

Human memory FOXP3⁺ Tregs secrete IL-17 ex vivo and constitutively express the T_H17 lineage-specific transcription factor ROR γ T

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Recent studies have suggested a close relationship between CD4⁺FOXP3⁺ regulatory T cells (Tregs) and proinflammatory IL-17-producing T helper cells (T_H17) expressing the lineage-specific transcription factor ROR γ T. We report here the unexpected finding that human memory Tregs secrete IL-17 ex vivo and constitutively express ROR γ T. IL-17-secreting Tregs share some phenotypic and functional features with conventional T_H17 cells, expressing high levels of CCR4 and CCR6 and low levels of CXCR3. However, unlike conventional T_H17 cells, they express low levels of CD161 and mostly fail to cosecrete IL-22 and TNF- α ex vivo. Ex vivo secretion of IL-17 and constitutive expression of ROR γ T by human memory Tregs suggest that, in addition to their well-known suppressive functions, these cells likely play additional, as yet undescribed, proinflammatory functions.

CD4⁺ T cells encompass different subsets that express lineage-specific transcription factors and play different roles, not only in initiating and supporting the development of immune responses, but also in orchestrating and regulating them. T helper type 1 (T_H1) cells, which differentiate in the presence of IL-12 and IFN- γ , express the transcription factor T-bet and assure protection against intracellular pathogens and cancer (1). T_H2 cells differentiate in the presence of and produce IL-4, express the transcription factor GATA-3, and are involved in allergic reactions and protection against extracellular parasites (1). Whereas the T_H1 and T_H2 subsets have been described since the late 1980s (2), 2 additional CD4⁺ T cell subsets, regulatory/suppressor T cells (Tregs) and T helper IL-17-producing cells (T_H17), have been described more recently. Tregs are characterized by the expression of the lineage-specific transcription factor FOXP3, which is involved both in their development and in their suppressor functions (3, 4), and they express high levels of CD25 and low levels of CD127 (5–7). At variance with T_H1 and T_H2 cells that are memory cells, human FOXP3⁺ Tregs exist both at the naïve (NnTreg) and the memory (MTreg) stages (8). FOXP3⁺ Tregs are suppressive ex vivo, anergic, and fail to secrete IL-2 or IFN- γ (4, 9). They play a major role in the control of self-tolerance—their depletion or functional alteration resulting in the development of autoimmune diseases—and are involved in the regulation of T cell homeostasis as well as of immune responses to pathogens, alloantigens, and cancer (10). Finally, the most recently defined subset, T_H17 cells, produce IL-17A and IL-17F, 2 members of the IL-17 family that have similar proinflammatory functions, including recruitment and activation of neutrophils, and trigger the production of other proinflammatory chemokines and cytokines by monocytes, endothelial cells, and epithelial cells (11, 12). Several studies have suggested that T_H17 cells are involved in protection against extracellular bacteria and fungi but also are involved in the pathogenesis of certain autoimmune diseases, including experimental autoimmune encephalitis and collagen-induced arthritis (13, 14). The nuclear hormone receptor RA-related orphan receptor γ T (ROR γ T), originally defined as a thymic

specific isoform of ROR γ (15), has been identified as the lineage-specific transcription factor for T_H17 cells, required for their generation (16, 17).

Initial studies assessing the differentiation of Tregs and T_H17 cells from murine naïve CD4⁺ T cells have reported reciprocally regulated and mutually exclusive differentiation programs for the 2 subsets (18, 19). Recently, however, several reports have suggested that the relationship between Tregs and T_H17 cells could be more complex than anticipated, involving plasticity of the respective differentiation programs (20) and in vivo equilibrium of the 2 subsets (21).

In this study, by assessing ex vivo cytokine production by CD4⁺ T cells from healthy donors, we found, unexpectedly, that a significant proportion of circulating memory FOXP3⁺ Tregs secrete IL-17 and express high levels of ROR γ T ex vivo. Together, our findings shed light on the close relationship between human Tregs and T_H17 cells and suggest that, in addition to their well-recognized immune suppressive functions, memory FOXP3⁺ Treg cells likely play additional, as yet undescribed proinflammatory functions.

