

MHC class II–dependent B cell APC function is required for induction of CNS autoimmunity independent of myelin-specific antibodies

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Whether B cells serve as antigen-presenting cells (APCs) for activation of pathogenic T cells in the multiple sclerosis model experimental autoimmune encephalomyelitis (EAE) is unclear. To evaluate their role as APCs, we engineered mice selectively deficient in MHC II on B cells (B–MHC II^{−/−}), and to distinguish this function from antibody production, we created transgenic (Tg) mice that express the myelin oligodendrocyte glycoprotein (MOG)–specific B cell receptor (BCR; IgH^{MOG-mem}) but cannot secrete antibodies. B–MHC II^{−/−} mice were resistant to EAE induced by recombinant human MOG (rhMOG), a T cell– and B cell–dependent autoantigen, and exhibited diminished Th1 and Th17 responses, suggesting a role for B cell APC function. In comparison, selective B cell IL–6 deficiency reduced EAE susceptibility and Th17 responses alone. Administration of MOG-specific antibodies only partially restored EAE susceptibility in B–MHC II^{−/−} mice. In the absence of antibodies, IgH^{MOG-mem} mice, but not mice expressing a BCR of irrelevant specificity, were fully susceptible to acute rhMOG-induced EAE, also demonstrating the importance of BCR specificity. Spontaneous opticospinal EAE and meningeal follicle-like structures were observed in IgH^{MOG-mem} mice crossed with MOG-specific TCR Tg mice. Thus, B cells provide a critical cellular function in pathogenesis of central nervous system autoimmunity independent of their humoral involvement, findings which may be relevant to B cell–targeted therapies.

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Abbreviations used: Ag, antigen; CNS, central nervous system; EAE, experimental autoimmune encephalomyelitis; LFB, Luxol fast blue; MACS, magnetic-activated cell sorting; MOG, myelin oligodendrocyte glycoprotein; MS, multiple sclerosis; Mtb, *Mycobacterium tuberculosis*; NP, nitrophenyl; OSE, opticospinal EAE; rhMOG, recombinant human MOG; rmMOG, recombinant mouse MOG; SPMS, secondary progressive MS; Tfh cell, T follicular helper cell; Tg, transgenic.

Evidence supports roles for B cells and antibodies in the pathogenesis of multiple sclerosis (MS), a central nervous system (CNS) inflammatory demyelinating disease (von Büdingen

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et al., 2011). B cells and plasma cells are frequently observed in active inflammatory MS lesions, and myelin-specific antibodies have been detected in areas of CNS demyelination (Meinl et al., 2006). Identification of oligoclonal antibodies in cerebrospinal fluid is also used to confirm the diagnosis of MS (Blennow et al.,

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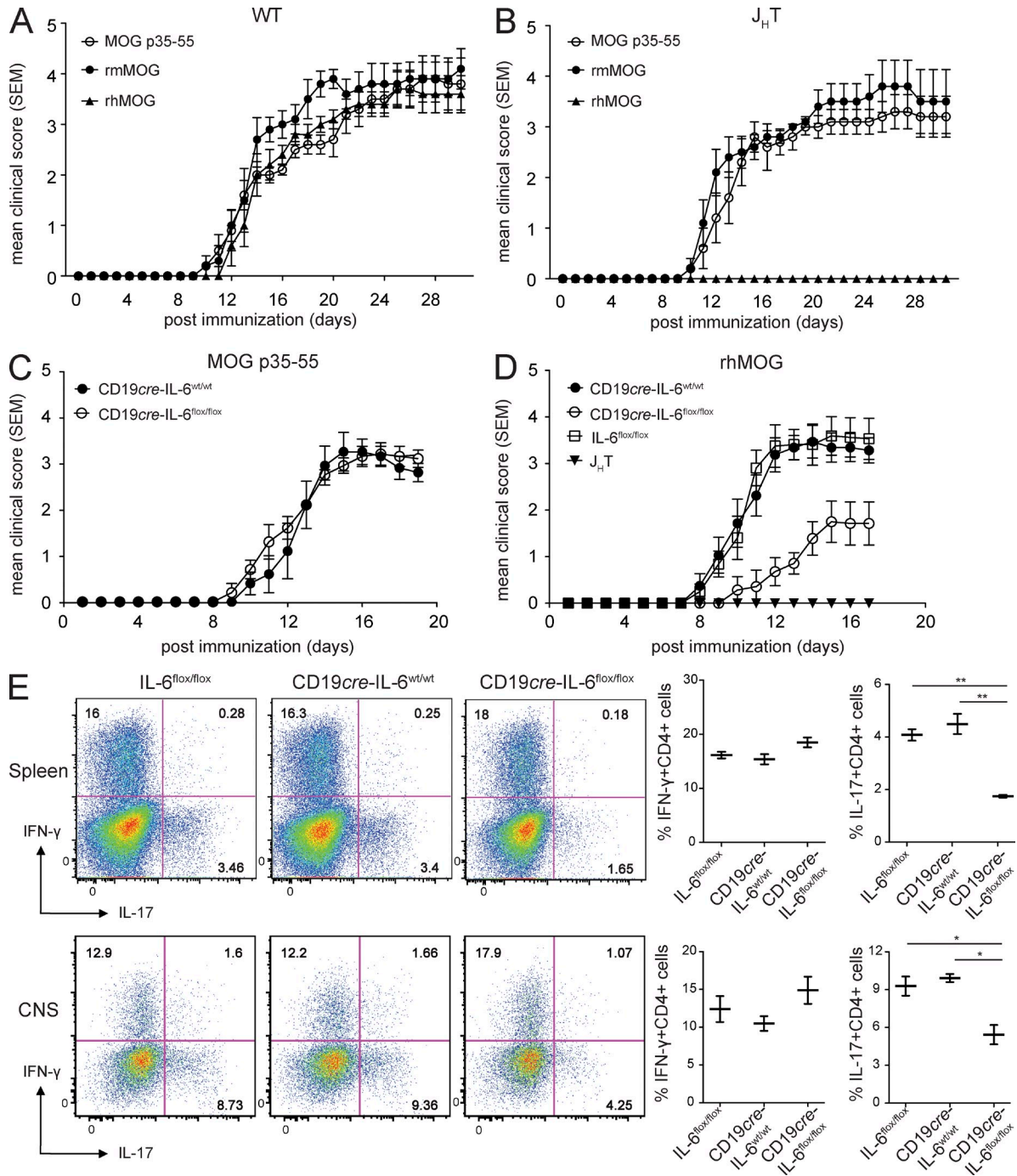


Figure 1. B cell IL-6 production was required for induction of EAE by rhMOG. (A and B) EAE was induced in C57BL/6J WT (A) or B cell-deficient $J_H T$ mice (B) by injecting MOG p35-55 (50 μ g s.c.), rmMOG (100 μ g s.c.), or rhMOG (100 μ g s.c.) emulsified in CFA along with heat-killed Mtb (200 μ g) and *B. pertussis* toxin (200 ng i.v.) on days 0 and 2, and survival was monitored. Results shown are representative of five independent experiments of $n = 5$ mice/group. (C and D) EAE was induced in mice harboring IL-6-deficient B cells (CD19cre-IL-6^{flox/flox}) or WT B cells (CD19cre-IL-6^{wt/wt} or IL-6^{flox/flox}) or mice lacking B cells ($J_H T$) with MOG p35-55 or rhMOG as in A, and survival was monitored. (E) Lymphocytes were isolated from the spleen and CNS of the indicated mouse strains after disease onset and stimulated with PMA + ionomycin for 6 h, and IL-17 and IFN- γ production was assessed by flow cytometry. Flow cytometry plots from a representative mouse are shown (left), and graphs (right) show quantitative data from five to six mice/group. Results are representative of two independent experiments ($n = 5$ mice/group [C]; $n = 7-8$ mice/group [D]). For all EAE experiments, mean disease score \pm SEM is shown. For all experiments: *, $P < 0.05$; **, $P < 0.01$ by Mann-Whitney *U* test.

Table 1. Evaluation of clinical signs and histological EAE in WT, B cell-deficient ($J_{H}T$), B-MHC II^{+/+}, and B-MHC II^{-/-} mice

Mice and Ag	Incidence	Onset	Mean maximal clinical score	Cumulative score	Mean number of foci (\pm SEM) ^a		
					Meninges	Parenchyma	Total
WT							
MOG p35–55	5/5	11.8 (\pm 0.6)	3.5 (\pm 0.5)	295.0	126 (\pm 69)	137 (\pm 69)	263 (\pm 138)
rmMOG	5/5	12.0 (\pm 0.6)	4.0 (\pm 0.3)	336.5	122 (\pm 42)	149 (\pm 45)	271 (\pm 87)
rhMOG	5/5	12.8 (\pm 0.5)	3.5 (\pm 0.4)	291.0	154 (\pm 47)	150 (\pm 10)	304 (\pm 57)
$J_{H}T$							
MOG p35–55	5/5	11.8 (\pm 0.7)	3.5 (\pm 0.3)	288.0	98 (\pm 25)	128 (\pm 55)	226 (\pm 80)
rmMOG	5/5	11.2 (\pm 0.4)	3.9 (\pm 0.5)	325.0	103 (\pm 18)	125 (\pm 24)	228 (\pm 42)
rhMOG	0/5	-	0.0 (\pm 0.0)	0.0	0.0 (\pm 0.0)	0.0 (\pm 0.0)	0.0 (\pm 0.0)
B-MHC II^{+/+}							
MOG p35–55	5/5	8.8 (\pm 1.3)	3.2 (\pm 0.3)	113.5	178 (\pm 13)	147 (\pm 8)	325 (\pm 21)
rmMOG	5/5	8.4 (\pm 0.5)	3.4 (\pm 0.2)	124.0	151 (\pm 43)	131 (\pm 19)	282 (\pm 62)
rhMOG	5/5	8.2 (\pm 0.8)	3.7 (\pm 0.2)	136.5	109 (\pm 17)	148 (\pm 27)	257 (\pm 34)
B-MHC II^{-/-}							
MOG p35–55	5/5	10.2 (\pm 0.8)	2.9 (\pm 0.2)	95.0	142 (\pm 22)	177 (\pm 43)	319 (\pm 65)
rmMOG	5/5	8.8 (\pm 0.4)	2.8 (\pm 0.1)	95.5	107 (\pm 33)	87 (\pm 7)	194 (\pm 40)
rhMOG	0/5	-	0.0 (\pm 0.0)	0.0	0.0 (\pm 0.0)	0.0 (\pm 0.0)	0.0 (\pm 0.0)

Onset indicates mean day of disease onset (paralytic EAE) \pm SEM. Clinical score shows mean maximal score of paralytic EAE of diseased mice \pm SEM. Cumulative score shows sum of the daily scores of all animals from day 0 until the end of the experiment. Results are representative of five separate experiments (five mice/group/experiment).

^aThe infiltration in the brain and spinal cord is quantified as averaged number of meningeal and parenchymal inflammatory foci (>10 clustered inflammatory cells) per CNS tissue section (five mice/group).

1994). Further, it has been suggested that the meningeal B cell follicles detected in progressive MS may contribute to detrimental humoral immunity (Magliozzi et al., 2007). Although it was the emphasis on humoral autoimmunity that provided the impetus to test B cell depletion in MS, the clinical benefit observed in recent clinical MS trials that tested anti-CD20 agents was not associated with reduction in serum or cerebrospinal fluid Ig titers, and oligoclonal antibodies were unchanged (Cross et al., 2006; Martin et al., 2009). Besides serving as the source for antibody-secreting plasma cells, B cells express MHC class II (MHC II) molecules constitutively, and via up-regulation of co-stimulatory molecules, they can participate as APCs for activation of antigen (Ag)-specific T cells (Lanzavecchia, 1985; van der Veen et al., 1992; Constant et al., 1995b). In this regard, reduction of proinflammatory Th17 cells has been observed in CD20-mediated B cell depletion in MS (Bar-Or et al., 2010). Together, these observations suggest that, independent of their potential humoral participation, there is prominent cellular involvement of B cells in MS pathogenesis.

