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Suppressive role of hepatic dendritic cells in concanavalin A-induced hepatitis

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Introduction

It has been established recently that dendritic cells (DC) in the liver have unique properties in terms of their phenotype, morphology and function. In contrast to conventional (c) DC (CD11c⁺ CD11b⁺ B220⁻) present in the spleen and other organs, hepatic DC are thought to be plasmacytoid (p) DC (CD11c⁺ CD11b⁻ B220⁺) [1-3]. Furthermore, hepatic DC are able to mediate certain immunotolerance phenomena [4,5]. These properties are quite different from those of cDC [6,7]. Thus, cDC have the potential to mediate immunostimulatory functions with regard to antigen presentation. However, a recent report shows that hepatic DC also mediate immunostimulatory function, resulting in fibrosis [8].

In light of these findings, we conducted experiments to elucidate further the characteristics of hepatic DC using a murine model of concanavalin A (Con A)-induced acute autoimmune hepatitis. The effector cells responsible for Con A-induced hepatitis are primarily CD4⁺ T cells [9-11], natural killer (NK) T cells [12,13] and neutrophils [14],

Summary

Concanavalin A (Con A)-induced hepatitis is a mouse model of acute autoimmune hepatitis. The aim of this study was to investigate the role of hepatic dendritic cells (DC) in the immune modulation of tissue damage. Almost all hepatic DC were plasmacytoid DC (CD11c⁺ **I-Alow B220**⁺ **); however, conventional DC were CD11c**⁺ **I-Ahigh B220– . At an early stage (3–6 h) after Con A administration, the number of DC in both the liver and spleen decreased, increasing thereafter (12–24 h) in parallel with hepatic failure. The hepatic CD11c**⁺ **DC population contained many CD11b**- **cells, while the majority of splenic CD11c**⁺ **DC were CD11b**⁺ **. After Con A administration, the proportion of I-A**⁺ **and CD11b**⁺ **cells within the CD11c**⁺ **DC population tended to increase in the liver, but not in the spleen. Similarly, expression of the activation markers CD80, CD86 and CD40 by CD11c**⁺ **DC increased in the liver, but not in the spleen. Next, adoptive transfer of DC isolated from the liver and spleen was performed 3 h after Con A administration to examine the immunomodulatory function of DC. Only hepatic DC had the ability to suppress hepatic failure. Analysis of cytokine production and subsequent identification of the effector cells showed that hepatic DC achieved this by suppressing the production of interleukin (IL)-12 and IL-2, rather than modulating effector cell function.**

Keywords: autoimmune hepatitis, dendritic cells, immunosuppression, liver, plasmacytoid DC

> although an association with the function of regulatory T cells (T_{res}) cells has also been reported [15,16]. However, the precise role played by hepatic DC in Con A-induced autoimmune hepatitis is unknown. In particular, it is unclear whether hepatic DC have an immunosuppressive or immunostimulatory function in this model. In the present study, we confirmed the phenotype and morphology of hepatic DC, and showed that hepatic DC have an immunosuppressive effect on Con A-induced hepatitis.

Materials and methods

Mice

C57BL/6(B6) mice (female; 6–12 weeks old) were purchased from Charles River Japan (Yokohama, Japan). The mice were kept in a room under constant temperature $(25 \pm 2^{\circ}C)$ and humidity (50–70%) and a 12-h light–dark cycle (lights on from 08:00 to 20:00 h). They were fed under specific pathogen-free conditions in the animal facility of Niigata University (Niigata, Japan).

Induction of Con A-induced hepatitis

Con A (type IV; Sigma-Aldrich, St Louis, MO, USA) was dissolved in sterilized phosphate-buffered saline (PBS) and injected intravenously at a dose of 15 mg/kg [12]. Sera from individual mice were obtained 3, 6, 12 and 24 h after Con A injection. Serum aminotransferase [alanine aminotransferase (ALT) and aspartate aminotransferase (AST)] activity was measured using standard photometric methods. All experiments were approved by the Animal Ethics Committee of Niigata University.

