

Increased Circulating Pro-Inflammatory Cytokines and Th17 Lymphocytes in Hashimoto's Thyroiditis

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Context: Th17 lymphocytes play an important role in different chronic inflammatory and autoimmune conditions.

Aim: The aim of the study was to explore the status of Th17 cells in patients with autoimmune thyroid diseases (AITD).

Design: We assessed the serum levels and *in vitro* synthesis of IL-17 and IL-22 and of different cytokines (IL-6, IL-15, and IL-23) involved in the differentiation of Th17 cells in the peripheral blood and thyroid glands of 26 patients with AITD, eight with Graves' disease, and 18 with Hashimoto's thyroiditis (HT) as well as 10 healthy controls.

Results: We found enhanced levels of T cells synthesizing IL-17 and IL-22 in the peripheral blood from AITD patients, mainly in those with HT. In addition, a stronger expression of IL-17 and IL-22 and an enhanced number of IL-23R⁺ cells was detected in thyroid glands from HT patients compared with Graves' disease or controls. Furthermore, increased concentrations of IL-6 and IL-15 were detected in sera from HT patients, whereas serum levels of IL-23 tended to be higher in these patients. Finally, an enhanced *in vitro* differentiation of T lymphocytes into Th17 cells induced by IL-23/IL-6 was observed in AITD patients. Accordingly, a strong induction of RORC2 gene was detected in lymphocytes from HT patients when stimulated with IL-23.

Conclusion: Our results indicate that there is an increased differentiation of Th17 lymphocytes and an enhanced synthesis of Th17 cytokines in AITD, mainly in HT. These phenomena may have an important role in the pathogenesis of thyroid autoimmunity. (*J Clin Endocrinol Metab* 95: 953–962, 2010)

CD4⁺ T helper lymphocytes play a key role in the pathogenesis of inflammatory and autoimmune diseases via the production of distinctive sets of cytokines (1). On the basis of their pattern of cytokine synthesis, CD4⁺ T helper cells were originally classified into Th1 and Th2 lymphocytes, which are involved in the cellular and humoral immune responses, respectively (2). A third subset of CD4⁺ cells (Th3 lymphocytes) mainly synthesize TGF- β and are considered as regulatory cells (3). Additional lymphocyte subsets have been identified, including

Th9 (4) and Th17 cells (5–7). The latter lymphocytes are characterized by the synthesis of IL-17A, IL-17F, IL-21, and IL-22 (8) and express the transcription factors: retinoic acid receptor-related orphan receptor C2 (RORC2) and signal transducers and activators of transcription-3 (Stat-3), which have a critical role in their development (9, 10). Th17 cells are involved in the pathogenesis of chronic inflammatory conditions (11), including asthma (12), rheumatoid arthritis (13), psoriasis (14), and systemic lupus erythematosus (15).

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Abbreviations: Ab, Antibody; AITD, autoimmune thyroid diseases; FT₄, free T₄; GD, Graves' disease; HT, Hashimoto's thyroiditis; IFN, interferon; mAb, monoclonal antibody; PBMC, peripheral blood mononuclear cell; TFC, thyroidal follicular cell; Tg, thyroglobulin; TMC, thyroid mononuclear cell; TPO, antithyroperoxidase; Treg, T regulatory; TSH, thyroid-stimulating hormone; TSHR, thyroid-stimulating hormone receptor.

Several cytokines are involved in the differentiation of human Th17 cells, mainly IL-21 and IL-23 as well as several proinflammatory cytokines, including IL-1 β and TNF- α (16–19). On the contrary, other cytokines exert a negative regulatory role on Th17 cell development, including interferon (IFN)- γ , and IL-27 (20). On the other hand, TGF- β seems to have a dual role, inhibiting the generation of Th17 cells in the absence of other cytokines and promoting its differentiation in combination with IL-23, IL-6 and, likely, IL-1 β and TNF- α (21, 22). The proinflammatory cytokine IL-15 is also able to induce the synthesis of IL-17 by Th17 cells (23). Th17 cells synthesize IL-17A (also called IL-17) that exhibits a strong proinflammatory activity, inducing the synthesis of chemokines and proinflammatory cytokines (8, 11, 24, 25). Accordingly, the synthesis of IL-17 is increased in different inflammatory autoimmune conditions such as collagen-induced arthritis (26), rheumatoid arthritis (27), systemic sclerosis (28), psoriasis (29), and systemic lupus erythematosus (30). On the other hand, IL-22, mainly synthesized by Th7 cells, also has an important role in the pathogenesis of different inflammatory conditions (31).

Human autoimmune thyroid diseases (AITD), mainly including Hashimoto's thyroiditis (HT) and Graves' disease (GD), are characterized by reactivity to self-thyroid antigens (32). It has been proposed that the interaction of thyroidal follicular cells (TFCs), antigen-presenting cells, and autoreactive T cells results in an autoimmune response against thyroid antigens, which can be mediated by Th1 or Th2 lymphocytes (33). In the case of a Th1 response, a heavy/strong inflammatory infiltrate with destruction of thyroid gland occurs (HT), whereas in the case of a Th2 response, stimulatory anti-TSH receptor antibodies, an enhanced function of the gland and a variable degree of inflammation are observed (GD) (33–35). Autoreactive T lymphocytes and TFCs are able to synthesize different chemokines and cytokines, which favor the expansion of the intrathyroidal lymphocyte pool, exacerbating the autoimmune process (36–38).

