Melanoma Cell Adhesion Molecule Expressing Helper T Cells in CNS Inflammatory Demyelinating Diseases

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Abstract

Background and Objective
To elucidate the relationship between melanoma cell adhesion molecule (MCAM)-expressing lymphocytes and pathogenesis of CNS inflammatory demyelinating diseases (IDDs).

Methods
Patients with multiple sclerosis (MS) (n = 72) and neuromyelitis optica spectrum disorder (NMOSD, n = 29) were included. We analyzed the frequency and absolute numbers of MCAM⁺ lymphocytes (memory helper T [mTh] cells, naive helper T cells, CD8⁺ T cells, and B cells) in the peripheral blood (PB) and the CSF of patients with MS and NMOSD, treated with/without disease-modifying drugs (DMDs) or steroids, using flow cytometry.

Results
The frequency of MCAM⁺ cells was higher in the mTh cell subset than that in other lymphocyte subsets. A significant increase in the frequency and the absolute number of MCAM⁺ mTh cells was observed in the PB of patients with NMOSD, whereas no increase was observed in the PB of patients with MS. The frequency of CSF MCAM⁺ mTh cells was higher in relapsing patients with MS and NMOSD than that in the control group. Although there was no difference in the frequencies of MCAM⁺ lymphocytes among the DMD-treated groups, fingolimod decreased the absolute number of MCAM⁺ lymphocytes.

Discussion
MCAM⁺ mTh cells were elevated in the CSF of relapsing patients with MS and in both the PB and CSF of patients with NMOSD. These results indicate that MCAM contributes to the pathogenesis of MS and NMOSD through different mechanisms. MCAM could be a therapeutic target of CNS IDDs, and further study is needed to elucidate the underlying mechanism of MCAM in CNS IDD pathogenesis.
Multiple sclerosis (MS) is the most common CNS inflammatory demyelinating disease (IDD). Clinical and experimental evidence suggests an autoimmune-mediated etiology for MS. Previous studies have suggested that T helper 1 (Th1) and T helper 17 (Th17) cells are pathogenic in MS, and other inflammatory diseases. In addition, the frequency of MCAM⁺ Th cells increases in the peripheral blood (PB) and the CSF and accumulates in CNS lesions of patients with MS. An MCAM blockade ameliorates EAE severity. Therefore, the role of MCAM in CNS IDD pathogenesis is increasingly attracting attention. However, studies on MCAM in MS or neuromyelitis optica spectrum disorder (NMOSD) are limited.

In this study, we evaluated the frequency and the absolute number of MCAM⁺ lymphocytes in the PB and the CSF of CNS IDDs to elucidate the relationship between MCAM-expressing lymphocytes and the pathogenesis of CNS IDDs.

**Methods**

**Participants**
In total, 72 patients with MS and 29 with NMOSD were included in this study. In addition, 23 patients with myasthenia gravis (MG), 34 with noninflammatory neurologic disorder (NIND), and 25 healthy subjects (HCs) were included as the controls. NIND included stroke, chronic headache, neurodegenerative disorders, and epilepsy. The patient profile among patients with MS, NMOSD, MG, NIND, and HCs was similar, as shown in Table 1. The mean ages of patients with MS, NMOSD, MG, NIND, and HCs were 42.0 (range 19–74), 51.5 (27–77), 52.8 (20–74), 49.0 (18–82), and 43.1 (25–80) years, respectively, and the male-to-female ratios were 26/46, 5/24, 9/14, 11/23, and 11/14, respectively. The relapse phase (active wrinkle) to remission ratio in MS, NMOSD, and MG were 30/36 (the other 6 patients presented with secondary progressive MS), 14/15, and 15/8, respectively, and the disease duration was 12.5 ± 8.4, 10.4 ± 10.4, and 8.9 ± 12.5 (mean ± SD) years, respectively. We also compared MCAM expression in each lymphocyte subset among the untreated patients with MS (n = 25) and those treated with different DMDs (interferon [IFN]–β [n = 12], dimethyl fumarate [DMF] [n = 11], fingolimod [FTY] [n = 18], and NTZ [n = 6]). At the time of examination, 25 of 29 patients with NMOSD and 19 of 23 patients with MG were being treated with oral prednisolone (PSL; mean dose, 12.3 mg/day and 10.6 mg/day, respectively). Patients with NIND did not receive any immunotherapy. None of these patients had received pulse steroid therapy, IV immunoglobulin, or plasma exchange >1 month before blood sampling. We analyzed the MCAM expression >3 months after initiation of any DMDs. CSF and PB paired samples were obtained from 10 patients with MS (8 relapse and 2 remission), 6 with NMOSD (4 relapse and 2 remission), and 15 with NIND. The diagnoses of MS and NMOSD were established according to the 2010 McDonald criteria and the international consensus diagnostic criteria for NMOSD, respectively. All patients were recruited from Tokyo Women’s Medical University Hospital, Japan.

