

and BDCA-2, respectively, and mDCs are identified as BDCA-1⁺ and CD19⁻ cells (10, 11). Langerin is a C-type lectin expressed in LCs from the skin and the small intestine, and Langerin⁺ LCs are involved in immunological surveillance at the surface (12).

In the periphery, immature DCs such as LCs capture antigens and, under the influence of inflammatory stimuli, subsequently migrate. LCs represents a population of DCs only found in epithelia. Although several chemokines have been reported to attract immature DCs *in vitro*, macrophage inflammatory protein-3 α (MIP-3 α , also known as CCL20) is the main chemokine expressed in epithelium and recruiting LCs to mucosal surfaces (13). Indeed, it has been demonstrated recently that the receptor for MIP-3 α , namely CCR6, is expressed in LCs (13, 14).

Although DCs surely play an important role in the pathogenesis of the cholangiopathy in PBC, more studies are needed to explain the bile duct-specific autoimmunity. The aim of this study is to characterize periductal DCs in PBC and to clarify the interaction between BECs and DCs in terms of the periductal cytokine milieu and biliary innate immunity. This study may open avenues for understanding and eventually the control of the specific functions of biliary DCs.

Materials and methods

Patients and tissue preparation

A total of 72 needle or wedge liver biopsy specimens were obtained from the patients with PBC and the controls including chronic viral hepatitis (CVH) associated with hepatitis C virus (HCV), hepatic sarcoidosis, drug-induced liver injury and no significant histopathological changes (so-called histologically 'normal liver') (Table 1). Liver tissue specimens were fixed in 10% neutral-buffer formalin. The histopathological diagnosis was established by a histological review by at least two pathologists with consideration of clinical and laboratory data. All PBC patients were examined before ursodeoxycholic acid therapy, and AMA was detected by immunofluorescence microscopy in all PBC cases. The diagnosis of drug-induced liver injury was clinicopathologically performed.

Table 1. Hepatobiliary diseases examined in this study

Patients	Case number	Average age (years)	Male/female	Histological stage
Primary biliary cirrhosis	26	59	2/24	I/II/III = 14/9/3
HCV-related chronic hepatitis	18	62	6/12	F1/F2/F3 = 6/8/4
Hepatic sarcoidosis	6	52	3/3	
Drug-induced liver injury	14	49	5/9	
Normal liver	8	60	3/5	

HCV, hepatitis C virus.

Benzbromarone, tiquizium bromide, analgesic agent, Chinese medicines (two cases), protein and health food were speculated as a causative drug by lymphocyte stimulation test in seven patients. According to Councils for International Organizations of Medical Sciences/Roussel Uclaf Causality Assessment Method (15) scoring system, 'unlikely', 'probable' and 'possible' were two, five and seven cases respectively. Liver histology also confirmed specific findings such as perivenular zonal necrosis, cholestasis, granulomatous changes and eosinophil/neutrophil infiltration, suggesting drug-induced liver injury.

More than 10 sections, 3 μ m thick, were prepared from each paraffin block; three of them were stained with haematoxylin and eosin, Gomori's reticulum and orcein stain for histological diagnosis and evaluation. The remaining sections were processed for immunohistochemistry. In addition to fixed specimens, fresh surgical specimens were available in seven cases each of PBC and HCV-related CVH, and used for immunohistochemistry. Informed consent to conduct research was obtained from these patients. This study was approved by the Kanazawa University Ethics Committee.

Immunohistochemistry and immunocytochemistry

After deparaffinization, the sections were incubated in Target Retrieval Solution (Dako Japan, Tokyo, Japan) at 95 °C for 20 min for the pretreatment of tissue before immunostaining. Following endogenous peroxidase blocking in methanolic-hydrogen peroxide for 20 min and incubation in normal goat serum (diluted 1:10; Vector Lab, Burlingame, CA, USA) for 20 min, the sections were incubated at 4 °C overnight with a primary mouse monoclonal antibody against human Langerin (clone 12D6; 2.5 μ g/ml; Abcam Japan, Tokyo, Japan), human BDCA-2 (clone 104C12.08; 5 μ g/ml; AbCys S.A., Paris, France) or human MIP-3 α (clone 308B7.06; 5 μ g/ml; Dendritics S.A., Lyon, France) and then Dako Envision-HRP or the CSA system (Dako Japan) was used. After the benzidine reaction, sections were weakly counterstained with haematoxylin. No positive staining was obtained when the primary monoclonal antibody was replaced with an isotype-matched, non-immunized immunoglobulin as a negative control of the staining procedures.

The simultaneous detection of Langerin and CCR6 was evaluated by double immunohistochemical staining. After staining for Langerin using a benzidine reaction, CCR6 staining was performed using human CCR6 antibody (rabbit polyclonal, 5 μ g/ml, MBL, Nagoya, Japan) and HistoGreen (AbCys S.A.). Moreover, the simultaneous detection of BDCA-1 and CD19 was assessed using frozen sections. After fixation of frozen 5 μ m sections in cold acetone for 10 min and the blocking of endogenous peroxidase, human CD19 antibody (clone HD37, 2 μ g/ml, Dako Japan) was added and then Dako Envision-HRP was used. After the benzidine reaction, BDCA-1 staining

was performed using human BDCA-1 antibody (clone L161; 5 µg/ml, GeneTex, Irvine, CA, USA), Histofine simple stain-AP (Nichirei Bioscience, Tokyo, Japan) and Vector Blue (Vector Lab).

Histological examination

The BDCA-2⁺ cells, BDCA-1⁺ and CD19⁻ cells, and Langerin⁺ cells were evaluated as pDCs, mDCs and LCs respectively. In a primary survey, we examined all portal tracts in each specimen and, for counting DCs, selected one or two representative portal tracts containing interlobular bile ducts (16) and expressed the results as the number of DCs per bile duct. For MIP-3α, the immunoreactivity in bile ducts was semiquantitatively graded as follows: 0, absence of expression; 1, weak constitutive expression; 2, moderate expression; and 3, strong expression. The final indices of PBC and the controls were defined as the mean of individual cases. In PBC, some interlobular bile ducts appeared normal, while others showed variable biliary epithelial damage including CNSDC. Therefore, the former and the latter were evaluated separately as normal and damaged bile ducts, respectively, in this study.

