

damage-associated molecules, such as HMGB1, heat shock protein and ATP, leads to the release of inflammatory cytokines. IL-1, TNF- $\alpha$  and, to a lesser extent, IL-6 stimulate the HPA axis at the level of vagus afferent nerves, the hypothalamus, the pituitary and the adrenal glands to release GC as well as the SNS to release NA, providing a negative feedback loop to stop inflammation. Cytokines produced in the periphery activate primary afferent nerves, such as the vagus nerves, enter the brain through the areas with a poorly developed blood-brain barrier, such as the circumventricular region or are actively transported. In addition, neurons, glial cells and endothelial cells produce these cytokines in the CNS.<sup>153</sup> In contrast to HPA axis, inflammatory cytokines have negative effects on the HPG axis, leading to the reduction of gonadal functions.<sup>154</sup>

The effects of cytokines on the CNS are not limited to the HPA axis and SNS, but are also involved in behavior induced by sickness including changes in behaviour that occur in ill patients, and even in depression. Systemic or intrathecal administration of IL-1 $\beta$  or TNF- $\alpha$  induces signs of behavior resulting from sickness, such as decreased motor activity, social withdrawal, altered cognition and fatigue. Although administration of IL-6 does not induce these behavioral changes, LPS-induced sickness behaviors are reduced in IL-6<sup>-/-</sup> mice, suggesting its involvement in these behavioral changes, although the degree is less compared with IL-1 $\beta$  or TNF- $\alpha$ . In contrast, anti-inflammatory cytokines, such as IL-10, attenuate LPS-induced sickness behaviors.<sup>155-158</sup>

Type I interferon, IFN- $\alpha$  and IFN- $\beta$ , are used for the treatment of hepatitis C and MS, respectively. These cytokines show neuropsychiatric complications, including sleep disorders and depression, which serves as evidence of the cytokine-mediated modulation of neural activities. The potential link between inflammatory cytokines and depression is tryptophan metabolism. Tryptophan is an essential amino acid and is a source of serotonin, as it is metabolized to serotonin and kynurenine. In the latter pathway, tryptophan is metabolized by tryptophan 2,3 dioxygenase (TDO) and indoleamine 2,3 dioxygenase (IDO) to kynurenine, and then metabolized either to 3-hydroxykynurenine or kynurenic acid, an antagonist of NMDA receptors. 3-Hydroxykynurenine is further metabolized to 3-hydroxyanthranilic acid and quinolinic acid, an agonist of NMDA receptors. TDO is primarily located in the liver and is activated by GC, whereas IDO is widely expressed and is activated by inflammatory cytokines and downregulated by IL-4.<sup>153,159</sup> In patients treated

with type I IFN, plasma levels of kyurenic acid as well as serotonin are decreased, suggesting that the behavior and depression caused by inflammatory cytokines including IFN might be a result of an alteration in glutamatergic neurotransmission.

Although much of the focus on T cells in a variety of pathogenic conditions, has been on classical immune-mediated inflammation, including infection and autoimmune disorders, in diseases newly related to inflammation, such as ischemia, neurodegenerative and psychiatric disorders, a neuroprotective role has emerged as an important task for T cells. The production of neurotrophic factors, such as brain-derived neurotrophic factor (BDNF) from T cells,<sup>160</sup> reduced learning capacity in T cell-deficient mice and its restoration by passive T cell transfer<sup>161</sup> and enhanced hippocampal neurogenesis by T cells,<sup>162,163</sup> suggest a fundamental function of T cells in the maintenance of cognitive functions. Anti-inflammatory cytokines, such as IL-4 and TGF- $\beta$ , are detectable in the CNS, and IL-4 is downregulated in a mouse model of Alzheimer's disease.<sup>164</sup> These cytokines, in addition to BDNF, might contribute to the maintenance of homeostasis of the CNS. Although the precise mechanisms remain elusive, T cells might serve as important players in the maintenance of neuronal integrity.

#### Future directions

Acute stress responses in the autonomic and peripheral nervous systems amplify local immune responses to eliminate pathogens and other dangerous occurrences. Subsequent to these initial responses, the neuroendocrine and autonomic systems act to inhibit immune responses and terminate inflammation. In contrast, chronically sustained stress induces unusual conditions, such as inadequate secretion of GC, as well as resistance to GC, increased sympathetic tone propelling the RAAS, functional loss of sympathetic nerve fibers at the inflammation site and a local  $\beta$  to  $\alpha$  adrenergic shift.<sup>165,166</sup> Further studies to clarify the consequences of stress on chronic inflammatory conditions will provide novel strategies for the control of complex pathogenic conditions including autoimmune diseases, and neurodegenerative and psychiatric disorders.

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## Mucosal-Associated Invariant T Cells Promote Inflammation and Exacerbate Disease in Murine Models of Arthritis

Asako Chiba, Ryohsuke Tajima, Chiharu Tomi, Yusei Miyazaki, Takashi Yamamura, and Sachiko Miyake

**Objective.** The function of mucosal-associated invariant T (MAIT) cells remains largely unknown. We previously reported an immunoregulatory role of MAIT cells in an animal model of multiple sclerosis. The aim of this study was to use animal models to determine whether MAIT cells are involved in the pathogenesis of arthritis.

**Methods.**  $MR1^{-/-}$  and  $MR1^{+/+}$  DBA/1J mice were immunized with bovine type II collagen (CII) in complete Freund's adjuvant to trigger collagen-induced arthritis (CIA). To assess CII-specific T cell recall responses, lymph node cells from mice with CIA were challenged with CII *ex vivo*, and cytokine production and proliferation were evaluated. Serum levels of CII-specific antibodies were measured by enzyme-linked immunosorbent assay. Collagen antibody-induced arthritis (CAIA) was induced in  $MR1^{-/-}$  and  $MR1^{+/+}$  C57BL/6 mice by injection of anti-CII antibodies followed by injection of lipopolysaccharide. To demonstrate the involvement of MAIT cells in arthritis, we induced CAIA in  $MR1^{-/-}$  C57BL/6 mice that had been reconstituted with adoptively transferred MAIT cells. MAIT cell activation in response to cytokine stimulation was investigated.

**Results.** The severity of CIA was reduced in  $MR1^{-/-}$  DBA/1J mice. However, T and B cell responses

to CII were comparable in  $MR1^{-/-}$  and  $MR1^{+/+}$  DBA/1J mice.  $MR1^{-/-}$  C57BL/6 mice were less susceptible to CAIA, and reconstitution with MAIT cells induced severe arthritis in  $MR1^{-/-}$  C57BL/6 mice, demonstrating an effector role of MAIT cells in arthritis. MAIT cells became activated upon stimulation with interleukin-23 (IL-23) or IL-1 $\beta$  in the absence of T cell receptor stimuli.

**Conclusion.** These results indicate that MAIT cells exacerbate arthritis by enhancing the inflammation.

Rheumatoid arthritis (RA) is an autoimmune disease characterized by chronic inflammation in the joints. It has been suggested that environmental factors influence autoimmunity, and in particular, increasing evidence highlights the important role of gut flora in the development of autoimmune diseases (1), including arthritis. For example, differences in the intestinal microbiota of patients with early RA have been described, and tetracycline treatment was shown to reduce disease activity in RA (2,3). In addition, oral vancomycin treatment significantly decreased the severity of adjuvant-induced arthritis (4). More recently, it was demonstrated that germ-free conditions strongly inhibit arthritis in the K/BxN arthritis model and that the introduction of segmented filamentous bacteria induced severe arthritis in germ-free K/BxN mice (5). Thus, mucosal immunity plays an important role in the development and progression of arthritis.

Natural killer (NK) cells, invariant NK T (iNKT) cells,  $\gamma/\delta$  T cells, mucosal-associated invariant T (MAIT) cells, B-1 B cells, and marginal-zone B cells are categorized as innate-like lymphocytes. Such lymphocytes reside in unique locations, including the marginal zone of the spleen and epithelial and mucosal tissues and rapidly exert effector functions in the absence of clonal expansion (6–15). Therefore, these innate-like lymphocytes are thought to play important roles in “first-line” im-

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Asako Chiba, MD, PhD, Ryohsuke Tajima, MS, Chiharu Tomi, Yusei Miyazaki MD, PhD, Takashi Yamamura MD, PhD, Sachiko Miyake MD, PhD: National Institute of Neuroscience, National Centre of Neurology and Psychiatry, Tokyo, Japan.

Address correspondence to Sachiko Miyake MD, PhD, Department of Immunology, National Institute of Neuroscience, National Centre of Neurology and Psychiatry, 4-1-1 Ogawahigashi, Kodaira, Tokyo 187-8502, Japan. E-mail: miyake@ncnp.go.jp.

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mune responses against exogenous stimuli. As MAIT cells are preferentially located in the gut lamina propria, there is a growing interest in the function of MAIT cells in various types of immune responses, including autoimmunity (16–20).

