Requirement for endocytic antigen processing and influence of invariant chain and H-2M deficiencies in CNS autoimmunity

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The role of processing in antigen (Ag) presentation and T cell activation in experimental allergic encephalomyelitis (EAE) was evaluated in wild-type mice, mice that selectively express either Ii p31 or p41, and mice completely deficient in Ii or H-2M. We demonstrate that processing of myelin oligodendrocyte glycoprotein (MOG) is required for presentation of the dominant encephalitogenic MOG epitope, p35-55. Ii p31- and p41-expressing mice developed EAE with similar incidence to wild-type mice, although p41 mice had a more severe course. Ag-presenting cells (APCs) from Ii- or H-2M–deficient mice could present p35-55, but not MOG, demonstrating that these APCs could not process native MOG. Ii- and H-2M–deficient mice were not susceptible to EAE by immunization with p35-55 or MOG or by adoptive transfer of encephalitogenic T cells. However, CD4+ T cells from p35-55–immunized H-2M–deficient mice proliferated, secreted IFN-γ, and transferred EAE to wild-type, but not H-2M–deficient, mice. Thus, EAE resistance in H-2M–deficient mice is not due to an inability of APCs to present p35-55, or an intrinsic defect in the encephalitogenic T cell repertoire, but reflects a defect in APC function. Our results indicate that processing is required for initial Ag presentation and CNS T cell activation and suggest that autopathogenic peptides of CNS autoantigen may not be readily available for presentation without processing.


Introduction

Experimental allergic encephalomyelitis (EAE) is an inflammatory demyelinating CNS disease that serves as a model for multiple sclerosis (MS) and other organ-specific T cell–mediated autoimmune diseases (1, 2). Activated CD4+ Th1 cells that recognize CNS self-antigens (autoantigens) mediate EAE and are thought to have a central role in MS pathogenesis (2). Activation of EAE-Th cells, which occurs when a specific Ag is presented in association with MHC class II molecules expressed on the surface of antigen presenting cells (APCs), may be required at different stages in the pathogenesis of CNS inflammatory disease (3–5). Ag presentation by APCs outside the CNS can lead to T cell activation, a prerequisite for T cell entry into the CNS (3, 4). Ag presentation by resident CNS accessory cells, which are nonprofessional APCs, may be necessary for recognition of CNS autoantigen and T cell activation during the initial CNS inflammation (3) and T cell activation during progression from acute to chronic and relapsing stages of CNS demyelinating disease (5). In general, Ag presentation by APCs requires “processing,” which involves proteolytic Ag degradation in endocytic/lysosomal compartments and association of peptide cleavage products with class II molecules before display on the surface of APCs (6). In vitro studies have established that endocytic processing of the native form of the CNS autoantigen, myelin basic protein (MBP), is necessary for presentation to encephalitogenic CD4+ T cells (7). If peptides of myelin autoantigens are not readily accessible for Ag presentation in normal CNS, endocytic processing by CNS APCs may be required for initial Ag presentation and T cell activation in vivo.

Invariant chain (Ii) and H-2M (HLA-DM) are molecules involved in class II biosynthesis and endocytic processing (8). Ii serves as a chaperone, directing transport of newly formed class II α and β chains to the endosomal/lysosomal compartment (8). Two isoforms of Ii are expressed, p31 and p41 (9). Although Ii p31 and p41 both facilitate class II assembly and prevent premature Ag binding to class II molecules (8), some in vitro data have indicated that Ii p41 may modulate the proteolytic environment of the endocytic compartment and confer selective advantage in Ag presentation to CD4+ T cells (10–12). As Ii p31 and p41 are proteolytically cleaved, the residual peptide
fragment, CLIP (class II-associated invariant chain peptide), remains associated to class II molecules (13). H-2M, a nonclassical class II molecule, facilitates the removal of CLIP, permitting the binding of peptide Ag’s for T cell presentation (14). Deficiencies in either Ii or H-2M, can cause defects in Ag processing (8), and have been instrumental in evaluating the requirements for intracellular peptide loading and Ag presentation in vivo (15–20).