Cultured human biliary epithelial cells

Two cultured human BEC lines that were isolated from the explanted livers of two PBC patients were established in this study. These lines were incubated with a culture medium composed of D-MEM/F-12, Nu-Serum (Becton Dickinson, Bedford, MA, USA), ITS+ (Becton Dickinson), 5 µM forskolin (Wako, Osaka, Japan), 12.5 mg/ml of bovine pituitary extract (Gibco, Rockville, MD, USA), 1 µM dexamethasone (Sigma, St Louis, MO, USA), 5 µM triiodo-thyronine (Sigma), 5 mg/ml of glucose (Sigma), 25 mM sodium bicarbonate (Sigma), 1% antibiotics antimycotic, 20 ng/ml of human epidermal growth factor (Gibco) and 10 ng/ml of human hepatocyte growth factor (Gibco) (17). These cell lines had been confirmed to be BECs by the expression of a biliary-type cytokeratin, CK7. BECs were used between passages 6 and 10 for this study.

Treatment with cytokines and pathogen-associated molecular patterns

Cultured BECs were stimulated with recombinant human cytokines including interleukin (IL)-1β, IL-4, IL-6, IL-17, tumour necrosis factor (TNF)-α and interferon (IFN)-γ (1000 U/ml, PeproTech, London, UK) and TLR-related pathogen-associated molecular patterns (PAMPs) including Pam3CSK4 (TLR1/2 ligand, 100 ng/ml, InvivoGen, San Diego, CA, USA), polyinosinic-polycytidylic acid [poly(I:C), a synthetic analogue of viral dsRNA, TLR3 ligand, 25 µg/ml, InvivoGen] or lipopolysaccharide (LPS) (ultrapure grade, TLR4 ligand, 1 µg/ml, InvivoGen) for 3 and 24 h and used for the preparation of samples for

polymerase chain reaction (PCR) analysis and immunocytochemistry respectively.

Isolation of RNA and real-time polymerase chain reaction

For the evaluation of MIP-3α mRNA, total RNA was extracted from cultured BECs using the RNeasy Total RNA System (Qiagen, Hilden, Germany) following the manufacturer's instructions. Then, 1 µg of total RNA was reverse-transcribed with an oligo-(dT) primer and ReverTra Ace (Toyobo, Osaka, Japan) to synthesize a cDNA template for PCR. To carry out relative quantification, real-time quantitative PCR was performed for measurements according to a standard protocol using the Brilliant II SYBR Green QPCR Reagents and Mx300P QPCR system (Stratagene Japan, Tokyo, Japan) and the relative gene expression was calculated using the comparative cycle threshold method. Specific primers were as follows: MIP-3α forward, 5'-GCGCAAATCCAAAACAGACT-3', and reverse, 5'-CAAGTCCAGTGAGGCACAAA-3', and glyceraldehyde 3 phosphate dehydrogenase (GAPDH, internal positive control), forward, 5'-GGCCTCCAAGGAGTAAGACC-3', and reverse, 5'-AGGGGTCTACATGGCAACTG-3'.

In vitro chemotaxis assay

The MIP-3α-induced chemoattractant activity of human LCs was determined by an invasive assay. Human LCs were prepared according to the previous report (18). Briefly, monocytes obtained from peripheral blood mononuclear cells using Monocyte Isolation Kit II (Miltenyi Biotec K.K., Tokyo, Japan) were cultured in a complete medium supplemented with granulocyte macrophage-colony stimulating factor (GM-CSF) (100 ng/ml, PeproTech), IL-4 (10 ng/ml, PeproTech) and transforming growth factor (TGF)-β1 (10 ng/ml, PeproTech). At days 2 and 4, a fresh medium supplemented with GM-CSF and TGF-β1 was added. At days 5–7 of culture, cells were resuspended in 96-well plates assembled with the Collagen I Cell Invasion Assay (Trevigen, Gaithersburg, MD, USA). Induction to LCs was confirmed by a Langerin positivity. The supernatants of IL-1β- or Pam3CSK4-stimulated BECs for 2 days were added to the medium of LCs at a concentration of 5% and the degree of LC migration was evaluated according to the manufacturer's instruction. To confirm the MIP-3α-mediated process, LCs were incubated with additional human recombinant MIP-3α (1000 U/ml, PeproTech) or sufficient neutralizing anti-MIP-3α antibody (5 µg/ml, R&D systems, Minneapolis, MN, USA) in the culture. Normal rabbit IgG (5 µg/ml, R&D systems) was used as a negative control.

Statistical analysis

Data were analysed using Welch's *t*-test. *P* < 0.05 was considered statistically significant.

Results

Detection of plasmacytoid dendritic cells, myeloid dendritic cells and Langerhans cells in liver sections

Immunohistochemistry revealed that BDCA-2⁺ pDCs were mainly scattered in the portal tracts and, in particular, prominent in inflamed portal tracts in the cases of PBC, CVH and drug-induced liver injury (Fig. 1A and B). In contrast, in normal livers and hepatic sarcoidosis, there were few BDCA-2⁺ cells in the portal tracts. The distribution of BDCA-2⁺ cells was not anatomically correlated with bile ducts, irrespective of diseased livers and those attaching bile ducts were hardly found (Fig. 1A and B). Double immunolabelling of BDCA-1 and CD19 revealed that BDCA-1⁺ and CD19⁻ mDCs were also present in the portal tracts (Fig. 1C and D). In contrast to pDCs, mDCs were mostly distributed around and within bile ducts and especially prominent in PBC (Fig. 1C and D). Langerin⁺ LCs as well as mDCs were found around and within bile ducts and several Langerin⁺ LCs were rather accentuated around the damaged interlobular bile ducts in PBC (Fig. 2A). In CVH and drug-induced liver injury, several Langerin⁺ LCs were scattered within portal tracts, but no peribiliary accentuation was seen in the controls including drug-induced liver injury (Fig. 2B and C). In hepatic sarcoidosis and normal liver, Langerin⁺ LCs were almost unrecognizable (2D). Double staining of Langerin and PAS revealed that several LCs were

located within PAS-positive biliary basement membranes, demonstrating that LCs had infiltrated the biliary epithelial layer (Fig. 2E). Moreover, double immunostaining of Langerin and CCR6 revealed that these LCs express CCR6 (Fig. 2F).

Evaluation of Langerhans cells

Because the primary survey revealed that the distribution of LCs was closely associated with bile ducts, the number of Langerin⁺ LCs attaching to or embedding in bile ducts was calculated in each bile duct. As shown in Figure 3, the number of Langerin⁺ LCs in damaged bile ducts in PBC was relatively large in comparison with that in normal bile ducts in PBC or interlobular bile ducts in the controls ($P < 0.05$).

