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Thiamine Deficiency Promotes T Cell Infiltration in Experimental Autoimmune Encephalomyelitis: the Involvement of CCL2

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Abstract

Multiple sclerosis (MS) is a complex multifactorial disease that results from the interplay between environmental factors and a susceptible genetic background. Experimental autoimmune encephalomyelitis (EAE) has been widely used to investigate the mechanisms underlying MS pathogenesis. Chemokines, such as C-C Chemokine Ligand 2 (CCL2), are involved in the development of EAE. We have previously shown that thiamine deficiency (TD) induced CCL2 in neurons. We hypothesized that TD may affect the pathogenesis of EAE. In this study, EAE was induced in C57BL/6J mice by the injection of myelin oligodendroglial glycoprotein (MOG) peptide 35–55 with/without TD. TD aggravated the development of EAE which was indicated by clinical scores and pathological alterations in the spinal cord. TD also accelerated the development of EAE in an adoptive transfer EAE model. TD caused microglial activation and a drastic increase (up 140%) in leukocyte infiltration in the spinal cord of the EAE mice; specifically TD increased Th-1 and Th-17 cells. TD upregulated the expression of CCL2 and its receptor CCR2 in the spinal cord of EAE mice. Cells in peripheral lymph node and spleen isolated from MOG-primed TD mice showed much stronger proliferative responses to MOG. CCL2 stimulated the proliferation and migration of T lymphocytes *in vitro*. Our results suggested that TD exacerbated the development of EAE through activating CCL2 and inducing pathological inflammation.

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Conflicts of interest

The authors have no conflicts of interest to declare.

Keywords

TD; EAE; MOG; infiltration of T lymphocyte; inflammation

Introduction

Human multiple sclerosis (MS) is an autoimmune disease, in which T lymphocytes specific for myelin antigens initiate an inflammatory reaction in the central nervous system (CNS), leading to demyelination and subsequent axonal injury (1–3). The etiology of MS is unclear. Recent genome-wide association studies (GWAS) indicate that genetic factors play a critical role (4). Epidemiological studies demonstrate that environmental influences also contribute to MS pathogenesis; these influences include infection, immunisations, physical and emotional stresses, climate, diet, and occupational exposures (5–6). Mapping the distribution of MS reveals a high prevalence of the disease in high-latitude areas, suggesting a positive relationship between vitamin D deficiency and MS (7–8). Vitamin D supplements have been considered a potential therapeutic strategy for MS patients. Vitamin D3 supplementation has been found to prevent or reduce MS (9–10). A recent study showed that serum levels of vitamin B12 and foliate were lower in MS patients (11). Other studies, however, did not support an association between B12 deficiency and MS (12). Interestingly, patients receiving vitamin B12 supplementation showed remarkable improvement in clinical outcomes (13). It is necessary to conduct a prospective study to determine whether treatment with supplements at an early stage of the disease can alter biomarker levels and change the course of the disease (14).

Experimental autoimmune encephalomyelitis (EAE) is a classic model for autoimmune demyelination of the CNS and has been widely used to explore pathogenic mechanisms of MS (15). The generation of myelin protein-reactive T cells is an immunological hallmark of both EAE and MS. The disease is initiated when activated autoreactive T cells cross the blood brain barrier (BBB) to reach the CNS. Upon encountering the antigens, these cells recruit inflammatory effector cells to the CNS, resulting in devastating demyelination and axon destruction (16–17). EAE can be induced either by immunizing animals with myelin components (actively induced EAE) or by transferring encephalitogenic T cells (adoptive transfer EAE) (1, 18–19). Animals are differentially susceptible to EAE (20). C57BL/6 mice develop a monophasic disease upon myelin oligodendroglial glycoprotein (MOG) challenge, showing extensive demyelination and inflammation in the CNS and are a commonly used EAE model (21).

Chemokines or chemoattractant cytokines are crucial during inflammatory responses for the recruitment of leukocyte populations to sites of tissue damage (22). Virtually all cell types, including many tumor cells, express chemokines and chemokine receptors (23–24). Chemokines play a major role in the immune response because they regulate the migration and activation of leukocytes. C-C Chemokine Ligand 2 (CCL2) was first identified as a potent chemotaxin for monocytes in response to proinflammatory stimuli (25). CCL2 regulates the migration and activation of monocytes, T cells, NK cells and basophils (26–28). CCL2 binds solely to C-C chemokine receptor (CCR) 2, a seven-transmembrane-

spanning protein that is functionally linked to downstream signaling pathways through heterotrimeric G proteins. In addition to being a receptor to CCL2, CCR2 serves as the receptor for four other ligands, MCP-2, -3, -4 and -5 (29). CCL2 and CCR2 have been implied in EAE (30–36). For example, anti-CCL2 antibody blocks relapses of adoptive transfer EAE in SJL mice (37). Mice that lack CCR2 fail to develop EAE after active immunization and are resistant to induction of EAE by the adoptive transfer of primed T cells from syngeneic wild-type mice (29, 38).

Thiamine is a cofactor of key enzymes in glucose metabolism (39). Thiamine deficiency (TD) causes mild impairment of oxidative metabolism and induces Wernicke–Korsakoff’s syndrome (WKS) in humans. TD causes regionally selective neuronal death, mitochondrial dysfunction, energy shortage and chronic oxidative stress in the brains of humans and animals (40–41). TD in animals has been used to model some neurodegenerative diseases (41). Subclinical TD is common in the elder population (42). However, it is unclear whether the status of thiamine will affect MS and EAE. Our previous studies demonstrated that TD increased CCL2 expression in the CNS (41). Here we demonstrated that TD enhanced the EAE severity by activating T cells reaction and increasing CCL2 expression in the spinal cord.

Materials and Methods

Peptides and antibodies

Rat MOG35–55 peptides were obtained from Biosynth International (Naperville, IL, USA) and purified by high performance liquid chromatography, and the purity of the peptide was >95%. The sequence of MOG35–55 was MEVGWYRSPFSRVVHLYRNGK. Purified Hamster anti-mouse CD3e, Fluorescein Isothiocyanate (FITC) rat anti-mouse CD4, APC rat Anti-mouse CD8a, APC rat anti-mouse IL-4 and PE rat anti-mouse IL-17A antibodies were purchased from BD Pharmingen (Basel, Switzerland); Goat anti-mouse ionized calcium-binding adaptor molecule-1 (IBA1) antibody was obtained from abcam (Cambridge, UK); Rat anti-mouse CD45, Rabbit anti-mouse CCL2 antibody and CCL2 (rat recombinant) was purchased from AbD Serotec (Raleigh, NC, USA). FITC rat anti-mouse IFN- γ antibody was purchased from eBioscienc (San Diego, CA, USA); rabbit anti-mouse CCR2 antibody was purchased from Abcam (Cambridge, UK). 2-(1-benzyl-indazol-3-yl) methoxy)-2-methyl propionic acid (bindarit) was synthesized by and obtained from Angelini (Angelini Research Center-ACRAF, Italy).

