In multiple sclerosis (MS) patients who carry the Class II major histocompatibility (MHC) type HLA-DR2, T cells specific for amino acids 95-116 in the proteolipid protein (PLP) are activated and clonally expanded. However, it remains unclear whether these autoreactive T cells play a pathogenic role or, rather, protect against the central nervous system (CNS) damage. We have addressed this issue, using mice transgenic for the human MHC class II region carrying the HLA-DR2 (DRB1*1502) haplotype.

After stimulating cultured lymph node cells repeatedly with PLP95-116, we generated 2 HLA-DR2–restricted, PLP95-116–specific T-cell lines (TCLs) from the transgenic mice immunized with this portion of PLP. The TCLs were CD4+ and produced T-helper 1 (Th1) cytokines in response to the peptide. These TCLs were adoptively transferred into RAG-2–/– mice expressing HLA-DR2 (DRB1*1502) molecules. Mice receiving 1 of the TCLs developed a neurological disorder manifested ataxic movement without apparent paresis on day 3, 4, or 5 after cell transfer. Histological examination revealed inflammatory foci primarily restricted to the cerebrum and cerebellum, in association with scattered demyelinating lesions in the deep cerebral cortex. These results support a pathogenic role for PLP95-116–specific T cells in HLA-DR2+ MS patients, and shed light on the possible correlation between autoimmune target epitope and disease phenotype in human CNS autoimmune diseases.

bral cortex in immunodeficient mice generated by mating the HLA-DR2 (DRB1*1502) transgenic mice with recombination activation gene knockout mice (RAG-2−/− mice) (14). These results support a pathogenic role of PLP95-116-specific T cells in HLA-DR2−/− MS patients, and indicate a possible correlation between target epitope and disease phenotype in human CNS autoimmune diseases.

**Methods**

**Mice.** C57BL/6J (B6) and RAG-2−/− mice with an H-2b background (14) were purchased from CLEA Japan Inc. (Tokyo, Japan) and Taconic Farms (Germantown, New York, USA), respectively. Generation of DRA transgenic mice with a B6 background (DRA-B6) (15) and DRB1*1502 transgenic mice with a B10.RQB3 background (DRB1*1502-B10.RQB3) (16) have been described previously. DRB1*1502 transgenic mice with a B6 background (DRB1*1502–B6) were obtained by backcrossing the DRB1*1502–B10.RQB3 mice to B6 mice for 6 generations. After crossing DRA-B6 and DRB1*1502–B6 mice, HLA-DR2 (HLA-DRB1*1502) transgenic mice (HLA-DR2–B6) were selected from their F1 progeny. HLA-DR2–B6 mice were further crossed twice with RAG-2−/− mice; RAG-2−/− mice expressing HLA-DR2 (DRB1*1502) (HLA-DR2/RAG-2−/−) were selected from the N2 progeny. Both HLA-DR2–B6 and HLA-DR2/RAG-2−/− mice were identified using PCR as described below. All the mice were kept under specific pathogen−free conditions. Female mice (6–12 weeks of age) heterozygous for the DRA or DRB1*1502 genes (or both together) were used.

**PCR.** A pair of primers were used for each gene as follows. DRA gene: 5′-GACACGTGATCATCCAGGCGC-3′ and 5′-GACCCATGATGCTTGACCAC-3′; DRB1*1502 gene: 5′-CTCAAGAGGGGATGCTATTTCCTC-3′ and 5′-TGATGAAAGCATCTTTAACCAACACCA-3′; RAG-2 gene: 5′-CCACCTCTGTTATACCAGCAAC-3′ and 5′-GTCTGCCTAAAGAGACCCCC-3′; pMC1 neo/poly(A)+ fragment disrupting endogenous RAG-2 gene in RAG-2−/− mice: 5′-TACACCTTACCTCCATTTGCGCC-3′ and 5′-TATGTTCCTATTAGCGGTTCCC-3′. The genomic DNA extracted from mouse tail tips was amplified by PCR for 30 cycles by using the temperature profile that follows. DRA and DRB1*1502 genes: denaturation at 94°C for 1 minute, annealing at 66°C for 30 seconds, and extension at 72°C for 30 seconds. RAG-2 gene and pMC1 neo/poly(A)+ fragment: denaturation at 94°C for 1 minute, annealing at 62°C for 30 seconds, and denaturation at 72°C for 30 seconds. Amplified PCR products were analyzed on 6% polyacrylamide gels stained with ethidium bromide.

