post-test. **P < 0.01, ***P < 0.001, ****P < 0.0001.

Figure 4. Effect of CTLA-4 expression by Tregs on modulation of CD80/CD86 and inhibition of fast-phase LIR. Rag–/– hosts were reconstituted with flow-sorted WT or Cta-4–/– Tregs. Cta-4–/– and WT Tregs were purified from mixed BM chimeras generated with an 80:20 ratio of Cta-4–/–/WT BM. Fully reconstituted chimeras (≥3 months after irradiation) were treated with IL-2/JES6-1 at –5, –4, and –3 days to expand Treg numbers before harvest at day 0. Treg reconstitution was performed as described in Figure 1. (A) MFI of CD80 (left) and CD86 (right) on migratory DCs from pLN (n = 4 per group). Bars represent mean ± SEM with individual values indicated by the open circles. Data are from 1 of 2 independent experiments. (B) 1 × 10⁶ CFSE-labeled WT polyclonal CD4+ T cells were transferred into Rag–/– mice, either unreeconstituted or reconstituted with WT or Cta-4–/–deficient Tregs (n = 3 per group). All groups were treated with second daily IL-2/JES6-1 throughout the course of the experiment. One group of Rag–/– mice reconstituted with WT Tregs received water supplemented with 5 mg/ml 1-MT from 3 days before CFSE-labeled cell transfer until organ harvest. Shown are representative CFSE division profiles from the pLN and spleen at day 7 after transfer. (B) The ratio of fully divided CFSE− T cells (>7 divisions) to CFSE+ T cells (0–3 divisions) within adoptively transferred CD4+ T cells was calculated for pLN and spleen, and summarized for all LIP experiments performed, totaling 6 independent experiments for each group. Statistical analysis was performed using one-way ANOVA with a Newman-Keuls post-test. **P < 0.01, ***P < 0.001, ****P < 0.0001.

highest Treg dose failed to normalize the Treg/DC ratio (Supplemental Figure 4C). When the degree of Treg reconstitution was adjusted to reflect the Treg/DC ratio in individual lymphoid organs, an inverse relationship with total costimulation (MFI multiplied by DC number relative to WT) was apparent for all organs (Figure 3B). Normal levels of CD80 were achieved at approximately 50% of normal Treg/DC ratio in LN and spleen, consistent with the highly activated phenotype of reconstituting Tregs (Supplemental Figure 4A). We next correlated DC expression of costimulation at the time of T cell transfer with the rapid spontaneous T cell proliferative response shown in Figure 1. The size of the T cell response (indicated by the ratio of CFSE− cells [≥7 divisions] to CFSE+ T cells [0–3 divisions]) in Treg-reconstituted Rag–/– mice was normalized against Rag–/– controls in order to allow comparison between organs in which baseline proliferation differed. This analysis was limited to pLN and mLN, since the spleen contains a mixture of T cells that have undergone activation and proliferation in situ and divided T cells that have migrated from other sites. When plotted against expression of CD80 and CD86 in Treg-reconstituted mice (also normalized against expression in Rag–/– controls), a strong positive correlation between CD80/CD86 expression and T cell proliferation was observed (Figure 3C).
Taken together, these data suggest that, under steady-state conditions, the capacity of Tregs to control costimulation is a function of the ratio of Tregs to DCs. As the Treg/DC ratio drops below normal, costimulation increases and fast-phase LIP ensues as individual CD4+ T cells receive sufficient TCR/CD28-dependent stimulation to divide. Thus Tregs can be considered as a fail-safe against autoimmunity, backing up thymic negative selection of high-affinity self-reactivity by fine-tuning costimulation to control the peripheral T cell activation threshold.

CTLA-4 expression by Tregs is required for modulation of costimulation and fast-phase LIP. Treg-dependent downregulation of CD80/CD86 expression in vitro requires CTLA-4 (28, 29). To test whether Tregs also use CTLA-4 in vivo, we compared the function of CTLA-4-deficient and WT Tregs in reconstituted Rag-/- mice. To generate populations of WT and CTLA-4-deficient Tregs with equivalent activation status at the time of adoptive transfer, we made mixed WT/CTLA-4-/- BM chimeras in which the dominant effect of WT CTLA-4-/- Tregs prevented the spontaneous activation and immunopathology present in CTLA-4-/- mice (31). CTLA-4-/- and WT Tregs isolated from these chimeras had comparable purity and Foxp3 expression levels (Supplemental Figure 5, A and B). Reconstitution of Rag-/- and WT Tregs isolated from these chimeras had comparable purity and Foxp3 expression levels (Supplemental Figure 5, A and B). Reconstitution of Rag-/- mice with CTLA-4-deficient Tregs failed to reduce expression of either CD80 or CD86 (Figure 4A). This was not due to suboptimal Treg reconstitution, as the number of Foxp3+ Tregs was higher in the CTLA-4-deficient group (Supplemental Figure 5C), most likely due to the absence of the cell-intrinsic CTLA-4–mediated negative signal. Involvement of CTLA-4 in Treg-dependent downregulation of CD80/CD86 expression, combined with the approximately 2-fold increase in CTLA-4 levels in reconstituting Tregs (Supplemental Figure 4A), may explain why only 50% of the normal number of Tregs was required to reduce costimulation to WT levels (Figure 3B).