Animal models

Active immunization model of EAE—C57BL/6J mice were obtained from Shanghai SLAC Laboratory Animal Co. Ltd (Shanghai, China). The procedure for animal surgery was performed in accordance with the Guidelines of Animal Care and Use Committee of the Institute for Nutritional Sciences, Shanghai Institutes for Biological Sciences, Chinese Academy of Sciences. Induction of EAE was performed as previously described (43). Briefly, mice were subcutaneously injected at two sites with 100 μ g of rat MOG peptide 35–55 (MEVGWYRSPFSRVVHLYRNGK; >95% purity; Bio-Synthesis, Naperville, IL, USA), emulsified in complete Freund’s adjuvant containing 400 μ g of Mycobacterium tuberculosis

(Difco Laboratories, Detroit, MI, USA). On the same day (day 0) and on day 2 p.i., mice were intravenously injected with 200 ng of pertussis toxin (Sigma-Aldrich, St Louis, MO, USA). All mice were weighed, examined and graded daily for neurological signs in a blinded manner as follows: 0, no disease; 1, decreased tail tone or slightly clumsy gait; 2, tail atony and moderately clumsy gait and/or poor righting ability; 3, limb weakness; 4, limb paralysis; and 5, moribund state. Average disease scores were assessed daily. Additionally, in the EAE model, we documented the weight changes during the disease course. Only mice with a score of at least 2 for more than two consecutive days were judged to have fully developed EAE. The maximum clinical score achieved by each animal during the 30-day-observation period was used to calculate average maximum clinical scores for each experimental group. To study the time course of disease development, average clinical scores were calculated daily for each group of mice and plotted. MOG35–55 induced monophasic EAE was monitored for 30 days. Animals were euthanized if scores were worse than grade 4.

Adoptive transfer EAE model—To prepare encephalitogenic cells for adoptive transfer of EAE, mice were immunized with MOG35–55 as described above. Spleens and lymph nodes were collected and cell suspensions were prepared after 10 days of induction. Cells (6×10^6 /ml) were cultured in RPMI 1640 medium (supplemented with 10% fetal bovine serum, 2mM L-glutamine, 1 mM sodium pyruvate, 100 Iu ml^{-1} penicillin/streptomycin and 2×10^{-5} M 2-ME (Life Technologies, Carlsbad, CA, USA)), with MOG35–55 (20 μ g/ml) and IL-12 (30 ng/ml) (R&D Systems, Minneapolis, MN, USA). After 3 days in culture, the cells were harvested, washed in phosphate-buffered saline (PBS) and injected into recipient mice intravenously that were irradiated sublethally (500 rad) within 16 hours before cell injection. All mice were weighed, examined and scored daily after cell transfer (44).

TD induction—The animal TD model has been previously described. (40, 45–46). Briefly, adult male C57BL/6J mice were housed in a controlled environment (23°C and 53% humidity). The animals were fed with either a control diet or a thiamine-deficient diet (ICN Nutrition Biomedicals, Cleveland, OH, USA) *ad libitum* at the 0 day of EAE.

Treatment of bindarit—2-((1-benzyl-indazol-3-yl) methoxy)-2-methyl propionic acid (bindarit) is a small synthetic indazolic derivative that preferentially inhibits transcription of CCL2 (47). Bindarit has been shown some clinical efficacy in treating a broad array of experimental inflammatory, autoimmune and vascular disorders; it also had some success in recent clinical trials for diabetic nephropathy and lupus nephritis (48).

The method for bindarit treatment in animals has been previously described (48). Briefly, bindarit was prepared as a suspension in dimethyl sulfoxide (DMSO) at a concentration of 40 mg/ml. Then mice were given daily i.p. injection of bindarit (or vehicle DMSO) at 200 mg/kg for three consecutive days, beginning one day before MOG immunization (day –1), then injections every other day. This schedule was designed to minimize trauma associated with daily injections at times of peak neurologic disease and physical compromise.

Immunohistochemistry and immunofluorescence staining

For immunohistochemical (IHC) analysis of spinal cord tissues, mice were euthanized at the peak of EAE by intracardiac perfusion with ice-cold PBS, followed by 4% paraformaldehyde solution, under anesthesia. Spinal cords were rapidly dissected and sectioned at a thickness of 25 μm . The sections were rinsed in PBS, incubated with 0.3% hydrogen peroxide, blocked by the incubation with 10% bovine serum albumin at 37°C for 1 hour, then incubated overnight at 4°C with a primary antibody (rat anti-mouse CD45 antibody, 1/1,000; Goat anti-mouse IBA1 antibody, 1/1,000). The sections were then incubated with appropriate biotinylated secondary antibodies at 37°C for 1 hour and treated with diaminobenzidine. All antibodies were diluted in 1% bovine serum albumin in PBS. Negative controls were performed by the incubation of preimmune IgG. For detecting inflammatory infiltrates, the sections were stained with hematoxylin and eosin (HE).

For immunocytofluorescence staining, tissue sections or cells from lymph nodes were rinsed in PBS, blocked by incubation with 1% bovine serum albumin at 37°C for 1 hour, then incubated overnight at 4°C with primary antibodies (rabbit anti-CCL2 polyclonal antibody, 1/200; rat anti-mouse CD4 antibody, 1/50; rat anti-mouse CD8a, 1/50). The sections were incubated with appropriate FITC secondary antibodies at 37°C for 1 hour. The bright field images were taken on a BX51 Olympus microscope (Olympus Corporation, Tokyo, Japan); Immunofluorescent images were recorded using a Zeiss LSM 510 Meta confocal microscope (Carl Zeiss MicroImaging Inc., Thornwood, NY, USA). For the quantification, five sections from each mouse were used for cell counting. Cells were counted using ImageJ (US National Institutes of Health) in a designated area. Data represent mean \pm SD of 5 mice for each group.

T cell proliferation

To examine the proliferation of T cells, we isolated lymph nodes and spleen from MOG35–55-immunized mice and cultured T cells in a 96-well plate (1×10^5 per well) in the presence of MOG35–55 (0, 0.8, 4, 20 and 100 $\mu\text{g}/\text{ml}$), CCL2 (20 $\mu\text{g}/\text{ml}$) or Con A (10 $\mu\text{g}/\text{ml}$) (Sigma-Aldrich). Cells were maintained in RPMI 1640 medium supplemented with 10% fetal bovine serum (Life Technologies), 2 mM L-glutamine, 1mM sodium pyruvate, 100 IU/ml penicillin/streptomycin and 2×10^{-5} M 2-ME (Life Technologies) for 72 hours. Cell proliferation was determined using an AMR PLUS kit (Lonza Rockland, Rockland, ME, USA) according to manufacturer's instruction. The absorbance was analyzed with a luminometer (Bio-Tek, Atlanta, GA, USA).