MAIT cells are restricted by a nonpolymorphic class IB major histocompatibility complex (MHC) molecule, the class I MHC-related molecule (MR1), and express an invariant T cell receptor (TCR)  $\alpha$ -chain: V $_{\alpha}$ 7.2–J $_{\alpha}$ 33 in humans and V $_{\alpha}$ 19–J $_{\alpha}$ 33 in mice. The invariant TCR $\alpha$  chain associates with a limited set of V $_{\beta}$  chains (14,21,22). MAIT cells are selected in the thymus in an MR1-dependent manner, but, interestingly, MAIT cells require B cells as well as commensal flora for their peripheral expansion (14,23). Our group previously demonstrated a protective role of MAIT cells against autoimmune encephalomyelitis (EAE), an animal model of human multiple sclerosis. The suppression of EAE was accompanied by increased production of interleukin-10 (IL-10) by B cells, which was induced in part by ICOS costimulation (17). Because the invariant V $_{\alpha}$ 7.2–J $_{\alpha}$ 33 TCR is highly expressed in central nervous system lesions of multiple sclerosis patients, human MAIT cells may also be involved in the pathogenesis of multiple sclerosis (16).

In addition to their regulatory function, MAIT cells also possess proinflammatory functions like other innate-like lymphocytes. Le Bourhis et al (20) demonstrated that MAIT cells display antimicrobial capacity. Both human and mouse MAIT cells are activated by *Escherichia coli*-infected antigen-presenting cells in an MR1-dependent manner. MAIT cells show a protective role against *Mycobacterium abscessus* or *E coli* infections in mice. Human MAIT cells are capable of producing interferon- $\gamma$  (IFN $\gamma$ ) and IL-17 and are found in *Mycobacterium tuberculosis*-infected lung tissues. Thus, MAIT cells play an antimicrobial function under these infectious conditions. Although accumulating evidence suggests that certain subsets of innate-like lymphocytes, such as NK cells, iNKT cells, and  $\gamma/\delta$  T cells, are involved in the pathogenesis of arthritis in animal models of the disease, the role of MAIT cells in arthritis remains unknown (24–31).

We report herein that MAIT cells play a pathogenic role in murine models of arthritis. The disease severity of collagen-induced arthritis (CIA) in MAIT cell-deficient MR1 $^{-/-}$  DBA/1J mice was ameliorated compared with that of MR1 $^{+/+}$  DBA/1J mice. However, T cell responses to type II collagen (CII) and CII-specific serum antibody levels were comparable between CIA-induced MR1 $^{-/-}$  and MR1 $^{+/+}$  DBA/1J mice. We found that MR1 $^{-/-}$  C57BL/6J mice are much less suscep-

tible to collagen antibody-induced arthritis (CAIA) as compared to MR1 $^{+/+}$  C57BL/6J mice. MR1 $^{-/-}$  C57BL/6J mice reconstituted with adoptively transferred MAIT cells developed severe arthritis, suggesting that MAIT cells may be one of the effectors contributing to inflammation in arthritis. Finally, we investigated the cytokine-producing capacity of MAIT cells. No differences in IFN $\gamma$  production by liver mononuclear cells (LMNCs) from MR1 $^{-/-}$  C57BL/6J and MR1 $^{+/+}$  C57BL/6J mice were observed upon TCR stimulation, but the level of IL-17 produced by LMNCs from MR1 $^{+/+}$  C57BL/6J mice was much higher than that produced by cells from MR1 $^{-/-}$  C57BL/6J mice. We further demonstrated that sorted murine MAIT cells produce IL-17 upon TCR engagement. Surprisingly, IL-17 production by MAIT cells was observed after exposure to IL-23 without TCR stimulation, and IL-1 $\beta$  alone induced proliferation of MAIT cells, indicating that MAIT cells may be activated by cytokines and may enhance the inflammation in arthritis.

## MATERIALS AND METHODS

**Mice.** DBA/1J mice were purchased from the Oriental Yeast Company. C57BL/6J mice were obtained from CLEA Laboratory Animal Corporation. MR1 $^{-/-}$  mice (14) were provided by S. Gilfillan (Department of Pathology and Immunology, Washington University School of Medicine, St. Louis, MO), and V $_{\alpha}$ 19i-transgenic mice (32) on a C57BL/6J background were provided by M. Shimamura (University of Tsukuba, Ibaraki, Japan). MR1 $^{-/-}$  mice were backcrossed to DBA/1J mice for 10 generations to obtain MR1 $^{-/-}$  DBA/1J mice. V $_{\alpha}$ 19i-transgenic CD1d1 $^{-/-}$  C57BL/6J mice were generated by backcrossing V $_{\alpha}$ 19i-transgenic mice with CD1d1 $^{-/-}$  C57BL/6J mice for 7 generations. Mice were maintained under specific pathogen-free conditions in accordance with institutional guidelines and used in the experiments at 7–12 weeks of age.

**Induction of CIA.** Both MR1 $^{-/-}$  DBA/1J mice and their littermate controls (MR1 $^{+/+}$  DBA/1J mice) (n = 5–6 per group; ages 7–8 weeks old) were immunized intradermally at the base of the tail with 150  $\mu$ g of CII (Collagen Research Center) emulsified with an equal volume of complete Freund's adjuvant containing 250  $\mu$ g of heat-killed *Mycobacterium tuberculosis* H37Ra (Difco). Three weeks after the primary immunization, mice were given an intradermal booster injection of 150  $\mu$ g of CII emulsified in incomplete Freund's adjuvant (Difco).

**Induction of CAIA.** MR1 $^{-/-}$  C57BL/6J mice and their littermate controls (MR1 $^{+/+}$  C57BL/6J mice) were injected intravenously with a mixture of anti-CII monoclonal antibodies (mAb) (Arthrogen-CIA mAb, 2 mg; Chondrex) followed 2 days later by an intraperitoneal injection of 50  $\mu$ g of lipopolysaccharide.

**Clinical assessment of arthritis.** Mice were examined for signs of joint inflammation, which was scored on a scale of 0–4, where 0 = no change, 1 = significant swelling and redness

of 1 digit, 2 = mild swelling and erythema of the limb or swelling of  $\geq 2$  digits, 3 = marked swelling and erythema of the limb, and 4 = maximal swelling and redness of the limb and later, ankylosis. The average macroscopic score was expressed as a cumulative value for all paws, with a maximum possible score of 16.

**Histopathologic assessment.** Arthritic mice were killed, and all 4 paws were fixed in buffered formalin, decalcified, embedded in paraffin, sectioned, and then stained with hematoxylin and eosin. Histologic assessment of joint inflammation was scored on a scale of 0–3 as follows: 0 = normal joint, 1 = mild arthritis (minimal synovitis without cartilage/bone erosion), 2 = moderate arthritis (synovitis and erosion but joint architecture maintained), and 3 = severe arthritis (synovitis, erosion, and loss of joint integrity). The average of the macroscopic scores was expressed as a cumulative value for all paws, with a maximum possible score of 12.

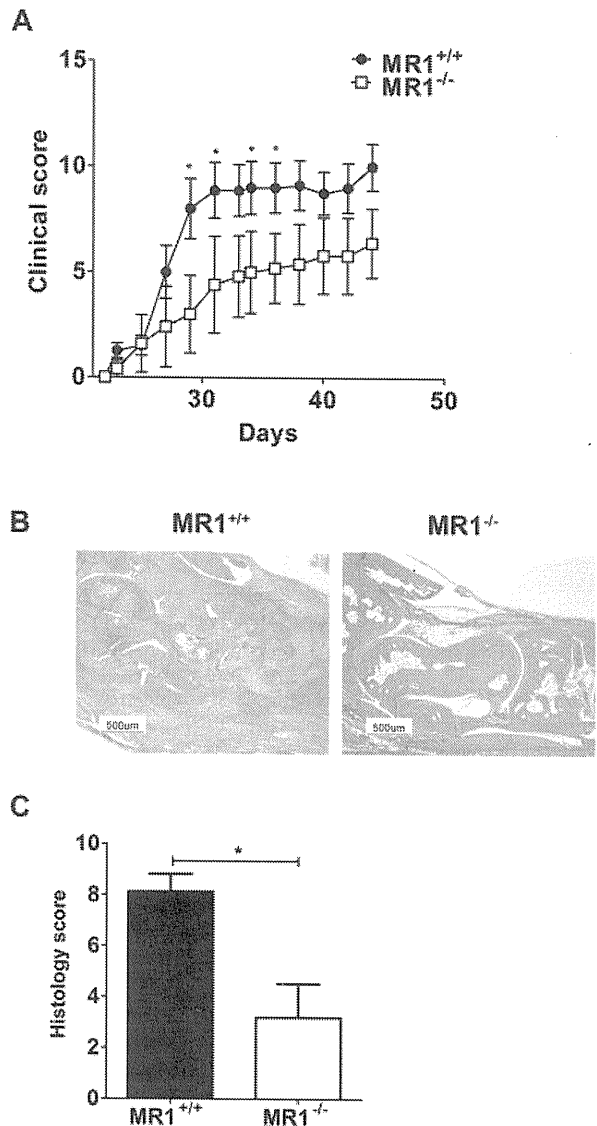
**CII-specific T cell response.** Lymph node cells were collected on days 35–42 after immunization and suspended in complete RPMI 1640 medium (Life Technologies) containing 1% syngeneic mouse serum. The cells were cultured for 72 hours in 96-well flat-bottomed plates at a density of  $1 \times 10^6$ /well in the presence of CII. Proliferative responses were measured using a  $\beta$ -1205 counter (Pharmacia) to detect the incorporation of  $^3\text{H}$ -thymidine (1  $\mu\text{Ci}$ /well) during the final 16 hours of culture.

