

HHS Public Access

Author manuscript *Cell Immunol.* Author manuscript; available in PMC 2019 January 01.

Published in final edited form as:

Cell Immunol. 2018 January ; 323: 9-18. doi:10.1016/j.cellimm.2017.08.008.

High PD-L1/CD86 MFI ratio and IL-10 secretion characterize human regulatory dendritic cells generated for clinical testing in organ transplantation

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Abstract

Human regulatory dendritic cells (DCreg) were generated from CD14 immunobead-purified or elutriated monocytes in the presence of vitamin D3 and IL-10. They exhibited similar, low levels of costimulatory CD80 and CD86, but comparatively high levels of co-inhibitory programed death ligand-1 (PD-L1) and IL-10 production compared to control immature DC (iDC). Following Tolllike receptor 4 ligation, unlike control iDC, DCreg resisted phenotypic and functional maturation and further upregulated PD-L1:CD86 expression. Whereas LPS-stimulated control iDC (mature DC; matDC) secreted pro-inflammatory tumor necrosis factor a, but no IL-10, the converse was observed for LPS-stimulated DCreg. DCreg weakly stimulated naïve and memory allogeneic CD4⁺ and CD8⁺ T cell proliferation and IFN_γ, IL-17A and perforin/granzyme B production in MLR. Their stimulatory function was enhanced however, by blocking PD-1 ligation. Highthroughput T cell receptor (TCR) sequencing revealed that, among circulating T cell subsets, memory CD8⁺ T cells contained the most alloreactive TCR clonotypes and that, while matDC expanded these alloreactive memory CD8 TCR clonotypes, DCreg induced more attenuated responses. These findings demonstrate the feasibility of generating highly-purified GMP-grade DCreg for systemic infusion, their influence on the alloreactive T cell response, and a key mechanistic role of the PD1 pathway.

Keywords

human; dendritic cells; immune regulation; cell therapy

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1. Introduction

Currently, there is major interest in the potential of adoptive cell therapy with innate or adaptive immune cells for the control of allograft rejection [1, 2] and reduced patient dependence on pharmacologic immunosuppression. Based on extensive preclinical studies, both regulatory myeloid cells and regulatory T cells (Treg) offer considerable promise. Thus, dendritic cells (DC) with regulatory function (DCreg) maintain self-tolerance in the normal steady-state [3], terminate memory T cell responses [4] and regulate auto- and alloimmunity [5]. Ex vivo generation of DC in the presence of specific pharmacologic agents or anti-inflammatory cytokines inhibits their maturation, confers resistance to maturation and promotes their tolerogenicity [6]. These DCreg subvert naïve and memory T cell responses by various mechanisms [7]. Moreover, their administration as cell-based vectors inhibits autoimmune disorders [8], graft-versus-host disease [9] and organ transplant rejection in experimental animals [7, 10].

When infused into rodent allograft recipients before transplantation, with or without immunosuppressive agents, DCreg of donor origin promote long-term graft survival or tolerance [7]. Similar results have been obtained in mice or rats by infusing autologous DC [10, 11]. More recently, using a clinically-relevant, nonhuman primate (NHP) renal allograft model, we have shown [12] that pre-transplant infusion of DCreg of donor origin, together with a short course of minimal immunosuppression, prolongs graft survival, without evidence of host sensitization. Collectively, these findings raise expectations of the therapeutic potential of DCreg for promotion or restoration of tolerance in the clinic [13, 14].

Recently, phase 1 safety trials of autologous monocyte-derived DCreg, including those pulsed with autoAg and administered locally to patients with certain autoimmune diseases have been conducted [15–18]. The results indicate that DCreg are well-tolerated, and a phase 1/2 trial of autologous DCreg in live donor renal transplantation (NCT02252055) has begun as part of The ONE Study (www.onestudy.org) [19].

