Distinct T cell signatures define subsets of patients with multiple sclerosis

ABSTRACT

Objective: We investigated T cell responses to myelin proteins in the blood of healthy controls and 2 groups of patients with relapsing-remitting multiple sclerosis (RRMS) who exhibited lesions either predominantly in the brain or predominantly in the spinal cord in order to assess whether distinct neuroinflammatory patterns were associated with different myelin protein-specific T cell effector function profiles and whether these profiles differed from healthy controls.

Methods: Peripheral blood mononuclear cells were obtained from patients with brain-predominant RRMS, patients with spinal cord-predominant RRMS, and age-matched healthy controls and analyzed by enzyme-linked immunosorbent spot assays to quantify interferon gamma-secreting (Th1) and interleukin 17-secreting (Th17) cells responding directly ex vivo to myelin basic protein (MBP) and myelin oligodendrocyte glycoprotein (MOG).

Results: Although MBP and MOG elicited different responses, patients with multiple sclerosis (MS) who had spinal cord-predominant lesions exhibited significantly higher Th17:Th1 ratios in response to both MBP and MOG compared to patients with brain-predominant MS. Incorporating the cytokine responses to both antigens into logistic regression models showed that these cytokine responses were able to provide good discrimination between patients with distinct neuroinflammatory patterns.

Conclusions: Our findings suggest that the localization of lesions within the brain vs the spinal cord in patients with MS is associated with different effector T cell responses to myelin proteins. Further investigation of the relationship between T cell effector function, antigen specificities, and lesion sites may reveal features of pathogenic pathways that are distinct to patients with different neuroinflammatory patterns.

Neurrol Neuroimmunol Neuroinflamm 2016;3:e278; doi: 10.1212/NXI.0000000000000278

GLOSSARY

AUC = area under the curve; BRI = Benaroya Research Institute; EAE = experimental autoimmune encephalomyelitis; HC = healthy control; IFN-γ = interferon gamma; IL-17 = interleukin 17; MBP = myelin basic protein; MOG = myelin oligodendrocyte glycoprotein; MS = multiple sclerosis; PBMC = peripheral blood mononuclear cell; RRMS = relapsing-remitting multiple sclerosis.

Inflammatory lesions and plaques of demyelination are hallmark features of multiple sclerosis (MS). Most patients with relapsing-remitting MS (RRMS) exhibit lesions disseminated predominantly in the brain. In up to 5% of patients with RRMS, however, lesions are localized primarily within the spinal cord instead of the brain. Spinal cord lesions often affect motor function and mobility, typically resulting in a higher degree of disability.

In experimental autoimmune encephalomyelitis (EAE), several studies have linked different immune mediators with preferential induction of brain vs spinal cord inflammation. For example, expression of neutrophil chemoattractants induced by interleukin 17 (IL-17) signaling has been associated with brain inflammation, while interferon gamma (IFN-γ) signaling has been
shown to promote spinal cord inflammation. We previously demonstrated that parenchymal brain inflammation in EAE was enhanced when the ratio of IL-17–secreting T cells (Th17 cells) to IFN-γ–secreting T cells (Th1 cells) infiltrating the CNS was ≥1. When the Th17:Th1 was ≤1, parenchymal inflammation was largely restricted to the spinal cord. Therefore, we hypothesized that the Th17:Th1 ratio of myelin protein–specific T cells may influence the extent of lesion burden in the brain vs spinal cord in patients with MS. While previous comparisons between healthy controls (HCs) and patients with MS of the frequency of myelin antigen–specific IFN-γ- and/or IL-17–producing T cells in peripheral blood mononuclear cells (PBMCs) have been contradictory, no study has investigated T cell responses to myelin antigens in patients stratified according to neuroinflammatory patterns. We investigated the frequency of T cells secreting these cytokines following stimulation with myelin basic protein (MBP) or myelin oligodendrocyte glycoprotein (MOG) in PBMCs from patients with RRMS who had lesions localized predominantly in the brain or the spinal cord. Strikingly, the cytokine responses differed when triggered by exposure to different myelin antigens, and the combined responses to both myelin antigens defined distinct T cell signatures that distinguished patients with MS who had different neuroinflammatory patterns. These results suggest that regional lesion localization may be associated with myelin-specific T cells with different properties.

