Lymphocytic profiling in thyroid cancer provides clues for failure of tumor immunity

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Abstract
Thyroid cancers are usually surrounded by a significant number of immune-reactive cells. Tumor-associated lymphocytes as well as background lymphocytic thyroiditis are frequently mentioned in pathology reports of patients who have undergone surgery for thyroid cancer. The nature of this lymphocytic reaction is not well understood. The fact that cancer can survive in this adverse microenvironment is indicative of immune regulation. We characterized the lymphocytic infiltration that accompanies thyroid cancer and compared it with that present in thyroid autoimmunity. We found that double-negative (DN) T cells were significantly more abundant in thyroid cancer than in thyroid autoimmunity. Although FOXP3 T regulatory T cells were also present, DN T cells were the dominant cell type, associated with thyroid cancer. Furthermore, upon stimulation, the DN T cells associated with cancer remained unchanged, while the few (<5%) DN T cells associated with thyroid autoimmunity increased in numbers (>20%). CD25 expression on DN T cells remained unchanged after stimulation, which indicates that the increase in the absolute number of DN T cells in thyroid autoimmunity was at the expense of inactivation of single-positive T cells. We concluded that in the setting of thyroid cancer, DN T cells appear to suppress tumor immunity. In contrast, in thyroid autoimmunity, DN T cells were barely present and only increased at the expense of inactivated, single-positive T cells upon induction. Together, these findings indicate that thyroid cancer-associated DN T cells might regulate proliferation and effector function of T cells and thereby contribute to tumor tolerance and active avoidance of tumor immunity.

Key Words
- Hashimoto thyroiditis
- thyroid cancer
- autoimmunity
- tumor immunity

Introduction
Highly immunogenic tumors can spontaneously prime protective immunity, a phenomenon known as tumor immunity (Prehn & Main 1957). However, poorly immunogenic tumors do not spontaneously induce tumor immunity nor do they prime functional T-cell responses (Turk et al. 2004). Unfortunately, such tumors represent a large proportion of cancers in humans, where antitumor T-cell responses are often detected but do not control progression and/or may favor tumor growth (Romero et al. 1998, Valmori et al. 2002). In autoimmunity, on the other hand, immunity is induced to target self. Autoantigen-driven affinity maturation of B and T cells perfects the immune response to its most lethal form. Memory immune cells persist for life and facilitate chronic
target destruction even under immunosuppressive conditions (Jaume 2011).

Thyroid cancer and thyroid autoimmunity seem to be situated at opposite extremes of the immune response spectrum. In thyroid cancer, the immune response seems tolerant, allowing for tumor growth. In thyroid autoimmunity, the immune response is destructive, leading to complete thyroid failure (Paparodis et al. 2014). Although known to immunologists for decades, it has re-emerged in recent years that more effective immunotherapy of cancer is associated with autoimmunity (Korman et al. 2005, Gogas et al. 2006). For example, the appearance of autoantibodies or clinical manifestations of autoimmunity during treatment with interferon α are associated with statistically significant improvements in relapse-free survival and overall survival of patients with melanoma (Gogas et al. 2006). In cancer development, progressive accumulation of genetic abnormalities renders cells malignant. The immune system seems to be allowing or even promoting cancer progression for some tumors (Balkwill et al. 2005). While immune regulation in cancer seems to support development and progression, immune dysregulation in autoimmunity leads to tissue destruction and elimination of the target.

Thyroid cancers are usually surrounded by a significant number of immune-reactive cells. Tumor-associated lymphocytes as well as lymphocytic infiltration organized in germinal centers (GC; chronic lymphocytic thyroiditis or Hashimoto thyroiditis) are frequently mentioned in pathology reports of patients who have undergone surgery for thyroid cancer (Chen et al. 2013). The nature of this lymphocytic reaction is not well understood. The fact that cancer can survive in this adverse microenvironment is indicative of immune regulation.

Suppression of immune responses by regulatory T cells (Tregs) is critical for the induction and maintenance of self-tolerance. Tregs have been shown to be involved in the downregulation of immune responses in autoimmunity, transplant rejection, and tumor immunity (Sakaguchi 2000, Baecher-Allan & Anderson 2006). A variety of T-cell subsets possess immunoregulatory properties. The thymus-derived Treg (naturally occurring CD4+ CD25+ FOXP3+, Abbas et al. 2013) cells are the most extensively studied subset of Tregs and their role has been investigated in a wide range of animal models and in humans (Bennett et al. 2001, Kriegel et al. 2004). However, inducible Tregs such as T-regulatory type 1 (Tr1) cells, T-helper 3 (Th3) cells, CD8+ CD28− T cells, and CD3+ CD4+ CD8− double-negative (DN) T cells also exhibit the ability to inhibit immune responses (Groux et al. 1997, Wildin et al. 2001, Chang et al. 2002, Faria & Weiner 2006, Zhang et al. 2006).