We previously described that patients with AITD show an altered proportion and a defective function of T regulatory (Treg) cells in both the peripheral blood and thyroid gland, (39). Because there is a reciprocal negative regulation between Treg and Th17 lymphocytes, we hypothesized that patients with AITD show an expansion of Th17 cells and that these cells are involved in the pathogenesis of this condition. In this work, we found enhanced levels of Th17 cells and Th17 cytokines in patients with AITD, mainly those with HT. In addition, an increased capability of peripheral blood lymphocytes to differentiate *in vitro* into Th17 cells, with a strong expression of the RORC2 gene, was observed in these patients. These data suggest

TABLE 1. Clinical features of AITD patients included in the study (PBMC assays)

	HT	GD	Range
n	13	5	
Gender (M/F)	2/11	2/3	
Age (yr)	45.36 \pm 13.20	45.80 \pm 16.63	
TSH (aeU/ml)	9.75 \pm 5.49	0.33 \pm 0.63	0.3–4.94
Free T ₄ (ng/dl)	0.83 \pm 0.13	1.52 \pm 1.30	0.7–1.48
TPO-Ab (U/ml)	584.09 \pm 498.00	543.0 \pm 514.64	<344
Tg-Ab (U/ml)	1056.3 \pm 2543.3	<0.02	<100
TSHR-Ab (U/liter)	0.054 \pm 0.052	13.71 \pm 16.6	<2

Data correspond to the arithmetic mean \pm sd. M, Male; F, female.

that Th17 cells may be involved in the pathogenesis of AITD.

Patients and Methods

Individuals and samples

Eighteen AITD patients (32) (13 with HT and five with GD) were included in the study. Three of the five patients with GD had ophthalmopathy. Peripheral blood samples were obtained from all patients, and complete thyroid work-up was performed in them. Main clinical data of these patients are shown in Table 1. Serum free T₄ (FT₄), TSH levels, and titers of anti-TSH receptor (TSHR), anti-thyroglobulin (Tg) and antithyroperoxidase (TPO) antibodies (Abs) were measured as previously described (40). Ten age- and sex-matched healthy subjects were included as controls.

Thyroid glands were obtained from five patients with HT and three with GD who had relapsed after antithyroid drug treatment and were euthyroid under carbimazole therapy at the time of surgery; two of these patients had ophthalmopathy. In addition, normal thyroid tissue was obtained from patients undergoing parathyroidectomy. These patients did not have evidence of thyroid or autoimmune disease (Table 2). Thyroid specimens were frozen, embedded in OCT medium (Ames, Miles Laboratories, Elkhart, IN) and stored at -80°C . All samples were taken in accordance with the regulations and approval of the Institutional Review Board of Hospital Universitario de la Princesa. In all cases, a written informed consent was obtained, and the local hospital bioethical committee approved this study.

Immunohistochemistry

Thyroid tissue sections (5.0 μm thick) were mounted on glass slides (Dako Cytomation, Copenhagen, Denmark) and stained by an immunoperoxidase method as previously described (41). Briefly, endogenous peroxidase was inactivated, and Fc receptors blocked with diluted human AB serum. Then sections were incubated with anti-IL-17 (R&D Systems, Wiesbaden-Nordenstadt, Germany), anti-IL-22 (Santa Cruz Biotechnology Inc., Heidelberg, Germany), anti-IL-18R (R&D Systems), goat anti-human polyclonal Ab, or an anti-IL-23R monoclonal antibody (mAb; R&D Systems). Next, sections were incubated with the proper horseradish peroxidase-conjugated secondary antibody (Envision; Dako Cytomation), washed, and developed with 3,3'-diaminobenzidine. Finally, sections were counterstained with Carazzi's hematoxylin and evaluated by at least two independent investigators.

TABLE 2. Clinical features of AITD patients included in the study (thyroid gland assays)

	Normal	HT	GD	Range
n	3	5	3	
Gender (M/F)	0/3	0/5	0/3	
Age (yr)	53.0 ± 10.8	44.0 ± 9.82	40.0 ± 15.7	
TSH (aeU/ml)	0.9 ± 1.1	4.5 ± 4.1	0.3 ± 0.7	0.3–4.94
Free T ₄ (ng/dl)	0.9 ± 0.1	0.9 ± 0.11	1.3 ± 0.3	0.7–1.48
TPO-Ab (U/ml)	Negative	296.50 ± 336.76	215.78 ± 296.45	<344
Tg-Ab (U/ml)	Negative	2168.8 ± 2821.8	<0.02	<100
TSHR-Ab (U/liter)	Negative	Negative	15.3 ± 17.2	<2

Data correspond to the arithmetic mean ± SD. M, Male; F, female.

Cell isolation

Peripheral blood mononuclear cells (PBMCs) were isolated by density-gradient centrifugation over Ficoll-Hypaque cushions (Biochrom AG, Berlin, Germany). Cell viability (trypan blue dye exclusion) was always greater than 95%.

To isolate thyroid mononuclear cells (TMCs), thyroid specimens were minced and then digested with collagenase (1.0 mg/ml; Roche Molecular Biochemicals, Roche, Mannheim, Germany) in Hanks' balanced-salt solution (BioWhittaker, Lonza, Verviers, Belgium) for 1 h at 37 C and passed through a steel mesh. Then cells were isolated by density-gradient centrifugation over Ficoll-Hypaque cushions, washed twice with PBS, and resuspended in complete RPMI 1640 culture medium (Life Technologies, Inc., Gaithersburg, MD). Cell viability was always higher than 95%.

In vitro stimulation of T cells

For the detection of Th17 lymphocytes, PBMCs or TMCs were incubated in complete RPMI 1640 culture medium in the presence of 50 ng/ml phorbol myristate acetate (PMA; Sigma-Aldrich St. Louis, MO) and 1.0 μg/ml ionomycin (Sigma-Aldrich) or 100 ng/ml recombinant human IL-15 (PeproTech EC Ltd., London, UK) for 5 and 6 h, respectively, and with the addition of 5.0 μg/ml of Brefeldin A (Sigma-Aldrich). Then IL-17⁺ cells were detected by flow cytometry, as described below.

