# Analysis of T cell receptor V $\beta$ diversity in peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T lymphocytes in patients with autoimmune thyroid diseases

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# Summary

Autoimmune thyroid diseases are characterized by intrathyroidal infiltration of CD4<sup>+</sup> and CD8<sup>+</sup> T lymphocytes reactive to self-thyroid antigens. Early studies analysing T cell receptor (TCR) Va gene usage have shown oligoclonal expansion of intrathyroidal T lymphocytes but not peripheral blood T cells. However, TCR VB diversity of the isolated CD4<sup>+</sup> and CD8<sup>+</sup> T cell compartments in the peripheral blood has not been characterized fully in these patients. We performed complementarity-determining region 3 (CDR3) spectratyping as well as flow cytometric analysis for the TCR V $\beta$  repertoire in peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T cells from 13 patients with Graves' disease and 17 patients with Hashimoto's thyroiditis. Polyclonal TCR V $\beta$  repertoire was demonstrated by flow cytometry in both diseases. In contrast, CDR3 spectratyping showed significantly higher skewing of TCR V $\beta$  in peripheral CD8<sup>+</sup> T cells but not CD4<sup>+</sup> T cells among patients with Hashimoto's thyroiditis compared with healthy adults. We found trends towards a more skewed CDR3 size distribution in those patients having disease longer than 5 years and requiring thyroid hormone replacement. Patients with Graves' disease exhibited no skewing both in CD4<sup>+</sup> and CD8<sup>+</sup> T cells. These findings indicate that clonal expansion of CD8<sup>+</sup> T cells in Hashimoto's thyroiditis can be detected in peripheral blood and may support the role of CD8<sup>+</sup> T cells in cell-mediated autoimmune attacks on the thyroid gland in Hashimoto's thyroiditis.

**Keywords:** autoimmune, Graves' disease, Hashimoto's thyroiditis, TCR  $V\beta$  repertoire, spectratyping

# Introduction

Autoimmune thyroid diseases, including Graves' disease and Hashimoto's thyroiditis, are prototypes of human organspecific autoimmune diseases. Hyperthyroidism in Graves' disease is caused by thyroid-stimulating immunoglobulins that bind to and stimulate the thyroid-stimulating hormone receptor, whereas hypothyroidism in Hashimoto's thyroiditis is associated with autoantibodies to thyroperoxidase and thyroglobulins [1]. Variable numbers of locally infiltrating T and B lymphocytes may be involved in the production of such autoantibodies [2,3]. Indeed, approximately 10% of activated T cells infiltrating the thyroid gland in those patients has been shown to proliferate in response to thyroid cell antigens [4]. Analysis of the T cell receptor (TCR) repertoire is one of the most reliable tools to detect clonally expanded antigen-driven T cell populations. Although there is some controversy, studies of TCR V $\alpha$  usage in patients with autoimmune thyroid diseases have demonstrated that clonally expanded T cells existed in the thyroid gland, and that TCR V gene expressions by intrathyroidal T cells were different from those found in peripheral T cells from the same patients [5–7].

A sensitive method to study TCR repertoire is to determine size distribution of cDNA for the complementaritydetermining region 3 (CDR3) of TCR V $\beta$ ; that is, the hypervariable region generated by genetic rearrangements during T cell maturation in the thymus [8]. This CDR3 spectratyping allows us to determine clonal dominance or restriction within T cell populations. Using this method, in combination with flow cytometric analysis for TCR V $\beta$ , we analysed the entire TCR V $\beta$  repertoire both in peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T cells among patients with autoimmune thyroid diseases and found significant skewing of TCR V $\beta$ usage in circulating CD8<sup>+</sup> T cells from Hashimoto's thyroiditis. We also discuss the correlation between the TCR V $\beta$  restriction and clinical features in those patients.

# Table 1. Patient characteristics.

	Graves' disease	Hashimoto's thyroiditis	Normal range
No. of cases	13	17	
Sex (no. male/female)	4/9	2/15	
Age* (years, mean $\pm$ s.d.)	$15.6 \pm 4.2$	$21.0 \pm 6.8$	
(no. < 15 years/ $\geq$ 15 years)	4/9	4/13	
Disease duration (years, mean $\pm$ s.d.)	$3.9 \pm 5.0$	$7.9 \pm 6.1$	
(no. < 5 years/ $\geq$ 5 years)	8/5	6/11	
Serum FT3* (pg/ml)	$6.14 \pm 3.60$	$2.86 \pm 0.38$	2.20-4.30
Serum FT4* (ng/dl)	$2.03 \pm 0.9$	$1.17 \pm 0.27$	0.80-1.80
Serum TSH* (µU/ml)	$0.27 \pm 0.79$	$6.89 \pm 16.30$	0.27-4.65
AbTg* (IU/ml)	$749.3 \pm 1037.3$	$499.3 \pm 739.1$	$\leq 44.0$
AbTPO* (IU/ml)	$315.6 \pm 220.5$	$238.4 \pm 251.8$	$\leq 7.2$

\*Data at sample collection. FT3, free triiodothyronine; FT4, free thyroxine; TSH, thyroid-stimulating hormone; AbTg, anti-thyroglobulin antibody; AbTPO, anti-thyroid peroxidase antibody; s.d., standard deviation.