In this investigation we evaluated the role of Ag processing in EAE by examining the T cell response to myelin oligodendrocyte glycoprotein (MOG) in mice that selectively express Ii p31 or p41 and mice that are completely deficient in Ii or H-2M molecules. We established that processing of native MOG by APC is required for presentation of the dominant encephalitogenic MOG peptide, p35-55. Although both Ii p31 and p41 mice were susceptible to EAE, Ii p41 mice sustained a more severe course. Our in vivo studies utilizing mice completely deficient in Ii or H-2M indicate that processing is required for Ag presentation during the induction of T cell–mediated CNS inflammation. Results in this study indicate that peptides of this CNS autoantigen may not be available for class II restricted Ag presentation in normal CNS without processing. Thus, reagents that selectively interfere with endocytic processing could be beneficial for immunotherapy in Ag-specific CNS autoimmune disease.

Methods

Mice. Characterization of mice deficient in Ii, Ii p31, Ii p41, and H-2M has been described (15, 18, 21, 22). Deficiency in each of these genes has been bred onto the C57BL/6 background for four generations. Wild-type C57BL/6 mice, Ii-deficient mice, and H-2M–deficient mice used in adoptive transfer studies were purchased from The Jackson Laboratory (Bar Harbor, Maine, USA).

Antigens, antibodies, and recombinant cytokines. Mouse MOG peptide 35-55 (MEVGWYRSPFSRVVHLYRNGK) (Hopkinton, Massachusetts, USA) was synthesized and HPLC purified by QCB Inc. (Detroit, Michigan, USA). Immediately thereafter and fixed in 0.05% paraformaldehyde, 2 mM glutamine, 100 µg/ml penicillin, and 100 µg/ml streptomycin. A total of 1 × 10⁴ MOG p35-55–specific CD4⁺ T cells were cultured with 5 × 10⁵ γ-irradiated splenic APC and MOG peptide p35-55 or intact MOG in 0.2 ml culture media in 96-well microtiter plates (Falcon Labware, Oxnard, California, USA). At 48 hours, each well was pulsed with 1 µCi [³H]thymidine and harvested 16 hours later. In APC fixation experiments, 5 × 10⁴ splenic APCs were treated with 0.05% paraformaldehyde for 30 minutes at room temperature, washed three times, and cultured with 1 × 10⁴ MOG p35-55–specific T cells in the presence of either p35-55 or intact MOG. For primary proliferative responses, 5 × 10⁴ spleen cells removed from immunized mice were cultured in 0.2 ml media with p35-55 or native MOG. Cultures were pulsed at 72 hours with 1 µCi [³H]thymidine and harvested 16 hours later. In experiments using chloroquine, APCs were incubated with 75 µM chloroquine for 75 minutes and removed. After incubation with Ag for 30 minutes, MOG-specific T cells were added, pulsed at 72 hours, and harvested 16 hours later.

Cytokine analysis. Cell culture supernatants were collected at 24-hour incubation for cytokine analysis. Quantitative ELISA for IL-2, IL-4, and IFN-γ were performed using paired mAb’s specific for corresponding cytokines per manufacturer’s recommendations (PharMingen, San Diego, California, USA).

RT-PCR. Mice were perfused with PBS during euthanasia to remove circulating blood cells. Total RNA was then extracted with TRIZOL (GIBCO BRL; Life Technologies Inc., Rockville, Maryland, USA). RT and PCR were performed using the Access RT-PCR system from Promega Corp. (Madison, Wisconsin, USA) using one cycle: 48°C for 45 minutes followed by 94°C for 2 minutes. The following oligonucleotide primers (designed from published murine cDNA sequences; refs. 9, 24) were purchased from Operon (Alameda, California, USA): CIITA, 5′-(CCCTGCGTGTGATGGATGTC) and 5′-(GTGTCCTCTACTCGTCTTCAG); Ii, 5′-GAGGCTAGACCATGGATGAC-3′ and 5′-(3′AGATGCTTGATATTCTTGCCC-3′); H-2M, 5′-CTCAGGAGATGGTGATGGGAAAGT-3′ and 5′-GTGTCAGGCTTACATGCTGCTG-3′; class II β-chain, 5′-GCTACCTCCAACAGGGGAG-3′ and 5′-GCTCTTGGCAGGGATGCT-3′; β-actin, 5′-TTGTGATGGTGGGAAATG-GTCAG-3′ and 5′-TTTGTGATCCACCGATTTCC-3′. Primers for internal CIITA detect an expected 635-bp fragment; primers for Ii detect an expected 490-bp fragment; primers for an H-2M detect a 320-bp fragment; and primers for β-actin detect a 510-bp fragment. PCR was 1134