PBMCs and TMCs were also cultured for 5 d with plate-bound anti-CD3 (T3b; 1.0 μg/ml) and anti-CD28 (10 μg/ml; Immunotech, Marseille, France) mAbs in the presence of anti-IFN-γ (5.0 μg/ml) and anti-IL-4 (5.0 μg/ml) blocking mAbs (R&D Systems), and different combinations of recombinant human IL-23 (20 ng/ml; R&D Systems), TGF-β (10 ng/ml; Peprotech), or IL-6 (20 ng/ml; Peprotech). Then cells were incubated with PMA and ionomycin, for 4 h in the presence of 5.0 μg/ml Brefeldin A (Sigma-Aldrich).

Flow cytometry analysis

Cells were stimulated with PMA/ionomycin, washed, and immunostained with a CD3-Peridinin Chlorophyll Protein Complex (PerCP) and a CD8-fluorescein isothiocyanate mAb (BD PharMingen, San Diego, CA), fixed, and permeabilized and then incubated with a Phycoerythrin conjugated anti-IL-22 (R&D Systems) and an Alexa fluor 647-conjugated anti-IL-17 mAb (eBioscience, San Diego, CA). Isotype-matched Ab controls (BD PharMingen) were used in all procedures. Cells (1–2 × 10⁵) were analyzed with a FACSCalibur flow cytometer (Becton Dickinson, San Jose, CA), and results were expressed as the percent of CD3⁺CD8⁻ cells expressing IL-17 or IL-22.

RNA isolation and real-time RT-PCR

Total RNA was isolated with the Ultraspec kit of Biotech Laboratories (Houston, TX). Reverse transcription was performed with random hexamers and the Moloney murine leukemia virus reverse transcriptase (Promega Biotech Ibérica, S.L., Madrid, Spain) in a GeneAmp PCR system 9700 thermal cycler (Perkin Elmer, Waltham, MA). Real-time RT-PCR was performed by duplicate using the StepOnePlus real-time PCR system using Power SYBR Green PCR master mix (Applied Biosystems, Foster City, CA) and using the following primers: IL-17, sense, 5'-CCA CCT CAC CTT GGA AT-3'; antisense, 5'-GGA CCA GGA TCT CTT GCT-3', IL-22, sense, 5'-TGA TAA CAA CAC AGA CGT TCG-3', antisense, 5'-CTG CAT ATA AGG CTG GAA CC-3'. As internal control, we used Histone H3, with the following primers: sense, 5'-AAA GCCGCT CGC AAG AGT GCG-3'; antisense, 5'-ACT TGC CTC CTG CAA AGC AC-3'. Gene-specific amplification was confirmed by a single peak in melting curve analysis. Data were analyzed with the StepOne software version 2.0 (Applied Biosystems). In the case of RORC2, cDNA products were amplified by PCR with the following primers: sense, 5'-GCT AGG TGC AGA GCT TCA GG-3', antisense, 5'-GGT GAT AAC CCC GTA GTG GA-3'. Amplification cycles (35) were as follows: 30 sec at 95 C, 30 sec at 57 C, and 45 sec at 72 C. Next, samples were incubated at 72 C for 7 min and cooled to 4 C. Finally, amplicons were electrophoresed on 2% agarose gel containing ethidium bromide (Bio-Rad) and visualized with an UV transilluminator.

Cytokine quantification

Levels of IL-6, IL-15, IL-17, IL-22, and IL-23 were determined by ELISA (ImmunoTools GmbH, Friesoythe, Germany; and R&D Systems) using a Tecan Sunrise ELISA reader (Tecan, Spain). All determinations were performed by duplicate and the lower detection limits for IL-6, IL-15, IL-17, IL-22, and IL-23 were 0.70, 3.9, 15.0, 2.7, and 6.8 pg/ml, respectively.

To quantify the *in vitro* synthesis of cytokines (IL-17 and IL-22), PBMCs or TMCs were stimulated with anti-CD3/CD28 mAbs and/or PMA/ionomycin or with 100 ng/ml of recombinant human IL-15 in the presence or absence of blocking anti-IFN-γ/IL-4 mAbs and recombinant cytokines (IL-23, TGF-β, IL-6), as described above. At the end of cell culture, cytokine concentration in cell-free supernatants was determined by ELISA.

Statistical analysis

Data are showed as the arithmetic mean ± SD or mean ± SEM. For continuous variables, normality was assessed by the Kolmogorov-Smirnov test. Data were analyzed using both parametric and non-