### METHODS

**Patients and HCs.** Cryopreserved PBMCs were obtained from the Immune-Mediated Disease Registry maintained at the Benaroya Research Institute (BRI) from patients with RRMS exhibiting lesions primarily in the brain (“brain-predominant,” n = 29) or predominantly in the spinal cord (“spinal cord–predominant”); a total of 16 matching our criteria were available in the registry) based on MRI at the time of blood draw. Only patients with MRI lesions typical of MS in morphology and location (infratentorial, juxtaocular, periventricular, spinal cord) were included (see table e-1 at Neurology.org/nn for patient and HC demographics). Brain scans were graded as follows: 0 = normal brain, 1 = all nonspecific lesions, 2 = 1 lesion, 3 = 2–4 lesions, 4 = 5–8 lesions, and 5 = >8 typical MS lesions. Spinal cord scans were graded by the cumulative area of spinal cord segments affected by MS lesions. A priori definition of brain-predominant MS required a brain grade of 3–4 with ≤0.5 affected spinal cord segments or brain grade of 5 with ≤1 affected cord segment. Spinal cord–predominant MS was defined as brain grade of 1 with ≥2 affected cord segments, brain grade of 2 with ≥2.5 affected cord segments, brain grade of 3 with ≥6 affected cord segments, or brain grade 4 with ≥12 affected cord segments. Characteristics of MRI scans for the 2 cohorts are shown in table 1. No differences between patients with MS who had brain-predominant vs spinal cord–predominant lesions were seen in the distribution of 33 different HLA-DRB1 alleles (including HLA-DRB1*1501 and DRB1*0401) and 15 HLA-DQB1 alleles that are routinely analyzed when patients enter the registry (data not shown). Only patients who were within 5 years of diagnosis of MS and not treated with disease-modifying agents for at least 3 months before time of blood draw were selected for the study. Historically, only 3 brain-predominant patients and zero spinal cord–predominant patients had ever been on immune-modifying therapy. Only 3 brain-predominant and 2 spinal cord–predominant patients exhibited clinical signs at time of blood draw. Neuromyelitis optica was ruled out in spinal cord–predominant patients based on lesion characteristics observed on MRI; patients with spinal cord lesions affecting >3 spinal cord segments were excluded. Consistent with a diagnosis of MS, all spinal cord–predominant patients exhibited some brain lesions. Age-, ethnicity- and sex-matched HCs (n = 45) had no history of autoimmunity or signs of infection at time of blood draw. Demographics of the patients and controls including Expanded Disability Status Scale scores are shown in table e-1.

### Table 1

<p>| MRI-based criteria used to identify patients with brain-predominant and spinal cord–predominant MS |
|-------------------------------------------------|-------------------------------------------------|-------------------------------------------------|-------------------------------------------------|-------------------------------------------------|-------------------------------------------------|</p>
<table>
<thead>
<tr>
<th>MRI brain grade</th>
<th>No.</th>
<th>Median no. of spinal cord segments with lesions</th>
<th>Range in no. of spinal cord segments with lesions</th>
<th>No.</th>
<th>Median no. of spinal cord segments with lesions</th>
<th>Range in no. of spinal cord segments with lesions</th>
</tr>
</thead>
<tbody>
<tr>
<td>5</td>
<td>17</td>
<td>0</td>
<td>0–1.0</td>
<td>0</td>
<td>NA</td>
<td>NA</td>
</tr>
<tr>
<td>4</td>
<td>7</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>12</td>
<td>NA</td>
</tr>
<tr>
<td>3</td>
<td>5</td>
<td>0</td>
<td>0–0.5</td>
<td>4</td>
<td>7.5</td>
<td>6–11.0</td>
</tr>
<tr>
<td>2</td>
<td>0</td>
<td>NA</td>
<td>NA</td>
<td>3</td>
<td>3.5</td>
<td>2.5–7.0</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>NA</td>
<td>NA</td>
<td>8</td>
<td>5</td>
<td>2.0–8.0</td>
</tr>
</tbody>
</table>

Abbreviations: MS = multiple sclerosis; NA = not applicable.

Patients with MS were initially stratified according to severity of lesion burden in the brain as described in the methods section and then evaluated for the number of affected spinal cord segments. The combinations of brain and spinal cord features used to define patients with brain-predominant and spinal cord–predominant MS are indicated.
Standard protocol approvals, registrations, and patient consents. Written informed consent was obtained under protocols approved by the institutional review board at BRI before inclusion in this study.