Results

Human Memory FOXP3⁺ Tregs Secrete IL-17 ex Vivo. Human natural CD4⁺ Tregs are characterized by ex vivo expression of the specific transcription factor FOXP3. As assessed by intracellular staining using specific antibodies, about 5% of total circulating CD4⁺ T cells from healthy donors expressed FOXP3 ex vivo. CD4⁺FOXP3⁺ Tregs included a population of CD45RA⁺ cells corresponding to NnTregs (8), as well as CD45RA⁻ MTregs (Fig. 1A). Most FOXP3⁺ Tregs, both naïve and memory, expressed CD25 and were CD127^{low}. To assess ex vivo cytokine production with respect to differentiation stage and FOXP3 expression, we stimulated purified CD4⁺ T cells with phorbol 12-myristate 13-acetate (PMA)/ionomycin and then stained them with CD45RA-specific, FOXP3-specific, and cytokine-specific antibodies. As expected, significant proportions of memory but not naïve CD4⁺ T cells secreted IFN- γ or IL-17 ex vivo (Fig. 1B). Consistent with the reported inability of FOXP3⁺ Tregs to secrete IFN- γ , most memory cells secreting IFN- γ ex vivo were FOXP3⁻ (Fig. 1C). Unexpectedly, however, a significant proportion (up to 20%) of memory CD4⁺ T cells secreting IL-17 ex vivo were FOXP3⁺ Tregs.

To confirm and quantify ex vivo secretion of IL-17 by MTregs, we isolated CD4⁺ conventional T cell and Treg subsets from

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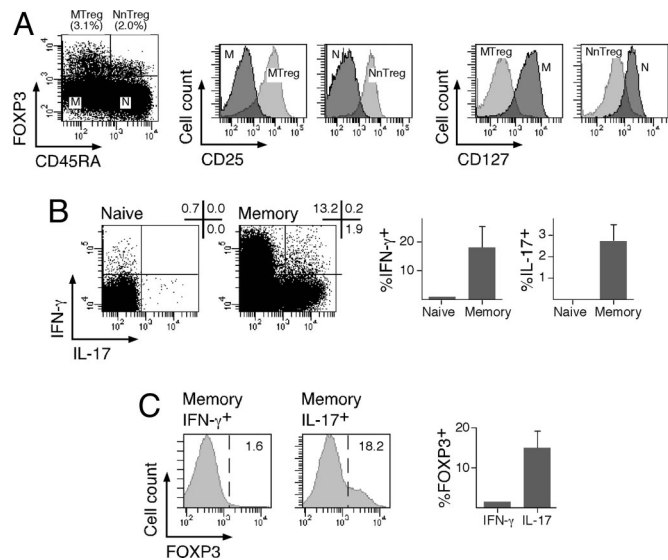


Fig. 1. A fraction of $CD4^+$ T cells producing IL-17 ex vivo express FOXP3. (A) $CD4^+$ T cells isolated from PBMCs of healthy donors were stained with anti-FOXP3, anti-CD25, anti-CD45RA, and anti-CD127 antibodies and analyzed by flow cytometry. Expression of FOXP3 and CD45RA defines memory (M; FOXP3⁺CD45RA⁻) and naïve (N; FOXP3⁻CD45RA⁺) conventional $CD4^+$ T cells and memory (MTreg; FOXP3⁺CD45RA⁻) and naïve (NnTreg; FOXP3⁻CD45RA⁺) Tregs. Histograms show the expression of CD25 and CD127 gated on the defined $CD4^+$ T cell populations. Results are shown for 1 of 3 donors. (B and C) Enriched $CD4^+$ T cells were stimulated for 6 h with PMA/ionomycin and stained with antibodies specific for CD45RA, FOXP3, IL-17, and IFN- γ . (B) The proportion of IL-17-secreting and IFN- γ -secreting cells is shown gated on total naïve (CD45RA⁺) and memory (CD45RA⁻) $CD4^+$ T cells. Dot plots for 1 donor and data for 3 donors are shown (mean \pm SD). (C) The proportion of FOXP3-expressing cells is shown gated on IL-17-secreting and IFN- γ -secreting cells as indicated. Histograms for 1 donor and data for 3 donors are shown (mean \pm SD).