Flow cytometry

T cells ($1 \times 10^6/\text{ml}$) obtained from lymph nodes were washed and resuspended in PBS. Cells were stained for surface markers with specific primary antibodies and appropriate fluorescein isothiocyanate-conjugated (FITC) secondary antibodies in fluorescence-activated cell sorting (FACS) buffer at 4°C for 40 min. Cells were washed twice and resuspended in the 200–400 μl of PBS for flow cytometry analysis as previously described (49–50). The cell sorting was performed with a FACSCalibur (BD Biosciences, San Diego, CA, USA) equipped with CellQuest software (BD Biosciences). Data were analyzed with FlowJo software (Tree Star, San Carlos, CA, USA). For intracellular staining, cells were

maintained in a 6-well plate (2×10^6 per well) and treated with MOG35–55 (20 $\mu\text{g}/\text{ml}$) for 72 hours. An inhibitor of protein transport Brefeldin A (1: 10 dilution; BD Biosciences, San Jose, CA, USA) was added to the cultures during the last 4–5 hours, then cells were collected, fixed and permeabilized using Cytofix/Cytoperm (BD Biosciences, San Jose, CA, USA), followed by the staining using fluorescence labeled rat anti-mouse antibodies directed against IFN- γ , IL-4 or IL-17. Cells were then analyzed with FACSCalibur. Isotype-matched antibodies were used as controls. For the CCL2 binding experiments, draining lymph node cells were incubated with IgG, CCL2 or CCL2 plus anti-CCL2 antibody for 60 min. The cells were then incubated with appropriate FITC antibodies and analyzed with FACSCalibur.

Transwell migration assay

Chemotaxis assays were performed with purified lymph node T cells using uncoated 8- μm transwell filters (Corning Costar, Corning, NY) as previously described (51). Briefly, T cells ($2 \times 10^5/\text{ml}$) suspended in RPMI 1640 containing 1% FBS were added to the upper chamber; CCL2 (5 $\mu\text{g}/\text{ml}$) as a T cell chemokine was added to the lower chamber. Cells were allowed to migrate for 4 hours at 37°C in 5% CO₂ atmosphere. The transwell filters were collected and fixed with methanol for 10 min. then stained with Diff-Quick (Baxter Diagnostics, Inc., Deerfield, IL) for 20 min. The migrated T cells were counted under a microscope at $\times 400$ magnification (Axiovert 200M; Carl Zeiss). Data are expressed as mean \pm SD for duplicate filters from four independent experiments.

Statistical analysis

Statistical analysis was assessed by ANOVA followed by Student–Newman–Keuls analyses. An unpaired t test was used for the analysis of quantitative data of cell counting. Data were presented as means \pm SD. Difference in which $p < 0.05$ was considered statistically significant.

Results

TD exacerbates EAE in active immunization model

We investigated the effects of TD on EAE using an active immunization model of EAE. The EAE mice developed a monophasic disease characterized by ascending paralysis 10–18 days after immunization, and EAE mice with TD showed markedly more severe neurologic dysfunction. Table 1 summarizes the neurobehavioral features of MOG35–55-(MOG) induced EAE mice with/without TD. In our EAE mouse model (MOG injection), 11 out of 16 (69%) mice developed fatal EAE, whereas in EAE plus TD model, 13 out of 16 (81%) mice developed fatal EAE (Table 1). The onset of EAE was earlier in EAE plus TD model (13 days) compared to EAE mouse model (16 days). The neurological signs of EAE in mice were quantified and expressed as clinical scores and presented in Fig. 1A. Mice with TD alone revealed neurological alterations, such as clumsy gait or poor righting ability after 25 days. The clinical score for EAE model peaked at day 20 and then slightly declined; however, in EAE plus TD model, the clinical score was significantly higher and kept increasing after day 20 (Fig. 1A). Additionally, in EAE plus TD model, mice had a more severe loss of body weight (Fig. 1B).

We next performed histopathological analysis on spinal cords of EAE mice with or without TD. Inflammatory cell infiltration in lumbosacral enlargement was examined by hematoxylin and eosin staining (Fig. 1C) and the infiltrated cells were quantified (Fig. 1D). Compared to controls, the number of infiltrated cells increased by 283% and 613% in EAE mice and EAE plus TD mice, respectively (Fig 1D).

To determine the identity of inflammatory cells, we performed immunohistochemical analysis of CD45 and Iba1 on the spinal cord (Fig. 2). CD45 is a marker for the mononuclear cells from the circulation, while Iba1 is a marker for microglia in the CNS. In the EAE plus TD model, there were much more CD45+ cells (139% increase) compared to the EAE model. EAE activated microglia in the spinal cord; the active microglia displayed a larger cell body and thicker processes. TD significantly increased the number of active microglia in EAE mice (Figs 2C and D). We further examined the infiltrated lymphocytes in the spinal cord by IHC of CD4 and CD8. CD4+ and CD8+ cells were only observed in EAE mice and there was no CD4+ and CD8+ T lymphocytes in the spinal cord of wild-type mice and TD mice. TD drastically increased CD4+ and CD8+ cells in EAE mice (Fig. 3). We have also performed FACS to test the proportion of CD4 and CD8 cells obtained from spleen and PLN in different group mice on the onset and peak of EAE. The results showed there were not obvious difference between different group (Supplementary Fig. 1).

TD potentiates antigen-stimulated proliferation of T lymphocytes

We hypothesized that TD increased the proliferation of T lymphocytes in response to the antigen. To test this hypothesis, we examined the antigen-stimulated proliferation of T lymphocytes isolated from the peripheral lymph node and spleen of control, TD mice, and EAE with/without TD mice on the onset and peak of EAE. Although other cells may be present, T lymphocytes are major cells in peripheral lymph node. MOG did not stimulate the proliferation of lymphocytes isolated from control and TD mice, but significantly increased the proliferation of lymphocytes isolated from EAE mice (Fig. 4). More importantly, the lymphocytes isolated from EAE mice plus TD mice displayed a much stronger and concentration-dependent response to MOG-mediated proliferation.

TD aggravates EAE in adoptive transfer model

We hypothesized that the antigen-specific T cells isolated from EAE plus TD would cause more severe EAE. To test this hypothesis, we transferred encephalitogenic cells from mice treated with MOG with/without TD for 10 days to recipient mice that were irradiated sublethally. The encephalitogenic cells isolated from MOG-treated mice caused EAE which was shown by the clinical score and the loss of body weight; while the encephalitogenic cells isolated from MOG plus TD-treated mice induced more severe EAE in recipient mice which was indicated by higher clinical scores and a greater loss of body weight (Fig 5A). Next, we transferred encephalitogenic cells isolated from MOG-treated mice to recipient mice that were treated with/without TD. In the recipient mice that were treated with TD, the encephalitogenic cells caused a quicker and more severe development of EAE as determined by the clinical scores (Fig. 5B). Consistently, the encephalitogenic cells induced greater body weight loss in the recipient mice that were treated with TD (Fig 5B).