Expression of macrophage inflammatory protein-3 α in liver sections

The expression of MIP-3 α was basically found in infiltrating mononuclear cells, periportal hepatocytes and bile ducts and particularly prominent in the damaged bile ducts in PBC (Fig. 4A). Some MIP-3 α -positive mononuclear cells and periportal hepatocytes were found in the control diseased livers as well as PBC, but were faint or lacking in normal livers (Fig. 4B). Moreover, the semiquantitative evaluation also showed the

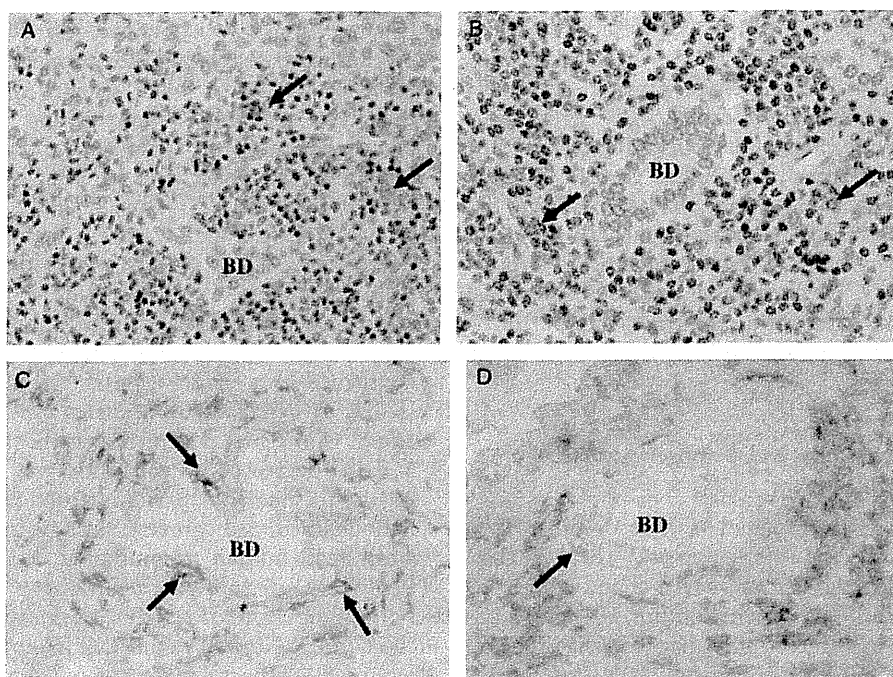


Fig. 1. Detection of plasmacytoid dendritic cells (pDCs) and myeloid dendritic cells (mDCs) in liver sections. Blood dendritic cell antigen (BDCA)-2⁺ pDCs (arrows) are scattered in inflamed portal tracts of primary biliary cirrhosis (PBC) (A) and chronic viral hepatitis (CVH) (B) patients, but their distribution is not related to bile ducts (BD). BDCA-1 (blue)-positive and CD19 (brown)-negative mDCs are mostly distributed around and within BD in PBC (C), but are rare in CVH (D). (A) and (B) show the immunohistochemistry for BDCA-2 using formalin-fixed, paraffin-embedded sections. (C) and (D) show double immunohistochemistry for BDCA-1 (brown) and CD19 (blue) using frozen sections.

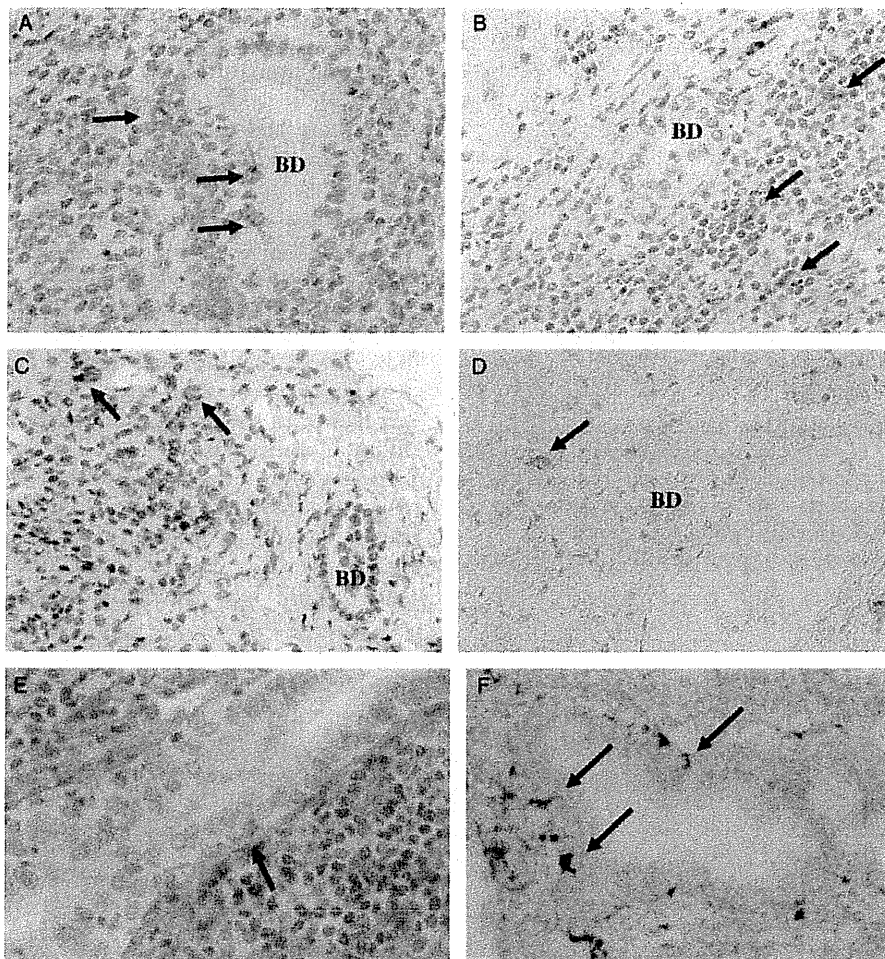


Fig. 2. Detection of Langerhans cells (LCs) in liver sections. (A) Langerin⁺ LCs are found around and within bile ducts (BD) showing cholangitis in primary biliary cirrhosis (PBC). In chronic viral hepatitis (CVH) (B) and drug-induced liver injury (C), several Langerin⁺ LCs are scattered in the portal tracts, but periductal accumulation is not found. (D) In normal liver, Langerin⁺ LCs are very rare. (E) Some Langerin⁺ LCs are located within PAS-positive biliary basement membranes in PBC. Arrowheads denote Langerin⁺ LCs. (F) Langerin⁺ LCs (brown) express CCR6 (green). Arrows and 'BD' denote representative positive cells and BD, respectively, in each figure. (A, B, C and D) Immunohistochemistry for Langerin using formalin-fixed, paraffin-embedded sections. (E) Double immunohistochemistry for Langerin and histochemistry of PAS staining. (F) Double immunohistochemistry for Langerin and CCR6.

expression of MIP-3 α to be statistically significant in the damage bile ducts of PBC, compared with normal bile ducts of PBC and interlobular bile ducts of the controls ($P < 0.05$) (Fig. 4C).