In preparation for first-in-human testing of donor-derived (allogeneic) DCreg in organ transplantation, we have characterized DCreg generated in vitamin (Vit)D3 and IL-10 from circulating monocytes isolated by either immunobead selection or elutriation. We show that adequate numbers of highly-purified DCreg that meet release criteria can be generated readily from elutriated monocytes under good manufacturing practice (GMP) conditions to meet envisioned target doses for systemic infusion. These DCreg resist maturation,- an important pre-requisite for their adoptive transfer into prospective graft recipients. They also express high levels of co-inhibitory programed death ligand-1 (PD-L1) relative to co-stimulatory CD86. Furthermore, PD1 pathway ligation plays a key role in regulating their allogeneic T cell stimulatory function.

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2. Materials and Methods

2.1. DC generation and stimulation

Monocytes were isolated from either normal human buffy coat PBMC (LeukoPaks; Central Blood Bank of Pittsburgh) using CD14-specific immunobeads (Miltenyi, San Diego, CA), or obtained as the elutriated monocyte fraction of normal human volunteer leukapheresis products (Institute for Transfusion Medicine, Pittsburgh, PA) under an IRB-approved protocol. Immature DC (imDC) were generated as described [20], except that those generated from elutriated monocytes were cultured in DC serum-free media (Corning) without antibiotics using GMP grade reagents. DCreg were generated by addition of VitD3 (20mM; Sigma, St. Louis, MO) on days 0 and 4 and rhIL-10 (60ng/ml; Peprotech, Rocky Hill, NJ) on day 4 of culture. Lipopolysaccharide (LPS; *Salmonella minnesota* R595; 0.5 μ g/ml, Enzo Life Sciences, Farmingdale, NY) was added to DC cultures on day 6 to evaluate their response to a potent maturation-inducing agent. Responses to soluble CD40 ligand (sCD40L; MegaCD40L, 100ng/ml, Enzo Life Sciences) or a pro-inflammatory cytokine cocktail (PCC) consisting of IL-6, TNFa, IL-1 β (all 10ng/ml; Peprotech) and prostaglandin (PG)E₂ (1 μ g/ml; Sigma) were also evaluated. Upon harvest, DC were washed thoroughly (1× PBS) before phenotypic and functional analyses.

2.2. Flow cytometric analysis

Cell surface marker and intracellular cytokine staining was performed as described [20]. Details of the panel of monoclonal Abs used are shown in Supplementary Table 1. Flow cytometry was performed on a BD Fortessa cytometer and data analyzed using FlowJo software (FlowJo, LLC Ashland, OR).

2.3. T cell proliferation assays

Allogeneic pan T cells were isolated (Miltenyi), labeled with carboxyfluorescein succinimidyl ester (CFSE; 2 μ M; Invitrogen, Waltham, MA) and co-cultured for 5 days at 37°C in 5% CO₂ in air with DC (1DC:10T cell ratio). T cell proliferation was estimated by CFSE dilution. For some MLR, anti-PD-L1 (10 μ g/ml, Biolegend, San Diego, CA) and anti-PD-1 blocking Abs (10 μ g/ml; Affymetrix/eBioscience), anti-IL-6R blocking Ab (400 μ g/ml; Genentech, San Francisco, CA), anti-IL-10R blocking Ab (10 μ g/ml; Biolegend) or IgG1 isotype control (10 μ g/ml; Affymetrix/eBioscience) was added at the start of culture.

2.4. Cytokine cytometric bead array (CBA) assay

Cytokines in culture supernatants were quantified using BD Bioscience (San Jose, CA) Human Cytometric Bead Array (CBA) Flex kits, according to the manufacturer's instructions.

2.5. Clonotypic analysis of alloreactive T cells

Naïve (CD45RA⁺) or memory (CD45RO⁺) CD4⁺ and CD8⁺ T cells were isolated from healthy controls by FACS. Sorted subsets were CFSE-labeled and incubated with allogeneic DCreg or matDC (1DC:10T cell ratio) in 5-day CFSE-MLR. Controls consisted of unstimulated T cell subsets or DC alone. Genomic DNA was extracted using the Qiagen