**Reagents.** The human PLP95-116 (AVRQIFGDYKT-TICGKLSDLATV) and MBP143-168 (GVDAGGTL-SKIFKLLGRDRSRSGPMA) peptides were purchased from the Peptide Institute, Inc. (Osaka, Japan) and Kurabo Industries, Ltd. (Osaka, Japan), respectively. Incomplete Freund’s adjuvant and heat-killed Mycobacterium tuberculosis H37Ra were purchased from Difco Laboratories (Detroit, Michigan, USA). Recombinant human IL-2 (rhIL-2) was a gift of Shionogi Pharmaceutical Co. Ltd. (Osaka, Japan). Anti-NK1.1 mAb (PK136) was purified from culture supernatants using protein A column chromatography.

**Generation and propagation of PLP95-116-specific TCLs.** HLA-DR2–B6 mice were immunized in the footpads of both hind feet and in the tail base with 200 µL of emulsion containing 200 µg of PLP95-116 in incomplete Freund’s adjuvant supplemented with 500 µg of M. tuberculosis. Ten days after immunization, popliteal and inguinal lymph nodes were removed, and the single-cell suspensions were prepared at 2 × 10^6 cells/mL in complete medium (RPMI-1640 supplemented with 10% FCS, 10 mM HEPES buffer, 5 × 10^-3 M 2-mercaptoethanol, 2 mM l-glutamine, and 100 U/mL of penicillin and 100 µg/mL of streptomycin). Two hundred microliters of the suspension was added to 96-well plates and then stimulated with 25 µg/mL of PLP95-116. The cells were fed with the complete medium supplemented with 7.5% T-cell growth factor medium (supernatant of concanavalin A-stimulated rat splenocyte cultures) every 3 days. An aliquot of each TCL was assayed for antigen specificity on day 14. The TCLs that were reactive to PLP95-116 were further expanded and were restimulated every 2 weeks with the peptide in the presence of irradiated (35 Gy) syngeneic splenocyte cultures as APCs.

**Proliferation and inhibition assays.** Splenocytes (4 × 10^5/well) from HLA-DR2–B6 or DRA-B6 mice, or PBMCs (2 × 10^5/well) from healthy human subjects were activated with various antigens in the presence of PE-labeled anti–HLA-DR mAbs. (a) Splenocytes from B6, DRA-B6, DRB1*1502-B6, and HLA-DR2–B6 mice were stained with PE-labeled anti–HLA-DR (L243) or FITC-labeled anti–HLA-DRβ (TU36) mAb. The bars marked M1 indicate the population that stained significantly with the mAb, whereas M2 shows the positive population selected on more strict criteria. (b) Splenocytes from HLA-DR2–B6 mice were doubly stained with PE-conjugated anti–HLA-DRα (L243) and FITC-conjugated anti–HLA-DRβ (TU36) mAbs.
Figure 2
Proliferative response of TCLs B3-4 and B3-23 generated from HLA-DR2–B6 mice immunized with PLP95-116. Peptide-specific proliferation of TCLs B3-4 (a) and B3-23 (b) were evaluated as described in Methods. By using splenocytes from HLA-DR2–B6 (HLA-DR2*, DRα/1-EB2*, I-Aβ*) or DRA-B6 (DRα/1-EB2*, I-Aβ*) mice as APCs, T-cell proliferative response to control peptide MBP143-168 or PLP95-116 (25 μg/mL) was measured. Blocking effects of anti–HLA-DR mAb (G46-6) or isotype-matched control IgG2a (G155-178) were also examined as indicated. Data represent mean ± SD of the cpm obtained by triplicate assays in 4 independent experiments.