Next, we assessed the extent to which fast-phase LIP was controlled by CTLA-4-dependent downregulation of costimulation. Compared with WT Tregs, CTLA-4-deficient Tregs were significantly impaired in their ability to inhibit fast-phase LIP in Rag-/- mice (Figure 4, B and C), although some residual suppressive activity of CTLA-4-deficient Tregs was apparent, particularly in pLN. The mechanism mediating such residual suppres-
sion in our model is currently unknown. However, our finding that deletion of a single molecule, CTLA-4, in Tregs simultaneously impaired their ability to control DC costimulation and to prevent fast-phase LIP supports a model in which downregulation of DC costimulation is a major mechanism used by Tregs to control fast-phase LIP.

We also tested whether CTLA-4–dependent induction of the immunomodulatory enzyme indoleamine 2,3-dioxygenase (IDO) was involved in Treg-dependent suppression, as previously reported (32, 33). Blocking IDO function by administering 1-methyl-DL-tryptophan (1-MT) (34) did not significantly impair Treg-dependent inhibition of fast-phase LIP (Figure 4, B and C), indicating no detectable role for IDO in our experimental model.

Figure 6. Effect of expansion of Treg populations in immunosufficient mice on DC costimulation and T cell proliferation. WT mice were treated with IL-2/JES6-1 complexes at days 0, 1, and 2, and lymphoid organs were harvested at day 5. (A) Representative plots of Foxp3 versus CD25 expression by CD4+ T cells from pLN and spleen. Numbers indicate the percentage of cells within the gate. (B) Absolute number of Foxp3+ Tregs in pLN and spleen of untreated versus IL-2/JES6-1–treated WT mice. (C) MFI of CD80 and CD86 expression by migratory DCs in pLN (left panels) and CD8+ and CD11b+ subsets of splenic DCs (right panels). Data in A–C are representative of 3 independent experiments with n = 5 per group. (D) Proliferation of adoptively transferred 5C.C7 T cells in response to 1 μg i.v. peptide in WT hosts pretreated with IL-2/JES6-1 complexes at days −5, −4, and −3. Left panel: representative CFSE division profiles from pLN and spleen at day 3 after T cell transfer for IL-2/JES6-1 treated hosts (unfilled histograms) compared with untreated WT hosts (filled histograms). Right panel: MFI of CFSE expression by adoptively transferred 5C.C7 T cells (n = 5 mice per group). Data are from a single experiment. Statistical analysis of B–D was performed using 2-tailed unpaired t tests. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.

Treg reconstitution normalizes enhanced CD4+ T cell responses to foreign antigen. If Tregs mediate at least part of their suppressive function by downregulating costimulation, then they should also inhibit high-affinity responses. To test this, we substituted as a read-out the well-defined response of 5C.C7 TCR transgenic (tg) CD4+ T cells to their cognate antigen, moth cytochrome c (MCC) (35, 36). We predicted that higher constitutive expression of CD80/CD86 by DCs in unmanipulated Rag–/– hosts (Figure 2A) should enhance the naive 5C.C7 T cell response, particularly at suboptimal antigen doses. 5C.C7 TCR tg T cells showed increased proliferation to MCC peptide ligand in Rag–/– hosts compared with WT hosts, most pronounced at lower peptide doses (Figure 5A). Treg reconstitution reduced 5C.C7 proliferation to peptide in the pLN to the level seen in
Effect of conventional T cells on costimulation and fast-phase LIP. (A) Rag\(^{-/-}\) mice were reconstituted with 2.5 × 10^6 Tregs or naive conventional T cells at a dose of 1.25 × 10^6 or 2.5 × 10^6 (n = 4 per group). Naive conventional T cells were flow-sorted as CD3^+ CD4^+ CD25^− CD38^− CD44^−. Expression of CD80 (left panel) and CD86 (right panel) by migratory DCs in pLN is shown. Data are representative of 2 independent experiments. (B) 10^4 CFSE-labeled polyclonal CD4^+ T cells were transferred into unreconstituted, Treg-reconstituted, or conventional T cell–reconstituted Rag\(^{-/-}\) mice. Second daily IL-2/JES6-1 treatment was continued for all groups until lymphoid organ harvest 7 days after CFSE-labeled cell transfer. Adoptively transferred CD4^+ WT T cells were identified based on differential expression of CD45 alleles. Left panel: Representative CFSE division profiles from the pLN day 7 after transfer. Right panel: Ratio of CFSE^− T cells (>7 divisions) to CFSE^+ T cells (0–3 divisions) within adoptively transferred T cells was calculated (n = 6 per group). Data are pooled from 2 independent experiments. (C) Proliferation of adoptively transferred 5C.C7 T cells in response to 1 μg i.v. peptide in Rag\(^{-/-}\) hosts, either unreconstituted or reconstituted with 2.5 × 10^6 Tregs or conventional T cells (n = 4 per group). Left panel: Representative CFSE division profiles in pLN of Rag\(^{-/-}\) hosts (unfilled histograms) compared with WT controls (filled histograms) 3 days after adoptive transfer of 5C.C7 T cells. Right panel: MFI of CFSE expression by adoptively transferred 5C.C7 T cells. Data are representative of 2 independent experiments. Statistical analysis of A–C was performed using one-way ANOVA with a Newman-Keuls post-test. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.