**Measurement of CII-specific total IgG, IgG1, and IgG2a.** Bovine CII (1 mg/ml) was coated onto enzyme-linked immunosorbent assay (ELISA) plates (Sumitomo Bakelite) overnight at 4°C. After blocking with 1% bovine serum albumin in PBS, serially diluted serum samples were added to CII-coated wells. For detection of anti-CII antibodies, the plates were incubated with biotin-labeled anti-IgG1 and anti-IgG2a (SouthernBiotech) or anti-IgG antibody (CN/Cappel) for 1 hour and were then incubated with streptavidin-peroxidase. After adding substrate, the reaction was evaluated as the optical density values at 450 nm ( $\text{OD}_{450}$ ).

**Adoptive transfer and in vitro stimulation of  $V_{\alpha}19i$  T cells.** LMNCs were purified from  $V_{\alpha}19i$ -transgenic  $\text{CD1d1}^{-/-}$   $\text{C57BL/6J}$  mice by use of Percoll density-gradient centrifugation, and erythrocytes and B cells were depleted with phycoerythrin (PE)-conjugated anti-Ter-119 and PE-conjugated anti-CD19 (BD) followed by separation with anti-PE-conjugated magnetic-activated cell sorter beads (Miltenyi Biotec). Cells were stained with fluorescein isothiocyanate-conjugated anti-TCR $\beta$  and PerCP-Cy5.5 anti-NK1.1 (BD), and TCR $\beta$ + NK1.1+ cells were sorted using a FACSAria cell sorter (BD). The purity of isolated NK1.1+ T cells (MAIT cells) was  $>95\%$ , as assessed by flow cytometry.

In adoptive transfer experiments,  $5 \times 10^5$  MAIT cells or NK1.1- T cells (T cells) were injected intravenously into naive  $\text{MR1}^{-/-}$   $\text{C57BL/6}$  recipient mice 1 day before administration of CII mAb. LMNCs or sorted MAIT cells were resuspended in RPMI 1640 medium supplemented with 10% fetal bovine serum, 2 mM L-glutamine, 50 units/ml of penicillin/streptomycin, and 55  $\mu\text{M}$   $\beta$ -mercaptoethanol (Life Technologies) and stimulated with immobilized anti-CD3 mAb (2C11, 1  $\mu\text{g}/\text{ml}$ ) and/or the following cytokines: IL-1 $\beta$ , tumor necrosis factor  $\alpha$  (TNF $\alpha$ ), IL-6, and transforming growth factor  $\beta$  (TGF $\beta$ ) (all from PeproTech) and IL-23 (R&D Systems).

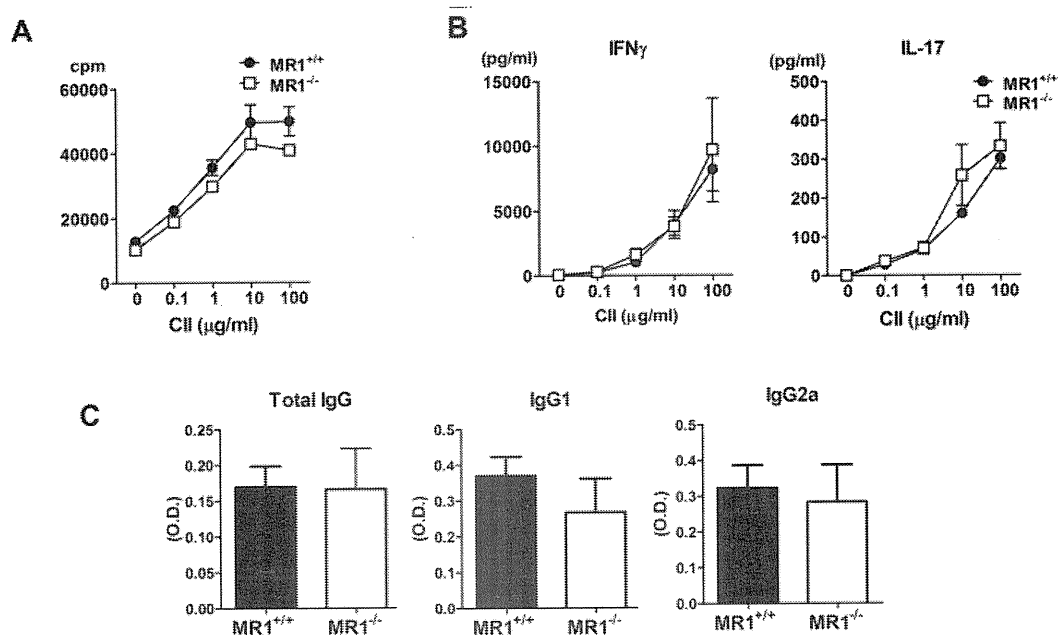
**Detection of cytokines.** Cytokine levels in the culture supernatant were determined using a sandwich ELISA. The



**Figure 1.** Amelioration of collagen-induced arthritis (CIA) in  $\text{MR1}^{-/-}$  mice. **A**, Clinical scores for CIA in  $\text{MR1}^{-/-}$  DBA/1J mice and in  $\text{MR1}^{+/+}$  DBA/1J mice. Values are the mean  $\pm$  SEM of 5–8 mice per group. \* =  $P < 0.05$  versus  $\text{MR1}^{-/-}$  DBA/1J mice. **B**, Representative histologic sections of the joints of  $\text{MR1}^{+/+}$  DBA/1J mice and  $\text{MR1}^{-/-}$  DBA/1J mice. Hematoxylin and eosin stained; original magnification  $\times 40$ . **C**, Histology scores in  $\text{MR1}^{-/-}$  DBA/1J mice and in  $\text{MR1}^{+/+}$  DBA/1J mice, expressed as the sum of the scores in the 4 paws. Results from a single representative experiment of 2 similar experiments performed are shown. Values are the mean  $\pm$  SEM. \* =  $P < 0.05$ .

ELISA antibodies for IFN $\gamma$  were purchased from BD. Levels of IL-17 were determined using an IL-17 ELISA kit (R&D Systems).

**Statistical analysis.** Clinical or pathologic scores for CIA and CAIA in the various groups of mice are presented as



**Figure 2.** Type II collagen (CII) responses in MR1<sup>-/-</sup> DBA/1J mice. A and B, Inguinal lymph node cells from MR1<sup>-/-</sup> DBA/1J mice and MR1<sup>+/+</sup> DBA/1J mice with collagen-induced arthritis were incubated for 48 hours in the presence of CII. Proliferative responses were determined by the uptake of <sup>3</sup>H-thymidine (A), and the levels of interferon- $\gamma$  (IFN $\gamma$ ) and interleukin-17 (IL-17) in culture supernatants were measured by enzyme-linked immunosorbent assay (B). C, CII-specific antibody levels in individual serum samples obtained at the end of the experiment were analyzed as described in Materials and Methods. Results from a single representative experiment of 2 similar experiments performed are shown. Values are the mean  $\pm$  SEM of 5–8 mice per group. OD = optical density

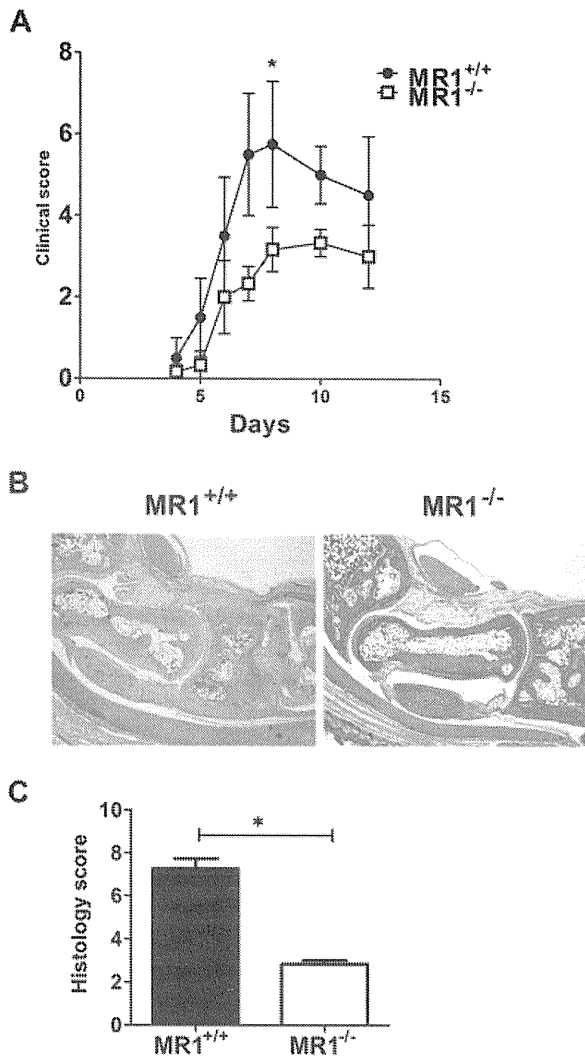
the mean  $\pm$  SEM clinical score for the group, and statistical differences were analyzed with a nonparametric Mann-Whitney U test. Data for cytokines and proliferation were analyzed with an unpaired *t*-test.