CD4⁺ T cells have been classified into distinct subsets namely Th1 (characterized by the production of interferon- γ), Th2 (characterized by the production of interleukin 4) and Th17 (characterized by the production of interleukin 17). IFN- γ -producing Th1 cells and IL-17-producing Th17 cells have been implicated in EAE induction. We examined the relative populations of Th1, Th2 and Th17 cells in the lymph node using flow cytometry after 10 days of induction. There was a significant increase in the interferon- γ producing cells (23% more) and interleukin producing cells (56% more) in EAE plus TD model compared to EAE only model (Fig. 5C). However, there was no difference in IL4 producing cells among all experimental groups (Fig. 5C).

CCL2 is involved in TD-induced exacerbation of EAE

We have demonstrated that the expression of C-C Chemokine Ligand 2 (CCL2) was selectively induced by TD in the brain (52). We hypothesized that CCL2 may mediate TD-induced exacerbation of EAE. To test this hypothesis, we examined the expression of CCL2 in the spinal cord with IHC. Both MOG and TD treatment increased the number of cells expressing CCL2, but MOG plus TD further increased CCL2⁺ cells in comparison to MOG or TD-treated group alone (Figs 6A and B). To confirm that CCL2 can regulate T lymphocytes infiltration, we first determined whether CCL2 can directly bind to T cells. We treated lymph node T cells with IgG, CCL2 or CCL2 plus anti-CCL2 antibody for 60 min, and then determined the amount of CCL2 bound to T cells by flow cytometry. Anti-CCL2 antibody blocked the binding of CCL2 to T cells, indicating that CCL2 specifically bound to T cells (Figs 6C and D). More importantly, CCL2 increased the proliferation and migration of T lymphocytes *in vitro* (Figs. 6E and F).

To confirm the involvement of CCL2, we treated mice with bindarit, an inhibitor of CCL2 synthesis. Bindarit inhibited the expression of CCL2 in the spinal cord of (Supplementary Fig. 2C and D). Bindarit had therapeutic effects on mice of EAE plus TD, lowering the clinical scores and alleviating the loss of body weight (Supplementary Fig. 2A and B). Mice suffered EAE plus TD showed a rapid progression of EAE, reaching a maximum mean clinical score of 2.5. Bindarit-treated mice with EAE plus TD had a slower progression and a maximum mean clinical score of 1.5.

TD increases the expression CCR2 in T lymphocytes

CCR2 is a chemokine receptor that responds predominantly to CCL2. We examined the expression of CCR2 in lymphocytes isolated from lymph nodes by flow cytometry. The expression of CCR2 on lymphocytes was confirmed by isolated T cells from LN (Supplementary Figure 3). Approximately 30% of lymphocytes expressed CCR2 and CCR2 was observed in both CD4⁺ and CD8⁺ cells (data not shown). We next examined the effect of MOG and TD on CCR2 expression in CD4⁺ and CD8⁺ T lymphocytes in the spinal cord. MOG significantly increased the number of CCR2-positive CD4⁺ or CD8⁺ T lymphocytes in the spinal cord and TD potentiated the effect of MOG (Figs 7 and 8).

Discussion

Multiple sclerosis (MS) is considered a T lymphocyte-mediated CNS autoimmune disease that results from the interplay between environmental factors and a susceptible genetic background (15, 53–54). Epidemiological studies show that environmental influences, such as diet, play an important role in the development of MS (5). Vitamin deficiency has been implicated in the pathogenesis of MS. For example, there is a correlation between vitamin D deficiency and MS (7–8). Vitamin D supplements have been considered a potential therapeutic strategy for multiple sclerosis patients (9–10). MS and vitamin B12 deficiency share some common inflammatory, neurodegenerative and pathophysiological characteristics and a decreased level of vitamin B12 has been demonstrated in MS patients (55–56). The supplementation with vitamin B12 improved clinical outcomes in MS patients (13). In an animal model of MS, both nicotinic acid and fumaric acid esters have been shown to improve neurological function, and this effect was accompanied by a significant reduction in inflammatory infiltrates (57). Fumaric esters have recently been evaluated in Phase III trials for MS (58–60) and nicotinic acid has been shown to have neuroprotective and anti-inflammatory effects during acute ischemic stroke (61).

Thiamine (vitamin B1) deficiency (TD) produces a mild impairment of oxidative metabolism and TD in animals has been used to model the diminished metabolism and reduced activity of the thiamine-dependent mitochondrial enzymes that occur in the brain of aging-related neurodegenerative disorders (40, 62). In this study, we show that TD exacerbates experimental autoimmune encephalomyelitis (EAE) which is a widely used model to study the pathogenesis of MS (63). Two models of EAE were used in our study; an active immunization model was achieved by subcutaneous injection of myelin antigen (MOG) and an adoptive transfer model which transferred purified MOG-specific T cells obtained from EAE mice into naive mice. In both active and adoptive transfer EAE models, TD drastically increases the progression of EAE which is indicated by a significant worse clinical score and a more severe loss of body weight (Figs 1 and 5). The mice of TD alone showed neurological alterations, such as clumsy gait or poor righting ability after 25 days.

The clinical severity of EAE is directly associated with T cell activation (64). We show that TD increases the number of infiltrated cells and active microglia in the spinal cord. We further demonstrate that TD increases CD4⁺ and CD8⁺ T lymphocytes in the spinal cord. These results indicated that TD exacerbates the development of EAE by inducing T activation. Neuro-antigen-specific CD4⁺ T cells can initiate and sustain neuroinflammation and pathology in EAE (15). Depending on the major cytokines produced, CD4⁺ T cells have been classified into distinct subsets namely Th1, Th2 and Th17. Recently, many studies have been undertaken to identify the T cell subsets that are involved in tissue-specific autoimmune diseases. Interferon- γ -expressing Th1 cells were initially considered to be the effector CD4⁺ T cell subset that induced EAE. Later, Th17 cells were also reported to cause more severe EAE; animal models that were deficient in IL-17A or the IL-17RA receptor were more resistant to EAE (65–67). We demonstrate that TD significantly increases the number of CD4⁺ cells and specifically the number of Th1 and Th17 cells but not Th2 cells (Figs 3 and 5). These results are consistent with the role of these T cells in EAE. TD also increases the number of CD8⁺T cells (Fig. 3). The role of CD8⁺T cells in EAE is unclear,

although some studies indicate a potential pathogenic action for CNS-targeted CD8+T cells (68–69).

