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# IL-21 and TGF- $\beta$ are required for differentiation of human T<sub>H</sub>17 cells

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The recent discovery of CD4<sup>+</sup> T cells characterized by secretion of interleukin (IL)-17 (T<sub>H</sub>17 cells) and the naturally occurring regulatory FOXP3<sup>+</sup> CD4 T cell (nT<sub>reg</sub>) has had a major impact on our understanding of immune processes not readily explained by the T<sub>H</sub>1/T<sub>H</sub>2 paradigm. T<sub>H</sub>17 and nT<sub>reg</sub> cells have been implicated in the pathogenesis of human autoimmune diseases, including multiple sclerosis, rheumatoid arthritis, inflammatory bowel disease and psoriasis<sup>1,2</sup>. Our recent data and the work of others demonstrated that transforming growth factor-β (TGF-β) and IL-6 are responsible for the differentiation of naive mouse T cells into T<sub>H</sub>17 cells, and it has been proposed that IL-23 may have a critical role in stabilization of the T<sub>H</sub>17 phenotype<sup>3-5</sup>. A second pathway has been discovered in which a combination of TGF-β and IL-21 is capable of inducing differentiation of mouse T<sub>H</sub>17 cells in the absence of IL-6 (refs 6-8). However, TGF-β and IL-6 are not capable of differentiating human T<sub>H</sub>17 cells<sup>2,9</sup> and it has been suggested that TGF- $\beta$  may in fact suppress the generation of human  $T_H17$  cells<sup>10</sup>. Instead, it has been recently shown that the cytokines IL-1\beta, IL-6 and IL-23 are capable of driving IL-17 secretion in short-term CD4<sup>+</sup> T cell lines isolated from human peripheral blood<sup>11</sup>, although the factors required for differentiation of naive human CD4 to T<sub>H</sub>17 cells are still unknown. Here we confirm that whereas IL-1β and IL-6 induce IL-17A secretion from human central memory CD4<sup>+</sup> T cells, TGF-β and IL-21 uniquely promote the differentiation of human naive CD4<sup>+</sup> T cells into T<sub>H</sub>17 cells accompanied by expression of the transcription factor RORC2. These data will allow the investigation of this new population of T<sub>H</sub>17 cells in human inflammatory disease.

To better understand regulation of IL-17A secretion from human CD4 $^+$  T cells, we used a strategy that would allow us to evaluate the effects of various combinations of cytokine on expansion of  $T_{\rm H}17$  cells from memory T cells versus differentiation of naive CD4 $^+$  lymphocytes into  $T_{\rm H}17$  cells. Specifically, we used high-speed flow cytometry for sorting these two distinct populations of CD4 $^+$  T cells from the peripheral blood of healthy subjects: CD4 $^+$ CD25 $^-$ CD62L $^+$ CD45RA $^{\rm hi}$  cells highly enriched for naive T cells and CD4 $^+$ CD25 $^-$ CD62L $^+$ CD45RA $^-$  cells enriched for central memory T cells ( $T_{\rm CM}$ ; Fig. 1a). All cells enriched for a naive or a central memory phenotype expressed the chemokine receptor CCR7 (data not shown). These two T cell populations were then stimulated with plate-bound anti-CD3 and soluble anti-CD28 monoclonal antibodies for 7 days in serum-free medium containing different combinations of cytokines implicated in CD4 $^+$   $T_{\rm H}17$  cell differentiation.

As reported previously, the cytokine IL-1 $\beta$  induced the greatest amount of IL-17A secretion from T<sub>CM</sub> (Fig. 1b). The addition of IL-6 alone had little effect on induction of IL-17A, and when added with IL-1 $\beta$  had no additive or synergistic effect on IL-17A production.

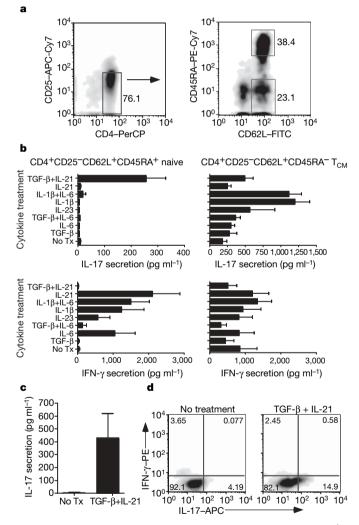


Figure 1 | TGF-β and IL-21 promote  $T_H17$  differentiation from naive CD4<sup>+</sup> T cells. a, CD4<sup>+</sup> T cells were sorted into populations enriched for naive or central memory T helper cells. b, IFN-γ and IL-17A secretion is shown from T cells stimulated for 7 days in the presence of the indicated cytokines. Standard deviation using T cells from three unrelated subjects is represented. TGF-β- and IL-21-induced IL-17 secretion is highly significant (P < 0.01). Tx, treatment. c, IL-17 secretion from naive T cells from six different donors is represented (mean  $\pm$  s.e.m.). d, Intracellular expression of IL-17 and IFN-γ from one of five experiments is shown.