Detection of macrophage inflammatory protein-3 α in cultured biliary epithelial cells

The mRNA of MIP-3 α was constantly expressed in cultured BECs under normal conditions. Moreover, its expression was upregulated by several cytokines including IL-1 β , TNF- α and IL-17 and all PAMPs examined in this study (Fig. 5A). In addition to this molecular analysis, immunocytochemistry also demonstrated that MIP-3 α protein was strongly expressed in cytokine- and PAMP-stimulated BECs, compared with non-stimulated BECs (Fig. 5B and C).

Chemotactic effects of macrophage inflammatory protein-3 α to Langerhans cells

The degree of LCs that transmigrated in an assay chamber containing the culture medium of IL-1 β - or PamCSK4-treated BECs was larger than that of the non-treated supernatant and these increased migrations were inhibited by neutralizing MIP-3 α antibody, suggesting that the MIP-3 α secreted by BECs chemoattracted LCs (Fig. 6).

Discussion

Dendritic cells form a network comprising different populations that initiate and differentially regulate immune responses and regulate the generation of inflammatory cells including T cells and are known as a regulator of continuous chronic inflammation. Although

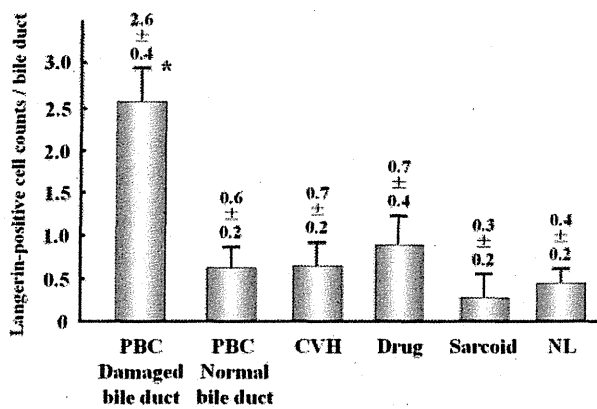


Fig. 3. Semiquantitative evaluation of Langerhans cells (LCs). The number of Langerin⁺ LCs attached to damaged bile ducts in primary biliary cirrhosis (PBC) was relatively large in comparison with that attached to normal bile ducts in PBC or interlobular bile ducts in chronic viral hepatitis (CVH), drug-induced liver injury (Drug), hepatic sarcoidosis (Sarcoid) and normal livers (NL). Bars indicate the mean ± SEM. **P* < 0.05.

studies have already demonstrated that extensively inflamed biliary mucosa of PBC contain several DCs including immature DCs (8), S-100⁺ DCs (7) and LCs (19, 20), it is necessary to investigate the extent of infiltration and the mechanism of migration to understand the immune response forming cholangiopathy, especially chronic cholangitis, in PBC. The present study revealed that periductal DCs were mostly mDCs and that the major representative type of mDCs, Langerin⁺ LCs, were attached to or embedded in the damaged bile ducts of PBC patients. These results suggest that LCs existing around and within bile ducts play an important role as APCs presenting bile duct-derived autoantigens. LCs are a population of immature DCs involved in antigen processing in the skin and intestinal mucosa and can be found in diseased tissues (13). The presence and function of LCs closely associated with bile ducts, therefore, are suggested to be of importance in the establishment of bile duct-specific autoimmune phenomena in PBC.

Because LCs possess chemokine receptors (CCR4 and CCR6), their ligands including CCL17, CCL22 and

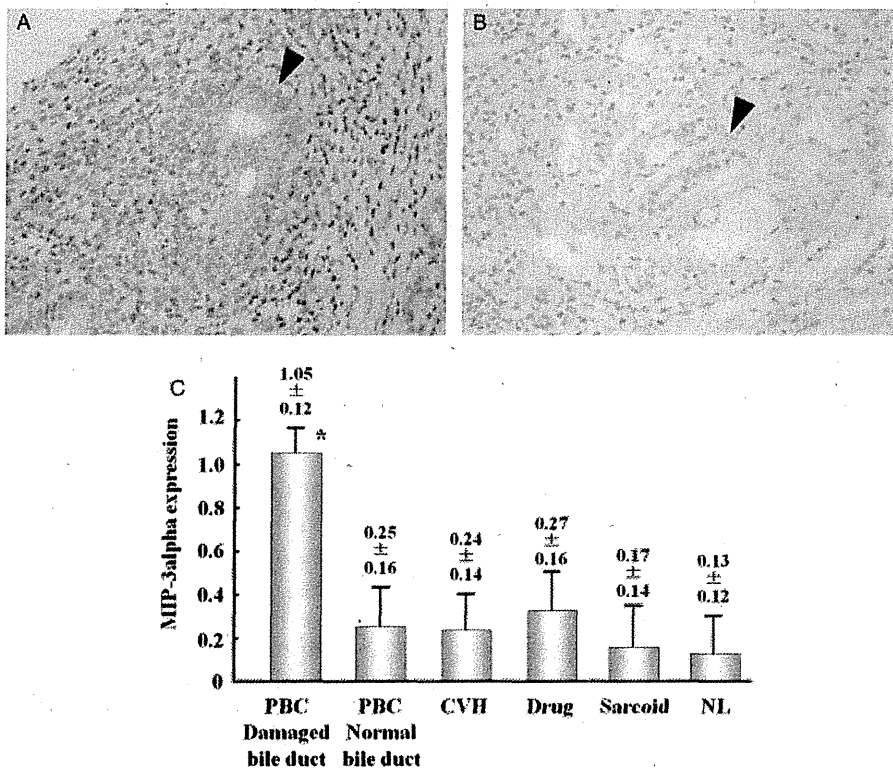


Fig. 4. Detection of macrophage inflammatory protein-3α (MIP-3α) in liver sections. MIP-3α was expressed in infiltrating mononuclear cells and bile ducts. In primary biliary cirrhosis (PBC), the damaged bile ducts show strong MIP-3α expression (A, arrowheads). In chronic viral hepatitis (CVH), a few MIP-3α⁺ mononuclear cells were found, but bile ducts show no expression (B, arrowheads). (A) and (B) show serial sections of Figures 2A and B respectively. Semiquantitative evaluation also shows that the degree of MIP-3α expression in the damaged bile ducts of PBC is significantly higher than that in normal bile ducts in PBC or interlobular bile ducts in chronic viral hepatitis (CVH), drug-induced liver injury (Drug), hepatic sarcoidosis (Sarcoid) and normal livers (NL) (C, *P* < 0.05). Bars indicate the mean ± SEM. **P* < 0.05. (A and B) Immunohistochemistry for MIP-3α using formalin-fixed, paraffin-embedded sections.

