

Impaired expression of MHC class II molecules in response to interferon-gamma (IFN- γ) on human thymoma neoplastic epithelial cells

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SUMMARY

A human thymoma is a neoplasm derived from the thymic epithelial cell, and is well known for its association with autoimmune diseases, especially myasthenia gravis. The neoplastic epithelial cells of thymoma clearly retain thymic epithelial functions, but the development of T cells in thymoma is somewhat impaired. In this study, we quantified by flow cytometry the *in vitro* expression of MHC molecules on neoplastic epithelial cells precultured with IFN- γ . While MHC class I expression was comparable with that on normal thymic epithelial cells, the level of MHC class II molecules on neoplastic epithelial cells was lower than in controls, and also varied greatly from case to case. Additionally, there was a significant positive correlation between the expression level of MHC class II and the proportion of mature CD3⁺ cells in the CD4⁺CD8⁻ subset. Thus, accumulation of CD3⁻CD4⁺CD8⁻ cells in thymoma may result from impaired expression of the MHC class II molecules, suggesting that the function of the neoplastic epithelial cells might determine the maturation and the positively selected repertoire of T cells in thymomas.

Keywords thymoma thymic epithelial cell MHC HLA-DR T cell differentiation

INTRODUCTION

A human thymoma is a neoplasm derived from thymic epithelial cells [1], and is well known for its association with autoimmune diseases, especially with myasthenia gravis. One of its histological characteristics is the coexistence of large numbers of lymphocytes. The ratio of lymphocytes to neoplastic epithelial cells varies greatly from tumour to tumour, thus thymomas are classified as predominantly lymphocytic, predominantly epithelial, or mixed type [2,3]. The lymphocytes in thymomas are of T cell lineage [4-6], and a considerable proportion are CD4⁺CD8⁺, so called double-positive cells, exactly as in the normal thymic cortex [7-9]. In the murine and the human thymus, CD3⁻CD4⁻CD8⁻ cells differentiate to CD3⁺CD4⁺CD8⁻ or CD3⁺CD4⁻CD8⁺ cells via CD4⁺CD8⁺ cells. Recently, we have shown that the neoplastic epithelial cells from thymoma were able to induce the differentiation of CD4⁺CD8⁺ cells from CD3⁻CD4⁻CD8⁻CD34⁺ T cell precursors in an *in vitro* culture [10]. Furthermore, mature CD4 single-positive cells expressing CD69 have also been found in

thymoma and shown to be capable of proliferating in response to a mitogen [11]. These observations suggest that T cell development takes place even in the neoplastic environment of a thymoma.

We and another group have identified an intermediate CD3⁻CD4⁺CD8⁻ stage, so called immature CD4 single-positive cells, between the CD3⁻CD4⁻CD8⁻ and the CD3⁻CD4⁺CD8⁺ stages [12,13]. Interestingly, these CD3⁻CD4⁺CD8⁻ cells were shown to accumulate more in thymomas than in normal thymus [14,15], and did not express CD69 [11], a marker of positive selection of thymocytes, suggesting that T cell development is somewhat impaired in thymomas.

Thymic stromal cells are known to govern T cell development. The interaction between the T cell antigen receptor (TCR) on thymocytes and MHC antigen on thymic epithelium is believed to play an important role in selection of mature thymocytes [16-18]. Previous immunohistochemical analyses have demonstrated a reduced expression of MHC class II molecules on thymoma epithelial cells [19-22]. Thus, it seems reasonable to speculate that the abnormal T cell development in thymomas might be accompanied by the altered selection of newly generated T cells, and this in turn might lead to autoimmune disease [15]. To examine the relation between the function of the neoplastic epithelial cells and T cell development in thymomas, we quantitatively evaluated

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the surface marker expression of lymphocytes and the expression level of the MHC molecules on neoplastic epithelial cells by flow cytometry. The results of this study clearly demonstrate a correlation between the expression level of MHC class II molecules and the maturation of CD4⁺CD8⁻ cells in the tumour.