In this study, we characterized the lymphocytic infiltration that accompanies human papillary thyroid cancer (PTC; the most common form of thyroid cancer) and compared it with that present in human autoimmune thyroid disease (Hashimoto thyroiditis, the most common form of thyroid autoimmunity). Ex vivo aspirations of human thyroid glands affected by both diseases and controls were carried out. Lymphocytes were isolated and either analyzed immediately by flow cytometry or placed in culture and stimulated after 24 h. Thyroid tissue specimens were also studied. We found that the lymphocytic microenvironment was critically different in both thyroid diseases and that immunity in thyroid cancer was under the control of an unexpected T cell player.

Subjects and methods

Thyroid subjects

The UW Thyroid Multidisciplinary Clinic is a large referral site for thyroid diseases. Patients referred for thyroid surgery include those with positive or suspicious cytology for malignancy on fine-needle aspiration (FNA) and those with goiter associated with compressive symptoms (such as dysphagia, shortness of breath or hoarseness). Hashimoto thyroiditis per se is not an indication for thyroid surgery; however, some patients develop compressive symptoms that require surgical intervention. For this study, we identified patients with cytology-proven thyroid cancer and Hashimoto thyroiditis. Patients undergoing thyroid surgery had their thyroids ex vivo aspirated in the operating room. A tissue sample from each patient was snap frozen in liquid nitrogen and stored for further analysis. Postoperative histology confirmed the presence of thyroid cancer and Hashimoto thyroiditis. For those patients with histology of Hashimoto’s, we confirmed the diagnosis (hypothyroidism with abnormally high TSH and low free T4 levels) had been made before surgery. We only included Hashimoto’s patients in whom hypothyroidism was not due to previous thyroid surgery or radioactive iodine treatment, but rather the pathological outcome of chronic thyroiditis causing thyroid failure. Established pathological characteristics were followed by our academic pathologists for the diagnosis of thyroid cancer and Hashimoto’s. The diagnosis of Hashimoto thyroiditis relied on the presence of both thyroid follicles accompanied by lymphocytic infiltrates and the formation of GC (Fig. 1 top right as an example). Representative thyroid cancer sample coexisting with lymphocytic infiltration is shown in Fig. 1 top left. Patients diagnosed with either PTC (n=11) or hypothyroid
Figure 1
Surgical pathology. Hematoxylin/eosin and immunofluorescence staining. 40× magnification of representative surgical pathology specimens from patients with hypothyroid Hashimoto thyroiditis and papillary thyroid carcinoma (PTC) with tumor-associated lymphocytes (TAL) stained with hematoxylin/eosin (top). PTC, papillary thyroid carcinoma; GC, germinal center-like structures. Images at the same magnification and representative specimens immunostained with DAPI (red for nuclei), anti-CD8 in green, and merged images of both are subsequently shown. Inset (A) is a close-up of CD8+ stained cells.
Hashimoto thyroiditis (Hash, n=7) were studied. Controls (Ctr, n=7) included patients undergoing thyroid surgery for none of these conditions (i.e., symptomatic multinodular goiters). The collection of patient’s tissue/data and subsequent analysis was approved by University of Wisconsin Human Subjects Institutional Review Board. Patients were invited to participate and sign consent pre-operatively to be part of this study.