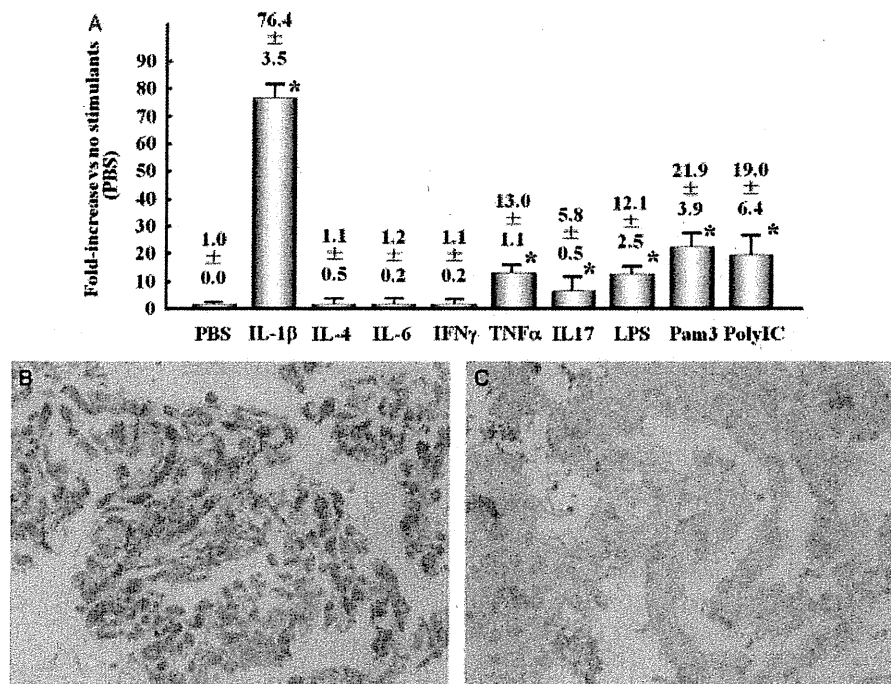


Fig. 5. Detection of macrophage inflammatory protein-3α (MIP-3α) mRNA in cultured human biliary epithelial cells (BECs). A: Quantitative analyses using real-time polymerase chain reaction in two independent experiments revealed that IL-1β, TNF-α, IL-17, LPS, Pam3CSK4 and poly(I:C) significantly upregulated MIP-3α mRNA expression. There were no differences between the two BEC lines. Data are shown relative (fold-increase) to the value for unstimulated cells (PBS). Bars indicate the mean ± SEM. **P* < 0.05. (B and C) Immunocytochemistry also demonstrated that MIP-3α protein was strongly expressed in the LPS-stimulated BECs (B), compared with the unstimulated BECs (C). IL, interleukin; LPS, lipopolysaccharide; poly(I:C), polyinosinic-polycytidylic acid; TNF, tumour necrosis factor;

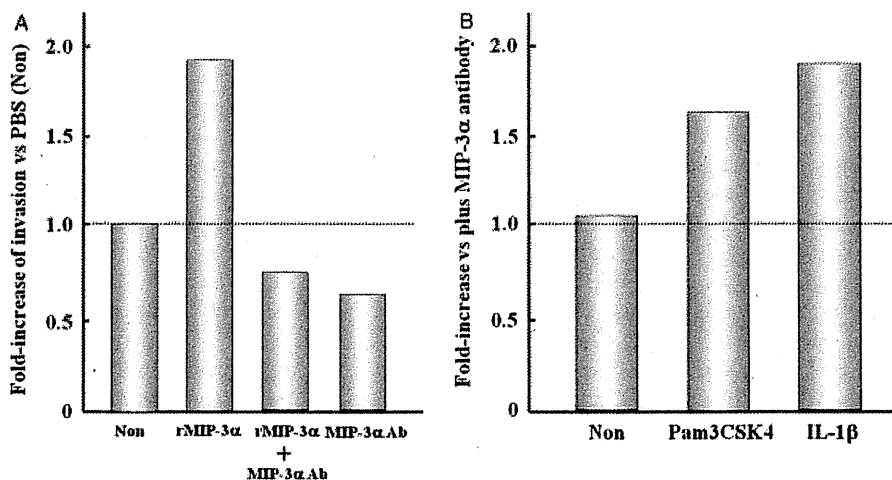


Fig. 6. Chemotactic effects of recombinant human macrophage inflammatory protein-3α (rMIP-3α) and secreted MIP-3α from cultured human biliary epithelial cells (BECs) on Langerhans cells (LCs). The representative data are shown in two independent experiments. (A) Data are shown as a relative degree of cell invasion of LCs vs that without rMIP-3α or MIP-3 antibody (Non). The migration of LCs was induced by rMIP-3α and inhibited by an additional neutralizing MIP-3α antibody (rMIP-3α + MIP-3α Ab). More precisely, the MIP-3α antibody decreased the migration at a level lower than the standard level (Non), suggesting that the standard medium for LCs originally contained MIP-3α. (B) Cultured BECs were stimulated with Pam3CSK4 or interleukin (IL)-1β for 2 days and these supernatants were added to a medium for LCs in the assay chamber at a concentration of 5%. Data are shown as the relative fold-increase vs those with additional neutralizing MIP-3α Ab, because the standard medium for LCs contains MIP-3α. Consequently, the additional supernatant of Pam3CSK4- or IL-1β-treated BECs upregulated the migration of LCs, compared with the non-treated medium (Non).