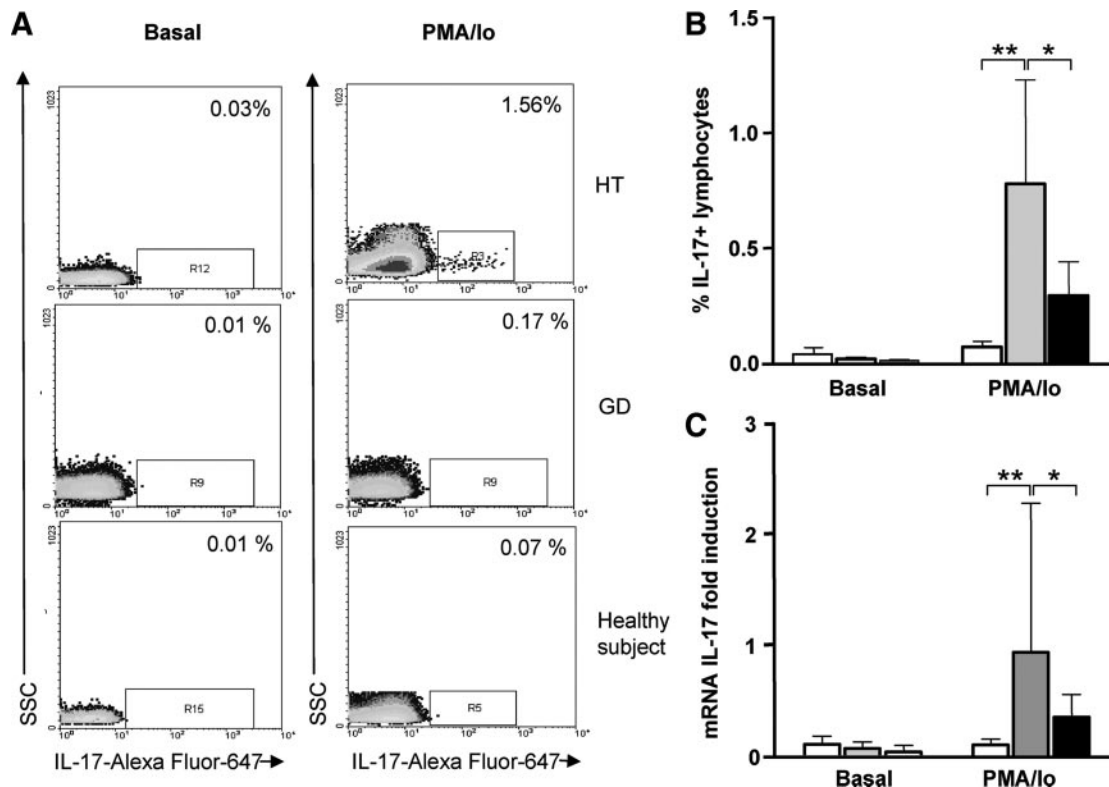


FIG. 1. IL-17 expressing cells in the peripheral blood from AITD patients. PBMCs from HT, GD, and healthy controls were incubated with PMA/ionomycin, stained for intracellular IL-17 and analyzed by flow cytometry. A, Representative dot plots from HT and GD patients and a control subject are shown. Numbers correspond to the percentage of CD3⁺CD8⁻IL-17⁺ cells. B, Percentages of IL-17⁺ cells in the peripheral blood of HT and GD patients, and healthy controls. Data correspond to the arithmetic mean \pm SEM. White bars correspond to healthy controls, gray bars to HT, and black bars to GD. C, Levels of IL-17 mRNA in PBMCs from AITD patients and controls pretreated or not with PMA/ionomycin. IL-17 mRNA was detected by real-time PCR analysis. Data correspond to the arithmetic mean \pm SEM of mRNA fold induction. White bars correspond to healthy controls, gray bars to HT, and black bars to GD. *, $P < 0.05$; **, $P < 0.01$. No significant differences were detected between GD patients and controls.

parametric tests. Mean group values were compared by one-way ANOVA and nonparametric ANOVA (Kruskal-Wallis test), and *post hoc* comparisons were carried out using the Bonferroni's and Dunn's test, respectively. Paired *t* test and Wilcoxon sum rank test were used to analyze two related samples. Analyses were performed using SPSS version 16.0 for Windows (SPSS Inc., Chicago, IL) and GraphPad Prism software version 5.0 (San Diego, CA).

Results

Th17 cells in the PBMCs from AITD patients

We first analyzed by flow cytometry the proportion of CD3⁺CD8⁻IL-17⁺ lymphocytes in PBMCs from AITD patients (Fig. 1A). An increased percentage of Th17 cells was observed in HT patients compared with GD patients or healthy subjects (Fig. 1B). In addition, the percentage of IL-17⁺ cells tended to be higher in GD patients compared with healthy subjects (Fig. 1B). However, no apparent differences were found between GD patients with and without ophthalmopathy (data not shown).

Real-time RT-PCR analysis showed enhanced levels of IL-17 mRNA in the PBMCs from HT patients, with lower expression in GD patients and healthy subjects (Fig. 1C

and data not shown). However, IL-17 was not detected by ELISA in sera from any individual included in the study (data not shown). In addition, a very low number of IL-17⁺ cells was detected in nonstimulated freshly isolated PBMCs from the three groups studied, with no significant differences among them (Fig. 1, A and B). Similar results were observed when the IL-17 mRNA was detected by real-time PCR (Fig. 1C).

In agreement with the above data, we found a modest but significant increased percentage of CD3⁺CD8⁻IL-22⁺ T cells in the PBMCs from HT patients compared with GD or control individuals ($P < 0.05$ in both cases, Fig. 2, A and B). Accordingly, serum levels of IL-22 were significantly higher in HT patients compared with both GD patients and healthy volunteers ($P < 0.05$ in both cases, Fig. 2, A and C). However, additional flow cytometry analysis revealed that in most individuals only a small percentage of IL-17⁺ cells also synthesized IL-22 and vice versa (data not shown).