# Materials and methods

### Patients

We studied 30 Japanese patients affected with autoimmune thyroid diseases: 13 with Graves' disease and 17 with Hashimoto's thyroiditis (Table 1). All but three patients with Hashimoto's thyroiditis showed disease onset before 18 years of age. The diagnosis was based on clinical data and thyroid ultrasound imaging, in which all patients showed moderately heterogeneous, reduced echogenicity. Two patients had other autoimmune diseases: glomerulonephritis associated with anti-neutrophil cytoplasmic antibodies existed in patient 11 of Graves' disease, and type 1 diabetes in patient 3 of Hashimoto's thyroiditis. No patients underwent fineneedle biopsy. All patients with Graves' disease were treated with anti-thyroid drugs (methimazole or propylthiouracil) because of hyperthyroidism, and euthyroidism was achieved in six patients at the time of sample collection. Because of hypothyroidism, nine of 17 patients with Hashimoto's thyroiditis received levothyroxine, and one patient remained hypothyroid at the time of sampling (patient 3). Disease duration of Hashimoto's thyroiditis patients with and without levothyroxine requirement was  $8.3 \pm 7.0$  and  $7.5 \pm 5.4$  respectively. No patient was under treatment with immunomodulant therapies. Approval for the study was obtained from the Human Research Committee of Kanazawa University Graduate School of Medical Science, and informed consent was provided according to the Declaration of Helsinki.

### Cell preparations

Peripheral blood mononuclear cells (PBMCs) were isolated by Ficoll-Hypaque gradient centrifugation from the patients and controls. CD4<sup>+</sup> and CD8<sup>+</sup> T cells were purified by positive selection from PBMCs using monoclonal antibody (mAb)coated magnetic beads according to the manufacturer's instructions (Becton Dickinson, San Diego, CA, USA) [9]. The purity of each isolated cell population exceeded 90% routinely, as assessed by flow cytometric analysis.

# The CDR3 spectratyping

Total RNA was extracted from CD4+ and CD8+ T cells with Trizol reagent (Invitrogen, Carlsbad, CA, USA), and firststrand cDNA was generated from 2 µg total RNA with random hexamers and RAV-2 reverse transcriptase (Takara Bio Inc., Shiga, Japan). CDR3 size distribution assay was performed as described [10,11]. Briefly, each TCR VB fragment was amplified with one of the 25 V $\beta$ -specific primers and a 6-fluorescein phosphoramidite-labelled CB primer recognizing both C\u00df1 and C\u00df2. Thirty-five cycles of amplification (94°C for 1 min, 55°C for 1 min, and 72°C for 1 min) were used. One µl of 1:10 diluted polymerase chain reaction (PCR) products was mixed with 12 µl of deionized formamide and 0.5 µl of size standard and heated at 95°C for 3 min. The size distribution of each fluorescent PCR product was determined by electrophoresis in performance optimized polymer 4 on the Applied Biosystems 310 automated sequencer, and data were analysed using GeneScan software (Applied Biosystems, Foster City, CA, USA). To assess each individual profile as normal or skewed, we used a previously reported complexity scoring system [12,13]; that is, a complexity score = (sum of all the peak heights/sum of the major peak heights) × (number of the major peaks). Major peaks were defined as those on the spectratype histogram whose amplitude was at least 10% of the sum of all the peak heights. The mean complexity scores of healthy adult controls was 5.06 (range: 4.89-5.19), as reported previously [13]. The number of TCR V $\beta$  subfamilies with skewed CDR size pattern, as shown by a complex score lower than 4, was determined for each subject [13,14]. Analysis of differences among data groups was performed using the Student's t-test for unpaired samples. Values of P less than 0.05 were considered significant.