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Histological evaluation of tissue. At the completion of the experiment on day 30, mice were anesthetized by isoflurane inhalation and perfused with PBS containing 4% (vol/vol) paraformaldehyde. Fixed CNS tissues were embedded in paraffin wax, cut in 5-µm sections, and stained with hematoxylin and eosin. Stained sections were examined in a blind fashion as described previously (23).

Proliferation assays. Cells were cultured in serum-free medium, X-Vivo 20 (BioWhittaker Inc., Walkersville, Maryland, USA) supplemented with 5 × 10⁻⁵ M 2-mercaptoethanol, 2 mM glutamine, 100 µg/ml penicillin, and 100 µg/ml streptomycin. A total of 1 × 10⁴ MOG p35-55–specific CD4⁺ T cells were cultured with 5 × 10⁵ γ-irradiated splenic APC and MOG peptide p35-55 or intact MOG in 0.2 ml culture media in 96-well microtiter plates (Falcon Labware, Oxnard, California, USA). At 48 hours, each well was pulsed with 1 µCi [³H]thymidine and harvested 16 hours later. In APC fixation experiments, 5 × 10⁴ splenic APCs were treated with 0.05% paraformaldehyde for 30 minutes at room temperature, washed three times, and cultured with 1 × 10⁴ MOG p35-55–specific T cells in the presence of either p35-55 or intact MOG. For primary proliferative responses, 5 × 10⁴ spleen cells removed from immunized mice were cultured in 0.2 ml media with p35-55 or native MOG. Cultures were pulsed at 72 hours with 1 µCi [³H]thymidine and harvested 16 hours later. In experiments using chloroquine, APCs were incubated with 75 µM chloroquine for 75 minutes and removed. After incubation with Ag for 30 minutes, MOG-specific T cells were added, pulsed at 72 hours, and harvested 16 hours later.

Cytokine analysis. Cell culture supernatants were collected at 24-hour incubation for cytokine analysis. Quantitative ELISA for IL-2, IL-4, and IFN-γ were performed using paired mAb’s specific for corresponding cytokines per manufacturer’s recommendations (PharMingen, San Diego, California, USA).

RT-PCR. Mice were perfused with PBS during euthanasia to remove circulating blood cells. Total RNA was then extracted with TRIZOL (GIBCO BRL; Life Technologies Inc., Rockville, Maryland, USA). RT and PCR were performed using the Access RT-PCR system from Promega Corp. (Madison, Wisconsin, USA) using one cycle: 48°C for 45 minutes followed by 94°C for 2 minutes. The following oligonucleotide primers (designed from published murine cDNA sequences; refs. 9, 24) were purchased from Operon (Alameda, California, USA): CIITA, 5′-(CCCTGCGTGTGATGGATGTC) and 5′-(GTGTCCTCTACTCGTCTTCAG); Ii, 5′-GAGGCTAGACCATGGATGAC-3′ and 5′-(3′AGATGCTTGATATTCTTGCCC-3′); H-2M, 5′-CTCAGGAGATGGTGATGGGAAAGT-3′ and 5′-GTGTCAGGCTTACATGCTGCTG-3′; class II β-chain, 5′-GCTACCTCCAACAGGGGAG-3′ and 5′-GCTCTTGGCAGGGATGCT-3′; β-actin, 5′-TTGTGATGGTGGGAAATG-GTCAG-3′ and 5′-TTTGTGATCCACCGATTTCC-3′. Primers for internal CIITA detect an expected 635-bp fragment; primers for Ii detect an expected 490-bp fragment; primers for an H-2M detect a 320-bp fragment; and primers for β-actin detect a 510-bp fragment. PCR was
performed using 35 cycles of amplification: 94°C for 30 seconds, 62°C for 60 seconds, and 72°C for 120 seconds.