## RESULTS

**Amelioration of CIA in MR1<sup>-/-</sup> mice.** To investigate whether MAIT cells play a role in the pathogenesis of arthritis, we first evaluated the involvement of MAIT cells in CIA using MR1<sup>-/-</sup> mice lacking MAIT cells. Because DBA/1J mice bearing the H-2q haplotype are the most susceptible strain for CIA, MR1<sup>-/-</sup> C57BL/6J mice were backcrossed to DBA/1J mice for 10 generations to obtain MR1<sup>-/-</sup> DBA/1J mice. Both MR1<sup>-/-</sup> DBA/1J mice and littermate MR1<sup>+/+</sup> DBA/1J mice were immunized with CII to induce CIA, and the clinical severity of arthritis was evaluated by visual scoring of each paw. As shown in Figure 1A, the clinical scores in MR1<sup>-/-</sup> DBA/1J mice were reduced in comparison to those in MR1<sup>+/+</sup> DBA/1J mice. Histologic examination of the joints of the 4 paws 44 days after CIA induction showed less cell infiltration, cartilage erosion, and bone

destruction in MR1<sup>-/-</sup> DBA/1J mice than in the MR1<sup>+/+</sup> DBA/1J mice (Figure 1B). Quantification of the histologic severity of arthritis revealed that MR1<sup>-/-</sup> DBA/1J mice developed milder joint inflammation than MR1<sup>+/+</sup> DBA/1J mice (Figure 1C). These results suggest that MAIT cells contribute to the exacerbation of the disease course of CIA.

**CII responses in MR1<sup>-/-</sup> DBA/1J mice.** As the presence of MAIT cells augmented the severity of CIA, we next asked whether MAIT cells influence the CII-specific responses of T and B cells. Lymph node cells from CIA-induced animals were rechallenged with CII *ex vivo*. As shown in Figure 2A, the proliferative responses of lymph node cells upon stimulation with CII were similar in the two groups. Lymph node cells from both MR1<sup>-/-</sup> DBA/1J mice and MR1<sup>+/+</sup> DBA/1J mice produced comparable amounts of IL-17 and IFN $\gamma$  in response to CII in a dose-dependent manner (Figure 2B). We also evaluated CII-specific immunoglobulin levels in serum obtained 35–42 days after arthritis induction. We observed a trend of reduced levels of CII-specific IgG1 in MR1<sup>-/-</sup> DBA/1J mice compared to the

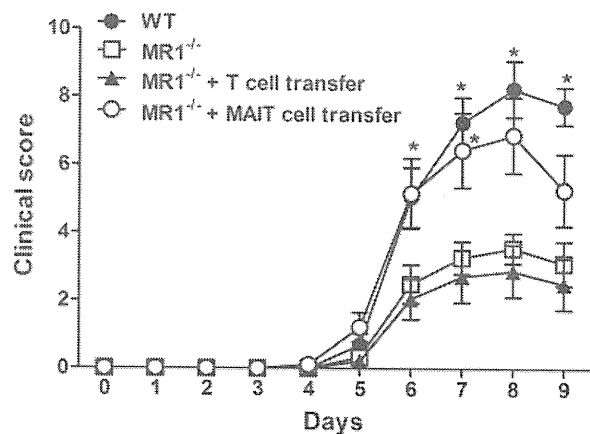


**Figure 3.** Amelioration of collagen antibody-induced arthritis (CAIA) in MR1<sup>-/-</sup> mice. **A**, Clinical scores for CAIA in MR1<sup>-/-</sup> C57BL/6J mice and MR1<sup>+/+</sup> C57BL/6J mice. Values are the mean ± SEM of 4–6 mice per group. \* = *P* < 0.05 versus MR1<sup>-/-</sup> C57BL/6J mice. **B**, Representative histologic sections of the joints of MR1<sup>+/+</sup> C57BL/6J mice and MR1<sup>-/-</sup> C57BL/6J mice. Hematoxylin and eosin stained; original magnification × 40. **C**, Histology scores in MR1<sup>-/-</sup> C57BL/6J mice and in MR1<sup>+/+</sup> C57BL/6J mice, expressed as the sum of the scores in the 4 paws. Results from a single representative experiment of 2 similar experiments performed are shown. Values are the mean ± SEM. \* = *P* < 0.05.

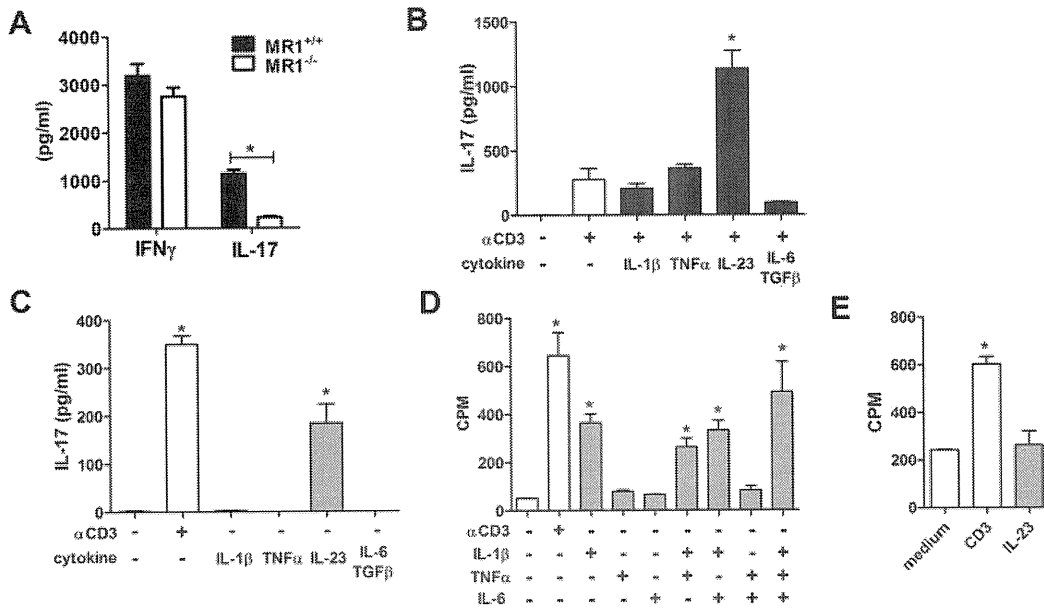
levels in MR1<sup>+/+</sup> DBA/1J mice, but the difference did not reach statistical significance (Figure 2C). These results indicate that the presence of MAIT cells has little effect on CII-specific responses.

**Amelioration of CAIA in MR1<sup>-/-</sup> mice.** The CIA model requires both adaptive and innate immune responses for disease development, and T cells and B cells responding to CII are the major players in the initiation of the disease. Although we observed significant differences in both the clinical and pathologic severity of arthritis when comparing MR1<sup>-/-</sup> DBA/1J mice to MR1<sup>+/+</sup> DBA/1J mice (Figure 1), the CII-specific responses of T and B cells appeared not to depend on the presence of MAIT cells (Figure 2). Thus, we hypothesized that MAIT cells may influence the effector phase of arthritis. To test this hypothesis, we induced CAIA in MR1<sup>-/-</sup> and MR1<sup>+/+</sup> C57BL/6J mice. By 7 days after injection of anti-CII mAb, MR1<sup>+/+</sup> C57BL/6J mice had developed severe arthritis, as assessed by clinical scores (Figure 3A). In contrast, the clinical scores in the MR1<sup>-/-</sup> C57BL/6J mice were lower compared to those in the MR1<sup>+/+</sup> C57BL/6J mice. Histologic assessment 10 days after arthritis induction revealed severe arthritis with leukocyte infiltration, synovial hyperplasia, pannus formation, cartilage erosion, and bone destruction in MR1<sup>+/+</sup> C57BL/6J mice, whereas these features were milder in MR1<sup>-/-</sup> C57BL/6J mice (Figures 3B and C).