DNeasy Blood & Tissue Kit (Cat No./ID: 69506; Qiagen, Gaithersburg, MD), quantified and shipped to Adaptive Biotechnologies (Seattle, WA) for T cell receptor (TCR) β complementarity determining region (CDR3) amplification, sequencing and data analysis. Briefly, a multiplexed two-step polymerase chain reaction (PCR) method was employed using a mixture of forward primers specific to TCR V β gene segments and reverse primers specific to TCR J β gene segments. V, D and J gene definitions were based on annotation in accordance with the IMGT database. TCR libraries were sequenced using an Illumina instrument according to the manufacturer's instructions. The set of observed biological TCR β CDR3 sequences were normalized to correct for residual multiplex PCR amplification bias and quantified against a set of synthetic TCR β CDR3 sequence analogues with proprietary software (Adaptive Biotechnologies).

2.6. Statistical analyses

Results are expressed as means ± SEM. Differences between means were determined using a paired Student 't'-test, Wilcoxon signed-rank test or Mann-Whitney U test, as appropriate. 'P' values 0.05 were considered significant. For sequencing analysis, summary statistics were obtained from the immunoSEQ Analyzer toolset. Repertoire diversity was measured as one minus normalized Shannon's entropy, commonly referred to as "clonality". Clonal expansion was measured as a statistically significant change in clonal abundance using Fisher's exact test. The Benjamini-Hochberg Procedure was used to control the false discovery rate with a 'P' value <0.01 for the two-tailed test considered significant.

3. Results

3.1. DCreg generation, purity and yield

Initially, we generated DCreg in small-scale cultures from CD14 immunobead-purified monocytes isolated from 10 separate healthy individual buffy coats. We compared the results with those obtained subsequently following large scale-up production of DCreg from monocytes elutriated from 4 individual healthy donor leukapheresis products under GMP conditions. Table 1 shows the much greater number of DCreg generated under GMP scale-up conditions. The mean theoretical yield of DCreg from a single whole leukapheresis product was 593 [range 290–820] $\pm 266 \times 10^6$. This number of DCreg from a single donor leukapheresis product would be sufficient to infuse from 4.1 to 11.7 (mean 8.5) $\times 10^6$ /kg into a 70 kg recipient, a dose range equivalent to/in excess of that which we have shown [12] can prolong graft survival in NHP (3.5–10.10⁶/kg). DCreg recovered from both sources of monocytes readily met release criteria: >80% viability, >85% purity (lin⁻ HLA-DR⁺ CD11c ⁺) and <0.3% contaminating T or B cells (release criteria <1% of each) (Fig. 1).

3.2. DCreg are phenotypically immature and resist maturation following exposure to proinflammatory stimuli

We compared the cell surface phenotype of DC populations generated from bead-isolated or elutriated monocytes. HLA-DR, CD11c, CD40, CD86, and PD-L1/2 were expressed on >90% of imDC and DCreg (Fig. 2A). However, DCreg expressed comparatively low levels (MFI) of HLA-DR and CD86, but much higher levels of PD-L1 compared with imDC (Fig. 2B). An important prerequisite of DCreg generated for therapeutic use, is the ability to resist

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maturation under pro-inflammatory conditions. Expression of HLA-DR, CD40, CD86, CD83 and CD14 on unstimulated DCreg were very similar to those on LPS-stimulated DCreg (Fig. 2B), indicating they resisted maturation induced by a potent Toll-like receptor agonist. The DCreg also resisted phenotypic maturation when cultured with a PCC or sCD40L (Supplementary Fig. 1). Interestingly, LPS-stimulated DCreg displayed similar, high levels of PD-L1 to unstimulated DCreg (Fig. 2B). Furthermore, although PD-L2 was expressed at lower levels than PD-L1, it followed the same pattern of expression as PD-L1 (LPS-stimulated DCreg compared with DCreg; Fig. 2B).

3.3. DCreg express a high PD-L1:CD86 ratio

The PD-L1:CD86 MFI ratio expressed by DCreg was significantly higher than that expressed by imDC (Fig. 2C). Importantly, this elevated PD-L1:CD86 ratio was maintained on DCreg generated from bead-isolated or elutriated monocytes following exposure to LPS (Fig. 2C), the PCC or sCD40L (Supplementary Fig. 1) and remained significantly higher than that observed for matDC. Similar observations were made for PD-L2 (Supplementary Fig. 2).