Distinguishing the separate cellular and humoral contributions of B cells in the pathogenesis of MS, and in its model experimental autoimmune encephalomyelitis (EAE), has been challenging. As myelin-specific T cells are essential for initiation of CNS inflammation and clinical manifestations, EAE is T cell dependent (Zamvil and Steinman, 1990). B cells are not required in many EAE models, in particular those that are induced by encephalitogenic peptides of myelin proteins. In this regard, immunization with myelin oligodendrocyte

glycoprotein (MOG) peptide 35–55 (p35–55) induces EAE in B cell-deficient mice and does not elicit strong humoral responses in WT mice (Fillatreau et al., 2002; Lyons et al., 2002). In contrast, immunization of WT mice with recombinant human MOG (rhMOG) protein induces B cell activation and production of anti-MOG antibodies, which are pathogenic only in association with T cell-mediated CNS inflammation (Lyons et al., 2002; Oliver et al., 2003; Marta et al., 2005). As B cell-deficient mice are resistant to EAE induced by immunization with rhMOG, this model is considered both T cell and B cell dependent. Therefore, this EAE model is well suited to evaluate the dual cellular and humoral roles of B cells in CNS autoimmunity. It is recognized that B cells are capable of processing native Ags and are very efficient APCs when they recognize the same Ag as the responding T cells (Constant et al., 1995a,b). When MOG-specific BCR knockin (IgH^{MOG-ki}) mice (Litzenburger et al., 1998), which are capable of secreting all Ig isotypes, were crossed to MOG-specific TCR (TCR^{MOG}) mice (Bettelli et al., 2003), the progeny developed a spontaneous form of EAE that affected primarily the optic nerves and spinal cord (Bettelli et al., 2006a; Krishnamoorthy et al., 2006). Spontaneous opticospinal EAE (OSE) was associated with development of MOG-specific Th17 responses and signs of B cell activation, findings suggesting a role for B cell APC function (Bettelli et al., 2006a; Krishnamoorthy et al., 2006). Meningeal B cell aggregates containing germinal center-like structures resembling the ectopic B cell follicles identified in the leptomeninges in secondary progressive MS (SPMS) patients were also

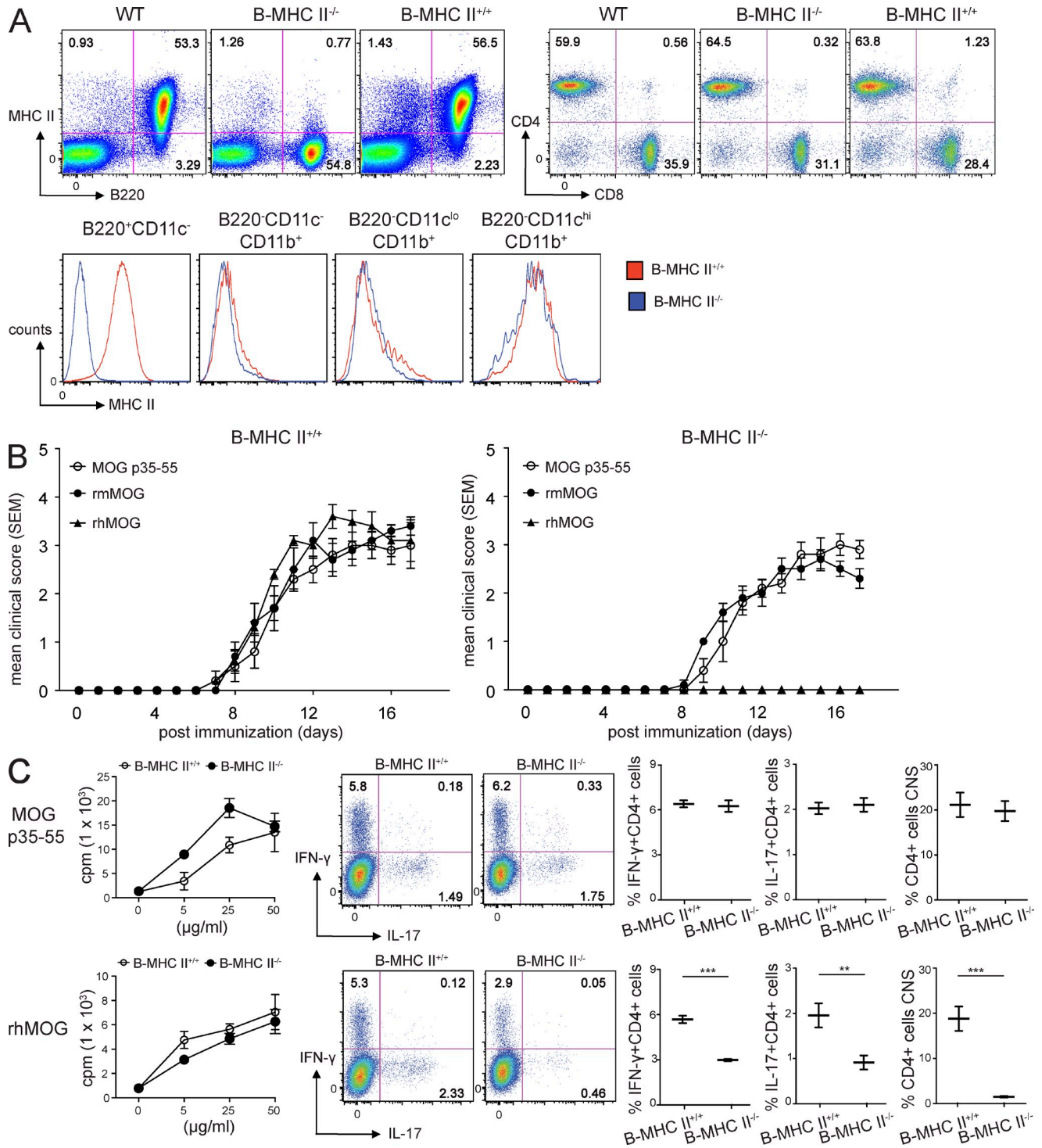


Figure 2. B cell MHC II expression promoted proinflammatory T cell cytokine production in rhMOG-induced EAE. Mixed BM chimera mice containing MHC II-deficient B cells were generated by reconstituting C57BL/6J (WT) mice with mixed BM from J_HT + MHC II-deficient mice (MHC II^{-/-}) mice. (A) B and T cell reconstitution of secondary lymphoid organs was evaluated by FACS analysis of spleen cells 6–8 wk after BM transplantation. Panels show frequencies of MHC II-expressing cells (left) and CD4⁺ and CD8⁺ cells (right) from B-MHC II^{-/-} and B-MHC II^{+/+} chimeras. FACS plots and histograms are representative of five independent experiments (*n* = 3 mice/group). (B) EAE was induced in B-MHC II^{+/+} or B-MHC II^{-/-} chimeras by injecting MOG p35–55 (50 μg s.c.), rmMOG (100 μg s.c.), or rhMOG (100 μg s.c.) emulsified in CFA along with heat-killed Mtb (200 μg) and *B. pertussis* toxin (200 ng i.v.) on days 0 and 2. Survival was monitored daily. Results shown are representative of five independent experiments of *n* = 5 mice/group. Mean disease score ± SEM is shown. (C) Analyses were performed 14 d after immunization. Spleen cells of B-MHC II^{+/+} and B-MHC II^{-/-} mice were cultured with the indicated concentrations of MOG p35–55 or rhMOG for 48 h, and T cell proliferation was measured 18 h later by incorporation of [³H]thymidine

observed in mice with spontaneous OSE (Bettelli et al., 2006a). Furthermore, IgG1 was elevated in IgH^{MOG-ki} × TCR^{MOG} mice (Bettelli et al., 2006a; Krishnamoorthy et al., 2006). Thus, similar to studies of MOG protein-induced EAE, the role of B cell APC function was not clearly distinguished from humoral immunity in OSE.

The goal of this investigation was to distinguish the role of APC function by B cells from the production and participation of myelin-specific antibodies in EAE pathogenesis. To this end, we used novel murine models that permitted us to evaluate requirements for B cell MHC II expression, BCR specificity and myelin-specific antibodies. First, we created mice that were selectively MHC II deficient on B cells (B-MHC II^{-/-} mice). These B-MHC II^{-/-} mice were susceptible to MOG p35–55-induced EAE. However, in comparison with mice containing B cells that expressed MHC II (B-MHC II^{+/+}), which were susceptible to EAE induced by rhMOG, B-MHC II^{-/-} mice were completely resistant to clinical or histological EAE, and they exhibited reduced Th1 and Th17 responses. Thus, B cell MHC II expression was required for EAE induction, which provided support for B cell APC function. Furthermore, administration of pathogenic MOG-specific antibodies only partially restored EAE susceptibility. To study B cell APC function in the absence of any possible contribution of antibodies, we created transgenic (Tg) mice containing B cells that expressed cell surface MOG-specific BCR but could not secrete antibodies. These MOG-specific “membrane-only” Tg (IgH^{MOG-mem})/B cell-deficient (J_HT; IgH^{MOG-mem}/J_HT) mice were fully susceptible to acute EAE induced by rhMOG, although their chronic course was slightly milder than that in either WT or IgH^{MOG-ki} mice. Spontaneous OSE was observed in the absence of MOG-specific antibodies when IgH^{MOG-mem}/J_HT mice were crossed to MOG-specific TCR (TCR^{MOG})/B cell-deficient (J_HT; TCR^{MOG}/J_HT) mice. In comparison with OSE in IgH^{MOG-ki} × TCR^{MOG} mice, the onset of OSE was delayed and severity was less in IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice. Our results demonstrate that in models that require participation of B cells, their cellular function is necessary and sufficient for induction of CNS autoimmunity.

RESULTS

B cell IL-6 production promotes proinflammatory T cell polarization in rhMOG-induced EAE

Immunization with rhMOG induces EAE in C57BL/6J WT mice (Fig. 1 A and Table 1), but not in B cell-deficient (J_HT) mice (Fig. 1 B and Table 1) or in B cell-depleted mice (not depicted). In contrast, MOG p35–55 and recombinant mouse MOG (rmMOG) can induce EAE in either WT mice (Fig. 1 A) or B cell-deficient mice (Fig. 1 B; Hjelmström et al., 1998;

Fillatreu et al., 2002; Oliver et al., 2003). Thus, only rhMOG-induced EAE is considered B cell dependent (Lyons et al., 1999; Oliver et al., 2003; Marta et al., 2005).

When serving as APCs, B cells can produce proinflammatory cytokines that promote development and activation of encephalitogenic T cells (Weber et al., 2010). As IL-6 has a key role in Th17 differentiation (Okuda et al., 1999; Bettelli et al., 2006b), we first addressed whether selective IL-6 production by B cells is required in rhMOG-induced EAE. No difference in disease course was observed between mice containing selective B cell IL-6 deficiency (CD19^{cre}-IL-6^{flox/flox}) and control (CD19^{cre}-IL-6^{wt/wt}) mice that were immunized with MOG p35–55 (Fig. 1 C). In contrast, when immunized with rhMOG, mice containing IL-6-deficient B cells developed less severe EAE in comparison with control (CD19^{cre}-IL-6^{wt/wt} and IL-6^{flox/flox}) mice (Fig. 1 D) and exhibited reduced IL-17 production in both the periphery and the CNS (Fig. 1 E). These results confirmed the importance of B cell IL-6 production in a model of B cell-dependent CNS autoimmunity.