circulating lymphocytes of healthy donors by cell sorting based on the expression of CD25 and CD45RA. The purified populations were assessed for purity (routinely >97%), FOXP3 and CD127 expression (Fig. 2A), and for cytokine production after stimulation with PMA/ionomycin. Isolated FOXP3⁺ MTregs secreted IL-17 ex vivo (Fig. 2B and C). In addition, unlike IL-17-producing T cells in conventional memory $CD4^+$ T cells, which contained a significant proportion of cells coproducing IFN- γ , the large majority of IL-17-producing cells in Tregs did not produce IFN- γ . The proportion of IL-17-secreting MTregs was higher compared with conventional memory $CD4^+$ T cells. Accordingly, *IL17* mRNA levels as well as the levels of secreted IL-17 were higher in stimulated MTregs than in conventional memory $CD4^+$ T cells (Fig. 2D and E). Because the intensity of FOXP3 expression in IL-17⁺ MTregs was slightly lower than in IL-17⁻ MTregs (Fig. 2C), it was important to assess the suppressive function of the identified population. Direct ex vivo assessment of suppressive functions was unfeasible because of the low frequency of the identified population together with the lack of methods allowing the isolation of living IL-17-secreting cells. Therefore, to directly address the suppressive function of FOXP3⁺ IL-17-producing Tregs, we cloned ex vivo-sorted FOXP3⁺CD25⁺CD127^{low}CD4⁺ T cells, isolated clones that had maintained the ex vivo phenotype of the defined population (FOXP3 expression and IL-17 secretion), and assessed their suppressor function (see *SI Materials and Methods*). As shown in Fig. S1, FOXP3⁺ IL-17-producing clones derived from Tregs exhibited high suppressive activity.

Memory FOXP3⁺ Tregs Express ROR γ t ex Vivo. Because IL-17 secretion has been associated with expression of ROR γ t, ex vivo

secretion of IL-17 by MTregs prompted us to assess the expression of ROR γ t in conventional $CD4^+$ T cell and Treg isolated populations ex vivo. Remarkably, by using RT-PCR, we detected ex vivo *RORC* mRNA expression in unstimulated MTregs and, at lower levels, in conventional memory but not in naïve $CD4^+$ T cells (Fig. 3A). To confirm and quantify the increased expression of *RORC* in MTregs, we performed quantitative PCR analysis (Fig. 3B). The results confirmed ex vivo expression of *RORC* in MTregs at levels about 100-fold higher than those found in conventional naïve $CD4^+$ T cells and 4-fold higher than in conventional memory $CD4^+$ T cells. For internal comparison, we also assessed expression of *FOXP3*, *T-bet*, and *GATA3* in ex vivo isolated populations by quantitative PCR (Fig. 3C). As expected, both naïve and memory Tregs expressed highly increased levels of *FOXP3* compared with conventional $CD4^+$ T cells. Ex vivo expression of *T-bet* was slightly higher in conventional memory $CD4^+$ T cells compared with all other populations, and ex vivo expression of *GATA3* was slightly more elevated in Tregs (both NnTregs and MTregs) than in conventional $CD4^+$ T cells. Finally, we assessed ex vivo ROR γ t expression with respect to IL-17 production in isolated populations by intracellular staining after stimulation with PMA/ionomycin, using ROR γ / γ t-specific antibodies that very recently became available. As illustrated in Fig. 3D, this analysis confirmed the presence, among MTregs, of ROR γ t-expressing/IL-17-producing cells, which were found in increased proportions compared with conventional memory $CD4^+$ T cells. No significant levels of ROR γ t-expressing/IL-17-producing cells were detectable in naïve populations.