designated MN (HLA-DR2 [DRB1*1502]/4) and YM ([HLA-DR1]/8) were irradiated (35 Gy for mouse splenocytes and 40 Gy for human PBMCs) and used as APCs. T-line cells (4 × 10^4/well) were incubated for 72 hours with or without peptide in the presence of APCs. The cultures were pulsed with 1 μCi of [3H]thymidine for the last 16 hours of the incubation. Cell incorporation of [3H]thymidine was counted with a 1205 Betaplate counter (Pharmacia, Uppsala, Sweden). Mean cpm of the triplicate cultures was calculated, and TCLs that showed both a stimulation index greater than 3.0 and a change in cpm of more than 500 in the presence of the peptide were defined as peptide-specific. Blocking assays were performed with anti–HLA-DR mAb (G46-6) or isotype-matched mouse IgG2a (G155-178) added at a final concentration of 10 μg/mL.

ELISA. Supernatants were collected from T-cell cultures 48 hours after peptide stimulation. The concentration of cytokines (IFN-γ, IL-2, TNF-α, and IL-4) in the supernatants was measured by sandwich ELISA using pairs of relevant anti-cytokine mAbs according to the protocol recommended by PharMingen (San Diego, California, USA).

Adaptive transfer experiments. HLA-DR2/RAG-2−/− and RAG-2−/− mice were intravenously injected with 500 μg of anti-NK1.1 mAb for in vivo depletion of natural killer (NK) cells. The next day, the mice were irradiated (3.5 Gy) and injected via tail vein with T-line cells (5 × 10^6) generated from HLA-DR2–B6 mice. The line cells had been stimulated with PLP95-116 in the presence of syngeneic splenocytes for 3 days. Immediately after cell transfer, 250 ng of pertussis toxin (Seikagaku Kogyo Co. Ltd., Tokyo, Japan) was intravenously injected. All the mice were observed daily for neurological signs.

Flow cytometry. Splenocytes or T-line cells were first incubated with Fc Block (anti-mouse FcRγ/III mAb) (PharMingen) for 5 minutes, and then incubated for 30 minutes with mAbs in combination as follows. Splenocytes were incubated with either FITC-conjugated anti–HLA-DRβ (TÜ36) and PE-conjugated anti–HLA-DRα (L243), or FITC-conjugated anti–I-Abβ (25-9-17) and PE-conjugated anti–I-Aγβ (AF6-12.1). T-line cells were incubated with either FITC-labeled anti-CD4 (RM4-5) and PE-labeled anti-CD8 (53-6.7); FITC-labeled anti-adhesion molecule mAb (anti-LFA-1 [M17/4], anti-CD44 [IM7], anti-α4 integrin [M17/4], anti-ICAM-1 [3E2], or anti-L-selectin [MEL-14]) and PE-labeled anti-CD4 (RM4-5); or FITC-labeled T-cell receptor (TCR) Vβ chain–specific mAb (anti-Vβ2, -3, -4, -5, -6, -7, -8, -9, -10, -11, -12, -13, or -14) and PE-labeled mAb recognizing the common determinant of TCR Vβ chains (H57-597). All mAbs were purchased from PharMingen, except anti–HLA-DRα (Becton Dickinson Immunocytometry Systems, Mountain View, California, USA). After intensive washing, samples were suspended in PBS containing 0.5 μg/mL of propidium iodide (Wako Pure Chemical Industries, Ltd., Osaka, Japan) and analyzed using FACSort with CellQuest software (both from Becton Dickinson Immunocytometry Systems).