**Augmentation of arthritis in MR1<sup>-/-</sup> mice by adoptive transfer of MAIT cells.** To demonstrate that MAIT cells actually enhance disease severity in the



**Figure 4.** Augmentation of arthritis by adoptive transfer of mucosal-associated invariant T (MAIT) cells in MR1<sup>-/-</sup> mice. MR1<sup>-/-</sup> C57BL/6J mice received  $5 \times 10^5$  NK1.1+TCR $\beta$ + T cells (MAIT cells) or an equal number of NK1.1–TCR $\beta$ + cells (T cells) from V $\alpha$ 19-transgenic CD1d1<sup>-/-</sup> mice. One day later, collagen antibody-induced arthritis was induced in wild-type (WT) C57BL/6J mice, MR1<sup>-/-</sup> C57BL/6J mice, and MR1<sup>-/-</sup> C57BL/6J mice reconstituted with T cells or MAIT cells. Results pooled from 2 similar experiments performed are shown. Values are the mean ± SEM of 8–10 mice per group. \* = *P* < 0.05 versus MR1<sup>-/-</sup> C57BL/6J mice.



**Figure 5.** Cytokine-mediated mucosal-associated invariant T (MAIT) cell activation. **A**, Liver mononuclear cells from MR1<sup>+/+</sup> C57BL/6J mice and MR1<sup>-/-</sup> C57BL/6J mice were stimulated for 48 hours with immobilized anti-CD3 ( $\alpha$ CD3) monoclonal antibody (mAb). The levels of interferon- $\gamma$  (IFN $\gamma$ ) and interleukin-17 (IL-17) in culture supernatants were measured by enzyme-linked immunosorbent assay (ELISA). \* =  $P < 0.05$ . **B**, MAIT cells were stimulated for 48 hours with immobilized anti-CD3 mAb, with or without IL-1 $\beta$ , tumor necrosis factor  $\alpha$  (TNF $\alpha$ ), IL-23, or IL-6 plus transforming growth factor  $\beta$  (TGF $\beta$ ), and the levels of IL-17 were measured by ELISA. \* =  $P < 0.05$  versus anti-CD3 mAb stimulation alone. **C**, MAIT cells were stimulated with immobilized anti-CD3 mAb or the indicated cytokines, and IL-17 levels were measured. \* =  $P < 0.05$  versus unstimulated control. **D** and **E**, Proliferative responses after 48 hours of exposure to the indicated cytokines were determined as the uptake of <sup>3</sup>H-thymidine. Results from a single representative experiment of 2 similar experiments performed are shown. \* =  $P < 0.05$  versus unstimulated control. Values in A–E are the mean  $\pm$  SEM.

CAIA model, we performed adoptive transfer experiments. Most NK1.1<sup>+</sup> TCR $\beta$  T cells within liver lymphocytes from CD1d1<sup>+/+</sup> mice are iNKT cells, and we and other investigators previously demonstrated that the NK1.1<sup>+</sup> TCR $\beta$  T cell population in V $\alpha$ 19i-transgenic CD1d1<sup>-/-</sup> mice is highly enriched in V $\alpha$ 19i TCR $\beta$  T cells (15,17). Thus, to obtain MAIT cells, we isolated NK1.1<sup>+</sup> TCR $\beta$  T cells from V $\alpha$ 19i-transgenic CD1d1<sup>-/-</sup> mice. We adoptively transferred these MAIT cells into MR1<sup>-/-</sup> C57BL/6J mice, and 1 day later, we injected these mice with anti-CII mAb to induce CAIA. MR1<sup>-/-</sup> C57BL/6J mice reconstituted with MAIT cells developed severe arthritis at a level similar to that of wild-type (WT) C57BL/6J mice (Figure 4). However, the transfer of an equal number of T cells into MR1<sup>-/-</sup> C57BL/6J mice had little effect on the clinical arthritis scores. Taken together, these results suggest that the presence of MAIT cells augmented arthritis mainly by enhancing the inflammation in arthritis.

**Cytokine-mediated MAIT cell activation.** To understand the mechanism by which MAIT cells exacerbate

the disease course of arthritis, we first compared the cytokine-producing capacity of T cells from MR1<sup>-/-</sup> and WT C57BL/6J mice. Upon anti-CD3 mAb stimulation, LMNCs from MR1<sup>-/-</sup> and WT C57BL/6J mice produced comparable amounts of IFN $\gamma$ . However, the level of IL-17 was lower in MR1<sup>-/-</sup> C57BL/6J mice than in WT C57BL/6J mice (Figure 5A).

It was recently demonstrated that human MAIT cells express the Th17-associated transcription factor retinoic acid receptor–related orphan nuclear receptor (ROR) and produce high levels of IL-17 (33). We therefore sought to determine whether mouse MAIT cells produce IL-17, which is known to play a pathogenic role in arthritis. MAIT cells were sorted from LMNCs obtained from V $\alpha$ 19i-transgenic CD1d1<sup>-/-</sup> mice and were stimulated *ex vivo* with anti-CD3 mAb. As previously shown (34), MAIT cells produced large amounts of IL-17. In addition, IL-17 production by anti-CD3 mAb-stimulated MAIT cells was augmented in the presence of IL-23 (Figure 5B).

Innate-like lymphocytes such as iNKT cells and

$\gamma/\delta$  T cells are known to be activated by cytokines directly, without TCR stimulation. A combination of IL-12 and IL-18 activates iNKT cells to produce IFN $\gamma$ , and IL-1 together with IL-23 induces IL-17 production by  $\gamma/\delta$  T cells (31,35,36). We therefore next asked whether MAIT cells are activated directly by cytokines. MAIT cells were incubated with various cytokines without TCR stimulation, and cytokine concentrations in the culture supernatants were evaluated. Surprisingly, MAIT cells produced high levels of IL-17 after exposure to IL-23 in the absence of TCR stimulation (Figure 5C).

Inflammatory cytokines such as IL-1 $\beta$ , TNF $\alpha$ , and IL-6 play critical roles in arthritis models and in human RA. Therefore, we next tested whether MAIT cells could be activated by these cytokines. As shown in Figure 5D, IL-1 $\beta$  induced robust proliferation of MAIT cells, although cytokine production was not observed after exposure to these cytokines, including IL-1 $\beta$  (data not shown). In addition, IL-23 did not induce proliferation of MAIT cells (Figure 5E). Thus, in the absence of TCR stimuli, IL-1 $\beta$  induced the proliferation of MAIT cells and IL-23 promoted the production of IL-17 by MAIT cells.

## DISCUSSION

Previous studies by our group as well as others revealed that iNKT cells play pathogenic roles in CIA and CAIA by inducing a Th1 or Th17 shift of auto-immune T cells and by augmenting the inflammation in arthritis (25–27). In the present study, we demonstrated that MAIT cells contribute to the severity of CIA and CAIA mostly by augmenting joint inflammation during the effector phase of arthritis. MR1 $^{-/-}$  mice were originally generated on the 129P2 background. Although MR1 $^{-/-}$  mice were backcrossed onto C57BL/6 or DBA/1J, we are not able to exclude the possibility that some residual sequence from the 129P2 mice affects the arthritis susceptibility of MR1 $^{-/-}$  C57BL/6 and MR1 $^{-/-}$  DBA/1J mice. However, since the reconstitution of MAIT cells induced severe CAIA in MR1 $^{-/-}$  C57BL/6 mice, the phenotype observed in MR1 $^{-/-}$  mice seems to be dependent on the lack of MAIT cells.

It has been revealed that there are CD1d-restricted T cells that are different from iNKT cells and do not express an invariant TCR $\alpha$  chain (V $\alpha$ 14–J $\alpha$ 18 in mice and V $\alpha$ 24–J $\alpha$ 18 in humans). Such CD1d-restricted T cells are called type II NKT cells and possess different functions from iNKT cells. Recently, CD1d-restricted NKT cells, which recognize murine type II collagen peptide 707–721, were reported to suppress CIA (37). It is not known whether there are distinct subsets with

different functions among MAIT cells or whether there are other T cells that are restricted by the MR1 molecule. As adoptively transferred V $\alpha$ 19i T cells augmented CAIA in MR1 $^{-/-}$  mice, MAIT cells include the population that enhances the inflammation in arthritis.

It was recently shown that IL-17-producing  $\gamma/\delta$  T cells were observed in the joints of mice with CIA and that blocking a certain subset of IL-17-producing  $\gamma/\delta$  T cells suppressed CIA (29). However,  $\gamma/\delta$  T cells have been shown to be dispensable for the development of CIA (38). In addition, anti-CII-specific antibody levels were comparable between  $\gamma/\delta$  T cell-deficient and wild-type mice. These findings suggest that MAIT cells and  $\gamma/\delta$  T cells share similar roles in arthritis and that both are involved mainly in the effector phase of arthritis. It is known that  $\gamma/\delta$  T cells as well as iNKT cells are increased during CIA. Because MAIT cells share similar features with  $\gamma/\delta$  T cells and iNKT cells, MAIT cells may also be increased during CIA.