MIP-3 α (CCL20) could migrate LCs. Dieu-Nosjean *et al.* (13) reported that MIP-3 α plays a central role in recruiting LC precursors into the epithelium during inflammation. Langerin⁺ DCs, moreover, were localized subepithelially adjacent to MIP-3 α -expressing epithelium (21) and, in the colon of ulcerative colitis patients, MIP-3 α plays an important role in the disease's pathogenesis (22). Interestingly, in the present study, MIP-3 α -expressing BECs and Langerin⁺ LCs were frequently co-localized in the same bile ducts showing cholangitis. A previous study demonstrated that MIP-3 α was produced by keratinocytes and that the migration of LC precursors induced by the supernatant of activated skin keratinocytes was completely blocked with an antibody against MIP-3 α (13). Therefore, the presence of BECs producing MIP-3 α suggests that BECs play an important role in the migration of LCs in biliary layers of PBC patients.

Recently obtained evidence indicates that bile duct-related acquired immunity is closely associated with the periductal cytokine milieu and biliary innate immunity (4, 23–26). We have reported previously that human BECs possess receptors for some cytokines (IL-1 β , IL-6, IL-4, IFN- γ , TNF- α and IL-17) and TLRs for some PAMPs [LPS, Pam3CSK4 and poly(I:C)] (24, 27), suggesting that they could show a response to these cytokines and PAMPs and induce various immunological reactions. In fact, the present study demonstrated that IL-17 as well as inflammatory cytokines (IL-1 β and TNF- α) significantly upregulated the expression of MIP-3 α . We demonstrated previously that IL-17-positive cells (Th17 cells) were accumulated around the damaged bile ducts in PBC and that BECs possessed the ability to produce Th17-inducible cytokines including IL-1 β as a result of the innate immune response (27). Our previous studies, moreover, have found that the expression of IL-1 β and TNF- α was increased in pathologic BECs of PBC patients (27, 28). The present study demonstrated that the expression of MIP-3 α was selectively restricted at sites of inflammation including cholangitis in PBC, suggesting the production of MIP-3 α to be closely associated with periductal cytokines including IL-1 β , TNF- α and IL-17. In addition to these cytokine milieus, the damaged bile ducts and/or portal tracts show an increased susceptibility to PAMPs and enhanced production and secretion of innate inflammatory mediators in PBC (23, 29). All PAMPs examined in this study significantly upregulated the production of MIP-3 α in human BECs. Therefore, BEC-derived MIP-3 α production caused by PBC-specific cytokine and innate immune responses facilitates the migration of LCs as biliary epithelium-specific APCs, followed by the aggravating and continuous cholangitis in PBC.

In summary, this study provides the following key findings: the frequent clustering of mDCs, in particular LCs, around or within the damaged bile ducts in PBC; a significant positive correlation between the frequency of LCs and the degree of MIP-3 α expression in bile ducts; and the induction of MIP-3 α expression in BECs caused

by the biliary innate immune and cytokine responses. These findings suggest that LCs existing around or within biliary epithelial layers are important as APCs in the damaged bile ducts in PBC and directed in their migration by BEC-derived MIP-3 α induced via a PBC-specific periductal cytokine milieu and biliary innate immunity. An understanding of the specific functions of LCs and MIP-3 α in biliary inflammatory and immune disorders might provide new strategies to attenuate the cholangiopathy in PBC.

Acknowledgements

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Pathogenesis of IgG4-related disease

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Purpose of review

To review studies that have examined underlying genetic and immunological aspects of IgG4-related disease.

Recent findings

Genetic studies have suggested that several human leukocyte antigen (HLA) and non-HLA haplotypes/genotypes are associated with susceptibility to IgG4-related disease or to disease relapse after steroid therapy. Among several autoantibodies identified so far, autoantibodies against lactoferrin and carbonic anhydrase II are most frequently detected in serum of IgG4-disease patients. However, it has not been well clarified whether or not those autoantibodies belong to an IgG4 subclass. Studies that have demonstrated molecular mimicry between *Helicobacter pylori* and constituents of pancreatic epithelial cells suggest that gastric *H. pylori* infection triggers autoimmune pancreatitis in genetically predisposed individuals through antibody cross-reactivity. Recently, T-helper 2 immune reaction has been suggested to be predominant in IgG4-related disease. Interestingly, regulatory immune reactions are activated in IgG4-related disease, and regulatory cytokines interleukin-10 and transforming growth factor- β have been suggested, respectively, to play important roles in IgG4 class switch and fibroplasia.

Summary

Autoimmunity has been considered the most probable pathogenesis of IgG4-related disease, but has not been completely proved so far. A breakthrough study to detect a specific autoantigen, autoantibody, or pathogen is necessary.

Keywords

autoimmune pancreatitis, cytokine, IgG4, pathology, regulatory T cells

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Introduction

IgG4-related disease is a unique inflammatory condition characterized by tumorous swelling of affected organs and high serum IgG4 concentrations [1]. Histological characteristics are diffuse lymphoplasmacytic infiltration, occasional eosinophils, storiform fibrosis, obliterative phlebitis, and infiltration by numerous IgG4-bearing plasma cells [2*]. That autoimmunity may participate in pathogenesis is suggested, particularly by serologic abnormalities (e.g. presence of autoantibodies and elevated levels of γ -globulin) and by a dramatic response to steroid therapy [3,4**]. However, the detailed pathogenetic mechanism and underlying immune response have not been well clarified. Especially mysterious is why IgG4 is specifically elevated in this disease.

In this article, we review reports that have examined underlying genetic and immunological aspects of IgG4-related disease. A caveat is that most studies on the pathogenesis of IgG4-related disease have examined only pancreatitis and sialodacryoadenitis.

Genetic backgrounds

Several genetic susceptibility factors for IgG4-related disease have been identified. Most studies so far have examined only Asian patients with autoimmune pancreatitis (AIP), a pancreatic manifestation of IgG4-related disease. It is still unknown whether or not these data apply to Western people because susceptibility factors often vary among different ethnic groups. In 2002, Kawa *et al.* [5] reported that the frequencies of human leukocyte antigen (HLA) serotypes DRB1*0405 and DQB1*0401 were significantly higher in Japanese patients with AIP than in healthy individuals and in patients with chronic calcifying pancreatitis. In contrast, a Korean study [6] could not show significant associations between HLA type and development of AIP in Korean patients. Interestingly, this study revealed that the nonaspartic acid at DQB1 57 was significantly associated with relapse of AIP. All Korean patients who experienced a relapse during or after steroid treatment had nonaspartic acid at DQB1 57 in contrast to 30% of AIP patients without a relapse [6]. However, this result could not be confirmed in Japanese patients [7*].