Results
Native MOG requires APC processing for presentation of the dominant encephalitogenic MOG peptide, p35-55. II- and H-2M–deficient mice have separate defects in maturation of MHC class II molecules. APC from H-2M–deficient mice express normal levels of mature (compact) class II molecules that are loaded with CLIP and have defects in processing native Ag (17, 18). Although II-deficient APCs have defects in expression of class II molecules and processing native Ag’s, they are remarkably efficient at presenting certain peptide Ag’s (15, 16). To examine whether endocytic processing of native MOG is required for presentation of the dominant encephalitogenic determinant, p35-55 (23), we first tested whether splenic APCs from wild-type C57BL/6 mice, II-deficient mice, and H-2M–deficient mice were capable of presenting p35-55 and intact MOG to II-Aβ–restricted p35-55–specific T cells. As shown in Figure 1a, II-deficient and wild-type APCs presented p35-55 equally well. H-2M–deficient APCs, which express class II molecules bound with CLIP, required more than one log higher p35-55 concentration to stimulate proliferation, which is consistent with other observations using H-2M–deficient APCs for presentation of exogenous peptide Ag’s (17, 18). In contrast, neither II-deficient nor H-2M–deficient APCs were very effective in presenting native MOG to p35-55–specific T cells compared with wild-type APC (Figure 1b). That II- and H-2M–deficient APCs could present p35-55, but not MOG, suggested that processing is required for presentation of native MOG. When tested at the same time, we observed that paraformaldehyde-fixed wild-type APCs (e), stimulated T cell proliferation using native MOG. For chloroquine treatment (d), APCs were treated with 75 µM chloroquine for 75 minutes before the addition of Ag and T cells (see Methods).
CNS expression of Ii and H-2M mRNA is upregulated during EAE. When stimulated by IFN-γ, astroglia cells cultured in vitro upregulate Ii and H-2M and present CNS autoantigen to encephalitogenic T cells (7). These results suggested that expression of class II processing elements might be important in CNS Ag presentation. Thus, we examined Ii and H-2M expression in the CNS of wild-type mice with and without EAE. Ii expression was detected in the brain of mice with EAE, but was not detected in normal brain (Figure 2). Ii message was barely detectable in normal spinal cord, but was clearly upregulated in EAE. Constitutive H-2Ma expression was detected in the spinal cord and brain of control mice, although there was marked H-2Ma upregulation in these tissues during EAE. Constitutive levels of Ii and H-2M mRNAs were detected in the spleens of both normal and EAE mice.

Ii p31 and p41 mice are susceptible to EAE, but mice completely deficient in Ii or H-2M are resistant. Our observations that Ii- and H-2M–deficient APCs could present p35-55, but not native MOG, enabled us to use Ii- and H-2M–deficient mouse strains to evaluate the requirement for processing in Ag presentation in EAE. Ii-deficient and H-2M–deficient mice were first examined for EAE susceptibility by direct immunization with p35-55 or rMOG (Table 1). In two separate experiments, none of the Ii-deficient or H-2M–deficient mice developed EAE (Table 1). However, after the initial onset of acute EAE, Ii p41 mice showed a more severe disease course compared with either Ii p31 or wild-type mice (Figure 3). In fact, the clinical score of EAE in p41 mice was approximately two times greater than in p31 mice.

Table 1
Incidence and severity of EAE in wild-type, Ii-deficient, and H-2M–deficient mouse strains