Th17 lymphocytes in thyroid glands from AITD patients

Immunohistochemical analysis of thyroid sections revealed a variable staining for IL-17 and IL-22, with a

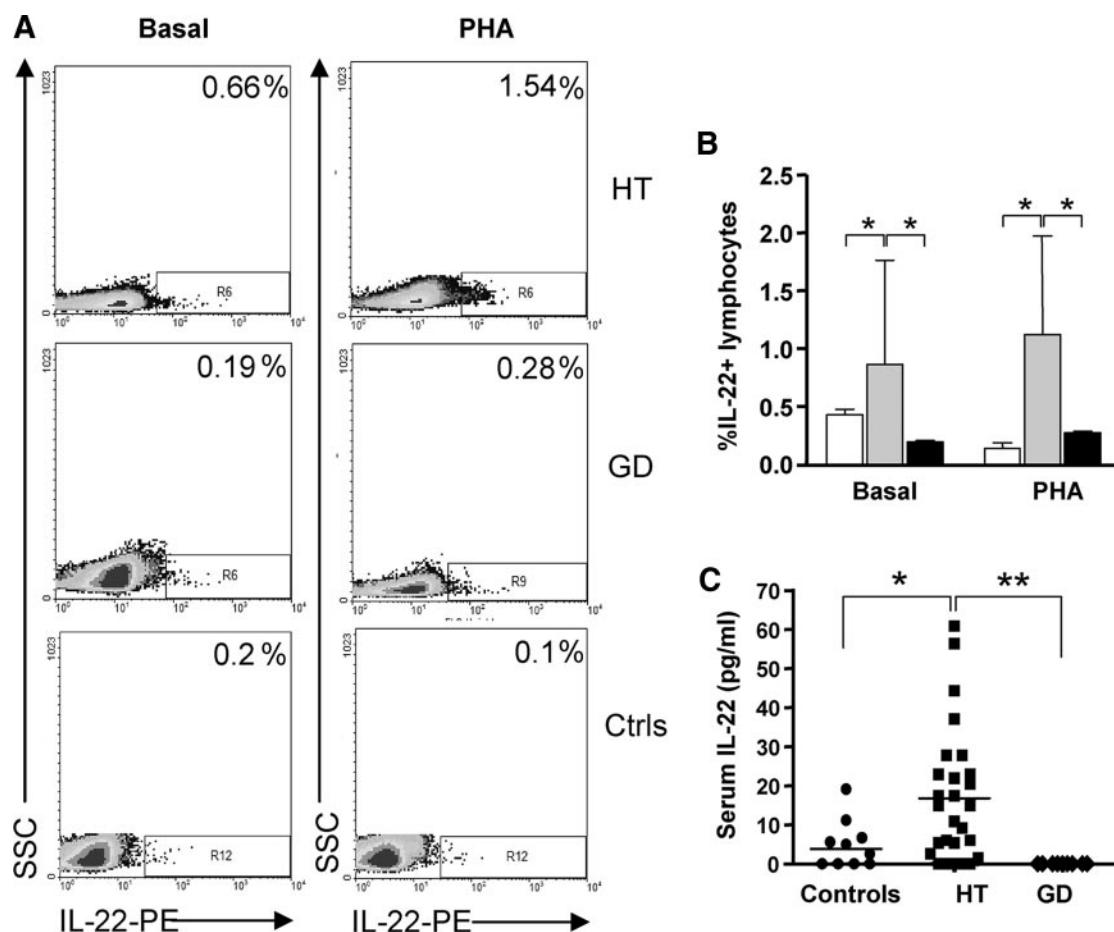


FIG. 2. IL-22 expressing cells in the peripheral blood from AITD patients. PBMCs from HT, GD, and healthy controls were incubated with PMA/ionomycin, stained for intracellular IL-22 and analyzed by flow cytometry. A, Representative dot plots from HT and GD patients and a control subject. Numbers correspond to the percentage of CD3⁺CD8⁻IL-22⁺ cells. B, Data of the percent of IL-22⁺ cells in the peripheral blood of HT and GD patients, and healthy controls are represented as the arithmetic mean \pm SEM. White bars correspond to healthy controls, gray bars to HT, and black bars to GD. C, Serum levels of IL-22 were determined by ELISA in samples from AITD patients and healthy controls. *, $P < 0.05$; **, $P < 0.01$. No significant differences were detected between GD patients and controls.

stronger expression of these cytokines in patients with HT with active lesions, compared with GD and healthy controls (Fig. 3A). As expected, in HT glands IL-17⁺ and IL-22⁺ cells were mainly detected into the inflammatory cell infiltrate. IL-17 and IL-22 were almost undetectable in thyroid glands from GD patients and controls (Fig. 3A and data not shown). Accordingly, IL-17 and IL-22 mRNAs were detected in TMCs from HT but not GD patients or healthy controls (Fig. 3, B and C, and data not shown). In this regard, IL-15 induced a modest but significant differentiation of Th17 cells in the PBMCs from HT patients, whereas a similar effect was exerted by IL-23, alone or in combination with TGF- β (Fig. 3B). In addition, IL-6 alone showed a greater effect on IL-17 cell differentiation compared with IL-15 or IL-23, and this effect was not augmented by the addition of TGF- β (Fig. 3B). On the other hand, IL-15 did not induce the synthesis of IL-22 on TMCs from HT patients, whereas IL-23 or IL-23/TGF- β exerted this effect (Fig. 3C). Finally, IL-6 and TGF- β exerted a

similar effect on the induction of IL-22⁺ cells than IL-23/TGF- β (Fig. 3C).

Additional immunohistochemical analysis showed a prominent expression of both IL-18 receptor and IL-23 receptor, which have been reported to be expressed by Th17 cells in the inflammatory cell infiltrates from HT thyroid specimens (Fig. 3D). In contrast, these cytokine receptors were not detected in GD or normal glands (data not shown).