Detection of intracellular cytokine. T-line cells (10^6/well) were stimulated in 48-well plates with 25 μg/mL of PLP95-116, 50 IU/mL of rhIL-2, or both, in the presence of APCs (2.5 × 10^5/well) for 48 hours. GolgiPlug (brefeldin A; PharMingen) was added for the last 10 hours of the incubation to disrupt Golgi function. The detection of intracellular IFN-γ was performed using a Cytofix/Cytoperm kit (PharMingen). Briefly, cells preincubated with Fc Block® were stained with PE-labeled anti-CD4 (YM5-1) and FITC-labeled anti–HLA-DRβ (TÜ36) and PE-conjugated anti–HLA-DRα (L243), or FITC-conjugated anti–I-Abβ (25-9-17) and PE-conjugated anti–I-Aγβ (AF6-12.1). T-line cells were incubated with either FITC-labeled anti-CD4 (RM4-5) and PE-labeled anti-CD8 (53-6.7); FITC-labeled anti-adhesion molecule mAb (anti-LFA-1 [M17/4], anti-CD44 [IM7], anti-α4 integrin [M17/4], anti-ICAM-1 [3E2], or anti-L-selectin [MEL-14]) and PE-labeled anti-CD4 (RM4-5); or FITC-labeled T-cell receptor (TCR) Vβ chain–specific mAb (anti-Vβ2, -3, -4, -5, -6, -7, -8, -9, -10, -11, -12, -13, or -14) and PE-labeled mAb recognizing the common determinant of TCR Vβ chains (H57-597). All mAbs were purchased from PharMingen, except anti–HLA-DRα (Becton Dickinson Immunocytometry Systems, Mountain View, California, USA). After intensive washing, samples were suspended in PBS containing 0.5 μg/mL of propidium iodide (Wako Pure Chemical Industries, Ltd., Osaka, Japan) and analyzed using FACSort with CellQuest software (both from Becton Dickinson Immunocytometry Systems).

Histology. Four to 6 days after cell transfer, the moribund HLA-DR2/RAG-2−/− mice were anesthetized and
per fused through the heart with 0.1 M phosphate buffer followed by 4% parformaldehyde. The brains and spinal cords were removed and fixed in the same fixative for 3 days. The paraffin-embedded tissues were sectioned at a thickness of 5 μm. Sections were stained with hematoxylin and eosin (H&E) or Luxol fast blue for conventional light microscopy.

Results

Expression of HLA-DR2 molecules in HLA-DR2–B6 mice. We produced HLA-DR2–B6 mice by mating DRA–B6 mice (15) with DRB1*1502–B6 mice that were generated from DRB1*1502–B10.RQB3 mice (16). Previous studies had ascertained that DRA–B6 and DRB1*1502–B10.RQB3 mice express DRα and DRβ chains, respectively, on immunocompetent cells. It is of note that DRα or DRβ chain alone could not be expressed on the cell surface, but either of the chains could appear on the cell surface in the context of xenogeneic heterodimer complexed with an endogenous I-Eβ or I-α chain. We first analyzed expression of DRα and DRβ chains on splenocytes from DRA–B6, DRB1*1502–B6, and HLA-DR2–B6 mice. The cells were stained with PE-labeled anti–HLA-DRα (L243) or FITC-labeled anti–HLA-DRβ mAb (TÜ36) for single-color analysis (Figure 1a). Consistent with a previous report (17), a proportion of the splenocytes from DRA–B6 mice (18.3 ± 1.0%; n = 3) were stained with the anti–HLA-DRα mAb, indicating that the DRα chain appears on the cell surface in the form of a DRα/DRβ heterodimer. In contrast, DRβ cells were not detected in DRB1*1502–B6 mice. This is explained by the fact that mice with a B6 background lack endogenous I-Eβ, which is necessary for formation of a xenogeneic heterodimer. On the other hand, the splenocytes from HLA-DR2–B6 mice were stained not only with the anti–HLA-DRα mAb, but also with the anti–HLA-DRβ mAb, implying that the DRβ chain is probably expressed in the form of a DRα/DRβ heterodimer (HLA-DR2 molecule). It is also noteworthy that the proportion of the cells that stained brightly with the anti–HLA-DRα mAb (M2 in Figure 1a) was significantly higher in samples from HLA-DR2–B6 mice than from DR-β-B6 mice (15.1 ± 2.8%, n = 6 vs. 3.9 ± 0.3%, n = 5, respectively). In contrast, the expression level of endogenous I-Aβ molecules in the transgenic mice was not altered by DRA or DRB1*1502 transgene (data not shown). The augmentation of DRα chain expression together with the expression of DRβ chain indicates that HLA-DR2 molecules are formed and expressed in HLA-DR2–B6 mice. Two-color analysis of HLA-DR2–B6 splenocytes (Figure 1b) further confirmed coexpression of DRα and DRβ chains in 32.0 ± 5.2% (n = 5) of the splenocytes. These data imply that HLA-DR2–B6 mice express 3 different MHC class II molecules: HLA-DR2, DRα/1-Eββ, and I-Aβ.