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Addition of IL-23 was also able to modestly enhance IL-17A secretion from T<sub>CM</sub>. However, IL-1β alone or together with IL-6 failed to induce IL-17A secretion from naive CD4<sup>+</sup> T cells. In marked contrast, a combination of TGF-β and IL-21 was uniquely able to induce T<sub>H</sub>17 differentiation. Whereas IL-21, IL-1β or IL-6 induced significant amounts of interferon- $\gamma$  (IFN- $\gamma$ ) secretion from naive T cells, the addition of TGF-β with IL-21 suppressed IFN-γ secretion and induced differentiation of T<sub>H</sub>17 cells. Whereas we did observe variability in the extent of T<sub>H</sub>17 differentiation among unrelated healthy donors (Fig. 1c), we always observed induction of IL-17A after differentiation in the presence of TGF-β with IL-21. Intracytoplasmic staining demonstrated, in agreement with enzyme-linked immunosorbent assay (ELISA) results, that the combination of TGF-β and IL-21 differentiated CD4<sup>+</sup> T cells that secreted only IL-17A and no IFN-γ (Fig. 1d). A recent study<sup>12</sup> has demonstrated that the pathogenicity of mouse IL-17-secreting T cells is influenced by whether they also secrete IL-10. Using intracytoplasmic staining, we failed to observe any IL-10 when naive CD4<sup>+</sup> T cells were differentiated in the presence of TGF-β and IL-21 (data not shown).

RORC2 is the human homologue of mouse RORγt—a transcription factor critical for the differentiation of mouse IL-17-secreting T cells. Thus, to understand the molecular mechanisms involved in differentiation of human T<sub>H</sub>17 cells, we used quantitative PCR with reverse transcription (RT-PCR) to evaluate messenger RNA levels of RORC2 and other molecules implicated in T<sub>H</sub>17 differentiation. The combination of TGF-β and IL-21 induced high levels of RORC2 (Fig. 2a), consistent with their ability to induce IL-17A secretion from naive human CD4<sup>+</sup> T cells. It was of particular interest that the combination of TGF-β and IL-6 that induces T<sub>H</sub>17 differentiation in mouse T cells also induced expression of RORC2 in naive human CD4 cells. Because this combination of cytokines did not, however, induce IL-17A secretion, these data indicate that expression of RORC2 in humans is not in itself sufficient to induce IL-17 production, and that another as yet unidentified transcription factor in combination with RORC2, perhaps the human homologue of mouse

 $ROR\alpha$  (ref. 13), may be required to induce IL-17A-secreting  $T_H17$  cells.

T-bet (also known as TBX21) is the master regulator for T<sub>H</sub>1 cells secreting IFN-γ, whereas GATA3 induces T<sub>H</sub>2, IL-4 secreting CD4 cells. Messenger RNA expression levels of Tbet were highly concordant with amounts of IFN-γ secretion and were consistent with our findings that whereas TGF-β and IL-21 induce T<sub>H</sub>17 cell differentiation with RORC2 expression, TGF-β suppresses the induction of T-bet by IL-21. Similarly, there was no induction of GATA3 with TGF-β and IL-21. The cytokines IL-6, IL-21 and IL-1β but not TGF-β induced IL-23 receptor upregulation in stimulated naive CD4<sup>+</sup> T cells. We also examined the expression of the T<sub>reg</sub> transcription factor FOXP3. As has been previously reported in both mouse and human systems, FOXP3 was induced by TGF-β1. Induction of FOXP3 was inhibited by IL-6 and to a greater extent by IL-21—transcription factors that induce RORC2. Thus, although the induction of RORC2 and FOXP3 transcription factors was highly similar between mouse and human naive CD4 cells, the induction of IL-17A by IL-6 in combination with TGF-β is discordant between the species.

We and others have shown previously that IL-21 secreted by mouse CD4<sup>+</sup> T cells can induce the secretion of IL-21 in an autocrine amplification loop<sup>6-8,14</sup>. Thus, we examined whether human IL-21 similarly induced IL-21 secretion from naive CD4<sup>+</sup> T cells; we also evaluated the effects of a combination of TGF-β with IL-21 and IL-1 $\beta$ , given the ability of these cytokines to induce IL-17 from naive and central memory CD4<sup>+</sup> T cells. Consistent with results in mice, IL-21 upregulated IL-21, although IL-1β induced even greater amounts of IL21 mRNA (Fig. 2b). In contrast to what has been observed in mice, IL-21 also increased IL22 mRNA levels in naive CD4<sup>+</sup> T cells in the absence of any exogenous IL-23. However, TGF-β suppressed the expression of IL21 and IL22 mRNA induced by IL-21 (Fig. 2b). These data further highlight similar yet subtle differences between human and mouse CD4<sup>+</sup> T cells, because whereas IL-21 induces IL21 and IL22, differentiation to T<sub>H</sub>17 cells with TGF- $\beta$  inhibits the expression of these cytokines.