Serum levels of proinflammatory cytokines are increased in HT patients and are able to induce Th17 differentiation

Then we analyzed the levels of cytokines involved in Th17 cell differentiation, and we found that HT patients had significant increased serum concentrations of IL-6 compared with healthy subjects ($P < 0.05$, Fig. 4A). In addition, serum levels of IL-15 were also significantly augmented in HT patients compared with GD patients and controls ($P < 0.01$ in both cases, Fig. 4B). Finally, al-

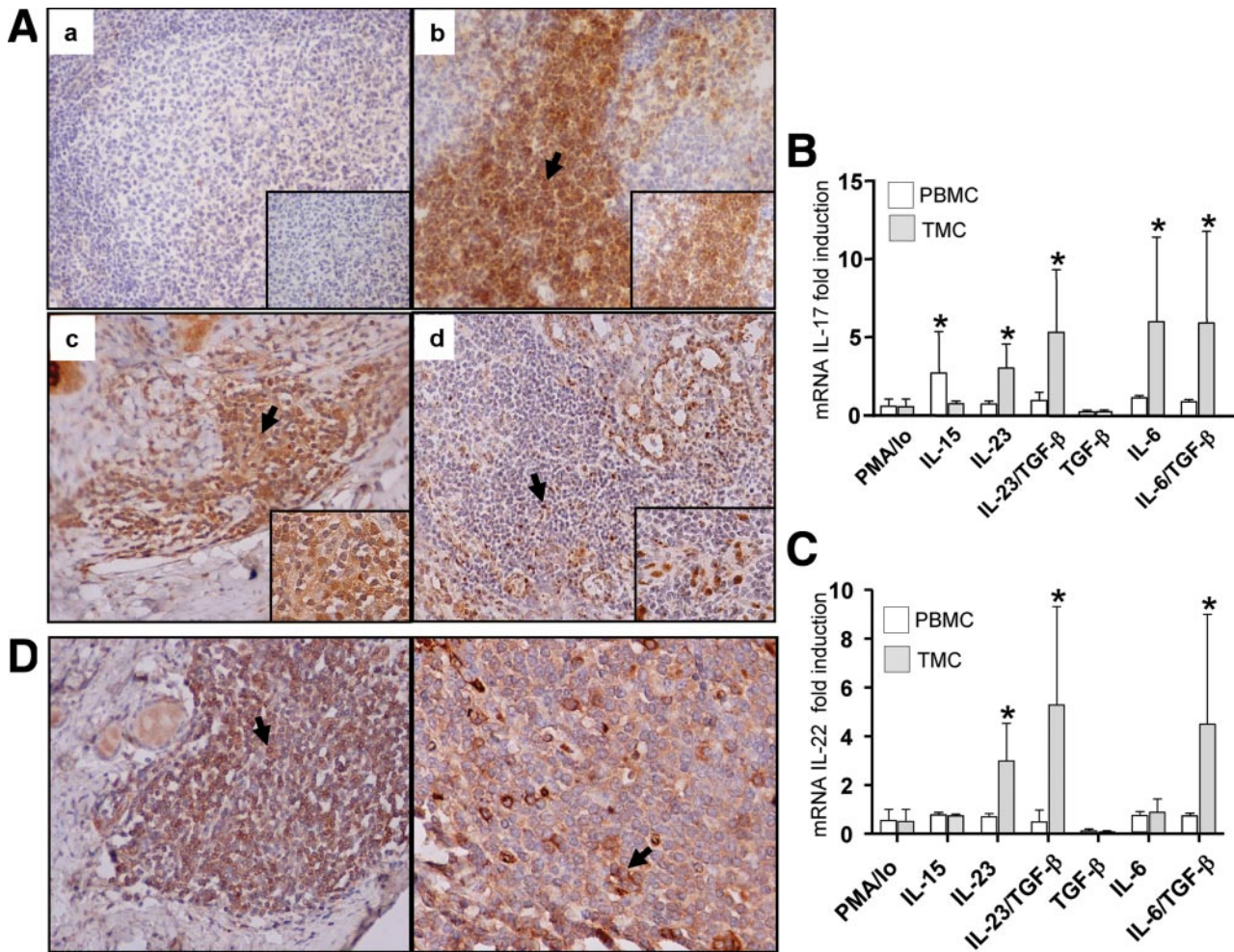


FIG. 3. IL-17 and IL-22 in thyroid tissue from AITD patients. A, Thyroid gland sections from an HT patient were stained for CD4, IL-17, or IL-22 by an immunohistochemical technique. a, Negative X63 control; b, CD4; c, IL-17; d, IL-22. Original magnification, $\times 250$ and $\times 500$ (insets). Arrows point to the inflammatory cell infiltrate. B and C, Detection of IL-17 and IL-22 mRNA in PBMCs and TMCs from HT patients after cell culture with the indicated stimuli. IL-17 and IL-22 mRNA was estimated by real-time RT-PCR. Data correspond to the arithmetic mean \pm SEM of five patients with HT and three with GD. D, Immunohistochemical detection of IL-18R (left panel) and IL-23R (right panel) in thyroid sections from a patient with HT. Original magnification, $\times 250$. *, $P < 0.05$ compared with PMA/ionomycin (Io).

though serum concentrations of IL-23 tended to be higher in HT patients in comparison with both GD patients and healthy subjects, this difference did not reach statistical significance (data not shown).

We also analyzed the *in vitro* differentiation of PBMCs and TMCs into IL-17- and IL-22-producing CD4⁺ T cells (Fig. 4C). When cells were incubated with IL-15 alone, a modest differentiation effect was observed in cells from AITD patients (Fig. 4D). In contrast, IL-23 exerted a greater Th17 differentiation effect, which was higher in HT and GD patients compared with healthy controls (Fig. 4D). Interestingly, TGF- β partially inhibited the inducing effect of IL-23, whereas IL-6, alone or in combination with TGF- β , enhanced the development of Th17 cells in HT and GD patients, and, to a lesser degree, healthy controls (Fig. 4D). These results were confirmed by RT-PCR analysis of the levels of IL-17 and IL-22 mRNA in the indicated cell cultures (Fig. 4E). In addition, different stimuli

(mainly IL-23 and IL-6) were able to induce the release of IL-17 and IL-22 by cultured cells from AITD patients (data not shown). In contrast, all these stimuli were unable to induce the secretion of IFN- γ (data not shown).