Tissue staining

Frozen tissue sections were stained first with hematoxylin and eosin and used for basic histology and for correlation with specific immunohistochemistry/fluorescence studies. Anti-CD19 for identification of B-cell lineage, anti-CD3 for T-cell lineage, and anti-CD45 common leukocyte antigen (all 1:100 from BD Pharmingen, San Jose, CA, USA) were used as primary antibodies for immunohistochemistry. Secondary species-specific HRP-linked antibodies were developed using a commercial substrate (Vector, Burlingame, CA, USA). Histomorphometric microscopic studies were carried out in 15 or more randomly selected fields (as previously described in Mirocha et al. (2009)). Immunofluorescence was carried out on sections pre-treated for antigen retrieval with Retrivagen (BD Pharmingen) at 90°C for 10 min. Endogenous peroxidase activity was quenched with 0.3% H2O2 for 10 min in PBS. For reduction of background staining, the sections were treated for 1 h with 5% normal serum from the same species as the labeling antibody used. The samples were then incubated with primary antibody in a humid chamber at 4°C overnight. Primary antibodies used included anti-human CD4, and CD8 (all 1:100; BD Pharmingen). Secondary MAB, biotinylated and species-specific, were then stained with avidin-fluorescein isothiocyanate (Vector, Burlingame, CA, USA). Histomorphometric analyses were done on these individual cell populations. CD4, CD8, CD14, CD19, CD25, CD45, and iNKT (anti-CD3-allophycocyanin/HIT3a, anti-CD4-APC-cy7/RPA-T4, anti-CD8-PE-cy5/HIT8a, anti-CD14-PE/M5E2, anti-CD19-PE-cy5/HIB19, anti-CD25-PE/M-A251, anti-CD45-FITC/H130, anti-CD56-PE-cy7/B159, anti-iNKT FITC/6B11, TCRαβ-FITC/T10B9.1A-31, and TCRγδ-PE/B1; BD Biosciences (San Jose, CA, USA)) or isotype controls in serum-containing media. Freshly isolated single cells were incubated with antibodies for 20 min on ice for surface staining, washed, and fixed in 1% paraformaldehyde for 15 min. A subset of cells was permeabilized with cytofix/cytosperm fixation and permeabilization solution (BD Biosciences) and stained with fluorochrome-conjugated antibodies against human IL17, interferon γ (IFNγ), and Foxp3 (anti-IL17-PE/N49-653, anti-IFNγ-PE/25723.11, anti-Foxp3-Alexa Fluor 488/259D/C7; BD Biosciences). The cells were also stained with Hoechst 33342 (10 μg/ml for 2 h, Hoechst fluorescence, 350 nm excitation/450 nm emission, linear scale) to gate live cells containing 2n–4n cellular DNA.

Induction of lymphocytes

A minimum of three million cells from each freshly isolated thyroid ex vivo aspirate were incubated in RPMI1640 with 10% FCS for 18 h at 37°C in a 5% CO2 atmosphere. Then, cells were stimulated (induced) with phorbol 12-myristate 13-acetate (PMA, 50 ng/ml), ionomycin (1 μg/ml), and Golgi plug brefeldin A (1 μg/ml), (Sigma–Aldrich) for 6 h. The unstimulated cells served as controls.

Flow cytometry data analysis

All samples were prepared in triplicates and mean of the three samples was considered as individual data. In each replicate at least 10 000 cells were acquired. The samples were acquired in a LSRII flow cytometer (BD Biosciences). Analysis of fluorescence-activated cell sorting (FACS) data was done with FlowJo v. 7.6.5 software (TreeStar Inc. Ashland, OR, USA). Analysis of the cell population was carried out based on four step criteria. The first step was the gating of lymphocytes on the basis of forward- and side-scatter characteristics. The second step was the removal of dead cells, debris and doublets on the basis of DNA integrity using Hoechst 33342 stain. The third step was the gating of T cells and non-T cells on the basis of CD3 expression. From CD3-gated cells, CD4 vs CD8 positive cells were plotted, which allowed the identification of discrete CD4, CD8 and DN T cell populations. Subsequent analyses were done on these individual cell populations.
Statistical analysis

Since data were not normally distributed, data were transformed to natural logarithms or ranks. Individual end points were analyzed by group interactions. The SAS MIXED procedure (version 9.3, SAS Institute, Inc., Cary, NC, USA) was used for statistical analyses. Student’s *t*-test or the Mann–Whitney *U* test was used to identify the differences between the groups and controls. A probability of *P* ≤ 0.05 indicated that a difference was significant, and a probability of *P* > 0.05 to *P* ≤ 0.1 indicated that a trend towards significance was approached. Data are presented as the mean ± S.E.M.

Results

We propose the hypothesis that intra-thyroidal lymphocytes accompanying thyroid cancer may be reactive to the presence of the tumor and different from lymphocytes found in autoimmune thyroid disease. To address our hypothesis, we designed a systematic characterization of the lymphocytic microenvironment of human post-surgical thyroid specimens.