A critical event in the pathogenesis of EAE is the entry of both antigen-specific T lymphocytes and antigen-nonspecific mononuclear cells into the CNS. Chemokines are a key mediator that regulate the transmigration of T cells and monocytes across the blood–brain barrier (BBB) (70). CCL2, one of the first chemokines to be characterized, regulates the activity of monocytes, dendritic cells, and NK cells and plays an important role in innate immunity (35); it is also associated with pathological inflammation (71). CCR2 is a C-C chemokine receptor that responds predominantly to CCL2. In murine EAE, the expression of CCL2 mRNA in the brain and spinal cord was upregulated and may mediate the onset of EAE (31–32). We have previously demonstrated that neuronal CCL2 plays an important role in TD-induced microglia recruitment/activation and neurodegeneration in the brain (52). We show here that TD increases the expression of CCL2 and CCR2 in spinal cord of EAE in mice (Figs 6–8). In humans, CCR2 is expressed in monocytes and CD4+ T cells in the circulation (72). Our results show that CCR2 is expressed in both CD4+ and CD8+ T cells in C57BL6 mice. Our data indicate that CCL2 can directly activate T cells as it binds to T cells and promotes the proliferation and migration of T cells *in vitro* (Fig. 6). Bindarit, CCL2 synthesis inhibitor, suppresses the expression of CCL2 in the spinal cord and has therapeutic effects on EAE plus TD mice (Supplementary Fig. 2). These indicate that the involvement of CCL2.

Taken together, our results suggest that TD upregulates CCL2 and CCR2, and through its interaction with CCR2, CCL2 enhances the proliferation and recruitment of encephalitogenic T cells into the spinal cord which causes more pathological inflammation. The mechanisms underlying TD-induced CCL2/CCR2 upregulation are currently unknown. TD causes oxidative stress. Oxidative stress is detrimental to neurons and may promote inflammatory activities and the production/secretion of CCL2. Although a direct regulation cannot be ruled out, it is more likely TD modulates CCL2 expression through an indirect mechanism, such as oxidative stress. Regardless of the mechanisms, CCL2/CCR2 signaling plays an important role in TD-potentiated EAE.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations

MS Multiple Sclerosis

EAE	Experimental Autoimmune Encephalomyelitis
CNS	Central Nervous System
TD	Thiamine Deficiency
MOG	Myelin Oligodendroglial Glycoprotein
CCL2	C-C Chemokine Ligand 2
DMSO	Dimethyl sulfoxide
CCR2	C-C Chemokine Receptor 2
WKS	Wernicke–Korsakoff’s syndrome

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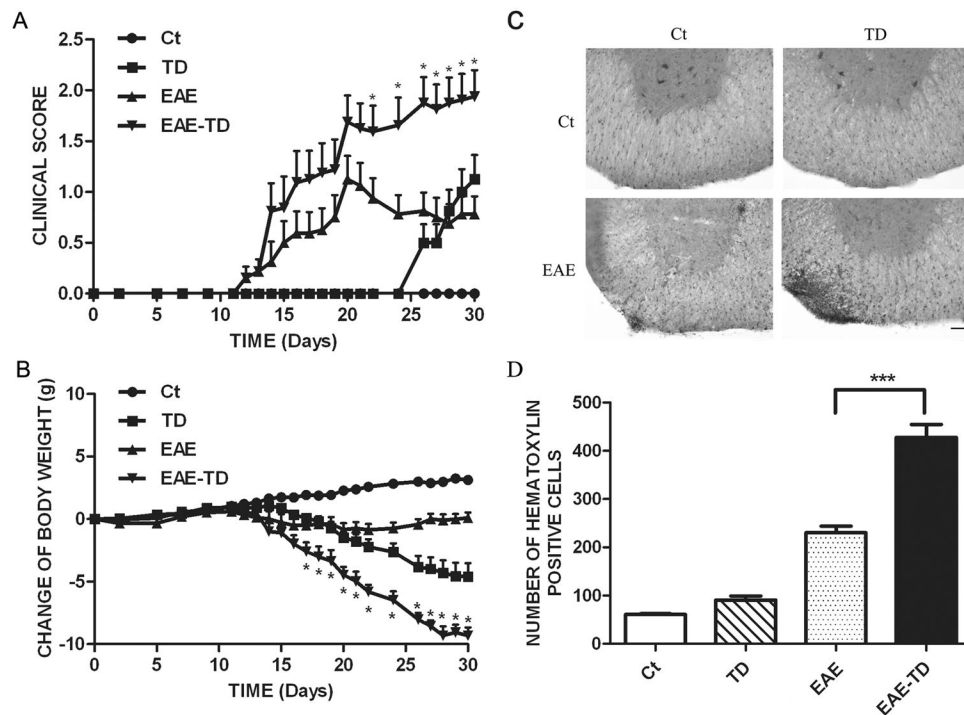


Figure 1. Effect of TD on the progression of EAE

EAE was induced in C57BL/6 mice by injection of MOG35-55 with or without TD as described under the Materials and Methods. The clinical score (A) and body weight (B) was determined in these mice as described under the Materials and Methods. Data were presented as mean \pm SEM; * $p < 0.05$, compared with EAE mice; $n = 20$ for each group. (C) The infiltration of inflammatory cells in the spinal cord was detected by hematoxylin and eosin staining on the peak of EAE. Bars = 50 μ m. (D) The number of hematoxylin positive cells in the spinal cord was quantified using ImageJ software in a designated area. Data were presented as mean \pm SD; *** $p < 0.001$, compared with EAE mice; $n = 5$ for each group.