B cell MHC II expression is required for rhMOG-induced EAE

Although cytokine production by accessory cells can promote proinflammatory T cell polarization, expression of MHC II on APCs is considered an obligate requirement for presentation and activation of Ag-specific CD4⁺ T cells (Schwartz, 1985). Thus, to determine the role of Ag presentation by B cells in EAE, we constructed mice containing MHC II-deficient B cells. Irradiated WT mice were reconstituted with mixed BM cells in a ratio of 80% J_HT and 20% MHC II^{-/-} (B-MHC II^{-/-}) or WT (B-MHC II^{+/+}) BM cells. Thus, in B-MHC II^{-/-} mice, MHC II deficiency was restricted to B cells, whereas other APCs, including DCs and monocytes, expressed MHC II (Fig. 2 A). The frequencies of B cells and CD4⁺ and CD8⁺ T cells were equivalent in BM chimeras containing either MHC II^{-/-} or MHC II^{+/+} B cells (Fig. 2 A). As shown in Fig. 2 B, BM chimera mice containing MHC II^{+/+} B cells developed EAE in response to immunization with rhMOG. In contrast, BM chimera mice containing MHC II-deficient B cells were resistant to clinical or histological EAE induction by rhMOG (Fig. 2 B and Table 1), although, like B cell-deficient J_HT or WT mice, BM chimera mice selectively deficient of MHC II on B cells did develop EAE in response to MOG p35–55 or rmMOG. Thus, EAE induction by rhMOG requires B cell MHC II expression.

Next, we examined how deficiency of MHC II on B cells influenced T cell activation and proinflammatory polarization in EAE induced by MOG p35–55 or rhMOG. Selective B cell MHC II deficiency did not influence proliferative responses to either Ag (Fig. 2 C), presumably because MHC

(cpm; left). Data are presented as means of triplicate values ± SD and are representative of at least three independent experiments. Single-cell suspensions from spleens and CNS of the indicated mice were stimulated with PMA + ionomycin for 6 h, and frequencies of IL-17⁺ CD4⁺ Th17 and IFN-γ⁺ CD4⁺ Th1 cells were assessed by flow cytometry. Representative FACS plots and quantification (mean ± SD) of *n* = 5 mice/group are shown (middle and right). For all experiments: **, *P* < 0.01; ***, *P* < 0.001 by Mann-Whitney *U* test.

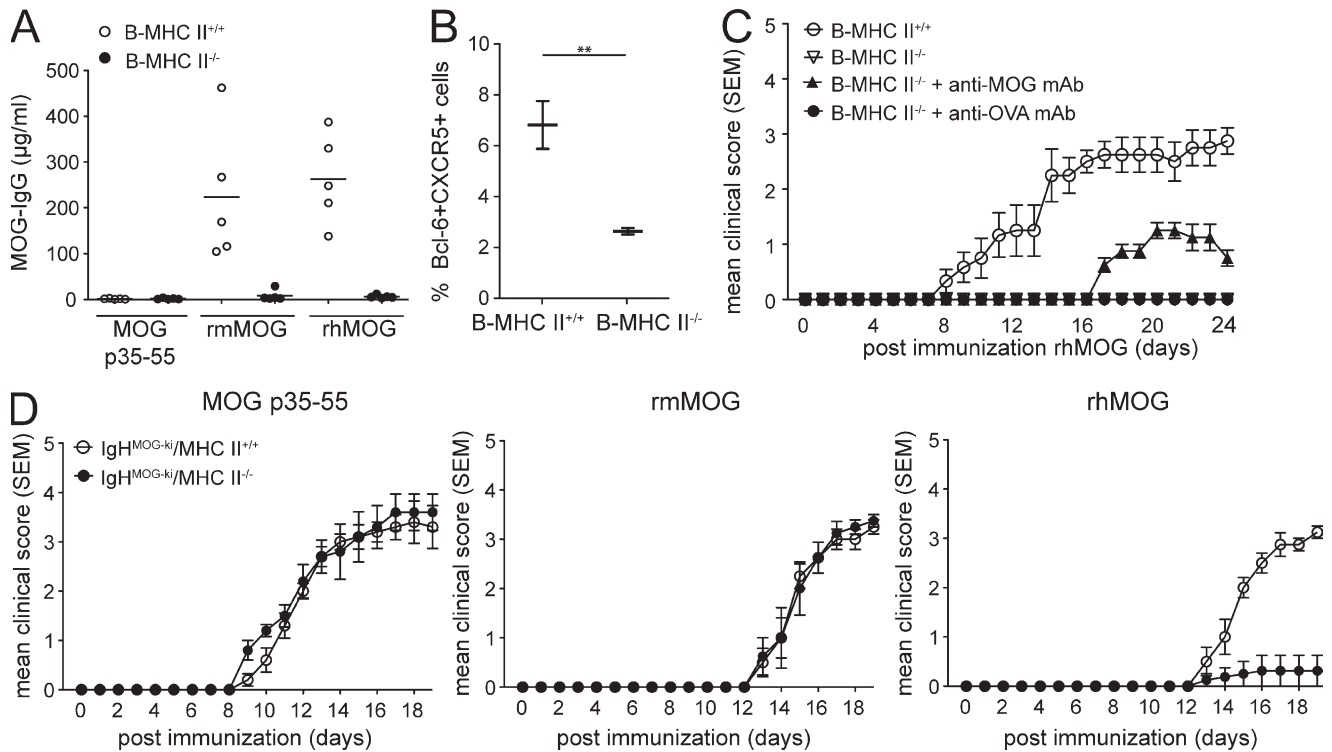


Figure 3. MOG-specific antibodies partially restored susceptibility to rhMOG-induced EAE in B-MHC II^{-/-} mice. (A) Sera were obtained from B-MHC II^{-/-} and B-MHC II^{+/+} mice on day 14 after immunization with MOG p35-55, rmMOG, or rhMOG. Total anti-MOG IgG levels ($\mu\text{g/ml}$) determined by ELISA (1:1,000 dilution) are shown ($n = 5$ mice/group). Horizontal bars indicate mean. (B) Numbers indicate the percentages of splenic Bcl-6⁺CXCR5⁺ Tfh cells (pregated on CD4⁺CD44⁺ICOS⁺PD-1⁺ T cells) evaluated by intracellular FACS staining 14 d after EAE induction with rhMOG. (C) B-MHC II^{-/-} and B-MHC II^{+/+} mice were immunized with rhMOG ($n = 4-6$ /mice/group), and clinical scores were monitored at the indicated days after immunization. When B-MHC II^{+/+} mice showed EAE signs (score ~ 2.0), B-MHC II^{-/-} mice received i.p. injections of 150 μg of either anti-MOG 8-18C5 mAb (closed triangles) or an equal amount of anti-OVA 1B7.11 mAb (closed circles) three times at 48-h intervals. Results are representative of two independent experiments. In one experiment, sera from IgH^{MOG-ki} mice and non-Tg littermate mice were used in lieu of 8-18C5 and 1B7.11 mAbs. (D) BM chimera mice with MHC II deficiency restricted to MOG-specific B cells from IgH^{MOG-ki} mice were generated by reconstituting C57BL/6J mice with mixed BM from J_HT + IgH^{MOG-ki} mice or J_HT + IgH^{MOG-ki} mice backcrossed onto the MHC II^{-/-} background. Chimera mice with MOG-specific MHC II-deficient B cells (IgH^{MOG-ki}/MHC II^{-/-}) or MOG-specific MHC II-competent B cells (IgH^{MOG-ki}/MHC II^{+/+}) were immunized with MOG p35-55, rmMOG, or rhMOG ($n = 5$ mice/group), and clinical scores were monitored at the indicated days after immunization. Data shown are representative of four independent experiments. Mean disease score \pm SEM is shown for all experiments displayed. For all experiments: **, $P < 0.01$ by Mann-Whitney U test.

II-competent myeloid cells also can process and present Ags. The contribution of MHC II expression by B cells in the development of pathogenic cells *in vivo* was also evaluated. No significant differences were observed in peripheral Th1 or Th17 responses in MOG p35-55-immunized BM chimera mice containing either MHC II^{+/+} or MHC II^{-/-} B cells. Also, there were no differences in the frequency of CNS-infiltrating T cells or the number of CNS infiltrates in these mice (Fig. 2 C and Table 1). In contrast, in response to immunization with rhMOG, the frequency of splenic CD4⁺ T cells expressing IFN- γ and IL-17A was significantly decreased in B-MHC II^{-/-} chimera mice. Furthermore, CNS inflammation was not detected in mice completely lacking MHC II only on B cells. Therefore, MHC II expression on B cells is critical for pathogenic T cell responses in rhMOG-induced EAE.

MOG-reactive antibodies partially restore EAE susceptibility to rhMOG in B-MHC II^{-/-} mice

Antibodies specific for rhMOG are known to be pathogenic (Marta et al., 2005). Although MHC II expression is required for B cell APC function, data indicate that MHC II is also necessary for germinal center formation and subsequent formation of long-lived plasma cells in T cell-dependent responses (Shimoda et al., 2006). Thus, we analyzed production of MOG-specific antibodies in our B-MHC II^{-/-} mice. Consistent with previous findings, MOG-specific antibodies were produced in control (B-MHC II^{+/+}) mice in response to rhMOG and rmMOG, but not to MOG p35-55 (Weber et al., 2010). In contrast, anti-MOG IgG was not detected in BM chimera mice containing MHC II-deficient B cells (Fig. 3 A). Interestingly, these data are consistent with the initial work that demonstrated the importance of MHC II in

Table 2. Passive transfer of MOG-specific antibodies partially restored EAE susceptibility in B-MHC II^{-/-} mice

Ag	Incidence	Onset	Mean maximal clinical score	Mean clinical score day 12	Mean clinical score day 18	Mean clinical score day 24	Cumulative score
B-MHC II^{+/+}							
rhMOG	6/6	15.0 (±1.2)	2.5 (±0.4)	1.3 (±0.5)	2.6 (±0.3)	2.9 (±0.2)	150.0
B-MHC II^{-/-}							
rhMOG	0/4	-	0.0 (±0.0)	0.0 (±0.0)	0.0 (±0.0)	0.0 (±0.0)	0.0
+anti-MOG mAb	4/4	3.0 (±0.0) p.i.	1.0 (±0.1)	0.0 (±0.0)	0.9 (±0.1)	0.9 (±0.1)	32.0
+anti-OVA mAb	0/4	-	0.0 (±0.0)	0.0 (±0.0)	0.0 (±0.0)	0.0 (±0.0)	0.0

p.i., post injection. Onset indicates mean day of disease onset (paralytic EAE) ± SEM. Clinical score shows mean maximal score of paralytic EAE of diseased mice ± SEM. Mean clinical score shows mean score of paralytic EAE of disease mice ± SEM. Cumulative score shows sum of the daily scores of all animals from day 0 until the end of the experiment.

humoral immunity (McDevitt et al., 1972). Absence of MHC II-expressing B cells was also associated with reduction of T follicular helper cells (Tfh cells; Deenick et al., 2010), a specialized T cell subset which provides help for B cell maturation and isotype switching (Fig. 3 B; Vinuesa et al., 2005; Deenick et al., 2010). As B-MHC II^{-/-} mice did not produce anti-MOG IgG, we tested whether antibodies could restore EAE susceptibility after immunization with rhMOG. Whereas transfer of isotype control OVA-specific mAb had no effect on EAE in B-MHC II^{-/-} mice immunized with rhMOG (Fig. 3 C and Table 2), administration of pathogenic anti-MOG mAb, 8-18C5, led to development of mild clinical disease, but did not restore EAE severity to the level observed in B-MHC II^{+/+} mice. Together, these results emphasize that the humoral contribution of B cells is insufficient to compensate for the lack of MHC II-dependent cellular function in MOG protein-induced EAE.