Subtle differences in thyroid autoimmunity and thyroid cancer lymphocytic milieu

Patients with FNA cytology consistent with Hashimoto thyroiditis (hypothyroid by history), those with cytology positive for PTC, and those undergoing thyroid surgery for symptomatic multi-nodular goiters (controls) were identified. Patients’ thyroid glands were *ex vivo* aspirated (for FACS) and small pieces of thyroid tissues snap frozen in liquid nitrogen (for staining) in the operating room. The frozen tissues were processed and stained first with hematoxylin/eosin (H&E, Fig. 1 top as examples) and then with immunostains for characterization of the lymphocytic profile of the different thyroid conditions. H&E staining was not helpful in distinguishing the differences in lymphocytic infiltration (Fig. 1 top). Immunostaining, however, showed subtle differences among the groups. H&E staining showed that patients with PTC had more dispersed lymphocytic infiltrates, while in the hypothyroid Hashimoto’s patients the lymphocytic infiltration appeared to be more organized in structures resembling lymph-node GC. Nevertheless, GC were sometimes present in both groups. In specimens from hypothyroid Hashimoto’s patients, many lymphocyte specimens from all patients of both groups. Percentages of CD3+CD4+ gated T cells sorted for the same immune markers are shown. Black bars represent results for specimens from papillary thyroid cancer (PTC) patients. White bars represent results for specimens from hypothyroid Hashimotos’ patients (Hash).

Figure 2

Flow cytometry of classic Tregs. (Top) Contours plots of CD3+CD4+ lymphocyte specimens from representative patients with papillary thyroid carcinoma (PTC) and hypothyroid Hashimoto thyroiditis (Hash). CD3+ and CD4+ gated T cells were sorted for CD25, Foxp3, and both (classic Tregs, CD3+CD4+CD25+Foxp3+). (Bottom) Bar graph of statistical analysis of
lymphocytes stained positive for CD8, while in specimens from patients with PTC, CD8-positive lymphocytes were less abundant (Fig. 1 DAPI for nuclei in red, CD8 in green, and merged images of both shown in bottom panels). Notably, many lymphocytes did not stain for either CD4 or CD8. Controls did not have measurable infiltrates and lymphocytes present were not different than what is expected in peripheral blood (not shown).

**FOXP3** Tregs were predominantly present in thyroid cancer but in low numbers

The combined expression of CD4, CD25, and FOXP3 identifies FOXP3$^+$ Tregs (Sakaguchi 2000). FOXP3$^+$ Tregs were more numerous in PTC specimens, scarce in hypothyroid Hashimoto’s ones and almost undetectable in controls (not shown). As expected from what we observed on histology, when *ex vivo*-extracted lymphocytes were analyzed by flow cytometry, in patients with PTC, the associated lymphocytic concentration of FOXP3$^+$ Tregs was higher than in patients with hypothyroid Hashimoto’s (Fig. 2). However, although the difference was significant, as a percentage of the total lymphocytic population, the FOXP3$^+$ Tregs represented <2.5%.

**CD3$^+$ CD4$^-$ CD8$^-$ T cells stand out as the dominant T cell population in thyroid cancer**

FACS analysis revealed another group of T cells never described in the setting of thyroid cancer. CD3$^+$ CD4$^-$ CD8$^-$ (double negative, DN) T cells were significantly more abundant in lymphocytic infiltrates accompanying thyroid cancer than in those present in hypothyroid Hashimoto’s (Fig. 3) and only traceable in controls (not shown). Overall, DN T cells were the dominant T cell type associated with thyroid cancer. DN T cells were 20 times more abundant than classic Tregs in this setting. Reciprocally, CD4$^+$ and CD8$^+$ T cell counts were also significantly lower in thyroid cancer as seen in the histological analysis, while natural killer T (NKT) cells only displayed a trend towards lower levels (Fig. 3). Upon PMA/Ionomycin stimulation/induction (Fig. 4), the DN T cells associated with the thyroid cancer remained unchanged while the few (<5%) DN T cells associated with hypothyroid Hashimoto’s increased in absolute numbers (>20%).