PATIENTS AND METHODS

Patients

The characteristics of the thymoma patients studied here are shown in Table 1. All of the cases were classified according to the clinicopathological staging system of Masaoka [23]. None of these patients had been treated with corticosteroids, induction chemotherapy or preoperative irradiation. Various autoantibodies, including anti-acetylcholine receptor antibody (α AChR), were examined in these patients. Although α AChR was positively detected in cases 2, 5, 6 and 7, myasthenic symptoms were found only in case 2. As controls, normal thymi were obtained from 21 children during open cardiac surgery under written informed consent when resection of a part of the thymus was necessary to perform the surgical procedures. The mean age of the control patients was 2.2 ± 4.2 years.

Monoclonal antibodies

For the analyses of lymphocytes, FITC-conjugated control mouse IgG and anti-CD3, PE-conjugated anti- κ -chain and anti-CD4, and biotin-conjugated anti-CD8 antibodies were purchased from Becton Dickinson (San Jose, CA). For the analyses of stromal cells, anti-HLA-ABC antibody was purchased from Pharmingen (San Diego, CA). Anti-HLA-DP/DQ/DR (CR3/43), anti-cytokeratin (MNF116) and FITC-conjugated anti-cytokeratin (MNF116) antibodies were purchased from Dako (Glostrup, Denmark).

Histological evaluation

Surgical specimens were fixed in 10% formaldehyde and then paraffin-embedded. They were classified as predominantly lymphocytic, predominantly epithelial, or mixed type according to the ratio of lymphocytes to epithelial cells by haematoxylin-eosin staining. By definition, the ratio of lymphocytes to epithelial cells was at least 2:1 in the predominantly lymphocytic thymomas and below 1:2 in the predominantly epithelial thymomas. Intermediate cases were classified as the mixed type.

Table 1. Characteristics of thymoma patients

Case	Age (years)	Sex	Stage	Type	MG	Positive autoantibodies
1	37	F	I	Ep		RF
2	28	M	IV	Mix	(+)	α AChR
3	51	M	II	Ly		TGHA, MCHA
4	37	M	II	Ly		
5	53	F	II	Mix		α AChR
6	42	M	III	Ep		α AChR
7	49	F	II	Mix		α AChR, α DNA
8	56	M	II	Mix		ANF
9	62	M	II	Mix		

Type, histological type; Ep, predominantly epithelial cell type; Ly, predominantly lymphocytic type; mix, mixed type; MG, myasthenia gravis; RF, rheumatoid factor; α AChR, anti-acetylcholine receptor antibody; TGHA, thyroglobulin haemagglutination; MCHA, microsome haemagglutination; α DNA, anti-DNA antibody; ANF, anti-nuclear factor.

Preparation of lymphocytes

Lymphocytes in the thymoma or thymus were isolated by mechanical teasing and pressing the tissue against stainless steel mesh. The viable lymphocytes were separated on a Ficoll-Hypaque (Lymphoprep; Nycomed, Oslo, Norway) density gradient to remove any erythrocytes and dead cells. These isolated lymphocytes were then counted and applied to the flow cytometric analyses.

Preparation of stromal cells

Stromal cells were isolated from the thymoma or normal child thymus tissue by 2-deoxyguanosine (2-dGuo; Sigma, St Louis, MO) treatment and trypsin-EDTA dispersion [24,25]. Briefly, fresh thymoma or thymus tissue was minced and the fragments were washed in PBS repeatedly. The tissue fragments were cultured for 5 days in the presence of 1.35 mM 2-dGuo in complete medium (RPMI 1640 in 10% fetal calf serum (FCS), 2 mM glutamine, 5 μ M 2-mercaptoethanol) to deplete the lymphocytes. Five days later, the 2-dGuo-treated fragments were rinsed with Ca²⁺Mg²⁺-free PBS and incubated in 0.25% trypsin-0.02% EDTA for 30 min at 37°C. The digested fragments were suspended vigorously, and trypsinization was stopped by the addition of an equal volume of 10% FCS-RPMI. After the dispersed cells were cultured for 4 days in complete medium, the adherent stromal cells were removed, followed by a further culture for 2 days in the presence of IFN- γ (1000 U/ml) in complete medium. The adherent cells were collected by vigorous suspension with 0.02% cold EDTA and then washed and applied to flow cytometric analysis.