Besides serving as APCs or a source of autoantibodies, data indicate that through their cell surface-specific BCRs, B cells can transfer their target Ag to other professional APCs, a process which may be MHC II independent (Harvey et al., 2007). Thus, we addressed whether MOG-specific B cells contribute to MOG protein-induced EAE independent of their expression of MHC II molecules. BM chimera mice containing only IgH^{MOG-ki} B cells that expressed MHC II (IgH^{MOG-ki}/MHC II^{+/+}) were susceptible to EAE induced by MOG p35–55, rmMOG, or rhMOG (Fig. 3 D). In contrast, BM chimera mice containing MHC II-deficient IgH^{MOG-ki} B cells (IgH^{MOG-ki}/MHC II^{-/-}) were susceptible to EAE induced by MOG p35–55 or rmMOG but not to EAE induced by rhMOG (Fig. 3 D). Therefore, these results do not support BCR-mediated Ag exchange in rhMOG-induced EAE.

Generation of IgH^{MOG-mem} Tg mice

In the experimental system we used thus far, disruption of cellular function (i.e., MHC II deficiency) of B cells was accompanied by a reduction in humoral autoimmunity. It was therefore important to generate a model that would permit examination of the role of MHC II-competent MOG-specific B cells without possible contribution of antibodies. Tg mice containing B cells that express membrane IgH only have been useful for studying APC function in systemic and

organ-specific autoimmune diseases (Chan et al., 1999; Wong et al., 2004). Thus, we created Tg mice containing B cells that express membrane MOG-specific Ig (IgH^{MOG-mem}) but are incapable of secreting Ig. We subcloned the rearranged heavy chain Ig gene from 8-18C5, the same gene used to create the IgH^{MOG-ki} mice, which are capable of secreting all Ig isotypes, into a construct containing the IgM^a constant region from which the exon encoding for secretion was excised (Fig. 4 A; see Materials and methods; Chan et al., 1999). Mice bearing the IgH^{MOG-mem} Tg were backcrossed onto the B cell-deficient J_HT strain (IgH^{MOG-mem}/J_HT; Chen et al., 1993) to prevent the development of B cells expressing endogenous Ig. More than 50% of Tg B cells in the founder line #3088, which was chosen for most of our subsequent experiments, bound MOG protein similarly to B cells in other founder lines and in IgH^{MOG-ki} mice (Fig. 4 B and not depicted). B cells in those Tg mice also expressed MHC II comparably with IgH^{MOG-ki} and WT mice (Fig. 4 C). B cells in IgH^{MOG-mem}/J_HT Tg mice expressed cell surface IgM only (not depicted). Comparable frequencies of CD4⁺, CD8⁺, naive, and effector/memory T cells were detected in IgH^{MOG-mem}/J_HT, WT, and IgH^{MOG-ki} mice (not depicted).

The absence of secreted antibody in IgH^{MOG-mem}/J_HT mice was confirmed in separate analyses of total and MOG-specific IgM, as well as total and MOG-specific IgG in naive, MOG p35–55-immunized, and MOG protein-immunized mice (Fig. 4, D and E). These antibodies were undetectable or observed only in trace amounts in IgH^{MOG-mem}/J_HT mice and were never significantly more than in B cell-deficient mice. In general, antibodies were 10²–10³ times lower than WT controls. In contrast, naive and immunized IgH^{MOG-ki} mice secreted much higher concentrations of MOG-specific IgM and IgG.

MOG-specific B cells, but not antibodies, are required for induction of B cell-dependent EAE

IgH^{MOG-mem}/J_HT mice were tested for susceptibility to EAE by immunization with MOG p35–55, rmMOG, and rhMOG. Onset, severity, and disease course in IgH^{MOG-mem}/J_HT, WT, and IgH^{MOG-ki} mice immunized with either MOG p35–55 or rmMOG were similar (Fig. 5 A and Table 3). Onset and initial clinical severity were also comparable when these mice were

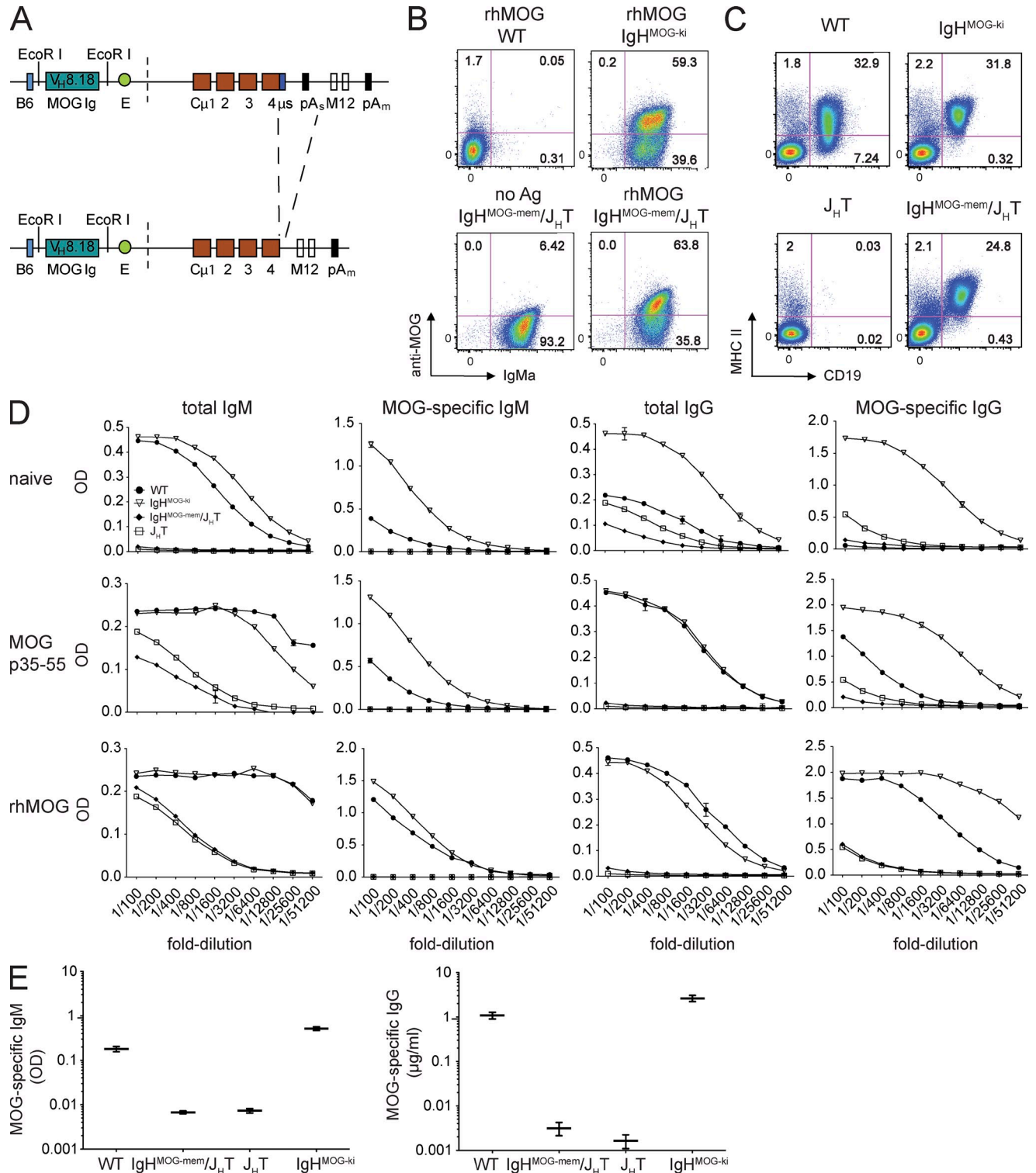


Figure 4. Generation and characterization of IgH^{MOG-mem} Tg mice. (A) Schematic representation of the construct used to generate IgH^{MOG-mem} Tg mice. The DNA directing secretion (μ s) and the transcription termination site (pA_s) have been deleted (Chan et al., 1999). V_H8.18 is the rearranged heavy chain V(D)J gene segment from the hybridoma 8-18C5 (Linnington et al., 1984), which was used to generate IgH^{MOG-ki} mice (Litzemberger et al., 1998). 8.18-C5 is known to recognize rhMOG₁₋₁₂₅ (Menge et al., 2007). E indicates the heavy chain intronic enhancer. Much of the switch region was deleted; a small segment of the residual switch region is indicated. (B) Spleen cells of WT, IgH^{MOG-ki}, J_HT, and IgH^{MOG-mem}/J_HT mice were analyzed for surface expression of IgM^a and binding of rhMOG by flow cytometry (gated on B220⁺ cells). Percentages of rhMOG-binding or -nonbinding IgM^a-positive B cells are indicated. (C) Spleen cells from WT, J_HT, IgH^{MOG-ki}, and IgH^{MOG-mem}/J_HT mice were analyzed by FACS for CD19 and MHC II expression. Data shown in B and C

immunized with rhMOG. However, the IgH^{MOG-mem}/J_HT mice appeared to have a slightly milder disease course after the acute stage, although this difference was not statistically significant. A similar EAE course was observed in another IgH^{MOG-mem}/J_HT Tg founder line after immunization with rhMOG (not depicted). There were no obvious differences in the extent of meningeal or parenchymal CNS infiltration among any of these mice with EAE (Fig. 5 B and Table 3).

Data from our IgH^{MOG-mem}/J_HT mice demonstrated that it was the cellular and not the humoral contribution that was required for development of acute disease in B cell-dependent EAE. We therefore examined the role of BCR specificity by comparing EAE induction in our IgH^{MOG-mem}/J_HT mice with Tg mice containing B cells that express a fixed cell surface IgH specific for nitrophenyl, an irrelevant Ag. Membrane-only nitrophenyl-specific IgH (B1-8; IgH^{B1-8-mem}/J_HT) mice differ from IgH^{MOG-mem}/J_HT mice only in the specificity of their rearranged heavy chain gene (Chan et al., 1999). In comparison with B cells from IgH^{MOG-mem}/J_HT, a much lower frequency of B cells from IgH^{B1-8-mem}/J_HT were capable of binding rhMOG (Fig. 5 C). As anticipated, B cells from IgH^{B1-8-mem}/J_HT mice could present MOG p35–55 to MOG-specific 2D2 cells but were inefficient in presentation of MOG protein (Fig. 5 D), which requires processing (Slavin et al., 2001; Bettelli et al., 2003). Like IgH^{MOG-mem}/J_HT mice, IgH^{B1-8-mem}/J_HT mice were susceptible to EAE induced by MOG p35–55 or rmMOG, independent of BCR specificity. However, IgH^{B1-8-mem}/J_HT mice developed a much milder EAE course than IgH^{MOG-mem}/J_HT mice after immunization with rhMOG (Fig. 5 E), which corresponded to a reduction in MOG-specific proinflammatory Th1 and Th17 cytokine responses in rhMOG-induced EAE (Fig. 5 F). These results clearly support an important role for CNS auto-Ag BCR specificity in B cell APC function in vivo.