Cell preparation

Mice were anaesthetized with isoflurane, perfused with 1500 IU/ml dispase II solution (Life Technologies Co., Carlsbad, CA, USA) via the portal vein, and the liver and the spleen removed. Hepatic DC were isolated using the following methods. Briefly, the liver was cut into fragments using scissors. These fragments were then incubated in dispase II solution for 15 min at 37°C in a shaking water-bath. The digested products were pressed through a 200-gauge stainless steel mesh and suspended in Eagle's minimum essential medium (MEM) supplemented with 5 mM HEPES and 10% fetal calf serum (FCS). After washing, the cell pellet was separated by 30% Percoll centrifugation to obtain mononuclear cells (including dendritic cells). The cells were collected and centrifuged in a discontinuous Percoll gradient at 960 *g* for 20 min. The DC-enriched fraction at the 40/50% interface was collected. Splenic dendritic cells were obtained by enzymatic digestion of the spleen and passage through a stainless steel mesh. Splenic DC were treated with 0·2% NaCl solution to remove red blood cells (RBC) and isolated using the Percoll discontinuous density gradient/centrifugation method.

For the adoptive transferred experiments, lymphocytes were isolated from mouse livers and spleens of DC-post injected Con A-treated mice. Hepatic lymphocytes were then prepared as described previously [13]. Briefly, the liver was pressed through a 200-gauge stainless steel mesh and suspended in Eagle's MEM supplemented with 5 mM HEPES and 2% FCS. After washing, the cells were resuspended in a 35% Percoll solution containing 100 U/ml heparin and centrifuged at 424 *g* for 15 min. The pellet was resuspended in RBC lysis solution [155 mM NH₄Cl, 10 mM KHCO₃, 1 mM ethylenediamine tetraacetic acid (EDTA)-2Na and 17 mM Tris, pH 7·3] and washed twice with medium.

Adoptive transfer of DC into mice

Purified CD11c⁺ cells from the liver and spleen of normal B6 mice were prepared using the magnetic bead sorting method incorporating biotin-conjugated anti-mouse CD11c monoclonal antibodies (mAb) (HL3) and streptavidin particles plus (BD Bioscience, San Diego, CA, USA) with > 90% purity. Hepatic or splenic CD11c⁺ cells from normal B6 mice were then transferred adoptively into Con A-injected mice $(3 \times 10^5 \text{ cells/mouse})$ by injection into the tail vein.

Flow cytometric analysis

The phenotype of the DC was determined by two- or threecolour immunofluorescence analysis [17]. The reagents used were anti-CD11c, anti-I-A $^{\text{b}}$ (AF6-120.1), anti-CD80 (16-10A1), anti-CD86 (GL1), anti-CD40 (3/23), anti-CD3e (145-2C11), anti-NK1·1 (PK136) (BD Bioscience), anti-CD11b (M1/70) (e-Bioscience Inc., San Diego, CA, USA) and anti-PDCA-1 (JF05-1C2·4.I) mAbs (Miltenyi Biotec GmbH, Bergisch Gladbach, Germany). All mAbs were conjugated to fluorescein isothiocyanate (FITC), phycoerythrin (PE) or biotin. The biotin-conjugated reagents were developed using TRI colour-conjugated streptavidin (Caltag Laboratories, Burlingame, CA, USA). To prevent nonspecific binding, anti-CD16/CD32 (2·4G2), mAb was added before staining with the labelled mAbs. The suspended mononuclear cells (MNCs; 5×10^5 -10⁶/tube) were stained with the mAbs and analysed using a fluorescence activated cell sorter (FACScan) flow cytometer (BD Bioscience). Dead cells were excluded by the use of appropriate forward-scatter, side-scatter and propidium iodide gating.

Morphological analysis

I-A^{high} CD11c⁺ cells and I-A^{low} CD11c⁺ cells were obtained by FACS Vantage (BC Bioscience) cell sorting and stained using the May–Grünwald–Giemsa method for morphological analysis.