Non-HLA genes have been also investigated regarding susceptibility to AIP. Cytotoxic T lymphocyte-associated antigen 4 (CTLA-4), expressed on activated T cells, plays an important role in the regulation of T-cell stimulation. Many single-nucleotide polymorphisms (SNPs) have been identified in *CTLA-4*, encoding CTLA-4, and some are associated with susceptibility to autoimmune diseases [8]. It was reported that the frequency of a *CTLA-4* 49A haplotype was significantly higher in Chinese AIP patients than in healthy individuals [9]. The -318C/+49A/CT60G haplotype increased susceptibility to AIP. The investigators also found that the tumor necrosis factor- α (TNF- α) promoter -863A haplotype was related to extrapancreatic involvement in Chinese patients with AIP [9]. A Japanese study [10] showed that a SNP at *CTLA-4* +6233 in the 3'-untranslated region contributes to disease susceptibility. A +6230G/G genotype increased susceptibility to AIP, whereas a +6230A haplotype was associated with AIP resistance [10]. In addition, the +49A/A and +6230A/A genotypes were related to an increased risk of relapse [10].

Fc receptor-like 3 (FCRL3) molecule, expressed on B cells, is known to augment autoantibody production in individuals with disease-susceptible genotypes [11]. *FCRL3* polymorphisms have been shown to contribute to susceptibility to autoimmune diseases. Umemura *et al.* [12] reported that the -110A/A genotype is associated with AIP in the Japanese population. Interestingly, serum IgG4 concentrations were positively correlated with the number of susceptible alleles [12]. The same group also examined SNPs in the gene encoding Toll-like receptor 4 (*TLR4*), but they were not associated with susceptibility to or relapse of AIP [13*].

Target organs

IgG4-related disease can develop in almost all organs. The pancreatic lesion (AIP) is the prototype of IgG4-related disease. After the landmark study by Hamano *et al.* [1], IgG4-related lesions histologically similar to AIP have been identified in various organs such as bile duct (sclerosing cholangitis) [14], salivary gland (chronic sclerosing sialadenitis) [15*], lacrimal gland (chronic sclerosing dacryoadenitis or Mikulicz's disease) [16*], retroperitoneum (retroperitoneal fibrosis) [17*], aorta (inflammatory aneurysm) [18], lung (nodular or diffuse inflammation) [19*], and kidney (tubulointerstitial nephritis) [20**]. Multiple lesions can develop in different organs in a patient simultaneously or metachronously. Interestingly, kidney lesions have always been associated with extrarenal disease in personally studied cohorts. This may suggest that kidneys are not a primary site of IgG4-related disease.

IgG4-related lesions in various organs are now considered one of the manifestations of systemic IgG4-related dis-

ease based on similar clinical and histological features [21*,22**]. However, clinicopathological features are not completely identical among manifestations. One of the differences is that the proportion of women is significantly higher in patients with sialadenitis or dacryoadenitis (approximately 50%) than in patients with involvement of other organs (15–25%) (Y. Zen, unpublished data). Although IgG4-related disease has been thought of as a single disease entity, different causative factors may be involved in different organs.

Potential autoantigens

Several groups have tried to identify autoantigens based on the hypothesis that IgG4-related disease is an autoimmune disorder. Nonspecific antinuclear antibodies are identified in more than half of patients with IgG4-related disease. Antibodies against lactoferrin and carbonic anhydrase (CA)-II are the most frequently detected autoantibodies in AIP (73 and 54%, respectively) [23]. Interestingly, a strong positive correlation between increase in serum IgG4 levels and anti-CA-II antibody levels has been reported in AIP [24]. More recently, it was reported that anti-CA-IV antibodies were detected in 10 of 29 (34%) patients with AIP [25]. Lactoferrin and carbonic anhydrases are expressed in some exocrine organs including the pancreas, salivary gland, bile duct, and renal tubules. The distribution of these antigens may be related to systemic manifestations of IgG4-related disease.

Pancreatic secretory trypsin inhibitor (PSTI) is another potential autoantigen. Antibodies against this molecule were detected in 30–40% of patients with AIP [26]. In a study by Asada *et al.* [26], 23 of 26 patients with AIP had antibodies against either CA-II or lactoferrin. Two of three patients in whom neither of these autoantibodies could be demonstrated had anti-PSTI antibodies. Interestingly, these autoantibodies were of the IgG1, not the IgG4 subclass [26]. A recent European study [27*] also revealed autoantibodies against trypsinogens, as well as anti-PSTI antibodies in AIP patients.

In 2010, Yamamoto *et al.* [28*] performed a proteomics study to identify potential autoantigens for IgG4-related diseases in 28 patients with IgG4-related dacryoadenitis, sialadenitis, pancreatitis (AIP), or tubulointerstitial nephritis. Immune complexes taken from patients' sera were divided into immunoglobulin and antigens and were analyzed by surface-enhanced laser desorption/ionization–time-of-flight mass spectrometry. They found an antigen, a 13.1-kDa protein that was a potential candidate autoantigen, in all samples from patients with IgG4-related diseases, but not in samples from patients in the normal and disease-control cohorts [28*].

To the best of our knowledge, IgG4 type autoantibodies have not been detected in IgG4-related disease. It remains to be answered whether IgG4 is an autoantibody in IgG4-related disease or is overexpressed secondarily in response to an unknown primary inflammatory stimulus. Aoki *et al.* [29] examined whether or not IgG4 from patients with IgG4-related disease can bind to human tissue. Histological sections of normal human tissue were incubated with patients' serum. IgG4 bound to a tissue antigen was then detected by secondary antibodies against human IgG4. Interestingly, IgG4 from the patients was found to be able to bind to the normal epithelia of the pancreatic ducts, bile ducts, gallbladder, and salivary gland ducts [29].

Involvement of *Helicobacter pylori*

The possible involvement of *Helicobacter pylori* in the pathogenesis of AIP was reported in 2005 [30]. Guarneri *et al.* [31] found a significant homology between human CA-II and α -carbonic anhydrase of *H. pylori*. The homologous segments contain the binding motif of the HLA molecule DRB1*0405. This study suggests that gastric *H. pylori* infection triggers AIP in genetically predisposed individuals through molecular mimicry [31]. More recently, Frulloni *et al.* [32**] found that 94% of patients with AIP had antibodies against plasminogen-binding protein of *H. pylori*. The amino acid sequence of plasminogen-binding protein exhibits homology with that of ubiquitin-protein ligase E3 component n-recogin 2, an enzyme expressed in pancreatic acinar cells. However, serum IgG4 levels were elevated in only 53% of patients in this study, suggesting that the cohort assessed might in substantial part represent non-IgG4-related AIP (type II AIP [4**]) [32**].