MOG-specific T and B cells cooperate in development of spontaneous OSE in the absence of MOG-specific antibodies

It was previously observed when TCR^{MOG} mice were crossed with IgH^{MOG-ki} mice, the progeny developed spontaneous paralysis, which was associated with meningeal and parenchymal inflammatory lesions within the optic nerves and spinal cord (OSE; Bettelli et al., 2006a; Krishnamoorthy et al., 2006), a lesion topography reminiscent of neuromyelitis optica. OSE was also associated with elevation in MOG-specific IgG1, and as IgH^{MOG-ki} B cells can also serve as APCs, it was not clear whether it was their potential role as APCs or production of MOG-specific autoantibodies that promoted development of OSE. Thus, to evaluate the participation of MOG-specific B cells, independent of MOG-specific antibodies, we crossed

IgH^{MOG-mem}/J_HT with TCR^{MOG}/J_HT mice and evaluated their progeny for spontaneous development of OSE. 55% of IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice and 78% of IgH^{MOG-ki} × TCR^{MOG} developed paralysis spontaneously (Table 4). In contrast, spontaneous paralysis was not observed in IgH^{B1-8-mem}/J_HT × TCR^{MOG}/J_HT, TCR^{MOG}/J_HT, or IgH^{MOG-ki} mice. Onset of paralysis occurred earlier and the clinical severity was greater in IgH^{MOG-ki} × TCR^{MOG} mice than in IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice, possibly reflecting participation of MOG-specific antibodies in the manifestation of OSE. Interestingly, the numbers of inflammatory lesions were similar in both IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT and IgH^{MOG-ki} × TCR^{MOG} mice. Prominent meningeal and parenchymal inflammatory foci with demyelination and formation of meningeal lymphoid follicle-like aggregates (Fig. 6, A–E) were observed in IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice, as was described previously in IgH^{MOG-ki} × TCR^{MOG} mice with spontaneous OSE (Bettelli et al., 2006a), and resembled similar structures observed in SPMS (Magliozzi et al., 2007). Meningeal foci in IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice were composed predominantly of B cells that infiltrate the parenchyma (Fig. 6 C). Relatively fewer meningeal and parenchymal T cells were detected in these areas (Fig. 6 D). Follicle-like organization was evident using a reticulin stain (Fig. 6 E). Meningeal inflammation and follicle-like structures were not detected in clinically normal IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice (Fig. 6, F–J). Our observations indicate that the APC function of MOG-specific B cells is sufficient to cooperate with MOG-specific T cells in development of OSE and further suggest that in the context of CNS inflammation, anti-MOG antibodies augment phenotypic expression of this spontaneous disease.

DISCUSSION

Recent MS trials indicate that B cell-depleting agents have a profound beneficial effect, underscoring the importance of B cells in MS pathogenesis (Bar-Or et al., 2008; Hauser et al., 2008; Marriott and O'Connor, 2010; Kappos et al., 2011). Although it was the appreciation of the potential humoral contribution of B cells that inspired testing B cell depletion in MS, the clinical response was not associated with reduction in antibody levels (Bar-Or et al., 2008; Hauser et al., 2008; Marriott and O'Connor, 2010; Kappos et al., 2011). B cells afford multiple functions. Besides serving as a source of antibody-producing plasma cells, B cells can participate as APCs (Lanzavecchia, 1985; Constant et al., 1995a,b; Chan et al., 1999) in Ag transport (Phan et al., 2007) and as regulatory cells (Matsushita et al., 2008). Although MHC II is expressed constitutively on B cells, it is unknown whether B cell Ag

are representative of five independent experiments. (D) Sera from individual naive and immunized (MOG p35–55, rhMOG) WT, J_HT, IgH^{MOG-ki}, and IgH^{MOG-mem}/J_HT mice were obtained on day 14 after immunization, and serum Ig levels were calculated. Total and MOG-specific IgM and IgG titers are expressed as mean OD values ± SD from one of three representative experiments (performed in triplicate) from 1:2 serial dilutions. (E) Quantification of MOG-specific IgM and IgG (*n* = 6 mice/group) determined by ELISA are expressed as means of OD values or µg/ml ± SD of one of three representative experiments (performed in triplicate), respectively, and are shown on the logarithmic y axis.

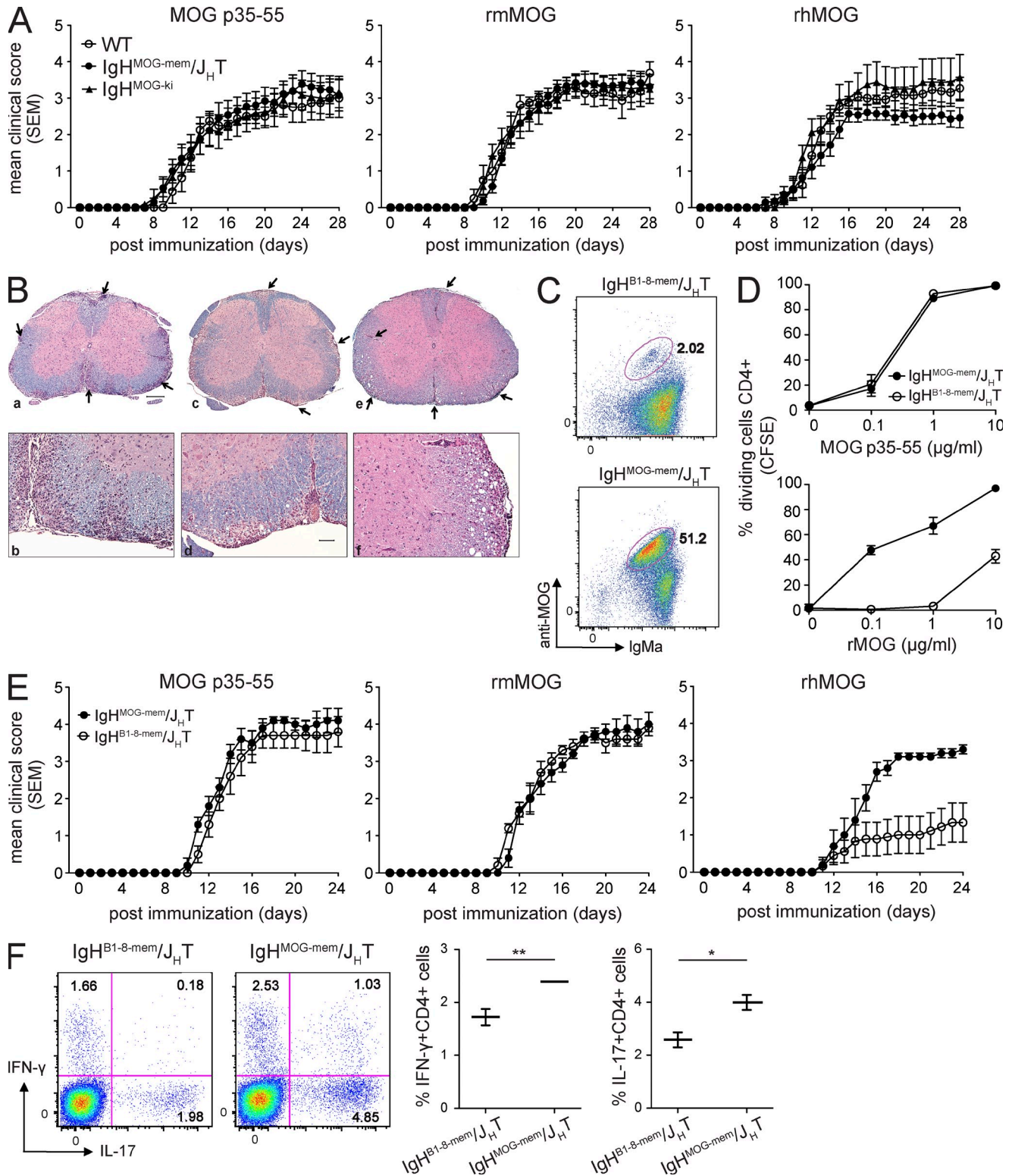


Figure 5. Myelin-specific BCR contributed to B cell APC function and susceptibility to CNS autoimmune disease independent of antibodies. (A) EAE was induced in WT, IgH^{MOG-mem}/J_HT, or IgH^{MOG-ki} mice by injecting MOG p35-55 (50 μg s.c.), rmMOG (100 μg s.c.), or rhMOG (100 μg s.c.) emulsified in CFA along with 200 μg heat-killed Mtb and *B. pertussis* toxin (200 ng i.v.) on days 0 and 2, and survival was monitored. Composite disease course of three experiments is shown (*n* = 3–6 mice per group/experiment). Mean disease score ± SEM is shown. (B) Spinal cord tissues from mice immunized with rhMOG were collected on day 28, fixed in neutral-buffered formalin, processed routinely, and embedded in paraffin (a, b: WT; c, d: IgH^{MOG-ki}; e, f: IgH^{MOG-mem}/J_HT). 5-μm sections were processed and stained with H&E (for assessment of inflammation) and LFB (for demyelination). Meningeal and parenchymal inflammatory/

Table 3. Evaluation of clinical signs and histological EAE in WT, IgH^{MOG-mem}/J_HT, and IgH^{MOG-ki} mice

Ag and mice	Incidence	Onset	Mean maximal clinical score	Mean number of foci (\pm SEM) ^a		
				Meninges	Parenchyma	Total
MOG p35–55						
WT	15/15	11.5 (\pm 0.4)	3.6 (\pm 0.3)	238 (\pm 53)	203 (\pm 39)	441 (\pm 92)
IgH ^{MOG-mem} /J _H T	18/18	11.4 (\pm 1.1)	3.7 (\pm 0.3)	117 (\pm 23)	117 (\pm 24)	234 (\pm 47)
IgH ^{MOG-ki}	15/15	11.6 (\pm 1.2)	3.8 (\pm 0.2)	118 (\pm 28)	138 (\pm 23)	256 (\pm 51)
rmMOG						
WT	15/15	11.3 (\pm 0.6)	3.9 (\pm 0.2)	160 (\pm 33)	199 (\pm 41)	339 (\pm 74)
IgH ^{MOG-mem} /J _H T	18/18	11.4 (\pm 0.2)	3.9 (\pm 0.2)	189 (\pm 18)	199 (\pm 12)	388 (\pm 30)
IgH ^{MOG-ki}	15/15	11.8 (\pm 0.8)	3.8 (\pm 0.3)	137 (\pm 16)	145 (\pm 24)	282 (\pm 30)
rhMOG						
WT	18/18	11.8 (\pm 0.5)	3.7 (\pm 0.2)	154 (\pm 47)	150 (\pm 10)	304 (\pm 57)
IgH ^{MOG-mem} /J _H T	18/18	11.8 (\pm 0.7)	3.0 (\pm 0.2)	118 (\pm 28)	139 (\pm 23)	257 (\pm 51)
IgH ^{MOG-ki}	17/18	10.4 (\pm 0.4)	4.2 (\pm 0.3)	146 (\pm 25)	139 (\pm 18)	285 (\pm 43)

Onset indicates mean day of disease onset (paralytic EAE) \pm SEM. Clinical score shows mean maximal score of paralytic EAE of diseased mice \pm SEM. Composite data from three independent experiments with five to six mice per group are indicated.