**IFNγ and IL17 are augmented in thyroid cancer, mostly from DN T cells**

IFNγ production from lymphocytes accompanying thyroid cancer was overall higher. A major contributor to
this IFNγ increase was the DN T cell population (over 40%, Fig. 5). A similar observation was made for IL17 production although CD4+ T cells were also contributors (Fig. 6). We interpreted the cytokine production as a regulatory phenomenon because it derived mostly from the DN T cells and reciprocally affected single-positive T cells (Figs 5 and 6). After PMA/ionomycin induction, there was a small reduction in IFNγ-producing CD4+ T cells that was more pronounced in the Hashimoto’s lymphocytes than in the thyroid-cancer-associated lymphocytes which rendered the difference between the two groups significant (Fig. 5). For IL17-producing CD4+ T cells that difference became almost significant as a consequence of both increased numbers in the thyroid cancer specimens and decreased numbers in Hashimoto’s ones (Fig. 6). The opposite was observed for IL17-producing DN T cells which were decreased after induction in PTC and increased in Hash (Fig. 6).

**Activated DN T cells in thyroid cancer and inactivated DN T cells in Hashimoto’s**

The highest expression of CD25 (α chain of the IL2 receptor and maker for activation) was found in DN T cells in the setting of thyroid cancer and remained almost unchanged (already maximized) after stimulation/induction (Fig. 7A). The expression of CD25 in DN T cells in the setting of hypothyroid Hashimoto’s also remained unchanged, which indicates that the increase in the absolute number of DN T cells (Fig. 4) was at the expense of inactivated single-positive T cells (mostly CD4+ T cells). PMA/ionomycin stimulation/induction is known to inhibit the expression of CD4 and CD8 in vitro (DiSanto et al. 1989, Jin et al. 2004). Therefore, the increase in DN T cells in Hashimoto’s upon induction is probably the consequence of loss of CD4/CD8 expression (inactivated single-positive T cells).

**Macrophages, B cells, and all other monocytes were not particularly affected**

We also looked at the potential differences in CD19 (B cells), CD14 (macrophages), and CD45 (common leukocyte antigen) expression between the two groups of patients. Under both conditions, the number of macrophages was similar as was the number of B cells (albeit proportionally high as compared with expected peripheral counts and controls, Fig. 7B). Induction did not significantly affect CD19 and CD14 expression under either disease condition. CD45 expression accounted for most of the lymphocytes studied in both groups. Induction decreased overall expression more in thyroid-cancer-associated lymphocytes, but the differences were not significant (Fig. 7C).

**Discussion**

Glands from patients with Hashimoto thyroiditis had lymphocytic infiltrates functionally different from
those accompanying thyroid cancers. Lymphocytes in Hashimoto’s were mainly effector cells. Lymphocytes accompanying thyroid cancer appeared to be under T cell regulation. CD4+ and CD8+ T cells were significantly decreased, while CD4− CD8− T cells were significantly increased. This, to our knowledge, never previously described phenomenon may be explained by a few possibilities. One could be that the decreased count of CD4+ and CD8+ T cells in patients with thyroid cancer is the result of T cell ‘exhaustion’. Indeed, T cell exhaustion has been described as a state of T cell dysfunction that arises during many chronic infections and cancer (Wherry 2011). T cells can recover from exhaustion by boosting immune responses (Wherry 2011). However, we did not observe recovery at least after short induction. Another possibility is that thyroid cancer negotiates its survival with the immune system by downregulating effector T cells while recruiting Tregs. The cancer ‘immunoediting’ hypothesis stresses the dual host-protective and tumor-promoting actions of immunity in the development of tumors (Schreiber et al. 2011). Specifically, it proposes three phases of immune modulation of cancer development (elimination, equilibrium, and escape) which help to explain our observations and give us a mechanistic framework for our hypothesis. Lymphocytes in Hashimoto’s are clearly capable of elimination of the target. As with anti-CTLA-4 immunotherapy, exacerbation of rampant autoimmunity may help to eliminate cancer (Mellman et al. 2011). Lymphocytes accompanying thyroid cancer on the opposite extreme appear to be in equilibrium with cancer. The immune response in thyroid cancer seems tolerant to the presence of the tumor as if it was regulated. We found higher numbers of FOXP3+ Tregs in the thyroid cancer immune microenvironment compared with Hashimoto’s. However, Hashimoto’s specimens were from patients with long-standing disease that required surgery either because of compressive symptoms (fibrosis) or pain (painful thyroiditis). Surprisingly, we found that the regulatory immune component in thyroid cancer was represented by an unexpected player. In both mice and humans, about 1–5% of all peripheral T cells are of DN phenotype. These T cells do

![Figure 5](http://erc.endocrinology-journals.org)