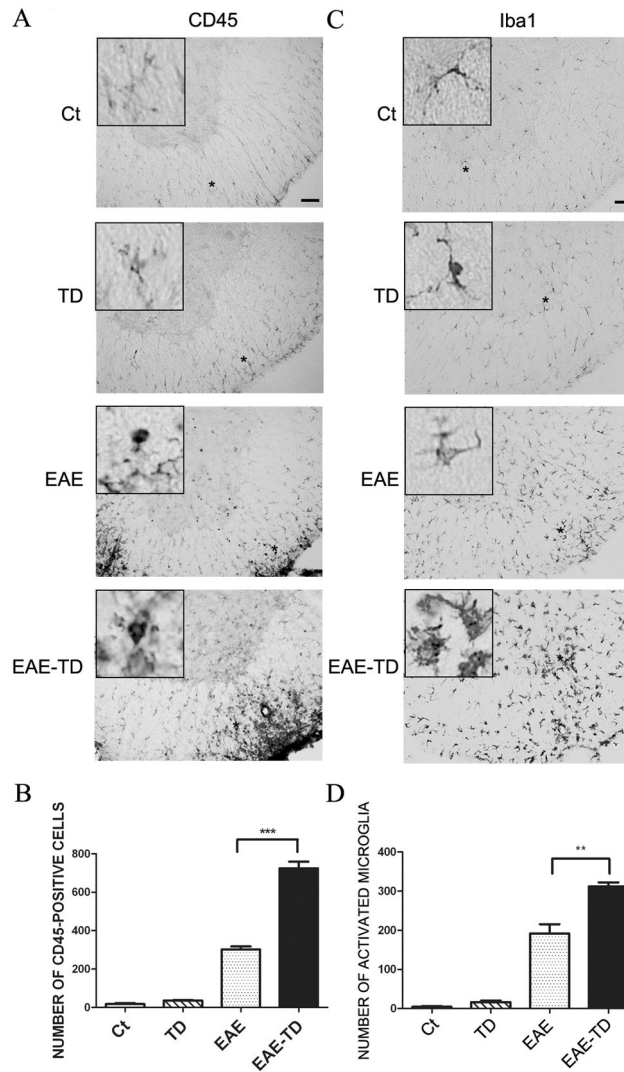


Figure 2. Effect of TD on microglial activation in EAE mice

EAE in C57BL/6 mice was induced by injection of MOG35-55 with or without TD. The activation of microglia was measured by immunohistochemical (IHC) staining with an antibody directed against CD45 (A) or Iba-1 (C) on day 20 after of MOG35-55 injection. Bars = 50 μ m. The number of CD45+ cells (B) and active microglia (D) was determined. Data were presented as mean \pm SD; ** p <0.01, *** p <0.001, compared with EAE mice; n = 5 for each group.

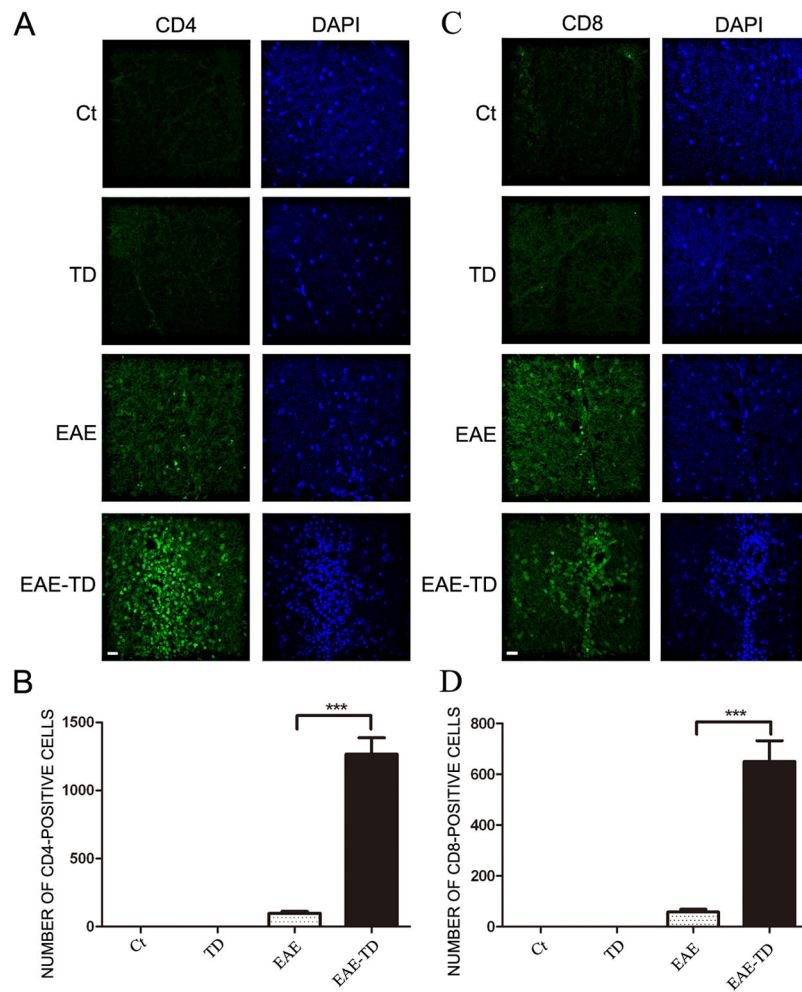


Figure 3. Effect of TD on the infiltration of CD4+ and CD8+ T lymphocytes in the spinal cord of EAE mice

EAE in C57BL/6 mice was induced by injection of MOG35-55 with or without TD. On day 20 after MOG35-55 injection, CD4+ and CD8+ T lymphocytes in the spinal cord were measured by IHC using specific antibodies (A and C). Bars = 20 μ m. The number of CD4+ and CD8+ T lymphocytes was quantified (B and D). Data were presented as mean \pm SD; n = 5 for each group. ***p<0.001, compared with EAE mice.

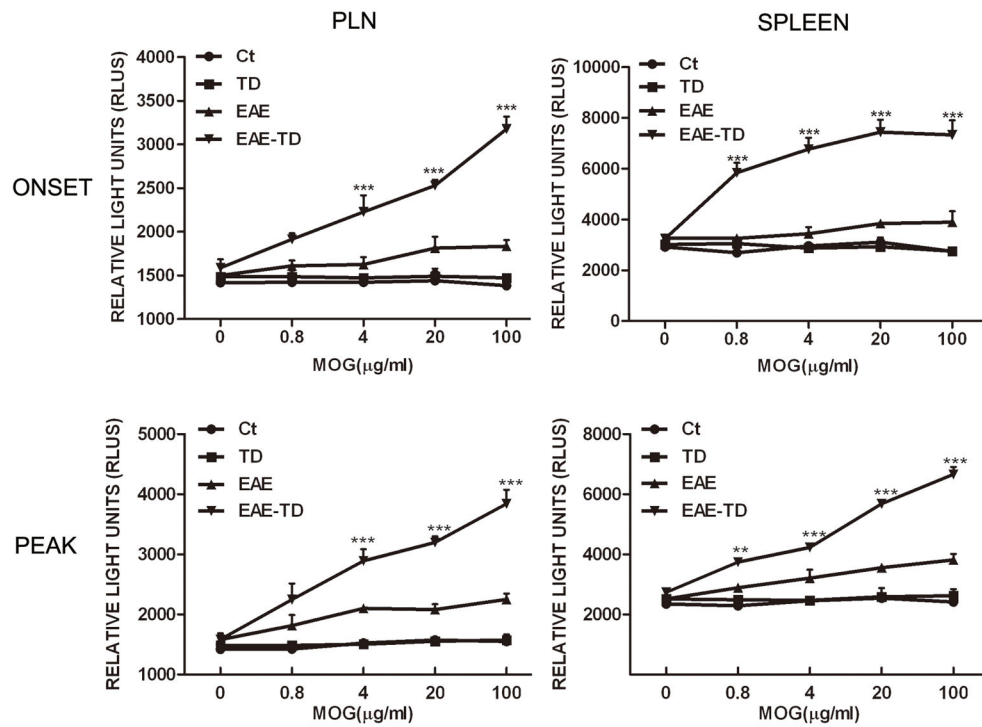


Figure 4. TD enhances the proliferation of T lymphocytes in response to MOG35–55
 EAE in C57BL/6 mice was induced by injection of MOG35–55 with or without TD. On the onset and peak of EAE, T lymphocytes were isolated from peripheral lymph node and spleen and challenged with MOG35–55 *in vitro*. Cell proliferation was determined as described under the Materials and Methods. Data were presented as mean ± SD; **p<0.01, ***p<0.001, compared with EAE group; n = 4 for each group.