^aThe infiltration in the brain and spinal cord is quantified as averaged number of meningeal and parenchymal inflammatory foci (>10 clustered inflammatory cells) per CNS tissue section (five mice/group).

presentation may be required for activation of pathogenic CNS auto-Ag-specific T cells in vivo in EAE pathogenesis. In this investigation, we have created models that permitted us to evaluate and distinguish the role of B cells as APCs in vivo from their contribution to humoral CNS autoimmunity. Our results clearly demonstrate how B cell MHC II expression can be pivotal for activation of proinflammatory Th1 and Th17 cells in vivo and susceptibility to CNS autoimmunity. By generation of Tg mice that expressed cell surface myelin-specific BCR, we provided unequivocal evidence that B cell APC function, in the absence of humoral autoimmunity, is sufficient to drive proinflammatory T cell activation in vivo and EAE pathogenesis. Although BCR specificity for MOG protein promoted activation of encephalitogenic proinflammatory T cells, expression of this myelin-specific BCR was insufficient to drive proinflammatory T cell differentiation in the absence of B cell MHC II molecules (Fig. 3 D), providing further evidence for the MHC II-dependent APC participation of B cells. Elimination of B cell APC function can paralyze T cell activation and histological and clinical manifestations of CNS autoimmunity, findings which provide insight regarding B–T communication in MS pathogenesis and the influence of B cell depletion in MS therapy.

With greater appreciation for the role of B cells in MS, it is important to study MS models that require B cells for development of CNS autoimmunity. There are several different murine EAE models, and each one has merits that permit investigation of specific aspects involved in pathogenesis of CNS autoimmunity. With identification of encephalitogenic T cell epitopes within individual myelin Ags, most studies use those corresponding encephalitogenic peptides for EAE induction. In this regard, MOG p35–55 is frequently used in EAE studies. In general, B cells recognize conformational determinants created by longer contiguous or noncontiguous amino acid sequences within native proteins (Roelants, 1972). Whereas MOG p35–55 activates encephalitogenic T cells (Kuchroo et al., 2002), it does not efficiently activate B cells or promote MOG-specific antibody production (Weber et al., 2010). In contrast, MOG protein efficiently leads to both B cell activation and production of anti-MOG antibodies (Weber et al., 2010), and it is recognized that rhMOG, but not rodent MOG isoforms, induces EAE in a B cell-dependent manner (Marta et al., 2005). Interestingly, this model is not DC dependent, as mice depleted of this myeloid APC subset were susceptible to EAE induced by MOG protein (Isaksson et al., 2012). The B cell dependence of rhMOG-induced EAE is attributed to the presence of proline 42 (P42) in its sequence

demyelinating foci are indicated by arrows. Bars: (a, c, and e) 200 μ m; (b, d, and f) 50 μ m. (C) Splenic B cells (B220⁺) from IgH^{MOG-mem}/J_HT and IgH^{B1-8-mem}/J_HT mice were stained for cell surface IgM^a, and capability to bind MOG protein and then analyzed by flow cytometry. (D) Purified CD4⁺ T cells from TCR^{MOG} mice were stained with CFSE and cultured with purified B cells from IgH^{MOG-mem}/J_HT or IgH^{B1-8-mem}/J_HT mice in the presence of various concentrations of MOG p35–55 or rhMOG. T cell proliferation (CFSE dilution) was analyzed by FACS. Data are presented as means of triplicate values \pm SD and are representative of two independent experiments. (E) EAE was induced in IgH^{MOG-mem}/J_HT or IgH^{B1-8-mem}/J_HT mice ($n = 5$ mice/group) as in A. Survival was monitored daily. Clinical EAE scores are shown as mean \pm SEM. EAE data shown are representative of five independent experiments. (F) Phenotype of splenic T cells was evaluated by intracellular FACS staining for IL-17A and IFN- γ (gated on CD4⁺ T cells) 14 d after disease induction. Representative FACS plots (left) and quantification (mean \pm SD; right; $n = 5$ mice/group) are shown. For all experiments: *, $P < 0.05$; **, $P < 0.01$ by Mann–Whitney U test.

Table 4. IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice developed spontaneous EAE

Mice	Incidence	Onset	Mean maximal clinical score	Mean number of foci (±SEM) ^a		
				Meninges	Parenchyma	Total
	%					
IgH ^{MOG-mem} /J _H T × TCR ^{MOG} /J _H T	55 (21/38)	60 (±4)	2.9 (±0.2)	43 (±25)	55 (±18)	98 (±43)
IgH ^{B1-8-mem} /J _H T × TCR ^{MOG} /J _H T	0 (0/22)	n/a	0.0	0.0	0.0	0.0
TCR ^{MOG} /J _H T	0 (0/52)	n/a	0.0	0.0	0.0	0.0
IgH ^{MOG-mem} /J _H T	0 (0/56)	n/a	0.0	0.0	0.0	0.0
IgH ^{MOG-ki}	0 (0/45)	n/a	0.0	0.0	0.0	0.0
IgH ^{MOG-ki} × TCR ^{MOG}	78 (38/49)	38 (±2)	3.8 (±0.1)	60 (±14)	51 (±14)	111 (±24)

n/a, not applicable. Onset indicates mean day of disease onset (paralytic EAE) ± SEM. Clinical score shows mean maximal score of paralytic EAE of diseased mice ± SEM.

^aThe infiltration in the brain and spinal cord is quantified as averaged number of meningeal and parenchymal inflammatory foci (>10 clustered inflammatory cells) per CNS tissue section (five mice/group).

(Oliver et al., 2003; Marta et al., 2005), as substitution of serine, the corresponding residue in rodent MOG, with P42 in rodent MOG protein confers B cell-dependent induction of EAE (Oliver et al., 2003; Marta et al., 2005). Use of rhMOG-induced EAE also permitted us to evaluate the role of B cell cytokine production in expansion of proinflammatory T cells in vivo. Selective deficiency in MHC II-dependent B cell APC function was associated with significant reductions in both Th1 and Th17 cells. As it is recognized that B cells are a source of IL-6, a key Th17-polarizing cytokine (Bettelli et al., 2006b), we investigated how B cell production of IL-6 itself influenced EAE susceptibility. Although selective B cell IL-6 deficiency did not significantly alter the course of MOG p35–55-induced EAE, onset of rhMOG-induced EAE was delayed, severity was attenuated and the frequency of Th17 cells only was significantly reduced in B cell IL-6-deficient mice. Similar results were obtained in mixed BM chimera mice containing IL-6-deficient B cells (unpublished data). These results underscore the importance of IL-6 production by activated B cells in Th17 differentiation in vivo in rhMOG-induced EAE. Interestingly, a recent study evaluated B cell IL-6 production in MOG p35–55-induced EAE and observed that B cell IL-6 production was not required for initiation of EAE, but that the disease course was reduced in mice containing IL-6-deficient B cells (Barr et al., 2012). Thus, B cell IL-6 deficiency may have a variable effect when EAE is induced by MOG p35–55, but it is clear that the impact of B cell IL-6 production is greater in rhMOG-induced EAE. As rhMOG is currently the only known B cell-dependent encephalitogen, it may be advantageous to examine this myelin protein when evaluating the pathogenic role of activated myelin-specific B cells in EAE.

In this study, selective MHC II deficiency in B cells was evaluated using mixed BM chimera mice reconstituted with B cells from either WT mice or from MHC II-deficient mice, ensuring in the latter a complete absence of B cell MHC II expression and MHC II-restricted Ag presentation. As an alternative, we also examined selective B cell MHC II deficiency in CD19-targeted MHC II-deficient (CD19^{cre}-MHC II^{flox/flox}) mice (Shimoda et al., 2006). It is known that

~95% of B cells in CD19^{cre}-MHC II^{flox/flox} mice are MHC II deficient (Shimoda et al., 2006). Although a minority of B cells expressed MHC II molecules, CD19^{cre}-MHC II^{flox/flox} mice were also resistant to EAE induced with rhMOG (unpublished data). Thus, with both approaches, the results were consistent in demonstrating the in vivo requirement for B cell MHC II expression for EAE induced by this myelin protein. Although selective MHC II deficiency eliminated B cell APC function, it was also associated with a reduction in MOG-specific antibodies. One could contend that the absence of B cell MHC II expression simply eliminated the humoral response, which itself was primarily responsible for EAE pathogenesis. In our study, B cell MHC II deficiency resulted in a reduction in Tfh cells, a subset of CD4⁺ T cells that participate in germinal center formation, generation of memory B cells, and long-lived plasma cells (Crotty, 2011). In this regard, B cell APC function is a prerequisite for humoral immunity as Ag presentation by responding B cells is required for development and maintenance of Tfh, especially when Ag is limiting (Deenick et al., 2010). Nevertheless, it should be recognized that transfer of MOG-specific antibodies alone, or Tg-driven secretion of MOG-specific antibodies alone, does not lead to CNS inflammation and EAE (Linington et al., 1988; Genain et al., 1995; Litzemberger et al., 1998), and, as we have observed, transfer of pathogenic anti-MOG antibodies into B-MHC II^{-/-} mice that were immunized with rhMOG led to a partial restoration of EAE. Furthermore, despite MHC II expression on myeloid APCs, including DCs, there was a reduction in development of MOG-reactive Th17 cells in mice selectively deficient in B cell MHC II expression, implying direct B-T interaction in vivo.

IgH “membrane-only” BCR Tg mice can distinguish B cell Ag presentation in vivo from humoral immunity (Chan et al., 1999). Although not targeted to a defined self-Ag, IgH^{B1-8-mem} Tg mice have served to distinguish the contribution of B cells in Ag presentation from their role in the generation of antibodies in murine lupus and arthritis (Chan et al., 1999; O’Neill et al., 2005). To completely dissociate B lymphocyte cellular function from the potential contribution of humoral autoimmunity in EAE pathogenesis, we generated