Characterization of PLP95-116–specific TCLs generated from HLA-DR2–B6 mice. Preliminary experiments have revealed that HLA-DR2–B6 mice mount a T-cell response to PLP95-116 after immunization with the peptide (data not shown), and that the response is further augmented if the mice are pretreated with anti-NK cell antibody (anti-NK1.1 mAb, PK136). However, the primary T cells that are reactive to the PLP peptide are highly heterogeneous. To focus on the HLA-DR2–restricted T cells, we attempted to generate TCLs recognizing PLP95-116 in the context of HLA-DR2 molecules. More than 100 TCLs were generated from PLP95-116–primed HLA-DR2–B6 mice, and were examined for reactivity to PLP95-116 in a proliferation assay on day 14. We selected 15 TCLs that were reactive to PLP95-116, and restimulated the lines with the peptide for further expansion. We selected 5 PLP95-116–specific TCLs that proliferated in response to PLP95-116 in the presence of control mAb but not in the presence of anti–HLA-DR mAb (G46-6) (data not shown). Two of the 5 TCLs, designated B3-4 and B3-23, were studied in more detail.

The specificity and MHC restriction of TCLs B3-4 and TCL B3-23 are shown in Figure 2. Both of these TCLs proliferated significantly in response to PLP95-116, but

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<td>Induction of encephalitis by TCL B3-23 in HLA-DR2/RAG-2−/− mice</td>
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Encephalitisogenicity of TCLs B3-4 and B3-23 was assessed by adoptive transfer of T-line cells into HLA-DR2/RAG-2−/− or RAG-2−/− mice, as described in Methods. The HLA-DR2/RAG-2−/− and RAG-2−/− mice were monitored for the appearance of neurological signs for 8 weeks and 3 weeks, respectively. *Two moribund mice sacrificed for histological examination are included.
not in response to MBP143-168 in the presence of irradiated syngeneic (HLA-DR2–B6) splenocytes; these proliferative responses were almost completely blocked with the anti–HLA-DR mAb. Because splenocytes from HLA-DR2–B6 mice express DRα1/1-Eβ6 and I-Aα molecules in addition to HLA-DR2 molecules, there still remained a possibility that PLP95-116 was presented in the context of these non–HLA-DR2 molecules. However, this possibility was excluded because DRα-B6 splenocytes expressing DRβ1/1-Eβ6 and I-Aα molecules could not present the peptide to the TCLs.

Analysis by flow cytometry revealed that both of the TCLs were CD4+CD8− and expressed high levels of LFA-1, CD44, and ICAM-1. In contrast, α4 integrin was moderately positive in TCL B3-4, but rarely detected in TCL B3-23. L-selectin was not detected in either of the TCLs. TCL B3-4 expressed Vβ2 and Vβ14 TCRs predominantly, whereas TCL B3-23 was enriched in Vβ6- and Vβ14-expressing T cells (data not shown).

As shown in Figure 3, TCL B3-4 produced a large amount of IFN-γ and IL-2, whereas TCL B3-23 produced IFN-γ but not IL-2. Neither TNF-α nor IL-4 was detected in the supernatants. These results indicate that these TCLs are composed of Th1-type T cells.