3.4. DCreg secrete high levels of IL-10 but not TNFa

We quantified levels of multiple cytokines in 7-day DC cultures and in 5-day allogeneic MLR supernatants. IL-6, IL-10 and TNFa production profiles of DC generated from beaded or elutriated monocytes was very similar, with no significant differences in cytokine concentrations between these populations (Fig. 3). Notably, IL-10 was produced by DCreg, but not by control imDC. IL-1β, IL-4 and IL-12p70 were not detected under any culture conditions. Cytokine levels in DCreg culture supernatants were also compared before and after LPS stimulation. Highly variable concentrations of IL-6, increased levels of TNFa and profoundly decreased concentrations of IL-10 (compared with DCreg) were observed in matDC cultures, whereas LPS-stimulated DCreg supernatants exhibited similar high levels of IL-10 and minimal TNFa (high IL-10:TNFa ratios) to those of control, unstimulated DCreg. LPS-stimulated DCreg exhibited similar, variable levels of IL-6 production to matDC (Fig. 3). Linear regression analysis (Excel) was performed to determine whether IL-6 production by bead-derived or elutriated monocyte-derived DCreg during co-culture with allogeneic T cells correlated with T cell proliferation or DCreg IL-10 production levels. No correlation was observed between IL-6 production levels and T cell proliferation or DCreg IL-10 production.

3.5. DCreg are weak stimulators of allogeneic T cell proliferation

DCreg induced significantly lower levels of CD4⁺ and CD8⁺ T cell proliferation than imDC and matDC (Fig. 4). Further, following LPS stimulation, DCreg induced significantly lower levels of CD4⁺ and CD8⁺ T cell proliferation than control LPS-stimulated matDC. These results characterize DCreg as weak allostimulatory antigen-presenting cells and further demonstrate their ability to resist functional maturation. The poor T cell stimulatory ability of DCreg compared to imDC and matDC was almost identical, further showing that DCreg with similar function can be generated from CD14⁺ or elutriated monocytes (Fig. 4).

3.6. DCreg induce minimal levels of proinflammatory cytokines and cytotoxic effector molecule production, but enhanced IL-10 production in MLR

As shown in Fig. 5A, IFN γ , IL-17A, IL-4 and granzyme B expression by alloactivated CD4⁺ and CD8⁺ T cells stimulated by DCreg was markedly attenuated compared with T cells stimulated with control matDC. Levels of secreted IL-12p70, IL-1 β , IL-6, TNF α and IFN γ in DCreg-MLR culture supernatants were much lower than those in matDC-MLR supernatants. IL-10 was the only cytokine present at increased levels in DCreg-MLR compared to matDC-MLR supernatants (Fig. 5B). LPS-stimulated DCreg were associated with lower levels of IL-12p70, IL-6, TNF α , IL-17A and IFN γ , and an increased concentration of secreted IL-10 compared to matDC.

3.7 High PD-L1 expression by DCreg attenuates allogeneic T cell proliferation and IFN γ production

The PD-L1/PD-L2-PD1 pathway plays an important role in maintaining T cell homeostasis and in induction/maintenance of peripheral tolerance [21, 22]. To ascertain whether PD-L1 expression at markedly elevated levels by DCreg was responsible for their poor T cell stimulatory function, we added PD-L1 and PD-1 blocking Abs at the start of DCregallogeneic T cell co-cultures. This significantly increased CD4⁺ and CD8⁺ T cell proliferation (Fig. 6A), markedly enhanced IFN γ secretion, and increased IL-6 and TNF α levels when compared to co-cultures to which no blocking Abs were added (Fig. 6B). To test the role of IL-10, that was secreted by unstimulated and LPS-stimulated DCreg, but not by imDC or matDC, anti-IL-10R Ab was added along with PD-L1 and PD-1 blocking Ab, but its addition did not further affect T cell proliferation. Thus, PD-L1 expression by DCreg plays a key role in restraining allogeneic T cell proliferation and IFN γ production.