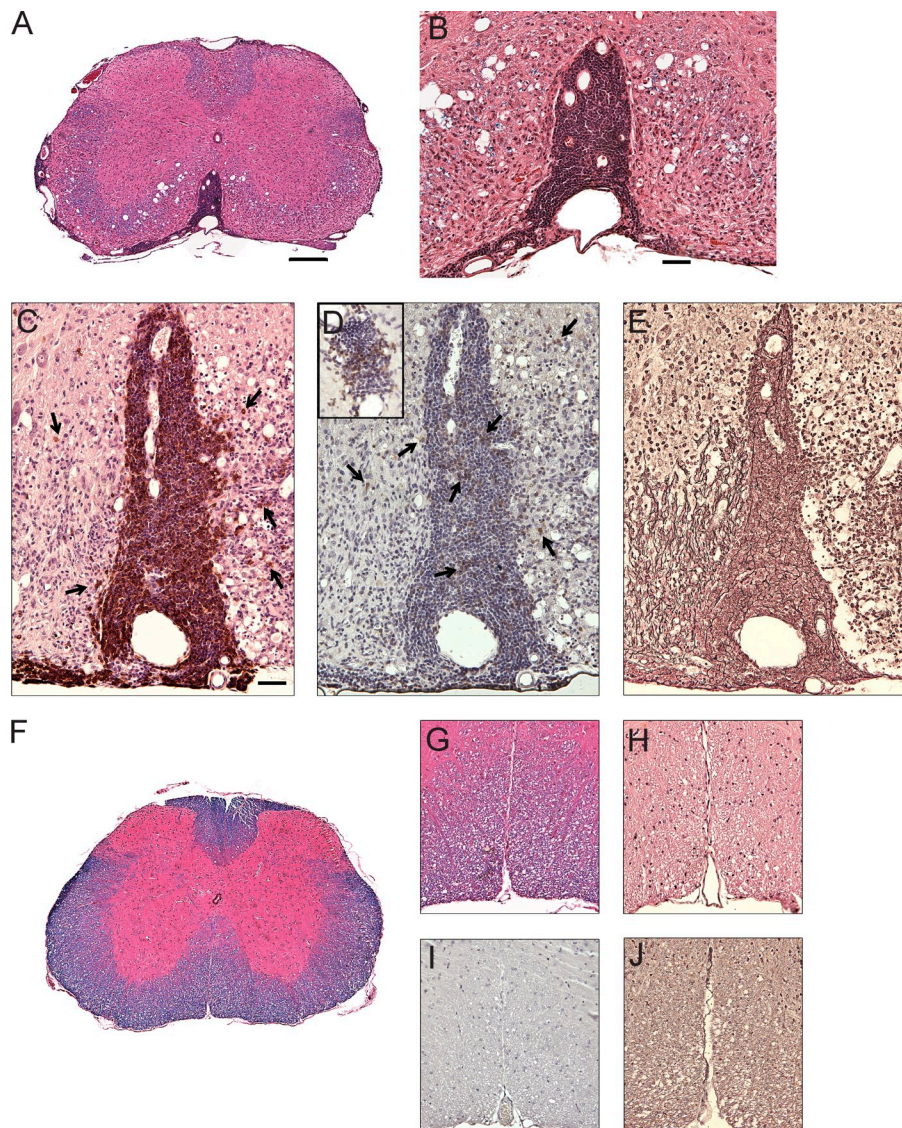


Figure 6. Meningeal B cell follicle-like aggregates were observed in $IgH^{MOG-mem}/J_H T \times TCR^{MOG}/J_H T$ mice that developed spontaneous OSE. (A–E) Spinal cords of $IgH^{MOG-mem}/J_H T \times TCR^{MOG}/J_H T$ mice that developed spontaneous OSE (A and B). (F–J) Spinal cord of a clinically normal $IgH^{MOG-mem}/J_H T \times TCR^{MOG}/J_H T$ mouse. (C and H) Immunostaining for B cells (CD45R). (C) Arrows indicate meningeal foci that contained B cells that infiltrated the parenchyma. (D and I) Immunostaining for T cells (CD3). (D) Arrows and inset show meningeal and parenchymal T cells. (E and J) Reticulin stain. (A, B, F, G) LFB/H&E; (C and H) H&E counterstain; (D and I) hematoxylin counterstain; (E and J) reticulin stain. Histological analysis of $IgH^{MOG-mem} \times 2D2/J_H T$ mice with and without spontaneous OSE is representative of three independent experiments with at least three mice per group. The entire CNS was sampled in a single slide for each mouse. Bars: (A and F) 200 μm ; (B and G–J) 50 μm ; (C–E) 70 μm ; (D, inset [same bar as in C]) 35 μm .

$IgH^{MOG-mem}/J_H T$ Tg mice. B cells from our $IgH^{MOG-mem}/J_H T$ mice efficiently presented MOG protein to MOG-specific 2D2 T cells and, despite the inability to secrete MOG-specific IgG or IgM, these mice were completely susceptible to acute clinical and histological EAE induced by rhMOG. Consistent with observations that B cells are more efficient APCs when they recognize the same target Ag as the responding T cells (Lanzavecchia, 1985; Constant et al., 1995a,b), we observed that B cells from $IgH^{MOG-mem}/J_H T$ mice presented rhMOG to 2D2 T cells and promoted proinflammatory Th1 and Th17 differentiation more efficiently than B cells from $IgH^{B1-8-mem}/J_H T$ mice, which bound rhMOG less effectively. Furthermore, in contrast to $IgH^{MOG-mem}/J_H T$ mice, $IgH^{B1-8-mem}/J_H T$ mice were less susceptible to EAE induction by rhMOG. Collectively, these in vitro and in vivo findings demonstrate how myelin-specific B cells can have a critical role in presentation to myelin-specific T cells in CNS autoimmunity. Although onset of acute EAE was similar

in $IgH^{MOG-mem}/J_H T$, IgH^{MOG-ki} , and WT mice, it was of particular interest that the chronic phase appeared less severe in $IgH^{MOG-mem}/J_H T$ mice than either IgH^{MOG-ki} or WT mice. Indeed, greater severity could reflect contribution of pathogenic MOG-specific antibodies elicited by immunization of IgH^{MOG-ki} or WT mice with rhMOG. However, given minor differences observed in the disease course, it may be experimentally challenging to dissect the contribution of anti-MOG antibodies in this model.

With the recent introduction of new agents for relapsing-remitting MS, the unmet need for successful therapy of progressive MS has become more evident (Fox et al., 2012; Brück et al., 2013). Understanding of the genetic and pathogenic mechanisms involved in relapsing-remitting MS may also be advancing more rapidly (Sawcer et al., 2011). Furthermore, it is recognized that there is a paucity of models for progressive MS (Amor et al., 1994; Bettelli et al., 2006a; Krishnamoorthy et al., 2006; Basso et al., 2008). Although models do not entirely

reflect spontaneous human disease, they can provide insight regarding specific pathogenic mechanisms. Meningeal B cell follicle-like structures have been observed in SPMS and in IgH^{MOG-ki} × TCR^{MOG} mice with OSE. IgH^{MOG-mem}/J_HT and IgH^{MOG-ki} mice are complementary; whereas IgH^{MOG-mem}/J_HT mice cannot produce any antibodies, IgH^{MOG-ki} mice can secrete all Ig isotypes. Here, by analysis of IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice, we observed it is the cellular participation of MOG-reactive B lymphocytes that is necessary and sufficient for the development of spontaneous OSE and formation of meningeal follicle-like structures. Clinical signs of OSE appeared less severe in IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice than in IgH^{MOG-ki} × TCR^{MOG} mice and the onset occurred later, suggesting that MOG-specific antibodies may augment the phenotypic expression. Thus, IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice should provide a foundation to evaluate which individual isotypes of MOG-specific antibodies cooperate with MOG-specific B and T cells in OSE. IgH^{MOG-mem} mice provide a powerful reagent to directly assess the contribution of select genes that may participate in B–T communication or development of OSE and formation of meningeal follicle-like structures without influence of soluble antibodies or potential antibody–Ag complexes. In summary, our observations in this study underscore the central importance of MHC II-dependent B cell Ag presentation in CNS autoimmunity. Use of IgH^{MOG-mem}/J_HT mice to study development of spontaneous OSE should provide further insight regarding peripheral and CNS involvement of myelin-specific B cells in MS progression.

MATERIALS AND METHODS

Mice. WT C57BL/6J CD45.1 mice as well as CD45.2 MHC II^{-/-} (B6.129S2-H2dAb1-Ea/J), IL-6^{-/-} (B6.129S2-Il6^{gm1Kopf}/J), and CD19^{cre/cre} (homozygous for cre cassette, B6.129P2(C)-Cd19^{tm1(cre)Cgn}/J) mice were purchased from the Jackson Laboratory. C57BL/6J B cell-deficient J_HT mice (B6.129P2-Igh-Jtm1Cgn/J; Gu et al., 1993) were provided by K. Rajewsky (Harvard University, Boston, MA), and C57BL/6J MOG p35–55-specific TCR Tg (2D2, TCR^{MOG}) mice (Bettelli et al., 2003) were provided by V.K. Kuchroo (Harvard University). C57BL/6J MOG-BCR knockin (IgH^{MOG-ki}, also referred to as Th) mice (Litzenburger et al., 1998) were provided by H. Wekerle. C57BL/6 IL-6^{flox/flox} mice generated by gene targeting technology (Quintana et al., 2013) were provided by J. Hidalgo. For selective deletion of floxed IL-6 gene in B lymphocytes, CD19^{cre/cre} mice were crossed with IL-6^{flox/flox} mice. Litters heterozygous for the cre cassette (CD19^{cre}-IL-6^{flox/flox}, CD19^{cre}-IL-6^{wt/wt}) were used for experiments. IgH^{B1-8} (IgH^{B1-8-mem}) mice (Chan et al., 1999) were backcrossed onto the C57BL/6J background and provided by J.-A. Lyons (University of Wisconsin, Milwaukee, WI) and A.H. Cross (Washington University School of Medicine in St. Louis, St. Louis, MO). IgH^{B1-8-mem} mice were crossed onto the J_HT background, and IgH^{MOG-ki} mice were crossed onto the MHC II^{-/-} background. Animals were housed in a specific pathogen-free barrier facility at the Parnassus Research Center of the University of California, San Francisco (UCSF). All breeding and experiments were reviewed and approved by the UCSF Institutional Animal Care and Use Committee.

Generation of IgH^{MOG-mem} and IgH^{MOG-mem} × TCR^{MOG} Tg mice. To generate IgH^{MOG-mem} mice, the genomic DNA fragment containing the rearranged heavy chain VDJ of the pathogenic MOG-specific Ig 8-18C5 mAb (Litzenburger et al., 1998) was inserted into the IgM⁺ constant region of a vector from which the secreted exon and polyadenylation site were excised

(IgM^{mem}), which permits expression of membrane-bound Ig only (Chan et al., 1999). Specifically, we excised the 3.6-kb fragment containing the rearranged heavy chain enhancer and rearranged VDJ gene segment of the pathogenic MOG-specific Ig by digestion with EcoRI. This fragment was then subcloned into the EcoRI sites in the IgM^{mem} only construct. Correct orientation was determined first by PCR and restriction analysis and then confirmed by DNA sequencing of the regions containing the integration sites. The IgH^{MOG-mem} construct was microinjected into C57BL/6J oocytes, which provided separate founders that were identified by PCR and confirmed by Southern blot. All founders contained B cells that expressed transgene-directed cell surface IgM⁺ and bound MOG. IgH^{MOG-mem} mice were crossed to C57BL/6J J_HT mice to generate IgH^{MOG-mem}/J_HT mice.

TCR^{MOG-mem} mice were crossed onto the J_HT background to generate TCR^{MOG}/J_HT mice. IgH^{MOG-mem}/J_HT and IgH^{B1-8-mem}/J_HT mice were intercrossed to TCR^{MOG}/J_HT mice to generate IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT and IgH^{B1-8-mem}/J_HT × TCR^{MOG}/J_HT mice. For comparison of IgH^{MOG-mem}/J_HT × TCR^{MOG}/J_HT mice, IgH^{MOG-ki} mice were crossed with TCR^{MOG} mice to generate IgH^{MOG-ki} × TCR^{MOG} mice (Bettelli et al., 2006a; Krishnamoorthy et al., 2006).

Generation of mixed BM chimera mice. 1,200 cGy γ -irradiation (Cs source) was administered to 6-wk-old recipient C57BL/6J mice. On the same day, recipients received 5×10^6 donor BM cells. All BM preparations were depleted of T cells by labeling with a biotinylated anti-Thy-1 (clone T24) before incubation with streptavidin-microbeads (Miltenyi Biotec) for negative selection via magnetic-activated cell sorting (MACS) column (Miltenyi Biotec). To restrict genetic MHC II deficiency to B cells, the BM inocula consisted of 80% BM from J_HT mice supplemented with 20% BM from MHC II-deficient mice. The control group received BM inocula consisting of 80% BM from J_HT mice supplemented with 20% BM from WT mice. In certain experiments, chimeras were generated with 80% BM from J_HT mice and 20% BM from either IgH^{MOG-ki}/MHC II^{-/-} or IgH^{MOG-ki} mice. BM chimeras were housed for at least 8 wk to reconstitute their peripheral lymphoid system before use in EAE experiments. FACS analysis confirmed hematopoietic reconstitution of sublethally irradiated C57BL/6J CD45.1 mice with CD45.2⁺ donor BM for expression of CD45.2 on peripheral blood mononuclear cells. Engraftment of donor CD45.2⁺ cells was $\geq 95\%$. The BM chimera containing MHC II-deficient B cells showed numbers of CD19⁺ splenocytes equivalent to numbers found in normal WT mice.

Ag. Mouse MOG p35–55 (MEVGWYRSPFSRVVHLYRNGK) was synthesized by Auspep. rmMOG was provided C.C.A. Bernard. rhMOG was provided by C.C.A. Bernard and D.E. Jenne. MOG proteins were synthesized, purified, and refolded as previously reported (Clements et al., 2003; Breithaupt et al., 2008).

