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Article

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HLA-DR2–restricted responses to proteolipid protein 95-116 peptide cause autoimmune encephalitis in transgenic mice

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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.

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Introduction

Multiple sclerosis (MS) is a chronic inflammatory disease of the central nervous system (CNS) that is histologically characterized by focal mononuclear cell infiltration with a varying extent of demyelination (1, 2). It is postulated that autoimmune T cells recognizing myelin autoantigens such as myelin basic protein (MBP) or proteolipid protein (PLP) play a central role in the pathogenesis of MS. This postulate is based on substantial evidence such as increased frequency of activated MBP- or PLP-specific T cells in peripheral blood and cerebrospinal fluid of MS patients (3–6), and significant complementarity-determining region 3 (CDR3) homology between T-cell infiltrates within MS plaques and MBP- or PLP-specific T-cell clones generated from MS patients (7, 8). Moreover, previous studies have identified the peptides MBP84-102 and PLP95-116 as immunodominant and possible encephalitogenic epitopes in MS patients with the *HLA-DR2* haplotype (9,

10). However, the role of such autoreactive T cells remains unclear. In fact, it is possible that some (but not all) autoreactive T cells may protect against CNS tissue damage by producing neurotrophic factors (11–13).

To address the issue regarding the function of autoimmune T cells, we have introduced *HLA-DR2* (*DRB1*1502*) transgenic mice that are able to mount a T-cell response to DR2-related epitopes in the context of *HLA-DR2* molecules. To explore the potential pathogenicity of PLP95-116 in *HLA-DR2*⁺ MS, we challenged the *HLA-DR2* (*DRB1*1502*) transgenic mice with PLP95-116 and generated 2 peptide-specific T-cell lines (TCLs) from the mice. These TCLs were *HLA-DR2*–restricted and produced T-helper type 1 (Th1) cytokine in response to the peptide in the presence of mouse or human antigen-presenting cells (APCs) expressing *HLA-DR2* (*DRB1*1502*) molecules. It was remarkable that transfer of 1 of the TCLs induced atypical autoimmune encephalitis involving the deep cere-

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-2^b 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'-GAACATGTGATCATCCAGGCCG-3' and 5'-GATCGGAGTATAGTTGGAGCGC-3'; *DRB1*1502* gene: 5'-CCTAAGAGGGAGTGTTCATTTCTTC-3' and 5'-TGTGAAGCTCTACCAACCCC-3'; *RAG-2* gene: 5'-CCACTCTTCGTTATCCAGC-3' and 5'-GTCCTCAAAGAGAACCCC-3'; pMC1 neo/poly(A)⁺ fragment disrupting endogenous *RAG-2* gene in *RAG-2^{-/-}* mice: 5'-TCATCTCACCTTGCTCCTGC-3' and 5'-TATGTCCTGATAGCGTCCG-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-TICGKGLSATV) and MBP143-168 (GVDAQGLSKIFKLGGRDSRSGSPMA) 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 × 10⁶ cells/mL in complete medium (RPMI-1640 supplemented with 10% FCS, 10 mM HEPES buffer, 5 × 10⁻⁵ 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 splenocytes as APCs.

Proliferation and inhibition assays. Splenocytes (4 × 10⁵/well) from *HLA-DR2*-B6 or *DRA*-B6 mice, or PBMCs (2 × 10⁵/well) from healthy human subjects