WT hosts (Figure 5B), consistent with the reduction in CD80/CD86 (Figure 2, C and D). This effect was abolished when Rag\(^{-/-}\) mice were reconstituted with CTLA-4–deficient Tregs (Figure 5C). Once again, inhibition of IDO had no effect on suppression by WT Tregs. Treg reconstitution similarly suppressed the 5C.C7 T cell response in the spleen but not in the mLN, consistent with the relatively low level of Treg reconstitution in the latter organ (not shown).

Steady-state regulation of DC costimulation by Tregs in immunosufficient mice. The data reported above indicate that Tregs downregulate CD80/CD86 expression during reconstitution of immunodeficient mice. To test whether they also downregulate costimulation in immunosufficient mice under steady-state conditions, we expanded Treg populations by administering IL-2 complexes to WT mice. IL-2 treatment increased Treg numbers (Figure 6, A and B) and concomitantly reduced expression of CD80 and CD86 (Figure 6C). After Treg expansion, antigen-specific T cell proliferation was also reduced (Figure 6D). Thus Tregs mediate tonic downregulation of costimulation under lymphosufficient and lymphopenic conditions.

Effect of conventional T cells on costimulation and fast-phase LIP. In the absence of Tregs, oligoclonal CD4^+ T cells reconstitute the immune system by means of fast-phase LIP. To model the downstream effects of ongoing LIP and compare them to Treg reconstitution, we adoptively transferred Rag\(^{-/-}\) hosts with naive conventional CD4^+ T cells and compared them to hosts reconstituted with 2.5 × 10^6 Tregs. All host mice were administered IL-2 complexes during reconstitution. Because conventional T cells produced generally larger but more variable numbers of reconstituting cells than Tregs, irrespective of whether they were provided with exogenous IL-2 (Supplemental Figure 6), each experiment included groups reconstituted with either 1.25 × 10^6 or 2.5 × 10^6 conventional T cells to bracket the size of the reconstituted Treg population (not shown). In marked contrast to the effect of Tregs, both doses of conventional T cells upregulated expression of costimulation on DCs in pLN (Figure 7A) and spleen (not shown). These opposing effects of Tregs and conventional T cells on costimulation appear to be specific for CD80/CD86, as we observed minimal changes in
expression of other costimulatory molecules under the same conditions (Supplemental Figure 7). In addition to increasing expression of CD80/CD86, T cell reconstitution also significantly increased fast-phase LIP in both pLN (Figure 7B) and spleen (not shown), but had no effect on 5C.C7 T cell responses to specific peptide (Figure 7C), possibly because the antigen-specific T cells in unreconstituted Rag–/– hosts were already proliferating at a maximal rate.