Memory FOXP3⁺ IL-17-Secreting Tregs Share some but Not all Phenotypic and Functional Features with Conventional T_H17 Cells. To further characterize IL-17-producing MTregs, we compared some of their phenotypic features to those recently described for conventional T_H17 cells. Recent studies have assessed the phenotype of T_H17 cells based on the expression of chemokine receptors. T_H17 cells were exclusively found in the fraction expressing CCR6, a receptor that mediates homing to mucosal tissues and skin (17). Within the CCR6⁺ fraction, cells producing IL-17 but not IFN- γ were found in the population expressing CCR4, previously reported to be preferentially associated with T_H2 cells, whereas those cosecreting IL-17 and IFN- γ (called T_H17/T_H1 cells) (22) were found in the population expressing CXCR3, associated with T_H1 cells. As illustrated in Fig. 4, in line with the ex vivo secretion of IL-17 in the absence of IFN- γ , MTregs were enriched in CCR6⁺ and CCR4⁺ cells but contained lower levels of CXCR3⁺ cells compared with conventional memory $CD4^+$ T cells. Cosmi et al. (23) have recently proposed that CD161, a lectin-like receptor expressed by a subset of human NK and T cells, is a marker of T_H17 cells, particularly in association with CCR6. To assess expression of CD161 in MTregs, we costained purified $CD4^+$ T cells with antibodies specific for CD45RA, CD25, CD161, and CCR6. A large proportion (>40%) of memory but only a few naïve $CD4^+$ T cells expressed CD161 (Fig. 5). In contrast to CCR6, the proportion of CD161-expressing cells was lower among MTregs compared with conventional memory $CD4^+$ T cells. To directly assess the correlation between expression of CCR6/CD161 and IL-17 secretion, we isolated 4 subpopulations of conventional memory $CD4^+$ T cells by cell sorting according to the expression of the 2 markers, and we assessed IL-17 (and IFN- γ) production after stimulation with PMA/ionomycin. We found a strict association between ex vivo expression of CCR6 and IL-17 secretion. In contrast, IL-17 secretion did not strictly segregate with CD161 expression, because the proportion of IL-17-secreting cells within the CCR6⁺CD161⁺ fraction was, on average, only increased 2-fold compared with that found among CCR6⁺CD161⁻

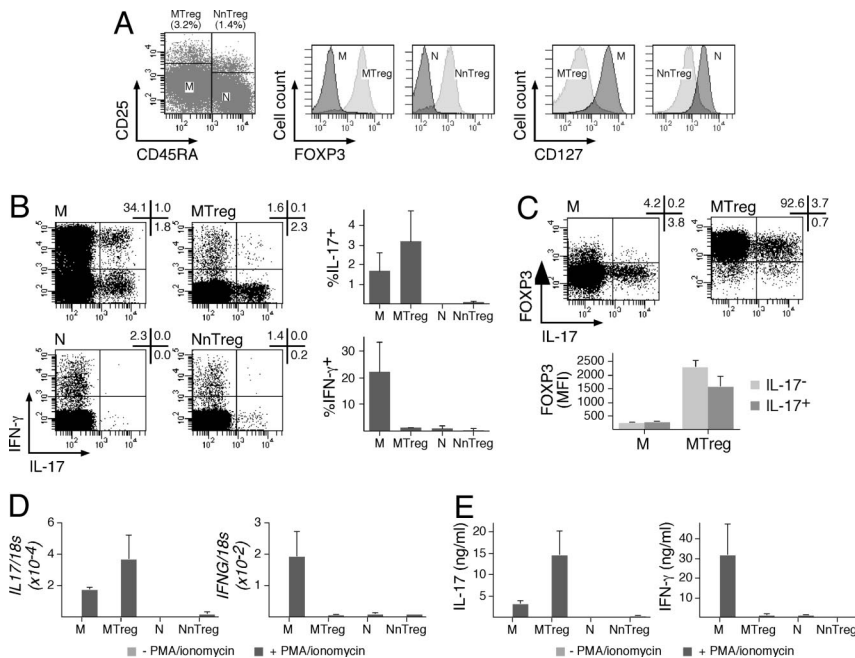


Fig. 2. Human FOXP3⁺ MTregs secrete IL-17 ex vivo. (A) Enriched CD4⁺ T cells were stained with anti-CD8, anti-CD25, and anti-CD45RA antibodies, and CD8⁻ cells were separated by flow cytometry cell sorting into memory (M; CD25⁻CD45RA⁻) and naïve (N; CD25⁻CD45RA⁺) conventional CD4⁺ T cells and memory (MTreg; CD25⁺CD45RA⁻) and naïve (NnTreg; CD25⁺CD45RA⁺) Tregs (dot plot). A fraction of sorted populations was stained with anti-FOXP3 and anti-CD127 antibodies and analyzed by flow cytometry (histograms). Results are shown for 1 of 3 donors. (B) The fraction of IL-17-secreting and IFN- γ -secreting cells among sorted CD4⁺ T cell populations (M, MTreg, N, and NnTreg) was determined by intracellular cytokine staining and flow cytometry analysis after 6 h of stimulation with PMA/ionomycin. Dot plots for 1 donor and data for 3 donors are shown (mean \pm SD). (C) Sorted M and MTreg cells were stimulated for 6 h with PMA/ionomycin and stained with anti-FOXP3 and anti-IL-17 antibodies and analyzed by flow cytometry. Dot plots for 1 of 3 donors are shown (Upper). Mean fluorescence intensity (MFI) of FOXP3 staining in IL-17⁺ and IL-17⁻ cells is shown for 3 donors (mean \pm SD; Lower). (D and E) Sorted CD4⁺ T cell populations were stimulated for 24 h in the absence or presence of PMA/ionomycin, and IL17 and IFNG mRNA levels were determined by quantitative PCR (D), and the quantity of IL-17 and IFN- γ secreted in the culture supernatant was measured by ELISA (E). Data are shown for 3 donors (mean \pm SD).