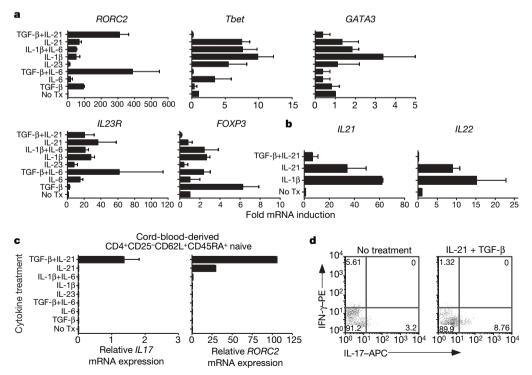


Figure 2 | TGF-β and IL-21 induce RORC2 in naive CD4<sup>+</sup> T cells. a, mRNA expression levels (fold-induction relative to T cells without exogenous cytokines) of *RORC2*, *Tbet*, *GATA3*, *IL23R* and *FOXP3* are shown after naive T cells were differentiated as indicated. b, Fold-induction  $\pm$  s.e.m. (n = 3) of *IL21* and *IL22* are represented. c, Mean expression  $\pm$  s.e.m. of *IL17A* and

*RORC2* are shown (n=3) for naive T cells obtained from cord blood. **d**, Shown is intracytoplasmic staining of IL-17 and IFN- $\gamma$  from cord blood naive T cells after 7 days of differentiation. Similar results were seen in another independent assay.

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To confirm the unique function of TGF- $\beta$  and IL-21 in the differentiation of  $T_H17$  cells from naive human CD4 $^+$  T cells, we sorted CD4 $^+$ CD25 $^-$ CD62L $^+$ CD45RA $^{hi}$  cells from human cord blood. As expected, a higher proportion of CD4 cells in the cord blood exhibited this naive phenotype relative to peripheral blood obtained from healthy adult subjects (data not shown). TGF- $\beta$  and IL-21 induced the upregulation of *IL17A* and *RORC2* mRNA (Fig. 2c). Although IL-21 alone modestly induced *RORC2*, only TGF- $\beta$  and IL-21 were able to induce *IL17A* mRNA. When given a very strong *in vitro* stimulus, naive CD4 $^+$  T cells sorted from cord blood secreted IL-17A protein (Fig. 2d). These data further indicate that TGF- $\beta$  and IL-21 are critical in the differentiation of both human and mouse  $T_H17$  cells.

Collectively, our data refine and extend our understanding of the regulation of IL-17A secretion from human CD4<sup>+</sup> T cells and define the conditions required for human T<sub>H</sub>17 cell differentiation. We confirm recent reports that IL-1β together with IL-6 (ref. 9) or IL-23 (ref. 2) can induce IL-17A secretion, which is most apparent in the human memory CD4<sup>+</sup> T cell subset. A combination of TGF-β plus IL-21 is required for the differentiation of T<sub>H</sub>17 cells from naive T cells. It is important to note that to observe T<sub>H</sub>17 differentiation from naive human  $\mathrm{CD4}^+$  T cells, appropriate amounts of both IL-21 and TGF-β are needed: addition of IL-21 alone must be sufficient to upregulate IFN-γ secretion, and the amount of TGF-β added must inhibit IL-21-induced IFN-γ secretion (see Methods for details). T<sub>H</sub>17 cells differentiated under these conditions are also notable for secretion of IL-17A in the absence of IFN-γ. In summary, our data together with previous reports in humans suggest that IL-1β and IL-6 induced during the early stages of an inflammatory response may act on memory T cells to promote IL-17 and IL-21 secretion, with induced IL-21 able to synergize with TGF-β to promote differentiation of T<sub>H</sub>17 cells from naive CD4<sup>+</sup> T cells. IL-23 may serve to expand or stabilize the phenotype of previously differentiated T<sub>H</sub>17 cells. These experiments allow for the characterization of human inflammatory T<sub>H</sub>17 responses associated with infection and autoimmune diseases.

#### **METHODS SUMMARY**

Cell sorting. PBMCs were obtained from the peripheral blood of healthy subjects or from cord blood (AllCells) in compliance with institutional review board (IRB) protocols. CD4<sup>+</sup> T cells were subsequently isolated by negative selection using magnetic beads (Miltenyi Biotech). Naive (CD25<sup>-</sup>CD62L<sup>+</sup>CD45RA<sup>hi</sup>) and central memory (CD25<sup>-</sup>CD62L<sup>+</sup>CD45RA<sup>-</sup>) CD4<sup>+</sup> T cells were obtained by staining with the following antibodies: CD62L–FITC, CD4–PerCP, CD45RA–PE–Cy7, CD25–APC–Cy7 (BD Pharmingen) and were sorted on a FACS Aria (BD Biosciences).