**Standard Protocol Approvals, Registrations, and Patient Consents**
This study was approved by the ethics committee of Tokyo Women’s Medical University School of Medicine. Written informed consent was obtained from each participant. This is a cross-sectional study conducted between February 2016 and September 2020.

**Cell Preparation and Flow Cytometry**
PB mononuclear cells (PBMCs) were freshly isolated through density-gradient centrifugation using Ficoll-Paque Plus (GE Healthcare, Oakville, ON, Canada) and immediately stored at −80°C until the time of analysis. They were stained against cell surface antigens, according to the manufacturer’s instructions. Fluorescent-conjugated monoclonal antibodies (mAbs) and isotype controls used in this study were as follows: CD3-PerCp-Cy5.5, CD4-PE-Cy7, CD45RA-PE-Cy7, CD19-APC-Cy7, MCAM (CD146)-APC (clone P1H12), α4-integrin (CD49d)- PE, and mouse IgG1-APC (BioLegend, San Diego, CA); CD8-fluorescein isothiocyanate (Beckman Coulter, Brea, CA); and IgG1x-PE (BD Pharmingen, San Jose, CA). Each lymphocyte subset was defined like in Figure 1 and Table 2. CSF cells were stained directly with the above-mentioned Abs. After incubation, cells were washed and resuspended in phosphate-buffered saline.
Cells were then measured using a Gallios Flow Cytometer (Beckman Coulter) and analyzed using Kaluza Analysis Software (Beckman Coulter). The absolute number of each lymphocyte subset was obtained by the dual-platform method.

### Statistical Analysis

Statistical analyses were performed with GraphPad Prism 7 (GraphPad Software, San Diego, CA). An unpaired *t* test or Mann-Whitney *U* test was used, as appropriate, for statistical analyses of data between 2 groups. One-way analysis of variance (ANOVA) or the Kruskal-Wallis test was used, as appropriate, to compare data from more than 2 groups. If the one-way ANOVA or Kruskal-Wallis test results were significant, post hoc analyses were performed using the Tukey test or the Dunn multiple comparison test, respectively. Correlations between 2 variables were examined using linear regression analyses. Statistical significance was indicated at *p* value <0.05.

### Data Availability

Anonymized data are available from the corresponding author on reasonable request from any qualified investigator for purposes of replicating the procedures and results.

### Results

#### The Frequency and Absolute Number of MCAM+ Lymphocytes in Patients With NMOSD Increased

First, we analyzed the frequency of MCAM+ cells in each lymphocyte subset (mTh cells, naive Th [nTh] cells, CD8+ T cells, and B cells) in the PB (Figure 1, A and B) of all 183 patients and HCs. Consistent with the results of previous studies, the frequency of MCAM+ lymphocytes was higher in mTh cells than that in other lymphocyte subsets,10,12 and most MCAM+ cells coexpressed CD49d.17 By contrast, only a few CD49d+ cells (1%–5%) coexpressed MCAM.