cells. Thus, lack of CD161 expression by Tregs was not at odds with their capacity to secrete IL-17 ex vivo.

Finally, the ability of MTregs to secrete IL-17 ex vivo

prompted us to assess other cytokines that have been reported to be coproduced by conventional T_H17 cells. With this aim, we stimulated purified MTregs and conventional memory CD4⁺ T cells with PMA/ionomycin ex vivo and costained them with antibodies against IL-17 and IL-22, TNF- α , or IL-21. As illustrated in Fig. 6, whereas up to 20% of conventional CD4⁺ T cells secreting IL-17 ex vivo coproduced IL-22, only a minority (<5%) of IL-17-secreting MTregs cosecreted IL-22. Similarly, whereas the majority (80%) of conventional memory CD4⁺ T cells secreting IL-17 coproduced TNF- α , only half of IL-17-secreting MTregs cosecreted TNF- α . Finally, only a minority of conventional CD4⁺ T cells secreting IL-17 ex vivo coproduced IL-21, and no significant proportions of IL-21-producing cells could be detected among MTregs. Collectively, these results show that MTregs able to secrete IL-17 ex vivo share some but not all phenotypic and functional features with conventional T_H17 cells, supporting the concept that these populations could at least partially colocalize in the same tissues and partially share effector functions.

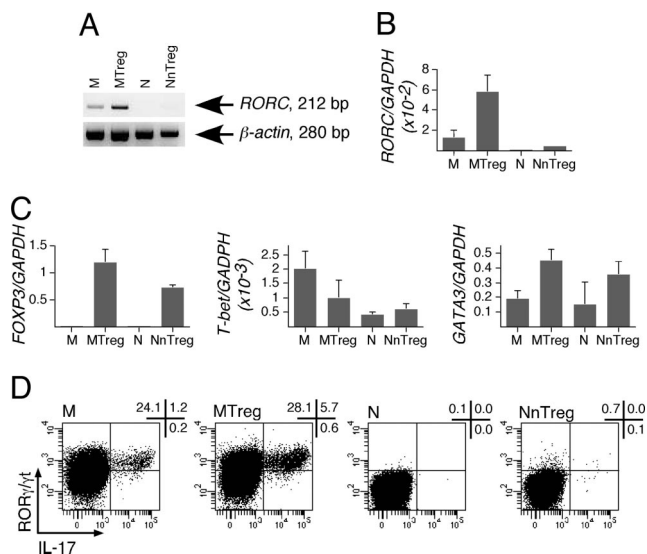


Fig. 3. Ex vivo expression of ROR γ t by MTregs. (A and B) CD4⁺ T cells were separated by flow cytometry as in Fig. 2A, and *RORC* mRNA expression in sorted populations was assessed by conventional (A) and quantitative (B) PCR. (C) *FOXP3*, *T-bet*, and *GATA3* mRNA expression was determined in sorted CD4⁺ T cell populations by quantitative PCR. (D) ROR γ t expression and IL-17 production by sorted populations were assessed by intracellular staining using specific antibodies after 6 h of stimulation with PMA/ionomycin. Results are shown for 1 of 3 donors.