Flow cytometry

For staining surface antigens on lymphocytes, 1×10^6 lymphocytes suspended in 300 μ l PBS were mixed with a combination of 5 μ l each of FITC-conjugated control mouse IgG or anti-CD3, PE-conjugated anti-CD4, and biotin-conjugated anti-CD8 antibodies. After incubation for 30 min at 4°C, the cells were washed and suspended in 300 μ l PBS. One microlitre of streptavidin-red 670 (Life Technologies, GIBCO BRL, Gaithersburg, MD) was added for three-colour flow cytometry. After incubation for 30 min at 4°C, the cells were washed and subjected to flow cytometric analysis. The cytometer used in this study was a FACScan (Becton Dickinson); 1×10^5 events were collected and analysed by the Cell Quest program.

For analysis of the stromal cells, 1×10^5 stromal cells suspended in 200 μ l PBS were mixed with 5 μ l anti-HLA-ABC or HLA-DP/DQ/DR MoAbs. After incubation for 30 min at 4°C, the cells were washed and suspended in 200 μ l PBS. PE-conjugated anti- κ -chain antibody (5 μ l) was added as a secondary antibody. After incubation for 30 min at 4°C, the cells were washed and suspended in 200 μ l PBS. The suspended cells were blocked with normal mouse serum for 10 min and fixed with 0.25% formaldehyde for 30 min at room temperature, washed and suspended in 200 μ l PBS. After permeabilization of the cells with 0.025% saponin (Sigma), 5 μ l of FITC conjugated anti-cytokeratin antibody were added and incubated for 30 min at 4°C; 3×10^4 events were collected and analysed by the Cell Quest program. The MHC expression level was judged by the *D* value, with the *p* value in Kolmogorov-Smirnov statistics. Pearson's correlation coefficient was calculated to examine the relationship between two variables.

RESULTS

Phenotypic characterization of the lymphocytes in thymomas

The result of the characterization of lymphocytes in thymomas is

Table 2. Phenotypic characterization of the lymphocytes in thymomas and normal infant thymi

	DP	4SP	8SP	3 ⁺ /4SP	3 ⁺ /8SP
Case 1	60	17	13	70	99
Case 2	63	19	14	60	100
Case 3	73	18	4	18	96
Case 4	56	29	9	35	100
Case 5	65	24	5	12	99
Case 6	44	39	6	48	97
Case 7	64	23	7	10	100
Case 8	69	24	2	22	96
Case 9	59	18	13	30	92
Thymoma	61.4 ± 8.3	23.4 ± 7.0	8.1 ± 4.4	33.9 ± 21.3	97.7 ± 2.7
Thymi (n = 17)	80.5 ± 4.2	9.6 ± 4.2	5.6 ± 2.1	88.8 ± 6.1	98.5 ± 1.8

The number indicates the percentage of cells in each population.

Results of all are shown as mean ± s.d.

DP, CD4⁺CD8⁺ cell; 4SP, CD4⁺CD8⁻ cell; 8SP, CD4⁻CD8⁺ cell; 3⁺/4SP, CD3⁺ cells among CD4⁺CD8⁻ cells; 3⁺/8SP, CD3⁺ cells among CD4⁻CD8⁺ cells.

shown in Table 2. CD4⁺CD8⁺ cells accounted for 61.4 ± 8.3% of all lymphocytes on average in nine thymomas. The mean proportions of CD4⁺CD8⁻ cells and CD4⁻CD8⁺ cells were 23.4 ± 7.0% and 8.1 ± 4.4%, respectively. In 17 normal child thymi, CD4⁺CD8⁺ cells accounted for 80.5 ± 4.2% of all lymphocytes on average. An example of the CD3 expression in each subset is shown in Fig. 1. In this case, the CD4⁻CD8⁺ subset consisted almost exclusively of mature CD3⁺ cells. In contrast, the proportion of mature CD3⁺ cells in the CD4⁺CD8⁻ subset was only 48%, and, typically, CD3-negative and -positive peaks were observed in

the CD4⁺CD8⁻ subset. This bimodal pattern is not usually seen in normal thymocytes. As shown in Table 2, a major proportion of the lymphocytes in the thymomas had a phenotype of cortical thymocytes, and immature CD3⁻CD4⁺CD8⁻ cells accumulated compared with the normal thymus samples, which is consistent with our previous report [14].