Cytokines and chemokines

In 2000, Okazaki *et al.* [23] serologically examined T-helper 1 (Th1)/Th2 balance in patients with AIP. Numbers of interferon- γ (IFN- γ)-producing peripheral CD4⁺ cells and secreted levels of IFN- γ were significantly increased compared with those in controls, whereas numbers of interleukin (IL)-4-producing cells and serum levels of IL-4 were not increased. Yamamoto *et al.* [33] similarly reported that IFN- γ /IL-4 ratios were significantly higher in IgG4-related dacryoadenitis and sialadenitis than in Sjögren's syndrome. In contrast, five recent studies provided data suggesting that the Th2-dominant immune response is more activated in IgG4-related disease [34,35*,36*,37*,38*]. Peripheral blood mononuclear cells collected from patients with IgG4-related disease produced predominantly Th2-type cytokines such as IL-4, IL-5, IL-10, and IL-13 after T-cell stimulation. Interestingly, peripheral blood mononuclear cells of one patient exhibited increased production of

IgG4, as well as IL-10 upon stimulation with TLR ligands [36*].

In 2007, we histopathologically examined cytokine production *in situ* in materials from patients with IgG4-related disease [39]. Quantitative real-time PCR using RNA extracted from frozen tissue of affected organs (pancreas, bile duct, salivary gland, and lacrimal gland) revealed significantly higher ratios of IL-4/IFN- γ (45.8-fold), IL-5/IFN- γ (18.7-fold), and IL-13/IFN- γ (20.7-fold) in IgG4-related disease tissues than in tissues from patients with classical autoimmune diseases [39]. Lymphocytes expressing IL-4 were clearly demonstrated by in-situ hybridization. Similarly, Th2-dominant cytokine production was also reported in salivary and lacrimal gland lesions [34]. Th2-type cytokines play an important role in inflammatory reaction in the organs affected by IgG4-related disease.

A chemokine expression profile in IgG4-related disease has not been well characterized. One study [40] showed that in IgG4-related pancreatitis and cholangitis expression levels of CXC chemokine receptor-5 (CXCR5) and CXC chemokine ligand-13 (CXCL13) mRNAs were higher than in chronic pancreatitis and primary sclerosing cholangitis. In addition, CXCR5⁺ or CXCL13⁺ lymphocytes were noted in periductal areas with or without parenchymal localization [40]. Another study [35*] showed expression of CD40 and its ligand in infiltrating lymphocytes and plasmacytes. Interaction of CXCR5-CXCL13 or of CD40-CD40 ligand may be involved in lymphocyte trafficking, B-cell activation, and immunoglobulin production in IgG4-related disease.

Role of regulatory T cells

Impaired function or a decreased number of regulatory T cells (Tregs) is believed to be an important step in the development of autoimmune disorders. However, of interest is that the number of Tregs is not decreased – rather, characteristically increased – in both tissue and blood of patients with IgG4-related disease. Our investigation revealed that mRNA of forkhead box P3 (Foxp3, a Tregs-specific transcriptional factor) in IgG4-related disease was expressed 36.4-fold higher than in classical autoimmune diseases. Two regulatory cytokines, IL-10 and transforming growth factor- β (TGF- β), were also significantly overexpressed (45.3-fold and 39.4-fold, respectively) [39]. Furthermore, CD4⁺CD25⁺Foxp3⁺ Tregs could be detected within affected tissues by immunohistochemistry, in numbers significantly higher than in autoimmune and nonautoimmune disease controls. Overexpression of IL-10 and TGF- β was also confirmed in IgG4-related tubulointerstitial nephritis [20**]. Koyabu *et al.* [41*] reported that the number of Foxp3⁺ cells was significantly correlated with the

number of IgG4⁺ plasma cells in IgG4-related cholangitis. Miyoshi *et al.* [42] examined the number of Tregs in the blood and reported that the number of CD4⁺CD25^{high} Tregs was significantly higher in patients with AIP than in patients with chronic pancreatitis and was correlated with the level of serum IgG4. The number of naive Tregs was significantly decreased. They speculated that hyporeaction of naive Tregs might be involved in the development of IgG4-related disease, whereas hyperreaction of CD4⁺CD25^{high} Tregs could reflect IgG4-related disease progression [42].

Histology-immunology correlations

Two key histological features of IgG4-related disease are infiltration of IgG4⁺ plasma cells and storiform fibrosis. IL-10 produced by Th2 cells or Tregs may participate in IgG4⁺ plasma cell infiltration because IL-10 has a potent function in directing B cells to produce IgG4. TGF- β secreted from Tregs and macrophages may induce fibroplasia, given the strong fibrogenic function of TGF- β and the expression of a TGF- β receptor in myofibroblasts in AIP [39,43]. On occasion, IgG4-related disease histologically shows eosinophilic infiltration. In addition to IgG4, serum eosinophilia and high serum IgE levels are frequently seen in patients with IgG4-related disease. These pathological and immunological features may be mediated by IL-4, IL-5, and IL-13 because these cytokines are important for class switch to IgE production and for eosinophil migration or activation. IgG4-related disease commonly develops in people with a predisposition to allergy [44,45]. About 20% of patients in our studied cohorts have allergic disorders (e.g. bronchial asthma or drug allergy). One attractive hypothesis is a possible involvement of chronic allergic reaction, but not autoimmunity, in the pathogenesis of IgG4-related disease. However, a specific allergen must be identified to confirm this hypothesis.

Conclusion

We have briefly described data on the pathogenesis of IgG4-related disease. Because most of the research is based on only AIP or sialadenitis/dacryoadenitis, it is still unknown whether the same immune reaction takes place in other organs. A point of particular importance is that potential autoantigens are not identified in nonglandular tissues (e.g. retroperitoneum or artery). Regarding Th1/Th2 balance, recent studies have accumulated evidence that suggests an involvement of Th2-dominant reaction in the pathogenesis of IgG4-related disease. Autoimmunity has been suggested as the most possible pathogenetic cause, but allergic cause is another attractive hypothesis given observed cytokine expression profiles and the frequent association of IgG4-related disease with a predisposition to allergy. A breakthrough study to detect

a specific autoantigen, autoantibody, pathogen, or allergen is necessary.

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Papers of particular interest, published within the annual period of review, have been highlighted as:

- of special interest
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Additional references related to this topic can also be found in the Current World Literature section in this issue (pp. 129–130).

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分冊 6 - 2

研究代表者 坪内 博仁

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V. 研究成果の刊行物・別刷
(平成23年度)

分冊 6 - 2

研究成果の刊行に関する一覧表（平成23年度）

雑 誌

発表者氏名	論文タイトル名	発表誌名	巻号	ページ	出版年
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