3.8. DCreg allostimulation induces more limited TCR clonotype expansion of memory CD8⁺ T cells than matDC

We used high throughput TCR immune-sequencing with DNA purified from flow-sorted naïve and memory CD4⁺ and CD8⁺ T cells of two healthy control subjects before and after allo-stimulation with either DCreg or matDC in CFSE-MLR (Fig. 7A). This allowed us assess the overall TCR clonal repertoire within resting T cells and the frequency of alloreactive TCR clonotypes before and after exposure to allogeneic DC. Circulating naïve CD4⁺ and CD8⁺ T cells displayed highly diverse (polyclonal) TCR repertoires, whereas memory T cells were less diverse, with CD8⁺ T cells being the most oligoclonal (Fig. 7B). Resting (unstimulated) memory CD8⁺ T cells (among all T cell subsets tested) contained most alloreactive TCR clonotypes that were shared and correlated positively with alloreactive TCR clonotypes after DCreg or matDC stimulation (Fig. 7C). For both subjects, allostimulation triggered clonal expansion, however DCreg induced more limited alloreactive memory CD8⁺ T cell clonal expansion. For both DC types, the existing repertoire of expanded clones was largely undisturbed, and there was no evidence of clones disappearing from the CD8⁺ memory repertoire upon stimulation.

3.9. Mock infusion of DCreg and their detection in allogeneic whole blood

DCreg harvested from two GMP scale-up preparations were placed (350×10^6) in infusion medium in a transfer bag, held at 4°C for 4 hr, then mock infusion performed slowly over 1 hr at RT using an infusion set (190 cm line) incorporating a 180 µm filter (CareFusion, San Diego CA). In each case, 80% of the 'infused' cells passed through the entire infusion set, including the filter. Additionally, we examined the presence and phenotype of DCreg following their addition to fresh allogeneic whole blood. As shown in Fig. 8, the allogeneic DCreg were readily detected using mAb directed against MHC class I Ag not shared with the 'recipient'.

4. Discussion

Crucial to the manufacture of human DCreg for infusion into organ allograft recipients,whether from healthy donors or patients with end-stage organ failure (recipient-derived DCreg), is their generation in sufficient purity and quantity under GMP conditions to achieve safe and effective target cell doses. DCreg must exhibit functional stability and resistance to maturation in the face of inflammatory stimuli to minimize any potential risk of undesired immunostimulatory effects in vivo (host sensitization). While there is no current consensus regarding the optimal choice of agents for generation of clinical grade DCreg [15–18, 23], our previous studies in NHP, including those given renal transplants, provide a strong rationale for generation of DCreg using VitD3 and IL-10. Thus, these DCreg induce functional unresponsiveness in alloreactive T cells and inhibit allograft rejection [12, 24]. While each agent alone inhibits DC maturation and confers tolerogenic properties on DC [6], we found by comparing VitD3 and IL-10 with other agents, that their combination resulted in the optimal yield of maturation-resistant DCreg. The present report shows that this approach readily allows the generation of human DCreg from monocytes purified either from peripheral blood buffy coats using CD14 immunobeads, or from monocytes elutriated from non-mobilized leukapheresis products. The immunophenotype, high level of purity (<1% T or B cell contamination) and function of DCreg that we generated from both sources was very similar. Since elutriation is the less expensive procedure, it is both an economically (and logistically) preferable choice for human DCreg generation in sufficient quantities for their systemic administration.