<table>
<thead>
<tr>
<th>Mice</th>
<th>Incidence</th>
<th>Mean maximum severity</th>
<th>Day of onset</th>
</tr>
</thead>
<tbody>
<tr>
<td>MOG p35-55</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C57BL/6</td>
<td>9/10</td>
<td>3.1 ± 0.3</td>
<td>17.2 ± 2.7</td>
</tr>
<tr>
<td>li–/–</td>
<td>0/10</td>
<td>N/A</td>
<td>N/A</td>
</tr>
<tr>
<td>li p31</td>
<td>0/10</td>
<td>3.5 ± 0.6</td>
<td>14.0 ± 0.9</td>
</tr>
<tr>
<td>li p41</td>
<td>10/10</td>
<td>3.6 ± 0.4</td>
<td>14.7 ± 1.7</td>
</tr>
<tr>
<td>H-2M–/–</td>
<td>0/10</td>
<td>N/A</td>
<td>N/A</td>
</tr>
<tr>
<td>rMOG</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C57BL/6</td>
<td>5/5</td>
<td>4.0 ± 0.6</td>
<td>17.8 ± 1.3</td>
</tr>
<tr>
<td>li–/–</td>
<td>0/3</td>
<td>N/A</td>
<td>N/A</td>
</tr>
<tr>
<td>li p31</td>
<td>5/5</td>
<td>4.2 ± 0.5</td>
<td>15.4 ± 1.0</td>
</tr>
<tr>
<td>li p41</td>
<td>5/5</td>
<td>4.7 ± 0.5</td>
<td>15.8 ± 1.2</td>
</tr>
<tr>
<td>H-2M–/–</td>
<td>0/5</td>
<td>N/A</td>
<td>N/A</td>
</tr>
</tbody>
</table>

*Mice were immunized subcutaneously in the footpads with 100 µg MOG p35-55 or 100 µg rMOG emulsified in an equal volume of CFA. Mice also received 150 ng PT immediately thereafter and again 48 hours later as described in Methods. *Mean maximum severity of affected animals. †Mean ± SEM.

Figure 3
H-2M– and Ii-deficient mice are resistant to EAE. Mean clinical score of mice immunized with (a) 100 µg pMOG35-55 or (b) 100 µg rMOG emulsified in an equal volume of CFA. Mice also received 150 ng PT at the time of immunization and again 48 hours later as described in Methods.
their CD4+ T cell compartments. Immunization of Ii-deficient mice with p35-55 did not stimulate a significant proliferative response (Figure 4a) or secretion of IFN-γ (Figure 4b), reflecting an inability to prime p35-55–specific CD4+ T cells. In contrast, T cells from p35-55–immunized H-2M–deficient mice proliferated nearly as well as T cells from wild-type and also secreted substantial IFN-γ. Thus, p35-55 elicited a Th1 response in H-2M–deficient, but not in Ii-deficient, mice. IL-4, a Th2 cytokine, was not detected in cultures from any mice tested (data not shown).

Lymphocytes from EAE-susceptible Ii p31 and p41 mice were also examined (Figure 4). Even though T cells from p35-55–immunized Ii p31 and p41 mice secreted less IL-2 than T cells from wild-type immunized mice, these IL-2 quantities were sufficient to support vigorous proliferation (data not shown). T cells from p35-55–immunized Ii p41 mice secreted similar levels of IFN-γ as T cells from wild-type mice, although T cells from p35-55–immunized p31 mice required nearly one log greater concentration of p35-55 to reach maximal detectable IFN-γ secretion.

Adoptive transfer of encephalitogenic MOG p35-55–specific T cells does not induce EAE in Ii-deficient or H-2M–deficient mice. As CLIP has a relatively high affinity for I-Ab (26), and it may be necessary for p35-55 to displace CLIP from class II molecules on H-2M–deficient APCs (17, 18), lack of EAE susceptibility in H-2M–recipient mice could reflect an inability of certain APCs to display sufficient p35-55–I-Ab complexes for adequate T cell activation. In contrast to H-2M–deficient APCs, Ii-deficient APCs express “empty” class II molecules (15, 16), and although Ii-deficient APCs may express fewer class II molecules, these APCs present p35-55 to encephalitogenic T cells as well as wild-type APCs (Figure 1).

To distinguish the influence of defects in Ag processing and presentation by APCs from alterations in selection of an encephalitogenic T cell repertoire, we examined whether activated wild-type encephalitogenic T cells could induce EAE when adoptively transferred into naive Ii- or H-2M–deficient mice. In contrast to wild-type recipient mice, H-2M– and Ii-deficient–recipient mice did not develop clinical EAE (Table 2) or histological signs of EAE (Figure 5).