We observed a significant decrease in IL-17 production by LMNCs upon stimulation with anti-CD3 mAb in MR1 $^{-/-}$  mice compared to WT control mice. As sorted MAIT cells produced high amounts of IL-17 after anti-CD3 mAb stimulation, the major source of IL-17 responsible for the difference between MR1 $^{-/-}$  and WT mice seems to be MAIT cells. Th17 cells and iNKT cells have been shown to produce IL-21, which enhanced IL-17 production or induced proliferation of IL-17-producing cells (39). It is not known whether MAIT cells produce IL-21, but MAIT cells might augment IL-17 production by other LMNCs, including  $\gamma/\delta$  T cells, through such mechanisms. Further studies to determine whether MAIT cells regulate  $\gamma/\delta$  T cells under both physiologic and pathologic conditions, including in the presence of arthritis, will be of interest.

The frequency of murine  $\gamma/\delta$  T cells is 1–5% in blood lymphocytes and 25–60% in gut lymphocytes. Human  $\gamma/\delta$  T cells also comprise up to 2–3% of peripheral T cells (9,10). Although the precise frequency of murine MAIT cells is not known, it has been speculated that MAIT cells may comprise up to 10% of double-negative T cells in the gut lamina propria and <2% of double-negative T cells in the mesenteric lymph nodes, indicating that the frequency of murine MAIT cells is much lower than that of mouse  $\gamma/\delta$  T cells (15). It has been suggested that  $\gamma/\delta$  T cells are the predominant source of IL-17 in the joints of CIA mice, but IL-17-producing  $\gamma/\delta$  T cells could not be detected in RA synovial tissue (31). Recently, Martin et al (23) revealed that human MAIT cells can be identified as V $\alpha$ 7.2+ CD161<sup>high</sup> T cells, which are abundant in blood. In addition, human MAIT cells produce IL-17 and express



tissue-homing chemokine receptors (23). An IL-17-producing CD161<sup>high</sup> T cell population has been described in human arthritic joints (40). Thus, it is possible that MAIT cells rather than  $\gamma/\delta$  T cells play a major role in the pathogenesis of human RA.

CD4<sup>+</sup> Th17 cells require IL-6/STAT-3 activation for the expression of ROR $\gamma$ t, which is a crucial transcription factor for IL-17 production (41). However, some innate-like lymphocyte subsets, such as iNKT cells,  $\gamma/\delta$  T cells, and lymphoid tissue-inducer (LTi)-like cells, are known to constitutively express ROR $\gamma$ t, IL-1 receptor type I, and IL-23R (42). In addition, these IL-17-producing innate-like lymphocytes, including LTi cells,  $\gamma/\delta$  T cells, and iNKT cells, secrete IL-17 when stimulated by IL-23 with or without IL-1 $\beta$ . In this study, we demonstrated cytokine-mediated activation of MAIT cells. MAIT cells produced IL-17 in response to IL-23. Moreover, IL-1 $\beta$  induced proliferation of MAIT cells. Thus, it is possible that MAIT cells may contribute to the disease progression of arthritis through another mechanism in addition to IL-17 production. In adoptive transfer experiments, MAIT cells augmented the disease severity of CIA in MR1-deficient mice. Thus, this result also indicates that MAIT cell-mediated exacerbation of arthritis may be induced by cytokines, without a requirement for TCR stimulation.

In EAE, disease suppression by MAIT cells was accompanied by a reduction in the production of cytokines, including IFN $\gamma$  and IL-17, by T cells and increased IL-10 production by B cells. Encephalitogenic T cells play a major role in EAE (43,44). EAE can be induced in naive mice by transferring myelin-reactive T cells. T cell-targeted therapies, including anti-very late activation antigen 4 treatment, have been shown to suppress EAE. Although CIA was reduced in MR1<sup>-/-</sup> DBA/1J mice, we observed a significant decrease in CII-specific IgG1 antibody levels in these mice as compared with their WT controls in some experiments (data not shown), suggesting the inhibition of Th1 responses by MAIT cells. Therefore, it is still possible that MAIT cells suppress Th1 response during the early induction phase of CIA. MAIT cells may be functionally plastic, and thus exert different functions depending on the pathologic condition. Arthritis involves massive cytokine production due to various types of immune cell activation. Since MAIT cells can be activated by inflammatory cytokines, MAIT cells may contribute to augment the immune response once overt inflammation occurs.

In summary, we have shown that MAIT cells contribute to the progression of arthritis by enhancing the inflammation in CIA and CAIA models. In addition, we demonstrated that MAIT cells could be activated by

cytokine stimulation even without TCR stimulation. We and others previously reported that, although iNKT cells play pathogenic roles in arthritis models, modulation of iNKT cell function by ligands successfully suppressed arthritis (45–47). The proportion of human MAIT cells appears to be much higher than that of mouse MAIT cells. Therefore, MAIT cells may play an important pathogenic role in human arthritis and MAIT cell-targeted therapy may hold promise as a new therapeutic intervention for arthritis, including RA.

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#### AUTHOR CONTRIBUTIONS

All authors were involved in drafting the article or revising it critically for important intellectual content, and all authors approved the final version to be published. Dr. Miyake had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

**Study conception and design.** Chiba, Tajima, Miyazaki, Miyake.

**Acquisition of data.** Chiba, Tajima, Tomi, Miyazaki.

**Analysis and interpretation of data.** Chiba, Tajima, Tomi, Miyazaki, Yamamura, Miyake.

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# Interleukin 6 signaling promotes anti-aquaporin 4 autoantibody production from plasmablasts in neuromyelitis optica

Norio Chihara<sup>a,b</sup>, Toshimasa Aranami<sup>a,c</sup>, Wakiro Sato<sup>a</sup>, Yusei Miyazaki<sup>a</sup>, Sachiko Miyake<sup>a,c</sup>, Tomoko Okamoto<sup>c,d</sup>, Masafumi Ogawa<sup>c,d</sup>, Tatsushi Toda<sup>b</sup>, and Takashi Yamamura<sup>a,c,1</sup>

<sup>a</sup>Department of Immunology, National Institute of Neuroscience, National Center of Neurology and Psychiatry (NCNP), Tokyo 187-8502, Japan; <sup>b</sup>Department of Neurology, Kobe University Graduate School of Medicine, Kobe 650-0017, Japan; and <sup>c</sup>Multiple Sclerosis Center and <sup>d</sup>Department of Neurology, National Center Hospital, NCNP, Tokyo 187-8551, Japan

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Neuromyelitis optica (NMO) is an inflammatory disease affecting the optic nerve and spinal cord, in which autoantibodies against aquaporin 4 (AQP4) water channel protein probably play a pathogenic role. Here we show that a B-cell subpopulation, exhibiting the CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> phenotype, is selectively increased in the peripheral blood of NMO patients and that anti-AQP4 antibodies (AQP4-Abs) are mainly produced by these cells in the blood of these patients. These B cells showed the morphological as well as the phenotypical characteristics of plasmablasts (PB) and were further expanded during NMO relapse. We also demonstrate that interleukin 6 (IL-6), shown to be increased in NMO, enhanced the survival of PB as well as their AQP4-Ab secretion, whereas the blockade of IL-6 receptor (IL-6R) signaling by anti-IL-6R antibody reduced the survival of PB in vitro. These results indicate that the IL-6-dependent B-cell subpopulation is involved in the pathogenesis of NMO, thereby providing a therapeutic strategy for targeting IL-6R signaling.

neuroinflammatory disease | autoimmunity | multiple sclerosis | central nervous system | IL-6 receptor blockade

Neuromyelitis optica (NMO) is an inflammatory demyelinating disorder characterized by recurrent attacks of severe optic neuritis and myelitis. Unlike the conventional form of multiple sclerosis (MS), the lesions of NMO tend to spare the cerebral white matter, especially during the early stage (1), and even a single episode of attack can cause serious neurological deficits such as total blindness and paraplegia. Accordingly, accumulation of irreversible damage to the central nervous system (CNS) along with rapid progression of disability is more frequently found in NMO compared with MS (2).

NMO can be distinguished from MS by clinical, neuroimaging, and serological criteria (3). It is now known that serum anti-aquaporin 4 (AQP4) autoantibodies can be used as a disease marker of NMO (1, 2). AQP4 is the most abundantly expressed water channel protein in the CNS and is highly expressed in the perimicrovessel astrocyte foot processes, glia limitans, and ependyma (4). Emerging clinical and pathological observations suggest that anti-AQP4 antibodies (AQP4-Abs) play a key role in the pathogenesis of NMO. Prior studies have documented a significant correlation of serum AQP4-Ab levels with the therapeutic efficacy of plasma exchange during clinical exacerbations of NMO (2, 5). In the CNS lesions of NMO, reduced expression of AQP4 on astrocytes is evident even during the early stage (6), which is followed by the occurrence of vasocentric destruction of astrocytes associated with perivascular deposition of complement and IgG (7).