In recent phase 1 studies of local adoptive transfer of DCreg into patients with autoimmune disease, doses in the range $1-10 \times 10^6$ have been administered either intradermally (type-1 diabetes), intradermally or arthroscopically (rheumatoid arthritis) or intraperitoneally (Crohn's disease) [15–18], with no evidence of significant adverse effects. Such doses of DCreg given locally can silence effector T cells and induce Ag-specific Treg in human volunteers [25, 26] and immunoregulatory and anti-inflammatory effects in patients with autoimmune disease [17]. Systemic infusion of related, donor-derived (allogeneic) regulatory myeloid cells (regulatory macrophages; 7 or 8×10^6 cells/kg, via central line) in renal transplant patients is also well-tolerated [27], with patients maintained on low dose tacrolimus immunosuppression three years post-transplant. The present findings demonstrate our ability to generate adequate numbers of monocyte-derived DCreg from a single, non-mobilized donor leukapheresis product under GMP conditions to allow infusion of up to $5 \times$

10⁶ DCreg /kg. This is within the therapeutic range that has proved efficacious in safely prolonging graft survival in minimally-immunosuppressed NHP transplant recipients [12].

The DCreg we generated expressed high levels of PD-L1, that played a key role in attenuating their T cell stimulatory ability. They also exhibited much higher PD-L1:CD86 ratios than control DC. This high ratio was preserved following DCreg stimulation with the potent myeloid DC activating agent LPS, a PCC or sCD40L. In addition, the DCreg secreted high levels of IL-10, but little or no IL-12. In addition to adequate cell number, viability, purity and sterility, each of which each constitute critical product release criteria, we propose that the PD-L1: CD86 ratio be considered as a potential additional release criterion, or provide the basis of a DCreg potency assay based on experience gained manufacturing these cells for use in patients.

Evidence from rodent and NHP studies suggests that, even though their survival may be limited in allogeneic hosts, donor-derived DCreg have the capacity to downregulate T cell responses via the direct, indirect and semi-direct pathways, leading to decreased alloAg-specific Tmem responses that may reflect clonal regulation/exhaustion [28, 29]. The human DCreg generated in the present study expressed high levels of PD-L1 that transmits an inhibitory signal via PD1 that, in turn, reduces proliferation of activated CD8⁺ T cells and promotes the development, maintenance and function of induced Treg [22]. Moreover, the DCreg also secreted anti-inflammatory IL-10, that potently suppresses DC, T cell and NK cell effector functions and promotes CD4⁺ type-1 Treg [30].

With respect to our TCR repertoire sequencing data, there is considerable heterogeneity within the human alloreactive memory CD8T cell pool that can be ascribed to genetic differences between individuals and epigenetic "instructions" gained by human T cells after exposure to viruses, bacteria, alloAg etc, throughout life. We also understand that these differences may contribute to heterogeneous modulation of alloreactive T cell clonotype responses by DCreg. Indeed, the results of our two TCR immunosequencing experiments suggest just this. Overall, they show that, compared with matDC, DCreg induce more limited alloreactive memory CD8⁺ T cell clonal expansion. Notably, donor-derived VitD3/ IL-10 DCreg attenuate alloreactive CD8⁺ T memory cell responses in renal-allografted NHP [12, 29]. We plan to set up TCR immunosequencing each time we infuse donor-derived DCreg into a transplant patient, and thus obtain the personalized mechanism/signature of donor-specific T cell hyporesponsiveness that we postulate will be induced.

5. Conclusion

In conclusion, DCreg meeting criteria for systemic infusion can be generated under GMP conditions and display properties that may be conducive to therapeutic efficacy in organ transplantation.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

The authors' work was supported by National Institutes of Health (NIH) Clinical Trial Planning Grant R34 AI 123033 (AWT), CTSI grant NIH U01 TR000005 (SR) and by the Starzl Transplantation Institute Hammill Family and Robert and Courtney Bost Funds. We thank Veronica Sutherland and Misty DeRiggi for technical support in the DCreg scale-up studies and acknowledge the generous participation of the volunteers without whom the study could not have been undertaken. The study used the University of Pittsburgh Cancer Institute CCSG-supported Immunologic Monitoring and Cellular Products Laboratory core facility, which is supported by NIH grant P30 CA047904-24.