As we observed that H-2M deficient APCs could present p35-55 to encephalitogenic T cells (Figure 1) and immunization of H-2M–deficient mice with p35-55 elicited proliferation and IFN-γ secretion (Figure 4), the reciprocal adoptive transfer using T cells from p35-55–immunized H-2M–deficient mice was performed. We observed that p35-55–specific cells isolated from H-2M–deficient mice induced EAE in wild-type, but not H-2M–deficient, recipient mice (Table 2). Although it is possible that Ii-deficient CNS APCs could not express sufficient MHC class II molecules, our results from H-2M–deficient mice indicate that it was the defect(s) in Ag processing, not an inability to present peptide Ag for T cell activation, that was responsible for the lack of EAE induction in H2-M–deficient mice. In contrast, lack of EAE in Ii-deficient mice reflected both a defect in Ag processing (Figure 1) and an inability to prime (Figure 3) an encephalitogenic p35-55–specific T cell repertoire.

Using a panel of Vβ-specific antibodies (Vβ 4, 5, 6, 8, 12, 13, 14, and 17) for FACS analysis, we examined the TCR repertoire of p35-55–specific encephalitogenic CD4+ T cells isolated from H-2M–deficient and wild-type mice. As reported previously (27), we observed that the repertoire of p35-55–specific T cells in wild-type C57BL/6 mice was heterogeneous (data not shown). In

**Table 2**

Adoptive transfer of encephalitogenic MOG35-55–specific T cells causes EAE in wild-type mice but not Ii-deficient or H-2M–deficient mice

<table>
<thead>
<tr>
<th>Donor cells</th>
<th>Recipient mice&lt;sup&gt;a&lt;/sup&gt;</th>
<th>Incidence</th>
<th>Mean maximum score</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild-type</td>
<td>Wild-type</td>
<td>9/13</td>
<td>3.6 ± 0.7</td>
</tr>
<tr>
<td>Wild-type</td>
<td>Ii–/–</td>
<td>0/10</td>
<td>0.0 ± 0.0</td>
</tr>
<tr>
<td>Wild-type</td>
<td>H-2Ma–/–</td>
<td>0/10</td>
<td>0.0 ± 0.0</td>
</tr>
<tr>
<td>H-2Ma–/–</td>
<td>Wild-type</td>
<td>7/10</td>
<td>3.2 ± 0.4</td>
</tr>
<tr>
<td>H-2Ma–/–</td>
<td>Ii–/–</td>
<td>0/5</td>
<td>0.0 ± 0.0</td>
</tr>
<tr>
<td>H-2Ma–/–</td>
<td>H-2Ma–/–</td>
<td>0/5</td>
<td>0.0 ± 0.0</td>
</tr>
</tbody>
</table>

*Recipient mice were injected intravenously with 5 × 10<sup>6</sup> encephalitogenic p35-55–specific T cells and monitored daily for clinical signs of EAE.*
addition, there was no significant difference in usage of TCR Vβ 4, 5, 6, 8, 12, 13 and 14 between wild-type and H-2M–deficient mice. Only one difference in TCR Vβ usage by p35-55–specific T cells was observed: Vβ 17 was used by 5.3% of wild-type T cells and 10.2% of the H-2M–deficient T cells. Thus, our results demonstrate that inability to induce EAE in H-2M–deficient mice reflects a defect in APC function and not development of a repertoire of encephalitogenic T cells.