On the other hand, recent studies have suggested that AQP4-Abs alone are incapable of causing the clinical and pathological features of NMO. In fact, Hinson et al. emphasized the role of cellular immunity in combination with AQP4-Abs by showing

that the attack severity of NMO was not correlated with serum AQP4-Ab levels (8). It was also demonstrated that direct injection of IgGs derived from NMO patients into the brains of naïve mice did not cause NMO-like lesions, although brain tissue destruction associated with leukocyte infiltration was elicited by coinjecting human complement (9). Other groups have shown that the passive transfer of IgGs from NMO patients to rats challenged with induction of experimental autoimmune encephalomyelitis (EAE) may cause a decrease in the expression of AQP4 in astrocytes along with worsening of clinical EAE (10–12). In contrast, the transfer of IgGs to unimmunized rats did not cause any pathology. These results suggest that induction of AQP4-Ab-mediated pathology in NMO depends on the presence of complement, leukocytes, and T cells.

Although AQP4-Ab-secreting cells are a potential target for therapy, detailed characteristics of AQP4-Ab-producing cells have not been clarified yet. Because some NMO patients have elevated serum anti-nuclear and anti-SS-A/SS-B Abs (1), as found in patients with systemic lupus erythematosus (SLE) or Sjögren syndrome, NMO might share common pathological mechanisms with these autoimmune diseases. Kikuchi et al. previously reported that CD180<sup>-</sup> B cells are activated B cells capable of producing autoantibodies in SLE (13). CD180 is a member of the leucine-rich repeat family of molecules with homology to Toll-like receptor 4 (14), which is highly expressed by naïve and memory B cells but not by plasma cells (15). Odendahl et al. demonstrated that CD27<sup>high</sup>CD38<sup>+</sup> B cells, capable of producing high-affinity IgG (16), are increased in the peripheral blood of SLE patients with some correlation to disease activity (17). Considering the phenotypes of autoantibody-producing cells reported in SLE, we analyzed the expression of CD27, CD38, and CD180 on CD19<sup>+</sup> B cells in the peripheral blood of NMO patients. We found that CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells were significantly increased in AQP4-Ab seropositive patients diagnosed with NMO or NMO spectrum disorder (1) compared with healthy subjects (HS) or MS patients. Notably, this B-cell subpopulation was found to be a major source of AQP4-Abs in the peripheral blood of AQP4-Ab seropositive patients and depended on interleukin-6 receptor (IL-6R) signaling for survival.

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<sup>1</sup>To whom correspondence should be addressed. E-mail: yamamura@ncnp.go.jp.

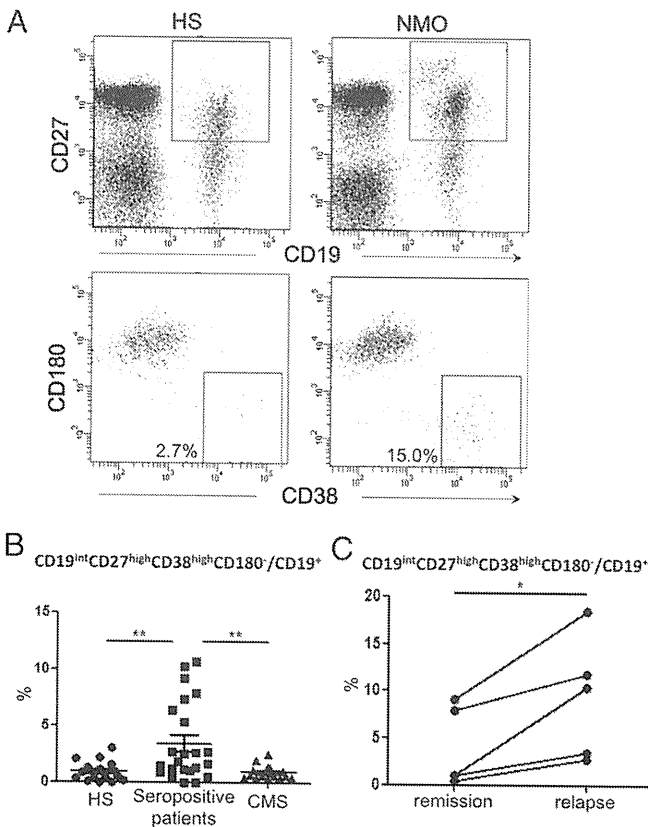
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## Results

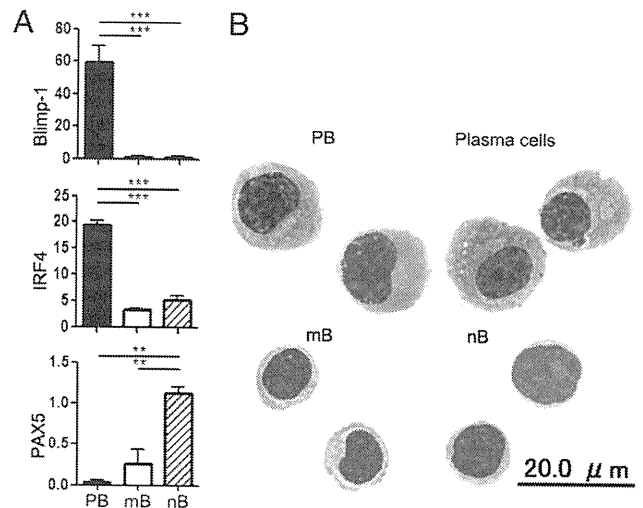
**CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B Cells Were Increased in the Peripheral Blood of NMO Patients.** Although AQP4-Abs are identified as IgGs (18), no prior study has focused on proportional changes of B-cell subsets in NMO. We therefore performed multicolor flow cytometric analysis of peripheral blood mononuclear cells (PBMC) derived from patients and controls. After starting the study, we soon noticed a remarkable expansion of a distinct B-cell subset in some patients with NMO. The expanded B cells were identified as a population of CD27<sup>high</sup>, CD38<sup>high</sup>, and CD180<sup>-</sup>, and showed lower expression of CD19 than other B cells (Fig. 1A). Notably, this population did not express the B-cell marker CD20 (Fig. S1). First, we collected samples from patients in remission and analyzed the pooled data. We found that the proportion of this subpopulation among CD19<sup>+</sup> B cells was significantly increased in AQP4-Ab seropositive patients with NMO or NMO spectrum disorder (Fig. 1B) compared with HS or CMS patients. There was no significant difference in the proportion of this B-cell subpopulation between those with typical NMO and those with NMO spectrum disorder. Furthermore, the frequency of this B-cell subpopulation was correlated with the serum AQP4-Ab titer (Fig. S2). Comparison of paired samples obtained from the same patients during relapse and in

remission showed that the CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells further increased during relapse (Fig. 1C). In contrast, the frequencies of CD27<sup>-</sup> naive B cells (nB) and CD27<sup>+</sup>CD38<sup>-low</sup> memory B cells (mB) were not altered in AQP4-Ab seropositive patients compared with controls (Fig. S3). The large majority of seropositive patients were treated with corticosteroids. However, the frequency of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells among CD19<sup>+</sup> B cells was not correlated with the daily corticosteroid dose given to patients (Fig. S4). Moreover, the increase in cells in NMO patients was still evident compared with that in CMS patients similarly treated with corticosteroids (Fig. S5). Taken together, the selective increase in CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells in seropositive patients was thought to reflect their role in the pathogenesis of NMO but not to be an effect of the corticosteroid treatment.

**Expanded Cells Resemble Early Plasma Cells in Gene Expression and Morphology.** To gain insights into the developmental stage of the CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells, we quantified the mRNA expression of B-cell-associated transcription factors in sorted cell populations. Compared with nB and mB, this population showed much higher expression of B-lymphocyte-induced maturation protein 1 (Blimp-1) and IFN regulatory factor 4 (IRF4), which are essential for the regulation of plasma cell differentiation (19, 20) (Fig. 2A). In contrast, the expression of paired box gene 5 (PAX5), known to be down-regulated in early plasma cell differentiation (21), was reciprocally reduced in the B-cell subset. This gene expression pattern is very similar to that of plasma cells. However, it was notable that the cells of interest expressed CD19, which is not detected in mature plasma cells. Moreover, only 40% of this population expressed the most reliable plasma cell marker CD138 (22). Morphological analysis also confirmed the similarity of this population to plasma cells: they exhibit eccentric nucleus, perinuclear hof region, and abundant cytoplasm. However, they possess a larger nucleus with a lower extent of chromatin clumping compared with CD138<sup>+</sup> plasma cells derived from HS (Fig. 2B). Notably, the CD138<sup>+</sup> population among CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells in NMO patients was



**Fig. 1.** CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> B cells increased in NMO patients. (A) A flow cytometric scheme for the analysis of B-cell subpopulations. PBMC from HS and NMO in remission were stained with fluorescence-conjugated anti-CD19, -CD27, -CD38, and -CD180 mAbs. CD19<sup>+</sup>CD27<sup>+</sup> cells were partitioned (Upper) and analyzed for expression of CD38 and CD180 (Lower). Values represent the percentage of CD38<sup>high</sup>CD180<sup>-</sup> cells within CD19<sup>+</sup>CD27<sup>+</sup> cells. (B) Analysis of the pooled data derived from patients in clinical remission. This shows the percentages of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells within CD19<sup>+</sup> cells from HS, seropositive patients, and CMS patients (\*\**P* < 0.01; Tukey's post hoc test). (C) Comparison of remission and relapse of NMO. Data obtained from the same patients are connected with lines (\**P* < 0.05; Wilcoxon signed rank test).