MHC expression on the neoplastic epithelial cells in vitro

MHC expression on the neoplastic epithelial cells *in vitro* in response to exogenously administered IFN-γ was examined by flow cytometry. Using two-colour flow cytometry, the epithelial cells from the entire sample of stromal cells were identified by anti-cytokeratin antibody and gated as cytokeratin-positive cells. Four normal child thymi were examined as positive controls. Figure 2a shows the HLA-ABC (MHC class I) expression and the percentage of CD3⁺ cells among CD4⁻CD8⁺ cells of a normal thymus and thymomas. MHC class I was distinctly expressed and most CD4⁻CD8⁺ cells expressed CD3 in these cases. The results of all cases are shown in Fig. 2b. Thus, MHC class I expression of thymoma epithelial cells was comparable to normal thymus. By contrast, HLA-DP/DQ/DR (MHC class II) expression and the percentage of CD3⁺ cells among CD4⁺CD8⁻ cells of a normal thymus and typical thymoma cases are shown in Fig. 3a. MHC class II expression on thymoma epithelial cells varied greatly from case to case. The results of all cases are shown in Fig. 3b. MHC class II expression on neoplastic thymic epithelial cells was significantly lower than normal infant thymus.

Relationship between HLA-DP/DQ/DR expression and the maturity of the CD4⁺CD8⁻ subset

Since there was a wide variation in the proportion of CD3⁺ cells among the CD4⁺CD8⁻ subset as well as in MHC class II

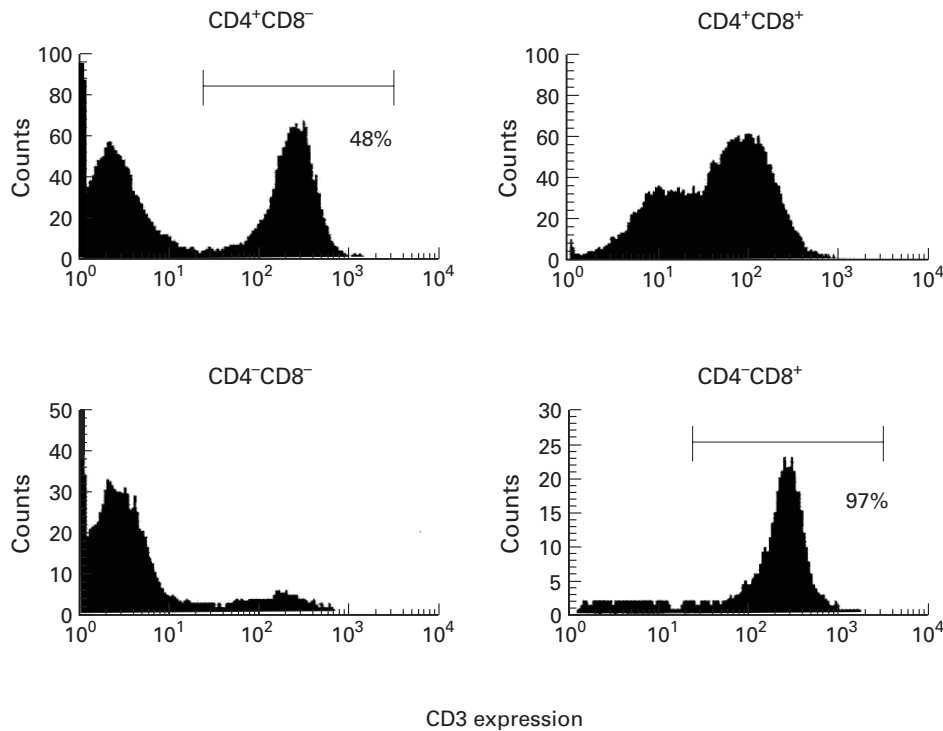


Fig. 1. A representative result of the CD3 expression in each subset defined by CD4 and CD8 expressions (case 6).