Abbreviations

DC(s)	dendritic cell(s)
DCreg	regulatory DC
GLP	good manufacturing practice
imDC	immature DC
matDC	mature DC
PCC	pro-inflammatory cytokine cocktail
PD-(L)1	programed death (ligand)1

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Highlights

- GMP grade regulatory dendritic cells can be generated from human blood monocytes in adequate numbers for systemic infusion
- DCreg exhibit high levels of PD-L1 expression and IL-10 production
- DCreg resist phenotypic and functional maturation
- Blocking PD-1 ligation upregulates their T cell stimulatory function
- High throughput TCR sequencing shows DCreg induce attenuated memory T cell responses

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Fig. 1.

Human regulatory dendritic cells (DCreg) generated from bead-isolated or elutriated monocytes display highly purity, with very low levels of T cell contamination. Flow cytometric analysis of HLA-DR, CD3 and CD4⁺ versus CD8⁺ cells is shown before and after DCreg generation from (A) CD14⁺ bead-isolated monocytes (upper panels) or (B) elutriated monocytes (lower panels). (C) shows CD11c expression at the start and end of cultures (from elutriated monocytes). Data are representative of 2 individual normal donors.

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Fig. 2.

Cell surface phenotype of DCreg. Percent positive cells and mean fluorescence intensity (MFI) were determined for MHC class II (HLA-DR), the DC markers CD11c, CD1c and CD209 (= DC-SIGN; data not shown), T cell costimulatory (CD40, CD80 and CD86) and co-inhibitory (PD-L1 [CD274] and PD-L2 [CD273]) molecules, DC maturation (CD83) and myeloid cell (CD14) marker, and secondary lymphoid tissue homing receptor (CCR7; CD197). (A) Representative examples of cell surface marker expression by immature (im)DC and regulatory DC (DCreg) generated from CD14⁺ bead-isolated monocytes (upper panels) or elutriated monocytes (lower panels). (B) Cell surface marker expression (MFI) by

DC populations generated from bead-isolated monocytes (n=9 individuals) or elutriated monocytes (n=7 individuals). Significances of differences were determined using Wilcoxon t-test. *p<0.01; [†]p<0.05' NS = not significant. (C) PDL-1:CD86 MFI ratio of DC cultured from bead-isolated monocytes (n=8; left panel) or elutriated monocytes (n=8, right panel). Significances of differences were determined using Wilcoxon t-test. [†]p<0.05; NS = not significant. Data are means ±1SEM. matDC, LPS-stimulated imDC; DCreg+LPS, DCreg stimulated with LPS.



Fig. 3.

Cytokine secretion by DCreg. DC populations were generated from (A) CD14⁺ beadisolated (n=8 individuals) or (B) elutriated monocytes (n=7 individuals) then cultured overnight in the absence or presence of LPS. Cytokine levels in supernatants were determined by cytometric bead array assay. Data are means \pm 1SEM. imDC, immature DC; DCreg, regulatory DC; matDC, imDC stimulated with LPS; DCreg + LPS, DCreg stimulated with LPS.[†], p<0.05; NS = not significant.

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Fig. 4.

Proliferation of allogeneic T cells stimulated with DCreg. DC were generated from (A) bead-isolated monocytes (n=7 individuals) or (B) elutriated monocytes (n=7 individuals). CFSE-labeled allogeneic T cells were co-cultured for 5 days with various populations of DC (1DC:10T cells). imDC or DCreg were stimulated for 18 to 20 hr with LPS (0.2µg/ml) to generate matDC and DCregLPS, respectively. T cell proliferation was measured by CFSE dilution. Data are means ±1SEM. Significances of differences were determined using Wilcoxon t-test. [†], p<0.05; NS = not significant.



Fig. 5.

Cytokine expression by allogeneic T cells co-cultured with DCreg. (A) Intracellular cytokine expression by allogeneic CD4⁺ (upper panels) and CD8⁺ T cells (lower panels) stimulated with DCreg or matDC generated from bead-isolated monocytes. CFSE-labeled allogeneic pan T cells were co-cultured for 5 days with DC (1DC:10T cells). The DC-stimulated T cells were harvested, then stimulated for 5 h with aCD2,CD3,CD28 microbeads in the presence of Golgi-plug. (B) Cytokine production in MLR cultures of allogeneic T cells stimulated with DC populations generated from bead-isolated monocytes (n=5 individuals;

upper panels) or elutriated monocytes (n=7 individuals; lower panels). Data are means $\pm 1SEM.~^{\dagger},~p{<}0.05.$

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Fig. 6.