Immunohistochemistry

Liver and spleen tissues were dissected from B6 mice at 6–12 weeks of age. Frozen tissue sections $(3 \mu m)$ in thickness) of each tissue were dried without fixation and stored at 4°C until use. For immunohistochemistry, the sections were blocked with 10% FCS in PBS for 30 min, and treated with 10-fold diluted PE-conjugated anti-mouse CD11c Armenian hamster monoclonal antibody (IgG1 λ 2; BD Bioscience) and 20-fold diluted anti-mouse I-A/I-E rat monoclonal antibody (IgG2b k; Biolegend, Inc., San Diego, CA, USA). Colour was detected by using 800-fold diluted Cy3-conjugated secondary antibodies against Armenian hamster immunoglobulin (Ig)G (Jackson Laboratories, Inc., PA, USA) and 200-fold diluted Alexa488-conjugated secondary antibodies against rat IgG (Molecular Probes Inc., Eugene, OR, USA) in PBS. As a negative control for the staining, sections were treated without primary antibodies. After washing with PBS, counterstaining was performed with Hoechst 33342 (Sigma-Aldrich) at 1μ g/ml. Serial sections were stained routinely with haematoxylin and eosin (H&E).

Intracellular forkhead box P3 (FoxP3) staining

For intracellular FoxP3 staining, MNCs were first stained with FITC-conjugated anti-CD4 (RM4-5). Cells were then fixed and permeabilized using the mouse FoxP3 buffer set (BD Bioscience). After permeabilization, the cells were stained intracellularly using an anti-FoxP3 (MF23) mAb and analysed using a FACScan.

Measurement of serum cytokine concentrations

The concentration of tumour necrosis factor (TNF)- α , interleukin (IL)-2, interferon (IFN)- γ and IL-12p70 was measured in the serum from each mouse using cytometric bead array (CBA) kits (BD Biosciences) according to the manufacturer's instructions.

Statistical analysis

Differences between the results obtained in normal mice and experimental mice were analysed using one-way analysis of variance (anova). A *P*-value of <0·05 was considered significant.

Results

Characterization of CD11c⁺ **DC in the liver and spleen**

First, DC in the liver and spleen of normal mice were characterized by two-colour immunofluorescence tests (Fig. 1a). These DC were detected as CD11c⁺. CD11c⁺ DC in the liver were mainly I- $A^{low} B220⁺$, but those in the spleen were a mixture of (CD11c⁺) I-A^{low} B220⁺ and $(CD11c^+)$ I-A^{high} B220⁻. CD11c⁺ I-A^{low} B220⁺ DC in the liver were characterized as hepatic pDC. In a second experiment, morphological staining was performed using DC purified by cell sorting (Fig. 1b). Irrespective of the organ of origin, I-Ahigh CD11 c^+ cDC were large, with a DC morphology, while I-A low CD11c⁺ pDC were small, with a plasmacytoidlike morphology.

Con A-induced hepatitis and the number of intrahepatic DC

AST and ALT levels in the sera of Con A-treated mice increased after 12–24 h (Fig. 2a). During this time, DC were also identified in the immunofluorescence experiments (Fig. 2b). DC were identified as CD11c⁺, irrespective of the organ of origin. During the early stages (3–6 h after administration), the absolute number of DC in the liver decreased; however, the number increased during the later stages (12–24 h). A similar pattern was observed for splenic DC.

Fig. 1. Characterization of dendritic cells (DC) in the liver and spleen. (a) Phenotype and (b) morphology. DC phenotype was characterized by immunofluorescence analysis. The expression of I-A and B220 on CD11 c^+ DC was examined. I-A^{high} CD11 c^+ and I-A^{low} CD11 c^+ cells were purified using a cell sorter and then stained using the Giemsa method. Representative data of four experiments are depicted.

Data showing the identification of DC are presented in Fig. 3. There were few CD11 c^+ I-A^{high} DC in the liver, but were abundant in the spleen. However, the results clearly show that the number of DC decreased in the early stages (3–6 h) after Con A administration.