Enzyme-linked immunosorbent spot assays. All assays were performed in a blinded manner. Ninety-six-well plates (MSP4W10; Millipore Corp., Billerica, MA) were coated with 3 μg/mL anti-IFN-γ (clone MD-1) or anti-IL-17A (clone eBio64CAP17) antibodies (eBioscience, San Diego, CA). PBMCs were thawed at 37°C, resuspended in room-temperature fetal bovine serum, overlaid with phosphate-buffered saline, pelleted, washed, and resuspended in complete RPMI (Roswell Park Memorial Institute medium) media with 10% human serum. PBMCs were plated at 5 × 10⁶ cells/well (IFN-γ assay) or 1 × 10⁶ cells/well (IL-17A assay) either alone (background) or with 50 μg/mL human MBP (cat. no. MO689; Sigma-Aldrich, St. Louis, MO), 50 μg/mL recombinant human MOG (residues 30–154, cat. no. 228-11134-3; RayBiotech, Norcross, GA), or tetanus toxoid (5 μg/mL; Astarte, Bothell, WA). Myelin antigen–specific responses were assayed in triplicate, media alone, and tetanus toxoid–specific responses in duplicate, and single wells of PBMCs were stimulated with anti-CD3/anti-CD28 antibodies (5 ng/mL; eBioscience). Plates were incubated for 24 (IFN-γ) or 48 (IL-17A) hours at 37°C, washed, and incubated with biotinylated detection antibodies (eBioscience) for 1 to 2 hours, followed by streptavidin alkaline phosphatase (cat. no. 7100-04; SouthernBiotech, Birmingham, AL) for 1 hour. Plates were developed using the Vector Blue AP substrate kit (Vector Laboratories, Burlingame, CA); spots were counted and analyzed using the CTL ImmunoSpot analyzer (Cellular Technology Ltd., Shaker Heights, OH). Background spots were subtracted before calculating the frequency of specific spots. In some experiments, PBMCs were incubated with 10–50 μg/mL of pan-MHC class I blocking antibody (clone W6/32; eBioscience) or 10 μg/mL of anti-HLA-DR (clone L243) and anti-HLA-DQ (clone SPV3) on ice for 30 to 60 minutes before adding myelin antigen. Low interassay variance was demonstrated by repetitive analyses of 10 HCs for their responses to tetanus toxoid in 4 independent assays. Quadruplicate values for the 10 HCs had an R² range from 0.85 to 0.95, with a mean correlation R² value of 0.91 (Spearman rank coefficient correlation), demonstrating very good replication of results.

Statistics. Analyses were performed using GraphPad Prism 6 (GraphPad Software, La Jolla, CA). The Mann–Whitney test was used to compare frequencies of myelin-specific cells and ratios of Th1:Th17 cells between groups. A Wilcoxon test was used for interantigen comparisons of Th1 and Th17 responses from the same patients. Two-way analysis of variance with Tukey multiple comparison test was used for MHC class I- and class II-blocking experiments. To test for normal distribution, a D’Agostino and Pearson omnibus normality test was utilized. A p value of <0.05 was considered to be statistically significant. A multiple imputation procedure was used to address the missing MOG-specific cytokine values in the logistic regression models, and 95% confidence intervals for these models were calculated from a nonparametric bootstrap (resampling) procedure.