**Effect of Treg depletion in immunosufficient mice.** Spontaneous immune reconstitution from a state of partial lymphopenia is likely to proceed in the presence of a mixed Treg/conventional T cell compartment, with each population exerting their essentially opposing effects. To model this situation, we used DEREG (depletion of regulatory T cell) mice expressing the diphtheria toxin receptor (DT receptor) under the control of the FoxP3 promoter (37). After administration of 2 doses of 1 μg DT on days 0 and 1, Treg frequency reached a minimum on day 2, while expression of CD80 and CD86 increased (Figure 8A and Supplemental Figure 8, A and B). The extent of CD80/CD86 upregulation on day 2 was similar to that seen previously in immunodeficient mice (Figure 2). Between days 2 and 4, a further increase in costimulation was seen, although Treg frequency was either stable or increased. This finding is consistent with a secondary, positive effect of conventional T cells undergoing slow-phase homeostatic LIP. In addition to undergoing fast-phase LIP, adoptively transferred CD4+ T cells undergo a slower homeostatic form of CD28-independent, IL-7–dependent proliferation in immunodeficient hosts, producing 1 or 2 cell divisions within 7 days of adoptive transfer (19, 25). This type of homeostatic proliferation maintains a highly diverse T cell repertoire of naive cells (39, 40). In the experiment shown in Figure 7B, it appeared that reconstitution with conventional T cells, but not Tregs, inhibited slow-phase LIP. This was confirmed by analysis of T cell phenotype, which showed that the proportion of cells with a naïve CD44+CD62L+ phenotype was lower in hosts reconstituted with conventional T cells compared with Tregs (Figure 9, A and B). In order to study the slow-phase LIP response in a more sensitive model, we examined spontaneous proliferation of T cells isolated from Cd28–/– mice, which do not undergo fast-phase LIP (19, 25). Conventional T cells strongly inhibited slow-phase LIP, whereas Tregs had very little effect (Figure 9C), likely due to their lower expression of IL-7R (41). This, thus, provided a more sensitive model for examining slow-phase LIP. When CD4+ T cells were adoptively transferred to DEREG mice, Tregs inhibited costimulation in a dose-dependent manner (Figure 9D).

**Figure 8. Effect of Treg depletion in immunosufficient animals on expression of costimulation.** (A and B) DEREG mice were treated with 1 μg DT i.p. at days 0 and 1. MFI of CD80 (upper panels) and CD86 expression (lower panels) by migratory DCs from pLN (left) or splenic DCs (right; unbroken lines) graphed against Tregs as a percentage of CD4+ T cells (dotted lines) between days 0 and 3 (A), and at days 0 and 3–7 (B). Data in A and B are mean ± SEM (n = 3–4 per group) and are each from a single experiment. (C) WT mice were treated with 500 μg anti-CD25 (clone PC61) 18 hours before harvest of spleens. Controls received either PBS, or 500 μg of anti-Vα11 (clone R8B) (n = 2–4 per group). MFI of CD80 and CD86 expression by unfractionated CD11c+B220– splenic DCs from a single experiment is shown. (D) WT mice were irradiated with 700 cGy, and spleens were harvested at 0, 3, or 24 hours. MFI of CD80 and CD86 expression by unfractionated CD11c+B220– splenic DCs is shown (n = 5–7 per group). Data are from a single experiment. Statistical analysis of A–D was performed using one-way ANOVA with a Newman-Keuls post-test. Bars represent mean ± SEM with individual values indicated by the open circles. *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.
ing repopulation of the lymphoid pool via slow homeostatic proliferation, while conventional T cells in the absence of Tregs undergo fast-phase LIP, which in turn inhibits homeostatic division.

Treg reconstitution in a murine model of allogeneic BMT. Having demonstrated the effectiveness of Treg reconstitution in normalizing both DC costimulation and T cell responses in lymphopenic Rag−/− mice, we next tested the relevance of these findings to BMT, in which the immune system reconstitutes a transiently lymphopenic environment produced by the BMT conditioning regimen. We performed allogeneic parent into F1 BMT, in which T cell depleted BM (TCDBM) from B10.BR mice was transplanted into lethally irradiated C57BL/6 × B10.BR/F1 mice (hereafter designated [B6.BR]/F1). BMT recipients were reconstituted with either allogeneic donor-type Tregs (B10.BR) or syngeneic host-type Tregs ([B6.BR]/F1). For Treg reconstitution, recipients were adoptively transferred with 2.5 × 10^6 Tregs at the time of BM transfer and treated second daily (on days 0, 2, 4, 6, and 8) with IL-2/JES6-1 to support Treg survival and expansion, as described for Rag−/− mice. Donor-derived Tregs could be clearly identified at day 7 after BMT in Treg-reconstituted mice but not in controls that received BM alone (Figure 10, A and B). Both allogeneic B10.BR and syngeneic [B6.BR]/F1 Tregs were capable of reconstituting irradiated [B6.BR]/F1 hosts. Mice that received BM alone contained a small population of residual host Tregs, which was not increased by treatment with IL-2/JES6-1 complexes.

The effect of Treg reconstitution on DC phenotype following BMT was then tested by examining migratory DCs in the pLN, which at day 7 after irradiation primarily consists of a radio-resistant population comprising Langerhans cells and a small subset of dermal DCs (42, 43). The rapid increase in DC costimulation observed at 24 hours after irradiation (Figure 8D) was maintained at day 7 after BMT, with expression of CD80 and CD86 on pLN migratory DCs increasing by 4-fold and 3-fold, respectively, compared with DCs from nonirradiated controls (Figure 10C). Reconstitution with either B10.BR or [B6.BR]/F1 Tregs reduced expression of both CD80 and CD86 to a level that did not significantly differ from nonirradiated WT controls. Consistent with the failure of IL-2/JES6-1 to increase the size of the residual Treg pool, treatment of irradiated BMT controls with IL-2/JES6-1 had no effect on expression of CD80/CD86 (Figure 10C). Expression of CD80/CD86 by DCs negatively correlated with Treg frequency for all treatment groups (Figure 10D), replicating the results in the Rag−/− model.