Characterization of DC in the liver and spleen

Next, the phenotypes of the DC isolated from the liver and spleen were compared (Fig. 4) after two-colour staining for CD11c and I-A (or CD11b or PDCA-1). The expression of I-A (or CD11b or PDCA-1) on CD11c DC was identified using gated analysis. The expression of I-A on liver DC

Fig. 2. Kinetics of tissue damage and the number of dendritic cells (DC) after concanavalin A administration. (a) The levels of aspartate aminotransferase (AST) and alanine aminotransferase (ALT) in sera. (b) The absolute numbers of CD11c⁺ DC cells in the liver and spleen. The mean \pm 1 standard error were produced by five experiments.

increased from 8·3% to 29·5% (Fig. 4a). The expression of CD11b also increased after administration of Con A (from 15·2% to 50·9%). In contrast, the expression of PDCA-1 decreased.

The same experiments were performed using spleen DC (Fig. 4b). A striking difference was seen in the expression of I-A and CD11b on splenic DC, which showed high expression of both molecules $(I-A^+, 88.9\%; CD11b^+, 75.3\%).$ This high I-A expression increased with time (up to 24 h), but CD11b levels remained static. The expression of PDCA-1 was very low in splenic DC, and tended to decrease with time.

Next, the costimulatory molecules expressed by DC were characterized further (Fig. 5) using two-colour staining for CD11c and CD80 (or CD86 or CD40). The expression of CD80, CD86 and CD40 by hepatic CD11c⁺ DC was determined both before and after Con A administration (Fig. 5a). In the late stages $(12-24 h)$, the expression of all these markers was increased (indicated by arrows).

The expression of CD80, CD86 and CD40 on spleen DC was high compared with that on liver DC (89·4%, 82·7% and 93·4%, respectively), and remained constant after Con A administration.

Adoptive transfer of DC into mice with Con A-induced hepatitis

The immunomodulatory function of hepatic or splenic DC was then investigated (Fig. 6). $CD11c⁺ DC$ were isolated by magnetic cell sorting and injected intravenously into mice 3 h after Con A administration (Fig. 6a). Purified CD11c⁺ DC in the liver contained > 60% pDC. However, those in the spleen contained no pDC (data not shown). The indicated parameters were then examined after 18 h. Figure 6b shows that hepatic DC, but not splenic DC, suppressed hepatic failure.

Histological examination of the liver and immunohistochemical staining of hepatic DC was then conducted (Fig. 7). The administration of Con $A + PBS$ (at 18 h) induced inflammatory cell infiltration into the liver and haemorrhage (Fig. 7b). However, this was attenuated by the adoptive transfer of hepatic DC (Fig. 7c).

Staining for I-A and CD11c showed that CD11c⁺ DC in the liver were primarily I-A⁻ (Fig. 7g,j,m). These CD11 c^+ I-A- DC disappeared after the administration of Con A (Fig. 7h,k,n). After adoptive transfer of hepatic DC, CD11c⁺ I-A- DC accumulated in the liver (Fig. 7i,l,o). These results showed that adoptively transferred DC infiltrated into the liver and were able to suppress hepatic failure.

Further characterization of cytokines and effector cells

The serum levels of proinflammatory cytokines (TNF- α and IL-12) and T helper type 1 (Th1)-type cytokines (IFN-g and IL-2) were measured in mouse serum 18 h after the adoptive

Fig. 3. Variations in the proportion of I-A⁺ CD11c⁺ dendritic cells (DC) in the liver and spleen after concanavalin A administration. Two-colour staining for CD11c and I-A was conducted. Data are representative of three independent experiments.

transfer of hepatic DC (Fig. 8). The levels of TNF- α were comparable before and after adoptive transfer of DC, but IL-12p70 levels decreased after the transfer of hepatic DC. IFN-g levels were also comparable, but IL-2 levels tended to decrease.

The major effector cells involved in hepatic damage during Con A-induced hepatitis are NK T cells. Therefore, the proportion of NK T cells was analysed 18 h after adoptive transfer (Fig. 9). NK T cells were abundant in the liver of normal mice (12·4%), but their numbers were reduced after Con A administration (2·7%). This did not change after the adoptive transfer of hepatic or splenic DC. CD4⁺ T cells are also involved in Con A-induced hepatitis, and respond directly to Con A. The proportion of CD4⁺ T cells was also down-regulated after Con A administration (from 20·4% to 13·0%). This was not changed by the adoptive transfer of hepatic DC or splenic DC.