# Flow cytometric analysis for TCR V $\beta$ repertoire

Three-colour immunofluorescence analysis was used to study TCR V $\beta$  repertoire distribution, as described previously [11]. Briefly, after washing twice in phosphatebuffered saline, peripheral blood samples were incubated with appropriate phycoerythrin (PE)-conjugated mAbs with specificity for TCR V $\beta$  1-23 (Immunotech, Marseille, France), fluorescein isothiocyanate-conjugated anti-CD8 (Becton Dickinson) and R-PE-Cy5-conjugated anti-CD4 (Dako, Glostrup, Denmark) mAbs. After lysis of erythrocytes and washing, stained cells were analysed with a fluorescence activated cell sorter Calibur flow cytometer using CellQuest software (BD Bioscience, Tokyo, Japan). TCR V $\beta$ expression was represented as a percentage of CD4<sup>+</sup> or CD8<sup>+</sup> cells for each family.

# Three-dimensional graphic display of TCR diversity

Qualitative alterations of TCR V $\beta$  repertoire obtained by CDR3 spectratyping were combined with the quantity of specific V $\beta^+$  CD4<sup>+</sup> and CD8<sup>+</sup> T cells for each V $\beta$  subfamily and plotted as landscape columns, as described previously [11,15].

# Results

# Skewed CDR3 size pattern in Hashimoto's thyroiditis but not Graves' disease

In healthy controls, the majority of V $\beta$  subfamilies exhibited a Gaussian curve with six peaks or more, reflecting a diverse TCR repertoire. Consistent with previous reports [16], CD8+ T cells exhibited a more skewed CDR3 profile than CD4+ T cells regardless of the presence of disease, probably because of age-related CD8<sup>+</sup> T cell clonal expansion (Fig. 1a-c). All 25 different V $\beta$  segments were amplified from CD4<sup>+</sup> and CD8<sup>+</sup> T cells obtained from each patient's sample. As shown in Fig. 1b and 1c, patients with Graves' disease showed a diverse distribution in the majority of their V $\beta$  subfamilies both in CD4<sup>+</sup> and CD8<sup>+</sup> T cells. In contrast, the frequency of skewed TCR V $\beta$  subfamilies of patients with Hashimoto's thyroiditis was significantly higher than that of Graves' disease and healthy controls in CD8+ T cells but not CD4+ T cells. No evidence was found for preferential skewing of particular V $\beta$  subfamilies in patients with Hashimoto's thyroiditis. Because none of the patients had clinical and laboratory evidence of acute infection at the time of sample collection, we conclude that these alterations reflect a stable state of the TCR repertoire in these patients.

Fig. 1. CDR3 spectratyping of T cell receptor (TCR) V $\beta$ . (a) CDR3 size distribution. Each TCR V $\beta$  fragment was amplified from cDNA with one of 25 V $\beta$ -specific primers. The size distribution of polymerase chain reaction (PCR) products was determined by an automated sequencer and GeneScan software. Shown are the results of CDR3 size distribution in CD4<sup>+</sup> and CD8<sup>+</sup> T cells from patient 16 of Hashimoto's thyroiditis. (b) CDR3 complexity scores. Complexity scores for each TCR V $\beta$ were shown. (c) Frequency of skewed TCR V $\beta$ repertoire. Shown are the mean (±standard deviation) numbers of skewed TCR V $\beta$ obtained from CD4+ and CD8+ T cells of the control and patients with Graves' disease and with Hashimoto's thyroiditis. GD, Graves' disease; HT, Hashimoto's thyroiditis. \*P < 0.05; \*\*P < 0.01.





Fig. 2. Frequency of skewed T cell receptor (TCR) V $\beta$  in subgroups of Hashimoto's thyroiditis. Shown are the mean (±standard deviation) numbers of skewed TCR V $\beta$  obtained from CD4<sup>+</sup> and CD8<sup>+</sup> T cells in each subgroup divided based on (a) age, (b) disease duration and (c) levothyroxine requirement of Hashimoto's thyroiditis. Tx, treatment.

To elucidate further the characteristics of skewing of TCR  $V\beta$  usage in Hashimoto's thyroiditis, those patients were divided into different subgroups based on age, disease duration and levothyroxine requirement (Fig. 2a-c). Although clonal T cell expansion can be seen in healthy individuals with age [16], no difference was observed between patients 14 years of age or younger and those older. In contrast, there were trends towards more skewing of CDR3 size distribution in CD8<sup>+</sup> T cells from patients having disease longer than 5 years and requiring replacement therapy of levothyroxine. Of note, six of nine patients with Hashimoto's thyroiditis who showed a more skewed pattern in CD8+ T cells (no. of skewed TCR V $\beta$ , 6.5  $\pm$  3.8) and having disease longer than 5 years required replacement therapy. Numbers of skewed TCR V $\beta$  did not correlate with the thyroid hormones levels or autoantibody titres in patients with Hashimoto's thyroiditis (data not shown).