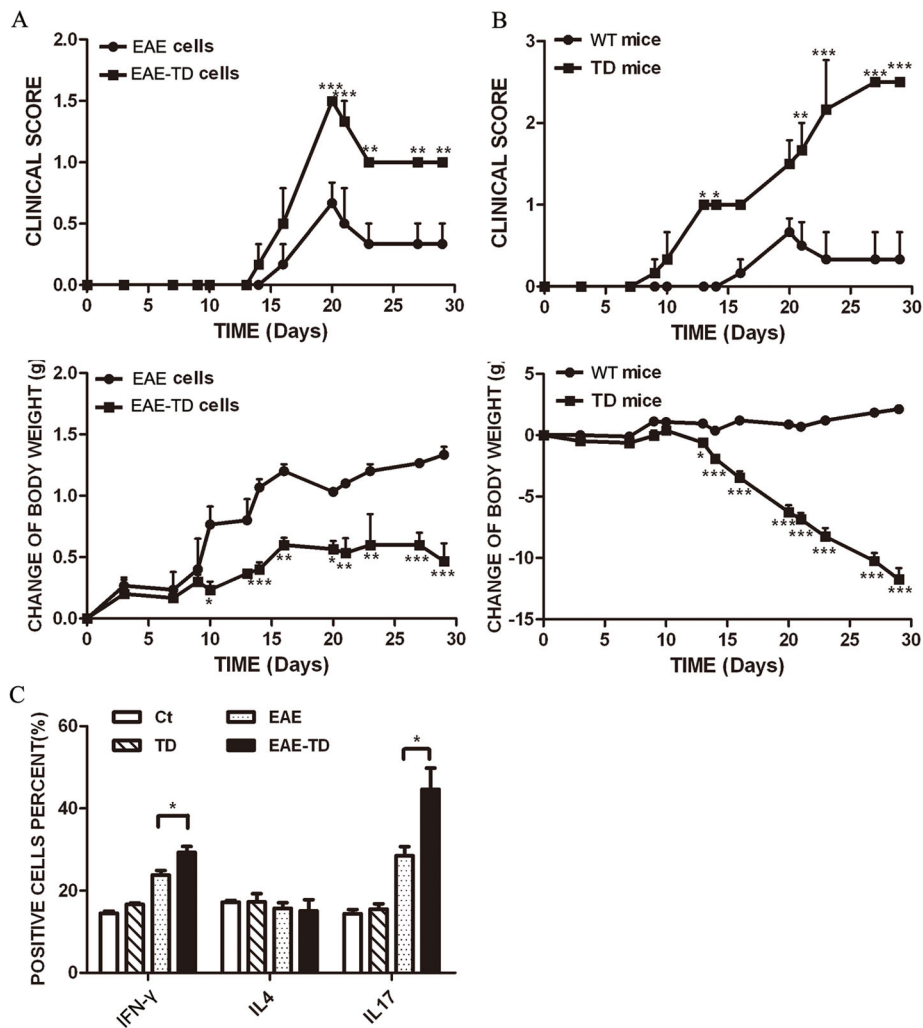


Figure 5. Effect of TD on adoptive EAE model

A: EAE was induced in C57BL/6 mice by injection of MOG35-55 with or without TD. Encephalitogenic cells were collected from these mice as described under the Materials and Methods and injected into recipient mice (wild-type). The recipient mice were irradiated sublethally (500 rad) 16 hours before the injection. The clinical scores for EAE and body weight were determined as described under the Materials and Methods. **B:** EAE was induced in C57BL/6 mice by injection of MOG35-55. The encephalitogenic cells were prepared from EAE mice and injected into recipient mice with/without TD. The clinical scores for EAE and body weight were determined as described under the Materials and Methods. Data were presented as mean ± SD; *p<0.05, **p<0.01, ***p<0.001, compared with control group. n = 8 for each group. **C:** Lymph node cells were isolated from EAE mice with or without TD after 10 days of induction. The percentage of IFN-γ, IL-4 and IL-17 producing cells were measured by FACS. Data were presented as mean ± SD; *p<0.05, compared with EAE group. n = 4 for each group.