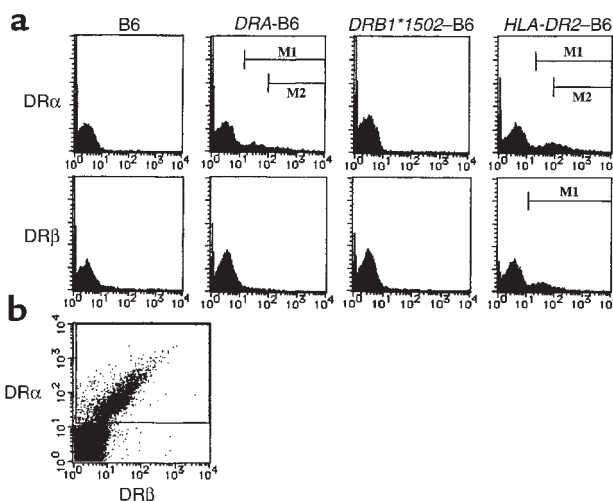


Figure 1

Expression of DRα/I-Eβ^b and HLA-DR2 molecules in *HLA-DR2*-B6 mice. (a) Splenocytes from B6, *DRA*-B6, *DRB1*1502*-B6, and *HLA-DR2*-B6 mice were stained with PE-labeled anti-*HLA-DRα* (L243) or FITC-labeled anti-*HLA-DRβ* (TÜ36) 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* (TÜ36) mAbs.

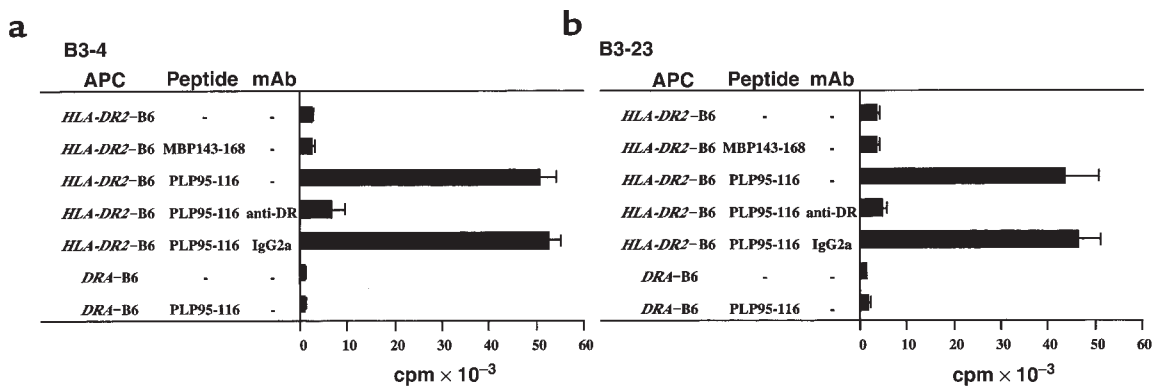


Figure 2

Proliferative response of T cells B3-4 and B3-23 generated from *HLA-DR2-B6* mice immunized with PLP95-116. Peptide-specific proliferation of T cells B3-4 (a) and B3-23 (b) were evaluated as described in Methods. By using splenocytes from *HLA-DR2-B6* (*HLA-DR2*⁺, *DRα*/*I-Eβ*⁺, *I-A*⁺) or *DRα-B6* (*DRα*/*I-Eβ*⁺, *I-A*⁺) 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-DRI*/8) were irradiated (35 Gy for mouse splenocytes and 40 Gy for human PBMCs) and used as APCs. T-line cells (4 × 10⁴/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 Beta-plate counter (Pharmacia, Uppsala, Sweden). Mean cpm of the triplicate cultures was calculated, and T cells 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).

Adoptive 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 × 10⁶) 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 FcRγII/III mAb) (PharMingen) for 5 minutes, and then incubated for 30 minutes with mAbs in combination as follows. Spleno-

cytes were incubated with either FITC-conjugated anti-*HLA-DRβ* (TÜ36) and PE-conjugated anti-*HLA-DRα* (L243), or FITC-conjugated anti-*I-Aβ*^b (25-9-17) and PE-conjugated anti-*I-Aα*^b (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 (10⁶/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 × 10⁶/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 for 30 minutes, then fixed and permeabilized with Cytofix/Cytoperm[®] solution for 20 minutes. The permeabilized cells were stained with FITC-labeled anti-mouse IFN-γ (XMG1.2) or isotype-matched rat IgG1 (R3-34) for 30 minutes. Samples were analyzed by flow cytometry as described above.