Expression of RORC2 gene in cells from AITD patients

Finally, we assessed the expression, at mRNA level, of the RORC2 transcription factor, which plays a critical role in the differentiation of Th17 lymphocytes. As shown in Fig. 5, a strong induction of RORC2 mRNA was observed in cells from HT patients when they were stimulated with IL-23. In contrast, when cells were stimulated with TGF- β alone, no induction of RORC2 was observed, but a prominently expression of Foxp3 was detected (Fig. 5, and data not shown). Furthermore, the addition of TGF- β to IL-23 did not enhance the effect of the latter cytokine on RORC2 induction. On the other hand, IL-15

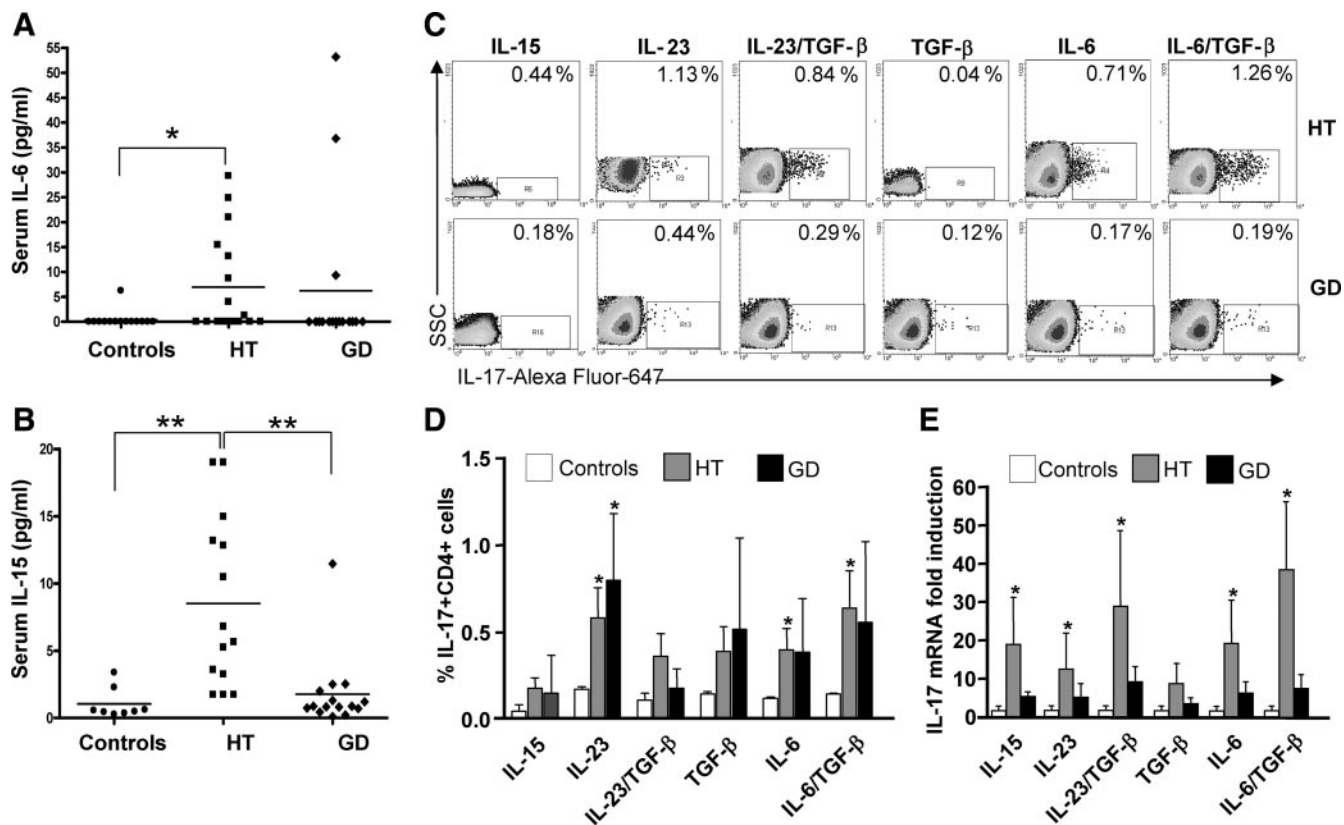


FIG. 4. Serum levels of proinflammatory cytokines are increased in HT and are able to induce the differentiation of Th17 cells *in vitro*. A and B, Serum levels of IL-6 and IL-15 were determined by ELISA in serum samples from AITD patients and controls. Mean group values were compared by one-way ANOVA and nonparametric ANOVA (Kruskal-Wallis test) and *post hoc* comparisons were carried out using the Bonferroni's and Dunn's test as appropriate. *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$. C, *In vitro* differentiation of PBMCs into Th17 cells in patients with AITD (n = 13 with HT, n = 5 with GD). PBMCs were cultured under the indicated stimuli, and then IL-17⁺ cells were analyzed by flow cytometry. Numbers correspond to the percent of IL-17⁺ cells. D, Bars represent arithmetic mean \pm SEM of the percent of IL-17⁺ cells in AITD patients and controls. E, IL-17 mRNA levels in the cell cultures indicated in C. IL-17 mRNA was determined by real-time RT-PCR. *, $P < 0.05$ compared with controls.

as well as IL-6 was also able to induce RORC2 gene expression in cells from HT patients. However, the addition of TGF-β to IL-6 did not enhance the effect of this cytokine on RORC2 expression. Finally, in PBMCs from GD patients, only IL-15 induced the expression of RORC2, whereas the other cytokines tested did not show any significant effect (Fig. 5). As expected, the different stimuli used did not induce the expression of the Th1- and Th2-specific transcription factors: Th1-specific T box transcription factor (Tbet), and GATA binding protein 3, respectively (data not shown).