Discussion

We have reported here the unexpected finding that human MTregs secrete IL-17 and express high levels of ROR γ t ex vivo. Whereas previous studies have documented conversion of murine and human Tregs into T_H17 cells after stimulation under various conditions (24–26), the ability of Tregs to secrete IL-17 ex vivo was not appreciated to date. Similarly, expression of ROR γ t in human Tregs was not known before the present report. This finding, however, is consistent with the recent observation by Lochner et al. (21) that up to 50% of murine ROR γ t⁺ T α β cells express FOXP3 and are functionally Treg cells. In addition, forced expression of ROR γ t in murine and

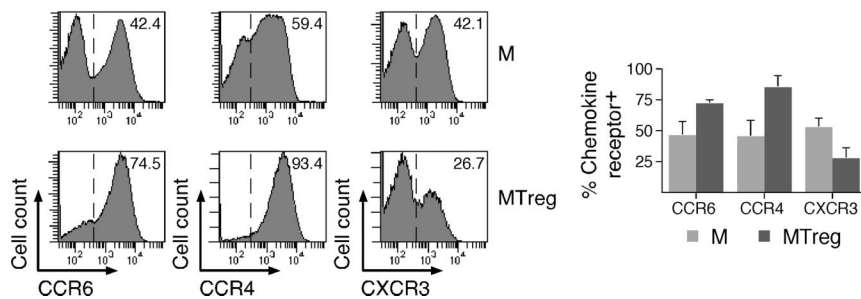


Fig. 4. CD4⁺ T cells were stained with antibodies specific for CD25 and CD45RA as well as for CCR4, CCR6, or CXCR3. Chemokine receptor expression was analyzed gated on CD25⁺CD45RA⁻ (M) and CD25⁺CD45RA⁺ (MTreg) cells. Histograms for 1 donor and data for 3 donors are shown (mean ± SD).

human T cells has been shown to inhibit IL-2 expression by competing with NFAT for binding to DNA (15, 27), consistent with the inability of Tregs to secrete IL-2 in response to T cell receptor (TCR)-mediated stimulation (9).

Our findings raise several questions. First, what factors/mechanisms are responsible for ROR γ t expression in Tregs? A likely candidate is TGF- β , which promotes the generation of human and murine Tregs and has clearly been shown to mediate, in association with IL-6, the generation of murine T_H17 cells (18), although its role in the generation of human T_H17 cells is debated (28–31). In line with this hypothesis, stimulation of CD4⁺ T cells with TGF- β has been shown to induce expression of both FOXP3 and ROR γ t (32, 33). It is, however, noteworthy that although NnTregs and MTregs express similar levels of FOXP3, expression of ROR γ t in MTregs is significantly higher than in NnTregs, which fail to produce significant levels of IL-17 *ex vivo*. Thus, expression of ROR γ t in Treg cells may be more difficult to achieve compared with expression of FOXP3, and only becomes significant at late differentiation stages.

Our finding that FOXP3⁺ MTregs secrete IL-17 *ex vivo* was unexpected, also taking into account recent studies reporting that FOXP3 inhibits IL-17 secretion by antagonizing ROR γ t (33, 34). However, the findings confirm recent data from Osorio et al. (25) that some murine Tregs can produce IL-17 while retaining FOXP3 expression.

Other important questions relate to the biological significance of ROR γ t expression and IL-17 production in Tregs. Because human T_H17 and IL-17-secreting Treg populations have partially common differentiation and migratory pathways, our findings support the concept that these 2 populations may not only

coexist in the same tissues, but may share some proinflammatory functions. IL-17-secreting Tregs present at mucosal sites may participate in early responses to infection by attracting mediators of innate immunity. Additionally, it must be taken into account that nuclear hormone receptors, such as ROR γ t, perform diverse functions in development and homeostasis by regulating cell growth, differentiation, and apoptosis. In addition to T_H17 cells, ROR γ t expression is found in double-positive thymocytes; in a subset of intestinal lamina propria T lymphocytes, most of which constitutively produce IL-17; in intestinal CD3⁻ lymphoid tissue-inducer cells, also producing IL-17 and clustered in crypt-patches that are precursor structures of lymphoid follicles (16, 35); as well as in subpopulations of NK cells and $\gamma\delta$ T cells (36–38). Thus, MTregs share ROR γ t expression and IL-17 production not only with proinflammatory T_H17 cells but also with other populations involved in innate immunity and lymphogenesis, strongly suggesting additional as yet undescribed roles of Tregs in these processes. Finally, because Tregs are believed to be autoreactive, their ability to secrete IL-17 after TCR-mediated stimulation could indicate a physiologic role for these cells in tissue homeostasis. Future studies addressing these issues will undoubtedly contribute to further elucidate the mechanisms regulating the balance between inflammation and tolerance under physiological conditions as well as in a variety of pathological conditions.