RESULTS MBP-specific Th1 but not Th17 responses differ between subsets of patients with MS. To test our hypothesis that myelin antigen–specific Th1:Th17 ratios would differ for patients with MS who had brain-predominant vs spinal cord–predominant lesions, we analyzed PBMCs from HCs and patients with RRMS who exhibited lesions either predominantly in the brain or in the spinal cord (see methods and table 1 for patient classification). All patients were within 5 years of diagnosis of MS and the average number of years since first experiencing a symptom associated with MS was similar for the patient subsets: 5.9 years for the brain-predominant group and 5.2 years for the spinal cord–predominant group. We analyzed PBMCs in a blinded manner using enzyme-linked immunosorbent spot to determine directly ex vivo the number of cells that secreted IFN-γ and IL-17 following stimulation with MBP. A significantly higher frequency of MBP-specific IFN-γ⁺ cells was detected in brain-predominant patients compared to both HCs and spinal cord–predominant patients (figure 1A). Unexpectedly, few PBMCs from the patients with spinal cord–predominant MS produced IFN-γ in response to MBP. Approximately half (7/16) of the spinal cord–predominant patients generated no MBP-specific IFN-γ⁺ response above background, even though comparable IFN-γ⁺ responses to stimulation with tetanus toxoid or anti-CD3/CD28 antibodies were observed among all groups (figure e-1). The number of MBP-specific spots was reduced 46% to 78% when PBMCs were preincubated with pan anti-human MHC class II–blocking antibodies, but no decrease was observed following preincubation with pan-MHC class I–blocking antibodies (figure e-2), indicating that the myelin antigen–specific responses detected in these assays reflect CD4⁺ T cell activity. No correlation was seen between the number of MBP-specific spots and spots detected after anti-CD3/CD28 stimulation, indicating that variation in T cell number among PBMC samples did not influence quantification of antigen-specific responses. Of note, the frequency of MBP-specific IL-17⁺ cells was not different among the 3 groups (figure 1B). Therefore, the MBP-specific Th17:Th1 ratio was significantly higher for patients with spinal cord–predominant MS compared to both patients with brain-predominant MS and HCs because of the minimal MBP-specific IFN-γ⁺ response observed for the spinal cord–predominant patient group (figure 1C). T cells exhibit different effector responses to MOG vs MBP. We next analyzed T cell responses to human MOG using PBMC samples from the same blood draw as the PBMCs used to analyze MBP-specific responses. Additional samples from the same blood draw were available for 16 of 45 HCs, 16 of 29 brain-predominant, and 13 of 16 spinal cord–predominant patients with MS. Similar to MBP-specific responses, patients with brain-predominant MS tended toward a higher frequency of MOG-
specific IFN-γ+ spots compared to both HCs and spinal cord–predominant patients (figure 2A). However, the MOG-specific responses in patients with spinal cord–predominant MS differed from their MBP-specific responses. In contrast to their minimal MBP-specific IFN-γ+ responses, the frequency of MOG-specific IFN-γ+ cells was similar between patients with spinal cord–predominant MS and HCs. Furthermore, the frequency of MOG-specific IL-17+ cells in patients with spinal cord–predominant MS was significantly higher compared to patients with brain-predominant MS. The MOG-specific Th17:Th1 ratio was again significantly higher for spinal cord–predominant compared to brain-predominant patients with MS (figure 2C), although the factors that caused these higher ratios differed between the MBP- and MOG-specific responses. For MBP-specific responses, the major influence was the low IFN-γ response detected for patients with spinal cord–predominant MS. For MOG-specific responses, increased IL-17+ cells in patients with spinal cord–predominant MS and increased IFN-γ+ cells in patients with brain-predominant MS were the driving factors in Th17:Th1 ratios.

A comparison of myelin protein specificities among Th1 and Th17 cells demonstrated that Th1 cells were predominantly MOG-specific in all 3 groups (figure 3). In contrast, Th17 cells were not biased toward MOG specificity. Patients with spinal cord–predominant MS and HCs exhibited a more even distribution between MBP and MOG.
The relative frequencies of MBP- and MOG-specific T cells differ among Th1 and Th17 effector subsets

**A. Healthy controls**

For IFN-γ and IL-17 PBMC responses, the frequencies of MBP-specific vs MOG-specific cells are compared for each individual in (A) healthy control, (B) brain-predominant, or (C) spinal cord-predominant multiple sclerosis patient groups. Asterisks indicate significant differences in the frequencies of MBP vs MOG-specific responses. *p < 0.05, **p < 0.01, ***p < 0.001, Wilcoxon test, 2-sided. IFN = interferon; IL = interleukin; MBP = myelin basic protein; MOG = myelin oligodendrocyte glycoprotein; PBMCs = peripheral blood mononuclear cells.

**DISCUSSION** We hypothesized from our studies in EAE that differences in effector T cell frequencies would be observed between subsets of patients with MS that were stratified according to their lesion burden in the brain vs the spinal cord, and that this

specificities in Th17 cells, while Th17 cells were predominantly MBP-specific among patients with brain-predominant MS. These data reveal group-specific differences in the effector T cell responses elicited by distinct myelin proteins.