Histology. Four to 6 days after cell transfer, the moribund *HLA-DR2/RAG-2*^{-/-} mice were anesthetized and

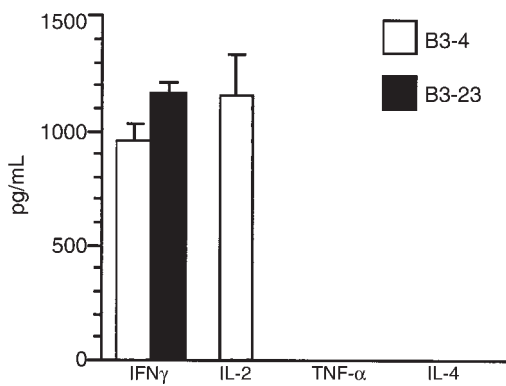


Figure 3

Cytokine production profile of T cells B3-4 and B3-23. T cells B3-4 and B3-23 were cultured for 48 hours with or without PLP95-116 in the presence of syngeneic splenocytes, and the culture supernatants were collected. Shown are the levels of IFN- γ , IL-2, TNF- α , and IL-4 detected in the supernatants of PLP95-116-stimulated cultures (no cytokine could be detected in the supernatants of the cultures without PLP95-116). Data represent mean \pm SD of the values obtained by duplicate assays in 3 independent experiments.

perfused through the heart with 0.1 M phosphate buffer followed by 4% paraformaldehyde. 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 μ m. 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 DR α and DR β chains, respectively, on immunocompetent cells. It is of note that DR α or DR β 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 xenogenic heterodimer complexed with an endogenous I-E β or I-E α chain. We first analyzed expression of DR α and DR β chains on splenocytes from DRA-B6, DRB1*1502-B6, and HLA-DR2-B6 mice. The cells were stained with PE-labeled anti-HLA-DR α (L243) or FITC-labeled anti-HLA-DR β 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 \pm 1.0%; n = 3) were stained with the anti-HLA-DR α mAb, indicating that the DR α chain appears on the cell surface in the form of a DR α /I-E β heterodimer. In contrast, DR β ⁺ cells were not detected in DRB1*1502-B6 mice. This is explained by the fact that mice with a B6 background lack endogenous I-E α , which is necessary for formation of a xenogenic heterodimer. On the other hand, the splenocytes from HLA-DR2-B6 mice were stained

not only with the anti-HLA-DR α mAb, but also with the anti-HLA-DR β mAb, implying that the DR β chain is probably expressed in the form of a DR α /DR β heterodimer (HLA-DR2 molecule). It is also noteworthy that the proportion of the cells that stained brightly with the anti-HLA-DR α mAb (M2 in Figure 1a) was significantly higher in samples from HLA-DR2-B6 mice than from DRA-B6 mice (15.1 \pm 2.8%, n = 6 vs. 3.9 \pm 0.3%, n = 3, respectively). In contrast, the expression level of endogenous I-A^b molecules in the transgenic mice was not altered by DRA or DRB1*1502 transgene (data not shown). The augmentation of DR α chain expression together with the expression of DR β 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 DR α and DR β chains in 32.0 \pm 5.2% (n = 5) of the splenocytes. These data imply that HLA-DR2-B6 mice express 3 different MHC class II molecules: HLA-DR2, DR α /I-E β , and I-A^b.

Characterization of PLP95-116-specific T cells 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 T cells recognizing PLP95-116 in the context of HLA-DR2 molecules. More than 100 T cells 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 T cells that were reactive to PLP95-116, and restimulated the lines with the peptide for further expansion. We selected 5 PLP95-116-specific T cells 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 T cells, designated B3-4 and B3-23, were studied in more detail.