Next, we compared the frequency and the absolute number of MCAM+ cells in each lymphocyte subset in the PB among the disease groups. FTY- and DMF-treated patients with MS were excluded from the absolute number analysis because treatment with these DMDs modified the number of lymphocytes, as described in eTables 1 and 2, links.lww.com/NXI/A559. In addition, the absolute number of MCAM+ nTh cells was not calculated because the absolute number of cells in this subset, compared with that in other subsets, was extremely low. Thus, the frequency and the absolute number of MCAM+ mTh lymphocytes were significantly higher in patients with NMOSD than those in other patients, including patients with MS and NIND (Figure 1, C and D). No difference was observed in either the frequency or the absolute number of cells in each lymphocyte subset between patients with MS and those with MG or NIND. In addition, no correlation between the frequency or the absolute number of MCAM+ cells in each lymphocyte subset and activity or severity of CNS IDDs was observed. Moreover, there was no difference in the frequency and the absolute number of CD49d+ lymphocytes among the disease groups (data not shown).
Effect of DMDs or Steroids on the Frequency and Absolute Number of MCAM⁺ Lymphocytes in Patients With CNS IDD

To clarify the effect of DMDs or steroids on MCAM⁺ lymphocytes, we examined the frequency and the absolute number of MCAM⁺ lymphocytes between untreated patients with CNS IDD and patients with CNS IDD treated with DMDs or steroids. In FTY-treated patients with MS, the frequencies and absolute numbers of some lymphocyte subsets decreased, as shown in eTables 1 and 2, links.lww.com/NXI/A559. Although there was no difference in the frequency of MCAM⁺ lymphocytes among the patients treated with different DMDs (Figure 2A), the absolute number of MCAM⁺ mTh cells, CD8⁺ T cells, and B cells was significantly decreased in FTY-treated patients compared with that in other DMD-treated and untreated patients with MS (Figure 2B).
Although there was no difference in the PSL dosage between patients with NMOSD and MG (mean dosage, 12.3 and 10.6 mg/day, respectively), a significant difference was observed in the frequency of MCAM⁺ mTh cells between PSL-treated patients with NMOSD and MG (p < 0.05, Figure 2C). This result suggested that the increase in the frequency of MCAM⁺ mTh cells in NMOSD was independent of the effects of the steroid.

MCAM⁺ mTh Cells Were Enriched in the CSF of Patients With CNS IDD

Finally, we examined the frequency and the absolute number of MCAM⁺ lymphocytes in the CSF of patients with CNS IDD and NIND (Figure 3, A and B). The frequency of CSF MCAM⁺ mTh cells was significantly higher in patients with MS and NMOSD than that in those with NIND. As shown in Figure 3A, the difference in the frequency of MCAM⁺ mTh cells between the PB and CSF of patients with MS was significant, whereas that between the PB and CSF of patients with NMOSD was not significant. Furthermore, the absolute number of CSF MCAM⁺ mTh cells and CD8⁺ T cells was significantly higher in patients with NMOSD than that in patients with NIND (Figure 3B). We speculated that the frequency of MCAM⁺ mTh cells in the PB reflected that of CSF MCAM⁺ mTh cells. Therefore, we examined the correlation between the frequencies of MCAM⁺ mTh cells in the PB and CSF. Although analysis of patients with NMOSD and NIND revealed a strong correlation (r² = 0.91, p < 0.001; Figure 3C, left), the MS-independent analysis showed no significant correlation between these frequencies (r² = 0.22, p > 0.05; Figure 3C, right). Similar to the PB, most of the MCAM⁺ mTh cells in the CSF coexpressed CD49d (data not shown). These data suggested that MCAM⁺ mTh cells might contribute to the pathogenesis of both MS and NMOSD through different mechanisms.