Finally, T_{reg} cells (CD4⁺ FoxP3⁺) were examined (Fig. 9, bottom panel). The proportion of T_{reg} cells did not change significantly after the adoptive transfer of hepatic DC or splenic DC compared with that in the controls (Con $A + PBS$).

Discussion

In the present study, we first characterized the phenotype and morphology of DC in the livers of normal mice. Almost all CD11 c^+ DC in the liver were I-A $^{\rm low}$ B220⁺. This phenotype was quite different from that of splenic DC (I-A^{high} B220⁻). In other words, hepatic DC had a phenotype of pDC, which sometimes show immunosuppressive functions [4,5]. We then purified I-A^{high} CD11c⁺ cells and I-A^{low} CD11c⁺ by cell sorting and compared their morphology. Irrespective of the organ of origin, I-A^{high} CD11 c^+ cells were large and highly dendritic in nature, whereas I-Alow CD11c⁺ cells were small and less dendritic in nature. These results suggest that hepatic DC comprise pDC rather than cDC, which is in agreement with earlier studies [1–3].

An acute Con A-induced autoimmune hepatitis model was used to examine the function of hepatic DC. First, the variation in the number and proportion of DC in the liver and spleen was examined. The level of transaminases in mouse serum indicated that hepatic damage occurred as early as 3 h after Con A administration. At this early stage (3–6 h), the number of DC in the liver and spleen tended to decrease. However, at a later stage (12–24 h), the numbers increased. A similar pattern was seen when the proportion (rather than discrete numbers) of DC in the liver and spleen was assessed.

The expression of I-A and CD11b on CD11c⁺ DC in the livers of normal mice was low, but increased after Con A administration. The level of I-A and CD11b was extremely high in splenic DC, independent of Con A administration. The expression of the pDC marker, PDCA-1 [18], was also assessed. The proportion of PDCA-1⁺ cells within the hepatic DC population was slightly higher (17%) than in the splenic DC population (11%) in normal mice, although expression tended to decrease in hepatic DC after Con A administration. pDC are relatively abundant in mouse liver [4] and produce IFN- α [19]. They have also been implicated in hepatic tolerance [20]. The costimulatory molecules CD80, CD86 and CD40, expressed by DC, are associated with antigen presentation and activation. The results of this study showed that the expression of costimulatory molecules by CD11c⁺ DC in the liver increased 12–24 h after Con A administration. Activated DC in the liver also produced large amounts of the proinflammatory TNF- α , IL-6 and macrophage inflammatory protein (MIP)-1 α (data not shown). Thus, it was speculated that activated hepatic DC modulate liver damage by secreting certain cytokines.

Adoptive transfer experiments were conducted to determine directly the function of hepatic DC. Cell transfer was performed 3 h after Con A administration. The results showed that that only hepatic DC had the potential to suppress hepatic damage induced by Con A.All these phenomena were also confirmed by histology and immunohistochemical staining. Because hepatic damage is mediated by killer molecules, the levels of cytokines with killer activity, including TNF- α , IL-12 and IFN- γ , were examined. The most prominent change observed after the transfer of hepatic DC was a reduction in IL-12p70 levels. Transfer of splenic DC did not evoke such an effect. In other words, the ability of hepatic DC

Fig. 4. Further characterization of dendritic cell (DC) phenotype in (a) liver and (b) spleen. Two-colour staining for CD11c and I-A (or CD11b or PDCA-1) was conducted. Gated analysis was used to examine the proportion of DC expressing I-A, CD11b or PDCA-1. Significant changes are indicated by arrows. Representative data of five experiments are depicted.

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(a) Liver

Fig. 5. Further characterization of dendritic cell (DC) phenotype in (a) liver and (b) spleen. Two-colour staining for CD11c and CD80 (or CD86 or CD40) was conducted. Gated analysis was used to examine the proportion of DC expressing CD80, CD86 and CD40. Representative data of five experiments are depicted. Significant changes are indicated by arrows.