Materials and Methods

Samples, Cell Purification, and Sorting. Peripheral blood samples were obtained from the Etablissement Français du Sang Pays de la Loire (Nantes, France). Peripheral blood mononuclear cells (PBMCs) were isolated by density gradient sedimentation using LSM 1077 lymphocyte separation medium (PAA

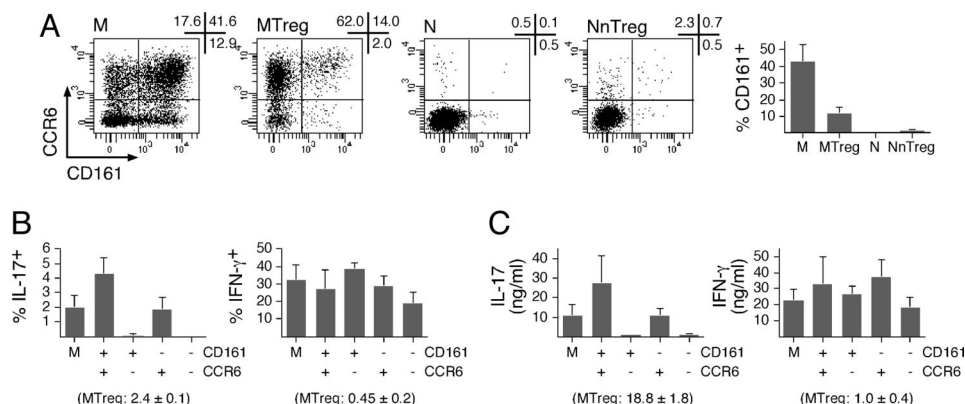


Fig. 5. Low proportions of MTregs express CD161. (A) CD4⁺ T cells were stained with antibodies specific for CD25, CD45RA, CCR6, and CD161. Dot plots from 1 donor are shown gated on the indicated population defined as in Fig. 2A. Data for 3 donors are shown (mean ± SD). (B and C) CD4⁺ T cells were stained with anti-CD25, anti-CD45RA, anti-CCR6, and anti-CD161 antibodies, and conventional memory cells (CD25⁻CD45RA⁻) were sorted by flow cytometry either as 1 population or separated into 4 populations according to the expression of CCR6 and CD161. Sorted populations were stimulated, and cytokine production was assessed as in Fig. 2B (B) and Fig. 2E (C) (mean ± SD, *n* = 2).

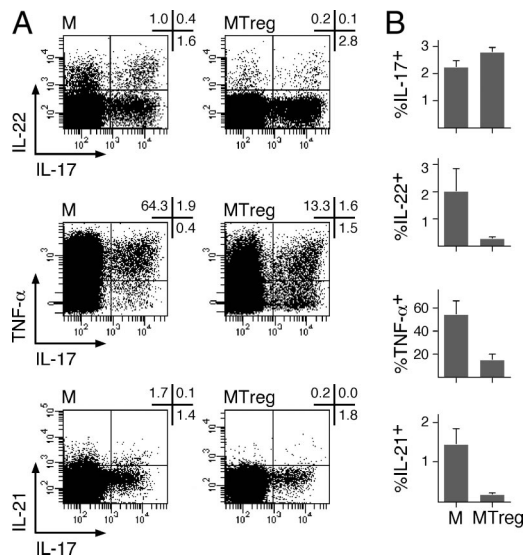


Fig. 6. Cytokine production by IL-17-producing MTregs. M and MTreg populations were sorted by flow cytometry as in Fig. 2A, and IL-17 and IL-22, TNF- α , or IL-21 production was determined by intracellular cytokine staining after 6 h of stimulation with PMA/ionomycin. Dot plots for 1 donor (A) and data for 3 donors (B; mean \pm SD) are shown.