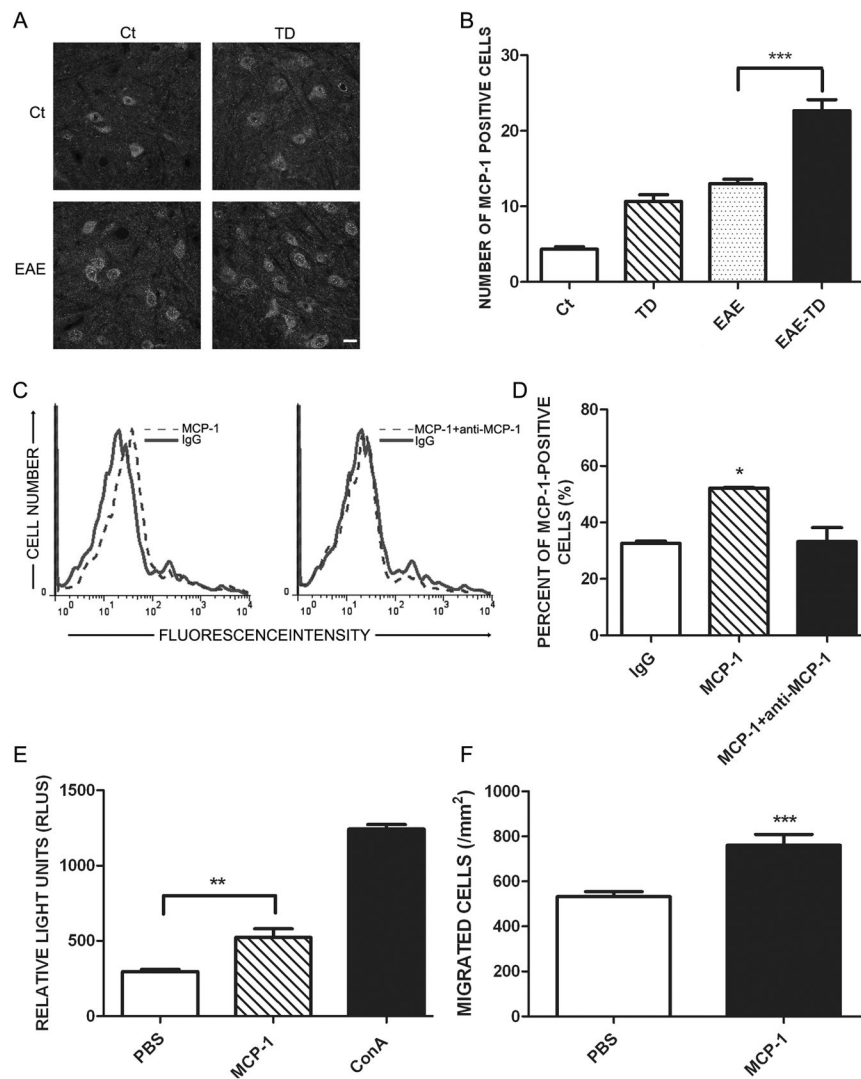


Figure 6. Effect of TD on the expression of CCL2 in spinal cord of EAE mice
A: EAE was induced in C57BL/6 mice by injection of MOG35-55 with or without TD. The expression of CCL2 in the spinal cord was determined by IHC on day 20 after MOG35-55 injection. Bar = 20 μ m. **B:** The number of CCL2-positive cells was quantified as described under the Materials and Methods. n = 5 for each group. **C:** Lymph node cells were isolated and incubated with CCL2 with/without anti-CCL2 antibody. The binding of CCL2 to T lymphocytes was evaluated by FACS. **D:** The binding of CCL2 to T lymphocytes was quantified by FlowJo software. **E:** Peripheral T lymphocytes were isolated from lymph nodes and treated with CCL2 (20 μ g/ml) for 72 hours. Cell proliferation was determined using ViaLight Plus Kit. ConA was used as positive control for the stimulation of T cell proliferation. **F:** T cells were treated with CCL2 (5 μ g/ml) for 4 hours. CCL2-induced T-cell migration was measured by a Transwell Migration assay. Data were presented as mean \pm SD; *p<0.05, **p<0.01, ***p<0.001, compared with control group. The experiment was replicated four times.

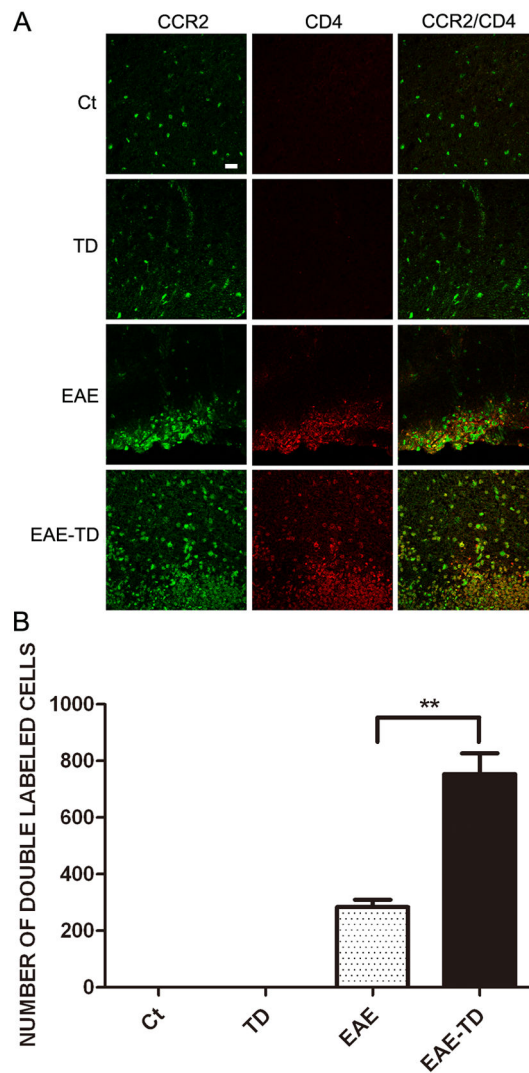


Figure 7. Effect of TD on CCR2-positive CD4 cells in spinal cord of EAE mice

A: EAE was induced in C57BL/6 mice by injection of MOG35-55 with or without TD. The expression of CCR2 on CD4⁺ T lymphocytes on the lumbar spinal cord of EAE mice was examined with double immunofluorescent staining using antibodies directed against CCR2 or CD4. Bars = 20 μ m. **B:** CCR2⁺ and CD4⁺ cells were quantified as described under the Materials and Methods. Data were presented as mean \pm SD; **p<0.01, compared with EAE group. n = 5 for each group.

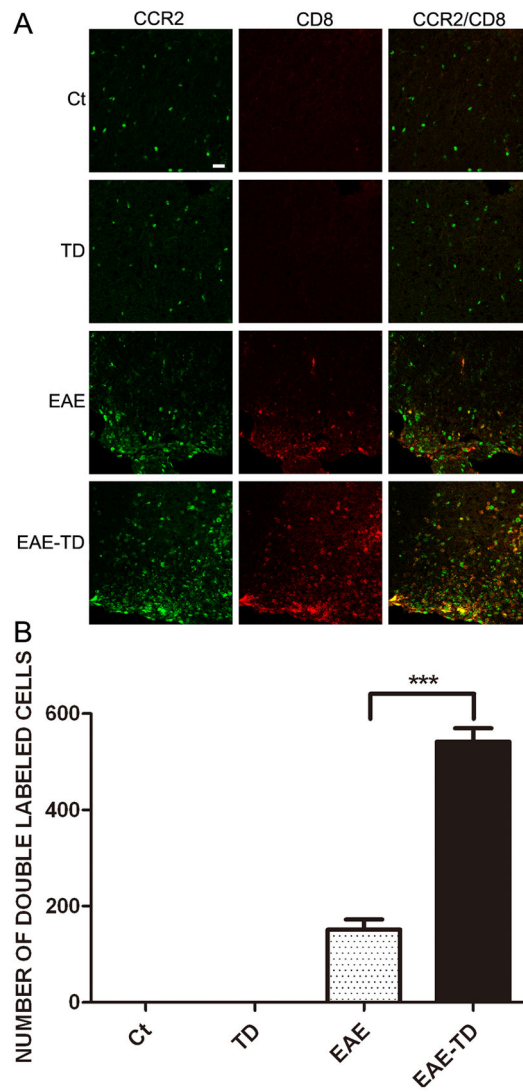


Figure 8. Effect of TD on CCR2-positive CD8 cells in spinal cord of EAE mice
 The notations are as Figure 8. Data were presented as mean ± SD; ***p<0.001, compared with EAE group. n = 5 for each group.

Table 1

EAE in wild-type and TD mice

Mouse genotype	Incidence	Days of EAE onset (mean \pm S.D.)	Maximum clinical score (mean \pm S.D.)
Wild type	11 of 16 (69%)	16.09 (\pm 2.91)	1.55 (\pm 0.55)
TD	13 of 16 (81%)	13.61 (\pm 3.50)	2.38 (\pm 0.46)*

Results are cumulative data from three different experiments,

* P<0.05, as compared to corresponding wild-type controls.