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

Table 1

Induction of encephalitis by T cell B3-23 in HLA-DR2/RAG-2^{-/-} mice

T cell	Strain	Incidence	Onset day	Mortality
B3-23	HLA-DR2/RAG-2 ^{-/-}	4/7	4.0 (3-5)	4/7 ^A
B3-23	RAG-2 ^{-/-}	0/6	-	0/6
B3-4	HLA-DR2/RAG-2 ^{-/-}	0/8	-	0/8

Encephalitogenicity of T cells 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. ^ATwo 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 *DR α /I-E β ^b* and *I-A^b* 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 *DRA-B6* splenocytes expressing *DR α /I-E β ^b* and *I-A^b* 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, $\alpha 4$ 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 *V β 2* and *V β 14* TCRs predominantly, whereas TCL B3-23 was enriched in *V β 6*- and *V β 14*-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-DRI/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, 5×10^6 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

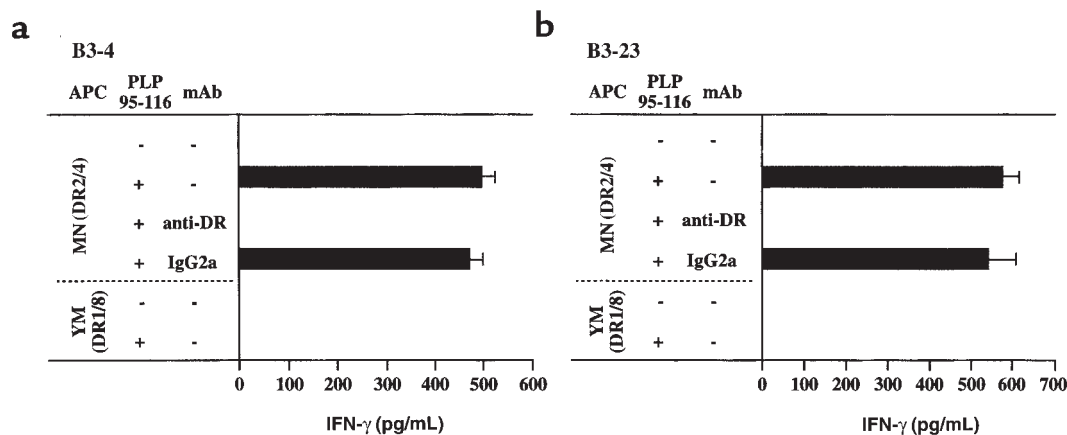


Figure 4

IFN- γ production by TCLs B3-4 and B3-23 in the presence of human PBMCs. TCLs B3-4 (**a**) and B3-23 (**b**) were cultured for 48 hours with or without PLP95-116 in the presence of human PBMCs from subject MN (*HLA-DR2 [DRB1*1502]/4*) or YM (*[HLA-DRI]/8*), and the supernatants were collected for measurement of *IFN- γ* . Blocking effects of anti-*HLA-DR* mAb (G46-6) or isotype-matched control IgG2a (G155-178) were also examined as indicated. Data represent mean \pm SD of the values obtained by duplicate assays in 4 independent experiments.