Laboratories). CD4⁺ T cells were enriched by positive selection from PBMCs by magnetic cell sorting (Miltenyi Biotec). For flow cytometry cell sorting, enriched CD4⁺ T cells were stained with fluorescent-conjugated antibodies specific for CD8 (BD Biosciences), CD45RA (BD Biosciences), and CD25 (Beckman Coulter). After gating on CD8⁻ lymphocytes, cells were separated into memory (M; CD25⁻CD45RA⁻) and naive (N; CD25⁻CD45RA⁺) conventional CD4⁺ T cells and memory (MTregs; CD25⁺CD45RA⁻) and naive (NnTregs; CD25⁺CD45RA⁺) Tregs to high purity (>97%) using a FACSAria (BD Biosciences). A fraction of sorted populations was stained with antibodies specific for CD127 and FOXP3 (eBiosciences) and analyzed by flow cytometry. In some experiments, CD4⁺ T cells were stained with antibodies specific for CD8, CD45RA, CCR6 (BD Biosciences), CD161 (BD Biosciences), and CD25, and conventional memory CD4⁺ T cells were sorted into CD161⁺CCR6⁺, CD161⁺CCR6⁻, CD161⁻CCR6⁺ and CD161⁻CCR6⁻ populations.

Phenotypic Analysis and Assessment of Cytokine Production. Conventional CD4⁺ T cell and Treg, naive and memory populations were assessed pheno-

typically by staining of total CD4⁺ T cells with antibodies specific for CD45RA, CD25, and CD127 together with antibodies specific for FOXP3, CCR4 (BD Biosciences), CXCR3 (BD Biosciences), or CCR6 and CD161, as indicated, and analysis by flow cytometry using FACSAria or an LSR II device (BD Biosciences). Cytokine production by enriched total CD4⁺ T cells was assessed ex vivo by intracellular staining after stimulation with PMA (100 ng/mL; Sigma-Aldrich) and ionomycin (1 μ g/mL; Sigma-Aldrich) for 6 h. Brefeldin A (10 μ g/mL; Sigma-Aldrich) was added 1 h after the beginning of the incubation. Cells were then fixed, permeabilized, and stained with antibodies specific for FOXP3, IL-17 (eBiosciences), and IFN- γ (BD Biosciences), and were analyzed by flow cytometry. Cytokine production by sorted CD4⁺ T cell populations was assessed ex vivo after stimulation with PMA and ionomycin in a 6-h intracellular cytokine staining assay using antibodies specific for IL-17, IFN- γ , and IL-22 (R & D Systems), TNF- α (BD Biosciences), IL-21 (BD Biosciences), or ROR γ / γ t (eBiosciences), as indicated. Cytokine production by CD4⁺ T cell populations was also assessed ex vivo in a secretion assay. Briefly, sorted CD4⁺ T cell populations (5×10^5 per well) were stimulated in the absence or presence of PMA and ionomycin in round-bottom 96-well plates. After 24 h, secreted IL-17 and IFN- γ were measured in the culture supernatant by ELISA (R & D Systems and Invitrogen, respectively), and IL-17 and IFN- γ mRNA levels in cells were determined by real-time quantitative RT-PCR as detailed below.

Conventional and Real-Time Quantitative RT-PCR. RNA was prepared from ex vivo-sorted CD4⁺ T cell populations (5 to 10×10^5 cells) by using an RNeasy Mini Kit (Qiagen). cDNA synthesis was performed by using Promega Reverse Transcription System A3500 (Promega). Semiquantitative PCR was performed by using GoTaq Flexi DNA Polymerase (Promega). cDNA integrity was tested by amplification of β -actin mRNA in a 35-cycle PCR, and RORC mRNA expression was assessed by using the following primers: forward primer, 5'-TTTCCGAGGATGAGATTGC-3'; reverse primer: 5'-CTTCCACATGCTGGC-TACA-3'. Quantitative real-time PCR was performed with a TaqMan assay on an ABI 7000 system (Applied Biosystems) using Assays-on-Demand Gene Expression probes for *RORC* (Hs01076112.m1), *FOXP3* (Hs00203958.m1), *T-bet* (*Tbx21*; Hs00203436.m1), *GATA3* (Hs00231122.m1), *IL17* (*IL17A*; Hs99999082.m1), and *IFNG* (Hs99999041.m1) (Applied Biosystems). For control of input RNA, we used either Assays-on-Demand Gene Expression probes for *18s* rRNA (Hs99999901.s1) or Taqman probe (FAM-5'-AAGGTGAAGTCCG-GAGTCAACGGATTTG-3'-TAMRA) and primers (5'-CCACATCGCTCAGACAC-CAT-3' and 5'-CCAGGCGCCCAATACG-3') for *GAPDH*, as indicated. Relative mRNA expression was calculated as $2^{-(Ct \text{ control RNA} - Ct \text{ test RNA})}$.

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