Treg reconstitution following BMT protects against GVHD. Finally, we tested if Treg reconstitution and the associated reduction in costimulatory molecules limits CD4+ T cell-mediated GVHD in BMT recipients. Previous studies have demonstrated that cotransfer of purified Tregs at the time of T cell infusion can reduce GVHD, albeit with variable efficacy (10, 16, 44, 45). Published effects of cotransfer range from delay in disease onset to near complete protection against GVHD, depending on a variety of factors including number, activation status, and specificity of the transferred T cells. In one report, a 2-day delay in T cell administration after Treg transfer significantly improved survival (17).

We transferred 2.5 × 10^6 conventional CD4+CD25+ T cells from B10.BR donors into [B6.BR]/F1 hosts on day 7 after irradiation, allowing time to achieve Treg reconstitution of BMT recipients prior to T cell transfer. Transfer of T cells into unreconsti-
against GVHD development, possibly due to the severe disease in the B10.BR into [B6.BR]F1 model. Median survival was 18.5 days after BMT (11.5 days after T cell transfer) (Figure 10F) — not significantly different from the group that received T cells alone ($P = 0.20$). In contrast, Treg reconstitution prior to T cell transfer prevented GVHD in all animals, regardless of whether
tuted BMT recipients resulted in rapid, highly lethal GVHD in which mice failed to recover any of their irradiation-dependent weight loss (Figure 10E). The median survival of this group was 16 days after BMT, i.e., 9 days after T cell transfer (Figure 10F). Cotransfer of $2.5 \times 10^6$ allogeneic B10.BR Tregs at the time of T cell transfer (a 1:1 Treg/T cell ratio) had minimal protective effect against GVHD development, possibly due to the severe disease in the B10.BR into [B6.BR]F1 model. Median survival was 18.5 days after BMT (11.5 days after T cell transfer) (Figure 10F) — not significantly different from the group that received T cells alone ($P = 0.20$). In contrast, Treg reconstitution prior to T cell transfer prevented GVHD in all animals, regardless of whether

Figure 10. Effect of Treg reconstitution following BMT. (A) [B6.BR]F1 mice were lethally irradiated and transplanted with $5 \times 10^6$ TCD B10.BR BM cells. Mice received $2.5 \times 10^6$ CD4+CD25+ Tregs from B10.BR or [B6.BR]F1 donors at the time of BMT with IL-2/JES6-1 second daily. Controls included nonirradiated (WT) mice and irradiated mice that received BM alone (BMT) or BM and IL-2/JES6-1 (BMT + IL-2) ($n = 4$ /group). Top panel: Foxp3 and CD25 expression on total CD4+ T cells in pLN 7 days after BMT. Bottom panel: CD4+Foxp3+ Tregs were divided into host origin (CD45.2+), B10.BR donor origin (CD45.1+) and [B6.BR]F1 donor origin (CD45.1+/CD45.2+). (B) Foxp3' Treg frequency of total CD4+ T cells (top panel), and total CD45-expressing cells (bottom panel). (C) Expression of CD80 (left) and CD86 (right) on pLN migratory DCs at day 7 after BMT. Bars represent mean ± SEM with individual values indicated by open circles. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$, and ****$P < 0.0001$. (D) MFI of CD80 (left panel) and CD86 (right panel) plotted against pLN Treg frequency among CD45-expressing cells. A line of best fit with $R^2$ value is displayed. Data in A–D are representative of 2 independent experiments. (E and F) BMT recipients were administered $2.5 \times 10^6$ B10.BR CD4+CD25+ T cells at day 7 after BMT. Groups included BMT recipients reconstituted with B10.BR or [B6.BR]F1 Tregs prior to T cell transfer, BMT recipients that received T cells alone, BMT recipients in which 1:1 B10.BR Tregs and T cells were cotransferred at day 7, and BMT recipients that did not receive T cells ($n = 9$–10/group). Shown are mouse weights (E) and a Kaplan-Meier survival analysis (F). Data are pooled from 2 independent experiments.
transferred Tregs were allogeneic donor type (B10.BR) or syngeneic host type ([B6.BR]F1). Mice reconstituted with Tregs (either allogeneic or syngeneic) prior to T cell transfer showed similar weight changes to those in controls that received BMT but no T cell transfer (Figure 10E). Importantly, no deaths were recorded in Treg-reconstituted mice (Figure 10F). It should be noted that mice with high starting weights but no GVHD typically recovered to only 85%-90% of their preirradiation weight (Supplemental Figure 9A). The 100% survival rate in Treg-reconstituted mice represented a highly significant improvement to survival when compared with the Treg/T cell cotransfer protocol (P = 0.0009 for B10.BR Treg reconstitution, P = 0.0005 for [B6.BR]F1 Treg reconstitution). The protective effect was not due to IL-2/JES6-1 treatment of Treg-reconstituted mice, as second daily IL-2/JES6-1 treatment failed to prolong survival in the absence of Treg cell reconstitution (Supplemental Figure 9, B and C).