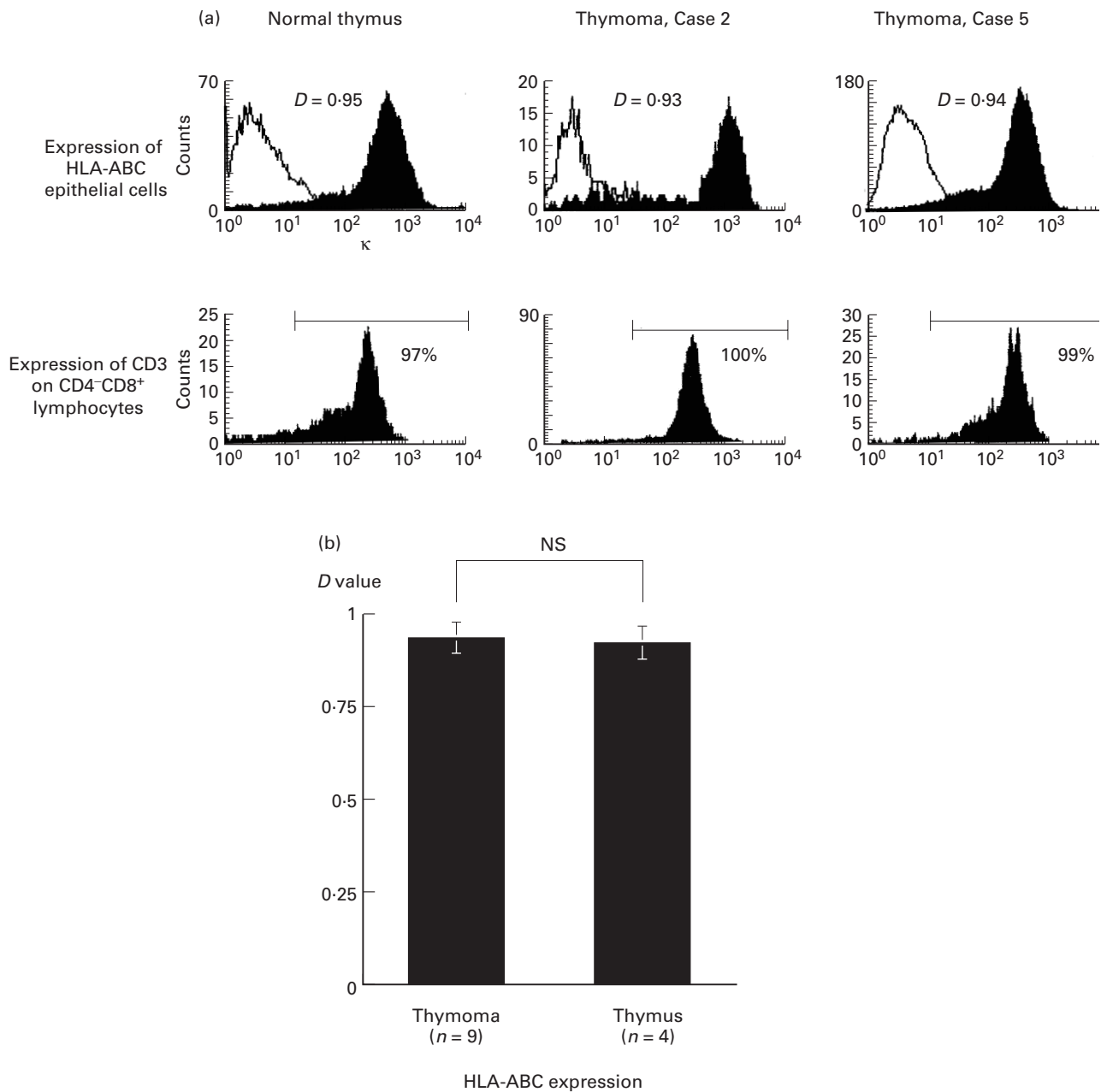


Fig. 2. (a) HLA-ABC (MHC class I) expression on epithelial cells (upper) and CD3 expression among CD4⁺CD8⁺ lymphocytes (lower) in typical cases. Solid lines show the negative control stains. The expression level was judged by *D* value in Kolmogorov–Smirnov statistics for MHC and the percentage for CD3. A normal thymus is presented as a positive control. (b) HLA-ABC expression of all thymoma cases and normal thymi. Thymoma: $D = 0.93 \pm 0.05$ versus normal thymus: $D = 0.92 \pm 0.05$, not significant (NS).

expression on neoplastic epithelial cells, we examined the relationship between these two variables. Interestingly, there was a significant positive correlation between MHC class II expression and the proportion of CD3⁺ cells among the CD4⁺CD8⁺ subset, as shown in Fig. 4. The patients with autoantibodies are indicated with open circles and there is no definitive tendency.

DISCUSSION

The subset of CD3⁺CD4⁺CD8⁺ cells has recently been recognized as an intermediate population between CD3⁺CD4⁺CD8⁺ and

CD4⁺CD8⁺ cells during T cell development in the thymus [12,13]. One of the characteristic features of lymphocytes in thymoma is a prominent accumulation of these immature CD4 single-positive cells, compared with a normal thymus [14,15]. These observations seem to indicate inefficient differentiation of T cells from CD3⁺CD4⁺CD8⁺ cells to CD4⁺CD8⁺ cells, or from CD4⁺CD8⁺ cells to CD3⁺CD4⁺CD8⁺ cells in thymomas. Because MHC molecules have an important role in T cell development in the thymus, we quantified their expression level on the neoplastic epithelial cells in response to exogenous IFN- γ . A preliminary experiment revealed that MHC class II expression