To confirm HLA-DR2 restriction of the TCLs, PBMCs from healthy subject MN (HLA-DR2 [DRB1*1502]/4) and YM (HLA-DR1/8) were used as APCs in the assays. These TCLs did not show a significant proliferative response to PLP95-116 in the presence of the human PBMCs (data not shown). In contrast, they produced comparable levels of IFN-γ in response to PLP95-116 only when HLA-DR2–expressing PBMCs from MN were used as APCs; their IFN-γ production was completely blocked by the anti–HLA-DR mAb (Figure 4). We further analyzed the TCLs for intracellular IFN-γ synthesis in response to PLP95-116. As shown in Figure 5, the vast majority (∼95%) of CD4+ T-line cells from TCLs B3-4 and B3-23 produced high levels of IFN-γ in response to PLP95-116 when syngeneic (HLA-DR2–B6) splenocytes were used as APCs. In contrast, when we used human PBMCs from the HLA-DR2– (DRB1*1502−) subject (MN), IFN-γ production was much weaker — only about 20% of the cells were defined as positive. It is possible that the poor response was due to a partial species barrier between murine accessory or costimulatory molecules and the human ligands. Therefore, we performed the assays in the presence of exogenous rhIL-2. We found that the T-line cells vigorously produced IFN-γ with positive staining for approximately 80% of the cells. This indicates that the large majority of the line cells would recognize PLP95-116 presented by human APCs expressing HLA-DR2 molecules. Furthermore, this analysis excluded the possibility that each TCL comprises 2 subpopulations: 1 that responds to mouse APCs presenting PLP95-116, and another that is responsive to human APCs presenting PLP95-116.

**Induction of autoimmune encephalitis by an HLA-DR2–restricted, PLP95-116–specific TCL.** We next examined encephalitogenicity of the TCLs in adoptive transfer experiments. We have previously revealed that RAG-2−/− mice lacking T cells, B cells, and NKT cells become highly prone to passive experimental autoimmune encephalomyelitis (EAE) after eliminating NK cells in vivo (18). To enhance the chance of disease induction by the TCLs, we generated HLA-DR2/RAG-2−/− mice that are MHC-compatible with the TCLs and could serve as their recipients. After in vitro stimulation with PLP95-116 in the presence of syngeneic (HLA-DR2–B6) splenocytes for 3 days, 5 × 10⁶ T-line cells were intravenously transferred to the mice that had been injected with anti-NK1.1 mAb (PK136) 1 day before cell transfer. TCL B3-4 did not induce any neurological sign in the recipient mice during the observation period of 8 weeks (Table 1). In contrast, 4 of the 7 mice receiving TCL B3-23 developed neurological...
signs of cerebellar ataxia 3–5 days after transfer. Signs of EAE in the mice began with the turning of their heads and trunks to 1 side and progressed to axial rotatory movement. Two mice died 1 day after clinical onset. Because the other 2 symptomatic mice developed a comparable progression of disease, we sacrificed them for histological examination 1 day after clinical onset. It was of note that there was no sign of the tail atony or hind-limb paresis that is characteristic of conventional EAE models. In contrast, transfer of the pathogenic TCL into nontransgenic RAG-2–/– mice did not induce any clinical sign during the 3-week observation period. Based on these results, we presume that HLA-DR2–restricted recognition of endogenous PLP95-116 in vivo lay behind the neurological sign induced by TCL B3-23.

To confirm the CNS involvement, we performed pathological studies of the 2 moribund HLA-DR2/RAG-2/– mice receiving TCL B3-23. Examination of the CNS samples from these mice revealed meningeal and perivascular parenchymal inflammatory infiltrates (Figure 6, c–e, g) composed of mononuclear cells and neutrophils (Figure 6h). Infiltration was most prominent in the cerebrum and cerebellum. In addition, discrete inflammatory demyelinating lesions were scattered predominantly in the deep cerebral cortex (Figure 6, a, b, e, and f). By contrast, only very low levels of meningeal infiltrates were detected in the spinal cord. These histological findings were thought to correlate well with the clinical phenotype.