**Figure 5**

Interferon γ production by T cell type. Interferon γ (IFNγ) production from lymphocytes in patients with papillary thyroid cancer (PTC) and hypothyroid Hashimoto thyroiditis (Hash) is shown. (Left panels) Uninduced (untreated) lymphocyte FACS contour plots of representative patients (top) and bar graphs of statistical analysis of all T cells (bottom) by T cell type. (Right panels) Induced (PMA/Ionomycin treated) lymphocyte FACS contour plots of representative patients (top) and bar graphs of statistical analysis of all T cells (bottom) by T cell type. Black bars represent results for specimens from papillary thyroid cancer (PTC) patients. White bars represent results for specimens from hypothyroid Hashimoto’s patients (Hash).
CD4 or CD8 cell surface molecules and show a characteristic cytokine profile. Zhang and colleagues were the first, to our knowledge, to identify and characterize the immunoregulatory function of DN T cells. They demonstrated that murine DN T cells specifically eliminate activated anti-donor CD4<sup>+</sup>, CD8<sup>+</sup>T cells and that adoptive transfer of DN T cells prolongs skin and heart allograft survival in transplant models (Zhang et al. 2006). Human DN T cells can strongly suppress the proliferation of CD4<sup>+</sup> and CD8<sup>+</sup>T cells too. In our study, CD4<sup>+</sup> and CD8<sup>+</sup>T cells were significantly affected by the presence of DN T cells in thyroid cancer (Fig. 8). Moreover, human DN T cells are able to downregulate cytokine production of highly activated human effector T cells (Merims et al. 2011). As observed in our study, human DN T cells produce interferon γ (IFNγ) as an important mediator in the control of proliferating lymphocytes. In line with their ability to produce high levels of both IFNγ and IL-7 (as shown in our study), DN T cells have also been reported to play a key regulatory role in various models of chronic infections. DN T cells are the major responding T cell subset in the lungs of mice infected with live strain vaccine (LSV) of the intracellular bacterium Francisella tularensis. IL17-producing DN T cells do not eradicate the infection, but allow a chronic state of infection to progress. DN T-IL17-deficient mice die at day 8–10 post LSV acute infection (Cowley et al. 2010). This indicates that IL17-producing DN T cells allow survival by regulating the acute infection to transition into a chronic state.

In our study, we observed similar diminished single-positive (CD4 and CD8) T cell responses in thyroid cancer. DN T cells were significantly increased while CD4 and CD8 T cells were significantly decreased in thyroid cancer, as opposed to what we observed in thyroid autoimmune disease (Figs 3 and 8). Although FOXP3<sup>+</sup> Tregs were also present, DN T cells were the dominant cell type associated with thyroid cancer. Furthermore, upon stimulation/induction, the DN T cells associated with thyroid cancer remained unchanged while the few (<5%) DN T cells associated with thyroid autoimmune disease increased in...
numbers (>20%, Fig. 4). The CD25 expression on DN T cells in the setting of thyroid cancer remained unchanged after stimulation (Fig. 7A). CD25 expression on DN T cells in the setting of thyroid autoimmune disease also remained unchanged (Fig. 7A), which indicates that the increase in the absolute number of DN T cells was at the expense of inactivation of single-positive T cells.

We do not know why DN T cells populate the microenvironment of thyroid cancer. Fas death-receptor dysfunction is known to give rise to DN T cells in mice (Shirai et al. 1990). Mice with defective expression of Fas/APO-1/CD95 (lpr) or Fas-ligand (gld) have been widely used to investigate the developmental origin of DN T cells. These mice show a peculiar accumulation of DN T cells in the lymph nodes and other organs (Shirai et al. 1990, Watanabe et al. 1995, Mohamood et al. 2008). The clinical relevance of these observations has been confirmed in humans. Patients suffering from autoimmune lymphoproliferative syndrome (ALPS, also known as Canale-Smith syndrome) have mutations in the Fas apoptotic pathway (Fisher et al. 1995). Interestingly, these patients share similar clinical features with lpr and gld mice and a selective accumulation of DN T cells (Poppema et al. 2004, Teachey et al. 2005, Cerutti et al. 2007, Marlies et al. 2007, Fleisher 2008). Moreover, as in patients with ALPS, DN T cells found in the thyroid cancer microenvironment are all αβ-TCR positive.