#### Polyclonal TCR V $\beta$ repertoire by mAbs in both diseases

The relative TCR V $\beta$  usage of CD4<sup>+</sup> and CD8<sup>+</sup> T cells was also investigated using a panel of mAbs specific for 20 different V $\beta$  regions (Fig. 3a,b). None of the patients with Graves' disease and Hashimoto's thyroiditis exhibited a high percentage (> 30%) of T cells expressing a certain V $\beta$ . Although relative expansion of TCR V $\beta$  subfamilies (i.e. above the mean  $\pm$  2 standard deviations) was seen in both diseases, the frequency was not high either in CD4<sup>+</sup> or CD8<sup>+</sup> T cells, and predominance of expansions in CD8+ T cells was not seen, unlike in CDR3 spectratyping (Fig. 3b). In addition, flow cytometric analysis demonstrated that the VB subfamily having a skewed CDR3 size pattern in CD4+ and CD8+ T cells in a particular patient did not necessarily show relative expansion compared with controls. No significant differences in the percentage of peripheral CD3<sup>+</sup>, CD4<sup>+</sup> and CD8<sup>+</sup> T cells, the ratio of CD4<sup>+</sup> to CD8<sup>+</sup> T cells and the percentage expression of CD45RO+ memory marker in CD4+ and CD8+ T cells were found among Graves' disease, Hashimoto's thyroiditis and control subjects (data not shown).

# Combined qualitative and quantitative analysis

To assess the pattern of clonal CD8<sup>+</sup> T cell dominance in Hashimoto's thyroiditis in more detail, we used a combined analysis of TCR V $\beta$  repertoire distribution and CDR3 size distribution. Although such analysis provided a better visual

Fig. 3. Flow cytometric analysis of T cell receptor (TCR) V $\beta$ . (a) Expression profiles of TCR V $\beta$  subfamilies. Peripheral blood samples were stained with monoclonal antibodies (mAbs) for individual TCR V $\beta$  together with anti-CD4 and anti-CD8 mAbs. The percentage of each TCR V $\beta$  expression within CD4<sup>+</sup> and CD8<sup>+</sup> T cells was analysed by flow cytometry. (b) Frequency of expanded TCR V $\beta$  repertoire. Shown are the mean (±standard deviation) numbers of expanded TCR V $\beta$  obtained from CD4<sup>+</sup> and CD8<sup>+</sup> T cells of patients with Graves' disease and Hashimoto's thyroiditis.





Fig. 4. Combined display of flow cytometric analysis of T cell receptor (TCR) V $\beta$  and CDR3 spectratyping. Data from qualitative (CDR3 spectratyping) and quantitative (flow cytometric analysis for TCR V $\beta$ ) analysis are displayed visually as graphic landscape columns for CD4<sup>+</sup> and CD8<sup>+</sup> T cells.

representation, again there was no evidence for preferential expansion of particular V $\beta$  subfamilies (Fig. 4).

# Discussion

Early studies of intrathyroidal T lymphocytes from patients with Graves' disease and Hashimoto's thyroiditis demonstrated significant restriction in TCR V $\alpha$  but not in V $\beta$  gene usage [5,6]. Restricted usage of TCR V $\alpha$  as well as TCR V $\beta$  was also shown in T cells infiltrating extrathyroidal sites in addition to thyroid from Graves' disease [17]. On the contrary, other reports demonstrated no restriction of intrathyroidal TCR Vα subfamilies in Graves' disease [18–20]. In Hashimoto's thyroiditis, a marked restriction of V $\alpha$  usage in intrathyroidal CD8<sup>+</sup> T cell but not in CD4<sup>+</sup> T cell has been shown in a small-scale study [7]. These controversies over the presence of restricted TCR V $\alpha$  and V $\beta$  usage in T cells from patients with autoimmune thyroid diseases may be due to the different sensitivity of each method used for assessing the TCR repertoire. In addition, in most studies whole T lymphocytes but not isolated CD4+ and CD8+ T cells were utilized, resulting in possible underestimation of TCR restriction. On the other hand, no restriction has been demonstrated in peripheral blood T cells from patients with autoimmune thyroid diseases, even though intrathyroidal T cells showed restricted TCR Vα usage [5–7,17]. Because organ-specific autoimmune disorders may result from molecular mimicry between environmental antigens and self-antigens of target organs, thyroid-infiltrating lymphocytes are assumed to play a pathogenic role and are the most suitable source to assess TCR restriction in autoimmune thyroid diseases. However, the availability of such samples may be limited because thyroid

fine-needle aspiration and biopsies are performed rarely in those patients. Recent advances in molecular and cellular techniques analysing the TCR V $\beta$  repertoire may allow us to obtain a reliable assessment of autoimmune conditions in peripheral blood [21,22]. We therefore set out to study in more detail the possible presence of TCR V $\beta$  restriction among circulating CD4<sup>+</sup> and CD8<sup>+</sup> T cells from patients with autoimmune thyroid diseases.