**Effector T cell responses distinguish patients with spinal cord–predominant from brain-predominant MS.** To determine whether particular cytokine responses performed best in discriminating between our patient and HC groups, we performed pair-wise comparisons of antigen-specific responses for all individuals who were analyzed with both myelin antigens. Plotting the frequency of MBP-specific IFN-γ vs MOG-specific IL-17+ cells for each individual was most effective in separating the data points belonging to spinal cord–predominant patients from brain-predominant patients (figure 4A), highlighting these antigen-specific, effector T cell responses as being the most disparate between these 2 subsets of patients with MS.

We then generated logistic regression models, which compare 2 groups by generating for each individual in a group a single summary value, or “optimal marker” that optimizes the separation between the groups. Here, the optimal markers for each individual are derived from their MBP- and MOG-specific IFN-γ+ and IL-17+ responses. We employed the widely used bootstrap variance methodology to generate values for MOG-specific responses for subjects who were analyzed only for MBP responses because of lack of additional PBMC samples. Receiver operating characteristic analyses were performed that determine the true-positive rates and false-positive rates of group assignment based on the optimal markers generated by the models. Graphs of the true-positive vs false-positive rates for each 2-group comparison are shown in figure 4B. The area under the curve (AUC) reflects how well the 2 groups are separated based on their optimal marker values. A perfect separation between groups results in an AUC = 1, while no separation between groups results in an AUC = 0.5. An AUC value ≥0.7 is considered a fair separation while an AUC ≥0.9 is considered very good. The AUC for the comparison of patients with brain-predominant vs spinal cord–predominant MS was 0.83 (95% confidence interval 0.69–0.98, p < 0.0001). The AUC generated by comparison of HCs to patients with spinal cord–predominant MS was 0.70 (95% confidence interval 0.53–0.87, p < 0.03), while comparison of HCs to patients with brain-predominant MS resulted in the lowest AUC of 0.66 (95% confidence interval 0.50–0.81, p < 0.061). These analyses indicate that the combined IFN-γ+ and IL-17+ T cell responses to MBP and MOG provide a good means of discriminating between patients with spinal cord–predominant vs brain-predominant lesions, and that these responses might also distinguish patients with spinal cord–predominant MS from HCs. The responses were less robust in discriminating between patients with brain-predominant MS and HCs.
stratification may be necessary to observe differences in effector T cell frequencies between patients with MS and HCs. To test our hypothesis, we selected patients for analyses of cytokine responses in whom lesions were localized predominantly in the brain or in the spinal cord based on MRI. Our data provide the first evidence that the regional localization of lesions within the CNS in patients with MS is associated with different effector T cell responses to myelin proteins. Specifically, a low frequency of MBP-specific Th1 cells and a higher frequency of MOG-specific Th17 cells were associated with spinal cord–predominant compared to brain-predominant patients with MS in this study. Consistent with our hypothesis, there is no significant difference in myelin antigen–specific T cell responses if we combine the data from our 2 MS patient populations into one group and compare it to HCs (figure e-3).

Our finding that MBP and MOG elicited different effector T cell responses in both MS patient subsets and HCs was unexpected but may explain discrepancies in earlier studies that used different myelin antigens to stimulate PBMCs. Of note, individuals within a group exhibited a similar pattern of
responses to each myelin antigen, and these patterns differed for the 2 subsets of patients with MS. The ability of myelin antigen–specific T cell responses to distinguish these cohorts of patients with MS who had spinal cord–predominant lesions from patients who had brain-predominant lesions was illustrated by the receiver operating characteristic analyses of logistic regression models that incorporated all of the T cell response data. These analyses indicated that the ability to distinguish spinal cord–predominant vs brain-predominant MS was good, and the ability to distinguish patients with spinal cord–predominant MS from HCs was fair. Of note, discrimination between the HCs and patients with brain-predominant MS was the weakest. It will be important to conduct similar studies on additional patients with MS stratified according to their neuroinflammatory patterns.