A aberrant FAS expression is known to occur in thyroid cancer. FAS is expressed in most thyroid carcinomas, yet its cross-linking fails to induce apoptosis in thyroid cancer cell lines in vitro (Mitsiades et al. 2000). In fact, thyroid cancer cells simultaneously express their own FAS ligand, which may confer a protective immunomodulatory effect and specimens for the presence of CD14 and CD19 (uninduced-untreated, left and induced-PMA/Ionomycin treated, right). (C) Bar graphs of statistical analysis of lymphocyte specimens for the presence of CD45 (uninduced-untreated and induced-PMA/Ionomycin treated). Black bars represent results for specimens from papillary thyroid cancer (PTC) patients. White bars represent results for specimens from hypothyroid Hashimoto’s patients (Hash).

Figure 7
CD25, CD14, CD19, and CD45 in all lymphocytes. Flow cytometry of lymphocyte specimens from patients with papillary thyroid cancer (PTC) and hypothyroid Hashimoto thyroiditis (Hash) for specific immune markers. (A) Bar graphs of statistical analysis of lymphocyte specimens for the presence of CD25 (uninduced-untreated, left and induced-PMA/Ionomycin treated, right) by T cell type. (B) Bar graphs of statistical analysis of lymphocyte specimens for the presence of CD14 and CD19 (uninduced-untreated, left and induced-PMA/Ionomycin treated, right). (C) Bar graphs of statistical analysis of lymphocyte specimens for the presence of CD45 (uninduced-untreated and induced-PMA/Ionomycin treated). Black bars represent results for specimens from papillary thyroid cancer (PTC) patients. White bars represent results for specimens from hypothyroid Hashimoto’s patients (Hash).

Figure 8
DNT/CD8 and DNT/CD4 T cell ratios in individual patients. The proportion of DN T cells as a function of CD8⁺ (A) and CD4⁺ (B) T cells in intra-thyroidal lymphocytes of patients with papillary thyroid cancer (PTC, circles) and hypothyroid Hashimoto’s (Hash, squares) is shown.
clearly has no negative effect on their survival (Mitsiades et al. 1999). Moreover, FAS has been reported to be actually overexpressed in papillary thyroid carcinomas compared with paired normal thyroid tissues (Arscott et al. 1999). One study clearly showed that thyroid cancer cells can divert FAS signaling from apoptosis to proliferation and may actually exploit FAS expression to their advantage (Mitsiades et al. 2004). Another study found a germline single-nucleotide polymorphism in the FAS gene in thyroid cancer patients that might explain its increased expression (Basolo et al. 2004). Hence, aberrant FAS expression may provide the thyroid cancer hosting signal for DN T cell recruitment/development as in lpr/gld mice and ALPS in humans.

In conclusion, our study shows for the first time that DN T cells appear to downregulate proliferation and cytokine production of activated effector T cells coexisting that DN T cells appear to downregulate proliferation and effector function of T cells and thereby contribute to tumor tolerance and active avoidance of tumor immunity.

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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References

Arscott PL, Stokes T, Myc A, Giordano TJ, Thompson NW & Baker JR Jr 1999 Fas (CD95) expression is up-regulated on papillary thyroid carcinoma. Journal of Clinical Endocrinology and Metabolism 84 4246–4252. (doi:10.1210/jcem.84.11.6139)


Cowley SC, Meierovics AI, Freienger JA, Iwakura Y & Elkins KL 2010 Lung CD4<sup>+</sup>CD8<sup>−</sup> double negative T cells are prominent producers of IL-17A and IFN-γ during primary respiratory murine infection with Francisella tularensis live vaccine strain. Journal of Immunology 184 5791–5801. (doi:10.4049/jimmunol.1000362)


Jin YJ, Zhang X, Boursiquot JG & Burakoff SJ 2004 CD4<sup>+</sup> and CD8<sup>−</sup> T cell populations have distinct functional properties. Journal of Immunology 173 5495–5500. (doi:10.4049/jimmunol.173.9.5495)


Kriegl MA, Lohmann T, Gabler C, Blank N, Kalden JR & Lorenz HM 2004 Defective suppressor function of human CD4<sup>+</sup>CD25<sup>+</sup> regulatory T


Shirai T, Abe M, Yagita H, Okumura K, Morse HC III & Davidson WF 1990 The expanded populations of CD4+CD8− T cell receptor α/β+T cells associated with the lpr and gld mutations are CD2. *Journal of Immunology* **144** 3756–3761.


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