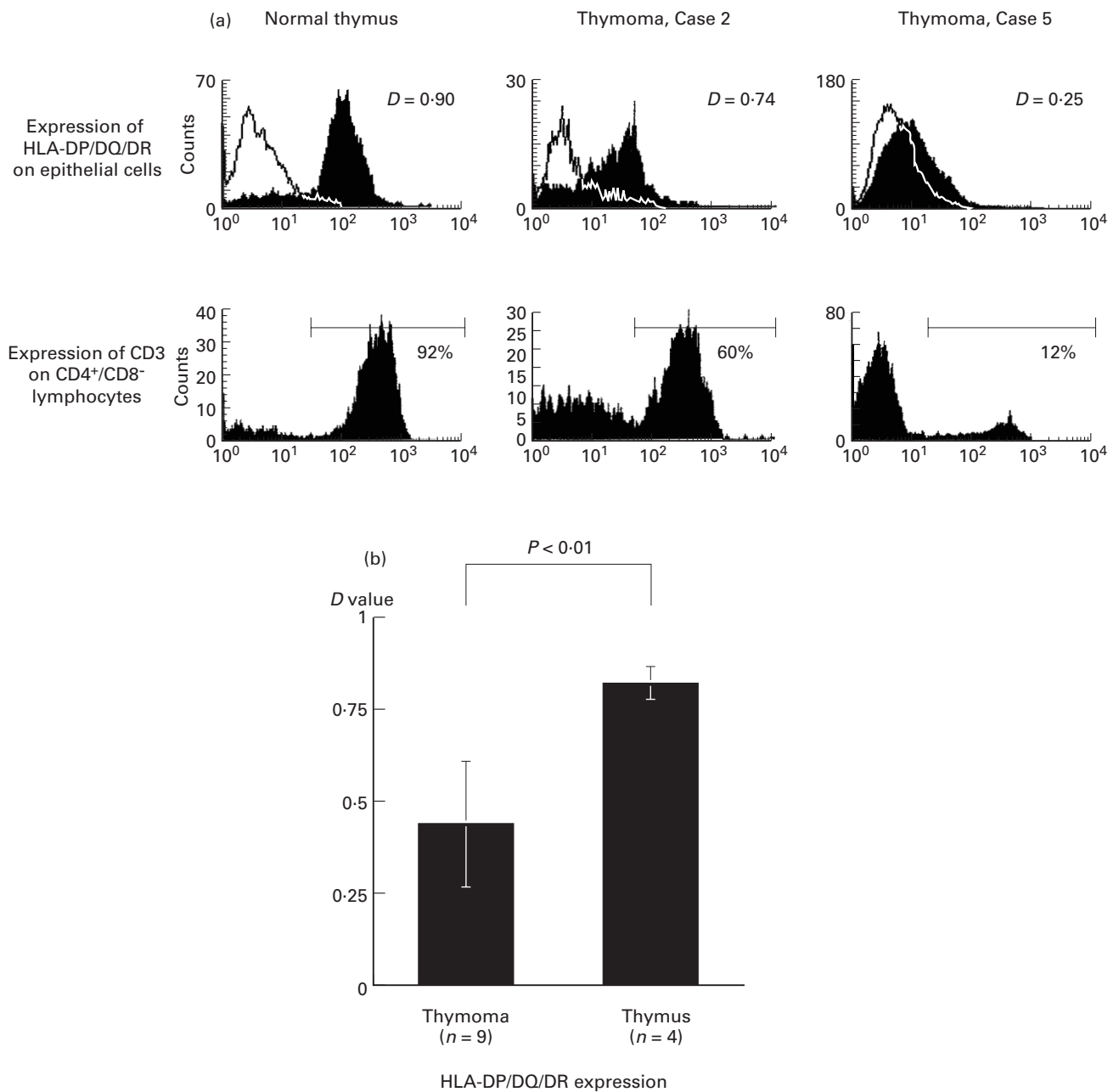


Fig. 3. (a) HLA-DP/DQ/DR (MHC class II) expression on epithelial cells (upper) and CD3 expression among CD4⁺CD8⁻ lymphocytes (lower) in typical cases. Solid lines show the negative control stains. The expression level was judged by *D* value in Kolmogorov–Smirnov statistics for MHC and the percentage for CD3. A normal thymus is presented as a positive control. (b) HLA-DP/DQ/DR expression of all thymoma cases and normal thymi. Thymoma: $D = 0.44 \pm 0.17$ versus normal thymus: $D = 0.82 \pm 0.08$; $P < 0.01$, significant by unpaired *t*-test.

on the neoplastic epithelial cells *in vitro* was reduced without stimulation by IFN- γ and restored with an administration of exogenous IFN- γ .

In our previous study, we evaluated MHC class II expression by immunohistochemistry and could not find a distinct difference [14]. MHC class II expression of all cases was also examined by immunohistochemistry