In the present study we demonstrated significantly higher skewing of TCR V $\beta$  in peripheral CD8<sup>+</sup> T cells by using CDR3 spectratyping among patients with Hashimoto's thyroiditis compared with healthy adults. Interestingly, the patients requiring thyroid hormone replacement who may have suffered from more severe autoimmune attacks on the thyroid gland tended to have more skewing of CDR3 size distribution than those without replacement therapy, which supports the idea that these TCR alterations in peripheral CD8<sup>+</sup> T cells may be reflected in the local events and may be involved in the development and maintenance of chronic thyroiditis. Similar results were obtained from the study of the TCR V $\beta$  repertoire by using CDR3 spectratyping of patients affected with myasthenia gravis, another organspecific autoimmune disorder, where persistent TCR expansions in peripheral blood T cells correlated with clinical severity [21]. Moreover, patients with disease longer than 5 years also tended towards more skewing of TCR VB compared with those with shorter disease duration in our crosssectional study. These findings contrast with the hypothesis of Davies et al. [23], that TCR restrictions seen in the thyroid aspirates from patients with autoimmune thyroid diseases represent an early disease marker. However, autoantigens may be sustained in the thyroid gland and cause chronic antigenic stimulation, probably resulting in the accumulation of antigen-specific T cell clones. A longitudinal examination will be needed to explain the persistence and progression of TCR restriction in Hashimoto's thyroiditis. In contrast, flow cytometric analysis for TCR V $\beta$  repertoire failed to show expansion or deletion of particular TCR V $\beta$ either in CD4<sup>+</sup> or CD8<sup>+</sup> T cells. Because the circulating pathogenic T cells in a large pool of peripheral T cells may be very small, it is reasonable to assume that they seldom reach the detection threshold of flow cytometric analysis.

It has been demonstrated recently that interferon- $\gamma$ inducible chemokines such as CXC chemokine ligand 10 (CXCL10) play an important role in the initial stage of autoimmune disorders involving endocrine gland [24]. Indeed, elevated levels of circulating CXCL10 have been reported in autoimmune thyroid diseases [25]. Both lymphocytes and thyroid follicular cells may account for production of CXCL10 [24,26]. Although we had no data concerning chemokines, it is tempting to speculate that CD8<sup>+</sup> T cells with skewed TCR V $\beta$  might be associated with secretion of CXCL10. Further studies are necessary in order to address these important issues.

In Graves' disease, CDR3 spectratyping and flow cytometric analysis of TCR V $\beta$  repertoire demonstrated no restricted TCR V $\beta$  usage either in peripheral CD4<sup>+</sup> or CD8<sup>+</sup> T cells regardless of the severity, duration of the disease or levels of the thyroid hormones. Although our methods were sensitive enough to detect clonal expansion in peripheral T cells in Hashimoto's thyroiditis, the possibility that specific antigendriven T cells were mildly expanded in the thyroid gland but undetectable in the circulation of patients with Graves' disease because of sensitivity problems cannot be ruled out. Alternatively, the absence of restricted TCR V $\beta$  usage could be ascribed to anti-thyroid drugs themselves, as the drugs have immunomodulant actions [27]. Further studies assessing TCR Vβ repertoire in intrathyroidal and peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T cells simultaneously will be useful to define the presence of clonal dominance in Graves' disease.

In summary, our studies provide evidence for clonal expansion of peripheral CD8<sup>+</sup> T cells from patients with Hashimoto's thyroiditis. Characterization of the clonal composition of such CD8<sup>+</sup> T cells that may be associated with cell-mediated autoimmune attacks on the thyroid gland will provide important insights into the pathogenesis of the disease and deserve close investigation.

#### Acknowledgements

We thank Ms Harumi Matsukawa and Ms Shizu Kouraba for excellent technical assistance, Dr Akira Asai for clinical samples and Dr Yoshihito Kasahara for insightful discussion. This work was supported by a Grant-in-Aid for Scientific Research from the Ministry of Education, Culture, Sports, Science and Technology of Japan and a grant from the Ministry of Health, Labour and Welfare of Japan, Tokyo.

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