Discussion
A major consideration in clinical settings of lymphopenia is the importance of achieving adequate immune reconstitution while avoiding complications due to oligoclonal T cell proliferation. The peripheral T cell pool expands to a greater size and expresses a more diverse TCR repertoire when Tregs are present during early immune reconstitution (5, 14, 39). Additionally, therapies that augment Treg numbers during lymphopenia have been demonstrated to be beneficial for the prevention of GVHD (9–16). However, it is not clear how Tregs mediate these beneficial effects.

Here, we have developed a mouse model to examine how Tregs control immune reconstitution. Using this model, we demonstrated that reconstitution of the Treg compartment prior to T cell transfer prevented rapid-phase proliferation of potentially pathogenic T cells, without impairing the slow-phase homeostatic form of LIP. In contrast, reconstitution with conventional CD4+ T cells enhanced subsequent fast-phase LIP and inhibited slow-phase immune reconstitution. Our data identify DC overexpression of the costimulatory molecules CD80/CD86 as the major driving force behind the fast-phase proliferative response in lymphopenia. The strong correlation between Treg cell number, DC expression of CD80/CD86, the extent of fast-phase LIP, and the magnitude of antigen-specific T cell responses in immunodefficient mice led us to hypothesize that Tregs play a critical role in controlling the steady-state level of DC costimulation under physiological conditions. We tested this function of Tregs in immunosufficient mice by demonstrating a similar relationship between Treg number and DC expression of CD80/CD86 following transient expansion or depleting of Tregs (summarized in Supplemental Figure 10). Deletion of a single molecule, CTLA-4, simultaneously impaired the ability of Tregs to control DC costimulation and prevent fast-phase LIP, supporting a model in which the 2 functions of Tregs are mechanistically linked. Our results indicate that CTLA-4–dependent downregulation of DC costimulation is a major mechanism used by Tregs to regulate the peripheral T cell–activation threshold. However, additional mechanisms must also be involved, particularly in LN where CTLA-4–deficient Tregs were still able to partially suppress fast-phase LIP. Finally, and most importantly for the potential application of our findings in clinical practice, we demonstrated that our findings in lymphopenic Rag–/– hosts were relevant to mice undergoing BMT; with Treg reconstitution similarly capable of mediating downregulation of CD80/CD86 and providing full protection against CD4+ T cell–mediated GVHD.

Our findings extend published data showing a CTLA-4–dependent reduction in expression of costimulatory molecules on DCs cocultured with Tregs (26–29) by demonstrating that CTLA-4 is also responsible for downregulation of CD80 and CD86 on DCs under steady-state conditions in vivo. Both secreted and cell-surface forms of CTLA-4 have been implicated in regulation of DC costimulation in vitro (46, 47), and cell-surface CTLA-4 expressed by high affinity conventional T cells was demonstrated to capture and internalize its ligands in vivo, a process termed transendocytosis. Whether Tregs also use transendocytosis in vivo is not yet known.

In our experiments, DC expression of CD80/CD86 was highly correlated with the ratio of Treg to DC. Our finding of the importance of the Treg/DC ratio is consistent with published studies in a variety of models in which induction of nonspecific CD4+ T cell lymphopenia leads to autoimmunity, despite a normal ratio of Tregs to conventional T cells (48–51). Our data suggest that immune reconstitution following cytoablative therapy is likely to benefit from early Treg reconstitution to normalize the ratio of Tregs to DCs, irrespective of the Treg/T cell ratio (2, 10). Indeed, in a murine model of GVHD, infusion of suboptimal Treg numbers was more effective at inhibiting GVHD if conventional T cell transfer was delayed by 2 days (17).

Recently, several studies have demonstrated that certain cytoablative treatments preferentially deplete conventional T cells, leaving Treg populations relatively intact (52, 53). Tregs are relatively resistant to cyclophosphamide, and the effectiveness of this treatment in preventing GVHD is critically dependent upon the presence of Tregs (52, 54). Additionally, treatment with antithymocyte globulin (ATG) preferentially depletes conventional T cells and, in low doses, actually promotes Treg expansion (53, 55). The mechanism by which such expansion inhibits GVHD (52) and promotes transplant tolerance (53) remains undetermined, but our results indicate that such an effect could readily be explained via Treg-dependent reduction in DC costimulation, thus limiting the activation of pathogenic T cells. Additionally, our results predict that early support for Treg reconstitution will be most crucial in patients receiving myeloablative conditioning rather than reduced-intensity conditioning, consistent with a recent report correlating Treg numbers and GVHD incidence after different conditioning for H SCT (13).