**Discussion**

Endocytic processing may be required at different stages in EAE pathogenesis. When exposed to native MOG outside the CNS, we have observed that processing by peripheral APCs is required for presentation and activation of p35-55–specific T cells. Transmigration across the blood brain barrier (BBB) does not require Ag presentation, as activated T cells, independent of their specificity, can enter the CNS (3, 4). However, recognition of CNS autoantigen is considered a prerequisite for the initiation of T cell–mediated CNS inflammation (3, 4, 28). How then are pathogenic epitopes of CNS autoantigen initially exposed to encephalitogenic Th1 cells? Is endocytic processing required at this stage? Processing independent presentation of CNS autoantigen has been described but is considered rare (29). During turnover of myelin in normal CNS, it is possible that myelin peptides could become accessible to APCs (30). Activated Th1 cells entering the CNS secrete IFN-γ, which causes class II upregulation on resident nonprofessional APCs. Thus if a sufficient pool of myelin peptides exists, autopathogenic epitopes of myelin proteins could be presented to CD4+ Th1 cells by resident CNS APCs. In this situation, endocytic processing may not be required for initial CNS Ag presentation. Alternatively, in normal CNS, a sufficient pool of encephalitogenic protein fragments may not exist or may not be accessible to CNS APC. Through secretion of proinflammatory cytokines, activated Th1 cells that penetrate the BBB may elicit changes that stimulate APCs to upregulate their endocytic machinery, enabling generation and presentation of encephalitogenic peptide products in association with class II molecules. In this regard, microglia (31, 32) and IFN-γ-stimulated astrocytes express cathepsin (Cat) S (J.C. Patarroyo and S.S. Zamvil, unpublished observation), a lysosomal cysteine protease that participates in endocytic processing (33) and MBP degradation (34). Our data in this report are consistent with this latter possibility. Ii-deficient or H-2M–deficient recipient mice did not develop EAE after adoptive transfer of activated wild-type or H-2M–deficient donor encephalitogenic Th1 cells, a situation that obviates the requirement for Ag presentation and T cell activation outside the CNS (2). Thus, even with a repertoire of encephalitogenic T cells, mice containing APCs with defects in Ag processing could not propagate an encephalitogenic CNS inflammatory response, although APCs from these mice could present encephalitogenic MOG peptide to autopathogenic CD4+ T cells.

Alternative splicing of Ii mRNA leads to the formation of p31 and p41 Ii isoforms (9). Ii p41 contains an additional exon, 6b, which encodes a 64–amino acid sequence that is located in the luminal Ii domain (9). Although controversial (21, 22, 35), several studies have indicated that the less abundant Ii p41 facilitates enhanced class II–restricted presentation of certain Ag’s and antigenic peptide fragments (10, 19, 36). Ii p41 may protect a subset of antigenic epitopes from excessive degradation by inhibiting the lysosomal cysteine protease, Cat L (11, 12). In contrast, more recent evidence indicates that p41 facilitates the maturation of Cat L in selected cell types and prevents its premature destruction by surrounding endocytic proteases (37). Despite differences, p31 and p41 both participate in endocytic localization and class II compact dimer formation, and mice genetically manipulated to express either p31 or p41 support thymic selection of CD4+ T cells and the development of a normal T cell repertoire (21, 22, 35). Although some studies have indicated that Ii p41 may enhance Ag presentation (10–12, 19, 36), this possibility had not been examined in vivo in an organ-specific autoimmune disease. We observed that the incidence of EAE was similar in Ii p31, Ii p41, and wild-type mice, although the disease course was more severe in Ii p41 mice. We also observed that there was a lower threshold for IFN-γ secretion by activated p35-55–specific T cells from Ii p41 mice, which could contribute to the disparity in disease course. Although multiple discrete encephalitogenic determinants have been identified for other CNS autoantigens (38) and repertoire diversification can occur after the initial phase of EAE (5), p35-55 is the only encephalitogenic region of MOG identified in C57BL/6 mice (23, 27). As p41 may confer selective advantage in presentation, one possibility is that p41 facilitates presentation of a cryptic MOG determinant(s), which could participate in the encephalitogenic T cell response in the later phase of acute EAE. We are currently investigating this possibility.

Several proteases that participate in the endocytic pathway have been identified (33). Although it is not...
clear whether different proteases are utilized for presentation of distinct T cell epitopes, strategies that alter expression of individual proteases may be useful for modifying Ag presentation and T cell activation in clinical diseases (33). In this regard, it was reported that APCs from mice deficient in Cat S have defects in Ag processing and were less susceptible to collagen-induced arthritis, an autoimmune disease mediated by pathogenic Th1 cells (39). In vivo administration of a cysteine protease inhibitor was also effective in preventing Ag presentation to T cells in a model of pulmonary hypersensitivity (40). Similarly, reagents that selectively interfere with the endocytic pathway may be applicable to treatment of CNS autoimmune inflammatory conditions, such as EAE or MS.

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