**Discussion**

We previously reported that the frequency of PLP95-116–specific T cells is significantly elevated in HLA-DR2+ MS patients in comparison with those who are HLA-DR2– (6). The association of PLP95-116 with HLA-DR2+ MS was later confirmed by Trotter et al. (10). Using whole PLP molecules, they further defined PLP95-117 as an immunodominant epitope for HLA-DR2+ MS. More recently, we reported evidence for transient or continuous activation/expansion of PLP95-116–specific T-cell clones in an HLA-DR2+ (DRB1*1501/DRB1*1502+) MS patient (19). Despite these observations, the in vivo role of PLP95-116–specific T cells has remained speculative. Here we demonstrate for the first time that HLA-DR2–restricted, (DRB1*1502–restricted) PLP95-116–specific T cells can cause autoimmune encephalitis in mice expressing HLA-DR2 (DRB1*1502) molecules. Because humans and rodents share identical amino acids in PLP95-116 residues (20), our results indicate that PLP95-116 can be a target epitope for autopathogenic T cells not only in HLA-DR2 (DRB1*1502) transgenic mice but also in HLA-DR2+ (DRB1*1502+) MS. Recent studies have demonstrated that transgenic mice expressing human HLA-DR molecules could develop encephalomyelitis after being challenged with myelin extract or peptide (21, 22). However, as encephalitogenic, DR-restricted TCLs or clones were not established from the transgenic mice, it
remained unclear if HLA-DR–restricted T cells alone could induce encephalomyelitis, or if other cell types such as B cells are necessary.

A number of studies have revealed that susceptibility to MS in Caucasian populations is positively linked with the DRB1*1501 allele (23). However, our study has revealed that a proportion of Japanese MS patients possess DRB1*1502 but not DRB1*1501, and that TCLs recognizing the DRB1*1501–related MBP or PLP peptides can be efficiently generated from DRB1*1502+ patients (Ohashi et al., unpublished data). Furthermore, DRB1*1502 was found to be present in the vast majority of patients with concentric sclerosis (Baló’s disease) (unpublished observation), which is probably a variant form of MS (24). Moreover, DRB1*1501 and DRB1*1502 differ only in the amino acid at position 86 of the DRβ chain (valine in DRB1*1501 and glycine in DRB1*1502) (25). Although the size of the P1 pocket in the peptide-binding groove differs between the DRB1*1501 and DRB1*1502 molecules (26, 27), MBP85-99 peptide could be presented to HLA-DR2-restricted (DRB1*1501–restricted) T cells also in the context of HLA-DR2 (DRB1*1502) molecules (28). These observations encouraged us to use HLA-DR2 (DRB1*1502) transgenic mice for establishing a model for HLA-DR2+ MS, or at least for relevant subtypes of MS.

It is of particular interest that the clinical and pathological features of EAE described herein are quite different from those of classical EAE. Whereas the classical EAE models are characterized by ascending paralysis caused by the lesions in the spinal cord white matter, the neurological sign in the HLA-DR2/RAG-2−/− mice receiving TCL B3-23 was represented by ataxia without paralysis, consistent with the predominant lesions in the cerebral cortex and cerebellum. Regarding the unique EAE phenotype in this study, it is relevant that some EAE models induced by PLP peptides exhibit similar clinical phenotypes, characterized by rotatory movement or ataxia (29–31). Just such an unusual form of EAE, displaying ataxia, was seen by Greer and colleagues in BALB/c, CBA/J, and C3H/HeJ mice when PLP peptides within PLP residues 178–232 used for active immunization, whereas encephalitogenic peptides within other PLP domains caused a typical form of EAE (31). Based on these results, Greer et al. speculated that there might be a correlation between target epitope and disease phenotype. Another relevant piece of information is that T cells recognizing MBP, a representative myelin antigen, and T-cells specific for a non-myelin antigen, S100β protein, have been shown to cause distinct disorders in the rat model of EAE (32). Very interestingly, S100β–specific T cells invaded not only white matter but also brain cortex, whereas MBP–specific T cells are primarily a cause of white matter lesions. Because distribution of MBP and S100β within the CNS could not account for this difference, Kojima et al. explained it by speculating that there is a differing capacity to process and present specific antigen/epitope among different regions of the CNS. These reports together with our own data suggest that a variety of neurological signs of MS may be finely dissected after identifying predominant target epitopes in each patient. In this regard, induction of cerebral cortical lesions induced by HLA-DR2–restricted, PLP95-116–specific T cells may be relevant to understanding why cortical lesions are seen in some MS patients but not in others (33). Further exploration of the correlation between epitope and disease phenotype in HLA-DR2 (DRB1*1502) transgenic mice may provide essential information for understanding anti-myelin autoimmunity.
Acknowledgments

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