Our results suggest that specific targeting of the mechanisms supporting rapid-phase LIP, independent of Treg reconstitution, may also improve clinical outcomes for patients with transient lymphopenia. One possible agent is CTLA-4–Ig, which blocks recognition of CD80/CD86 and has been shown to promote transplant acceptance when combined with ATG therapy (53). In addition, treatments that favor slow-phase over fast-phase LIP may also help to achieve complete immune reconstitution. For example, IL-7 treatment preferentially enhances slow-phase LIP while inhibiting fast-phase proliferation (56, 57), and has been shown to promote faster immune reconstitution without enhancing GVHD (58).
In stark contrast, antitumor responses may benefit from conditions favoring fast-phase LIP during immune reconstitution (59). In general, antitumor responses in mouse (60) and man (61) are enhanced by a reduction in absolute Treg numbers and thus the Treg/DC ratio. In contrast, the graft-versus-leukemia response is maintained in the presence of Tregs, suggesting that it may be of higher affinity and thus less sensitive to deprivation of costimulatory drive (11, 62). However, Treg-depleting strategies come at a cost to TCR diversity and the size of the peripheral lymphocyte pool (5, 11, 14, 39), impairing responses to foreign pathogens such as listeria (39) and MCMV (14). A contributing factor to this impairment appears to be autoimmune damage to sites of T cell expansion, namely pLN, spleen, and thymus (14).

Finally, in addition to their implications for immune reconstitution in patients with iatrogenic lymphopenia, our results may also be of relevance in understanding the initiation of autoimmune. In particular, the importance of the Treg/DC ratio in maintaining steady-state DC costimulation could explain the paradoxical link between autoimmunity and lymphopenia observed in animal models (48–51) and in several human immunodeficiency states (63).

Methods

Mice. All mice were bred and housed under SPF conditions in the Centenary Institute Animal Facility, apart from BALB/cWehi and C.B-17/lcr-scid animals raised under GF or SPF conditions at the Walter & Eliza Hall Institute SPF Unit (Kew, Australia). Rag-1–/– mice (64) were originally a gift from Lynn Corcoran (Walter & Eliza Hall Institute, Parkville, Australia). Rag-2–/– mice (65) were purchased from Taconic Biosciences Inc. DEREG mice (37) were provided by Stephen Alexander (Children’s Hospital, Westmead, Australia). Tg mice expressing the 5C.C7 TCR specific for the COOH-terminal epitope of MCC (35, 36) were bred on a Rag-2–/– background. Cd28–/– mice were originally purchased from The Jackson Laboratory. Ctla4–/– mice (66) obtained from Jim Allison (MD Anderson Cancer Center, Houston, Texas, United States) were maintained as homozygotes on Rag-1–/– and Rag-2–/– backgrounds. F1 crosses were performed to generate immunosufficient (Rag-1–/–Rag-2–/–) Ctla4–/– BM donors. Experimental mice were >8 weeks of age, with the exception of Ctla4–/– BM donors, which were used at 2–3 weeks of age due to early-onset lymphoproliferative disease in these mice.

Reconstitution of lymphopenic mice. Tregs for reconstitution were purified by 2-step magnetic bead enrichment of single-cell suspensions prepared from pooled peripheral and mesenteric LN and spleens. Step 1 involved staining with a mixture of hybridoma supernatants (anti-B220 [RA3-6B2], anti-CD8α [53–6.7], anti-CD11b [M1/70], and anti-erythroid lineage [Ter119]) (Centenary Institute) detected with anti-rat microbeads (Miltenyi Biotec), followed by negative selection to enrich for CD4+ T cells. In step 2, CD25-expressing Tregs were positively selected by staining with anti–CD25-PE (BD Biosciences) followed by anti-PE MACS beads (Miltenyi Biotec). FoxP3 was expressed by 90%–95% of the CD4+CD25+ population. Naive conventional T cells for reconstitution were purified from the bead-enriched CD4+ population by flow cytometric sorting for CD4+CD38+CD44 cells using an AriaII digital flow sorter (BD Biosciences). Treg contamination was <1%. Purified CD4+CD25+ Tregs or conventional CD4+CD38+CD44 cells from WT mice were adoptively transferred into syngeneic Rag-1–/– or Rag-2–/– hosts and expanded by second daily treatment with IL-2/JES6-1 complexes (24), consisting of 1 µg IL-2 complexed to 5 µg JES6-1 mAb.