**Fig. 2.** Resemblance of CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells to plasma cells. (A) mRNA expression of Blimp-1, IRF4, and PAX5. B-cell subpopulations [CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> (PB), CD27<sup>-</sup> naive (nB), CD27<sup>+</sup>CD38<sup>-low</sup> memory (mB)] were sorted by FACS and total RNA was extracted for qRT-PCR analysis. RNA levels were normalized to ACTB for each sample (\*\**P* < 0.01; \*\*\**P* < 0.001; Tukey's post hoc test). (B) May-Grünwald-Giemsa staining of B-cell subpopulations. PB (Upper Left), mB (Lower Left), and nB (Lower Right) from NMO are presented along with morphologically identified plasma cells (CD19<sup>int</sup>CD27<sup>high</sup>CD38<sup>high</sup>CD138<sup>+</sup>) from HS (Upper Right).

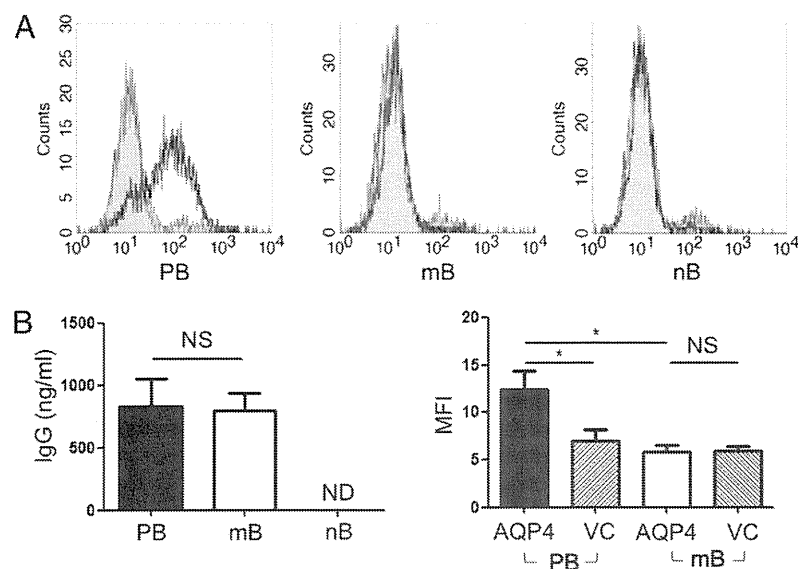
morphologically indistinguishable from the CD138<sup>-</sup> population in NMO patients or HS, indicating the immature characteristic of CD27<sup>high</sup>CD38<sup>high</sup>CD180<sup>-</sup> cells (Fig. S6). These phenotypical and morphological characteristics as well as the results of the quantitative real-time PCR (qRT-PCR) analysis indicate that this B-cell population is equivalent to plasmablasts (PB) (22–26). Hereafter, we use the term “PB” to distinguish this population from other B cells.

**Expression of B-Cell Cytokine Receptors on PB.** Prior studies have identified cytokines that are critically involved in the differentiation and/or survival of plasma cells, including IL-6 and B-cell-activating factor (BAFF). IL-6 induces B-cell differentiation into plasma cells, maintains early plasma cell survival, and enhances plasma cell IgG secretion (24). Besides, IL-6 is elevated in the cerebrospinal fluid (CSF) or peripheral blood of NMO patients compared with that of CMS patients and HS (27, 28). In a rodent autoimmunity model, IL-6 deficiency caused impaired autoantibody secretion by B cells (29). Given the potential role of IL-6 in NMO, we performed flow cytometry analysis for the expression of IL-6R. Results showed remarkable expression of IL-6R on PB, although it was only marginal or absent on mB and nB (Fig. S7). Because BAFF and A proliferation-inducing ligand (APRIL) can also promote the survival of PB (25, 26), we next evaluated the expression of the receptors for BAFF and APRIL, BAFF receptor (BAFF-R), B-cell maturation antigen (BCMA), and transmembrane activator and calcium modulator and cyclophilin ligand interactor (TACI). Expression of BCMA and TACI was selectively up-regulated in PB in parallel with IL-6R. In contrast, BAFF-R was up-regulated in mB and nB, but not in PB (Fig. S7).

**PB Is a Selective Source of AQP4-Abs in Peripheral Blood.** We were interested to know whether PB were capable of producing AQP4-Abs upon stimulation with cytokines and, therefore, examined the

ability of IL-6, BAFF, and APRIL to enhance AQP4-Ab secretion by PB. We cultured the isolated PB for 6 d in the presence or absence of each cytokine, and evaluated the presence of AQP4-Abs in the supernatants by measuring IgG binding to Chinese hamster ovary (CHO) cells transfected with the human AQP4 vector (CHO<sup>AQP4</sup>) or the vector control (CHO<sup>VC</sup>). We found that IL-6, but not BAFF or APRIL, could significantly enhance AQP4-Ab secretion from PB (Fig. S8), as assessed by specific IgG binding to CHO<sup>AQP4</sup>. Further study focusing on IL-6 showed that exogenous IL-6 promoted the production of AQP4-specific IgGs from PB (Fig. 3A), but not from the other B-cell subpopulations. Similar results were obtained from six independent experiments (Fig. S9), indicating that PB could be major AQP4-Ab producers in PBMC. In the absence of addition of IL-6, supernatants from PB did not show any significant reactivity to CHO<sup>AQP4</sup>. To further analyze the AQP4-Ab-secreting potential of each B-cell subpopulation, we next stimulated the cells with a combination of IL-6, IL-21, and anti-CD40 that efficiently induces B-cell differentiation and IgG production (30). This polyclonal stimulation induced the secretion of similar amounts of IgGs from mB and PB. However, only the supernatant of PB specifically reacted to CHO<sup>AQP4</sup> cell transfectants, indicating that AQP4-Ab-producing B cells were highly enriched in PB (Fig. 3B).

**Survival and Functions of PB Depend on IL-6 Signaling.** We evaluated the influence of IL-6, BAFF, and APRIL on the survival of PB after 2 d of in vitro culture (Fig. 4A). Among the added cytokines, only IL-6 was found to significantly promote the survival of PB (Fig. 4B). We also assessed the expression levels of X-box-binding protein 1 (XBP-1) in PB by qRT-PCR after 24 h of culture with or without IL-6. XBP-1 is a transcription factor critical for IgG secretion (31), and the splicing process of XBP-1 mRNA yields a more active and stable protein. We found that the expression of both unspliced [XBP-1(u)] and spliced [XBP-1(s)] forms of XBP-1 mRNA was augmented in PB by the ad-



**Fig. 3.** Production of AQP4-Abs by PB. (A) Using flow cytometry, we examined whether AQP4-Abs could be produced by PB, mB, or nB cells. FACS-sorted cells were cultured with IL-6 (1 ng/mL) for 6 d and supernatants were collected. Supernatant IgGs reactive to CHO<sup>AQP4</sup> (open histogram) and CHO<sup>VC</sup> cells (closed histogram) were detected by anti-human IgG secondary antibody. The supernatant from PB (Left), but not from mB or nB, contains IgGs reactive to CHO<sup>AQP4</sup>, indicating that only PB secrete AQP4-Abs after stimulation with IL-6. (B) Memory B cells (mB) produce IgGs but not AQP4-Abs. B-cell subpopulations were cultured in the presence of IL-6 (1 ng/mL), IL-21 (50 ng/mL), and anti-CD40 mAb (1 μg/mL) for 6 d. IgGs in the culture supernatants were measured by sandwich ELISA (Left) (each assay was performed in quadruplicate). Data from three patients are expressed as mean ± SD. The activity of AQP4-Abs in the culture supernatants from PB and mB was also measured by flow cytometry (Right). Aliquots of CHO<sup>AQP4</sup> cells (AQP4) and CHO<sup>VC</sup> cells (VC) ( $n = 4$  for each) were stained with the supernatant of PB or mB from every patient. Data are expressed as median fluorescence intensity values from the results of three patients (\* $P < 0.05$ ; Tukey's post hoc test). ND, not detected; NS, not significant.