### Table 2 Definition of Each Lymphocyte Subset

| Lymphocyte subset    | Marker                |
|----------------------|-----------------------|
| Memory helper T cells| CD3⁺ CD4⁺ CD45RA⁻     |
| Naive helper T cells | CD3⁺ CD4⁺ CD45RA⁺     |
| CD8⁺ cytotoxic T cells| CD3⁺ CD8⁺            |
| B cells              | CD3⁻ CD19⁺            |

**Figure 2** Frequency and Absolute Number of MCAM⁺ Lymphocytes in DMD- or Steroid-Treated Patients With CNS IDD

(A-a–C) Frequency of MCAM⁺ cells in each lymphocyte subset among the DMD- or steroid-treated patients with MS (IFN-β [n = 12], DMF [n = 11], FTY [n = 18], and NTZ [n = 6]) and NIND (n = 25). (A-a–B-c) Absolute number of MCAM⁺ cells in each lymphocyte subset among the DMD- or steroid-treated patients with MS (IFN-β [n = 9], DMF [n = 10], FTY [n = 18], and NTZ [n = 6]) and untreated patients with MS (n = 29). (B-a) Frequency of MCAM⁺ mTh cells in the PB of patients with NMOSD and MG treated with or without PSL. At the time of examination, 25 of the 29 patients with NMOSD and 19 of the 23 patients with MG were undergoing treatment with oral PSL. Dots represent individual samples. The lower, upper, and middle lines correspond to the 25th centile, 75th centile, and median, respectively. *p < 0.05, **p < 0.01, and ***p < 0.001. DMD = disease-modifying drug; DMF = dimethyl fumarate; FTY = fingolimod; IDD = inflammatory demyelinating disease; IFN = interferon; MCAM = melanoma cell adhesion molecule; MG = myasthenia gravis; mTh cell = memory helper T cell; NMOSD = neuromyelitis optica spectrum disorder; ns = not significant; NTZ = natalizumab; PB = peripheral blood; PSL = prednisolone.
Discussion

This study was designed to clarify the differences in MCAM expression using lymphocytes from patients with MS and NMOSD. The results showed that in patients with relapsing MS, the frequency of MCAM+ mTh cells increased only in the CSF, whereas in those with NMOSD, the frequency of MCAM+ mTh cells increased in both the PB and CSF.

Adhesion molecules play key roles in MS pathogenesis. α4-Integrin is the most important adhesion molecule that is involved in the migration of inflammatory lymphocytes to the CNS. NTZ is a widely used mAb against α4-integrin and one...
of the most effective DMDs used for MS treatment. Although there have been many studies on adhesion molecules in MS, NTZ is the only approved and effective drug that targets adhesion molecules.

MCAM is one of the adhesion molecules that is involved in the migration of lymphocytes to the CNS.12-14,18,20 MCAM is found on endothelial cells and 2%–3% of T cells. Laminin-411 is the ligand of MCAM.13,14 MCAM+ T cells have a proinflammatory phenotype; they secrete IL-17, IFN-γ, granulocyte macrophage–colony-stimulating factor, and granzyme B.10,11,18 The frequency of PB MCAM+ Th cells is increased in rheumatoid arthritis (RA), Behcet disease, and inflammatory bowel diseases, and MCAM expression is correlated with the activity and/or severity of these diseases.8-11 Moreover, in RA, MCAM+ Th cells accumulate in the synovial fluid, suggesting that MCAM+ Th cells can migrate to inflammatory sites from the blood owing to their ability of extravasation through cell-cell adhesion.9,23