in this study (data not shown), but we could not histologically detect a significant difference between case 2, which showed the highest expression of MHC class II, and case 5,

which showed the lowest *in vitro* by flow cytometric analysis. A proportion, but not all, of the neoplastic epithelial cells was positively stained with anti-HLA-DP/DQ/DR antibody in both of these cases, suggesting a heterogeneous expression of MHC class II molecules. Thus, it is difficult to analyse MHC class II expression in thymoma quantitatively using immunohistochemistry.

Although MHC class I molecules were expressed on the thymoma neoplastic epithelial cells at a level comparable to that on normal thymic epithelial cells, expression of MHC class II

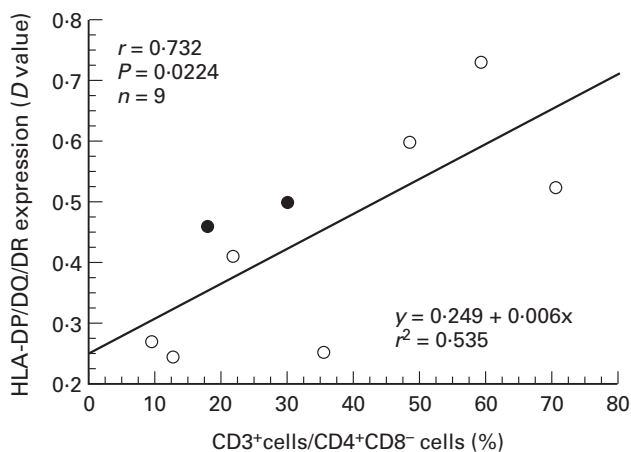


Fig. 4. The relationship between HLA-DP/DQ/DR expression and the proportion of CD3⁺ cells among the CD4⁺CD8⁻ subset. ○, Patients with positive autoantibodies.