Fig. 6. Adoptive transfer of dendritic cells (DC). (a) Time schedule and (b) modulation of tissue damage by adoptive transfer of DC. DC were transferred 3 h after concanavalin A administration. Mean \pm 1 standard error were produced by three experiments. **P* < 0·05.

to suppress liver damage may be related directly to their ability to suppress IL-12 secretion. IL-12 exerts its biological activity mainly by promoting a Th1 immune response, and its detrimental effect during Con A-induced hepatitis occurs via this mechanism [21–24]. IL-12 promotes increased IL-4 and IFN- γ production by invariant NK T cells in the livers of mice with Con A-induced hepatitis and aggravates the disease. In other words, a reduction in IL-12 levels suppresses the activation of NK T cells and effector T cells in the liver, and reduces the level of damage induced by Con A administration. The results of the present study support this mechanism, at least in part. The levels of TNF- α and IFN- γ remained unchanged even after the transfer of hepatic DC.

Other experiments also showed that the transfer of hepatic DC suppressed IL-2 production. There is a possibility that the growth of some effector cells, which respond to IL-2, might be suppressed by the transfer of hepatic DC into mice treated with Con A. Therefore, the effector cells responsible were identified in further experiments.

It is well established that several different effector cells are associated with hepatic damage during Con A-induced hepatitis. These are primarily CD4⁺ T cells [9–11] and NK T cells [12,13]. However, we did not observe suppression of CD4⁺ T cell or NK T cell numbers after the adoptive transfer of hepatic DC.

Finally, experiments were conducted to identify the suppressor cells involved. T_{reg} cells and NK T cells both have an immunosuppressive function. T_{rec} cells regulate autoreactive T cells, and their reduction or dysfunction causes certain autoimmune diseases in both animals and humans [25,26]. T_{rec} cells produce IL-10 and transforming growth factor $(TGF)-\beta$, and these cytokines suppress T cell effector function and inflammatory disease $[27,28]$. T_{reg} cells also express cytotoxic T lymphocyte antigen 4 (CTLA-4), which inhibits effector T cell proliferation via cell–cell interactions and also inhibits IL-2 production [29]. NK T cells also suppress certain autoimmune diseases, and their absence is associated with autoimmune phenomena [30–33]. There is a possibility that the transfer of hepatic DC might influence the induction of suppressor cells, rather than that of effector cells [15,16]. However, we did not detect any increase in T_{reg} cells or NK T cell numbers after the transfer of hepatic DC.

These results suggest that immunosuppression by hepatic DC was not due to their effects on effector cells or suppressor cells. Therefore, it appears that functional modulation of killer molecules may be the main mechanism by which hepatic DC suppress liver damage. In parallel with the assay of several cytokines, the level of IL-10 was also examined in sera. The level of IL-10 was not detectable with or without the transfer of hepatic DC in Con A-induced hepatitis (data not shown). We have to consider that some factors other than IL-10 might be used for the immunosuppression (down-regulation of IL-12 and IL-2) by hepatic DC in Con A-induced hepatitis.

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Disclosures

The authors have no conflicts of interest to declare.

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Fig. 7. Histology and immunohistochemical staining of the liver. (a, d, g, j, m) Control mice. (b, e, h, k, n) Mice treated with concanavalin A (Con A). (c, f, i, l, o) Mice treated with Con A plus adoptive transfer of liver dendritic cells (DC). (a–c) Haematoxylin and eosin staining. (d–o) Immunohistochemical staining. White arrowheads indicate the CD11c single positive cells. They were only seen in the control liver and the DC-post injected Con A-treated liver, but not in the Con A-treated liver. Representative data of three experiments are depicted.

Fig. 8. Cytokine production after adoptive transfer of dendritic cells (DC). The level of cytokines was determined 24 h after concanavalin A administration. Data represent the mean \pm 1 standard error from four experiments. **P* < 0·05.

Fig. 9. Identification of lymphocyte subsets by immunofluorescence analysis. Natural killer (NK) T cells were NK1·1⁺ CD3^{int}. CD4⁺ T cells were CD4⁺ CD8⁻. Regulatory T cells were CD4⁺ forkhead box P3⁺ . Data are representative of three independent experiments.

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