The MBP-specific IFN-\(\gamma\)^+ response was the most influential measure in determining the optimal marker values generated by the logistic regression models for each 2-group comparison, consistent with the fact that the MBP-specific IFN-\(\gamma\)^+ responses exhibited the greatest differences among the 3 groups. Paradoxically, patients with MS who had brain-predominant lesions exhibited significantly higher MBP-specific IFN-\(\gamma\)^+ responses compared to HCs while the MBP-specific IFN-\(\gamma\)^+ responses of patients with MS who had spinal cord–predominant lesions exhibited the opposite patterns and trended lower (\(p = 0.069\)) compared to HCs, highlighting the difference in T cell signatures seen in this study between patients with MS who exhibited distinct neuroinflammatory patterns. While other CNS antigens have been implicated in MS, these perturbations in myelin antigen specificities, and lesion sites may provide new insights into the pathogenic mechanisms that shape neuroinflammatory patterns in patients with MS.

One question that arises from our studies is whether T cell responses detected in PBMCs are indicative of effector T cell activity within the CNS, or whether peripheral T cell responses reflect the inverse of T cell activity within the CNS because the CNS acts as a “sink” for pathogenic T cells. A significantly higher frequency of myelin-specific IFN-\(\gamma\)^+ T cells was found in the CSF compared to PBMCs in patients with MS \(^{10,12,19}\); however, comparisons need to be performed for multiple antigen specificities in individual patients to determine whether there is an inverse relationship or direct correlation between frequencies of peripheral vs CSF myelin-specific T cells. It is of interest that patients with spinal cord–predominant MS exhibited a higher Th17:Th1 ratio compared to patients with brain-predominant MS for both myelin antigen specificities, although the ratio was influenced by different factors for each myelin protein. Our EAE studies showed that a lower Th17:Th1 ratio preferentially promoted spinal cord parenchymal inflammation, but this finding was based on T cells isolated from the CNS rather than the periphery. Therefore, if peripheral T cell responses in humans represent the inverse of responses that would be found in the CNS due to a “sink effect,” our findings in humans would be similar to those in mice with EAE.

The key finding from our study is that we identified T cell cytokine signatures based on production of IFN-\(\gamma\) and IL-17 by MBP- and MOG-specific T cells in PBMCs that were associated with patients with MS exhibiting different neuroinflammatory patterns. The higher Th17:Th1 ratio for both MBP- and MOG-specific T cells in patients with spinal cord–predominant MS could reflect differences in the nature of the pathogenic immune response that leads to different neuroinflammatory patterns in patients with MS. Alternatively, the local environment in the brain vs spinal cord may favor expansion of certain effector T cells specific for particular antigens over others. Future studies of new cohorts of patients with MS are warranted to confirm and extend the T cell signatures that we have described here. The ability to distinguish between patients with MS who have predominantly spinal cord vs brain lesions based on peripheral responses to myelin antigens would be valuable in the clinical setting if the type of neuroinflammatory pattern that a patient will develop can be predicted at the time of disease onset or in response to therapy. As spinal cord involvement in MS is a predictor of disability independent of the extent of brain involvement, \(^7\) prediction of increased lesion burden in the spinal cord may be beneficial. Further understanding of the relationships among T cell effector function, antigen specificities, and lesion sites may provide new insights into the pathogenic mechanisms that shape neuroinflammatory patterns in patients with MS.

**AUTHOR CONTRIBUTIONS**

M.C.J. and E.R.P. helped design the study, performed experiments, analyzed data and helped write the manuscript. A.J.S. analyzed data using logistic regression models. A.S.N. and M.K. evaluated patient MRIs. S.P. coordinated patient data. J.H.B helped design the study and interpret data. J.M.G. designed the study, oversaw the experiments, and wrote the manuscript.

**ACKNOWLEDGMENT**

The authors thank the BRI clinical core for the acquisition and handling of patient samples, Catriona Wagner for critical reading of the manuscript, and Michael Mason for consultation related to statistical analysis.

**STUDY FUNDING**

This research was supported by grants to J.M.G. from the National Institute for Allergy and Infectious Disease (R37 AI107494-01 and pilot grant support from U01 AI101990). Initial development of the immune-mediated disease registry at BRI was supported by a grant provided by the Life Science Discovery fund of Washington State.
REFERENCES


Distinct T cell signatures define subsets of patients with multiple sclerosis
Mark C. Johnson, Emily R. Pierson, Andrew J. Spieker, et al.
Neurol Neuroimmunol Neuroinflamm 2016;3;
DOI 10.1212/NXI.0000000000000278

This information is current as of August 23, 2016