The PD1 pathway plays a key role in DCreg function. (A) CFSE-labelled allogeneic T cells were co-cultured with DCreg for 5 days (1DC:10T cells). T cell proliferation was determined by CFSE dilution. PD-L1 and PD-1 blocking antibodies $(10\mu g/ml) \pm$ anti-IL-10R antibody $(10\mu g/ml)$ or IgG1 isotype control $(10\mu g/ml)$ were added at the start of culture. (B) Cytokine levels in MLR cultures. Data are means \pm 1SEM. Significances of differences were determined by using Wilcoxon t-test. *, p<0.01; [†], p<0.05. NS = not significant.

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Fig. 7.

Clonotypic analysis of alloreactive T cells stimulated with DCreg. (A) Gating strategy used to FACS sort naïve (N) and memory (M) CD4 and CD8 T cell subsets. Each sorted population was cultured with allogeneic DCreg or matDC in 5-day CFSE-MLR. DNA purified from each condition was amplified by PCR amplified and sequenced for the TCRVβ locus using the Adaptive Technologies platform. (B) TCR repertoire diversity within unstimulated T cell subsets (n=2 healthy individuals). (C) Comparison of shared alloreactive TCR clonotypes between unstimulated M CD8⁺ T cells (x-axis) and DC-stimulated M CD8⁺ T cells (y-axis). Dots on the axes represent the frequency of TCR clonotypes that were found in one population but not the other, while those close to the diagonal dotted line represent frequency equality. Significantly expanded clones are highlighted in orange. The elliptical dotted line signifies the threshold for statistical comparison. Representative data from Subject 1. (D) Numbers of alloreactive clones expanded after 5-day MLR stimulation with allogeneic DCreg or matDC. Data from both subjects are shown.



Fig. 8.

Detection of allogeneic DCreg in whole blood. DCreg were generated from elutriated monocytes of an A2 (MHC class 1) positive individual, incubated in 5% goat serum for 15 min on ice, then added to 50µL fresh whole blood (~ 5.10^5 WBC) of an A2 negative individual, so that the final % of allogeneic DCreg was 10% or 5%. An Ab cocktail containing FITC anti-A2 Ab, together with Abs against DC and costimulatory (CD86) and coinhibitory (PD-L1) markers was then added for 20 min at 4°C in the dark. Two ml of 1×BD lysis/fixation buffer was added. After 10 min at 4°C in the dark, the cells were washed thoroughly before flow cytometric analysis.

Table 1

Yield of DCreg from bead-isolated and elutriated monocytes

V					
LeukoPak ID	PBMC used for monocyte bead (×10 ⁶)	Bead-isolated monocytes (×10 ⁶)	Cultured DCreg _{Bead} recovered (×10 ⁶)	Total LeukoPak PBMC (×10 ⁶)	Theoretical # of cultured DCreg _{Bead} (×10 ⁶)
3747	200	50	6	546	23
3751	150	44	2	442	5
3747	190	72	4	546	10
6003	300	89	21	616	44
5058	170	43	13	976	73
774	160	34	5	564	17
<i>7</i> 72	300	130	34	333	37
1455	160	47	8	588	32
1928	170	69	13	453	33
1021	90	28	9	642	41
Mean(10)	170	49	8	555	33
SEM	21	10	3	54	9
В					
Leukopheres ID	Elutriated is monocytes (×10 ⁶)	Cultured DCreg _{Elut} recovered (×10 ⁶)	Total elutriate monocytes (×10 ⁶)	Theoretical # of cultured DCreg _{Elut} (×10 ⁶)	
CPL15-34	640	178	2900	812	1
CPL 15–39	620	255	2000	820	
CPL16-1	640	93	2000	290	
CPL16-15	960	350	2000	450	
Mean([,]	4) 715	219	2225	593	
SEI	M 82	55	225	133	1