Previous studies have revealed that MCAM is involved in EAE and MS pathogenesis.12-14,17,20 MCAM+ Th cells have the characteristics of pathogenic Th17 cells, such as the ability to produce both IFN-γ and IL-17, and a strong migration capacity.11,12,14,19 MCAM+ Th cells accumulate in the CSF and CNS lesions of patients with MS.12,17,19 Similarly, MCAM+ CD8+ T cells also accumulate in the CSF and CNS lesions of patients with relapsing MS.18 Moreover, an MCAM blockade restricted the migration of Th cells and CD8+ T cells from the blood to the CNS in the blood-brain barrier (BBB) model in vitro.12,14,18 Even in EAE, an MCAM blockade or depletion reduced the neurologic deficit.12,14,18 A previous study reported that the frequency of CSF CD49d+ Th cells decreased, whereas the frequency of MCAM+ mTh cells increased in patients with MS on long-term NTZ treatment compared with that in those without NTZ treatment, which suggests that MCAM+ mTh cell adhesion is not affected by NTZ alone.19 According to the results of previous studies, MCAM is involved in the pathogenesis of MS and EAE; however, studies on the role of MCAM in the pathogenesis of CNS IDDs are limited.

Contrary to our speculation and the results of previous studies,12,17 the frequency of MCAM+ mTh cells and CD8+ T cells was not increased in the PB of patients with MS. With regard to the CSF analysis, the frequency of MCAM+ mTh cells in the CSF was higher in the relapse phase (n = 8, mean frequency of 10.5%) than that in the remission phase (n = 2, mean frequency of 3.6%, Figure 3A). Although the number of patients was small, this finding suggests that the frequency of MCAM+ mTh cells might be a biomarker of disease activity in patients with MS. Of interest, several studies have shown that the frequency of Th17 cells is increased in the CSF of patients with active MS.22-25 Moreover, as well as MCAM+ Th cells, previous reports have indicated that CSF-accumulating Th17 cells also showed characteristics of pathogenic Th17 cells, and the frequency of this subset did not increase in the PB.22,25 These findings suggest that pathogenic Th17 cells do not necessarily increase in both the PB and CSF.

We speculated that, at first, an unknown cause, such as a genetic and/or environmental factor (viral infection, smoking, and low sun exposure), triggered autoimmune activation and induced BBB damage (or increased BBB permeability) in patients with MS.1,2,6-29 Subsequently, pathogenic lymphocytes, such as MCAM+ mTh cells, migrated from the PB to the CSF and the CNS, owing to their strong migration capacity,10-13,19,23 without a quantitative change in the PB. Then, inflammatory demyelination occurred. Unlike in MS, the frequency and the absolute number of MCAM+ lymphocytes (especially mTh cells) increased in both the PB and CSF in NMOSD. Thus, we speculated that the activation of MCAM-related pathogenic Th17 cell autoimmunity occurred not only in the CNS but also systemically, including in the PB in NMOSD, with a quantitative change in the PB. Our speculation is supported by previous studies in which more patients with NMOSD than those with MS tend to present with comorbid autoimmune disorders, such as systemic lupus erythematous, Sjögren syndrome, MG, and autoimmune thyroid diseases.16,30 Moreover, Th17 cell–related pathogenic autoimmunity has been shown in patients with NMOSD.31-34 A previous study demonstrated that AQP4-specific T cells in the PB of patients with NMOSD exhibited a Th17 bias that promoted neutrophil infiltration through IL-17.31 In addition, a meta-analysis showed that the proportion of PB Th17 cells was higher in patients with NMOSD than that in those with MS and/or in controls.33 Therefore, we speculated that these previously reported Th17 cells in NMOSD included MCAM+ Th cells.