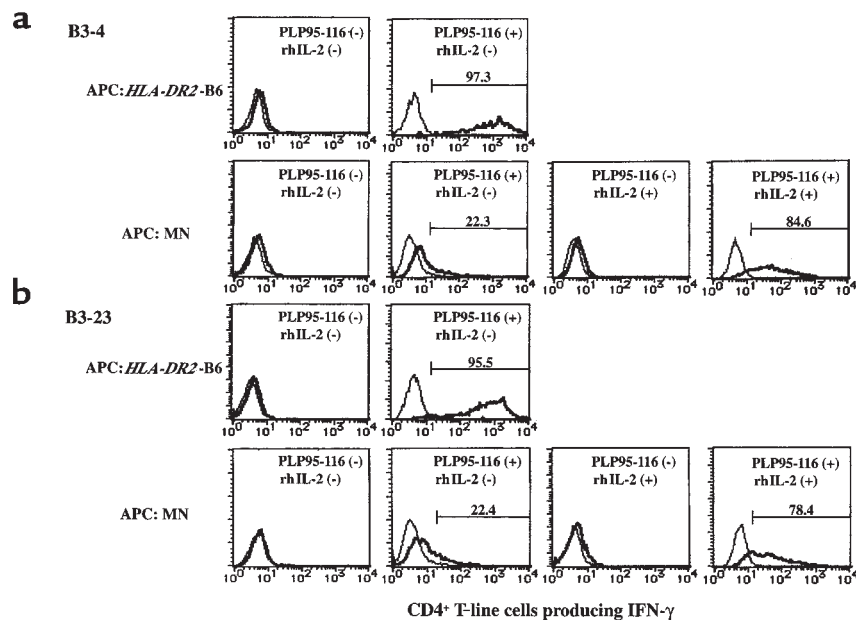


Figure 5

Flow cytometric detection of intracellular IFN- γ after stimulation with PLP95-116. Intracellular IFN- γ synthesis of T cells responding to PLP95-116 was examined using different sources of APCs: syngeneic splenocytes from *HLA-DR2-B6* mice, or human PBMCs (from subject MN expressing *HLA-DR2* [DRB1*1502]/4). The line cells B3-4 (a) and B3-23 (b) were incubated for 48 hours in the presence of the APCs with (+) or without (-) PLP95-116 (25 μ g/mL) and rhIL-2 (25 μ g/mL). GolgiPlug[®] (brefeldin A) was added for the last 10 hours of incubation, and the cells were stained for intracellular IFN- γ as described in Methods. Only the CD4⁺ T-cell population was gated for analysis, and the frequency (%) of the IFN- γ ⁺ population is shown (above bars). Note that exogenous rhIL-2 markedly enhanced IFN- γ production in the presence of the human APCs.

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 cerebrum 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.

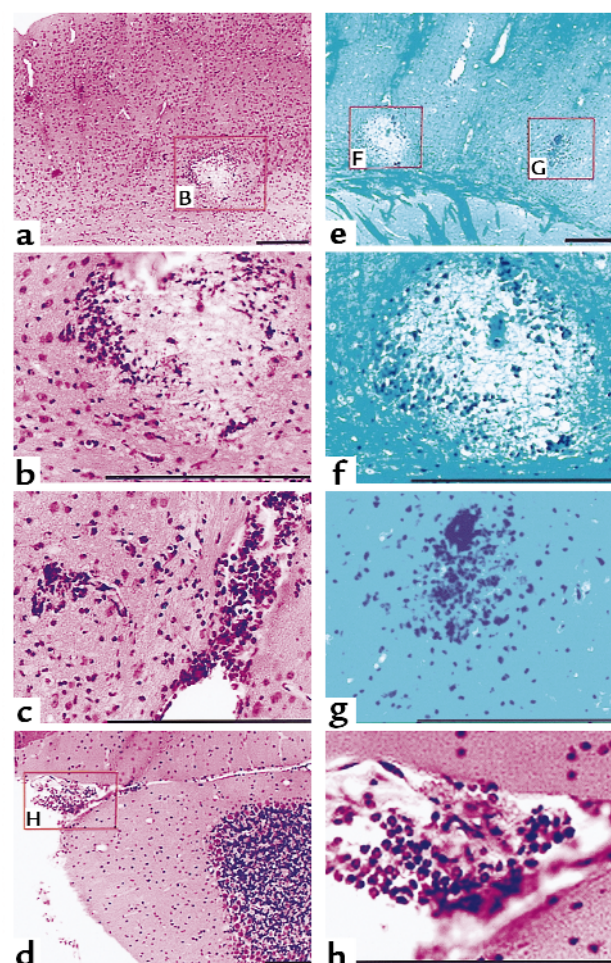


Figure 6

Histology of CNS sections from moribund *HLA-DR2/RAG-2*^{-/-} mice receiving TCL B3-23. Shown are representative inflammatory foci in the cerebrum and cerebellum from mice receiving TCL B3-23. (a-d and h) H&E staining. (e, f, and g) Luxol fast blue staining. Bars = 100 μm. Higher magnifications of the boxed areas B, H, F, and G are illustrated in b, h, f, and g, respectively. The inflammatory lesions include perivascular infiltrates in the cerebral cortex (g) and periventricular white matter (c) as well as inflammatory infiltrates within the ventricle (c) and cerebellar meninges (d). These infiltrates were primarily composed of mononuclear cells, but neutrophils were also identified (h). Some of the inflammatory foci located in the deep cerebral cortex (a and e) were accompanied by demyelinating lesions (b and f).

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