molecules was generally decreased and varied from case to case, consistent with previous reports [19–22]. In addition, interestingly, there was a significant correlation between the expression of MHC class II molecules and the proportion of CD3⁺ cells among the CD4⁺CD8⁻ subset in a thymoma. This result suggests that the proportion of CD3⁺ cells in the CD4⁺CD8⁻ subset reflects the expression level of MHC class II molecules on the thymoma neoplastic epithelial cells.

Recently, an affinity-avidity model was proposed to explain the mechanism of positive and negative selection processes of the T cell repertoire, based on genetically engineered murine models which enabled manipulation of the expression level of the MHC molecules [26–30]. According to this hypothesis, T cell that would normally be deleted by the normal level of MHC class II on the thymic epithelial cells might be positively selected when this level is reduced on the neoplastic epithelial cells. Thus, the T cells newly generated in a thymoma might include an altered TCR repertoire which can react to self-antigens and therefore might cause autoimmune diseases. This hypothesis would be supported if low MHC class II expression on thymoma epithelial cells correlated with the autoimmune associations. Since the presence of AChR-related antigen was reported, it is possible that AChR-specific T cells might be positively selected in the tumour [31]. Also, there must be many non-myasthenic patients with an elevated α AChR. However, we suppose that this autoimmunity is not restricted to neuromyopathy, because thymoma patients have various autoantibodies, as shown in Table 1.

Thymoma lacks a medullary structure and there may be insufficient contact with bone marrow-derived dendritic cells and macrophages that are believed to serve as the negatively selecting elements for autoreactive T cells generated in normal thymus. It has been reported that mice whose MHC class II antigen is expressed in thymic cortex only, and *relB*-deficient mice which lack thymic medulla, show impairment of thymic negative selection [32,33]. When the single-positive T cells are generated in the thymoma and exported to the periphery, they may contain autoreactive T cells.

Moreover, cell–cell interactions between thymocytes and thymic stromal cells through several adhesion molecules, such as vascular cell adhesion molecule-1 (VCAM-1)/fibronectin (FN)

to very late antigen-4,5 (VLA-4,5) or intercellular adhesion molecule-1 (ICAM-1) to leucocyte function-associated antigen-1 (LFA-1), have been shown to be involved in T cell development in the thymus [34]. Soluble factors such as IL-7 also participate in T cell development in thymus [35,36]. We cannot exclude defects in these accessory factors.

The signal transduction cascade of the IFN- γ receptor has been studied extensively, and Jak-1, Jak-2, Stat-1, and CIITA have been shown to be involved in the induction of MHC class II molecules [37,38]. The existence of a CIITA-independent class II regulatory pathway has also been reported [39]. Experiments to identify the molecules responsible for impairment of the IFN- γ receptor signal transduction cascade in thymoma neoplastic epithelial cells are currently being conducted in our laboratory.

In conclusion, MHC class II expression on neoplastic epithelial cells was generally lower than that on normal thymic epithelial cells *in vitro*. The level of MHC class II expression had a significant correlation with the proportion of CD3⁺ cells among the CD4⁺CD8⁻ subset. These results suggest that the degree of maturation of the CD4⁺CD8⁻ subset reflects the activity of the neoplastic cells in a thymoma. We suppose that the immaturity of the CD4⁺CD8⁻ subset in thymomas is caused by the lower expression of MHC class II molecules on neoplastic epithelial cells. It is expected that this impaired expression of MHC class II molecules on thymoma neoplastic epithelial cells might be a factor of pathogenesis of thymoma-associated autoimmune diseases [15]. Further investigation of the mechanism may elucidate the molecular pathogenesis of thymoma-associated autoimmune diseases.

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