T cell responses. For ILP, the bead-selected CD4+CD25+ population was CFSE labeled (5 × 106 cells/ml in RPMI + 1% FCS, 5 mM CFSE for 5 minutes at 37°C) and 1 × 105 cells were injected i.v. into syngeneic hosts. For antigen-specific T cell responses, pLN cells from 5C.C7 TCR tg Rag-2–/– mice were CFSE labeled and 1 × 106 CD4+ T cells were adoptively transferred into syngeneic hosts together with i.v. MCC peptide (residues 86–103). Where indicated, IDO was inhibited by the addition of 1-MT (Sigma-Aldrich) to drinking water at a concentration of 5 mg/ml, as previously described (34).

Manipulation of Treg number in immunosufficient mice. Tregs were expanded in WT mice by administering IL-2/JES6-1 at days 0, 1, and 2 before analyzing Treg number and DC phenotype on day 5. Tregs were depleted in DEREG mice by i.p. injection of 1 µg DT (Sigma-Aldrich) on days 0 and 1 (67). Alternatively, Tregs were depleted in WT mice by i.p. injection of 500 µg anti-CD25 mAb (clone PC61) 18 hours prior to organ harvest. Controls received either 500 µg anti-TCRαVα11 mAb (clone RR8) or PBS (Centenary Institute). For radiation-induced depletion of lymphocytes including Tregs, mice were irradiated with 700 cGy and lymphoid organs were harvested 3 or 24 hours later.

Generation and purification of CTLA-4-deficient Tregs. CTLA-4 mixed BM chimeras were generated by irradiating CD45.1+/– WT host mice with 2× 600 cGy 24 hours prior to adoptive transfer of 5 × 106 BM cells (4 × 106 CD45.1+/2Ctla4–/– plus 1 × 106 CD45.2+/– WT). After reconstitution for >3 months, Treg numbers were boosted by administration of IL-2/JES6-1 at days 0, 1, and 2 before lymphoid organ harvest on day 5. CD4+ T cells were enriched by one round of negative bead selection, as detailed above, and then flow sorted for CD4+CD25+ cells expressing CD45.1+/2 (Ctla4–/–) or CD45.2+/– (WT).

BM T and development of GVHD. [B6.BR]F1 hosts were lethally irradiated with 1,200 cGy (given as a split-dose of 2× 600 cGy separated by 3 hours) 24 hours prior to adoptive transfer of 5 × 106 BM cells from B10.BR donors. BM cells were TCD prior to transfer by magnetic bead depletion of T cells stained with hybridoma supernatants (anti-Thy1.2 and/or a mixture of anti-CD4 [GK1.5] and anti-CD8α [53–6.7]) and detected with anti-rat microbeads (Miltenyi Biotec). To reduce the incidence of postirradiation coagulopathy due to suspected vitamin K deficiency, mice were routinely treated with 1 mg vitamin K on day 1 prior to irradiation, and at 3 and 7 days after irradiation. Treg reconstitution was performed by transferring 2.5 × 106 B10.BR or [B6.BR]F1 CD4+CD25+ Tregs at the time of BM transfer and treating with IL-2/JES6-1 complexes at days 0, 2, 4, 6, and 8. GVHD was induced by transfer of 2.5 × 106 CD4+CD25– T cells 7 days after BM transfer. GVHD monitoring consisted of weighing 2–3 times per week, plus visual monitoring for activity, posture, fur texture, and skin integrity as previously described (68). Mice were euthanized if visual signs of GVHD were present and weight loss exceeded 15%, or if total weight loss reached 25%.

FACS analysis. For analysis of T cells, single-cell suspensions were prepared from pLN, mLN, or spleen. For analysis of DCs, organs were first enzymatically digested in 2 mg/ml collagenase IV (Sigma-Aldrich). mAbs used for cell-surface staining were all from BD Biosciences. Mice were first enzymatically digested in 2 mg/ml collagenase IV (Sigma-Aldrich). mAbs used for cell-surface staining were all from BD Biosciences.
staining kit (eBioscience) and anti-Foxp3 mAb (clone FJK-16s, eBioscience). Cells were analyzed using CantoII, LSRII, or LSRFortessa digital flow cytometers (all BD Biosciences). Data analysis was performed using FlowJo software (Tree Star Inc.).

Statistics. Statistical analysis was performed using GraphPad Prism version 5.0d. When comparing 2 groups, a 2-tailed unpaired t test was performed. Comparison of multiple groups was performed using one-way ANOVA with a Newman-Keuls post-test. Statistical analysis of cell–division ratios was performed on log-transformed data. Analysis of Kaplan-Meier survival curves was performed using a log-rank test with a Gehan-Breslow-Wilcoxon post-test. P values < 0.05 were considered significant.

Study approval. All experiments were performed with the consent of the University of Sydney or Sydney Local Health District Animal Ethics Committee.

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