Induction and assessment of EAE. C57BL/6J mice were injected s.c. with 50 μ g MOG p35–55 or 100 μ g rmMOG or 100 μ g rhMOG emulsified in CFA (DIFCO Laboratories) containing 200 μ g heat-killed *Mycobacterium tuberculosis* (Mtb) H37RA (DIFCO Laboratories) on day 0. Additionally, mice received i.v. 200 ng *Bordetella pertussis* toxin (List Biological Laboratories) in 0.2 ml PBS on days 0 and 2. Individual animals were observed daily, and clinical scores were assessed with a 0- to 5-point scoring system, as follows: 0 = no clinical disease, 1 = loss of tail tone only, 2 = mild monoparesis or paraparesis, 3 = severe paraparesis, 4 = paraplegia and/or quadriplegia, and 5 = moribund or death. Moribund mice were given disease severity scores of 5 and euthanized.

Generation of Ag-specific antibodies. IgH^{MOG-ki} mice were immunized with rmMOG (100 μ g/mouse). 14–16 d later, mice were bled via cardiac puncture. Blood was allowed to clot, and the serum was removed after centrifugation and frozen at -80°C until use. The presence of high-titer serum antibodies was confirmed by ELISA (described below). The 8-18C5 hybridoma cell line was provided by C. Linington. An IgG1 mouse anti-OVA hybridoma cell line (1B7.11) was purchased from UCSF hybridoma core.

Hybridoma cells were cultured in a BD Cell mAb Medium Serum-Free (BD) in large-scale flasks (Corning). mAbs were purified from hybridoma culture supernatants by the UCSF hybridoma core. The concentrations of the proteins were determined by means of the BCA protein assay (Thermo Fisher Scientific), according to the manufacturer's recommendations.

Antibody administration. B-MHC II^{+/+} and B-MHC II^{-/-} mice were immunized with 100 µg rhMOG emulsified in CFA. 200 ng *B. pertussis* toxin was injected on days 0 and 2. When B-MHC II^{+/+} mice showed clinical EAE signs (score ~2.0), B-MHC II^{-/-} mice received i.p. injections of 150 µg anti-MOG IgG1 mAb 8-18C5, an equal amount of isotype control OVA-specific mAb 1B7.11, or 250 µl IgH^{MOG-ki} serum, or non-Tg littermate mice serum three times at 48 h-intervals (Schluesener et al., 1987; Pöllinger et al., 2009). Sera were obtained through cardiac puncture from IgH^{MOG-ki} or non-Tg littermate mice.

Detection of anti-MOG antibodies. Total serum MOG-specific IgG was quantified using a commercially available anti-mouse MOG (1–125) IgG quantitative ELISA kit (Sensolyte; AnaSpec). Wells were precoated with rmMOG (1–125) protein. Results are expressed as IgG concentrations in µg/ml serum (Figs. 3 A and 4 E) or, to facilitate comparison with other Ig measurements, as OD of serial twofold dilutions (Fig. 4 D). Serum MOG-specific IgM antibodies were measured using a noncommercial ELISA. 96-Maxisorb plates (Costar) were precoated with rmMOG (1–117) protein (10 µg/ml in PBS), blocked with BSA (Sigma-Aldrich), and incubated with sera for 2 h at the indicated dilution. After washing, MOG-specific IgM retained by the plate-bound MOG was detected with horseradish-peroxidase-conjugated anti-mouse IgM (IMMUNO-TEK). Total IgM and IgG were measured using quantitative Ig ELISA kits (IMMUNO-TEK). Total serum IgG or IgM was captured by anti-mouse IgG or IgM antibodies immobilized on micro wells. OVA-coated plates were used as negative controls for non-specific binding. SOFTmax ELISA plate reader (450-nm wavelength) and software (Molecular Devices) were used for data analysis.

Proliferation assays. Cells were grown in RPMI supplemented with 10% FCS, β-mercaptoethanol, L-glutamine, and penicillin/streptomycin. For thymidine incorporation, either 2.5 × 10⁵ cells/ml of total splenocytes or 10⁴ cells/ml of purified CD4⁺ T cells plus 2.5 × 10⁵ cells/ml of purified B220⁺ B cells were cultured in 96-well plates in the presence of different concentrations of the rMOG proteins or MOG p35–55 (range 1–50 µg/ml). After 72 h, cells were pulsed with [³H]thymidine and harvested 16 h later. Thymidine incorporation is expressed as cpm. In certain experiments, T cell proliferation was assessed by CFSE dilution assay after gating on CD4⁺ T cells.

Cell isolation. Single-cell leukocyte suspensions from spleens and peripheral lymph nodes (paired axillary and inguinal) were generated by gentle dissection. CNS mononuclear cells were isolated from EAE mice at peak of disease after cardiac perfusion with PBS, as described previously (Zeine and Owens, 1992). In brief, minced CNS (spinal cord) tissues were digested with 2.5 mg/ml collagenase D (Roche) at 37°C for 45 min. Mononuclear cells were isolated by passing the tissue through 70-µm cell strainers (BD), followed by discontinuous Percoll gradient (70/30%; Sigma-Aldrich) centrifugation. Lymphocytes were collected from the 30:70% interface and washed. Total cell numbers were determined by counting on a hemocytometer, and viability was assessed by trypan blue exclusion.

MACS (Miltenyi Biotec) was used to purify lymphocyte populations according to the manufacturer's instructions. CD4⁺ T cells were isolated by depletion of non-CD4⁺ T cells (negative selection, kit #130-090-860). Naive B cells were isolated by depletion from CD43-expressing B cells (activated B cells, plasma cells and CD5⁺ B-1a cells) and non-B cells (kit #130-090-862) according to the manufacturer's instructions. When necessary, cells were enriched a second time using a fresh MACS column to obtain >95% cell purities, as assessed by flow cytometry.

Flow cytometric analysis. For two- to six-color immunofluorescence analysis, single-cell suspensions (10⁶ cells) were incubated with anti-mouse

FcR1IB/FcR1IIIA mAb (2.4G2; BD) to avoid nonspecific staining and were subsequently stained at 4°C using predetermined optimal concentrations of mAb for 30 min, as described previously (Weber et al., 2010). Blood erythrocytes were lysed before staining using FACS lysing solution (BD). Antibodies to the mouse proteins CD45R (B220), FITC/PE/APC/PE-Cy7/PerCP-Cy5.5/APC-Cy7 (RA3-6B2), CD4 FITC/PE (RM4-4), CD4 APC/PE-Cy7/PerCP-Cy5.5/APC-Cy7 (RM4-5), CD11c-APC (N418), CD11b-PE (ICRF44), CD44-PE (IM7), CD62L-APC (MEL-14), IgM-FITC (eB121-15F9), and IgD-FITC (11-26c) were purchased from eBioscience. MHC II I-Ab-FITC (AF6-120.1) was purchased from BD. Isotype- and concentration-matched control antibodies were used to assess nonspecific staining. In certain experiments, cells were fixed in 0.3% paraformaldehyde and kept at 4°C until analysis. Lymphocytes were examined by forward and side light scatter on a FACSCanto II flow cytometer using FACSDiva software (BD).

Intracellular cytokine staining. Lymphocytes were stimulated *in vitro* with 50 ng/ml PMA (Sigma-Aldrich) and 1 µg/ml ionomycin (Sigma-Aldrich) in the presence of 1 µl/ml brefeldin A (eBioscience) for 6 h before staining. FcRs were blocked before cell surface staining with anti-CD4-PerCP-Cy5.5 (RM4-5; eBioscience). After staining, cells were washed, fixed, and then permeabilized using the Cytotfix/Cytoperm Plus Fixation/Permeabilization kit (BD) according to the manufacturer's instructions. Permeabilized cells were then stained with anti-IL-17-PE-Cy7 (eBio17B7) and anti-IFN-γ-FITC (XMG1.2; all from eBioscience).

Bcl-6 staining. For detection of Bcl-6, FcRs were blocked followed by cell surface staining with anti-CD4-PE (GK1.5), anti-PD-1 (CD279)-PE-Cy7 (J43), and anti-ICOS (CD278)-FITC (C398.4A; all from eBioscience) and anti CXCR5-PE (2G8; BD). Bcl-6 staining was subsequently performed using the Foxp3/Transcription Factor Staining Buffer set (eBioscience) and anti-Bcl-6-PE (mGI191E; eBioscience).

Histology and immunohistochemistry. Brains, spinal cords, and optic nerves were removed and fixed in 10% neutral-buffered formalin, paraffin-embedded, and sectioned. Representative sections were stained with Luxol fast blue (LFB)-hematoxylin and eosin (H&E; for inflammation and demyelination) and reticulin preparation (for connective tissue) and examined by light microscopy. Meningeal and parenchymal inflammatory foci (>10 clustered inflammatory cells) were counted by a blinded observer (R.A. Sobel). Standard avidin-biotin immunohistochemical staining was performed on the sections with rabbit anti-mouse CD3 (Abcam) and rat anti-mouse CD45R (B220; BD), using reagents from Vector Laboratories. Normal mouse spleen tissue served as positive staining controls; negative controls included omission of the primary antibody. Neuropathologic analysis was performed by R.A. Sobel.

Statistical analysis. Data are presented as mean ± SEM or SD. For clinical scores, significance between groups was examined using the Mann-Whitney *U* test. A value of *P* < 0.05 was considered significant. All other statistical analysis was performed using a one-way multiple-range ANOVA test for multiple comparisons. A value of *P* < 0.01 was considered significant.

We thank Dr. Patricia Nelson (University of California, San Francisco) for helpful discussion and Kara Pekarek for technical assistance. We thank Drs. Jeri-Anne Lyons and Anne H. Cross for providing us with the IgH^{B1-B-mem} mice.

N. Molnarfi was supported by an advanced researcher fellowship from the Swiss National Science Foundation (SNF, PA00A-119532) and is a recipient of an advanced researcher exchange 2011 fellowship from the European Committee for Treatment and Research in Multiple Sclerosis (ECTRIMS) Foundation. U. Schulze-Toppoff is a fellow of the National Multiple Sclerosis Society (NMSS) and the Deutsche Forschungsgemeinschaft (DFG; SCHU 2587/1-1). M.S. Weber was supported by the NMSS (RG 4450A1/T). T. Prod'homme received fellowship support from the NMSS and Teva Neuroscience. M. Varrin-Doyer is a fellow of the NMSS. J. Hidalgo was supported by SAF2011-23272. Support for this work was provided to S.S. Zamvil by the National Institutes of Health (NIH; R01 AI-073737 and R01 NS-063008), the NMSS (RG 3913 and RG 4768), the Guthy Jackson Charitable Foundation, the Dana Foundation, and the Maisin Foundation and to M.J. Shlomchik by the NIH (R01 AI-043603).

The authors declare no competing financial interests.

Author contributions: N. Molnarfi, U. Schulze-Toppoff, and S.S. Zamvil designed research, analyzed data, and wrote the paper; M.S. Weber, J.C. Patarroyo, M. Varrin-Doyer, T. Prod'homme, and A.J. Slavin analyzed data, gave conceptual advice, and discussed the results at all stages; N. Molnarfi, U. Schulze-Toppoff, J.C. Patarroyo, and T. Prod'homme performed the experiments; M. Varrin-Doyer and A. Shetty assisted in the performance of experiments; R.A. Sobel performed the neuropathologic analysis; C. Linington, J. Hidalgo, D.E. Jenne, H. Wekerle, C.C.A. Bernard, and M.J. Shlomchik contributed new reagents; and S.S. Zamvil supervised the study. All authors read, commented on, and approved the final manuscript.

Submitted: 4 April 2013

Accepted: 11 November 2013

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