In this study, there was no difference in the frequencies of MCAM+ lymphocytes in patients with MS treated with different DMDs. However, although the number of patients treated with DMDs was small, the absolute number of MCAM+ lymphocytes was significantly lower in FTY-treated patients with MS than that in other DMD-treated or untreated patients with MS (Figure 2B). FTY is a sphingosine-1-phosphate (S1P) modulator that binds to the S1P receptor. This results in the sequestration of T cells and B cells in the lymph nodes; there is a reduction in the number of PB lymphocytes, and the migration of the lymphocytes from the PB to the CNS is inhibited.35-37 This study partly explains the mechanism underlying the therapeutic effects of FTY, inhibiting the migration of MCAM+ lymphocytes from the PB to the CNS. Other DMDs are effective in suppressing the disease activity; however, they do not reduce the number of PB MCAM+ lymphocytes. NTZ binds the α4-integrin subunit expressed on the lymphocyte and prevents the migration of the lymphocyte into the CNS; however, NTZ does not decrease the number of peripheral lymphocytes.38 Therefore, the results from this study are consistent with that for the mechanism of action for NTZ. The mechanisms for IFN-β and DMF are less understood.36,37 IFN-β does not change the frequency and the absolute number of Th17 cells; however, IFN-β inhibits the Th17 axis and enhances the production of anti-inflammatory cytokines such as...
IL-10 and IL-4. Therefore, it is understandable that IFN-β suppresses the disease activity without decreasing the number of MCAM+ lymphocytes. Similar to FTY, DMF reduces the PB lymphocytes, including the Th17 cells. In this study, the frequency and the absolute number of MCAM+ lymphocytes were unaltered in patients treated with DMF. One of the reasons could be the lack of effect on the MCAM expression in the lymphocytes. The small number of patients treated with DMF and/or the duration of DMF treatment in this study could also have contributed to this effect. Despite the lack of effects on the frequency and the number of MCAM+ lymphocytes, DMF could modulate the function of MCAM+ lymphocytes through mechanisms such as the upregulation of the nuclear factor erythroid-derived 2-related antioxidant pathway and the inhibition of the nuclear factor-κB translocation.

There are certain limitations in this study. First, we used cryopreserved PBMC and CSF samples; cryopreservation may result in the modification of the expression of MCAM. Of note, for consistency, we analyzed the expression of MCAM using cryopreserved samples from all patients. Second, the number of patients treated with DMDs was small and, therefore, not enough to speculate a relationship between DMDs and MCAM expression. Third, the number of patients with NMOSD and MG without steroid therapy was small. Thus, the possibility of the effects of steroid on MCAM expression must be considered. However, the frequency of MCAM+ mTh cells was significantly higher in PSL-treated patients with NMOSD than that in PSL-treated patients with MG, although there was no difference in PSL dosage between the groups. Therefore, we considered that the activation of MCAM-related Th17 cell autoimmunity occurred in both the periphery and the CNS in patients with NMOSD.

In conclusion, this study showed that MCAM+ mTh cells were elevated in only the CSF of relapsing patients with MS but in the PB and CSF of patients with NMOSD. These results suggest that MCAM+ Th17 cells contribute to the pathogenesis of both MS and NMOSD through different mechanisms. These findings suggest that MCAM could be a therapeutic target and biomarker of CNS IDDs, and further research is needed to elucidate the detailed role and underlying mechanism of MCAM expression in CNS IDD pathogenesis.

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Disclosure
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Appendix

| Name                | Location                                      | Contribution                                      |
|---------------------|-----------------------------------------------|--------------------------------------------------|
| Ryotaro Ikeguchi, MD, PhD | Tokyo Women’s Medical University, Japan        | Design and conceptualized the study; data collection; analyzed data; and drafting the manuscript for intellectual content |
| Yoko Shimizu, MD, PhD          | Tokyo Women’s Medical University, Japan        | Design and conceptualized the study; data collection; analyzed data; and drafting the manuscript for intellectual content |
| Akiko Kondo, MD            | Tokyo Women’s Medical University, Japan        | Data collection and analyzed data                 |
| Natsuki Kanda, MD          | Tokyo Women’s Medical University, Japan        | Data collection and analyzed data                 |
| Hayato So, MD              | Tokyo Women’s Medical University, Japan        | Data collection and analyzed data                 |
| Haruka Kojima, MD          | Tokyo Women’s Medical University, Japan        | Data collection and analyzed data                 |
| Kazuo Kitagawa, MD, PhD    | Tokyo Women’s Medical University, Japan        | Drafting and revision of the manuscript           |

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