Discussion

In this work, we explored the possible involvement of Th17 cells in AITD. Th17 cells participate in the resistance against different extracellular bacteria and fungi and act as a link between the innate and adaptive immune response (42). In addition, Th17 lymphocytes are involved in the pathogenesis of different inflammatory and autoimmune conditions, including rheumatoid arthritis, psoriasis, and asthma (12–14). In these conditions, Th17 cells mainly exert their pathogenic effect through the release of IL-17, IL-17F, and IL-22, which in turn induce the synthesis of chemokines and proinflammatory cytokines by resident cells (11, 43).

We herein detected a higher proportion of Th17 cells and raised levels of Th17 cytokines in the peripheral blood and thyroid gland from patients with AITD, mainly those with HT. In addition, the peripheral blood lymphocytes from these patients showed an increased induction of RORC2 gene and an enhanced capability to differentiate *in vitro* into Th17 cells. Finally, high levels of IL-6 and IL-15 were detected in the sera from patients with HT.

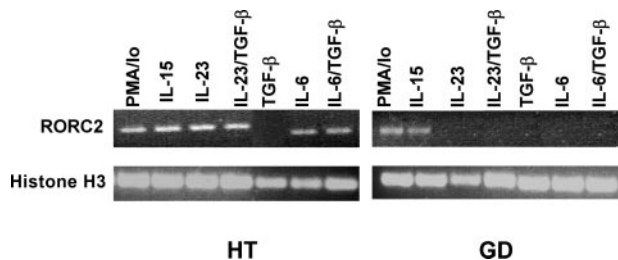


FIG. 5. Increased induction of RORC2 gene in PBMCs from AITD patients. PBMCs from patients with HT and GD were cultured under the indicated stimuli, and then the levels of RORC2 mRNA were determined by RT-PCR. Data from two representative patients, one with HT and one with GD, are shown.

Because it has been described that different cell subsets are able to synthesize IL-17 (44–47), we performed a flow cytometry analysis to detect CD3⁺CD8⁻IL-17⁺IL-22⁺ cells, which very likely correspond to classical Th17 lymphocytes. Our data indicate that patients with AITD, mainly those with HT, show enhanced levels of Th17 cells in their peripheral blood as well as the thyroid gland. We think that this is an interesting finding because it has been largely considered that HT is a Th1-mediated condition. Thus, our data suggest that, as in the case of rheumatoid arthritis or multiple sclerosis, Th17 may have a relevant role in the inflammatory phenomenon and tissue destruction seen in HT. This possibility adds an additional element to the complex pathogenesis of AITD, mainly because it has been proposed that there is a negative cross-regulation between Th1 cells and Th17 lymphocytes (48) and that these two cell subsets are under the control of Treg cells (49). In this regard, it has been recently reported that at least a subset of human Foxp3⁺ Treg lymphocytes is able to synthesize IL-17 as well as express RORC2 (50). Because we previously described that there is an increased proportion of Foxp3⁺ Treg cells in the peripheral blood and thyroid infiltrate of AITD patients (39), it is very likely that these cells correspond to the aforementioned Th17/Treg lymphocytes. However, our data on the defective function of Treg cells in AITD (39) suggest that in this condition, Th17/Treg cells mainly exert a proinflammatory effect, with a weak regulatory function.

The enhanced capability of freshly isolated PBMCs from HT patients to differentiate into Th17 cells is of interest. We think that this phenomenon could be related to the enhanced serum levels of IL-6 and IL-15 detected in these patients. In this regard, the important role of IL-6 in the induction of Th17 cell differentiation has been widely described (19). Likewise, the key effect of IL-15 on the development of Th17 lymphocytes (in combination with IL-6) has been also recently described (51). Although the cause of the enhanced levels of IL-6 and IL-15 in AITD remains to be determined, it is feasible that it could be related to genetic factors, mainly gene polymorphisms, which are associated with an enhanced transcription/translation rate. In this regard, different associations between single nucleotide polymorphisms of proinflammatory cytokine genes (e.g. TNF- α) have been described in patients with AITD (52).

Although to the best of our knowledge this is the first report on the detection of Th17 cells and cytokines in thyroid glands from AITD patients, there is a recent article on the levels of these cells in the peripheral blood (53). In that work, Th17 lymphocyte levels were higher in AITD patients compared with healthy controls, and those patients with intractable GD showed a higher proportion of

Th17 cells than patients with GD in remission. Even though we do not have a commendable explanation for our partially contradictory results, it is feasible that the apparent discrepancies between this study and our work could be due, at least in part, to the different genetic background of the individuals studied, the presence or absence of ophthalmopathy, and the stage of the disease of the patients. Furthermore, it is worth mentioning that we have studied only a small number of thyroid glands because of the limited availability of these specimens. In addition, it is feasible that the resident lymphocyte populations of the thyroid glands studied could be affected by the long treatment history of GD patients. In any case, we consider performing a longitudinal study of Th17 cells in patients with AITD of interest, mainly because it has been suggested that it could occur a Th1-Th2 shift during the evolution of GD (54). Thus, whether there are variations in the commitment or differentiation of Th17 cells during the evolution of HT or GD remains an interesting question.

In summary, our data suggest that there is an enhanced differentiation of Th17 lymphocytes in AITD patients, mainly in those with HT, and that these cells might participate in the pathogenesis of the inflammatory phenomenon and tissue damage observed in this condition.

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