Differentiation assays. Naive or central memory CD4 $^+$  T cells were stimulated with plate-bound anti-CD3 and soluble CD28 monoclonal antibodies in serum-free X-VIVO15 medium (Biowhittaker) and cytokines (IL-6, 25 ng ml $^{-1}$ ; TGF-β1, 5 ng ml $^{-1}$ ; IL-1β, 12.5 ng ml $^{-1}$ ; IL-21, 25 ng ml $^{-1}$ ; IL-23, 25 ng ml $^{-1}$ ) for a period of 7 days, at which point supernatants were collected and tested by ELISA for IFN-γ (BD Biosciences) or IL-17A (eBioscience) using paired

antibodies. We have observed that concentrations of TGF- $\beta$  ranging from 0.1 ng ml $^{-1}$  to 10 ng ml $^{-1}$  in the presence of IL-21 promote  $T_{\rm H}17$  differentiation, whereas 50 ng ml $^{-1}$  TGF- $\beta$  suppresses differentiation. Intracytoplasmic staining was performed using standard methodologies and anti-IL-17–APC (R&D Systems) and anti-IFN- $\gamma$ -PE or anti-IL-10–PE (BD Biosciences) antibodies. **Real-time PCR.** All primers and probes were obtained from Applied Biosystems and used according to standard methodologies.

**Full Methods** and any associated references are available in the online version of the paper at www.nature.com/nature.

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**Supplementary Information** is linked to the online version of the paper at www.nature.com/nature.

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#### **METHODS**

Differentiation assays. Plates (96-well U-bottom) were coated with 1.5 μg ml<sup>-1</sup> anti-CD3 monoclonal antibody (eBioscience, clone OKT3) in a volume of 50 µl of PBS and were incubated overnight (16 h) at 4 °C. For T cells isolated from cord blood, a 96-well plate was pre-coated overnight at 4 °C with 3 µg ml<sup>-1</sup> anti-CD3 monoclonal antibody (UCHT1, BD Biosciences). Antibody solution was then removed, plates were rinsed once with serum-free X-VIVO 15 medium (Biowhittaker), and naive or central memory CD4<sup>+</sup> T cells  $(5 \times 10^4 \text{ per well})$ were stimulated in serum-free X-VIVO15 medium with soluble CD28 (BD Biosciences, clone 28.2) monoclonal antibody (1.0 µg ml<sup>-1</sup>) and cytokines (IL-6, 25 ng ml<sup>-1</sup>; TGF-β1, 5 ng ml<sup>-1</sup>; IL-1β, 12.5 ng ml<sup>-1</sup>; IL-21, 25 ng ml<sup>-1</sup>; IL-23, 25 ng ml<sup>-1</sup>) for a period of 7 days in the absence of IL-2. The cytokines IL-6, IL-1β, IL-23 and TGFβ1 (catalogue number 240-B-002) were all obtained from R&D systems. The TGFβ1 was not acid-treated before addition. IL-21 was purchased from Cell Sciences (catalogue number CRI 172A). Supernatants were collected and tested by ELISA for IFN-γ (Endogen) or IL-17A (human IL-17A ELISA kit from eBioscience or human IL-17 duoset from R&D systems) using paired antibodies. We have observed that concentrations of TGF-\(\beta\)1 ranging from 0.1 ng ml<sup>-1</sup> to 10 ng ml<sup>-1</sup> in the presence of IL-21 promote T<sub>H</sub>17 differentiation, with lower doses of TGF\$\beta\$1 associated with induction of a higher proportion of IL-17-producing cells than higher doses; use of 50 ng ml TGF-β1 suppresses differentiation and IL-17 secretion. Intracytoplasmic staining was performed using standard methodologies and anti-IL-17-APC (R&D Systems) and anti-IFN-γ-PE or anti-IL-10-PE (BD Biosciences) antibodies. Cells were stimulated with phorbol 12-myristate 13-acetate (PMA; 10 ng ml<sup>-1</sup>), ionomycin (0.5 μg ml<sup>-1</sup>) and Golgistop for 5 h at 37 °C before intracellular staining. Real-time PCR. After removing supernatants from wells in differentiation assays, 250 µl per well of lysis buffer was added, at which point RNA was isolated using a RNeasy mini kit (Qiagen). Total RNA was converted to complementary DNA using Taqman Reverse transcription reagents (Applied Biosystems). Quantitative PCR was performed using a 7500 Fast Real-time PCR system (Applied Biosystems). All primers and probes were obtained from Applied Biosystems and used according to standard methodologies.